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Matt Whiteman *Editors*

Chemistry, Biochemistry and Pharmacology of Hydrogen Sulfide

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Editors

Chemistry, Biochemistry and Pharmacology of Hydrogen Sulfide

 Springer

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Part I

**Biochemistry, Molecular Biology and
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H₂S Synthesizing Enzymes: Biochemistry and Molecular Aspects

Caleb Weihao Huang and Philip Keith Moore

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Abstract

Hydrogen sulfide (H₂S) is a biologically active gas that is synthesized naturally by three enzymes, cystathionine γ -lyase (CSE), cystathionine β -synthetase (CBS) and 3-mercaptopyruvate sulfurtransferase (3-MST). These enzymes are constitutively present in a wide array of biological cells and tissues and their expression can be induced by a number of disease states. It is becoming

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increasingly clear that H₂S is an important mediator of a wide range of cell functions in health and in disease. This review therefore provides an overview of the biochemical and molecular regulation of H₂S synthesizing enzymes both in physiological conditions and their modulation in disease states with particular focus on their regulation in asthma, atherosclerosis and diabetes. The importance of small molecule inhibitors in the study of molecular pathways, the current use of common H₂S synthesizing enzyme inhibitors and the relevant characteristics of mice in which these enzymes have been genetically deleted will also be summarized. With a greater understanding of the molecular regulation of these enzymes in disease states, as well as the availability of novel small molecules with high specificity targeted towards H₂S producing enzymes, the potential to regulate the biological functions of this intriguing gas H₂S for therapeutic effect can perhaps be brought one step closer.

Keywords

H₂S synthesizing enzymes • CSE • CBS • 3-MST • Hydrogen sulfide

1 Introduction

Research into the biology of hydrogen sulfide (H₂S) over the last decade has exponentially increased our understanding of the way in which this gasotransmitter influences physiological and pathophysiological (i.e. disease) processes in a wide range of biological systems. With the characteristic and undoubtedly obnoxious odour of rotten eggs and an early history which appeared to mark H₂S as nothing more than a poisonous gas (Beauchamp et al. 1984), it is perhaps surprising that H₂S is synthesized endogenously in mammalian cells by both enzymatic and non-enzymatic mechanisms and that, once formed, H₂S produces the wide array of biological effects which are now being reported (Li et al. 2011). H₂S has therefore followed an interesting ‘development curve’ in the minds of researchers starting as an agent which was of toxicological (sometime devastatingly so) impact, via a gas of interest to pharmacologists to a molecule which now occupies a very much more central position in gas physiology and with growing therapeutic potential.

The last decade has seen multiple advances in our knowledge of the mechanisms by which H₂S is synthesized, how these are regulated, the tissue and cellular distribution of H₂S synthesizing enzymes in tissues and how this distribution changes in disease. These various areas form the central themes of this present review.

H₂S can be produced naturally in the body by at least three enzymes. Cystathionine γ -lyase (CSE), cystathionine β -synthetase (CBS) and 3-mercaptopyruvate sulfurtransferase (3-MST) are known to be significant producers of endogenous H₂S. Collectively they have been termed ‘H₂S synthesizing enzymes’. Both CSE and CBS are pyridoxal-5'-phosphate (PLP) dependent and utilize L-cysteine as substrate whilst 3-MST is non-PLP dependent and uses 3-mercaptopyruvate as substrate (Shibuya et al. 2009a; Steegborn

et al. 1999). Endogenous H₂S can also be produced non-enzymatically. For example, erythrocytes, supplemented with glucose and sulfur and containing electron carriers such as NADH, NADPH and reduced glutathione (GSH), spontaneously react with sulfur to produce H₂S (Searcy and Lee 1998). Iron-sulfur cluster containing proteins carrying Fe₂S₂, Fe₃S₄ or Fe₄S₄ clusters are another source of non-enzymatically generated H₂S. These include proteins such as ferredoxins and 'Rieske' proteins among others (Beinert et al. 1997). In the presence of reducing agents such as glutathione, H₂S can also be released from bound sulfur in, for example, persulfides in neurons and astrocytes (Ishigami et al. 2009). Although non-enzymatic routes for H₂S generation have been identified, it seems likely that the bulk, if not all, of the H₂S which is generated for the purpose of carrying out biological functions in the body is derived enzymatically. This no doubt explains the huge emphasis in recent years in understanding as much as possible about the biochemistry, molecular biology and physiological relevance of the H₂S synthesizing enzymes.

The list of reported biological effects of both endogenous and exogenously administered H₂S is far reaching and expanding on an almost weekly basis. Numerous molecular targets for this gas have also been identified. The biological effects of H₂S are outside the scope of this present review and are dealt with elsewhere in this volume. Rather, we will provide here an overview of the H₂S synthesizing enzymes. This will encompass a discussion of the way in which H₂S synthesizing enzymes are regulated in health and in disease, as well as a consideration of the limitations of the currently available H₂S synthesizing enzyme inhibitors which have been widely used to probe the biological roles of this gas.

2 H₂S Synthesizing Enzymes: The Beginnings

The detection of sulfide in brain tissue in 1989 prompted the suggestion that H₂S, produced endogenously, may have a role to play in physiology (Goodwin et al. 1989). The subsequent demonstration that H₂S produced by the enzyme CBS in the mammalian brain selectively enhanced NMDA receptor-mediated responses and facilitated the induction of hippocampal long-term potentiation led to the first proposed biological role of this gas as a neuromodulator (Abe and Kimura 1996). Shortly thereafter, the second H₂S producing enzyme, CSE, was found to be expressed in guinea pig ileum, rat portal vein and thoracic aorta, and the smooth muscle relaxant effect of H₂S was noted for the first time (Hosoki et al. 1997). This seminal work led to a number of detailed investigations which rapidly established the vascular (Zhao et al. 2001a, b) and gastrointestinal (Teague et al. 2002) muscle relaxant properties of this gas. Several years later, the third H₂S synthesizing enzyme (3-MST) was discovered when brain homogenates of CBS knockout mice were shown to produce H₂S at levels similar to that of wild-type mice in the absence of PLP, suggesting redundancy of CBS and the existence of yet another H₂S producing enzyme. This new enzyme, 3-MST, produced H₂S from 3-mercaptopyruvate, which is derived from cysteine in the presence of α -ketoglutarate. This latter reaction is catalysed by the enzyme cysteine

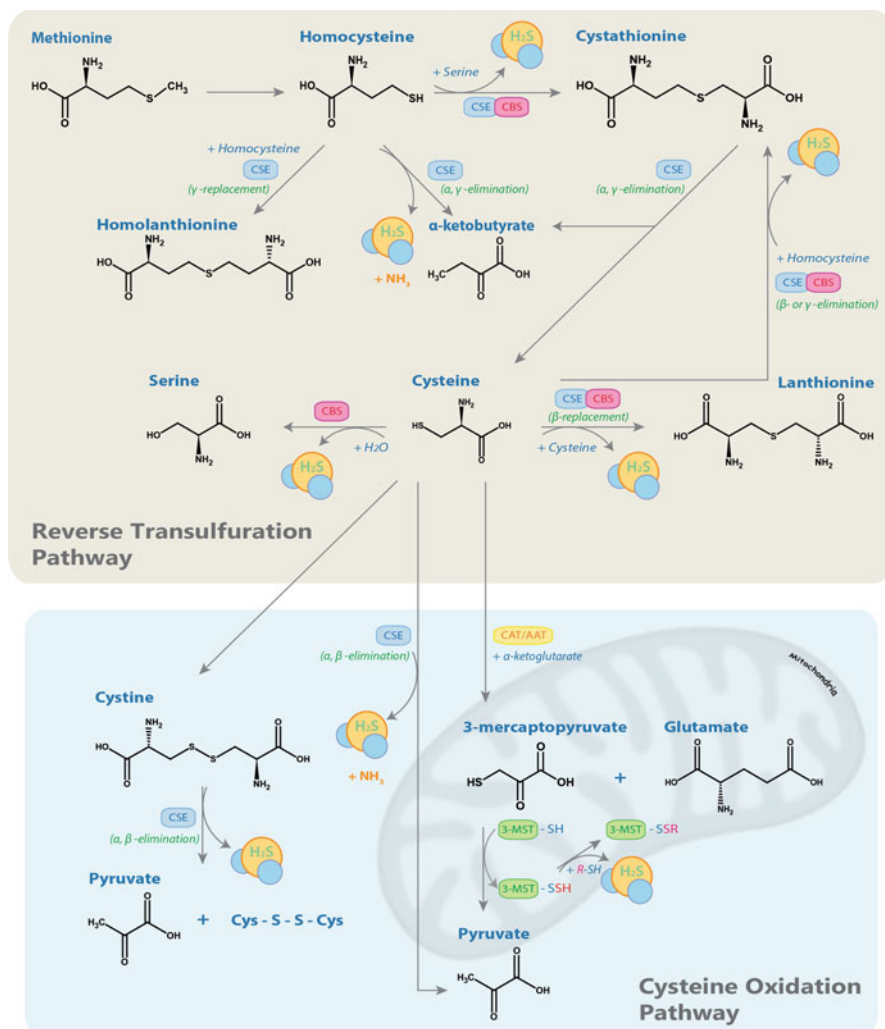


Fig. 1 Pathways for H₂S biosynthesis in the cell. The endogenous production of H₂S occurs via two main pathways—reverse transulfuration and cysteine oxidation which take place partly inside mitochondria. 3-MST, 3-mercaptopyruvate sulfurtransferase; CBS, cystathionine β -synthetase; CSE, cystathionine γ -lyase; H₂S, hydrogen sulfide; NH₃, ammonia

aminotransferase (CAT), which is identical with aspartate aminotransferase (AAT) (Fig. 1) (Shibuya et al. 2009b).

Over recent years, considerable attention has been focused on the biochemistry and molecular biology of the H₂S synthesizing enzymes. Much is now known about the mechanics of H₂S biosynthesis in terms of substrate specificity, cofactor requirement and tissue and cell distribution, and the molecular topology of these enzymes has also been described. Interested readers are referred to several excellent

reviews on this topic (e.g. Kabil and Banerjee 2014; Kabil et al. 2014; Nagahara et al. 2007; Singh and Banerjee 2011). In contrast, there have been few concerted attempts to review changes in the expression or activity of the H₂S synthesizing enzymes with disease states and this review will thence, in part, concentrate on this aspect.

3 Endogenous Synthesis of H₂S and Regulation of Its Synthesizing Enzymes Under Physiological Conditions

CBS catalyses the formation of cystathionine via two separate pathways. The first pathway involves linking homocysteine with serine (Stipanuk 1986) and the second pathway links homocysteine with cysteine (Chen et al. 2004). H₂S is produced only from the second reaction. Besides generating cystathionine, CBS also converts cysteine to serine and lanthionine, and in the process yields H₂S. These reactions are part of the reverse transsulfuration pathway of H₂S biosynthesis (See Fig. 1).

Although found in many organs including brain, kidney, liver, ileum, uterus, placenta (Patel et al. 2009) and pancreatic islets (Kaneko et al. 2006), mouse tissue-specific quantification of CBS revealed that this enzyme is predominantly expressed in the brain and kidney (Kabil et al. 2011a).

The regulation of CBS activity is dependent on the presence of several cofactors and activators. Besides requiring PLP to function, the NH₂ terminal end of CBS contains the binding site of a heme cofactor group that further regulates the activity of CBS by functioning as a redox-sensitive gas sensor (Banerjee and Zou 2005). Because CBS contains iron, the oxidation state of CBS can be toggled, viz. ferric (Fe³⁺) to ferrous (Fe²⁺) CBS by reducing agents such as NADPH in the presence of diflavin methionine synthase reductase (Kabil et al. 2011b). In the ferrous (Fe²⁺) state, CBS can be bound by other gasotransmitters like CO (Puranik et al. 2006) and NO (Taoka and Banerjee 2001), resulting in downregulation of the activity of CBS. Subsequently, the oxidation of the ligand-bound CBS (i.e. CO-CBS or NO-CBS) by air to the ferric (Fe³⁺) state restores the catalytic activity of CBS. Such a regulatory mechanism suggests a complex interplay between the different gasotransmitters in controlling endogenous H₂S biosynthesis at least via the CBS route. This would further imply that in conditions such as inflammation or hypoxia in which cellular NO and CO levels may be either increased or decreased, the precise effect on CBS activity and hence H₂S generation by this enzyme will be difficult to predict.

CBS activity is also regulated by allosteric binding of S-adenosylmethionine (SAM) (Finkelstein et al. 1975), which is a co-substrate that is involved in a multitude of other metabolic pathways including transmethylation, transsulfuration and aminopropylation (Cantoni 1952). Using mice in which the *MAT1A* gene (which encodes for SAM) has been disrupted and human hepatocellular carcinoma cells which have the *MAT1A* gene silenced, it was demonstrated that the binding of SAM to CBS stabilizes the protein against proteolysis (Prudova et al. 2006). In vitro, SAM increased CBS enzyme activity and thence H₂S production (Finkelstein et al. 1975).

CBS activity is not only regulated by cofactors and activators but also by its location within the cell. Using yeast cells as a model system, human CBS has been shown to be post-translationally modified by the small ubiquitin-like modifier-1 (SUMO-1) protein at its C-terminal regulatory domain. SUMOylation enables CBS to be shuttled into the nucleus where it associates with the nuclear scaffold (Kabil et al. 2006). SUMOylation of CBS was enhanced by human polycomb group protein 2 (hPc2). It has been proposed that nuclear localization of CBS occurs when local glutathione demand is high, since CBS generates cysteine, which is the rate-limiting amino acid in glutathione synthesis (Agrawal and Banerjee 2008). Coupled with the observation that regulation of CBS was correlated with the rate of proliferation in human and yeast cells through a redox-sensitive mechanism (Maclean et al. 2002), it is possible that the presence of CBS in the nucleus provides a source of an antioxidant (i.e. H₂S) in the cell when, and if, required. This possibility warrants further investigation to determine whether such a mechanism does indeed exist in cells either *in vitro* or *in vivo*.

Like CBS, cystathionine γ -lyase (CSE) uses homocysteine as substrate to produce H₂S, α -ketobutyrate, ammonia and homolanthionine. CSE can also generate H₂S from cysteine, forming cystathionine and pyruvate (See Fig. 1). Although both homocysteine and cysteine are substrates of CSE, at physiological concentrations of cysteine and homocysteine, kinetic simulations employing physiologically relevant concentrations of homocysteine and cysteine using the dithiobisnitrobenzene (DTNB) assay showed that α - and β -elimination of cysteine catalysed by CSE accounts for approximately 70 % of the H₂S produced while the α,γ -elimination of homocysteine contributes about 29 %. However, once homocysteine levels are elevated to those apparent in hyperhomocysteinemia, then about 90 % of the H₂S generated by CSE occurs via α,γ -elimination and γ -replacement reactions of homocysteine (See Fig. 1) (Chiku et al. 2009). This indicates that the identity of the main substrate for CSE-dependent generation of H₂S in cells is likely to depend on the relative availability of cysteine and homocysteine.

CSE is located in a wider range of tissues than CBS. CSE activity is several-fold higher in human brain than in mouse brain (Diwakar and Ravindranath 2007). The mouse small intestine, stomach (Ishii et al. 2004), rat portal vein and thoracic aorta (Hosoki et al. 1997), mouse pancreas (Kaneko et al. 2006), rat uterus, human placenta, myometrium, amnion and chorion (Patel et al. 2009) all express CSE. As compared to CBS, CSE is more abundant in mouse liver where it is the principal enzyme responsible for H₂S generation (Kabil et al. 2011a).

Like CBS, the activity of CSE is also regulated by a number of cofactors. CSE was initially reported to be activated physiologically by calcium-calmodulin after stimulation of muscarinic cholinergic receptors in vascular endothelial cells in culture, as determined by an increase in CSE activity and H₂S production. Through co-immunoprecipitation experiments, recombinant CSE was found to bind directly to calmodulin (Yang et al. 2008). Although Ca²⁺ has been suggested to activate CSE, the precise concentration of Ca²⁺ is critical as too much Ca²⁺ inhibits purified rat CSE-mediated H₂S production in the presence of PLP. At physiologically relevant Ca²⁺ concentrations (i.e. ~100 nM) H₂S was efficiently produced in the

presence of PLP. However, H₂S production was decreased at 300 nM Ca²⁺ and reduced even further at concentrations up to 3 μM. The involvement of calmodulin in regulating CSE activity however is not yet clear since a recent study showed that neither calmodulin nor a calmodulin antagonist, W-7, affected the production of H₂S by purified rat liver CSE (Mikami et al. 2013).

SUMO, besides decreasing CBS activity, also targets CSE in vitro, in that recombinant human CSE can also be SUMOylated (Agrawal and Banerjee 2008). However, whether the SUMOylation of CSE occurs within the cell and has a physiological effect in consequence is not clear. If this is indeed the case then the re-localization of both CBS and CSE into the nucleus may be a strategy to increase cysteine levels inside the nucleus when the demand for glutathione is increased such as during early phases of cell proliferation (Markovic et al. 2007) or when telomerase activity is high as observed in 3T3 fibroblast cells (Borrás et al. 2004).

As an important intracellular antioxidant protecting the cell against oxidative stress, physiological glutathione levels are well controlled via glutathione synthesis in the transsulfuration pathway. Without glutathione, numerous cell types including lymphocytes (Franco and Cidlowski 2009), embryos (Dalton et al. 2004) and hepatocytes (Ghibelli et al. 1999) undergo apoptosis due to excess reactive oxygen species (ROS) generation. Since activation transcription factor 4 (ATF4) controls expression of genes responsible for amino acid metabolism and redox status, ATF4 might, by increasing glutathione synthesis through the transsulfuration pathway, be a key cellular survival factor. Indeed, the absence of ATF4 in mouse embryonic fibroblasts resulted in a substantial fall in glutathione levels due to downregulation of CSE expression (Dickhout et al. 2012). Thus, if the amount of H₂S synthesizing enzymes in the nucleus are reduced, the levels of glutathione would subsequently be reduced, thereby affecting the fate of the cell.

The most recently discovered H₂S synthesizing enzyme 3-MST, unlike CBS and CSE, is PLP independent. 3-MST catalyses the conversion of 3-mercaptopyruvate to pyruvate by degrading cysteine. In this pathway, cysteine is first converted by cysteine aminotransferase (CAT), also known as aspartate aminotransferase (AAT), to 3-mercaptopyruvate via the incorporation of α-ketoglutarate into the reaction. 3-MST then forms a persulfide by transferring a sulfur from 3-mercaptopyruvate onto itself which in the presence of a reductant like thioredoxin produces pyruvate and H₂S (Nagahara et al. 2007; Yadav et al. 2013). Dihydrolipoic acid has also been identified to associate with 3-MST to release H₂S (Mikami et al. 2011a).

Like CBS and CSE, 3-MST is found in many tissues with high activity in the proximal tubular epithelium of the kidney, pericentral hepatocytes in the liver, cardiac cells and neuroglial cells (Shibuya et al. 2009b). Within the cell, 3-MST is located both in the cytoplasm as well as the mitochondria. However, because the concentration of cysteine is much higher in the mitochondria, as compared to the cytoplasm, it is plausible that most of the H₂S generated by 3-MST occurs in mitochondria (Nagahara et al. 1998).

H₂S production by 3-MST can be significantly inhibited by aspartate which is a potent inhibitor of CAT (Akagi 1982). H₂S produced by 3-MST and CAT in retinal neurons is also regulated by Ca²⁺. A negative feedback mechanism has been

proposed whereby H₂S produced after activation by Ca²⁺ in turn suppresses Ca²⁺ influx into photoreceptor cells by activating the vacuolar-type H(+)-ATPase (V-ATPase). The activated V-ATPase then releases protons into the synaptic cleft thereby suppressing Ca²⁺ channels found in the photoreceptor cells (Mikami et al. 2011b).

Unlike CBS and CSE, the activity of which is regulated for the most part by binding of other factors, 3-MST activity is probably regulated intrinsically by its redox state. Structural studies have shown that 3-MST contains catalytic site cysteines (Cys²⁴⁷, Cys¹⁵⁴ and Cys²⁶³) each of which is redox active (Nagahara 2013). Cys¹⁵⁴ is unique to rat 3-MST whilst Cys²⁶³ is conserved in mammalian 3-MST (Nagahara et al. 2007). The activity of 3-MST is regulated by both intermolecular and intramolecular redox-sensing switches as reviewed elsewhere (Nagahara 2013). Nevertheless, it is important to note that the redox-dependent modulation of 3-MST may not be relevant in human MST since the cysteine residues (i.e. Cys¹⁵⁴) which are involved in the cross-linking in rat MST are not conserved in the human enzyme (Nagahara et al. 2007).

4 Molecular Regulation of H₂S Synthesizing Enzymes

Hydrogen sulfide contributes to cell signalling by regulating the function of a number of molecular targets including ion channels (Sun et al. 2008; Zhao et al. 2001b), kinases (Hu et al. 2007) and transcription factors such as signal transducer and activator of transcription (STAT) (Calvert et al. 2009; Li et al. 2009a) as well as nuclear factor-kappaB (NFκB) (Li et al. 2011; Oh et al. 2006). However, the expression of the enzymes that are responsible for producing H₂S (i.e. CSE, CBS and 3-MST) are also frequently regulated at a transcriptional and subsequently at a protein level by multiple molecular factors. In many disease states, the expression of these enzymes is both inducible and dynamic. With a better understanding of how the H₂S synthesizing enzymes are regulated in disease states, a more targeted effort could be developed to identify drugs to affect endogenous production of H₂S in disease states with a view to restoring physiological H₂S levels.

To date, changes in expression (mRNA and/or protein) of H₂S synthesizing enzymes have been identified in a number of cells and tissues in a variety of different disease states and involving the intermediacy of a range of different signalling pathways (summarized in Table 1). Most evidence collected to date suggests a pivotal role for NFκB in the control of cellular CSE and CBS expression, and this interaction will therefore be described in some detail below.

In an early study seeking to identify the molecular targets of H₂S, it was shown that H₂S inhibited the activation of NFκB in lipopolysaccharide (LPS)-stimulated mouse macrophages. The mechanism is believed to be via increased heme-oxygenase-1 (HO-1) expression under the control of an ERK1/2-dependent pathway. Increased HO-1 expression then led to reduced IκB phosphorylation and degradation and impaired translocation of NFκB into the nucleus

Table 1 Examples of modulated H₂S producing enzymes in selected disease states

Disease/stress	Model	Factors/signalling involved	Location	Regulation of H ₂ S producing enzyme	Reference
Atherosclerosis	RAW264.7 macrophages	Akt signalling	-	↑ CSE (mRNA & protein)	Xu et al. (2014b)
	High-fat diet in mice	-	Liver, lung Liver, kidney	↓ CSE (protein) ↑ CBS (protein) ↓ 3-MST (protein)	Peh et al. (2014)
	High-fat diet in mice	-	Liver	↑ CSE & CBS (mRNA)	Hwang et al. (2013)
Asthma	Ovalbumin-treated mice	-	Lung	↓ CSE (protein)	Zhang et al. (2013)
	Ovalbumin-treated rats	-	Lung	↓ CSE (protein)	Chen et al. (2009)
Bowel syndromes	Visceral hyperalgesia in rats	Voltage-gated sodium channel	Colon	↑ CBS (protein)	Wang et al. (2012b)
	Chronic visceral hyperalgesia in rats	NFκB	Colon	↑ CBS (protein)	(Li et al. 2012)
Cancer	Colon cancer	-	Colon	↓ 3-MST (protein)	Ramasamy et al. (2006)
	Human hepatocellular carcinoma (HCC) cell lines	Sp1, PI3K/Akt (PKB)	-	↑ CSE (protein)	Yin et al. (2012)
	Down's syndrome megakaryocytic leukaemia cell line (CMK)	Sp1, Sp3	-	↑ CBS (mRNA)	Ge et al. (2003)
Diabetes	Streptozotocin-treated rats	-	Liver	↓ CSE (protein)	Manna et al. (2014)
	INS-1E cells, rat pancreatic islets	Glucose	-	↓ CSE (mRNA & protein)	Zhang et al. (2011)
	Streptozotocin-treated rats	Insulin	Liver	↑ CBS (mRNA)	Ratnam et al. (2002)

(continued)

Table 1 (continued)

Disease/stress	Model	Factors/signalling involved	Location	Regulation of H ₂ S producing enzyme	Reference
Down's syndrome	Patients with Down's syndrome	-	Astrocytes	↑ CBS (protein)	Ichinohe et al. (2005)
Hypoxia	U87-MG human glioblastoma cells, PC12 rat cells, rats exposed to hypoxic conditions	Hypoxia-inducible factors (HIFs)	Cerebral cortex	↑ CBS (mRNA & protein)	Takano et al. (2014)
Inflammation	LPS-treated mammalian cell lines (HEK-293, COS-7, J774.1A, RAW264.7) Temporomandibular joint pain in rats	NFκB Inhibition of voltage-gated potassium currents	- Trigeminal ganglion neuron	↑ CSE (mRNA & protein) ↑ CBS (mRNA & protein)	Wang et al. (2014) Miao et al. (2014)
Oxidative stress	LPS-induced inflammation in rats	Dexamethasone	Neutrophils	↓ CSE (protein)	Li et al. (2009b)
	LPS-induced inflammation in mice	-	Liver, kidney	↑ CSE (mRNA)	Li et al. (2005)
	Mammalian cells (A549, SMMC-7721)	H ₂ O ₂	-	↑ CSE (mRNA & protein)	Wang et al. (2012a)
Renal ischaemia	Rat mesangial cells	Platelet-derived growth factor (PDGF)-BB, STAT, Nrf2	-	↑ CSE (mRNA & protein)	Hassan et al. (2012)
	Renal ischaemia/reperfusion in rats	Sp1	Kidney	↓ CBS (mRNA & protein)	Wu et al. (2010)
Ulcerations	Gastric ulcers in rats	-	Stomach	↑ CSE & CBS (protein)	Wallace et al. (2007)

3-MST 3-mercaptopyruvate sulfurtransferase, *Akt/PKB* protein kinase B, *CBS* cystathionine β-synthetase, *CSE* cystathionine γ-lyase, *H₂S* hydrogen sulfide, *H₂O₂* hydrogen peroxide, *Nrf2* nuclear factor like 2, *PI3K* phosphoinositide 3-kinase, *Sp1* specificity protein 1, *STAT* signal transducer and activator of transcription

(Oh et al. 2006). Later studies confirmed that H₂S affects NFκB but by a different mechanism. In this case, H₂S was shown to sulfhydrate NFκB on the p65 subunit at the cysteine-38 residue. This molecular event then increased NFκB binding to a co-activator ribosomal protein, S3 (RPS3), which triggers the transcription of antiapoptotic genes (Sen et al. 2012). Interestingly, recent work has shown that NFκB also regulates CSE expression by binding to its promoter region in macrophages challenged with LPS (Wang et al. 2014). In a separate study, rats with chronic visceral hyperalgesia displayed upregulated colonic CBS expression. An inhibitor of the p65 subunit of NFκB, pyrrolidine dithiocarbamate (PDTC), reduced the expression of CBS, suggesting that activation of the p65 subunit of NFκB upregulates CBS expression (Li et al. 2012). Thus, it appears that H₂S may act to maintain the levels of CSE and CBS inside the cell in check during an inflammatory response by regulating the levels of NFκB.

Drugs have also been shown to regulate CSE expression. In the case of atherosclerosis, statins, a class of drugs used to lower cholesterol levels by inhibiting the enzyme 3-hydroxy-3-methyl-glutaryl-CoA reductase (HMG-CoA reductase), were recently reported to upregulate CSE transcription and subsequent H₂S generation. This effect was mediated by activation of Akt signalling (Xu et al. 2014b). In inflammation, dexamethasone, a glucocorticoid steroid, reduced CSE expression in neutrophils isolated from LPS-challenged rats. Dexamethasone, in this model, was proposed to inhibit the formation of pro-inflammatory H₂S, most probably through reduced NFκB-mediated CSE expression (Li et al. 2009b).

In certain disease conditions, the effect of the disease itself could also have a role to play in directly regulating the expression of H₂S synthesizing enzymes. For example in diabetes, high glucose significantly reduced CSE mRNA and CSE activity levels in freshly isolated rat pancreatic islets and in the rat pancreatic cell line, INS-1E. This was mediated by stimulating phosphorylation of the transcription factor SP1 via p38 MAPK activation which subsequently led to decreased CSE promoter activity (Zhang et al. 2011).

5 H₂S Synthesizing Enzymes in Diseases States

5.1 Asthma

Asthma is a chronic inflammatory disease characterized by shortness of breath, wheezing and recurring cough. It is caused by a combination of excess airway mucus production and reversible airflow obstruction. Immune cells such as eosinophils and neutrophils play a major role in this inflammatory state although different patients with severe asthma may exhibit different phenotypes that need to be properly defined if the most effective treatment is to be provided (Gibeon and Chung 2012). Current methods for distinguishing different phenotypes of asthma involve detecting sputum eosinophils (eosinophilic asthma), serum periostin levels which is a biomarker for eosinophilic asthma (Jia et al. 2012) and breath nitric oxide (NO) levels (Dweik et al. 2011). H₂S is one such potential biomarker of asthma

given that it was found to be elevated in the sputum of asthmatics (Chung 2014; Saito et al. 2013).

Endogenous serum H₂S levels were found to be altered in chronic obstructive pulmonary disease (COPD) in man almost a decade ago (Chen et al. 2005). This group reported that endogenous serum H₂S levels were higher in patients with COPD as compared to patients with acute exacerbation of COPD (AECOPD). Serum levels decreased in patients with stable COPD as the condition became clinically more severe. Subsequently, serum H₂S levels in asthmatic children were reportedly decreased compared with a matched healthy control group (Tian et al. 2012).

A more recent clinical study examined levels of H₂S in serum and sputum from 40 patients with varying degrees of severity of asthma and compared the data obtained with that of 15 healthy subjects (Saito et al. 2013). H₂S levels in sputum of asthma patients were significantly higher than those in sputum from healthy subjects. Serum H₂S concentration in asthmatics was also 10 times higher than that in sputum. Sputum measurements are more likely to be indicative of asthma as measurement of serum H₂S will be confounded by the presence of H₂S generated by non-respiratory tissues. Moreover, a positive correlation was identified between sputum H₂S levels and sputum neutrophil number. These observations raise the possibility that sputum H₂S concentration may perhaps be a biomarker of neutrophilic asthma.

Since endogenous H₂S levels are modulated during asthma, it is conceivable that the activity or expression of H₂S synthesizing enzymes may also be altered at one or other time in the course of the disease (See Table 1). To date, there have been few studies to investigate this possibility, although there are clues in the literature that this may be the case. For example, it has recently been shown, using mouse lung slices, that exogenous H₂S impeded airway contraction of smooth muscle cells by inhibiting intracellular Ca²⁺ release evoked by inositol-1,4,5-triphosphate (InsP3). This effect was also apparent when murine lung tissues was treated with L-cysteine (an endogenous H₂S precursor) which effect was reversed by the irreversible CSE inhibitor, DL-propargylglycine (PAG) (Castro-Piedras and Perez-Zoghbi 2013). More direct evidence comes from experiments using an ovalbumin (OVA)-induced mouse model of acute asthma. In this case, CSE expression was downregulated in the lungs of wild-type mice challenged with OVA, whilst mice in which CSE was knocked out (CSE^{-/-}) displayed augmented airway inflammation following ovalbumin treatment as indicated by higher levels of Th2 cytokines such as IL-5, IL-13 and eotaxin-1. Exogenous H₂S administration, in the form of NaHS, effectively 'rescued' CSE^{-/-} mice from the exacerbated symptoms of asthma strongly implying that an upregulated CSE/H₂S system has a protective role during the development, or perhaps the maintenance, of asthma in this model (Zhang et al. 2013). Whether such a mechanism also occurs in man is not yet known. Considerable evidence now exists, reviewed elsewhere in this volume, that H₂S exhibits both pro- and anti-inflammatory effects in a range of animal models. Thus, whether an H₂S donor administered may prove beneficial or deleterious in asthma remains an open question.

5.2 Atherosclerosis

H₂S exhibits pronounced vasodilator activity both *in vitro* and *in vivo*, and it is hence no surprise that deranged H₂S biosynthesis has been implicated in a number of cardiovascular diseases including atherosclerosis. H₂S dilates blood vessels by various mechanisms, including the opening of potassium-activated ATP channels (K_{ATP} channels) in vascular smooth muscle (Zhao et al. 2001a) as well as potentially via intracellular acidification by activating of the Cl⁻/HCO₃⁻ exchanger (Lee et al. 2007). Atherosclerosis is a chronic and complex inflammatory condition. Its pathogenesis involves an intricate tapestry of prolonged immune cell recruitment and cytokine/chemokine secretion coupled with the presence of reactive oxygen species (ROS), as well as vascular smooth muscle cell proliferation and migration (Ross 1999). Intriguingly, each of these separate events has been reported to be regulated by H₂S (Du et al. 2004; Yan et al. 2006). However based on the evidence available thus far, the link between H₂S and atherosclerosis and the involvement of the H₂S synthesizing enzymes in the process is far from clear.

In a recent study, Peh and colleagues reported reduced expression of CSE and CBE protein in liver, kidneys and lungs of mice fed a high-fat diet. 3-MST expression was also reduced in liver. In addition, CSE/CBS and 3-MST enzyme activity was also diminished as determined by *ex vivo* H₂S synthesis using either cysteine or 3-mercaptopyruvate as substrates. It should be noted that these animals showed no evidence of frank atherosclerosis in that plasma levels of serum amyloid A (SAA) and C-reactive protein (CRP) were normal and the histological appearance of blood vessels revealed no ongoing disease. Changes in the expression of these enzymes were therefore likely to be a consequence of fat feeding and not a reaction to vascular disease. If this is the case, then changes in H₂S synthesis may be presumed to occur before the onset of atherosclerosis (Peh et al. 2014). Whether measurement of these enzymes in susceptible individuals might be a surrogate biomarker of impending vascular damage due to atherosclerosis remains to be determined. Intriguingly, in a separate study, mice fed a high-fat diet leading to the development of fatty liver exhibited higher levels of CSE and CBS in the liver (Hwang et al. 2013). The mechanisms which control CSE, CBS and 3-MST expression in animals consuming fat in their diet warrant further study.

Other researchers have studied H₂S synthesizing enzyme expression and activity in apolipoprotein E (ApoE) knockout mice. These animals have increased plasma levels of cholesterol, triglyceride and low-density lipoprotein cholesterol, develop a frank atherosclerotic state and also show evidence of defective H₂S biosynthesis including lower production of H₂S in aorta and reduced plasma H₂S concentration. Interestingly, CSE mRNA expression in the aorta was actually increased. Administration of exogenous H₂S (NaHS) elevated plasma H₂S and reduced aortic CSE mRNA thereby suggesting an inverse correlation between CSE expression and H₂S levels in the aorta which may be a consequence of a positive feedback mechanism in which decreased H₂S production (due to vascular disease) is compensated for by increased CSE gene expression (Wang et al. 2009). Interestingly, elevated plasma H₂S has also been reported in patients with atherosclerosis (Peter et al. 2013), whilst

diminished production of endogenous H₂S which is apparent in CSE^{-/-} mice is associated with accelerated atherosclerosis (Mani et al. 2013). In this latter work, the rate of development of atherosclerosis in these animals could be reduced by replacement therapy with NaHS. The current consensus is therefore that lack of H₂S contributes to accelerated progression of atherosclerosis (Szabó et al. 2011; Xu et al. 2014a; Zhang et al. 2012).

5.3 Diabetes

Altered expression of H₂S synthesizing enzymes as well as endogenous H₂S levels has been observed in diabetic animals. In an early study, plasma H₂S and the expression of CSE and CBS were elevated in the pancreas of streptozotocin (STZ)-induced diabetic rats (Yusuf et al. 2005) which suggests a possible correlation between Type 1 diabetes (Schnedl et al. 1994) and increased pancreatic and liver H₂S synthesis. To evaluate more precisely the role of H₂S in diabetes, a subsequent study made use of CSE^{-/-} mice injected with STZ to provoke a diabetic state. Since the pathogenesis of diabetes is associated with reduced number, but increased activity, of K_{ATP} channels in β -pancreatic cells, these authors sought to compare the mass and K_{ATP} channel activity of β -pancreatic cells in CSE^{-/-} mice with that of wild-type mice (Yang et al. 2011). CSE deficiency protected mice from STZ-induced diabetes as well as the damage and dysfunction of pancreatic islets—as determined by measurement of the changes in insulin secretion and electrophysiological recording of K_{ATP} channel currents after induction of a diabetic state.

Curiously, other studies have reported diametrically opposite results. For example, it was previously noted that circulating H₂S levels were lower (c.f. control, non-diabetic animals) in both STZ-treated rats and non-obese diabetic (NOD) models of Type 1 diabetic mice (Brancaleone et al. 2008; Jain et al. 2010). A similar conclusion was reached from the examination of blood H₂S levels in humans with diabetes (Jain et al. 2010). CSE enzyme activity was also reduced in the liver of Type 1 diabetic rats as well as in peripheral blood mononuclear cells from Type 1 diabetic patients (Manna et al. 2014).

In contrast to Type 1 diabetes which is caused by the death of β -pancreatic cells, Type 2 diabetes arises, usually later in life, with the development of insulin resistance. To study the possible role of H₂S in this process, 6-month-old CSE^{-/-} mice were fed for 8 weeks with a high-fat diet in order to induce hyperglycaemia (Okamoto et al. 2013). The mice developed impaired glucose tolerance and decreased insulin content in their pancreatic cells as compared to their wild-type counterparts. It was noted that the lack of CSE in these animals promoted thioredoxin-binding protein-2 (TBP-2) gene expression which increases insulin insensitivity and causes glucose intolerance (Yoshihara et al. 2010). These results indicate that changes in expression or activity of CSE (and perhaps other H₂S synthesizing enzymes) are implicated in the outcome and severity of both Type 1 and 2 diabetes.

6 Frequently Used Inhibitors and Mouse Knockout Models of H₂S Synthesizing Enzymes

To study the biological effects of endogenously synthesized H₂S, as well as to study the characteristics and behaviour of the H₂S synthesizing enzymes, the use of selective inhibitors of H₂S synthesizing enzymes is crucial. Currently, several inhibitors of CSE and CBS have been used (See Fig. 2) although to date, pharmacological inhibitors of the enzyme 3-MST have yet to be identified. In this context, L-aspartate inhibits the production of H₂S by 3-MST indirectly by inhibiting CAT/AAT (Akagi 1982). Mice with genetic deletion of one or other of the H₂S synthesizing enzymes are also beginning to show their value to the field as well. Of the three known H₂S synthesizing enzymes only CSE^{-/-} mice have been widely used as 90 % of CBS^{-/-} mice died during the first 2 weeks of neonatal life because of growth retardation and severe hepatopathy (Watanabe et al. 1995) and only lately has it been possible to produce 3-MST knockout mice (Nagahara 2013). However, recently a non-neonatal fatal mouse model with the *cbs* gene inactivated but instead expresses low levels of the human CBS transgene and exhibits classical homocystinuria could provide a useful, alternative to a CBS^{-/-} mouse model (Maclean et al. 2010).

Some of the commonly used pharmacological inhibitors of CSE are DL-propargylglycine (PAG) (Marcotte and Walsh 1975) which irreversibly inhibits CSE by physically obstructing the access of the substrate to the active site of the enzyme (Sun et al. 2009) and the reversible inhibitor β-cyanoalanine (BCA) which has also been proposed as a competitive inhibitor of this enzyme (Pfeffer and Ressler 1967). However, these compounds have frequently been claimed to exhibit poor selectivity, require high concentrations and have limited ability to permeate the cell membrane (Szabó 2007). For example, PAG was reported many years ago to cause an irreversible inactivation of both aspartate aminotransferase (Tanase and Morino 1976) and alanine transaminase (Burnett et al. 1980). Aminoxyacetic acid (AOAA) is also very commonly used and is often stated to be a selective inhibitor of CBS (d'Emmanuele di Villa Bianca et al. 2009; Oh et al. 2006; Roy et al. 2012). However, experiments using recombinant human CSE and CBS enzymes have revealed that AOAA inhibits both CSE and CBS. Indeed, AOAA appeared to be

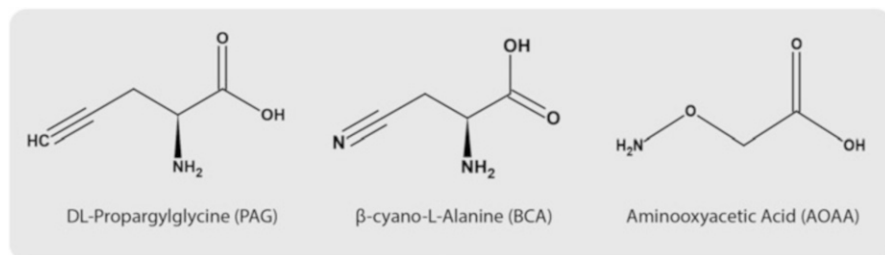


Fig. 2 Chemical structures of compounds that inhibit CSE or CBS

an even more potent inhibitor of CSE than CBS at least under these experimental conditions (Asimakopoulou et al. 2013). Moreover, AOAA, like PAG, is a general inhibitor of pyridoxal phosphate-dependent enzymes and has been reported to inhibit enzymes such as 4-aminobutyrate aminotransferase (GABA-T) (WALLACH 1961) and aspartate aminotransferase (Kauppinen et al. 1987).

7 Conclusion

Much has been learnt about the physiological and pathophysiological implications of H₂S since it was realized that this evanescent gas is formed naturally in cells and tissues in the 1990s. Indeed, research in the last two decades has thrown H₂S very much into the spotlight and it is now widely referred to as the so-called third gasotransmitter after CO and NO. Many mechanisms governing the regulation of H₂S synthesizing enzymes both in homeostatic and disease conditions have been elucidated. However, many questions still remain to be answered. For example, the quantitative contribution that each of the three H₂S synthesizing enzymes makes to the levels of H₂S found naturally in cells in health and disease remains unclear. This is important since, as detailed in this review, the presence and/or activity of such enzymes may be useful biomarkers of disease progression in some cases and it is critical to understand which H₂S synthesizing enzyme is important in which cell at which time point and how this is affected by disease. Moreover, whilst the biological significance of H₂S has been under intense scrutiny for more than two decades, there are still no reliably potent and selective inhibitors of any of the three H₂S synthesizing enzymes. For comparison, potent and selective inhibitors of both cyclooxygenase and nitric oxide synthase are available and have played crucial roles in understanding the complex biological roles of prostanoids and nitric oxide, respectively. Potent and targeted inhibitors of CSE, CBS and 3-MST could be expected to fill a similar niche. Finally, the precise intracellular signalling pathways which determine the degree of expression of H₂S synthesizing enzymes in healthy and in diseased cells have yet to be fully elucidated.

While many advances have been made, more selective and potent H₂S synthesizing enzyme inhibitors as well as cells in which the H₂S synthesizing enzymes (both alone and in combination) have been knocked out would certainly provide powerful tools to aid the understanding of the role of endogenous H₂S in biology. With the recent advent of high-throughput tandem-microwell assays that can screen enormous libraries of potential H₂S producing enzyme inhibitors (Zhou et al. 2013) as well as the introduction of 3-MST^{-/-} mice and novel transgenic mouse models of CBS-deficient homocystinuria, there is certainly much to look forward to in moving the field of H₂S biology forward towards its therapeutic potential.

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Part II

Molecular Mechanisms of Action

Persulfidation (S-sulfhydration) and H₂S

Milos R. Filipovic

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Abstract

The past decade has witnessed the discovery of hydrogen sulfide (H_2S) as a new signalling molecule. Its ability to act as a neurotransmitter, regulator of blood pressure, immunomodulator or anti-apoptotic agent, together with its great pharmacological potential, is now well established. Notwithstanding the growing body of evidence showing the biological roles of H_2S , the gap between the macroscopic descriptions and the actual mechanism(s) behind these processes is getting larger. The reactivity towards reactive oxygen and nitrogen species and/or metal centres cannot explain this plethora of biological effects. Therefore, a mechanism involving modification of protein cysteine residues to form protein persulfides is proposed. It is alternatively called S-sulfhydration. Persulfides are not particularly stable and show increased reactivity when compared to free thiols. Detection of protein persulfides is still facing methodological limitations, and mechanisms by which H_2S causes this modification are still largely scarce. Persulfidation of protein such as K_{ATP} could contribute to H_2S -induced vasodilation, while S-sulfhydration of GAPDH and NF- κ B inhibits apoptosis. H_2S regulates endoplasmic reticulum stress by causing persulfidation of PTP-1B. Several other proteins have been found to be regulated by this posttranslational modification of cysteine. This review article provides a critical overview of the current state of the literature addressing protein S-sulfhydration, with particular emphasis on the challenges and future research directions in this particular field.

Keywords

Hydrogen sulfide • Polysulfides • Sulfenic acids • Persulfidation • S-sulfhydration • S-nitrosation

1 H_2S as a Signalling Molecule

In order to maintain life, nature actually uses a limited number of chemical reactions, one of which is sulfur-based chemistry, mainly exploited for the control of intracellular redox homeostasis and redox-based signalling. Hydrogen sulfide (H_2S) is the simplest of the thiols found in the cells, and ever since the first report of its potential physiological role (Abe and Kimura 1996), there has been a growing literature on the subject of H_2S signalling. Very fast, H_2S joined the other two gases, nitric oxide (NO) and carbon monoxide (CO), as the third gasotransmitter (Wang 2002; Li et al. 2009; Mustafa et al. 2009a).

Numerous are the physiological functions assigned to be exclusively or partly regulated by H_2S , some of which are vasodilation (Yang et al. 2008; Mustafa et al. 2011), neurotransmission (Abe and Kimura 1996; Kimura et al. 2005), angiogenesis (Papapetropoulos et al. 2009; Szabó and Papapetropoulos 2011), inflammation (Li et al. 2005; Whiteman and Winyard 2011), hypoxia sensing

(Olson et al. 2008; Peng et al. 2010), etc. In addition, H₂S showed a tremendous pharmacological potential in preventing ischemia–reperfusion injury (Calvert et al. 2009, 2010). Furthermore, H₂S is able to induce suspended animation-like state in mice (Blackstone et al. 2005). Several pharmacological donors of H₂S have been developed with hope of their eventual use in disease treatment (Sparatore et al. 2008; Zhou et al. 2012; Szczesny et al. 2014; Zhao et al. 2013).

H₂S is produced by the action of at least three enzymes, cystathionine beta synthase (CBS), cystathionine gamma lyase (CSE) and mercaptopyruvate sulfurtransferase (MST) (Kabil et al. 2014; Kabil and Banerjee 2014). Differently expressed in different tissues (and even cellular compartments), these enzymes control H₂S production with different efficiencies. How are these enzymes regulated to maintain spatio-temporal production/distribution of H₂S is still unclear.

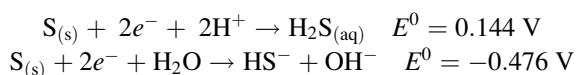
The intracellular levels of H₂S are also a matter of debate, with values spanning from nondetectable to >100 μM, although it seems more probable that the steady-state levels are at low micromolar and/or submicromolar levels, depending on the tissue (Olson 2012; Olson et al. 2014; Kabil et al. 2014). Conversely, the flux of H₂S production is huge, almost as that of glutathione, suggesting that the removal of H₂S is an efficient and tightly regulated process (Vitvitsky et al. 2012; Kabil et al. 2014; Kabil and Banerjee 2014).

Hydrogen sulfide is a weak acid and immediately ionizes in aqueous solution reaching the equilibrium between H₂S/HS[−]/S^{2−} species:



The pK_{a1} is around 6.9, while pK_{a2} is estimated to be >12, which means that under physiological conditions, approximately two thirds of H₂S are in the form of bisulfide (hydrogen(sulfide)(1-)), with negligible amounts of sulfide anion (sulfide(2-)). H₂S diffuses freely through the membranes (Mathai et al. 2009; Cuevasanta et al. 2012). Although the diffusion coefficient profile of H₂S is systematically lower than that of H₂O, the differences in the transmembrane Gibbs energy profiles are more dominant. Because of its hydrophobicity, H₂S experiences no barrier to permeation, so it can partition into the interior of the membrane readily (Riahi and Rowley 2014).

Sulfur atom in H₂S is in −2 oxidation state, but sulfur is very versatile in its ability to accept or donate electrons. It can cycle between −2 and +6 oxidation state, due to the six valence electrons and completely empty 3d orbital. Although the standard redox potential often cited in the literature for the two electron oxidation of H₂S to sulfur is 0.144 V (which makes H₂S a weaker reducing agent than cysteine or glutathione), another standard redox potential could be found for the alkaline conditions, which suggests that H₂S is a stronger reducing agent:



H₂S does not react readily with oxygen; however, the solutions of H₂S undergo oxidation, just like the solutions of other thiols (such as cysteine and glutathione).

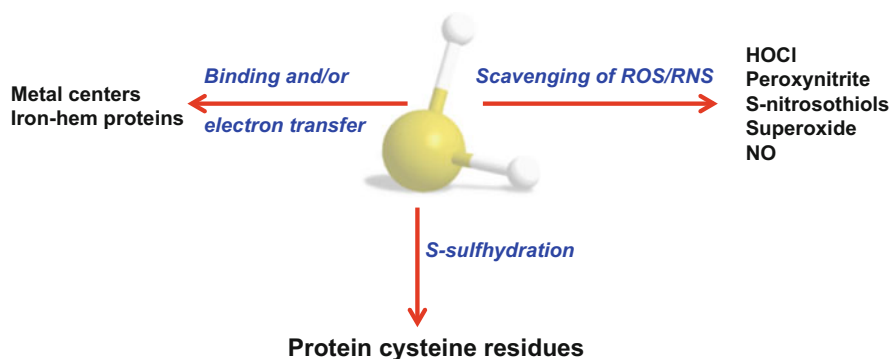


Fig. 1 Three main biochemical ways for direct H₂S signalling. Hydrogen sulfide can either react directly with reactive oxygen and nitrogen species (*arrow to the right side*) or react with metal centres by binding and/or electron transfer (*arrow to the left side*), or it can modulate cysteine residues of the proteins in a reaction called S-sulphydration (*central arrow*)

This process is believed to be catalysed by the traces of metal ions present in the solution (Kotronarou and Hoffmann 1991); therefore, the thorough cleaning of solutions from heavy metals and removal of oxygen can keep H₂S solution stable for a while (Wedmann et al. 2014). Oxidation products of H₂S in solution are polysulfides, sulfites, thiosulfites and eventually elemental sulfur.

In biological systems, reactivity of H₂S could be divided in three groups of reactions: (1) reaction with/scavenging of reactive oxygen (ROS) and reactive nitrogen species (RNS), (2) binding to and/or subsequent redox reactions with metal centres and (3) reaction with proteins, herein called persulfidation (alternatively S-sulphydration) (Fig. 1).

1.1 Reactions with ROS and RNS

H₂S reacts readily with hypochloric acid (HClO), produced by neutrophils, leading to the formation of polysulfides (Nagy and Winterbourn 2010). H₂S also reacts with peroxynitrite (Carballal et al. 2011; Filipovic et al. 2012a), in a reaction that generates thionitrate (HSNO₂) isomer, which can decompose and serve as an NO donor (Filipovic et al. 2012a). In addition, H₂S can also scavenge superoxide (Wedmann et al. 2014). Although the rates of these reactions are higher than those found for cysteine or glutathione, they are not so much higher that can overcome the difference in concentration (particularly when compared to glutathione which is present in millimolar steady-state levels) suggesting that H₂S cannot really serve as an antioxidant. Nonetheless, H₂S has been shown to have antioxidant and immediate protective effects in the cells exposed to ROS and RNS, something that cannot simply be explained by the modulation of proteins and gene expression. Since soluble macromolecules occupy a significant fraction of the total cell volume (Fulton 1982), within such crowded medium relative size and

shape of a molecule and probability of its successful diffusion, placement and effective contact with a potential target become crucial factors that significantly alter its reaction rates (Minton 1998). This speaks in favour of hydrogen sulfide, when compared to GSH, despite the difference in steady-state concentrations (Filipovic et al. 2012a).

H₂S reacts with NO signalling pathways as well (Whiteman et al. 2006; Ali et al. 2006; Yong et al. 2010, 2011; Filipovic et al. 2012b, 2013; Eberhardt et al. 2014). With protein S-nitrosothiols, H₂S reacts to form the smallest S-nitrosothiols, thionitrous acid (HSNO), which can freely diffuse through the membrane and serve as *trans*-nitrosating agent (Filipovic et al. 2012b). More importantly, H₂S can react directly with NO giving nitroxyl (HNO), the one-electron-reduced sibling of NO which possesses the signalling properties of its own (Eberhardt et al. 2014). 2 μM combination of NO and H₂S gives the same rate of HNO generation as ~1 mM Angeli's salt, the commonly used pharmacological source of nitroxyl. Eberhardt et al. (2014) recently showed that co-localization of H₂S and NO production facilitate intracellular generation of HNO which then activates transient receptor potential A1 channel (TRPA1) leading to the Ca²⁺ influx and to the release of calcitonin gene-related peptide (CGRP), the strongest known vasodilator.

1.2 Reactions with Metal Centres

One of the main biological targets for H₂S would be metal centres. H₂S can coordinate and then additionally reduce the metal centre. In haemproteins, polar active site favours the reduction, while nonpolar centres favour coordination only (Pietri et al. 2009). Indeed, the presence of positive charges around the haem centres leads to a complete change in the haems' reactivity, allowing the design of efficient catalysts for H₂S removal (Ivanovic and Filipovic 2012).

H₂S binds to both haem *a*₃ and Cu_B centres of cytochrome c oxidase, with K_i being 0.2 μM for the purified enzyme (Hill et al. 1984; Nicholls et al. 2013). This suggests that cytochrome c oxidase should be permanently inhibited under physiological conditions (considering the micromolar steady-state concentrations of H₂S), which is obviously not the case. In fact H₂S shows biphasic effects on whole cell respiration, stimulating it at low doses and then completely inhibiting it at higher concentrations (Koenitzer et al. 2007). Binding of H₂S to haem has an important function in mollusc, *Lucina pectinata*, where H₂S binds to haemoglobin I and is then transported to the symbiotic chemoautotrophic bacteria living in their gills (Ríos-González et al. 2014).

Reaction of oxyhaemoglobin with H₂S leads to the green pigment, sulphaemoglobin, which is in fact a chemically modified porphyrin centre. This subsequently affects oxygen-binding capacity of that haemoglobin. H₂S also binds and reduces cytochrome *c*, a reaction that in the presence of oxygen leads to superoxide production (Wedmann et al. 2014). In addition, Pálinkás et al. (2014) have recently investigated interactions of H₂S with human myeloperoxidase

(MPO), a major contributor to inflammatory oxidative stress, to show that H₂S inhibits the enzyme by reducing iron centre and by binding to the reduced Fe²⁺.

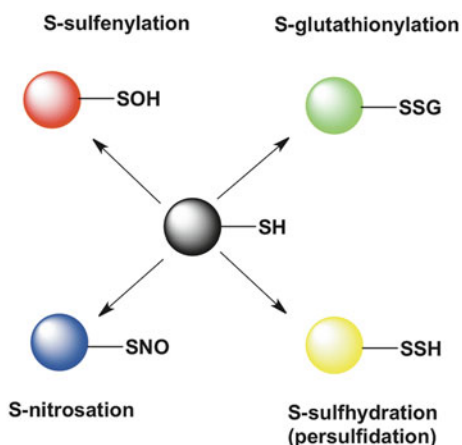
It is still, however, unclear to which extent is the coordination and/or reduction of metal centres involved in signalling by H₂S. Miljkovic et al. (2013) demonstrated that metal centres in mitochondria are responsible for the H₂S-stimulated haem centre-catalysed reduction of nitrite, a reaction which can explain the use of nitrite as an antidote for acute H₂S poisoning.

1.3 Protein Persulfides

The third way of direct H₂S signalling would be modulation of proteins by modification which is named S-sulfhydration, although the more correct term should be persulfidation. Protein persulfides add up to the list of oxidative posttranslational modifications (oxPTMs) of cysteine, such as S-nitrosation, S-sulfenylation and S-glutathionylation (Fig. 2). Modification of proteins by H₂S could explain the plethora of effects that H₂S exhibit and several proteins have been identified to be indeed controlled by this modification (Mustafa et al. 2009a, b, 2011; Paul and Snyder 2012). However, this field of research is at its beginning, and it is still facing difficulties/challenges such as the proper choice of detection method, understanding of the mechanism(s) by which persulfidation takes place and the actual impact it has on the cellular functions.

Based on the calculation of the bond energies of GSSG and GSSH, the latter has ~18 kJ mol⁻¹ lower bond energy (Filipovic et al. 2012b) so due to their inherent instability, there is limited information about persulfide reactivity to date. Francone et al. (2011) were among the first who reported preparation of the glutathione and papain persulfides, while Pan and Carroll (2013) successfully prepared persulfide on glutathione peroxidase 3. Zhang et al. (2014) also reported

Fig. 2 Oxidative posttranslational modifications of protein cysteine residues known to regulate protein structure/function



facile preparation of persulfide of bovine serum albumin (BSA), which can be used as a model to study protein persulfide reactivity.

Electronegativity of sulfur is almost identical to that of carbon; therefore in R-S-SH, the sulfur atom covalently bound to carbon could be considered as sulfane sulfur. Sulfane sulfur is sulfur with six valence electrons and a formal charge 0, often represented by S⁰. This sulfane sulfur could be a good target for nucleophilic attack, while the other sulfur atom, which is formally -1, could react with electrophiles. The pK_a of persulfides is lower than that of corresponding thiols, suggesting that at physiological conditions, majority of persulfide would be in deprotonated form (R-S-S⁻), making the persulfide “super” nucleophilic. This dramatically increases persulfides’ reactivity when compared to the corresponding thiols, as recently demonstrated for the reaction with H₂O₂ (Ida et al. 2014) and previously for the reduction of cytochrome c³⁺ (Francoleon et al. 2011).

Artaud and Galardon (2014) synthesized a persulfide analogue of the nitrosothiol SNAP, which opens up new possibilities for examining metastable low molecular weight (LMW) persulfides. The authors clearly demonstrate that spontaneous decay of LMW persulfides does not lead to H₂S release, but when mixed with other thiols, such as glutathione, an immediate H₂S release could be observed. Additionally, Bailey et al. (2014) characterized tritylhydrosulfide (TrtSSH), another LMW persulfide, to show that protonated form does not react with nucleophiles, while it readily reacts with electrophiles and reducing agents. Protonation leads to no change in substance stability, while the deprotonation stimulates decomposition with the elimination of elemental sulfur.

Based on the current literature data, the following characteristics of persulfides could be described:

1. Persulfides are metastable species which decompose in solution in a complex manner, leading to the formation of elemental sulfur, among other molecules.
2. Persulfides are much better nucleophiles than corresponding thiols which can explain their better reactivity.
3. Persulfides are better reducing agents than corresponding thiols.
4. Persulfides readily react with electrophiles.
5. Persulfides could transfer sulfane sulfur to other thiols, leading to *trans*-persulfidation (vide infra).

2 Persulfide Formation

Although there is a growing interest for persulfidation of proteins, only a few studies actually addressed the issue of the mechanism(s) behind protein S-sulfhydration. The original misconception (which tends to overtake this whole field) was that thiolate on the protein can react directly with H₂S to form protein persulfide (Fig. 3a). However, that reaction is impossible due to the thermodynamic constrains. Both sulfur atoms get oxidized in the reaction so the electrons will have to end up on protons, leading to elimination of hydrogen as a gas. Incubation of

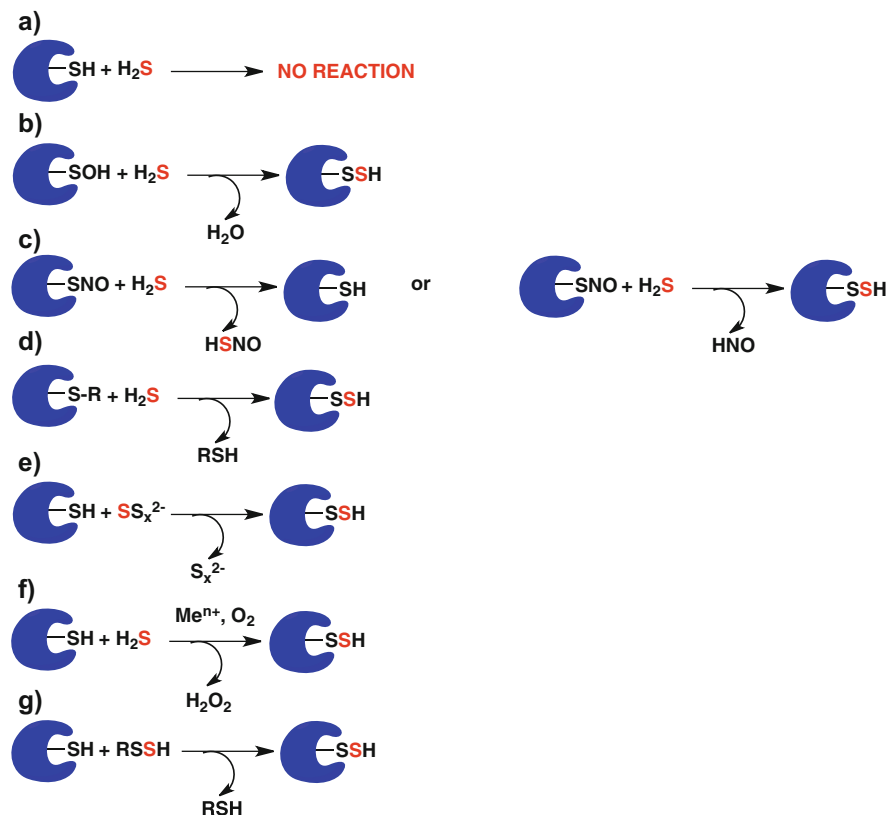


Fig. 3 Proposed reaction mechanisms for persulfide formation. (a) A direct reaction between protein thiols and H_2S is not possible, but H_2S can react with sulfenic acids (b). (c) Reaction of *S*-nitrosated cysteines with H_2S leads to the formation of HSNO, but depending on the protein environment surrounding the thiol, it is also possible to generate protein persulfides in this reaction. (d) H_2S could react with already existing inter- or intramolecular disulfides, while sulfane sulfur in polysulfides could react directly with protein thiols and give persulfide (e). (f) Metal centres could act as oxidants for the formation of protein persulfides from H_2S and protein thiols. (g) Persulfides could serve as carriers of sulfane sulfur and engage in “*trans-S*-sulfhydration” reaction

proteins, such as GAPDH, BSA or immunoglobulins with H_2S , led to no detectable protein *S*-sulfhydration (Zhang et al. 2014; Wedmann et al. 2014) confirming the theory.

So how are the proteins modified by H_2S ? Getting the answer(s) to this question is of the utmost importance for our understanding of H_2S signalling and also for the interpretation of the vast amount of data accumulated to date.

2.1 Enzymatic Generation of Persulfides

Despite the recent interest for protein persulfidation, sparked by the discovery of H₂S as a signalling molecule, protein persulfides formed as intermediates that facilitate sulfur delivery in several biosynthetic pathways have been known for a while (Mueller 2006). Persulfides are found to be formed in sulfurtransferases and cysteine desulfurases. Particularly interesting examples are the two enzymes involved in H₂S production and its oxidation, mercaptopyruvate sulfurtransferase (MST) and sulfide–quinone oxidoreductase (SQR), respectively.

MST is expressed in both mitochondria and cytoplasm and could be found in kidney cells, liver and cardiac cells, neuroglial cells, etc. (Kabil and Banerjee 2014; Kimura 2014). Recent studies suggest that MST is an important source of H₂S in some organisms and tissues (Mikami et al. 2011; Módis et al. 2013). Although it was known for a while that during the reaction protein persulfide is formed in the catalytic site (MST-Cys-S-SH), the mechanism of H₂S release has been only recently discovered. Yadav et al. (2013) reported the first crystal structure of MST with its cysteine residue in the form of persulfide. This allowed them to propose a detailed reaction mechanism, while kinetic analysis of the reaction led to conclusion that thioredoxin is likely to be the major physiological persulfide acceptor for MST (Fig. 4a).

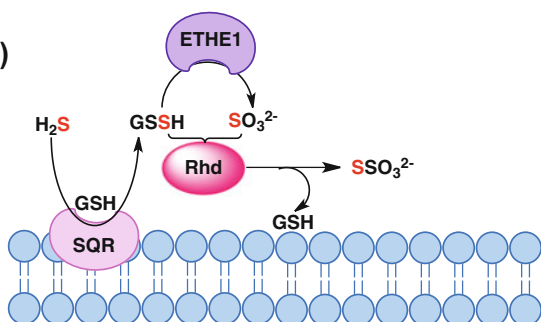
The same group, led by Ruma Banerjee, elucidated the full mechanism of SQR (Libiad et al. 2014). The sulfide oxidation pathway begins with SQR, and it also includes a sulfur dioxygenase, rhodanese and sulfite oxidase (Fig. 4b). By consuming H₂S and its persulfide products, SQR and sulfur dioxygenase are important switch-off regulators of sulfide signalling (Bouillaud and Blachier 2011). In the first step, SQR catalyses the oxidation of H₂S to sulfane sulfur, which remains covalently attached to the enzyme. In the second step, this sulfane sulfur could be transferred to sulfite, to form thiosulfate (Jackson et al. 2012), but recently, Libiad et al. (2014) showed that in fact glutathione is a more probable acceptor of sulfane sulfur. This leads to the generation of glutathione persulfide, which can be consumed by rhodanese to actually form thiosulfate rather than to use it as a substrate (Fig. 4b).

Recent work on reactive cysteine persulfides and S-polythiolation has shaken our understanding of the action of CBS and CSE (Ida et al. 2014). Namely, the authors demonstrated that persulfide formation by CSE and CBS-mediated CysSSCys metabolism are facile and more likely a source of biological persulfides (Fig. 4c). Therefore, the direct enzymatic production of per- and polysulfide that is highly prevalent has been suggested. In fact, the authors question the role of H₂S as a major signalling molecule suggesting that Cys-based persulfides may be the actual signalling species (Ono et al. 2014).

a)



b)



c)

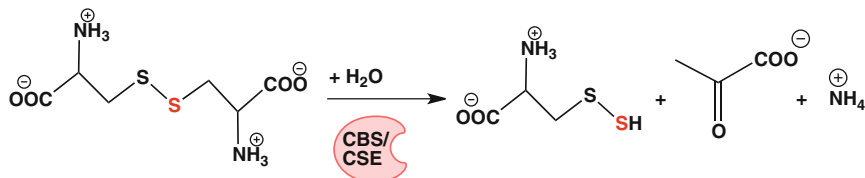


Fig. 4 Enzymatic production of protein persulfides. (a) Generation of protein persulfide on MST enzyme during its catalytic cycle. (b) Generation of glutathione persulfide during the enzymatic oxidation of H_2S . In the first step, SQR catalyses the oxidation of H_2S to sulfane sulfur with glutathione as probable acceptor of this sulfane sulfur. This leads to generation of glutathione persulfide, which can be consumed by rhodanese to actually form thiosulfate. (c) Cysteine persulfides are formed in the enzymatic cleavage of cystine by CBS or CSE

2.2 Direct Nonenzymatic Generation of Persulfides by H_2S

An obvious similarity between protein S-sulfhydration and S-glutathionylation could be drawn. Based on known mechanisms for the generation of glutathionylated proteins, it is possible to assume that the same reactions could lead to generation of protein persulfides.

2.2.1 Reaction with Sulfenic Acids

Protein S-sulfenylation (Fig. 2) is the reversible oxidation of protein thiols to sulfenic acids (R-SOH) that has been recognized as an important oxidative post-translational modification of cysteines (Paulsen and Carroll 2013; Gupta and Carroll 2014). Most recently, an extensive study was published identifying the

whole cell sulfenylome, i.e. site-specific mapping and quantification of protein S-sulfenylation in the cells (Yang et al. 2014). More than 1000 proteins have been characterized as modified by sulfenylation. Peroxiredoxins are the most abundant antioxidant enzymes in the cytosol, and their catalytic cycle relies on sulfenic acid formation (Poynton and Hampton 2014). Sulfenic acid formation is shown to be involved in H₂O₂-mediated inactivation of protein tyrosine phosphatases (PTPs) (Paulsen et al. 2011). Also, reversible sulfenylation is shown to switch on or off the activity of transcription factors such as OxyR, OhrR or Orp1-Yap1 (for extensive review, see Paulsen and Carroll 2012). Epidermal growth factor receptor and the phosphatases SHP2, PTEN and PTP1B were all found to be sulfenylated upon regular signalling by epidermal growth factor (Paulsen et al. 2011).

The biochemistry of sulfenic acid formation, reactivity and functions has been extensively reviewed recently (Paulsen and Carroll 2013; Gupta and Carroll 2014). Sulfenic acids are known to react with thiols to form disulfides. The same could be said for the reaction of sulfenic acids with H₂S (Fig. 3b). Namely, in the recent study, Zhang et al. (2014) showed that formation of sulfenic acids on GAPDH and subsequent reaction with H₂S do indeed lead to the formation of protein persulfides. Furthermore, they used sulfenic acid of bovine serum albumin (BSA), known as an example of a relatively stable protein sulfenic acid (Carballal et al. 2003), to successfully generate S-sulfhydrated BSA. Although the steady-state concentration of H₂S is orders of magnitude lower than that of glutathione (Olson et al. 2014; Kabil et al. 2014), which makes it difficult for H₂S to compete for the reaction with protein sulfenic acids, the high flux of H₂S generation (Vitvitsky et al. 2012), its free diffusion (Mathai et al. 2009; Cuevasanta et al. 2012) and therefore its ability to reach deeper parts of the proteins suggest that the reaction of H₂S with sulfenic acids could still be a major source of protein persulfide formation (Fig. 3b). Indeed, Zhang et al. (2014) find that intracellular persulfidation co-localizes with endoplasmic reticulum, an organelle rich with sulfenic acids.

It is worth noting that sulfenic acids can be formed from the thiols just by the presence of traces of metal ions in the buffers and oxygen (Paulsen and Carroll 2013). Considering that R-SOH could react with H₂S to form protein persulfides, these reaction steps could be an explanation for the misconception that H₂S reacts directly with thiols.

2.2.2 Reaction with S-Nitrosothiols

Protein S-nitrosation is considered by some to be the second most important posttranslational modification of proteins. The number of proteins found to be controlled by this modification is constantly increasing. To date, S-nitrosation has been implicated in the regulation of proteins involved in muscle contractility, neuronal transmission, host defence, cell trafficking, apoptosis, etc. (Hess and Stamler 2012; Seth and Stamler 2011; Foster et al. 2009; Lima et al. 2010). S-nitrosation of haemoglobin has been proposed to regulate its ability to release oxygen (Reynolds et al. 2013). S-nitrosation of ryanodine receptors regulates intracellular Ca²⁺ levels (Xu et al. 1998) as does S-nitrosation of transient receptor potential cation channels (TRPs) (Yoshida et al. 2006). The role of S-nitrosation in

controlling the protein function has been extensively reviewed elsewhere (Hess and Stamler 2012; Seth and Stamler 2011).

The mechanism of protein S-nitrosation is still the matter of debate (Broniowska and Hogg 2012). Like in the case of S-sulfhydration, direct reaction of NO with thiols to lead to the S-nitrosothiol formation is thermodynamically unfavourable. The mechanism usually involves either the reaction of thiols with higher nitrogen oxides, such as N_2O_3 , or reactions with metal centre-catalysed one-electron oxidation product of NO (Broniowska and Hogg 2012). *Trans*-nitrosation, transfer of NO^+ moiety, from one protein to another is also a subject of a lively debate. Cysteine and glutathione have been proposed as common carriers. However, H_2S also reacts with S-nitrosothiols leading to the formation of the smallest S-nitrosothiol, HSNO (Fig. 3c, Filipovic et al. 2012b). HSNO can freely diffuse through the membranes serving as a carrier of NO^+ and conveying further trans-nitrosation. In addition, HSNO is very unstable and highly reactive so it could react further with H_2S -forming polysulfides and/or sulfur, which could be a source of protein S-sulfhydration (Fig. 3c). The reaction of S-nitrosothiols with H_2S to directly give persulfides and nitroxyl (HNO) is thermodynamically unfavourable ($\Delta_{rxn1}G^\circ \approx +40 \text{ kJ mol}^{-1}$) (Filipovic et al. 2012b). Although it seems that *trans*-nitrosation is a generally favoured pathway, a recent computational study pointed out that the surrounding of S–NO bond within a protein could significantly affect the thermodynamic feasibility of the thiolation reaction, making it possible for certain proteins (depending on the surrounding of the cysteine residue) to get directly S-sulfhydrated (Fig. 3c, Talipov and Timerghazin 2013).

2.2.3 Reaction with Disulfides

Reaction of thiols with disulfides normally proceeds until the equilibrium is established (Moriarty-Craige and Jones 2004). Following the same logic, H_2S should be able to react with disulfides leading to persulfide formation (Fig. 3d). Protein disulfides are always formed between cysteine residues (of the same or different polypeptide chain), or they appear as mixed disulfides between the cysteine residues of the protein and glutathione (in S-glutathionylated proteins). Based on the calculation of the bond energies of GSSG and GSSH, the latter has $\sim 18 \text{ kJ mol}^{-1}$ lower bond energy (Filipovic et al. 2012b) which makes the reaction of H_2S with oxidized thiols very slow. Francoleon and colleagues demonstrated that glutathione persulfide could indeed be formed in a reaction mixture containing oxidized glutathione and H_2S , but the product was unstable (Francoleon et al. 2011). Formation of protein persulfides by the H_2S -induced reduction of intra- or intermolecular protein disulfides seems highly unlikely in the cells, due to the low levels of H_2S when compared to other thiols (such as glutathione or cysteine) and the very slow reaction rate. Zhang et al. (2014) did not observe any S-sulfhydration as a consequence of the reaction of H_2S with disulfides of BSA, and a study by Wedmann et al. (2014) confirmed this observation on immunoglobulins purified from human blood. The use of very high, non-physiological concentrations of H_2S or its donors in experimental setups could, nonetheless, overcome this limitation and lead to protein modification.

2.3 Nonenzymatic S-sulfhydrylation by Species Originating from H₂S

2.3.1 Reaction with Polysulfides

Polysulfides (HS_x⁻) are the products of incomplete H₂S oxidation on its way to total oxidation to elemental sulfur. Contrary to a widely spread opinion in a biological community that H₂S easily oxidizes, H₂S is known as a very persistent and difficult-to-get-rid-off pollutant in industrial processes. The oxidation that indeed does occur on air is most probably caused by the impurities present in the solutions such as traces of metal ions, which are inevitable contaminants of all standard buffer solutions (Kotronarou and Hoffmann 1991).

The length of polysulfide chain can be from 2 to 7. It is worth noting that older chemical literature clearly points out that H₂S₂ is a molecule that can be prepared under extreme conditions (Parsons and Walton 1921). It is a substance of pungent smell that cannot even stand the humidity from air, and it immediately decomposes to give H₂S and elemental sulfur (Parsons and Walton 1921). Therefore, assuming that H₂S₂ is an important player in the biological system would be wrong. If formed, H₂S₂ would immediately decompose.

Longer polysulfides are more stable, with $x = 4$ and/or 5 being the most abundant species. They are formed even in biological milieu and could be responsible for the part of the effects assigned to H₂S (Fig. 3e). Indeed, Kimura et al. (2013) demonstrated that polysulfides have a 300 times higher potency in activating TRP channels. In addition, Greiner et al. (2013) linked polysulfides to protein thiol oxidation suggesting that all sources of H₂S (salts and H₂S donors) inevitably contain polysulfides, which in turn oxidize cysteine residues leading to persulfide formation. All inorganic polysulfides with more than two sulfur atoms contain sulfane sulfur atoms, which could undergo nucleophilic attack by free protein thiols leading to the formation of protein persulfides. However, polysulfides could also serve as reducing agents. Francoleon et al. (2011) suggested that persulfides are much more potent reducing agents than H₂S, while Wedmann et al. (2014) recently showed that polysulfides are capable of completely cleaving intramolecular disulfides present in immunoglobulins. H₂S solutions prepared with care, however, could not do the same. Only when mixed with traces of metal ions, they achieved the same effect (Wedmann et al. 2014).

Although there is very little doubt that polysulfides are formed in the cells, it is still unclear how they could serve as signalling molecules. Signalling molecules should be able to achieve the effect at low doses, but at the same time, their production should be tightly controlled, and their reactivity should be specific, just like in the case of NO or H₂O₂. Formation of polysulfides cannot be controlled; it is guided by stochasticity as the reactions that polysulfide undergo are highly unpredictable and will largely depend on the availability of oxygen and metal centres on the one hand and the protein thiols/disulfides on the other hand. In addition, polysulfides are charged, and it is almost impossible to achieve their fully protonated form under physiological condition, making the diffusion through the membrane impossible without a facilitator.

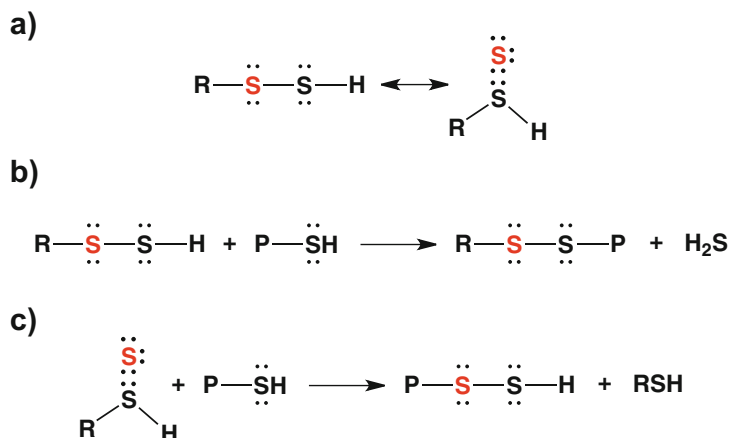
2.3.2 Reaction with Metal Centres and Generation of HS•

Although not studied into too much detail, metal centres could play an important role in catalysing formation of protein persulfides (Fig. 3f). Namely, some iron haem centres are able to oxidize hydrogen sulfide forming HS• (Miljkovic et al. 2013), which could in turn react with free thiols to finally generate protein persulfides (Zhang et al. 2014). Cytochrome c, for example, readily reacts with H₂S (Wedmann et al. 2014). Iron porphyrins in general could be a good source of HS•. HS• reacts further with H₂S, in a diffusion-controlled reaction, to give H₂S₂•⁻ (Das et al. 1999) which should be a powerful persulfidation agent when formed intracellularly, something that is yet to be confirmed. As demonstrated by Zhang et al. (2014), incubation of GAPDH with iron porphyrin and H₂S leads to the strongest generation of protein persulfides. This complemented with their observation that a large portion of intracellular S-sulphydration co-localizes with mitochondria, the organelle richest in metallo-proteins.

2.3.3 “Trans-persulfidation” by Polythiolated Cysteine or Glutathione

A recent study by Ida et al. (2014) suggested very high levels of circulatory LMW persulfides. Francoleon et al. (2011) noticed that treatment of papain with GSSH leads to the enzyme inhibition in the same manner as it does the persulfidation of this enzyme. The authors recently went a step further suggesting that cysteine persulfide and/or glutathione persulfide could be the main persulfidating agents in the cells (Ono et al. 2014). Cysteine persulfides could even be transported through the membrane. S-nitrosocysteine does the same and is considered as one of the main carriers of “NO⁺” moiety in *trans*-nitrosation reactions (Broniowska and Hogg 2012). Making a parallel with S-nitrosothiol chemistry, the term “*trans*-persulfidation” could be coined to describe this process (Fig. 3g).

The mechanism of this process is still unclear. As mentioned above, the persulfides would react with electrophiles rather than with nucleophiles (Artaud and Galardon 2014; Bailey et al. 2014), and when they do react with nucleophiles (such as other free thiols), this reaction leads to the release of H₂S (which represents the basis of one of the methods for persulfide detection: reduction with DTT). The reaction with nucleophiles could only go as an attack to sulfane sulfur, but that leads to the formation of a mixed disulfide and elimination of H₂S. Some literature data exist suggesting that persulfides could exist in the tautomeric thiosulfoxide form (Scheme 1), which would then act as a perfect donor of sulfane sulfur (Kutney and Turnbull 1982; Steudel et al. 1997). However, it is worth noting that although this may be true for the extreme case of F₂S₂, neither experimental nor computational data support the existence of this tautomeric form (Steudel et al. 1997). Therefore, the elucidation of this mechanism is to be done in some future studies.



Scheme 1 Possible mechanism of *trans*-persulfidation. (a) The tautomeric forms of protein or LMW persulfides. Sulfane sulfur is marked red. (b) Reaction of LMW persulfides with protein thiols leads to the H₂S elimination and formation of mixed disulfide. (c) Reaction of thiosulfoxide tautomer, however, could lead to *trans*-persulfidation

3 Detection of Protein Persulfides

Detection of protein *S*-sulphydration represents a certain challenge as the persulfide group exhibits the reactivity similar to other, free thiols (Flavin 1962; Heimer 1981; Mueller 2006; Pan and Carroll 2013; Zhang et al. 2014). Four distinctive approaches have been proposed (Fig. 5) for the persulfide detection, all of which are summarized here with a particular emphasis on the potential experimental problems that might occur.

The original method for protein persulfide detection was suggested by Mustafa et al. (2009b), and it was based on a premise that protein persulfides would not react with electrophilic thiol-blocking reagent *S*-methyl methanethiosulfonate (MMTS). In the subsequent step, persulfides were labelled with N-[6-(biotinamido)hexyl]-3'-(2'-pyridyldithio)propionamide (biotin-HPDP) (Fig. 5a). This method allowed the simultaneous labelling of *S*-sulphydration and *S*-nitrosation. Using this method, Mustafa and colleagues suggested that a large number of proteins were a target for H₂S signalling and that basal protein persulfidation is as high as 25 % (Mustafa et al. 2009b).

To date, this is the most used methodological approach in reporting protein *S*-sulphydration of different proteins. However, certain methodological limitations arise. Although MMTS has been widely used to study *S*-nitrosation (Forrester et al. 2009) and has been an efficient tool in trapping mixed disulfides *in vivo* (Peaper et al. 2005), Karala and Ruddock (2007) were able to show that *in vitro* MMTS treatment of both peptides and proteins resulted in the artificial formation of intramolecular and intermolecular protein disulfide bonds which could lead to

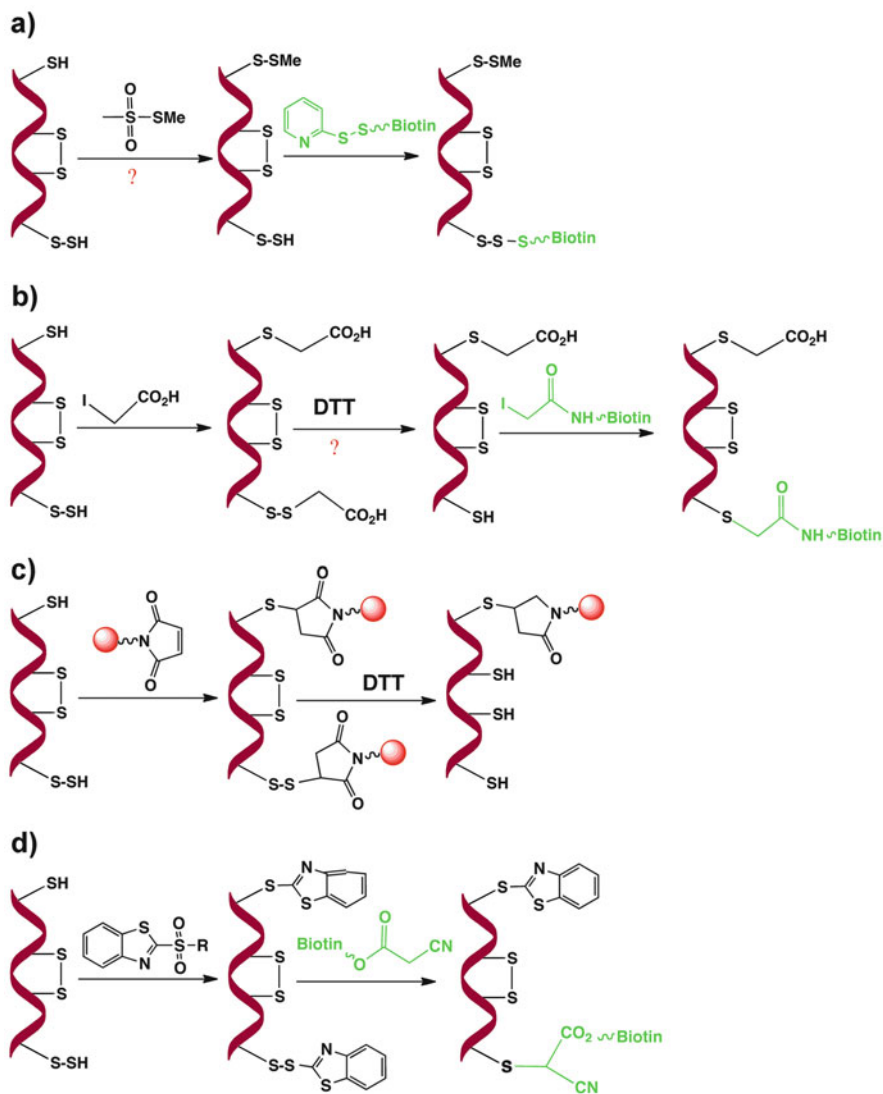


Fig. 5 An overview of the methods currently employed for the detection of protein persulfides. **(a)** The first method was based on a chemically wrong premise that protein persulfides would not react with electrophilic thiol-blocking reagent S-methyl methanethiosulfonate (MMTS). In the subsequent step, persulfides were labelled with N-[6-(biotinamido)hexyl]-3'-(2'-pyridyldithio) propionamide (biotin-HPDP). **(b)** In the second method, iodoacetic acid (IAA) is used to initially block both free thiols and protein persulfides. In the subsequent steps, alkylated persulfide is cleaved with DTT and then labelled with iodoacetamide-linked biotin (IAP). Although DTT would indeed cleave this adduct, it is unclear how this method distinguishes the persulfides from intra- and intermolecular disulfides and S-nitrosothiols, which would also be reduced by DTT. **(c)** In this method, both persulfide and free thiol would be blocked by the thiol fluorescently labelled N-ethyl maleimide (Cy5-conjugated maleimide). The adduct of persulfide and Cy5-maleimide is a disulfide that will be then cleaved by the DTT leading to a decrease of the in-gel fluorescence signal in the samples containing persulfides. **(d)** Finally, the use of the methylsulfonyl benzothiazole (MSBT) to block thiols and persulfides in the first step, followed by the tag switch with cyanoacetamide derivatives in the second step, leads to the efficient labelling of persulfides, by the method called tag-switch technique

general data misinterpretation. In addition, Pan and Carroll (2013) studied how persulfides react with both electrophilic and nucleophilic species, reaffirming the nucleophilic properties of the persulfide sulfane sulfur. They unambiguously showed that persulfides react with MMTS (and its analogue S-4-bromobenzyl methanethiosulfonate) as readily as free thiols, questioning the interpretation of the data obtained by modified biotin-switch technique (MBST) approach. Two possible models were proposed to explain the data generated by MBST: (1) free thiols may be incompletely blocked in the first MMTS alkylation step and subsequently react with the pyridyldisulfide biotin reagent; (2) alternatively or in addition, labelling may be achieved via stepwise thiol-disulfide exchange in a reaction catalysed by trace free thiols (RSH).

In their attempt to identify persulfidation of protein tyrosine phosphatase 1B (PTP1B), Krishnan et al. (2011) proposed an approach for persulfide detection based on a completely opposite chemical premise. Namely, they proposed that thiol-blocking reagent, iodoacetic acid (IAA), will react with both free thiols and protein persulfides (Fig. 5b), which is in agreement with the persulfides' reactivity similar to that of free thiols. In the subsequent steps, however, they proposed the cleavage of the alkylated persulfide with DTT and then labelling of that particular cysteine with iodoacetamide-linked biotin (IAP). Although DTT would indeed cleave this adduct, it is unclear how this method distinguishes the persulfides from intra- and intermolecular disulfides and S-nitrosothiols, which would also be reduced by DTT.

Sen et al. (2012) proposed an alternative method for the persulfide detection. This method is based on the fact that both persulfide and free thiol would be blocked by the thiol-blocking reagent N-ethyl maleimide. The authors used Cy5-conjugated maleimide in the first step followed by the used of DTT in the second (Fig. 5c). The adduct of persulfide and Cy5-maleimide is a disulfide that will be cleaved by the DTT leading to a decrease of the in-gel fluorescence signal in the samples containing persulfides. This method has its advantage of being relatively simple and available to every researcher (because the reagents are commercially available). The only limitation of the method is that it does not offer actual persulfide labelling which would allow wide proteomic analysis. Particular care should be taken to ensure that all thiols and persulfides are indeed blocked in the initial step, as that is the crucial step for the subsequent data interpretation.

Most recently, Zhang et al. (2014) proposed a different approach for protein persulfide detection, named tag-switch assay (Fig. 5d). This method was based on an idea that thiol-blocking reagent should be introduced in the first step, which would tag both free thiols and persulfides. If an appropriate tag is employed, the disulfide bond in persulfide adducts might show much enhanced reactivity to certain nucleophiles than common disulfides in proteins. Therefore, a tag-switching reagent (containing both the nucleophile and a reporting molecule such as biotin) could be introduced to label only the persulfide adducts. It should be noted that thiol adducts from the first step are thioethers, which are not expected to

react with the nucleophile. A major challenge in this technology was for the newly generated disulfide linkage from persulfide moieties to display a unique reactivity for a suitable nucleophile to an extent that distinguishes them from common disulfides. Using the methylsulfonyl benzothiazole (MSBT) to block thiols in the first step in combination with the tagged-cyanoacetate derivatives in the second step, persulfides could be efficiently labelled. Neither free thiols, intramolecular disulfides, S-glutathionylated or sulfenylated proteins were tagged by this approach (using BSA as a model protein). This method was further adapted for visualization of intracellular S-sulfhydration by fluorescence microscopy. Despite the lack of reactivity of BSA sulfenic acid derivative with the method, it is expected that sulfenic acids should react with cyanoacetate derivatives. Protein sulfenic acids are, in general, very unstable and prone to further oxidation, so it is improbable that any of them will remain in the solution after 30–45 min of initial incubation with the MSBT. Nonetheless, it is better to treat the cell lysates with dimedone, prior to the incubation with the first component of tag-switch assay, in order to eliminate any doubts of potential cross-reactivity (Park et al. 2015).

3.1 Detection of Sulfane Sulfurs

Although the methods for sulfane sulfur detection detect more than just protein persulfides, they could serve as a useful tool for the fast and easy proof-of-concept type of experiments. With recent advances in making these methods quantitative, it is also possible to assume that the results suggesting an increase or decrease of the sulfane sulfur levels do suggest the same trends for protein persulfides.

The traditional method for sulfane sulfur detection is cyanolysis (Wood 1987). The method is based on the reaction between sulfane sulfurs and cyanide under alkaline conditions ($\text{pH} > 8.5$). In this reaction, thiocyanate (SCN^-) is formed, which in the reaction with Fe^{3+} generates the $\text{Fe}(\text{SCN})^{2+}$ complex with characteristic absorbance maximum at 460 nm.

Recently, two new methods have been proposed. The first one is designed to be used for the detection of sulfane sulfur levels in the cells. Chen et al. (2013) based their discovery on an assumption that sulfane sulfurs are likely to react with the nucleophile components of a fluorescent probe, which could then undergo spontaneous and fast cyclization to release fluorophore. They successfully produced two probes, named SSP1 and SSP2, which could be easily used for intracellular detection of sulfane sulfur. The same group improved the probes designing a new DSP-3 fluorescence sensor for hydrogen polysulfides (Liu et al. 2014a).

Most recently, the group led by Ming Xian established isotope dilution mass spectrometric method for the quantification of sulfane sulfurs (Liu et al. 2014b). Accurate and reliable measurements of sulfane sulfurs in biological samples are required in order to understand the impact of H_2S signalling. The method that Liu et al. (2014b) proposed employs a triphenylphosphine derivative to capture sulfane

sulfurs as a stable phosphine sulfide product. The concentration of this product can be determined by isotope dilution mass spectrometry using a ¹³C₃-labelled phosphine sulfide as an internal standard. Using this method, the authors found that average concentrations of sulfane sulfur were 57.0 (liver), 150.9 (kidney), 46.0 (brain), 61.8 (heart), 56.1 (spleen) and 20.8 nmol/g (lungs).

4 Persulfidation in Action

As already mentioned, H₂S regulates the plethora of biological functions from neurotransmission and blood pressure to cardioprotection. To date, several key protein targets have been identified which do undergo oxPTM of the cysteine residue, suggesting that this modification can be responsible for some of the H₂S effects such as vasodilation, prevention of cell death and senescence, cell differentiation, etc.

4.1 S-sulfhydration of K_{ATP} Regulates Vasodilation

Seminal work by Yang et al. (2008) showed that CSE knockout mice develop hypertension, confirming previous assumptions about H₂S being the endogenous regulator of blood pressure. Several other studies have been published recently showing that majority of H₂S-induced vasodilation goes via its interaction with NO. This effect can be dual, via direct reaction with NO that leads to the nitroxyl (HNO) formation and subsequent activation of HNO–TRPA1–CGRP signalling cascade (Eberhardt et al. 2014) or by inhibiting the phosphodiesterase 5 and increasing the cGMP levels (Colleta et al. 2012). Nonetheless, the part of H₂S-induced vasodilation could be assigned to the activation of K_{ATP} channels. Persulfidation of C34 on the Kir6.1 subunit of K_{ATP} channel on smooth muscle cells prevents its association with ATP and promotes its binding to phosphatidylinositol-4,5-bisphosphate (PIP₂). This leads to the channel opening and K⁺ influx and subsequently to the smooth muscle cells' relaxation. Several other Ca²⁺ channels have been also implicated in the H₂S-induced vasodilation.

4.2 Persulfidation of Electrophilic Messengers and Cardioprotection

Recently, direct persulfidation of several electrophilic messengers (such as 8-nitro-cGMP) by sulfide has been reported offering additional mechanism for H₂S-mediated signalling (Nishida et al. 2012). Redox signalling by electrophilic by-products, such as nitrated cyclic nucleotides and nitro- or keto-derivatives of unsaturated fatty acids, all generated by the inflammation-related enzymes, ROS and/or NO, has attracted much interest lately (Nishida et al. 2014; Fujii and Akaike 2013). S-alkylation of cysteine residues by 8-nitroguanosine 3',5'-cyclic monophosphate mediates several redox signalling pathways. Nishida et al. (2012)

showed that HS^- could directly attack the above-mentioned electrophiles, forming derivatives that have biological effect of themselves. Namely, formation of 8-SH-cGMP blocks the S-guanylation of H-Ras, a modification which normally activates H-Ras to signal cell senescence as a response to stress (Fujii and Akaike 2013). The authors link the protective cardiovascular effects of H_2S and its donors to increased formation of 8-SH-cGMP and inhibition of H-Ras signalling. It is worth mentioning, however, that Terzić et al. (2014) showed that nucleophilic attack of the hydrosulfide anion to 8-nitro-cGMP cannot take place, as previously proposed. Instead, the formation of reactive species containing sulfane sulfur, like persulfides, is required (Terzić et al. 2014).

4.3 S-sulfhydration of GAPDH and NF- κ B Protects Against Apoptosis

GAPDH was the first protein characterized as S-sulfhydrated in the study that sparked the whole research field (Mustafa et al. 2009a). GAPDH has been known for a while as regulator of a cell death cascade (Hara et al. 2005). S-nitrosation of catalytic C152 abolishes its catalytic activity but makes it able to bind to Siah1, an E3 ubiquitin ligase. Siah1, which possesses a nuclear localization tag, helps the translocation of GAPDH to a nucleus where it enables Siah1 to degrade nuclear proteins, leading to cell death (Hara et al. 2005). Mustafa et al. showed that GAPDH is endogenously S-sulfhydrated at C150 which increases its enzymatic activity severalfold (Mustafa et al. 2009a). The authors also demonstrated that DTT treatment of GAPDH decreases its activity, suggesting that endogenous persulfidation regulates its function. CSE knockout mice showed ~35 % reduced activity of GAPDH when compared to control mice. Zhang and associates confirmed recently, using selective tag-switch assay, that GAPDH is indeed endogenously S-sulfhydrated (Zhang et al. 2014). The fact that S-sulfhydration of GAPDH increases its enzymatic activity means that it also prevents its interaction with Siah1 protecting the cells from apoptosis.

Nuclear factor- κ B (NF- κ B) is an anti-apoptotic transcription factor, which is under basal conditions kept in cytosol via interaction with the inhibitor- κ B α (Napetschnig and Wu 2013). During the inflammation, cells produce tumour necrosis factor- α (TNF- α), which could lead to cell death (Aggarwal et al. 2012). H_2S is known to have protective effects in inflammation but without distinctive mechanism that can explain it (Li et al. 2005; Whiteman and Winyard 2011). Sen and colleagues offered such mechanism in a study describing the persulfidation of NF- κ B (Sen et al. 2012). Namely, they showed that C38 of p65 subunit of NF- κ B is persulfidated which promotes its binding to the co-activator ribosomal protein S3, augmenting its binding to the promoters of anti-apoptotic genes (Fig. 6a). In addition, TNF- α stimulated transcription of CSE, increasing the total amount of S-sulfhydrated NF- κ B. Conversely, Du et al. 2014 showed recently that hydrogen sulfide suppresses oxidized low-density lipoprotein-induced macrophage inflammation by inhibiting NF- κ B. This study suggests that persulfidation of C38 on p65

in fact prevents the NF- κ B from leaving the cytosol, therefore completely inhibiting its DNA binding activity. Further studies are obviously needed to give more conclusive answer about the role of H₂S in inflammation.

4.4 S-sulfhydration of Parkin Protects Against Parkinson's Disease

Parkinson's disease (PD) is a neurodegenerative disease caused by the death of dopamine-generating cells in the substantia nigra (Shulman et al. 2011). One of the proteins considered responsible for this cell death is parkin. Parkin is an E3 ubiquitin ligase that catalyses ubiquitination of diverse substrates. Mutations in parkin, which lead to the loss of its activity, are one of the causes of PD (Shulman et al. 2011; Moore et al. 2005). Parkin has reactive cysteine residues, which can be subjected to oxidative posttranslational modifications. S-nitrosation of parkin, for example, inhibits its E3 ubiquitin ligase activity contributing to the Parkinson's disease (Chung et al. 2004).

Recently, Solomon Snyder and colleagues demonstrated that cysteines C59, C95 and C182 could undergo S-sulfhydration (Vandiver et al. 2013). This leads to the increase of parkin's activity (Fig. 6b). Persulfidation of parkin is markedly decreased in PD brains, whereas S-nitrosation is increased. Development of H₂S donors opens up a possibility of their use in the early treatment of PD. Increase of parkin's activity could salvage the neurons from the cell death by removing damaged proteins. Parkin is also an important regulator of mitophagy, leading to the removal of damaged mitochondria, particularly in ischemia–reperfusion injury. As H₂S is known to have great pharmacological potential in preventing ischemia–reperfusion injury, it is possible that part of this effect goes via persulfidation of parkin and increased removal of damaged mitochondria.

4.5 S-sulfhydration of PTP-1B Regulates ER Stress

Protein tyrosine phosphatases (PTPs) in conjunction with protein tyrosine kinases are important controllers of various biological functions. PTPs are particularly sensitive to oxidative posttranslational modifications of cysteine, as cysteine is present, is in the active site and is important for their function (van Montfort et al. 2003). H₂O₂ is known to modulate function of these enzymes by forming an inactive adduct which contains sulfenic acid in the active site (van Montfort et al. 2003). PTP-1B is one of the members of this class of enzymes, located in the cytoplasmic face of the endoplasmic reticulum, where it plays an important role in ER stress signalling (Paulsen et al. 2012).

Persulfidation of C215 leads to the loss of enzymatic activity, which increases the phosphorylation of Y619 and therefore the activation of PERK in response to ER stress (Krishnan et al. 2011). PERK activation leads to global inhibition of protein translation. The mechanism by which H₂S causes persulfidation of PTP-1B,

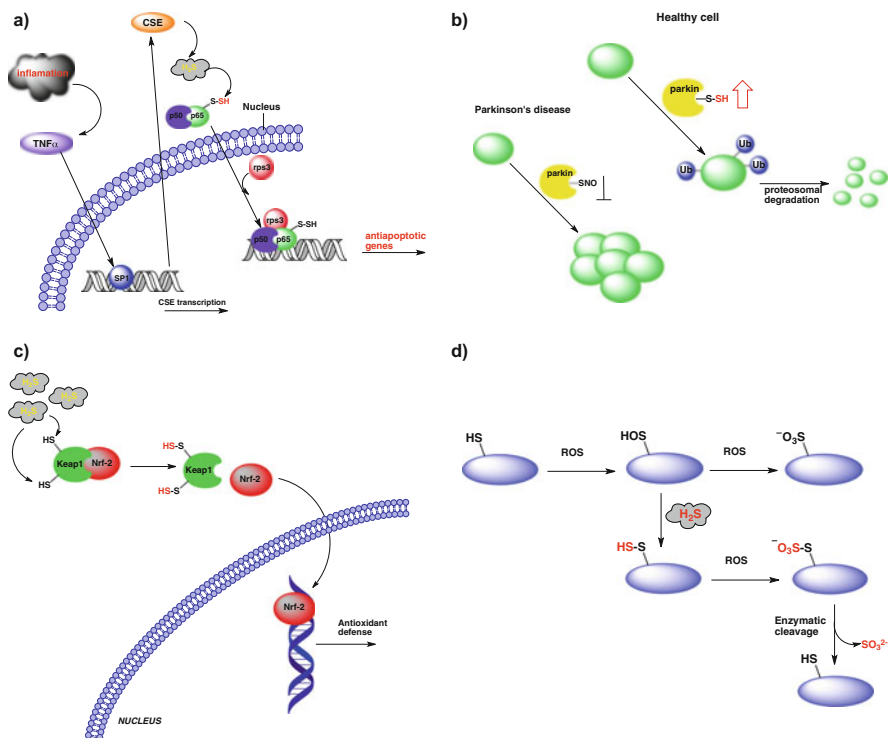


Fig. 6 Biological role of protein persulfidation. **(a)** As a response to inflammation, tumour necrosis factor alpha ($\text{TNF-}\alpha$) migrates to nucleus where it binds to the transcription factor specificity protein 1 (SP1) causing the transcription and overexpression of CSE. H_2S produced by CSE induces persulfidation of p65 subunit of NF- κ B which translocates to nucleus where its interaction with co-activator ribosomal protein S3 (rps3) is enhanced, resulting in the transcription of anti-apoptotic genes. **(b)** S-nitrosation of parkin inhibits its E3 ubiquitin ligase activity which leads to accumulation of toxic proteins. This is found in Parkinson's disease where H_2S production is reduced. Persulfidation of parkin, on the other hand, increases parkin's enzymatic activity, which could be used as a therapeutic approach for Parkinson's disease treatment. **(c)** Keap-1 keeps Nrf-2 in the cytoplasm where it can be ubiquitinated and proteolytically degraded. Persulfidation of critical cysteines on Keap1 disturbs its interaction with Nrf-2 leading to the release of Nrf-2 which can now move into the nucleus where it binds to antioxidant response element (ARE) causing the transcription of various antioxidant defence genes. **(d)** The hypothetical role of persulfidation in protection of protein function during oxidative stress. When exposed to hydrogen peroxide, proteins undergo oxidation to form sulfenic acids (P-SOH), sulfinic acids (P-SO₂H) and sulphonic acids (P-SO₃H), which cause the irreversible inactivation of the protein. Sulfenic acids could react with H_2S to form persulfides. In addition, protein persulfides could be formed by other mechanisms, and when exposed to ROS, they will readily react with it, forming P-S-SO₃⁻, which can be enzymatically cleaved to restore free thiol on the protein

as in many other cases, is not clear. In the case of PTP-1B though, it is possible that ER stress in fact leads to PTP-1B inactivation by initial formation of sulfenic acid in the active site, which then reacts with H_2S to give persulfide. Although both

modifications are inhibitory, persulfide could be then removed easier and the enzyme's activity restored. Therefore, H₂S could serve to protect the enzyme from a longer-lasting and potentially irreversible inhibition.

4.6 S-sulfhydration of Keap1 and p66Shc Increases Cellular Antioxidative Defence and Prevents Senescence

A major mechanism in which antioxidant enzymes are induced involves the activation of the antioxidant response element (ARE) by the oxidative-stress sensor protein Kelch-like ECH-associated protein 1 (Keap1) and the transcription factor nuclear factor (erythroid-derived 2)-like 2 (Nrf-2) (Hybertson et al. 2011; Kaspar et al. 2009). Under basal conditions, Keap1 sequesters Nrf-2 in the cytoplasm by binding to its Neh2 domain (Hybertson et al. 2011; Kaspar et al. 2009; Wakabayashi et al. 2004). Chemical inducers, such as sulforaphane, are known to react with Keap1 cysteine residues, thereby promoting Nrf-2 nuclear accumulation and hence ARE activation (Wakabayashi et al. 2004). A widely accepted model for Nrf-2 nuclear accumulation describes that a modification of the Keap-1 cysteines leads directly to the dissociation of the Keap1–Nrf-2 complex and to the translocation of Nrf-2 into the nucleus.

H₂S cardioprotective effects in ischemia–reperfusion injury are partly caused by Nrf-2 nuclear translocation and activation of antioxidant defence enzymes (Calvert et al. 2009, 2010). Recently, two independent studies suggested that Keap-1 gets S-sulfhydrated when cells were exposed to H₂S. Modification of C151 on Keap1 stimulates dissociation of Nrf-2 enabling its translocation to nucleus where it regulates expression of cytoprotective genes and delays cellular senescence (Fig. 6c; Yang et al. 2013). In second study, Hourihan et al. (2013) found that H₂S stabilizes Nrf2 via covalent modification of amino acids C226 and C613 in the Keap1. The authors show that H₂S leads to the production of H₂O₂, which inhibits Keap1 by stimulating the formation of an intramolecular disulfide bond between C226 and C613. The Keap1 C226 and C613 residues are also persulfidated. This may be explained either by the ability of H₂S to reduce C226–C613 disulfides originally formed by H₂O₂ or by direct reaction of H₂S with sulfenylated residues formed by H₂O₂. More importantly, the authors observed that Nrf-2 controls CBS, CSE and sulfide–quinone reductase-like enzyme, suggesting that a feedback loop exists between Nrf-2 and H₂S.

P66Shc belongs to the ShcA family of proteins whose members share three common functionally identical domains: the C-terminal Src homology 2 domain (SH2), the central collagen homology domain (CH1) and the N-terminal phosphotyrosine-binding domain (PTB) (Giorgio et al. 2005). In response to oxidative stress such as UV exposure or H₂O₂, p66Shc gets activated by phosphorylation at Ser-36. The activated p66Shc is then dephosphorylated and translocates to mitochondria, where it binds to cytochrome c helping in electron transport process (Giorgio et al. 2005). P66Shc^{-/-} mice show 30 % increase of the lifespan. It has been shown recently that sulfhydration of p66Shc impaired the association of PKC_{βII}

and p66Shc and attenuated H₂O₂-induced p66Shc phosphorylation, a critical step in p66Shc-mediated mitochondrial ROS generation (Xie et al. 2014). H₂S is known to have dramatic effects on inhibiting oxidative stress, something that cannot be simply explained by its direct redox chemistry. A study by Xie and associates suggests that H₂S may inhibit mitochondrial reactive oxygen species production via a p66Shc-dependent mechanism.

4.7 MEK1/PARP-1 Activation and DNA Damage Repair

The salvage of DNA damage is essential for normal cell function. DNA damage stimulates a complex and highly concerted DNA damage repair response, which includes poly(ADP-ribose)ation catalysed by poly(ADP-ribose)ation polymerases (PARPs). Upon DNA damage, PARPs bind to DNA strand breaks and catalyse poly(ADP-ribose)ation which attracts other DNA damage repair proteins (D'Amours et al. 1999). The activation of PARPs is regulated by several kinases, of which MEK/ERK signalling cascade plays an important role (Cohen-Armon et al. 2007). Zhao et al. (2014) reported recently that H₂S attenuates DNA damage in human endothelial cells by causing S-sulfhydration of cysteine 341 on MEK1. This facilitates the translocation of phosphorylated ERK1/ERK2 into nucleus where it activates PARP-1 and increases the DNA damage repair yield, protecting cells from senescence.

4.8 TRP Channel S-sulfhydration Regulates Osteogenic Differentiation

Bone marrow mesenchymal stem cells (BMMSCs) are nonhaematopoietic multipotent stem cells and play an important role in the maintenance of the bone marrow homeostasis (Prockop 1997; Pittenge et al. 1999). BMMSCs and BMMSC-derived osteoblasts are responsible for bone formation and balancing osteoclast-mediated bone resorption in order to maintain bone mineral density (BMD) (Pittenge et al. 1999). CBS-deficient patients exhibit a variety of phenotypes, including osteoporosis. Osteoporosis is characterized by low bone mass and deterioration of osseous microarchitecture, resulting in decreased bone strength and increased risk of fragility fractures. This phenotype is often observed in patients with hyperhomocysteinemia (Herrmann et al. 2005; Melton 2003).

It has been demonstrated recently that H₂S deficiency causes aberrant intracellular Ca²⁺ influx because of reduced persulfidation of cysteine residues on multiple TRP channels (Liu et al. 2014c). Decreased Ca²⁺ influx downregulates PKC-/Erk-mediated Wnt/beta-catenin signalling which controls osteogenic differentiation of BMMSCs. Therefore, the authors suggest that bone marrow mesenchymal stem cells produce H₂S in order to regulate their self-renewal and

osteogenic differentiation and that H₂S deficiency results in defects in BMMSC differentiation. This study is in agreement with the previous observation that hydrogen sulfide protects MC3T3-E1 osteoblastic cells against H₂O₂-induced oxidative damage (Xu et al. 2011) suggesting that development of new H₂S-releasing drugs could be a potential therapeutic route for the treatment of osteoporosis.

5 Placing Persulfidation in a Broader Biological Context and Future Directions

The biochemical properties of protein persulfides discussed here suggest a specific and enhanced reactivity, which can be used for the regulation of protein's function. As mentioned above, persulfidation increases the nucleophilicity of protein thiols, and, as shown in several examples of proteins such as GAPDH and parkin, it increases their enzymatic activity. Further, the increased reducing power of persulfides suggests that they could act as efficient antioxidants. For example, GSSH reacts rapidly with H₂O₂, neutralizing its toxic effects. In fact, one of the possible roles of protein persulfidation could be the protection of a particular protein from irreversible damage induced by ROS and/or RNS (Fig. 6d). Thiol oxidation, which initially starts with the formation of sulfenic acids (still reversible modification), could proceed further with the formation of irreversible sulfonic acids. H₂S could react with sulfenic acid preventing this oxidation. In addition, persulfidated protein, even when exposed to ROS/RNS, will form an adduct that could be cleaved by the action of certain enzymes restoring free thiol. Therefore, spatio-temporal distribution of sulfenylation vs. persulfidation could help us to understand the relative ratio and importance of these two oxPTMs of cysteine.

S-nitrosation of proteins is a very important posttranslational modification of proteins. S-nitrosation of parkin leads to its inactivation (Chung et al. 2004). Conversely, persulfidation of parkin stimulates the enzyme increasing its ligase activity, which prevents accumulation of toxic proteins (Vandiver et al. 2013). On the other hand, intracellular S-nitrosothiol levels were found to be lower in cells with lower H₂S production, suggesting its role in *trans*-nitrosation reactions as well (Filipovic et al. 2012a). Therefore, the understanding of the NO and H₂S crosstalk, with particular emphasis on S-nitrosation vs. persulfidation, should be one of the future tasks.

All these could be done with proper tools for persulfide detection. Further development of assays, which would allow easy labelling of protein persulfides and their subsequent proteomic analysis, would accelerate the progression of the field. It is worth mentioning that most of the protein persulfidation reports to date used the methodology, which has been shown to generate artefacts.

Very little is known about the mechanisms of protein persulfidation. *Trans*-persulfidation of proteins by low molecular weight persulfides is an exciting concept that warrants further exploring. The role of metal centres in facilitating

persulfidation is also of potential interest. Elucidation of all these mechanisms (the role of H₂S vs. polysulfides vs. LMW persulfide) could help us in getting a few steps closer to the understanding of actual contribution of different H₂S-producing enzymes in the regulation of the intracellular S-sulfhydration levels.

Finally, the question that warrants an equally important attention is the extent to which this modification is regulatory and to which it is just a consequence of stochastic events. Answering this could lead to the identification of specific targets that could later on be pharmacologically tempered with. Therefore, this field of research will remain one of the hot topics for many years to come.

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Physiological Roles of Hydrogen Sulfide and Polysulfides

Hideo Kimura

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Abstract

Hydrogen sulfide (H₂S) has been recognized as a signaling molecule as well as a cytoprotective molecule. H₂S modulates neurotransmission, regulates vascular tone, protects various tissues and organs, regulates inflammation, induces angiogenesis, and detects cellular oxygen levels. H₂S is produced from L-cysteine by cystathionine β-synthase (CBS), cystathionine γ-lyase (CSE), and

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3-mercaptopyruvate sulfurtransferase (3MST) together with cysteine aminotransferase (CAT). Recently, a novel pathway for the production of H₂S from D-cysteine was identified, involving D-amino acid oxidase (DAO) together with 3MST. Sulfuration (also called sulfhydration), which adds sulfur atoms to the cysteine residues of target proteins to modify protein activity, has been extensively studied as a mode of H₂S action. Recently, hydrogen polysulfides (H₂S_n, where $n = 3-7$; $n = 2$ is termed as persulfide) have been found to sulfurate target proteins in the brain, including transient receptor potential ankyrin 1 (TRPA1) channels, Kelch-like ECH-associating protein 1 (Keap1), and phosphatase and tensin homolog (PTEN), much more potently than H₂S. The physiological stimuli that trigger the production of H₂S and polysulfides, and the mechanisms maintaining their local levels, remain unknown. Understanding the regulation of H₂S_n (including H₂S) production, and the specific stimuli that induce their release, will provide new insight into the biology of H₂S and will provide novel avenues for therapeutic development in diseases involving H₂S-related substances.

Keywords

H₂S • H₂S_n • CBS • CSE • 3MST • Sulfuration • Sulfhydration • TRPA1 • Keap1 • Nrf2 • PTEN

Abbreviations

3MST	3-mercaptopyruvate sulfurtransferase
ATP	Adenosine triphosphate
CAT	Cysteine aminotransferase
CBS	Cystathionine β-synthase
CO	Carbon monoxide
CSE	Cystathionine γ-lyase
CysSSH	Cysteine persulfide
DAO	D-amino acid oxidase
DHLA	Dihydrolipoic acid
DTT	Dithiothreitol
eNOS	Endothelial nitric oxide synthetase
Fe	Iron
GAPDH	Glyceraldehydes-3-phosphate dehydrogenase
GCL	Glutamate cysteine ligase
GSH	Glutathione
GSNO	Nitrosoglutathione
GSSH	Glutathione persulfide
GSSSH	Glutathione trisulfide
H ₂ O ₂	Hydrogen peroxide
H ₂ S	Hydrogen sulfide

H ₂ S _n	Polysulfide
HPLC	High-performance liquid chromatography
HSNO	Thionitrous acid
Keap1	Kelch ECH-associating protein 1
LTP	Long-term potentiation
NaHS	Sodium hydrosulfide
NF-κB	Nuclear factor-κB
NMDA	N-methyl D,L-aspartate
nNOS	Neuronal nitric oxide synthetase
NO	Nitric oxide
NOS	Nitric oxide synthetase
Nrf2	Nuclear factor-erythroid 2-related factor 2
O ₂	Oxygen
PD	Parkinson's disease
PERK	Protein kinase RNA-like ER kinase
PLP	Pyridoxal 5'-phosphate
PTEN	Phosphatase and tensin homolog
PTP	Protein tyrosine phosphatase
ROS	Reactive oxygen species
SAM	S-adenosyl-L-methionine
SNAP	S-nitroso-N-acetyl-D,L-penicillamine
SNP	Sodium nitroprusside
SSNO ⁻	Nitrosopersulfide
TNFα	Tumore necrosis factor α
TRPA1	Transient receptor potential ankyrin 1
VEGFR2	Vascular endothelium growth factor receptor 2
γGCS	γ-glutamyl cysteine synthetase

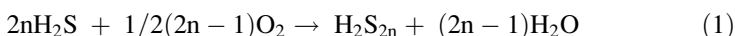
1 Introduction

Since Ramazzini's first description of hydrogen sulfide (H₂S) as a toxic gas in 1713, many papers have been published on H₂S toxicity. Enzymes involved in the H₂S-producing pathways were intensively studied from the 1950s to the 1970s (Meister et al. 1954; Cavallini et al. 1962; Braunstein et al. 1971). However, H₂S was thought to be a mere byproduct of metabolic pathways or simply a marker of enzyme activity. In 1989, measurements of the endogenous levels of sulfide in mammalian brains suggested a physiological role for this molecule (Goodwin et al. 1989; Warenycia et al. 1989). These initial studies by Goodwin et al. (1989) and Warenycia et al. (1989) overestimated the H₂S concentration, due to contaminant H₂S released from another cellular form of sulfur. Recent reevaluation confirmed the existence of H₂S in tissues, albeit at much lower concentrations than originally reported (Furne et al. 2008; Ishigami et al. 2009; Wintner et al. 2010). We have previously demonstrated a role for H₂S produced by cystathionine β-synthase

(CBS) as a neuromodulator in the brain and another role for H₂S produced by cystathionine γ -lyase (CSE) as a smooth muscle relaxant (Abe and Kimura 1996; Hosoki et al. 1997). The enzyme 3-mercaptopyruvate sulfurtransferase (3MST) also produces H₂S when 3MST is associated with thioredoxin (Shibuya et al. 2009; Mikami et al. 2011a; Yadav et al. 2013). Recently, we identified a novel pathway for the production of H₂S from D-cysteine, which involves the enzymes 3MST and D-amino acid oxidase (DAO) (Shibuya et al. 2013).

Various other effects of H₂S have been identified, including cytoprotective effects against oxidative stress and ischemia-reperfusion injury in various tissues and organs. H₂S protects neurons from oxidative stress by increasing the production of glutathione, a major intracellular antioxidant, and by scavenging reactive oxygen species (ROS) (Kimura and Kimura 2004; Whiteman et al. 2004; Kimura et al. 2010). The protective effects of H₂S against oxidative stress and ischemia-reperfusion injury have been demonstrated in the heart, the kidney, and in pancreatic islet β -cells (Elrod et al. 2007; Tripatara et al. 2008; Kaneko et al. 2009). H₂S induces angiogenesis by enhancing the expression of vascular endothelial growth factor (VEGF) and facilitating the phosphorylation of VEGF receptor 2 (VEGFR2); the regulation of nitric oxide synthase (NOS) may also be involved in this effect (Cai et al. 2007; Papapetropoulos et al. 2009; Polhemus et al. 2013). H₂S induces conditions similar to those found in hypoxia. H₂S metabolism is coupled with that of O₂, and H₂S concentrations are inversely related to those of O₂ in tissues. O₂-dependent H₂S inactivation may be effectively used as an O₂ sensor to accurately determine O₂ availability (Olson et al. 2006; Peng et al. 2010). H₂S exerts regulatory effects on inflammatory processes, thereby promoting the resolution of inflammation (Bhatia et al. 2005; Zanardo et al. 2006). H₂S also inhibits cancer development at various stages, but in colon cancer, H₂S is involved in promoting cellular bioenergetics, proliferation, and migration (Chattopadhyay et al. 2012; Szabo et al. 2013).

The mechanism by which H₂S exerts its actions has been proposed as sulfhydration or sulfuration: the addition of sulfur atoms to the cysteine residues of target proteins, inducing conformational changes to modify protein activity. It has been suggested that adenosine triphosphate (ATP)-dependent K⁺ channels, Kelch-like ECH-associated protein 1 (Keap1), nuclear factor kappa-B (NF- κ B), transient receptor potential ankyrin 1 (TRPA1), and parkin can be activated by sulfhydration or sulfuration (Mustafa et al. 2011; Ogawa et al. 2012; Sen et al. 2012; Vandiver et al. 2013; Yang et al. 2013). H₂S is sequentially oxidized to polysulfides with a varying number of sulfur atoms, until the number of sulfur atoms reaches eight; at that point, the sulfur molecules cyclize and separate from polysulfides (Nagy and Winterbourn 2010; Toohey 2011).



Recently, polysulfides (H₂S_{*n*}, where *n* = 3–7; when *n* = 2, the compound is termed as persulfide) were found to be present in the mammalian brain. Polysulfides

activate TRPA1 channels, facilitate the translocation of nuclear factor-erythroid 2-related factor 2 (Nrf2) to the nucleus via release of Nrf2 from sulfhydrated Keap1, and modulate the activity of PTEN; interestingly, polysulfides mediate these processes more efficiently than does H₂S (Greiner et al. 2013; Kimura et al. 2013; Koike et al. 2013). When considering whether these biological effects are mediated by H₂S or polysulfides, the efficiency of the effects and the chemical properties of the molecules suggest that polysulfides, rather than H₂S, sulfurate (or sulfhydrate) target proteins. H₂S may instead be involved in reducing reactions, such as the reduction of cysteine disulfide bonds.

2 Endogenous Free H₂S

Initial studies reported relatively high concentrations (50–160 μM) of H₂S in mammalian brains (Goodwin et al. 1989; Warenycia et al. 1989; Savage and Gould 1990). These studies used the methylene blue method, which is performed under strong acidic conditions. At pH values less than 5.4, H₂S is released from acid-labile sulfur, which consists mostly of the iron-sulfur complex at the active site of respiratory chain enzymes (Ishigami et al. 2009). Due to the localization of most enzymes containing an iron-sulfur complex to the mitochondrion, where the pH value is approximately 8, H₂S may not be released from iron-sulfur complexes under physiological conditions. Therefore, H₂S levels measured with the methylene blue method did not reflect free H₂S but instead reflected contaminant H₂S released from acid-labile sulfur.

The reevaluated endogenous H₂S concentrations are much lower than those previously measured, with recent studies suggesting a concentration range of 10 nM to 3 μM (Ishigami et al. 2009; Wintner et al. 2010; Levitt et al. 2011). These concentrations were measured during a steady state, and changes in H₂S concentration during cell activity are not well understood. Given that the K_m values of CSE and CBS for cysteine are 1.7 mM and 3.6 mM, respectively, and that the intracellular concentration of cysteine is approximately 100 μM (the exception being levels of approximately 1 mM in the kidney), the production of H₂S by CSE and CBS must occur relatively slowly, producing less amount of H₂S (Stipanuk et al. 2002; Chiku et al. 2009; Singh et al. 2009).

3 Regulation of H₂S Production

In order to function as a signaling molecule, H₂S concentrations must be properly regulated by physiological stimuli. There are at least two sources of H₂S: (1) H₂S-producing enzymes and (2) the intracellular H₂S stores.

3.1 H₂S-Producing Enzymes

Three enzymes are known to produce H₂S: the pyridoxal 5'-phosphate (PLP)-dependent CBS and CSE and the PLP-independent 3MST (Meister et al. 1954; Cavallini et al. 1962; Braunstein et al. 1971).

3.1.1 CBS

CBS produces H₂S via a β -replacement reaction involving L-cysteine. By catalyzing the β -replacement of L-cysteine with L-homocysteine, CBS produces H₂S with a 23-fold greater specific activity than that of H₂S produced from L-cysteine alone (Singh et al. 2009). Three sites on the CBS protein regulate its activity: (1) an S-adenosyl-L-methionine (SAM)-binding domain at the carboxy-terminus; (2) a heme group at the amino-terminus; and (3) cysteine 346 (Cys346) for S-glutathionylation. These three methods of CBS regulation are expanded upon in the following text (Fig. 1). (1) SAM enhances the H₂S-producing activity of CBS by approximately twofold (Abe and Kimura 1996). The SAM-binding domain covers the catalytic site of CBS in the absence of SAM, and binding to SAM thereby activates the enzyme (Shan et al. 2001). (2) CBS containing the Fe(II) form of heme is inhibited by the binding of carbon monoxide (CO) or nitric oxide (NO), while

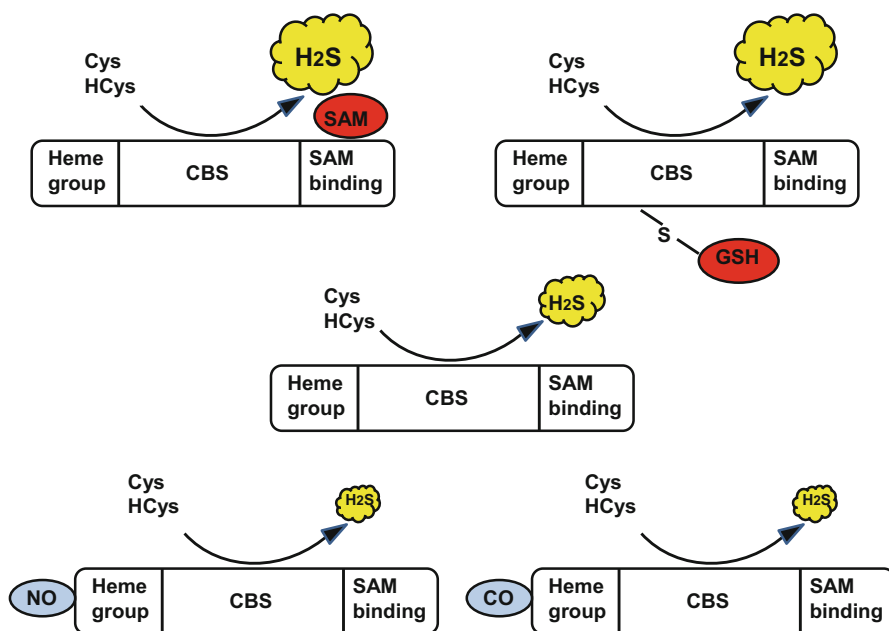


Fig. 1 The regulation of cystathionine β -synthase (CBS) activity. The activity of CBS is enhanced by the binding of S-adenosyl-L-methionine (SAM) to the SAM-binding domain located at the carboxy-terminus and by S-glutathionylation at Cys346. In contrast, the binding of nitric oxide (NO) or carbon monoxide (CO) to the heme group at the amino-terminus suppresses CBS activity

CBS activity is enhanced when Fe(II) is oxidized to Fe(III) (Taoka and Banerjee 2001). The regulation of H₂S production plays an important role in microcirculation in the brain (Morikawa et al. 2012). CO that is produced by heme oxygenase-2 in neurons suppresses the activity of CBS. Under hypoxic conditions, the production of CO is diminished, leading to the activation of CBS in astrocytes surrounding brain capillaries and the production of H₂S, which in turn vasodilates precapillary arterioles to restore blood flow. (3) Under conditions of oxidative stress, Cys346 of CBS is oxidized to a sulfenic acid, which binds to glutathione (Niu et al. 2014). CBS activity is enhanced by glutathionylation, increasing the production of cysteine and H₂S and in turn leading to increased glutathione production. Glutathionylation of CBS compensates for decreased levels of glutathione under oxidative stress.

The production of H₂S by CBS is enhanced by approximately 23-fold in the presence of homocysteine, by twofold when binding to SAM, and by threefold with glutathionylation. Therefore, the combination of these modifications to CBS may allow production of sufficient levels of H₂S to induce physiological responses despite low intracellular concentrations of cysteine (Abe and Kimura 1996; Singh et al. 2009; Niu et al. 2014).

3.1.2 CSE

CSE produces H₂S through an α,β -elimination reaction with L-cysteine (Chiku et al. 2009; Singh et al. 2009). It has been reported that CSE is regulated by Ca²⁺/calmodulin (Yang et al. 2008). However, the study by Yang and colleagues (2008) examined CSE activity in the presence of 1–2 mM Ca²⁺, which is the extracellular concentration of Ca²⁺. As CSE is a cytosolic enzyme, its activity should be examined in the presence of intracellular concentrations of Ca²⁺ (approximately 100 nM). Our recent reevaluation of the relationship between Ca²⁺ levels and CSE activity showed that the H₂S-producing activity of CSE is maximal at low Ca²⁺ concentrations and is suppressed by Ca²⁺ in the presence of PLP (Mikami et al. 2013). H₂S may be constitutively produced by CSE in cells under steady-state conditions in which the intracellular concentration of Ca²⁺ is maintained at low levels, whereas H₂S production is suppressed once the intracellular concentration of Ca²⁺ increases upon cell stimulation. Because calmodulin and its selective inhibitor W-7 do not alter CSE activity, it appears that calmodulin is not involved in the regulation of CSE by Ca²⁺ (Mikami et al. 2013).

In addition to regulation of CSE activity, the expression level of CSE is also transcriptionally regulated. The CSE gene promoter region has an SP1-binding site, which is activated by the multifunctional, pro-inflammatory cytokine tumor necrosis factor α (TNF α) (Ishii et al. 2004; Sen et al. 2012).

3.1.3 3MST

3MST produces H₂S by metabolizing 3-mercaptopyruvate (3MP); 3MP is synthesized through the metabolism of L-cysteine and α -ketoglutarate by cysteine aminotransferase (CAT; identical to aspartate aminotransferase) (Meister et al. 1954; Taniguchi and Kimura 1974; Ubuka et al. 1978; Cooper 1983; Shibuya et al. 2009). 3MST is localized to both the mitochondria and the cytosol; CAT has

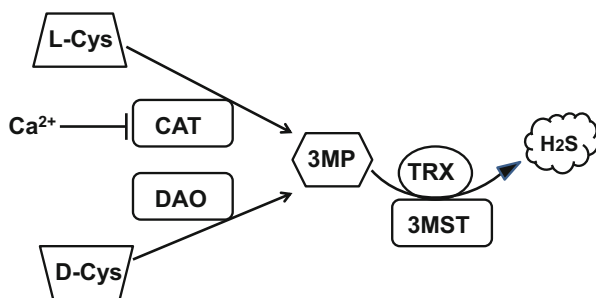


Fig. 2 Hydrogen sulfide (H₂S) production from L- and D-cysteine. L-Cysteine is metabolized to 3-mercaptopyruvate (3MP) by cysteine aminotransferase (CAT), which is regulated by Ca²⁺. Achiral 3MP is also produced from D-cysteine by D-amino acid oxidase (DAO). 3MP is metabolized to H₂S by 3-mercaptopyruvate sulfurtransferase (3MST) accompanied by thioredoxin (TRX)

both a mitochondrial and a cytosolic form, with these two CAT forms sharing 48 % identity in their amino acid sequences (Ubuka et al. 1978; Akagi 1982; Doyle et al. 1990; Nagahara et al. 1998; Shibuya et al. 2009). Unlike the cytosol, mitochondria contain a sufficiently high cysteine concentration (approximately 1 mM) for the 3MST/CAT pathway to produce H₂S (Griffith 1999; Tateishi et al. 1977) (Fig. 2).

3MST requires a reducing agent to produce H₂S from 3MP, while neither CBS nor CSE requires such a reducing agent. The endogenous reducing agents, thioredoxin and dihydrolipoic acid (DHLA), associate with 3MST to produce H₂S (Mikami et al. 2011a; Yadav et al. 2013). Thioredoxin is present at approximately 20 μM in cells and is four times as potent as dithiothreitol (DTT), and DHLA is found at approximately 40 μM in mitochondria and is as potent as DTT. Other endogenous reducing agents do not associate with 3MST to produce H₂S. Dithiols such as thioredoxin (which has two cysteine residues at the active site), DHLA, and DTT associate with 3MST to release H₂S (Mikami et al. 2011a).

The activity of CAT is suppressed by Ca²⁺, but as is the case with CSE, calmodulin is not involved in the regulation of CAT activity (Mikami et al. 2011b). Therefore, 3MP is supplied to 3MST under steady-state conditions, whereas the supply of 3MP is stopped once the intracellular concentrations of Ca²⁺ are increased.

3MST, together with DAO, produces H₂S from D-cysteine (Shibuya et al. 2013). DAO, which is located in peroxisomes, metabolizes D-cysteine to achiral 3MP (Fig. 2). Peroxisomes and mitochondria are in close vicinity to each other and can establish physical contact, and they exchange metabolites via a specific form of vesicular trafficking (Schumann and Subramani 2008). 3MP produced in peroxisomes is transported into mitochondria, where it is metabolized by 3MST to H₂S. This pathway is active in the brain and the kidney; this spatial restriction may be useful for the targeted production of H₂S in those specific organs following systemic delivery of D-cysteine. The 3MST/DAO pathway is highlighted in the

kidney, where 60 times more H₂S is produced from D-cysteine than from L-cysteine (Shibuya et al. 2013).

3.2 The Intracellular H₂S Store

Several factors change the activity of H₂S-producing enzymes, and H₂S levels may change according to physiological stimuli. H₂S can be provided from bound sulfane sulfur, which releases H₂S under physiological reducing conditions. The levels of bound sulfane sulfur vary among tissues, and the amount of H₂S released from brain homogenates by DTT is approximately 1.5 μmol/g protein (Ishigami et al. 2009). Lysates of neurons and astrocytes release H₂S in the presence of 3 mM glutathione and 100 μM cysteine at pH 8.4. The pH of astrocytes can reach 8.4 when the extracellular concentration of K⁺ is 10 mM, a level that can be induced by the excitation of nearby neurons. However, H₂S release from astrocytes has not been successfully detected; this is most likely due to the released H₂S being present at an insufficient concentration to be detected in perfusion medium (Ishigami et al. 2009).

4 Neuromodulation by H₂S and Polysulfides

H₂S facilitates the induction of hippocampal long-term potentiation (LTP) by enhancing the activity of N-methyl-D-aspartate (NMDA) receptors in the brain. LTP induction requires the activation of NMDA receptors, and reducing agents such as DTT reduce the cysteine disulfide bond at the hinge of the NMDA receptor ligand-binding domain to activate these receptors (Aizenman et al. 1989). In spite of having less reducing activity, the H₂S donor sodium hydrosulfide (NaHS) induces LTP more potently than does DTT, even when DTT is present at 10 times greater concentration than that of NaHS (Abe and Kimura 1996). It is therefore difficult to attribute the effects of H₂S on LTP to the reducing activity of H₂S alone.

Neurons are surrounded by astrocytes, which also express neurotransmitter receptors and are activated by neurotransmitters. The activated astrocytes, in turn, release gliotransmitters to modulate synaptic activity (Araque et al. 1999). Our previous research showed that NaHS activates transient receptor potential (TRP) channels to induce Ca²⁺ influx in astrocytes (Nagai et al. 2004). The involvement of this mechanism in the induction of LTP is discussed below.

In our previous research, we found that polysulfides activate TRP channels more potently than does H₂S (Nagai et al. 2006; Oosumi et al. 2010). High concentrations of NaHS (1–10 mM) activate TRPA1 channels in sensory neurons in the urinary bladder and dorsal ganglion (Streng et al. 2008; Ogawa et al. 2012). In these studies on sensory neurons, a portion of the NaHS-derived H₂S might be oxidized to polysulfides that activate TRPA1 channels. We found that polysulfides in the brain activated astrocytic TRPA1 channels to a much greater extent than did H₂S,

inducing a maximal response (Kimura et al. 2013). A comparison of K_m values revealed that the affinity of TRPA1 channels for polysulfides was approximately 1000 times greater than for H_2S (Kimura et al. 2013). The level of endogenous polysulfides in the brain was found to be approximately 20 μM , using HPLC analysis with monobromobimane (Kimura et al. 2013).

Ogawa and colleagues determined that Cys422 and Cys622 in the TRPA1 channel amino-terminus are sensitive to H_2S (Ogawa et al. 2010). Given that DTT abolishes the effects of high concentrations of H_2S , these two cysteine residues must be sulfurated by polysulfides generated by oxidation of H_2S . The idea that polysulfide, rather than H_2S , sulfurates TRPA1 channels is based on the fact that substances with the same oxidation state do not react with each other. The oxidation state of sulfur in H_2S and in cysteine residues is -2 . In contrast, the oxidation state of sulfur in polysulfides is 0, meaning that polysulfides easily react with cysteine residues to add sulfur for the generation of bound sulfane sulfur. It is also possible that one cysteine residue is sulfurated earlier than an adjacent cysteine, and the latter may react with the former to produce a cysteine disulfide bond (Fig. 3).

Shigetomi and colleagues recently demonstrated that the activated TRPA1 channels in astrocytes increase the intracellular concentrations of Ca^{2+} , which facilitate the release of the gliotransmitter D-serine from astrocytes; D-serine

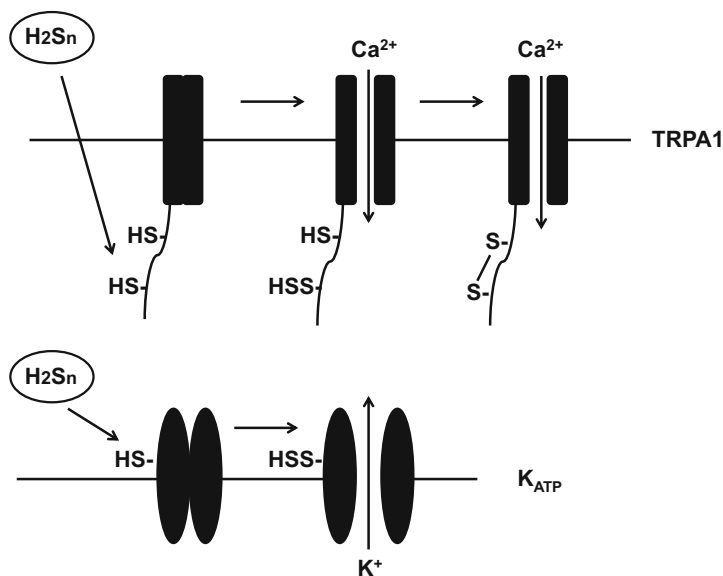


Fig. 3 The regulation of channel activity by sulfuration (sulfhydrylation). Transient receptor potential ankyrin 1 (TRPA1) channels and Adenosine triphosphate (ATP)-dependent K^+ (K_{ATP}) are activated by sulfuration (sulfhydrylation). H_2S has been proposed to sulfurate these channels, but polysulfides (H_2S_n) have much greater activity than H_2S . Thus, polysulfides have been proposed to be the molecular entity that induces sulfuration

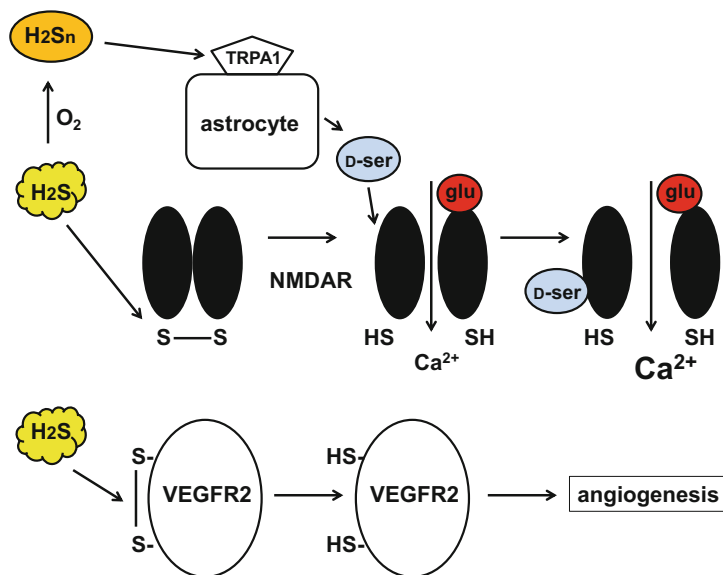


Fig. 4 H₂S enhances receptor activity by reducing a cysteine disulfide bond. H₂S enhances the activity of N-methyl-D-aspartate (NMDA) receptors by reducing the cysteine disulfide bond at the hinge of the ligand-binding domain. The activity of NMDA receptors is also enhanced by binding the gliotransmitter D-serine, released from astrocytes, which are activated by polysulfides (H₂S_n) through the activation of TRPA1 channels. H₂S also enhances the activity of vascular endothelial growth factor receptor 2 (VEGFR2) by reducing the cysteine disulfide bond between Cys1045 and Cys1024

activates NMDA receptors on postsynaptic neurons, leading to the induction of LTP (Shigetomi et al. 2012; 2013). A similar mechanism may be involved in the induction of LTP by H₂S and polysulfides. As described previously in this section, H₂S reduces the cysteine disulfide bond and enhances the activity of NMDA receptors, and polysulfides activate TRPA1 channels in astrocytes and induce D-serine release that enhances NMDA receptor activity. In contrast, DTT is not able to activate TRPA1 channels but reduces the cysteine disulfide bond in NMDA receptors. These findings explain the difference in LTP induction efficiency between H₂S/polysulfides and DTT (Abe and Kimura 1996; Kimura et al. 2013) (Fig. 4).

5 Cytoprotective Effects of H₂S and Polysulfides

H₂S protects various tissues/organs from oxidative stress or ischemia-reperfusion injury (Kimura and Kimura 2014; Elrod et al. 2007). Recently, polysulfides were found to have cytoprotective effect through a mechanism of sulfuration (sulfhydrating) target proteins (Kimura 2014).

5.1 Glutathione Production and ROS Scavenging

H₂S protects various tissues and organs from oxidative stress and ischemia-reperfusion injury. H₂S enhances the activity of the cysteine transporter and of the cystine/glutamate antiporter to increase the import of cystine even in the presence of high glutamate concentrations (Kimura and Kimura 2004). Cystine is reduced to cysteine inside cells and is used as a substrate for glutamate cysteine ligase (GCL) (previously known as γ -glutamylcysteine synthetase), which is a rate-limiting enzyme in glutathione production. H₂S enhances the activity of GCL. This enhancement only occurs when H₂S is applied extracellularly, not when GCL is directly exposed to H₂S (Kimura et al. 2010). This observation suggests that H₂S may activate a signaling pathway from the cell surface, which acts downstream to enhance the activity of GCL.

H₂S also scavenges ROS (Whiteman et al. 2004; Kimura et al. 2010). Considering the endogenous concentrations of H₂S (in the nM to μ M range) and glutathione (in the mM range) and the fact that H₂S is not a strong reducing agent compared to glutathione, the suppression of oxidative stress by H₂S may be due to the enhancement of GCL activity and subsequent glutathione production, rather than the direct scavenging of ROS.

5.2 Translocation of Nrf2 to the Nucleus

H₂S also upregulates the transcription of antioxidant genes to exert its cytoprotective effect. Keap1 binds Nrf2 and retains it in the cytoplasm, and the Keap1/Nrf2 complex is ubiquitinated and degraded by the proteasome under steady-state conditions. The cysteine residues of Keap1 serve as sensors for oxidative stress, and their modification causes a conformational change in Keap1 that releases Nrf2, which is then translocated to the nucleus and upregulates the transcription of antioxidant genes (Wakabayashi et al. 2004). H₂S protects cardiac myocytes from ischemia-reperfusion injury by preserving the structure and function of mitochondria (Elrod et al. 2007). The translocation of Nrf2 to the nucleus has been suggested as a mechanism for the H₂S-mediated protection of myocytes (Calvert et al. 2009). Another study suggested that H₂S sulfurates (sulfhydrates) Keap1 to release Nrf2 for translocation to the nucleus (Yang et al. 2013). However, as described in section 4, it is difficult for H₂S to sulfurate (sulfhydrate) cysteine residues in Keap1 because H₂S and cysteine residues share the same oxidation state. Instead, polysulfide has been suggested as the molecular entity that efficiently sulfurates (sulfhydrates) Keap1 to release Nrf2 for nuclear translocation (Koike et al. 2013).

5.3 Modification of Proteins by Sulfuration (Sulphydration)

Modification of enzyme activities by sulfuration was studied extensively from the 1960s to the 1980s (Kato et al. 1966; Branzoli and Massey 1974; Conner and Russell 1983). Mustafa et al. (2009) defined the process of sulphydration (sulfuration) as the reaction in which sulfur provided by H_2S attaches to reactive cysteine residues in target proteins. Sulfuration of glyceraldehyde 3-phosphate dehydrogenase (GAPDH) by H_2S increases its catalytic activity by sevenfold, and DTT treatment removes the added sulfur to diminish GAPDH activity (Mustafa et al. 2009). In contrast, nitrosylation of the same GAPDH cysteine residue abolishes the catalytic activity. Thus, sulfuration has been proposed to have the opposite effect to nitrosylation. Sulfuration of GAPDH by H_2S is supported by the observation that the activity of GAPDH is reduced by 35 % in CSE-knockout mice. Actin polymerization is also enhanced by H_2S , and this effect is reversed by DTT (Mustafa et al. 2009). ATP-dependent K^+ channels provide a further example of the regulation of protein function by sulfuration. ATP-dependent K^+ channels are sulfurated by H_2S at Cys43 on the Kir6.1 subunit; this sulfuration prevents association of the subunit with ATP, leading to activation of the K^+ channel (Mustafa et al. 2011) (Fig. 3).

NF- κ B, which plays a key role in regulating immune responses, is sulfurated by H_2S to increase its interaction with ribosomal protein S3, resulting in upregulation of antiapoptotic genes (Sen et al. 2012). Stress responses are also regulated by sulfuration. Sulfuration-mediated inhibition of protein tyrosine phosphatase (PTP) enables protein kinase RNA-like ER kinase (PERK) to retain its active state and phosphorylate eukaryotic translational initiation factor 2 α (eIF2 α), inhibiting global protein synthesis (Krishnan et al. 2011). In Parkinson's disease, which affects the substantia nigra and presents as motor abnormalities, the E3 ubiquitin ligase parkin is affected and causes accumulation of α -synuclein, a major component of Lewy bodies. In the brains of patients with Parkinson's disease, parkin activity is decreased in line with reduced activity, sulfuration of Cys95 and Cys59 in parkin is diminished, and nitrosylation of parkin, which inactivates it, is increased (Vandiver et al. 2013).

Tao et al. (2013) proposed a role for H_2S in the reduction of cysteine disulfide bonds rather than sulfuration. H_2S induces angiogenesis by promoting the migration and tube formation of vascular endothelial cells, with the effects of H_2S being mediated by VEGFR2 (Cai et al. 2007; Papapetropoulos et al. 2009). H_2S activates VEGFR2 by reducing a disulfide bond between Cys1045 and Cys1024. Mass spectrometry analysis showed that H_2S reduced a cysteine disulfide bond contained within a synthesized hexapeptide but did not sulfurate any of the 20 free amino acids, including cysteine (Tao et al. 2013). The sulfuration of cysteine residues was only transiently observed as an intermediate step during the reduction of the disulfide bond by H_2S ; the intermediate was immediately attacked by HS^- and reduced to cysteine (Fig. 4). These observations suggest that H_2S may be involved in reducing reactions such as the reduction of cysteine disulfide bonds, whereas polysulfides may mediate sulfuration. Because the reducing activity of H_2S is

weaker than that of cysteine and glutathione, the regulation of local concentrations of H₂S and the accessibility to the targets need to be clarified.

5.4 Glutathione Polysulfide

Massey and colleagues initially reported that glutathione persulfide (GSSH or GSS⁻) produced from glutathione trisulfide (GSSSG) reduces cytochrome c more efficiently than did glutathione (GSH) and that cysteine trisulfide has a similar effect (Massey et al. 1971). Francoleon and colleagues recently confirmed the reducing effect of GSSH on cytochrome c and found that GSSH was also able to suppress the activity of papain (Francoleon et al. 2011). GSSH was also found to scavenge H₂O₂ more efficiently than GSH (Ida et al. 2014).

Ida and colleagues (2014) proposed a mechanism of cysteine persulfide (CysSSH) and glutathione polysulfide production. CBS and CSE metabolize cystine to CysSSH, which produces GSSH in an exchange reaction with GSH. However, the physiological relevance of this pathway needs to be reevaluated. Although CSE has a high affinity for cystine (with a K_m value of 30–70 μM), CSE localizes to the cytosol where cystine is not present at a sufficient concentration to react with the enzyme. Cystine is an extracellular form of cysteine and is present at less than 0.2 μM in the cytosol, with only a few exceptions (such as the A549 cell line, containing approximately 12 μM cystine) (Stipanuk 1986; Ida et al. 2014). The extracellular concentration of cystine is only 40 μM in human blood (Brigham et al. 1960). The CSE- and CBS-catalyzed production of CysSSH and polysulfides was examined with cystine present at mM concentrations, which were far higher than the measured physiological concentration (Ida et al. 2014).

6 Cross Talk of H₂S with NO

The relaxation effect of H₂S on thoracic aorta smooth muscle was weak compared to the effects of H₂S on other smooth muscles found in the ileum and the portal vein; however, a synergistic effect was observed between H₂S and NO in thoracic aorta smooth muscle (Hosoki et al. 1997). A similar synergistic effect has been observed in the twitch response of the ileum (Teague et al. 2002). Cross-talk between H₂S and NO is also present in the production and transcription of an H₂S-producing enzyme. The NO donors nitroprusside (SNP) and S-nitroso-N-acetyl-DL-penicillamine (SNAP) enhance the production of H₂S and upregulate the transcription of CSE, respectively (Zhao et al. 2001).

Lefer and colleagues recently showed that H₂S activated endothelial NO synthetase (eNOS) to increase NO production, thereby providing a mechanism for the protection of the heart from ischemia-reperfusion injury (King et al. 2014) based on six observations outlined below. (1) The levels of H₂S and bound sulfane sulfur were reduced in CSE-knockout mice compared to wild-type mice. (2) The eNOS activation site was less phosphorylated in CSE-knockout mice than in wild-type

mice, whereas the eNOS inhibitory site was more greatly phosphorylated in CSE-knockout mice. (3) The levels of NO metabolites in the blood and heart were lower in CSE-knockout mice than wild-type mice. (4) Administration of H₂S activated eNOS and increased NO bioavailability in CSE-knockout mice. (5) Ischemia-reperfusion injury was exacerbated in CSE-knockout mice. (6) The administration of H₂S failed to protect against myocardial ischemia-reperfusion injury in eNOS-deficient mice. H₂S also protected against pressure overload-induced heart failure by a similar mechanism (Kondo et al. 2013). Lefer and colleagues obtained interesting data relating to the levels of cyclic guanosine monophosphate (cGMP) in plasma and in the heart. Although plasma cGMP levels were significantly lower in CSE-knockout mice than in wild-type mice, cardiac cGMP levels were not significantly different between CSE-knockout and wild-type mice (King et al. 2014). This finding suggests that plasma cGMP may play a more significant role than cardiac cGMP in protecting the heart against ischemia-reperfusion injury.

In contrast, H₂S suppressed the activity of neuronal NOS (nNOS) in submucosal as well as myenteric-ganglion neurons innervated the colon (Sha et al. 2014). Endogenously produced H₂S acts in an autocrine fashion to inhibit NO generation from nNOS in colonic circular smooth muscle cells. This knowledge comes from the observation that the endogenous generation of NO was significantly greater in muscle preparations from CSE-knockout mice than in those from wild-type mice. As H₂S exerts different NO production outcomes depending on the type of NOS expressed, a more detailed understanding of the mechanism of H₂S is required (King et al. 2014; Sha et al. 2014).

The chemical interaction between H₂S and NO produces nitrosothiol, which releases NO in the presence of Cu²⁺ (Whiteman et al. 2006). H₂S (HS⁻) releases NO from the NO intermediate or carrier, nitrosogluthathione (GSNO), potentiating vascular smooth muscle relaxation (Ondrias et al. 2008). Filipovic and colleagues reported that in the presence of human umbilical vein endothelial cells or Fe³⁺-porphyrins, nitrite and H₂S interact with each other to produce an intermediate compound, thionitrous acid (HSNO), which produces NO and HS[•], or produces nitroxyl (HNO) by a further reaction with H₂S (Filipovic et al. 2012). In contrast, Feelisch and colleagues reported that HSNO does not effectively release NO owing to a relatively strong S-N bond (Cortese-Krott et al. 2014). Feelisch and colleagues demonstrated that nitrosopersulfide (SSNO⁻ or HSSNO), which is more stable than HSNO, efficiently releases NO and polysulfides, resulting in more effective activation of soluble guanylyl cyclase and relaxation of smooth muscle compared to that observed with the parent nitrosothiol (Cortese-Krott et al. 2014). However, these studies were performed in the presence of high concentrations of NaHS (in the mM range). Thus, the physiological relevance of the interaction between H₂S and NO needs to be further examined.

7 Concluding Remarks

Almost two decades have passed since the first demonstration of a physiological role for H₂S, and various aspects of this molecule have been extensively studied. Although the steady-state levels of H₂S have recently been reevaluated as much lower than initially reported, the range of change in H₂S concentrations due to cellular activity and the physiological stimuli to induce such changes are not well understood. One of the approaches to address these problems is to understand the regulation of H₂S metabolic turnover (Kimura 2012; Kabil and Banerjee 2014). It has been proposed that a balance between H₂S production and degradation regulates its physiological levels. In terms of H₂S production, CBS activity is enhanced by S-adenosyl-L-methionine and glutathionylation, and the activities of CSE and CAT are regulated by Ca²⁺; H₂S degradation is performed by mitochondrial enzymes. However, the specific physiological stimuli required to effect changes in cellular H₂S levels have not been identified; identification of these specific stimuli is required to elucidate a role of H₂S as a signaling molecule.

Polysulfides have recently emerged as candidate molecules for the physiological mediators of H₂S signaling. Polysulfides activate channels, transcription factors, and enzymes by sulfuration (sulphydration) of these target molecules; polysulfides also have more potent antioxidant effects than H₂S (Greiner et al. 2013; Kimura et al. 2013; Koike et al. 2013; Ida et al. 2014). Polysulfides may be produced by both enzymatic and nonenzymatic pathways. Although a pathway for the enzymatic production of polysulfides has been proposed, the concentrations of substrate required are well outside the physiologically relevant range (Ida et al. 2014). It is also necessary to clarify how nonenzymatic production of polysulfides is regulated.

The cross-talk between H₂S and NO has also been extensively studied. A synergistic effect between both substances was initially reported, and subsequently, the transcriptional regulation of H₂S-producing enzymes by NO was identified (Hosoki et al. 1997; Zhao et al. 2001; Teague et al. 2002). Recent studies demonstrated the regulation of NO production by H₂S through the activation or inhibition of NOS isoforms (Kondo et al. 2013; Sha et al. 2014). Further study is necessary to clarify the mechanisms underlying the varying effects of H₂S on different NOS isoforms and tissue types. The chemical interaction between H₂S and NO produces HSNO, HNO, or HSSNO, and the rate of production was changed in the presence of cells (Filipovic et al. 2012; Cortese-Krott et al. 2014). Because these studies were performed in the presence of high concentrations of substrates, the interaction between H₂S and NO must be reexamined under physiological conditions. It is also necessary to clarify whether H₂S and NO are present in cells at the same time and if enzymes producing these molecules are activated under the same conditions.

By understanding the biochemical nature of these molecules, as well as their mechanisms of action and regulation, the physiological function and therapeutic potential of H₂S and related molecules will be unveiled.

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Part III

Cardiovascular and Urogenital Systems

H₂S and Blood Vessels: An Overview

Guangdong Yang and Rui Wang

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Abstract

The physiological and biomedical importance of hydrogen sulfide (H_2S) has been fully recognized in the cardiovascular system as well as in the rest of the body. In blood vessels, cystathionine γ -lyase (CSE) is a major H_2S -producing enzyme expressed in both smooth muscle and endothelium as well as periadventitial adipose tissues. Regulation of H_2S production from CSE is controlled by a complex integration of transcriptional, posttranscriptional, and posttranslational mechanisms in blood vessels. In smooth muscle cells, H_2S regulates cell apoptosis, phenotypic switch, relaxation and contraction, and calcification. In endothelial cells, H_2S controls cell proliferation, cellular senescence, oxidative stress, inflammation, etc. H_2S interacts with nitric oxide and acts as an endothelium-derived relaxing factor and an endothelium-derived hyperpolarizing factor. H_2S generated from periadventitial adipose tissues acts as an adipocyte-derived relaxing factor and modulates the vascular tone. Extensive evidence has demonstrated the beneficial roles of the CSE/ H_2S system in various blood vessel diseases, such as hypertension, atherosclerosis, and aortic aneurysm. The important roles signaling in the cardiovascular system merit further intensive and extensive investigation. H_2S -releasing agents and CSE activators will find their great applications in the prevention and treatment of blood vessel-related disorders.

Keywords

Hydrogen sulfide • Cystathionine γ -lyase • Blood vessel • Smooth muscle cells • Endothelial cells • Periadventitial adipose tissues • Blood vessel-related disorders

1 Hydrogen Sulfide Is a Gasotransmitter

The physiological and biomedical importance of hydrogen sulfide (H_2S) has been fully recognized in the cardiovascular system as well as in the rest of the body. The gasotransmitter identity of H_2S is validated against six criteria of gasotransmitters (Wang 2002, 2014):

1. H_2S is a small gas molecule. Once produced inside eukaryotes, it is partially dissolved in biological milieu in its free or bounded forms and partially dissociated to HS^- . It is occasionally squabbled over whether H_2S or nitric oxide (NO) in eukaryotes is still *gas*. Oxygen in the blood is a gas and this does not need verification by seeing oxygen gas bubbles. This is the same case with gasotransmitters. NO, carbon monoxide (CO), H_2S , and NH_3 are gas molecules in our bodies.
2. H_2S is freely permeable to the plasma membrane and intracellular organelle membranes. The same signal relay sequence for neurotransmitters is not required for H_2S signal since there will be no synaptic barrier or cognate

membrane receptors for the transmembrane movement of this gas molecule. The idea of a “gas channel” has been around for a while. No experimental evidence exists to date, however, which shows the reliance of transmembrane movement of H₂S on such “channels”. Furthermore, one has to distinguish the transmembrane movement of undissociated gas molecule from that of its dissociated ionic forms. Whereas ammonia gas (NH₃) freely permeates cell membranes, ammonium ion (NH₄⁺) passes the membrane through ion channels.

3. H₂S is endogenously generated in mammalian cells under both physiological and pathophysiological conditions. L-cysteine and homocysteine are the most important substrates of enzymatic H₂S production. Yes, H₂S is a metabolic product of reverse-transsulfuration pathway. But the production of H₂S is more than a metabolic need, more than a sulfur transfer phenomenon, and more than the degradation of cysteine or homocysteine. It interacts with different protein targets, alters the directions of multiple signaling pathways, and forms signaling webs and nets intracellularly and intercellularly. And H₂S does all these in response to the functional regulation as well as the metabolic needs of the body.
4. H₂S has well-defined specific functions at physiologically relevant concentrations. The effects of H₂S on the cardiovascular system, neuronal system, respiratory system, and gastrointestinal system, to name a few, have been extensively and convincingly documented. These effects of H₂S are realized at its physiologically relevant concentrations because decreasing endogenous H₂S level (the knockout or knockdown of H₂S-generating enzymes from the whole animal to tissue to cell levels) or increasing it (overexpression or knock-in of these enzymes) has been clearly correlated to the correspondingly functional changes in different systems.
5. The application of H₂S donors (fast releasing or slow releasing) has shown the similar effects as endogenous H₂S on different biological systems.
6. H₂S is involved in signal transduction and has specific cellular and molecular targets. It induces S-sulfhydration of numerous proteins. It regulates the levels and activities of traditional second messengers, such as cGMP, cAMP, and intracellular calcium. Its action and production are sensitive to cellular oxygen levels. Mitochondrial bioenergetics, endoplasmic reticulum stress, and gene transcription and translation in the nucleus are within the impact radius of H₂S.

It is relevant here to comment on the contextual connection of neurotransmitter and gasotransmitter to the conventional denotation of “transmitter”. A biological transmitter usually refers to a biological molecule that is generated in response to the homeostatic need and serves for “communication” to meet this need. Neurotransmitter and gasotransmitter both conduct the “communication” between the homeostatic needs and functional changes. The former does this via second messengers, and the latter directly interacts with its downstream signaling webs and nets.

2 Endogenous Production of H₂S in Blood Vessels

Different enzymes are involved in the production of H₂S from both vascular smooth muscle cells (SMCs) and endothelial cells (ECs). Cystathionine β-synthase (CBS) is critical for the transsulfuration of homocysteine to generate cystathionine and then to H₂S. Similar to CBS, cystathionine γ-lyase (CSE) is also a pyridoxal-5'-phosphate-dependent H₂S-generating enzyme. In the cardiovascular system, the transformation of L-cysteine to H₂S is mainly catalyzed by CSE with ammonium and pyruvate as two coproducts.

CSE gene in the cardiovascular system was cloned for the first time in 2001 (Zhao et al. 2001). This vascular CSE gene cloned from rat mesenteric artery tissues (GenBank #AB052882) shares the same sequence with that of rat liver CSE gene (GenBank #AY032875). Hosoki et al. (1997) detected CSE mRNA in rat thoracic aorta and portal vein (Hosoki et al. 1997). Zhao et al. showed CSE mRNA in rat mesenteric artery, tail artery, and pulmonary arteries (Zhao et al. 2001). The first Western blot study on CSE protein expression in the cardiovascular system was reported in 2006 (Yang et al. 2006) when CSE proteins were observed in human aorta vascular SMCs. In 2008, the expression of CSE proteins in vascular endothelium was reported (Yang et al. 2008). Following studies have demonstrated endothelial expression of CSE in mice, rats, and humans (Altaany et al. 2013; Papapetropoulos et al. 2009). To date, CSE proteins and/or mRNA have been detected in newborn pig cerebral microvessels (Leffler et al. 2011), mouse renal artery SMCs (Sen et al. 2012b), rat pulmonary artery SMCs and ECs (Sun et al. 2011; Chen et al. 2009), mouse pulmonary artery SMCs and ECs (Wang et al. 2011), human pulmonary artery SMCs (Kiss et al. 2008), bovine pulmonary artery SMCs and sea lion resistance pulmonary arteries (Olson et al. 2010), rat hepatic artery and portal vein (Siebert et al. 2008), and human internal mammary artery (Webb et al. 2008).

The detection of CBS proteins in blood vessels has been reported in hepatic artery and pulmonary artery. CBS proteins were observed in rat pulmonary artery rings (Sun et al. 2011) and bovine pulmonary artery endothelial cells (Olson et al. 2010). In rat hepatic artery and portal vein, both CSE and CBS proteins were localized, but interestingly the terminal branches of the hepatic afferent vessels only expressed CSE (Siebert et al. 2008).

Cysteine aminotransferase (CAT) and 3-mercaptopyruvate sulfurtransferase (MST) are other two enzymes involved in H₂S production in the cardiovascular system. Whereas CAT uses PLP as its cofactor, zinc is the cofactor of MST. The sequential reactions catalyzed by CAT and MST lead to the transformation of cysteine to 3-mercaptopyruvate (3-MP) to sulfane sulfur. This bound sulfur will need to be released or reduced to free H₂S (Wang 2012b). CAT protein was found in vascular ECs, and MST protein was localized in both ECs and SMCs of rat thoracic aortae (Shibuya et al. 2009). MST protein was also localized in bovine pulmonary artery SMCs and sea lion resistance pulmonary arteries (Olson et al. 2011).

3 Regulation of CSE-Mediated Vascular Production of H₂S

H₂S production through CSE is both tonic under resting conditions and phasic upon specific stimulations. The activation of muscarinic cholinergic receptor by acetylcholine in vascular ECs leads to the elevation of intracellular calcium. Subsequently, calcium-activated calmodulin stimulates CSE to produce H₂S in ECs (Yang et al. 2008). Testosterone is another endogenous CSE stimulator in vascular system (Bucci et al. 2009). Its vasorelaxant effect on rat aortic rings in vitro was inhibited by DL-propargylglycine (PPG) and β -cyano-L-alanine (BCA), two inhibitors of CSE. As well, H₂S production was increased by testosterone but inhibited by PPG and BCA (Bucci et al. 2009).

It has been shown that vascular endothelial growth factor (VEGF) stimulated H₂S production in cultured human umbilical vein endothelial cells (HUVECs) (Papapetropoulos et al. 2009). This effect of VEGF can be attributed to increased intracellular calcium and calcium-activated calmodulin in ECs. Whether VEGF affects CSE gene expression is unknown. Hassan et al. showed that platelet-derived growth factor-BB (PDGF-BB) upregulated CSE mRNA and protein levels in mesangial cells (Hassan et al. 2012). Hypoxia is another factor that leads to increased H₂S production in human placenta and rat liver, uterus, and fetal membranes (Patel et al. 2009). CSE gene expression in the vascular system may also be inhibited by insulin (Wang 2004).

CSE activities in vascular SMCs were increased by NO (Zhao et al. 2001). After incubating rat aortic tissue homogenates with a NO donor for 90 min, H₂S generation from the homogenates was significantly increased. One of the underlying mechanisms for this effect of NO is the upregulation of CSE expression, which was confirmed 6 h after incubating cultured vascular SMCs with a NO donor (Zhao et al. 2001). Similar observations were made by Patel et al. that NO donors increased both the expression and activity of CSE proteins in rat fetal membranes (Patel et al. 2009).

The regulation of CSE gene expression mostly occurs at the CSE promoter site. The specific protein 1 (Sp1) transcription factor, nuclear factor erythroid-2-related factor-2 (Nrf2), and farnesoid X receptor (FXR) ligand can all bind to the CSE promoter, hence stimulating CSE transcription (Zhang et al. 2011; Yang et al. 2011; Hassan et al. 2012; Renga et al. 2009). This transcriptional regulation mechanism can explain the inhibitory effect of microRNA 21 (miR-21) on CSE expression since miR-21 directly repressed the expression of Sp1 (Yang et al. 2012a, b). In contrast, increased CSE expression by TNF- α resulted from TNF- α -stimulated Sp1 binding to the CSE promoter (Sen et al. 2012a). In the case of Nrf2, it binds to an antioxidant-responsive element (ARE) to mediate the targeted gene transcription after Nrf2 is translocated into the nucleus. The CSE promoter contains an ARE sequence and this provides an oxidative stress-sensitive mechanism for regulating CSE expression. Hassan et al. showed that PDGF-BB-induced CSE expression in mesangial cells is inhibited by co-treatment with antioxidants or by Nrf2 knockout. Furthermore, stabilization of Nrf2 protein upregulated CSE protein expression (Hassan et al. 2012). The evidence for the interaction of FXR with the CSE

promotor was derived from HepG2 cells. The human CSE gene contains an FXR-responsive element in its 5'-flanking region. Treatment of HepG2 cells with an FXR ligand increased CSE expression and mutation of FXR blocks FXR ligand-induced CSE expression (Renga et al. 2009).

4 H₂S and SMCs

4.1 H₂S and SMC Apoptosis, Proliferation, and Migration

Abnormal SMC proliferation and apoptosis are among the causative factors for vascular remodeling. H₂S regulation of SMC proliferation and apoptosis has been extensively studied (Fig. 1). We previously demonstrated that exogenously applied H₂S or endogenous H₂S derived from overexpressed CSE gene inhibits proliferation and induces apoptosis of human aorta SMCs by activating ERK and caspase 3 (Yang et al. 2004, 2006). Du et al. also found that H₂S dose-dependently suppressed the proliferation of rat SMCs through the MAPK pathway (Du et al. 2004). Region-specific chromatin remodeling of MAPK signaling pathway-associated genes such as Ntf3, PcnA, and Pdgfr α can be regulated by H₂S, and Brg1 acts as a switch to turn these genes “on” in a spatially and temporally specific manner to inhibit SMC proliferation (Li et al. 2013). The hypoxia-induced proliferation of pulmonary artery SMCs is the main cause of pulmonary arterial hypertension, and H₂S was demonstrated to inhibit CoCl₂-induced pulmonary arterial SMC proliferation by the upregulation of cyclooxygenase-2 and prostacyclin (Li et al. 2014). Indeed, SMCs from CSE knockout mice displayed an increased proliferation rate in vitro and in vivo, and these cells were more susceptible to apoptosis induced by an oxidative stress inducer (H₂O₂) or a high dose of H₂S (100 μ M) (Yang et al. 2010). CSE knockout mice exhibited decreased endogenous H₂S level in the cardiovascular system and impaired endothelium-dependent vasorelaxation and age-dependent hypertension (Yang et al. 2008). The altered SMC proliferation in CSE knockout mice provides new insight into the pathogenesis of hypertension and underscores the importance of H₂S in homeostatic control of vascular integrity. Li et al. further found that the proliferation of cultured vascular SMCs isolated from wild-type mice was inhibited, but that from CSE gene knockout mice increased, by estrogen treatments, indicative of the interaction of H₂S and estrogen in regulating SMC proliferation (Li et al. 2012). In a rat model of pulmonary hypertension and pulmonary artery structural remodeling induced by high pulmonary blood flow, the inhibition of endogenous H₂S production by PPG markedly decreased the rate of pulmonary artery SMC apoptosis but supplements of H₂S donor increased pulmonary artery SMC apoptosis, as demonstrated by positive TUNEL staining (Li et al. 2009). Consistent with these in vivo findings, Baskar et al. observed that S-diclofenac, a novel molecule containing an H₂S-releasing dithiolthione moiety, stabilized p53 and induced the expressions of downstream proteins, such as p21, p53AIP1, and Bax, to repress SMC proliferation (Baskar et al. 2008).

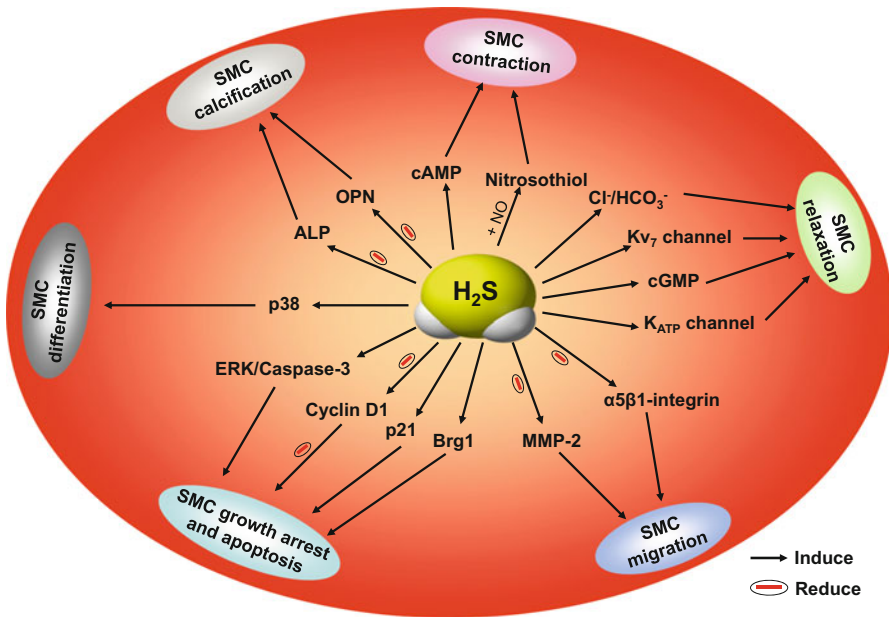


Fig. 1 H₂S signaling in SMC functions. H₂S maintains SMC differentiation by inducing p38 MAPK. H₂S stimulates SMC growth arrest and induces SMC apoptosis by activating ERK/caspase 3, p21, and/or Brg1 pathways but inactivating cyclin D1. H₂S also attenuates SMC migration by inhibiting the expressions of MMP-2 and $\alpha 5\beta 1$ -integrin. By activating Cl⁻/HCO₃⁻, K_{v7} channel, cGMP, and/or K_{ATP} channel, H₂S induces SMC relaxation, while H₂S mediates SMC contraction by enhancing cAMP production and interacting with NO to form nitrosothiol. H₂S also reduces SMC calcification by inhibiting the activities of ALP and OPN. Abbreviation used in this figure: ALP, alkaline phosphatase; Brg1, Brahma-related gene 1; cAMP, adenosine 3',5'-cyclic monophosphate; cGMP, cyclic guanosine monophosphate; ERK, extracellular-signal-regulated kinase; H₂S, hydrogen sulfide; K_{ATP} channel, ATP-sensitive potassium channel; K_{v7} channel, voltage-gated potassium channel subunit K_{v7}; MMP-2, matrix metalloproteinase 2; NO, nitric oxide; OPN, osteopontin; SMC, smooth muscle cell

Altered SMC proliferation and apoptosis have been considered as key events in vascular injury in diseases such as atherosclerosis and restenosis after invasive intervention. CSE knockout mice fed with atherogenic diet developed early fatty streak lesions in the aortic root and increased aortic intimal proliferation (Mani et al. 2013). Treatment of CSE knockout mice with NaHS inhibited the accelerated atherosclerosis development. By using a rat model of vascular remodeling induced by balloon injury, Meng et al. found that CSE expression and H₂S production are significantly reduced in the blood vessel during the development of neointimal formation after balloon injury and the administration of NaHS attenuated the development of neointimal hyperplasia by inhibiting SMC proliferation (Meng et al. 2007). NaHS induced a significant reduction in cell proliferation in the neointima. We also noticed that CSE deficiency in mice led to increased neointima formation in carotid arteries 4 weeks after ligation, which were attenuated by NaHS

administration (Yang et al. 2012a, b). All these data suggest that endogenous H₂S is critical for the inhibition of SMC proliferation during neointimal formation. In response to vascular injury, SMCs will first migrate from the tunica media to the intima, where they contribute to neointima formation (Thyberg 1998). Exogenously applied NaHS significantly inhibited SMC migration, and SMCs and aortic explants isolated from CSE knockout mice exhibited more migration and outgrowth compared with that from wild-type mice. SMCs became more elongated and spread in the absence of CSE. The interactions among $\alpha 5\beta 1$ -integrin, fibronectin, and MMP-2 promoted CSE deficiency-enhanced SMC adhesion and migration (Yang et al. 2012a, b). These studies provide further insight into the pathogenesis of proliferative cardiovascular disorders and underscore the protective effects of H₂S in maintaining vascular homeostasis.

4.2 H₂S and SMC Phenotype Modulation

Vascular SMCs are highly specialized cells whose contractile status regulates blood vessel tone, blood pressure, and blood flow distribution. In contrast to cardiac and skeletal muscle cells which exit cell cycle and undergo final differentiation, vascular SMCs are highly plastic and can switch their phenotypes between contractile and synthetic phenotypes in response to extracellular stimuli or damage (Owens 2007). SMC phenotypic switching is believed to play a key role in many cardiovascular diseases, such as hypertension, atherosclerosis, coronary heart diseases, postangioplasty restenosis, and transplantation arteriopathy (Thyberg 1998). The H₂S-regulated SMC phenotypic switch has been explored (Fig. 1). CSE expression and H₂S production were reduced in proliferated SMCs compared with differentiated SMCs in a Sp1-dependent manner (Yang et al. 2011). In the presence of 10 % serum, H₂S strikingly induced the expressions of SMC differentiation maker genes in proliferative human aorta SMCs. In addition, H₂S-stimulated SMC differentiation maker gene expressions were reversed by co-treatment of the cells with SB203580, a p38 MAPK inhibitor. In the absence of serum, exogenously applied H₂S did not change the expressions of SM-MHC and calponin in SMCs. However, the inhibition of endogenous H₂S production by PPG significantly repressed the expressions of SM myosin heavy chain and calponin (Yang et al. 2011). Induction of SMC differentiation maker gene expressions by H₂S in the presence of, but not absence, serum suggests that H₂S may coordinate the expression of proliferative and contractile proteins to induce differentiated SMC phenotype. All these data indicate that CSE/H₂S system is essential for the maintenance of SMC differentiation.

4.3 H₂S and SMC Relaxation and Contraction

H₂S-induced vasorelaxation is a well-known vascular event, and the most widely characterized cellular target for H₂S in SMCs is the ATP-sensitive K⁺ (K_{ATP})

channels (Fig. 1). In the vascular system, specifically in SMCs, K_{ATP} channels contribute significantly toward vasodilation in response to various vasoactive substances. The opening of K_{ATP} channels hyperpolarizes cell membrane and inactivates voltage-dependent L-type Ca²⁺ channels, leading to cell relaxation and blood vessel dilation by reducing intracellular free Ca²⁺ concentration (Wang 2012a, b). It has been previously shown that H₂S at physiologically relevant concentrations induced the relaxation of rat aortic tissue and transient reduction of blood pressure, and these vascular effects of H₂S were mediated by a direct stimulation of K_{ATP} channels and subsequent hyperpolarization of aortic SMCs (Zhao et al. 2001; Cheng et al. 2004; Wang 2014; Sun et al. 2011; Siebert et al. 2008). Using the whole-cell and single-channel patch-clamp technique, Tang et al. demonstrated that H₂S activates K_{ATP} channels and hyperpolarized cell membrane in rat mesenteric artery SMCs (Tang et al. 2005, 2010). H₂S enhanced the amplitude of whole-cell K_{ATP} currents and increased the open probability of single K_{ATP} channels. Furthermore, inhibition of endogenous H₂S production with PPG reduced whole-cell K_{ATP} currents. H₂S also causes the relaxation of human airway SMCs via stimulating sarcolemmal K_{ATP} channels (Fitzgerald et al. 2014). Other studies proved that H₂S interacts with the SUR subunits of the K_{ATP} channel complex to cause the channel to open, and the sulfhydryl groups located on the extracellular surface of the SUR subunits are potential targets for H₂S S-sulfhydration (Jiang et al. 2010). The deletion of extracellular cysteine 6 or 26 of SUR1 subunits caused the loss of channel sensitivity to H₂S. It appears that H₂S firstly breaks the disulfide bond between cysteine 6 and cysteine 26 and then caused their S-sulfhydrations. In addition to K_{ATP} channels, H₂S is reported to activate Kv7 voltage-gated potassium channels (particularly the Kv7.4 subtype) in SMCs, and the activation of Kv7 channel mediates a significant part of the vasorelaxing effects of H₂S (Martelli et al. 2013).

H₂S may induce SMC relaxation by altering intracellular pH. H₂S has been shown to decrease intracellular pH in a dose-dependent manner. Ionic exchangers, including Na⁺/H⁺ and Cl⁻/HCO₃³⁻ and Ca²⁺ ATPase, maintain the resting pH in SMCs between 7.1 and 7.2. Preincubation of SMCs with a selective inhibitor of Cl⁻/HCO₃³⁻, but not Na⁺/H⁺ exchanger inhibitor, prevented the drop of intracellular pH and vasorelaxation caused by H₂S, suggesting that Cl⁻/HCO₃³⁻ exchanger is involved in H₂S-induced relaxation in SMCs (Lee et al. 2007). Intracellular acidosis could activate K_{ATP} channel in SMCs and decrease vascular tone. As such, H₂S-induced opening of K_{ATP} channels may be partially triggered by intracellular acidification (Liu et al. 2011).

H₂S may also cause vasorelaxation by increasing cGMP level in SMCs. Incubation of cultured rat aortic SMCs with NaHS led to a concentration-dependent increase in cGMP levels. The NaHS-induced rise in cGMP was evident as early as 1 min, reached a maximum at 3 min, and remained elevated for at least 10 min. Blockade of CSE activity by PPG or BCA or knockdown of CSE mRNA by siRNA resulted in a significant reduction of cGMP accumulation (Bucci et al. 2010). In contrast, overexpression of CSE elevated intracellular cGMP level. Vascular tissue levels of cGMP in CSE knockout mice were lower than those in wild-type control

mice. Intracellular cGMP levels reflect the balance between the rate of cGMP synthesis via guanylyl cyclases and breakdown by phosphodiesterases. It appears that H₂S does not activate soluble guanylate cyclase, because H₂S-induced vasorelaxation is not inhibited by a soluble guanylate cyclase inhibitor (Coletta et al. 2012). Further studies showed that H₂S acts as an endogenous inhibitor of phosphodiesterase. In a cell-free assay, Bucci et al. demonstrated that NaHS at 10–30 nM significantly inhibits phosphodiesterase activity and causes a reduction in the breakdown of 5'-GMP (Bucci et al. 2012). H₂S also ameliorated the reduction in cGMP levels brought about by overexpression of phosphodiesterase 5A (Bucci et al. 2010, 2012). Because phosphodiesterases are involved in the degradation of both cAMP and cGMP, the researchers did not test the cellular effects of H₂S on cAMP level in SMCs. In contrast, another study by Lim et al. proved that NaHS significantly reverses forskolin-induced cAMP accumulation in SMCs (Lim et al. 2008). cAMP plays important roles in the regulation of mature contractile phenotype in SMCs. The effect of H₂S on SMC relaxation is actually biphasic depending on its concentration (Kubo et al. 2007; Liu et al. 2011). At higher level, H₂S produces vasorelaxation effect, while it induces vasoconstriction at lower concentration (Ali et al. 2006). To this end, the researchers observed that NaHS at a concentration range of 10–100 μM concentration-dependently reverses the vasodilation caused by isoprenaline and salbutamol (two β-adrenoceptor agonists) and forskolin (a selective adenylyl cyclase activator) in phenylephrine-precontracted rat aortic rings (Lim et al. 2008). Therefore, the authors proposed that the contractile effect of H₂S observed in isolated rat aorta is, at least partially, associated with reducing cAMP level in SMCs. In addition, H₂S may react with NO to form a compound, probably nitrosothiol, which leads to less NO bioavailability (Whiteman et al. 2006).

4.4 H₂S and SMC Calcification

Vascular calcification is implicated in the pathogenesis of various vascular diseases and resulted from passive precipitation of calcium and phosphate. Vascular calcification is now considered to be an active, regulative process similar to osteogenesis (McCarty and DiNicolantonio 2014). Calcified vessels have decreased capacity for vasodilatation and increased stiffness and promote a form of thrombus and atherosclerotic plaque rupture. Osteoblastic differentiation of SMCs is involved in the pathogenesis of vascular calcification. H₂S has been shown to ameliorate SMC calcification. In a rat vascular calcification model induced by the administration of vitamin D3 plus nicotine, aortic CSE expression and H₂S content were significantly reduced (Wu et al. 2006). Supplement of NaHS significantly reduced aortic calcium mineral deposits, OPN mRNA expression, and ALP activity, pointing to a regulatory role of CSE/H₂S pathway in the pathogenesis of vascular calcification. By using cultured SMCs, Zavaczki et al. explored the roles of H₂S in phosphate-induced osteoblastic transformation and mineralization (Zavaczki et al. 2011). H₂S inhibited calcium deposition in the extracellular matrix and suppressed the

induction of the genes involved in osteoblastic transformation of SMCs, including alkaline phosphatase, osteocalcin, and Cbfa1. H₂S also prevented phosphate uptake and phosphate-triggered upregulation of the sodium-dependent phosphate cotransporter. H₂S, regardless of its exogenous or endogenous origin, is a potent inhibitor of phosphate-induced calcification and osteoblastic differentiation of SMCs. In contrast, silencing CSE by siRNA and inhibition of CSE activity by PPG attenuated receptor activator of nuclear factor κ -B ligand-induced tartrate-resistant acid phosphatase type 5 activities and pit formation in RAW264.7 cells. Moreover, knockdown of CSE suppressed the expression of osteoclast differentiation markers. A large-scale proteomics study also identified that CSE acts in early stages of osteoclastogenesis (Itou et al. 2014). These results suggest that CSE is a potent inducer of calcium resorption in inflammatory cells.

5 H₂S and ECs

5.1 H₂S and EC Proliferation and Angiogenesis

H₂S significantly stimulates endothelial cell growth and angiogenesis (Wang 2012a, b; Polhemus and Lefer 2014; Liu et al. 2011) (Fig. 2). Several groups have demonstrated that H₂S significantly promotes cell growth and capillary-like structure formation of cultured ECs (Altaany et al. 2013; Papapetropoulou et al. 2009; Liu et al. 2011). Supplement of exogenous H₂S increased cellular infiltration and neovascularization in mouse Matrigel, enhanced the length of vascular network in the chick chorioallantoic membrane model, and promoted the formation of collateral vessels in ischemic hind limbs in rats (Papapetropoulou et al. 2009; Köhn et al. 2012a; Hofer 2007). A delayed wound healing was found in CSE knockout mice when compared with wild-type littermates (Papapetropoulou et al. 2009). Endothelial progenitor cells are a population of rare cells that circulate in the blood with the ability to differentiate into ECs. H₂S was shown to improve endothelial progenitor cell function in diabetic wound healing of type 2 diabetic mice. Conversely, PPG treatment reduced progenitor cell function and delayed wound healing (Liu et al. 2014). Under hypoxic condition, H₂S induced endothelial proliferation and migration by promoting VEGF and HIF-1 α expression and increasing HIF-1 α -binding activity (Liu et al. 2010). In contrast, H₂S decreased cell proliferation and capillary tube formation of EA.hy926 cells under hypoxia by inhibiting the expression of VEGF and HIF-1 α (Wu et al. 2012). The discrepancy of the aforementioned observations may be due to different hypoxic models, H₂S concentrations, and cell types. Pupo et al. showed that NaHS at 10 μ M did not exert any effect on cell migration and proliferation of normal human microvascular ECs (HMVECs) (Pupo et al. 2011). This apparent discrepancy may be ascribed to tissue specificity. Different from HMVECs, H₂S donors failed to induce cell proliferation but promote their migration of ECs obtained from human breast carcinoma (B-TEC). B-TECs pretreated with PPG showed drastically reduced migration

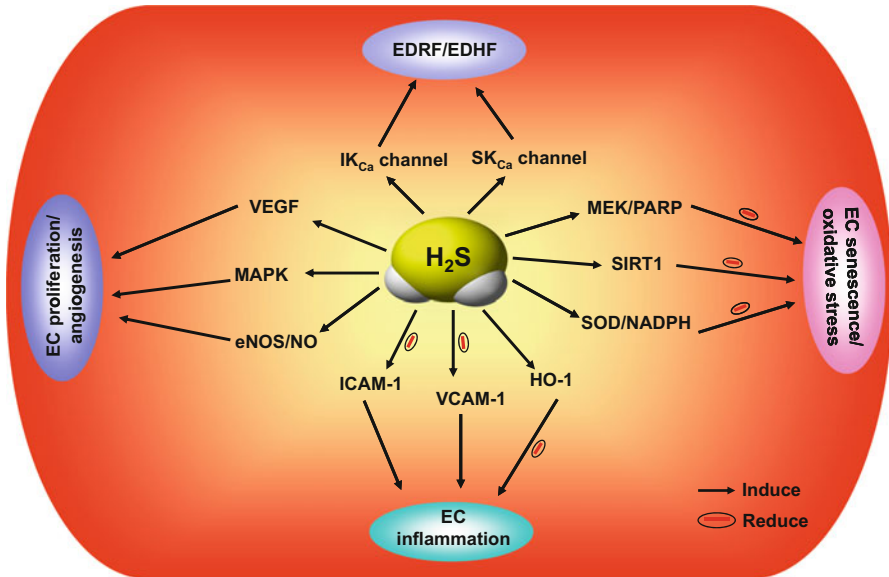


Fig. 2 H₂S signaling in EC functions. H₂S stimulates EC proliferation and angiogenesis by activating VEGF, MAPK, and/or eNOS/NO pathways. H₂S attenuates EC inflammation by inhibiting the expressions of ICAM-1 and VCAM-1 but stimulating HO-1. H₂S lowers EC senescence and oxidative stress by activating MEK/PARP, SIRT1, and/or SOD/NADPH pathways. H₂S acts as EDRF and EDHF at least through IK_{Ca} channel and SK_{Ca} channel. Abbreviation used in this figure: EDHF, endothelium-derived hyperpolarizing factor; EDRF, endothelium-derived relaxing factor; eNOS, endothelial nitric oxide synthase; HO-1, heme oxygenase-1; ICAM-1, intercellular adhesion molecule 1; IK_{Ca} channel, intermediate-conductance, calcium-activated potassium channel; MAPK, mitogen-activated protein kinase; NADPH, nicotinamide adenine dinucleotide phosphate-oxidase; NO, nitric oxide; PARP, poly (ADP-ribose) polymerase; SIRT1, sirtuin-1; SK_{Ca} channel, small-conductance calcium-activated potassium channel; SOD, superoxide dismutase; VCAM-1, vascular cell adhesion molecule 1; VEGF, vascular endothelial growth factor

induced by VEGF (Pupo et al. 2011). The authors concluded that H₂S plays a role in proangiogenic signaling of tumor-derived but not normal human ECs.

In addition to the direct stimulatory effect on EC growth and angiogenesis, H₂S was also reported to ameliorate stress-induced cell death of ECs. Exposure of primary human umbilical vein endothelium cells (HEVECs) to a high concentration of glucose (25 mM) resulted in the induction of apoptosis, but pretreatment with NaHS (50 μM) for 30 min attenuated the pro-apoptotic effect of 25 mM glucose. Further investigation of the apoptotic mechanisms in the cells demonstrated that high glucose upregulated the ratio of Bax/Bcl-2, activated caspase-3, increased the levels of reactive oxygen species and malondialdehyde, and suppressed superoxide dismutase activity (Guan et al. 2012). All these effects of glucose could be prevented by pretreatment with 50 μM NaHS. Pretreatment with NaHS (0.05–0.1 mM) attenuated methionine- or oxidized low-density lipoprotein-induced oxidative stress and cell death of ECs (Tyagi et al. 2009; Muellner et al. 2009).

Exposure of bEnd3 microvascular ECs to elevated extracellular glucose induced mitochondrial formation of ROS, and H₂S replacement protected against ROS formation, improved endothelial metabolic state, and maintained normal endothelial function (Suzuki et al. 2011). H₂S was also found to preserve the activities and protein levels of the antioxidant enzymes, superoxide dismutase, catalase, glutathione peroxidase, and glutathione-S-transferase in H₂O₂-exposed ECs (Wen et al. 2013). Zofenoprilat is a sulfhydryl-containing angiotensin-converting enzyme inhibitor, which can enhance CSE-dependent availability of H₂S. Zofenoprilat was shown to exert a protective effect on doxorubicin-induced endothelial damage without affecting its antitumor efficacy (Monti et al. 2013).

5.2 H₂S and EC Senescence and Oxidative Stress

Growing evidence shows that the progress of vascular aging alters cardiovascular function and subsequently increases the risk of cardiovascular diseases. Vascular aging has been largely associated with senescence of the vascular endothelium. Several lines of evidence point to the implication of H₂S signaling in the process of endothelial senescence (Fig. 2). Calorie restriction is reported to decelerate biological aging process, resulting in longer maintenance of youthful health and an increase in both median and maximum life span. Calorie restriction maintained normal H₂S level in vascular tissues from rats during aging, suggesting a protective role of H₂S in vascular aging (Predmore et al. 2010). Oxidative stress is a driving factor for vascular aging. H₂O₂ treatments of HUVECs lead to high rate of senescent cells, which was attenuated by NaHS incubation putatively through the modulation of SIRT1 activity (Suo et al. 2013). H₂S also improved the function of senescent HUVECs. Zhao et al. found that H₂S attenuates cellular senescence and DNA damage in HUVECs by MEK1 S-sulfhydration and PARP-1 activation (Zhao et al. 2014). In the presence of H₂S, activated PARP-1 recruits XRCC1 and DNA ligase III to DNA breaks to mediate DNA damage repair. AP39, a mitochondria-targeted H₂S donor, was shown to exert a concentration-dependent effect on mitochondrial activity in bEnd.3 murine microvascular ECs, as evidenced by the stimulation of mitochondrial electron transport and cellular bioenergetic function. Furthermore, AP39 pretreatment protected against glucose oxidase-induced mitochondrial DNA damage (Szczeny et al. 2014). A recent study showed that H₂S increases the life span of *Caenorhabditis elegans*. The life span-prolonging and health-promoting effects of H₂S in *C. elegans* are likely due to the antioxidant action (Qabazard et al. 2014). Similarly, compared with the lower passage of ECs, the higher passage of ECs had lower SOD activity and higher H₂O₂ level, whereas NaHS pretreatment reversed the changes of SOD activity and H₂O₂ level, indicating that H₂S delays senescence of HUVECs through lessening oxidative stress (Qi et al. 2012; Muellner et al. 2009). By using porcine pulmonary arterial ECs, Muzaffar et al. demonstrated that H₂S inhibited superoxide formation and upregulation of NADPH oxidase through the adenylyl cyclase-PKA pathway

(Muzaffar et al. 2008). In spite of these findings, the *in vivo* role of H₂S in regulating EC senescence and vascular aging remains unclear.

5.3 H₂S and EC Inflammation

The regulatory role of H₂S in inflammation involves the endothelium and its interaction with leukocytes (Whiteman and Winyard 2011; Zanardo et al. 2006). H₂S donors (NaHS and Na₂S) inhibited but PPG promoted aspirin-induced leukocyte adherence in mesenteric venules via the activation of K_{ATP} channels (Zanardo et al. 2006). H₂S also suppressed leukocyte infiltration in an air pouch model and carrageenan-induced paw edema, implicating a protective role of H₂S in acute inflammation by acting at the leukocyte–endothelium interface (Perna et al. 2013). Further study demonstrated that H₂S treatment of ECs decreased the expression of MCP-1, intercellular adhesion molecule-1 (ICAM-1), and vascular cell adhesion molecule-1 (VCAM-1) at the mRNA and protein levels. In an *in vitro* model entailing monocyte adhesion to an endothelial monolayer, H₂S prevented the increase in monocyte adhesion induced by tumor necrosis factor- α (TNF- α). Pan et al. also showed that H₂S dose-dependently suppressed TNF- α -induced mRNA and protein expressions of ICAM-1 and VCAM-1 in HUVECs, possibly through the upregulation of HO-1 (Pan et al. 2011). By using another H₂S donor, SPRC, the same group found that H₂S exerts anti-inflammatory effects on TNF- α -stimulated ECs through scavenging ROS, inhibiting JNK1/2/NF- κ B pathways, and attenuating adhesion molecule expression (Pan et al. 2012). NaHS was shown to reduce oxLDL-induced foam cell formation in macrophages and TNF- α -stimulated ICAM-1 expression in HUVECs (Zhao et al. 2011; Wang et al. 2013b). In contrast, Choi et al. showed that Korean red ginseng extracts inhibit the expression of inflammatory mediators, including IL-8 and IL-6, via reduced CSE expression and H₂S production in ECs (Choi et al. 2012). Supplement of exogenously applied H₂S reversed the Korean red ginseng extract-improved inflammation status in ECs.

H₂S protects vascular tissues from atherogenic damage by inhibiting adhesion molecule expression and suppressing monocyte adhesion to the activated endothelium (Wang et al. 2009). In spontaneously hypertensive rat (SHR), the expression of ICAM-1 and NF-kappaB p65 protein in aortic ECs was significantly higher, and NaHS treatment reduced blood pressure in SHR rats and downregulated the expressions of ICAM-1 and NF-kappaB p65 in aortic ECs. On the other hand, inhibition of H₂S production enhanced the expressions of ICAM-1 and NF-kappaB p65 protein in aortic ECs (Jin et al. 2008), suggesting that H₂S might attenuate the development of hypertension by suppressing endothelial inflammation reactions.

5.4 Interplay Between H₂S and NO in ECs

Both H₂S and NO are major gasotransmitters produced in ECs, and growing evidence showed that the “cross talk” between NO and H₂S mediates the

cardioprotective effect of H₂S (Jamroz-Wisniewska et al. 2014). Predmore et al. first showed that H₂S stimulated eNOS phosphorylation and NO production in ECs, and pharmacological inhibition of Akt, the kinase responsible for eNOS Ser 1177 phosphorylation, attenuated the stimulatory effect of H₂S on NO production (Predmore et al. 2011). Al Tanny et al. confirmed that H₂S promotes NO production in ECs via the activation of a cascade of phosphorylation events, starting from p38 MAPK to Akt to eNOS (Altaany et al. 2013). Deficiencies in H₂S signaling can directly impact on processes regulated by NO (Coletta et al. 2012). H₂S promotes EC tube formation, proliferation, and angiogenesis by both NO-dependent and NO-independent mechanisms. H₂S also modulates eNOS via S-sulfhydration and prevents eNOS coupling collapse and thus increases NO and decreases ONOO⁻ and O₂ levels in ECs (Al Taany et al. 2014). Exposure of ECs to H₂S increased intracellular cGMP in a PI3K/Akt and NO-dependent manner, and NO and H₂S are mutually required for the physiological control of vascular function (Coletta et al. 2012). H₂S may stimulate NO production by different mechanisms. Kida et al. showed that NaHS dose-dependently increased NO production in cultured ECs by releasing calcium from the intracellular store in endoplasmic reticulum. NaHS-induced eNOS phosphorylation and NO production were abolished by the ryanodine receptor inhibitor, inositol 1,4,5-triphosphate receptor inhibitor, and calcium chelator but not by the PI3K/Akt inhibitor and the absence of extracellular calcium (Kida et al. 2013). NaHS was also shown to increase eNOS expression and NO production in rat corpus, suggesting H₂S could exert its pro-erectile effects by augmenting NO pathway. In contrast, Na₂S, another H₂S donor, was shown to reduce the level of phospho-eNOS (serine 1177) and inhibit eNOS activity in cultured mouse aortic ECs (Chai et al. 2014). The discrepancy may be due to the difference of cell type and H₂S dose used. Further elucidation of the H₂S–NO relationship in the vascular biology would provide more insight into the vasodilator function of H₂S and improve our understanding of the pathogenic mechanisms for cardiovascular diseases.

5.5 H₂S Acts as an Endothelium-Derived Relaxing Factor (EDRF) and an Endothelium-Derived Hyperpolarizing Factor (EDHF)

In response to a variety of chemical and physical stimuli, ECs produce and release various vasoactive factors, including EDRF and EDHF. Both EDRF and EDHF relax vascular SMCs, causing blood vessel to expand in diameter (Wang 2012a, b). NO is a well-described EDRF, but other EDRFs are also produced and released from the endothelium. H₂S shares many features with NO. Accumulating evidence supports the concept that H₂S acts as both EDRF and EDHF (Baragatti et al. 2013; Han et al. 2013; Skovgaard et al. 2011; Wang 2009) (Fig. 2). Endogenously generated H₂S induces vasorelaxation in part through an endothelium-dependent mechanism (Yang et al. 2008; Zhao et al. 2001). The vasorelaxation elicited by H₂S is greater in small resistance arteries as compared with that in larger conduit arteries such as the aorta (Tang et al. 2013). Knocking out CSE expression attenuated

acetylcholine-induced membrane hyperpolarization of the isolated ECs and led to membrane depolarization of the whole vascular tissues. Different from EDRFs, the unique property of EDHF is its sole effect on hyperpolarizing vascular SMCs so as to close voltage-dependent calcium channels (Wang 2012a, b). The effect of EDHF is mainly mediated by the opening of small-conductance and/or intermediate-conductance K_{Ca} channels (Mustafa et al. 2011; Wang 2012a, b). Using electrophysiological microelectrode technique, Tang et al. demonstrated that the unique EC dependence of H_2S -induced SMC hyperpolarization was linked to the opening of IK_{Ca} and/or SK_{Ca} channels (Tang et al. 2013). The identification of H_2S as an EDRF/EDHF will not only help better understand the mechanisms underlying endothelium-dependent vasorelaxation of different types of vascular tissues but also shed light on devising novel therapeutic agents to deal with specific cardiovascular diseases.

6 H_2S and Periadventitial Adipose Tissue (PAT)

Adventitia is the outermost connective tissue covering a blood vessel, also called the tunica adventitia or the tunica externa. PAT, the major part of adventitia and defined as the accumulation of adipocytes around vascular structures, can be found in the proximity of virtually all blood vessels (Gollasch 2012). PAT is composed of various cells such as adipocytes, fibroblasts, and macrophages and can release various active agents, e.g., adipocyte-derived relaxing factor (ADRF), which play important roles in modulating the vascular tone. Recent studies have shown that H_2S can be generated from PAT and act as an ADRF, contributing to vascular relaxation (Fang et al. 2009). Immunohistochemical staining revealed the presence of CSE protein in PAT. In isolated PAT, H_2S production was reduced in an age-dependent manner. The CSE/ H_2S pathway was upregulated in PAT as a compensatory mechanism against the elevated blood pressure in hypertension (Bełtowski 2013). In consistent with this study, Köhn et al. showed that the inhibition of CSE activity by PPG reversed the anti-contractile effect of PAT in rat aorta (Köhn et al. 2012a, b). Statins are well-known drugs to reduce plasma LDL cholesterol, improve endothelial function, ameliorate oxidative stress, and maintain coagulation–fibrinolysis balance. It was recently found that atorvastatin, one kind of statins, increased H_2S production in PAT and H_2S mediated the protective role of statins in the cardiovascular system. Atorvastatin augmented the anti-contractile effect of PAT, most likely in an H_2S - and K_{ATP} channel-dependent manner, because its effect was abolished by CSE inhibitor PPG and K_{ATP} channel blocker glibenclamide (Wójcicka et al. 2011). A statin-induced increase in H_2S production may also contribute to the antiatherogenic effect of statins since H_2S inhibits platelet aggregation, vascular SMC proliferation, LDL oxidation, and local inflammatory reaction.

7 H₂S and Blood Vessel-Related Disorders

Blood vessel diseases are also called peripheral vascular diseases or artery diseases, including hypertension, atherosclerosis, aortic aneurysm, etc. H₂S plays an important protective role in various blood vessel diseases (Wang 2012a, b; Yang 2011).

7.1 H₂S and Hypertension

Hypertension or high blood pressure is a chronic medical condition in which the blood pressure is elevated. Hypertension results from a complex interaction of vasoactive factors with various types of cells in blood vessel walls. Recent studies have shown that CSE/H₂S system produces antihypertensive effects in different hypertensive models. An acute intravenous bolus injection of NaHS caused a transient fall of mean arterial blood pressure in anesthetized rats (Zhao et al. 2001). Exogenous H₂S administration lowers blood pressure and prevents the hypertrophy of intramyocardial arterioles and aortic thickening of spontaneously hypertensive rats (SHR) (Du et al. 2003; Zhao et al. 2008). Injection of a single dose of GYY4137 (a H₂S donor) alleviated L-NAME-induced hypertension in rats, and chronic treatment with GYY4137 successfully reduced blood pressure of SHR (Li et al. 2008a). ACS14, another H₂S-releasing donor, reduced blood pressure in buthionine sulfoximine-induced hypertensive rats (Rossoni et al. 2010). Inhibition of H₂S production by PPG worsened hypoxic pulmonary hypertension in rats. Direct evidence also demonstrated that complete deficiency of CSE in mice markedly reduced endogenous H₂S levels in the vascular system and led to age-dependent development of hypertension (Yang et al. 2008). Preeclampsia is a disorder of pregnancy characterized by high blood pressure and contributes to maternal and fetal morbidity and mortality worldwide. The cause and pathogenesis of preeclampsia has yet to be definitively uncovered. Wang et al. recently showed that plasma H₂S and CSE expression in the placenta were reduced in pregnancies complicated by preeclampsia in comparison with gestational age-matched controls (Wang et al. 2013a). Inhibition of CSE activity induces hypertension and causes placental abnormalities in pregnant mice owing to the inhibition of H₂S production. These discoveries suggest that a dysfunctional CSE/H₂S pathway may contribute to the pathogenesis of preeclampsia and targeting at the CSE/H₂S system would be an effective therapy for preeclampsia (Wang et al. 2013a, b).

7.2 H₂S and Atherosclerosis

Atherosclerosis is a chronic progressive pathological process in large- and medium-sized arteries, caused by the buildup of fatty/cholesterol plaques on the ECs of arteries, endothelial inflammation, and SMC proliferation. H₂S plays an anti-atherosclerotic role, and its deficiency leads to the development and progression of atherosclerosis (Zhang et al. 2012a, b; Mani et al. 2013, 2014; Xu et al. 2014;

Wang et al. 2009). Plasma H₂S level and aortic CSE activity was decreased in apolipoprotein E (ApoE) knockout mice with advanced atherosclerosis. Treatment of ApoE knockout mice with NaHS resulted in reduced atherosclerosis plaque, while inhibition of CSE activity enlarged plaque size (Wang et al. 2009). GYY4137, a H₂S donor, was also shown to decrease aortic atherosclerotic plaque formation and partially restored aortic endothelium-dependent relaxation in ApoE knockout mice (Li et al. 2008a; Liu et al. 2013; Qiao et al. 2010). Another study showed that onion extracts boosted endogenous production of H₂S and lessened atherosclerotic lesions in rats (Zhang et al. 2012a, b). In a high-fat and high-vitamin D1 diet-induced atherosclerotic rat model, H₂S was shown to slow down the development of atherosclerosis by improving the damage of vessels and inhibiting the expression of VEGF (Zhang et al. 2012a, b). Treatment with NaHS significantly inhibited arterial restenosis following balloon angioplasty in rabbits by reducing the intimal area and the intima/media ratio, while PPG treatment had a tendency to worsen the severe restenosis (Ma et al. 2012). Whether CSE deficiency impacts on the development of atherosclerosis has been directly addressed by knocking out the CSE gene in high-fat-fed mice. Decreased endogenous H₂S production in CSE knockout mice leads to the vascular remodeling and early development of atherosclerosis (Mani et al. 2013). The atherosclerotic plaque development is rescued by H₂S donor NaHS via reducing vessel intimal proliferation and inhibiting adhesion molecule expression. In contrast, antihypertensive (hydralazine), antioxidant (N-acetylcysteine), or lipid-lowering (ezetimibe) agents have no effect on high-fat-diet-induced plaque formation in CSE knockout mice, implying that hypertension, higher oxidative stress, and abnormal lipid profile do not play major roles in atherosclerosis development in CSE knockout mice. In addition, the knockout of CSE from ApoE knockout mice accelerated plaque formation even under normal diet, indicative of a potential therapeutic implication of endogenous H₂S (generated by CSE) in atherosclerosis. Interestingly, estrogen attenuates atherosclerosis development by stimulating H₂S production in female ovariectomized ApoE knockout mice fed with a high-fat diet, suggesting that H₂S mediates estrogen-induced vascular protection (Zhou et al. 2013; Fu et al. 2013).

7.3 H₂S and Aortic Aneurysms

Aortic aneurysms, including thoracic and abdominal aortic aneurysms, are the most life-threatening cardiovascular complication in Marfan syndrome, leading to aortic expansion, dissection, rupture, and sudden death. Enzymatic degradation of extracellular matrix (ECM) protein by matrix metalloproteinases (MMPs) leads to dilation of the aortic wall and constitutes the most prominent characters of aortic aneurysms. The regulatory role of H₂S in vascular remodeling during aortic aneurysms has not been explored yet. Early studies showed that H₂S is involved in vascular remodeling during the development of hypertension and neointimal formation. Rats with high pulmonary blood flow for 11 weeks showed a significant pulmonary hypertension and pulmonary artery collagen remodeling in association

with a decrease in lung tissue H₂S content (Li et al. 2008b). Supplement of exogenous H₂S lowered pulmonary artery collagen I and collagen III protein levels and normalized pulmonary hypertension, suggesting that the downregulation of H₂S is involved in the development of pulmonary artery collagen remodeling induced by high pulmonary blood flow. Vascular H₂S production was lower in SHR compared with Wistar Kyoto (WKY) rats, and the inhibitory role of H₂S on collagen generation was stronger in the SHR than in the WKY rats (Zhao et al. 2008). In a carotid artery-injured mouse model, H₂S was shown to mitigate vascular remodeling from endothelial damage by decreasing the expressions of TIMP-3 and MMP-9 (Vacek et al. 2010). In addition, we found that H₂S inhibited SMC migration and neointima formation by suppressing $\alpha 5\beta 1$ -integrin-dependent MMP-2 expression. The inhibitory roles of H₂S on the MMP/TIMP system suggest that H₂S can block excess degradation of ECM and maintain the normal structure of the aorta (Yang et al. 2012a, b). Future studies need to determine the role of H₂S in vascular degeneration and aortic aneurysm formation. The molecular mechanisms involved in the pathogenesis of aortic aneurysm formation as well as the mechanisms underlying CSE/H₂S system-regulated MMP activation are also waiting to be explored. All these endeavors will provide a new therapeutic avenue for the prevention and treatment of aortic aneurysms.

8 Prospective

Undoubtedly, the research scope and depth on H₂S signaling in the cardiovascular system, especially in blood vessel-related disorders, will continue to expand and deepen. The identification of new cellular targets and development of novel agents to enhance endogenous H₂S generation are highly expected. A better understanding of the roles of H₂S in vascular remodeling and the regulatory mechanisms for endogenous production of H₂S in blood vessel can provide insight into potential therapeutic interventions against blood vessel-associated disorders. The interaction of H₂S with numerous biological molecules, such as NO and CO, needs to be better characterized. At the end of the day, all these fundamental and mechanistic studies will become meaningless if the related discoveries cannot be translated from bench side to bedside so that human health and welfare will be improved. That day will not be too far away.

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Hydrogen Sulfide and Urogenital Tract

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and Raffaella Sorrentino

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Abstract

In this chapter the role played by H₂S in the physiopathology of urogenital tract revising animal and human data available in the current relevant literature is discussed. H₂S pathway has been demonstrated to be involved in the mechanism

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underlying penile erection in human and experimental animal. Both cystathionine- β synthase (CBS) and cystathionine- γ lyase (CSE) are expressed in the human corpus cavernosum and exogenous H_2S relaxes isolated human corpus cavernosum strips in an endothelium-independent manner. Hydrogen sulfide pathway also accounts for the direct vasodilatory effect operated by testosterone on isolated vessels. Convincing evidence suggests that H_2S can influence the cGMP pathway by inhibiting the phosphodiesterase 5 (PDE-5) activity. All these findings taken together suggest an important role for the H_2S pathway in human corpus cavernosum homeostasis. However, H_2S effect is not confined to human corpus cavernosum but also plays an important role in human bladder. Human bladder expresses mainly CBS and generates in vitro detectable amount of H_2S . In addition the bladder relaxant effect of the PDE-5 inhibitor sildenafil involves H_2S as mediator.

In conclusion the H_2S pathway is not only involved in penile erection but also plays a role in bladder homeostasis. In addition the finding that it involved in the mechanism of action of PDE-5 inhibitors strongly suggests that modulation of this pathway can represent a therapeutic target for the treatment of erectile dysfunction and bladder diseases.

Keywords

Bladder • Corpus cavernosum • Erectile dysfunction (ED) • Benign prostatic hyperplasia (BPH) • Lower urinary tract symptoms (LUTS) • PDE-5 inhibitor • CBS • CSE • 3MST

1 Introduction

Hydrogen sulfide (H_2S) presence in mammalian tissues has been known since the 1980s, but H_2S has been considered mainly as a metabolic waste product with no potential physiological activity. The first evidence, indicating H_2S as an endogenous mediator was published in 1996 by Abe and Kimura (1996) that described a role for this mediator in the brain. Some years, after this first evidence was published, many research groups have focused their interest on this new pathway as described in the other chapters. H_2S together with nitric oxide (NO) and carbon monoxide (CO) is now classified as gasotransmitter. Like the other two gasses, it has the ability to diffuse easily through the cellular membranes. Hydrogen sulfide can also interact with proteins such as ion channels or enzymes regulating their state either directly via chemical modification such as by sulfuration and sulphydration or indirectly via second messengers.

The role of H_2S in the homeostatic control of our body is now consistently supported by the literature, and in the other chapters of the present book, its synthesis, measurement, chemistry, and interaction have been reported. This chapter will deal with the role played by H_2S in the physiopathology of urogenital tract revising animal and human data available in the current relevant literature.

2 Hydrogen Sulfide in Male Sexual Function

2.1 The Male Reproductive System

Male reproductive system consists of a number of sex organs that concur to the human reproductive process localized around the pelvic region. Briefly, the main male sex organs are the penis and the testicles important for producing semen and sperm. In particular, the penis and the scrotum represent the external genital organs. The penile erectile apparatus consists of two vascularized paired corpora cavernosa. The corpus spongiosum together with the urethra is related to the ventral area of the penis. The corpora cavernosa, the corpus spongiosum, and the glans penis are composed of septa of smooth muscle and erectile tissue that enclose vascular cavities. The tunica albuginea forms a thick fibrous coat to the spongy tissue of the corpora cavernosa and corpus spongiosum, and it consists of two layers. The muscles involved in penile function are the ischiocavernosum and bulbo-spongiosus. The ischiocavernous muscle inserts into the medial and inferior surface of the corpora and increases penile turgidity during erection. In the flaccid state, the smooth muscle fibers are tonically contracted by the sympathetic system. During erection, trabeculae smooth muscle is relaxed, and this allows a cascade of events. The arterioles of the penis are dilated while the veins are passively compressed between the tunica albuginea and the peripheral sinusoids reducing the venous outflow. This process is known as the venous-occlusive mechanism, and it leads to erection that is sustained by an increase of intra-cavernous blood pressure of about 100 mmHg. The contraction of the ischiocavernous muscle further increases the intra-cavernous pressure leading to the rigid erection phase.

The main role of the scrotum is to hold and protect the testes. It also contains numerous nerves and blood vessels. Between the male internal genital organs, there is the epididymis, a whitish mass of tightly coiled tubes cupped against the testicles. It acts as a maturation and storage for sperm before they pass into the vas deferens, which carry sperm to the ampullary gland and prostatic ducts. Testosterone is the most important sexual hormone in males and is released by the testes. This hormone has a crucial role in the development of sperm and is also responsible for the development of physical characteristics in men such as facial hair and deep voice. Moreover the accessory glands, such as the seminal vesicles and the prostate gland, provide fluids that lubricate the duct system and nourish the sperm cells. In particular, the prostate gland surrounds the ejaculatory ducts at the base of the male urethra, just below the bladder. The prostate gland is responsible for the proof semen, a liquid mixture of sperm cells, prostate fluid, and seminal fluid. This gland is also responsible for making the semen milky in appearance by mixing calcium to the semen coming from seminal vesicle. The semen remains cloudy and clumpy until the prostatic pro fibrinolysis is formed into fibrinolysis and lysis of the fibrinogen from the seminal vesicle fluids occurs.

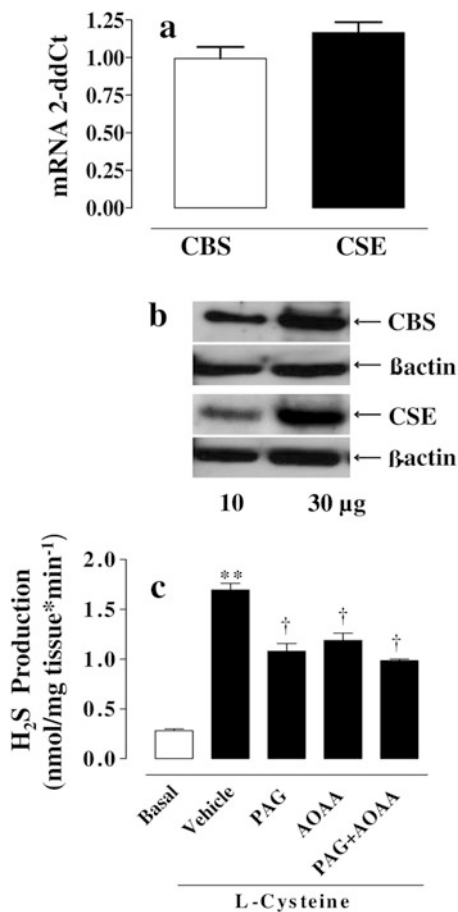
2.2 Erectile Function and Dysfunction

Erection is a vascular event which relies upon interaction of neural and humoral mechanisms at various levels. Indeed, since the penis receives innervations from sacral parasympathetic (pelvic), thoracolumbar sympathetic, and somatic (pudenda) nerves, the erection phenomenon requires participation of these three systems. Thus, erection is a consequence of a complex integration of several signals. In simple words, erection is essentially a spinal reflex that can be initiated by recruitment of penile afferents, but also by visual, olfactory, and imaginary stimuli. All these stimuli may contribute to the increase in intra-cavernous pressure (Cirino et al. 2006) and thus to penile erection. Erectile dysfunction (ED) is defined as the consistent or recurrent inability of a man to attain and/or maintain a penile erection sufficient for sexual activity (2nd International Consultation on Sexual Dysfunction-Paris, June 28th–July 1st, 2003). Of note, ED shares many of the risk factors that contribute to the development and the progression of cardiovascular diseases such as age, hypercholesterolemia, obesity, diabetes, and smoking (Brunner et al. 2005). Several clinical studies have shown that ED is often associated with cardiovascular disturbance and therefore it could be taken in account as an early sign of cardiovascular diseases (Dong et al. 2011; Nehra et al. 2012). This hypothesis relies on the fact that penile artery size is smaller as compared to coronary arteries. Therefore, it is feasible that the same level of endothelial dysfunction can cause a more significant reduction of blood flow in erectile tissues compared to that elicited in coronary circulation. Therefore, ED could be considered a prognostic factor for possible cardiovascular problems (Gandaglia et al. 2014).

2.3 Hydrogen Sulfide and Erectile Function and Dysfunction

The corpora cavernosa, as briefly reported above, are composed of sinusoids bearing a single layer of endothelial cells surrounded by multiple layers of smooth muscle cells. Thus, the corpora cavernosa are vascular organs. Among the vasodilator agents, it is undisputed that nitric oxide (NO) is considered one of the most important endogenous factors since it is released not only by endothelial cells but also from plexus nerve. Recently hydrogen sulfide (H_2S), as discussed in the other chapters, has been shown to possess an important role in the modulation of smooth muscle cell tone. In particular, it has been demonstrated that H_2S relaxes smooth muscle cells, and this finding together with others have lead to study the role of this mediator in the penile physiology. To date, the role of this gas in corpus cavernosum function has been addressed in animal models as well as in the human tissues (d'Emmanuele di Villa Bianca et al. 2011; Qiu et al. 2012). The first evidence was published in 2006 by Srilatha and coworkers. These authors showed that the intra-cavernous injection of sodium hydrogen sulfide (NaHS) resulted in a significant increase in penile length and cavernous pressure in primates measured by using laser Doppler flow meter and cutaneous probe. To acquire

Fig. 1 CBS and CSE: activity, Western blot analysis, and qRT-PCR of human penile tissue. (a) HCC-expressed mRNA for both CBS and CSE as determined by qRT-PCR. (b) Representative Western blot analysis for CBS and CSE. (c) HCC homogenate produced H₂S under basal conditions (*open bar*). Incubation of HCC homogenate with 10 mM L-Cys caused a significant increase in the H₂S production compared with basal values (***P* < 0.001). PAG (10 mM), 1 mM AOAA, or 10 mM PAG plus 1 mM AOAA significantly inhibited the L-Cys-induced increase in H₂S production ([†]*P* < 0.01). Data represent the mean SEM from 3 or 4 different human specimens (Reprinted with permission from d'Emmanuele di Villa Bianca et al. 2009)



further evidence on the role of H₂S in penile erection, a study on rats was also performed. Administration of dl-propargylglycine [PAG, a cystathione-γ lyase (CSE) inhibitor] to rats resulted in a significant reduction in cavernous nerve stimulation-evoked perfusion pressure. This study suggested a possible role for endogenous H₂S in facilitating nerve-mediated penile tumescence (Srilatha et al. 2006). In 2009, d'Emmanuele di Villa Bianca and coworkers demonstrated the role and function of H₂S in human. The authors showed that both cystathione-β synthase (CBS) and CSE, the main enzymes involved in H₂S synthesis, are expressed in the human corpus cavernosum (CC) (Fig. 1a, b). Moreover, by using human CC homogenates, as enzyme source, they confirmed that the enzymes present in human CC can efficiently convert L-Cys (the substrate) to H₂S (Fig. 1c). CBS and CSE are localized within muscular trabeculae and smooth muscle component of the penile artery (Fig. 2). Interestingly, the enzymes appear to be differently distributed within the penile structure. In particular, CSE but not

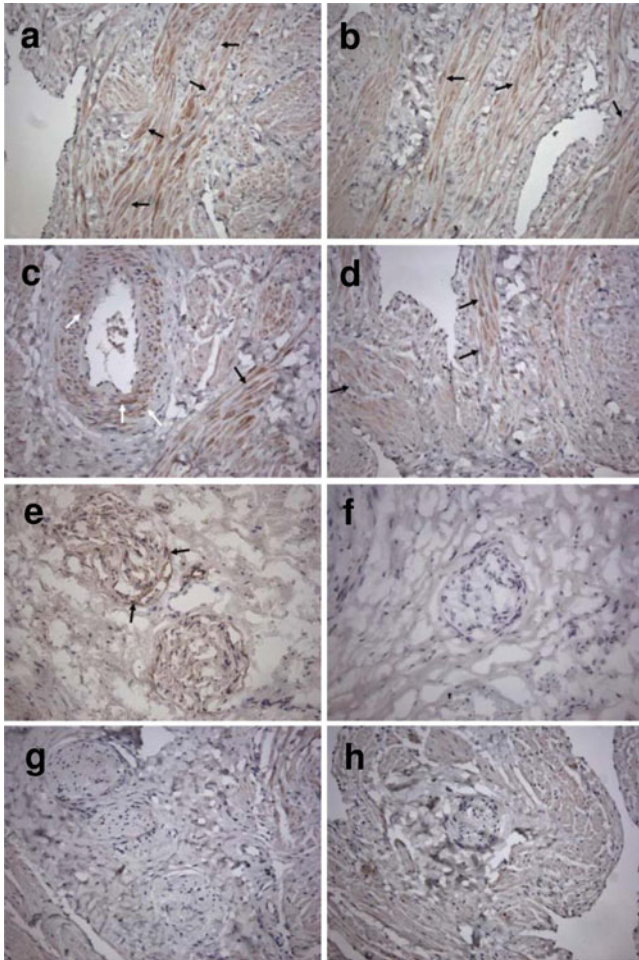
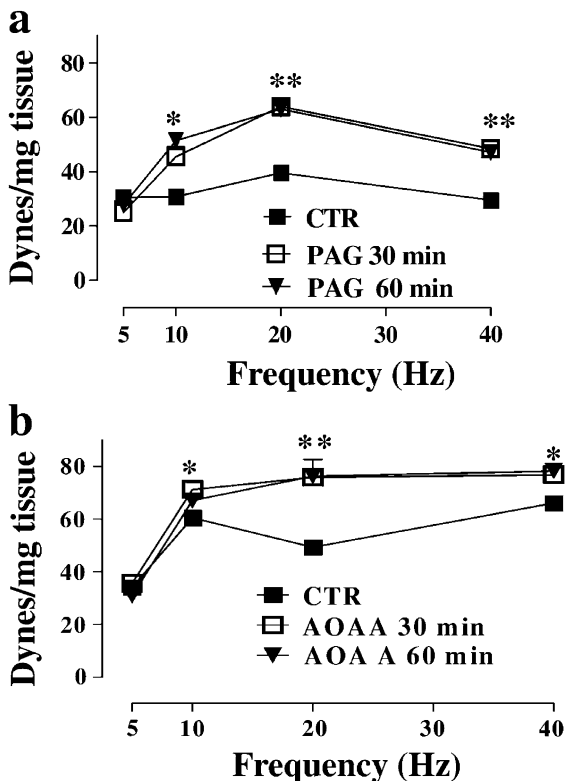


Fig. 2 Immunohistochemistry for CBS and CSE in HCC. Immunohistochemical detection of CSE and CBS in HCC tissue. Immunoreactivity and nuclear staining appear *brown* (DAB) and *blue* (hematoxylin counterstain), respectively. CSE was detected in trabecular muscular tissue (**a** and **b**, *black arrows*) and vascular smooth muscle cells (**c**, *white arrows*). Immunoreactivity for CBS was mostly observed in trabecular muscular tissue (**d**, *black arrows*). Results illustrated are from a single experiment and are representative of three different specimens. (Original magnification, 200.) Immunohistochemical detection of CSE and CBS in HCC nerve fibers. Immunoreactivity and nuclear staining appear *brown* (DAB) and *blue* (hematoxylin counterstain), respectively. CSE was detected in nerve fibers in cryostat (**e**, *arrows*) and not in paraffin (**g**) sections. Both cryostat (**f**) and paraffin (**h**) sections lacked immunoreactivity for CBS. Results illustrated are from a single experiment and are representative of three different specimens. (Original magnification, 200.) (Reprinted with permission from d'Emmanuele di Villa Bianca et al. 2009)

CBS is expressed in peripheral nerves (Fig. 2). Functional studies conducted by using human CC isolated strips demonstrated that both exogenous H₂S and L-Cys cause a concentration-dependent relaxation of human CC strips. This relaxant

Fig. 3 EFS of HCC strips. EFS caused a frequency-related increase in basal tone. Incubation of HCC strips with 10 mM PAG for 30 or 60 min (a) or with 1 mM AOAA for 30 or 60 min (b) significantly increased the EFS-induced contraction [$*P < 0.05$; $**P < 0.01$ vs. control (CTR)]. Tissue responses to EFS are expressed as force in dynes per milligram of tissue. Experiments were performed on three different specimens (Reprinted with permission from d'Emmanuele di Villa Bianca et al. 2009)



effect was inhibited by the CBS inhibitor, aminoxyacetic acid (AOAA) and only slightly reduced by the endothelial NO-synthase inhibitor, L-NAME. Electrophysiological experiments performed using peripheral nerve electrical field stimulation of human penile tissue demonstrated that H₂S pathway is involved in penile homeostasis. Indeed, EFS of human CC strips, under resting conditions, caused an increase in tension that was significantly potentiated by inhibiting CSE and/or CBS with the selective inhibitors PAG and/or AOAA (Fig. 3). The role of H₂S pathway in erectile function was confirmed also *in vivo* by using an experimental animal model of penile erection. Using this model, it was demonstrated that either NaHS or L-Cys administration cause an increase in the intra-cavernous pressure in anesthetized rats (d'Emmanuele di Villa Bianca et al. 2009).

These studies have revealed that there are important differences between the human and rat tissue for what concerns the H₂S pathway. In fact, in rats, only CSE is expressed. This finding is also confirmed by the fact that CSE but not CBS inhibition significantly decreased H₂S production when rat cavernosal tissue homogenates were used as enzyme source to generate H₂S from L-cysteine *in vitro*. Concerning the role of CSE and H₂S in rat corpus cavernosum, it has been also shown that CSE inhibition by PAG causes a significant increase in non-adrenergic non-cholinergic-induced relaxation leading to an enhanced

neurogenic relaxation of rat corpus cavernosum induced by EFS. The authors hypothesize that, since H₂S does not have a direct constrictor effect in corporal tissue, this effect might be due to either inhibition of NO synthesis by endogenous H₂S or to a direct chemical reaction between H₂S and NOS products (Ghasemi et al. 2012). Thus, while CSE in rats accounts for the production of H₂S, in human, the H₂S pathway is sustained by both CBS and CSE, which are also differently distributed in the human corpus cavernosum (Fig. 2) indicating that human and rat penile tissues are different for what concerns the H₂S pathway involvement in penile erectile mechanism(s).

2.4 Mechanism(s) of Action that Sustain the Effect of H₂S in Erection

2.4.1 Ion Channels and EDHF as H₂S Target

Ion channels are intimately involved in the biochemical events associated with smooth muscle function, and their activation/inactivation is tightly associated with tumescence/detumescence function during erection. As already reported in the previous chapters, H₂S activates within the vasculature the adenosine triphosphate (ATP)-activated potassium (K_{ATP}) channel. The mechanism of activation proposed involves S-sulfhydration by H₂S of the cysteine residues located on a specific subunit of the extracellular loop of the K_{ATP} channel complex leading to opening of this channel (Jiang et al. 2010; Mustafa et al. 2011). In particular, potassium channels are important in mediating physiologically relevant relaxation responses in human, rabbit, and rat corpus cavernosum strips (Christ et al. 1993; Mirone et al. 2000; Karicheti and Christ 2001; Spektor et al. 2002; Ruiz Rubio et al. 2004). In this context, it has been proposed that K_{ATP} channels could have an important role as modulator of corporal smooth muscle tone (Christ et al. 1993). Indeed, in diabetes, the K_{ATP} channels-mediated relaxation is significantly reduced in human CC strips (Venkateswarlu et al. 2002). Interestingly, in human CC strips glybenclamide, a selective K_{ATP} channel inhibitor, significantly impairs H₂S-induced relaxation (d'Emmanuele di Villa Bianca et al. 2009) indicating that these channels are involved in H₂S effect. K_{ATP} channels may also act by increasing cyclic adenosine monophosphate (cAMP) and thus cause relaxation. In fact, K_{ATP} channels are a physiologically important target of the adenylate cyclase/cAMP/PKA signaling pathway (Nelson et al. 2011), confirming indirectly the possible involvement of this channel as proposed by Srilatha et al. in 2007. In fact, the effect of NaHS in rabbit CC can be blocked by using MDL 12,330A, an adenylate cyclase inhibitor (Srilatha et al. 2007). On this basis, it is feasible to hypothesize that in diabetes, where there is a reduced activity of these channels, the H₂S pathways may be involved in ED associated with the diabetes. However, these issues need to be addressed more carefully in order to define the contribution of this mechanism to the physiology of erection as well as its involvement in ED.

Due to the ability of H₂S to directly interact with proteins, another channel that seems to be its target is the calcium-dependent potassium (K_{Ca}) channel.

The physiological role of this channel in human CCSM cells have been demonstrated (Christ et al. 1993). As mentioned before, NO is a key mediator in erectile function, and it derives from both peripheral nervous terminations and of course from endothelial cells. However, the existence of an unidentified endothelial factor that promotes smooth muscle hyperpolarization and relaxation, resistant to NO synthase (NOS) and cyclooxygenase (COX) inhibition, has been clearly established (Busse et al. 2002). This factor, named endothelial-derived hyperpolarizing factor (EDHF), has particular functional relevance in small arteries. This pathway involves the activation of two populations of endothelial potassium channels, the small/intermediate conductance and the large K_{Ca} channels (Grgic et al. 2009). It has been clearly demonstrated that in human penile resistance arteries, as opposite to human CC, there is a significant component in the relaxation response to acetylcholine that is resistant to NOS and COX inhibition attributed to EDHF-like activity (Angulo et al. 2003). This EDHF-like activity is impaired in human penile resistance arteries from diabetic patients. In this framework fits H_2S since there are significant data supporting H_2S as an EDHF (Wang 2003, 2009; d'Emmanuele di Villa Bianca et al. 2011; Tang et al. 2013).

Inhibitors of the phosphodiesterase-5 (PDE-5) are drugs largely used to treat men with ED to increase the half-life of cyclic guanosine monophosphate (cGMP) in the corpus cavernosum facilitating its relaxation and therefore penile erection. However, although sildenafil has been shown to be efficacious for treating ED, the percentage of efficacy in diabetic patients is clearly reduced when compared to nondiabetic men (Rendell et al. 1999; Price et al. 1998; Vickers and Satyanarayana 2002). Indeed, the presence of ED and diabetes in a patient represents a prognostic factor for a poor response to sildenafil but also with the other new PDE-5 inhibitors tadalafil (Saenz de Tejada et al. 2002) and vardenafil (Goldstein et al. 2003). The mechanism(s) responsible for this poor clinical outcome is not, as yet, understood. One hypothesis formulated is related to the impairment of EDHF in human penile resistance vessels of diabetes patients that is not restored by PDE-5 treatment. This hypothesis is supported by a study where it has been shown that dobesilate, a drug that specifically enhances endothelium-dependent relaxation attributed to EDHF, restores the endothelial function in penile arteries of diabetic patients (Angulo et al. 2003). Therefore, it is also feasible, and it needs to be explored the possibility, that the combination of an H_2S donor and a PDE-5 inhibitor may overcome the poor outcome of PDE-5 inhibitors in diabetic patients and could be beneficial in the oral treatment of diabetic ED.

Other H_2S target may be the calcium permeable channels. In fact, other authors have reported the ability of H_2S to interfere not only with potassium channels but also with calcium homeostasis particularly in neurons, cardiomyocytes, and endothelial cells. Although some effects of H_2S on calcium signals are secondary to K_{ATP} modulation, there is growing consensus about the existence of a direct effect of H_2S on Ca^{2+} -permeable channels (Munaron et al. 2013). The modulation/activation of these channels seems to be tightly related to the H_2S concentration. At the present stage, there are no data available in the current literature on CCSM cells in both animal models and human tissues.

2.4.2 Testosterone, Gender Difference, and Biosynthesis of H₂S

Testosterone (T), as already reported above, plays a critical role not only in human male sexual behavior, but several studies have indicated a relation between androgen levels and sexual interest, libido, and the frequencies of orgasm and nocturnal erections. Moreover, there is a growing body of evidence that low T levels are associated with an increased cardiovascular and cancer mortality (Hackett et al. 2014). Thus, T seems to have a protective effect in man as occurs for estrogen in woman. For example, T plasma levels negatively correlate with hypertension, diabetes, and severe coronary artery disease. While it is clear that T restoration to a physiological level is beneficial in hypogonadal subjects, the positive role of T treatment is questionable in men who are not clearly hypogonadal or eugonadal. Another important well-accepted concept is that T level correlates to ED (Isidori et al. 2014). In particular, patients with organic ED have lower free T levels than patients with psychogenic ED. Moreover, a strong positive correlation exists between free T levels and the degree of trabecular smooth muscle relaxation, as measured by resistive index at dynamic duplex ultrasound (Aversa et al. 2000). The role of T in regulating CCSM and penile arterial tone has been extensively explored in animal models. In this regards, a dual separate effect with or without the endothelium has been shown. The effect in presence of endothelium is related to NO; in particular, there are several evidences supporting a role for androgens in regulating the expression and activity of NOS isoforms in the corpus cavernosum in animal models (Traish et al. 2007). The increase in T plasma levels in response to sexual stimulation raised the question of its biological significance. To this aim, several indirect evidences support the presence of a local vascular effect of T in men. In fact, it has been demonstrated that penile erection in the healthy male is associated with a significant increase in T level in both circulating and in the cavernous plasma. Moreover, in the flaccid state, the T level in cavernous plasma is significantly lower when compared to the systemic concentration, and on this basis, the authors conclude that these differential evaluation could represent a diagnostic tool (Becker et al. 2000). Interestingly, during the penile tumescence and rigidity, T level in corpus cavernosum increases significantly in comparison to the flaccid state, and this increase is not observed in ED patients (Becker et al. 2001). These results further support the hypothesis that T, through the androgen receptor, has a direct effect on the cavernous smooth muscle. Androgen receptors are present in the human CC (Schultheiss et al. 2003), and T induces relaxation by activating smooth muscle K_{ATP} channels in human CC strips (Yildiz et al. 2009). This study was the first to report a direct non-genomic relaxant effect of T on human CC in vitro. The authors have demonstrated that T causes a rapid vasorelaxation partially mediated by an increase of potassium efflux through K_{ATP} channels, but not involving BK_{Ca}, voltage-dependent inward rectifier K channel, or voltage K channels. Thus, T-induced relaxation on smooth muscle CC seems to involve an endothelium-dependent (particularly through NO pathway) and -independent (probably by K channels) mechanism (Aversa et al. 2000). It has been confirmed that these channels are also involved in T-induced relaxation within the vasculature and that T effect involves H₂S pathway. In fact, it has been shown

that T causes an increase in H₂S level involving the L-Cys/H₂S pathway in rat aorta (Bucci et al. 2009). The increase of H₂S induced by T and its vasodilatory effect was prevented by the androgen receptor antagonist nilutamide in rat aorta rings indicating that the interaction with the androgen receptor is a key issue (Bucci et al. 2009). In this context, it has been recently shown the heat shock protein 90, which plays a role in the activation of androgen receptor, is also involved in CSE activation. In the same work, the presence of a similar mechanism for progesterone and 17- β -estradiol was ruled out. This latter observation well fit with the finding that H₂S levels in human blood, collected from male healthy volunteers, were higher than those in female samples (Brancaleone et al. 2014).

Thus, if T level is related to ED, it could well be that an impairment of T-mediated H₂S production plays a role in ED. On this specific subject, there is only one study available, performed using an in vivo rat model. In this work, the role of endogenous H₂S in ED induced by androgen deficiency has been investigated. Serum T levels were significantly reduced after 2 and 4 weeks from castration, and in this condition, the H₂S pathway was significantly impaired as well as there was a reduction in intra-cavernous pressure increase elicited by electrical stimulation. On the other hand, T replacement resulted efficacious to restore H₂S release and to improve intra-cavernous pressure in rats. These in vivo data support the hypothesis that H₂S pathway may be one of the mechanisms underlying androgen role in erection (Zuo et al. 2014). Moreover, it has been demonstrated that aging significantly impairs NO and H₂S level both in plasma and corpus cavernosum tissue. A reduction of the intra-cavernous pressure is countered by NaHS or sildenafil after 10 weeks of treatment. The link between T and H₂S was further confirmed by Srilatha and coauthors, who have shown a marked increase in T or estradiol levels after NaHS supplementation. These data support the idea that ED in aging may be also linked to a derangement in the H₂S pathway accompanied by low T levels (Srilatha et al. 2012).

All these findings support the importance of androgens in regulating smooth muscle function in the penis. A possible clinical application could be the use of a combination of PDE-5, T, and/or H₂S donors/substrate (such as L-Cysteine) to be used in patients nonresponder to PDE-5 therapy (Isidori et al. 2014).

2.4.3 Interaction Between H₂S and NO

There is a deep discussion in literature concerning the role of NO/cGMP pathway in H₂S effects, and this issue is under investigation in several anatomical district (see other chapters). For example, it has been demonstrated that CSE activity is upregulated by NO (Zhao et al. 2001), and it is partially inhibited by NOS blockade (Zhao and Wang 2002). The data available in the current literature demonstrated that H₂S (1) inhibits eNOS activity partly through inhibition of eNOS phosphorylation by reducing Akt phosphorylation (Geng et al. 2007), (2) stimulates the activity of several upstream kinase such as Akt and in turn activates eNOS by phosphorylation (Cai et al. 2007), (3) acts as PDE inhibitor (Bucci et al. 2010), (4) directly causes eNOS phosphorylation (Altaany et al. 2014), (5) is mutually required with NO in order to elicit angiogenesis and vasodilatation (Coletta

et al. 2012). Therefore, there are several evidence implying a cross talk at vascular level between NO and H₂S. As opposite, few studies are available on human CC tissues. The relaxing effect of H₂S on human CC strips has been demonstrated to be endothelium independent (d'Emmanuele di Villa Bianca et al. 2009), but this result does not exclude a possible indirect effect of H₂S since, in presence of the endothelium blockade of NOS, L-NAME causes a significant inhibition of H₂S-induced relaxation at higher concentration. This latter result is in line with a recent study by Meng and coworkers (Meng et al. 2013) demonstrating that incubation of rat CC tissues with NaHS leads to an increase in eNOS but not nNOS mRNA. In this study, the authors show that the increased mRNA expression of eNOS correlates with protein expression as well as with NO production. However, caveolin-1 expression, a dominant inhibitory interaction partner of eNOS, was not modified by H₂S. On this basis, the authors conclude that H₂S could be particularly useful in improving the clinical outcome of ED patients, whose erectile impairment involves a weakened function of endothelial-derived NO. However, whether H₂S increases the soluble guanylyl cyclase activity directly was not examined. The possible role of H₂S on modulating the NO/cGMP pathway has been, instead, suggested in vascular studies, where it has been shown that H₂S increases cGMP levels acting as a PDE inhibitor delaying cGMP degradation (Bucci et al. 2010; Coletta et al. 2012). Preliminary data indicate that an increase in cGMP level drive by PDE-5 can cause an increase in H₂S production in mice corpus cavernosum (Dikmen et al. 2013). Many aspects concerning the interaction between these two gasses are still matter of debate in the current relevant literature. A recent review addresses in depth the possible role of this cross talk in erectile function/dysfunction (Yetik-Anacak et al. 2014).

2.5 Hydrogen Sulfide and Vas Deferens

The vas deferens transports sperm from the epididymis to the ejaculatory ducts in anticipation of ejaculation. Due to the role of H₂S on erectile function, its presence and role in *vas deferens* have also been evaluated. Both CBS and CSE are functionally expressed in the vas deferens of rat, mice, and human. The endogenous H₂S causes a smooth muscle relaxation of vas deferens (Li et al. 2011). The same group has also demonstrated that NaHS-induced effect is mediated by BK_{Ca} channel. Indeed, they observed a consistent reversion of the relaxant effect of rat vas deferens-induced relaxation by performing a pharmacological modulation with iberiotoxin or tetraethylammonium. The authors also demonstrated that H₂S modulation of K_{Ca} channels requires a redox signaling. Indeed, N-ethylmaleimide, a sulfhydryl alkylation compound protecting thiols from oxidation, inhibited NaHS relaxation as opposite to DTT, a strong reducing agent, that did not affect the H₂S response of vas deferens. Besides, the presence of the BK_{Ca} channels in rat vas deferens smooth muscle cells was also confirmed. Finally, the involvement of the NO pathway, the transient receptor potential (TRP) channels and of the K_{ATP} channels were excluded (Li et al. 2012).

2.6 Hydrogen Sulfide and Human Prostate

Regarding the role of H₂S in the prostate, it has been reported by Guo and coworkers (Guo et al. 2012) that in human prostatic tissues (obtained from patients undergoing surgery for prostatic cancer) and cells, this pathway is physiologically present. The presence of H₂S pathway has been also confirmed by using biopsy from cancer-free human prostate (Gai et al. 2013). Moreover, Guo and coworkers have demonstrated that both cell activity and CBS/CSE protein levels are higher in the androgen-dependent prostate cancer cell LNCaP than in all the other cell lines evaluated and that dihydrotestosterone downregulates this activity. Varying H₂S levels, as well as CBS/CSE expression, in human prostate stromal and epithelial compartments have been also described. In particular, prostate epithelium expresses both CBS and CSE as opposite to stromal where only CSE is expressed. Overall, these data suggest that H₂S pathway can be involved in prostate cancer and benign prostatic hyperplasia. These results also further confirm that a link between H₂S and T exists. In physiological condition, T can modulate the H₂S synthesis contributing to erectile function. In pathologic condition, such as androgen-dependent prostate cancer, where H₂S pathway is over expressed, it may represent a potential therapeutic target (Guo et al. 2012). Indeed, H₂S does not alter significantly androgen receptor expression or its phosphorylation but inhibits androgen receptor transactivation probably at the DNA-binding level. This effect may involve a posttranslational regulation of androgen receptor by S-sulphydration leading to conformational change and alteration of protein function. Thus, H₂S may decrease the genomic effect associated with androgen receptor activation exhibiting an anti-proliferating effect, data that are in agreement with the higher cell proliferating rate associated with aging in prostate tissues of CSE knock-out mice (Zhao et al. 2014).

3 Hydrogen Sulfide in Female Sexual Function

There is very little published on the possible involvement of the H₂S pathway in female sexual physiology. In 2009, a pilot study has been published suggesting that H₂S pathway plays a physiological role in female sexual apparatus. In particular, the authors have studied the effect of exogenous H₂S in vaginal and clitoral cavernosal smooth muscle strips from New Zealand white female rabbits. By using H₂S donors and several inhibitors towards different enzymes and channels, the authors have shown that H₂S vasodilatory effect involves cAMP, (cyclic adenosine 3′/5′-monophosphate), NO-cGMP (cyclic guanosine monophosphate), and K_{ATP} channels. Of particular interest is the finding that inhibition of H₂S-induced relaxation was observed only when a combination of both adenylate- and guanylate- cyclase inhibitors was used indicating that both nucleotides may concur to H₂S effect. Thus, H₂S pathway appears to be involved also in female sexual responses. Moreover, these data further support the evidence that there exists an interplay between NO and H₂S pathways (Srilatha et al. 2009).

4 Hydrogen Sulfide and Bladder

4.1 Bladder Function

The urinary bladder has two important functions: storage of urine and emptying. Storage of urine occurs at low pressure, which implies that the bladder relaxes during the filling phase. The wall of the bladder is composed of 3 layers:

1. Outer layer of loose connective tissue, containing blood and lymphatic vessels and nerves.
2. Middle layer, consisting of a mass of interlacing smooth muscle fibers and elastic tissue—this is called the detrusor muscle.
3. Inner layer composed of transitional epithelium.

The storage and periodic elimination of urine are dependent upon the reciprocal activity of two functional units in the urinary tract: a reservoir (urinary bladder) and an outlet (bladder neck, smooth and striated muscle of the urethra). During urine storage, the outlet is closed, and the bladder is quiescent, allowing intravesical pressure to remain low over a wide range of bladder volumes. During voiding, the muscles of outlet relax and the bladder smooth muscles contract, raising intravesical pressure and inducing urine flow. These changes are coordinated by three sets of nerves (parasympathetic, sympathetic, and somatic) emerging from the sacral and thoracolumbar levels of spinal cord. Thus, when the bladder is filled, the relaxation of the wall stimulates the afferent fibers, and input is transmitted to the cortex eliciting the micturition reflex. During the micturition, the detrusor muscle and the longitudinal muscle of the neck and the urethral sphincter contract by activation of parasympathetic nerves. Contextually, an inhibitory input of the somatic nerves causes a relaxation of the external sphincter allowing the urine ejection. Different signaling molecules contribute to the fine tuning of the urinary bladder, acting both in autocrine and paracrine manner. The receptors involved are specifically distributed among different types of cells within the urinary bladder structure. Various neurotransmitters, including acetylcholine, norepinephrine, dopamine, serotonin, excitatory and inhibitory amino acids, adenosine triphosphate, nitric oxide, and neuropeptides, are implicated in the neuronal regulation of micturition (de Groat and Yoshimura 2001). Several disorders can affect the bladder such as cystitis, urinary stones, bladder cancer, urinary incontinence, hematuria, urinary retention, cystocele, bed-wetting, dysuria, overactive bladder, and lower urinary tract symptoms (LUTS). Disturbances of storage function may result in LUTS, such as urgency, frequency, and urge incontinence, the components of the overactive bladder syndrome (Abrams et al. 2002). The overactive bladder syndrome, which may be due to involuntary contractions of the smooth muscle of the bladder (detrusor) during the storage phase, is a common problem (Milsom et al. 2001). H₂S has been shown to be involved in the bladder function in different species such as trout, rat, pig, and human (Fusco et al. 2012; Gai et al. 2013; Fernandes et al. 2013a, b; Patacchini et al. 2004; Dombkowski et al. 2006).

4.2 Animal Studies

H₂S is endogenously produced in urinary bladder of trout, mice, and rat implying that the contribution of this pathway to bladder function is conserved in different species. Similarly, to human tissues, homogenates of bladder can generate detectable amount of H₂S in basal or stimulated conditions (Dombkowski et al. 2006; Matsunami et al. 2012; Gai et al. 2013). The expression of the enzymes responsible of H₂S production has been reported in bladder of rat, pig, and mice by using different methods. CSE and CBS were detected by immunohistochemistry in nerve fibers and are widely distributed within the smooth muscle layer of the pig bladder neck (Fernandes et al. 2013a). CBS, CSE, and 3MST have been also shown to be present in rat bladder (Gai et al. 2013). In mouse bladder, only the presence of CSE has been reported so far, (Matsunami et al. 2012). However, the presence of CBS cannot be excluded since it has not been evaluated (Matsunami et al. 2012).

In the urinary tract, H₂S donors caused smooth muscle relaxation and contraction depending on the species and experimental conditions used. NaHS (exogenous source of H₂S) or L-cysteine (the substrate) relaxes rat bladder strips in a concentration-dependent manner (Gai et al. 2013). This effect can be inhibited by incubating tissues with glybenclamide, a K_{ATP} channel inhibitor, CSE, and CBS blockers (Gai et al. 2013). In trout bladder, NaHS, as well as Na₂S (exogenous donor of H₂S less used in the current literature), inhibits spontaneous contractions and relaxes precontracted strips (Dombkowski et al. 2006). In trout bladder, the H₂S response does not appear to be mediated by K_{ATP} channels as occurs in human and rat since glybenclamide was ineffective. Similarly, the other types of K⁺ channels were not involved in H₂S-induced relaxation, as well. The rat urinary bladder can be indirectly contracted through H₂S stimulation of capsaicin-sensitive nerves as reported by Patacchini and coworkers (Patacchini et al. 2004, 2005). This discrepancy between rat and human could be due to the different derivation of the tissue (Kardong 2005). In rat, H₂S has a contractile effect that is mediated by the activation of sensory neurons in a ruthenium red-sensitive but not capsazepine-dependent manner. Thus, it is feasible that in this case, the molecular target of H₂S could be either a receptorial domain located on the transient receptor potential vanilloid receptor 1 (TRPV1) cation channel, independent from those bound by vanilloids and other TRPV1 activators, or another ruthenium red-sensitive TRP cation channels co-expressed on primary afferent neuron terminals (Patacchini et al. 2005). The super family of TRP includes the transient receptor potential ankyrin (TRPA) (Nilius et al. 2007), which has been found on capsaicin-sensitive primary sensory neurons (Story et al. 2003; Bautista et al. 2005). The TRPA1 activators allyl isothiocyanate and cinnamaldehyde contract rat urinary bladder through stimulation of the capsaicin-sensitive nerves (Andrade et al. 2006; Patacchini et al. 1990). Therefore, an action of H₂S also on TRPA1 in urinary bladder has been suggested. This hypothesis is based upon the following findings i) co-localization of TRPA1 and TRPV1 in rat bladder afferents ii) H₂S mimics the TRPA1 agonist cinnamaldehyde effect stimulating the micturition reflex after protamine sulfate pretreatment (Streng et al. 2008). Thus, TRPA1 represents a molecular target for H₂S in rat bladder.

H₂S-mediated O₂ sensing has been demonstrated in a variety of O₂-sensing tissues in vertebrate cardiovascular and respiratory systems, including smooth muscle in systemic and respiratory blood vessels and airways, carotid artery, adrenal medulla, and other peripheral as well as central chemoreceptors (Olson 2014). Concerning the possible involvement of this mechanism in urinary tract, it has been shown that H₂S is involved in O₂ sensing/signal transduction in the trout urinary bladder (Dombkowski et al. 2006) showing some similitude with mammalian tissue (Olson 2014).

An upregulation of CSE has been found in cyclophosphamide-induced cystitis in mouse, suggesting a role for the endogenous H₂S in the pathogenesis of this experimental model (Matsunami et al. 2012). In addition, a link between H₂S and Ca_v3.2 T-type channel has also been suggested in the cystitis-related nociceptive changes (Matsunami et al. 2012).

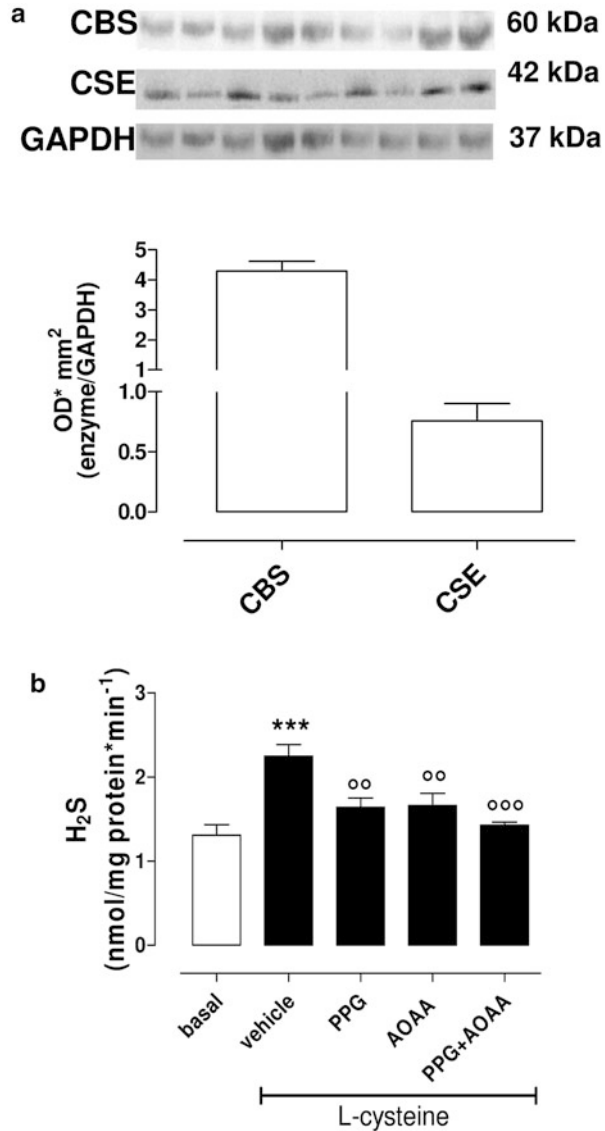
The control of the bladder neck tone is important during the voiding phase either in physiological or in pathological conditions. In this regard, CBS and CSE, as reported above, have been found expressed in nerve fibers in the pig bladder neck smooth muscle layer suggesting the involvement of H₂S as mediator released following muscle distension (Fernandes et al. 2013a). Both electrical field or GYY4137, an H₂S donor, relaxed the pig bladder neck strips through K_{ATP} channel activation as well as COX1-derived prostanoids (Fernandes et al. 2013a, b) indicating also a functional role at this level. These studies provided also evidence for a role of endogenous H₂S released from nerves involvement in the inhibitory transmission of the outflow region (Fernandes et al. 2013a). Additionally, H₂S promotes the release of inhibitory neuropeptide by the activation of TRPA1 and TRPV1 (Fernandes et al. 2013b). Therefore, at the present stage, K_{ATP} channels, TRPA1, and TRPV1 are considered feasible targets for H₂S in the bladder function.

4.3 Human Studies

Human bladder expresses all three enzymes responsible for the H₂S production namely CBS, CSE, and 3-MST (Fusco et al. 2012; Gai et al. 2013). Indeed, human bladder homogenate generates a basal level of production of H₂S that can be enhanced by incubation with L-cysteine (the substrate). This effect is reversed by either PAG (CSE inhibitor) or AOAA (CBS inhibitor) (Fig. 4). Functional studies performed using isolated human bladder strips have demonstrated that both sodium hydrogen sulfide (NaHS) or L-cysteine relax human bladder strips precontracted by carbachol. Thus, H₂S pathway is involved in the regulation of bladder homeostasis.

As discussed, K_{ATP} channels are activated by H₂S. It is known that urinary bladder smooth muscles express K_{ATP} channels, and they have been shown to be involved in the regulation of bladder contractility (Andersson 1992; Bonev and Nelson 1993). The presence of mRNA for sulfonylurea receptors has been demonstrated in both pig and human detrusor (Buckner et al. 2000). The detrusor smooth muscle expresses a variety of K⁺ channels that are responsible for the detrusor smooth muscle excitability and contractility (Andersson and Arner 2004;

Fig. 4 Cystathionine b-synthase (CBS) and cystathionine g-lyase (CSE) western blot, and activity in human bladder dome. (a) Representative western blot for CBS and CSE (*upper*) and densitometric analysis for CBS and CSE normalized to glyceraldehyde-3-phosphate dehydrogenase (GAPDH; *lower*) from nine patients; data were expressed as the mean plus or minus standard error of the mean (SE). (b) Production of hydrogen sulfide (H_2S) in human tissue homogenates from five patients under basal condition (*open bar*) and after incubation with L-cysteine 10 mM (*black bars*). L-cysteine caused a significant increase in H_2S production ($***p < 0.001$ vs. basal) that was significantly reverted by CSE inhibitor (D,L-propargylglycine [PPG]; 10 mM) and/or CBS inhibitor (aminooxyacetic acid [AOAA]; 1 mM) ($^{\circ\circ}p < 0.01$ and $^{\circ\circ\circ}p < 0.001$, respectively, vs. L-cysteine). Data were calculated as nanomoles per milligram of protein per minute and expressed as mean \pm SE (Reprinted with permission from Fusco et al. 2012)



Brading 2006; Christ and Hodges 2006; Gopalakrishnan and Shieh 2004). Interestingly, the density of K_{ATP} channels is significantly higher in detrusor smooth muscle (Petkov et al. 2001; Kajioka et al. 2008; Shieh et al. 2001). In this context, it has been shown that $\sim 1\%$ of the functional K_{ATP} channels need to be activated in order to inhibit detrusor smooth muscle action potentials and related phasic contractions (Petkov et al. 2001). Studies on isolated detrusor muscle from human and several animal species have demonstrated that K_{ATP} channel openers reduce not only spontaneous contractions but also those induced by electrical

stimulation, carbachol, and low external K^+ concentrations (Andersson 1993). Interestingly, the relaxant effect elicited by H_2S is inhibited by glybenclamide, a K_{ATP} channel inhibitor (Gai et al. 2013). Therefore, it is feasible to hypothesize that once released, H_2S causes relaxation of the detrusor muscle that involves K_{ATP} channel activation thereby contributing to the tonic regulation of the bladder tone. Bladder tone regulation is a key issue in LUTS. LUTS markedly increase with age in both males and females, and it represents a major problem in the elderly population. Recently, the PDE-5 inhibitor tadalafil has been approved by FDA for LUTS treatment. In addition, it has been reported that sildenafil, a well-known PDE-5 inhibitor, can directly relax human bladder strips and that this effect involves the H_2S pathway (Fusco et al. 2012). Indeed, sildenafil incubation of human bladder samples causes a time- and concentration-dependent increase in H_2S production (Fig. 5). In conclusion, the human data, available in the current

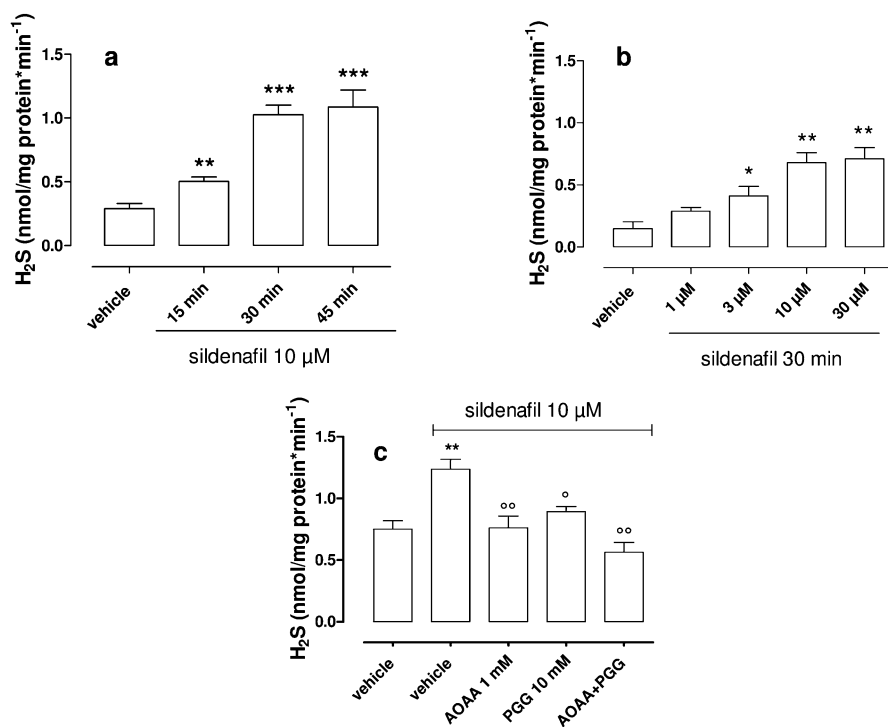


Fig. 5 Sildenafil-induced hydrogen sulfide (H_2S) production in human bladder dome. (a) Sildenafil 10 mM caused a time-dependent increase in H_2S production (*** p < 0.001 vs vehicle; ** p < 0.01 vs. 30 min and 45 min). (b) Sildenafil incubation for 30 min caused a concentration-dependent increase in H_2S production compared to vehicle (* p < 0.05 and ** p < 0.001 vs. vehicle). (c) Combination of CBS and/or CSE inhibitors (DL-propargylglycine [PPG] 10 mM and/or aminoxyacetic acid [AOAA] 1 mM) significantly inhibited sildenafil-induced H_2S production (° p < 0.05 and °° p < 0.01 vs sildenafil 10 mM; ** p < 0.01 vs. vehicle). Data were calculated as nanomoles per milligram of protein per minute and expressed as mean \pm SEM for five patients (Reprinted with permission from Fusco et al. 2012)

relevant literature, suggest that the H₂S pathway is involved not only in the physiology of human bladder function but also in LUTS as well as in overactive bladder. However, further studies are necessary in order to better define the role of this pathway.

5 Hydrogen Sulfide and Urethra

The urethra together with smooth muscles in the bladder controls the storage and the voiding of urine. The presence as well as the ability of CBS, CSE, and 3-MST to convert L-cysteine into H₂S has been reported both in human and rat urethra (Gai et al. 2013). On the functional side, H₂S has been shown to have no effect on the basal tone but to relax the human urethra strips (Gratzke et al. 2009). The relaxing effect on human urethra strips has been proposed to involve the TRPA1 receptors that are expressed on nerves fibers, urothelium, and interstitial cells (Gratzke et al. 2009).

6 Conclusions and Future Directions

On the basis of the findings present in the current literature, from non-mammalian and mammalian, including human, it is clear that H₂S is endogenously produced, and the enzymes responsible of its biosynthesis are constitutively present in the genitourinary tract. In addition, H₂S appears to be a phylogenetically ancient and versatile regulatory molecule. Nevertheless, the H₂S response in the genitourinary tract, as reported above, appears to be specie dependent. Therefore, more studies are needed in order to clearly define the role of this pathway in human since the translation of preclinical data to human is not always possible. For what concerns, drugs used in therapy that have been proposed to involve H₂S as mediator, there are some considerations that can be made. It is now well established that patients' response to a therapy is not equally effective, and who once were called nonresponders are now considered as a specific sub group. Indeed, it is now clear that genetics plays a great role in determining the nature of the drug responses. However, it is also true that, most likely, there are other therapeutic targets that have not as yet been defined within a specific pathology. In this regard, in both ED and in pathologies associated with bladder dysfunction, there is a need to find new therapies. Indeed, PDE-5 inhibitors often do not resolve the ED in diabetes patients or fail to act in a certain number of eligible patients with no comorbidities. In fact, the efficacy of PDE-5 inhibitor, which is a mainstay in the treatment of ED, is negatively associated with the nerve and endothelium damage. These latter features are associated with several pathologies such as cardiovascular disease, diabetes, obesity, and post-prostatectomy state and in turn lead to an impairment in the signaling of the NO/cGMP pathway. In this context, the H₂S pathway represents an attractive target since a link between PDE-5 inhibitors and H₂S pathway has been already shown. Indeed, tadalafil limits myocardial infarction through H₂S

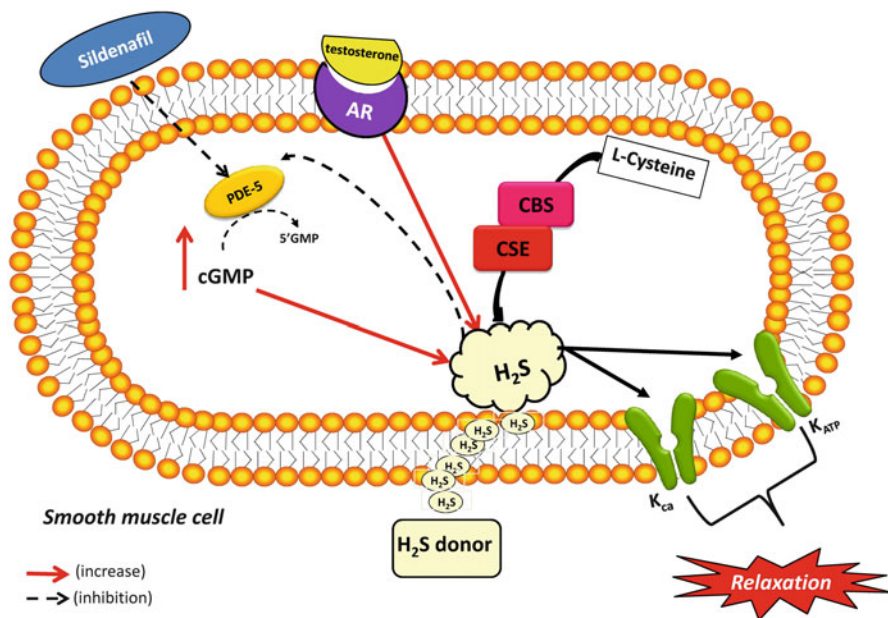


Fig. 6 Drugs proposed to involve hydrogen sulfide (H_2S) pathway in their mechanism of action. The cartoon reproduces a smooth muscle cell of the urogenital tract, where the involvement of H_2S in testosterone and sildenafil mechanism of action is depicted. Adenosine triphosphate-activated potassium channel (K_{ATP}); Androgen receptor AR; calcium-dependent potassium channel (K_{Ca}); cyclic guanosine monophosphate (cGMP); Cystathione- β synthase (CBS); Cystathione- γ lyase (CSE), phosphodiesterase-5 (PDE-5)

signaling (Salloum et al. 2009), and sildenafil causes an increase in H_2S production by CBS and CSE activities in human bladder (Fusco et al. 2012). A very preliminary attempt to develop a drug working on H_2S and cGMP pathways has been taken by Shukla and coworkers who have synthesized and characterized an H_2S -donating derivative of sildenafil (ACS6) (Shukla et al. 2009). For what concerns the bladder, it has to be stressed that the widely used antimuscarinic drugs often fails to alleviate the LUTS symptoms. Also in this case, by looking at the recent findings, highlighted in this chapter, the H_2S pathway represents an attractive therapeutic target that may allow to develop new drugs. In Fig. 6 are shown the drugs for which the involvement of H_2S in their mechanism of action has been proposed.

Another possible therapeutic approach that rises from what has been discussed is the possibility to modulate H_2S levels by exogenous supplementation. Intuitively, the easiest way should be to use H_2S donors that release the gas slowly, in order to reproduce as much as possible, the physiological conditions. Alternatively, the H_2S production could be enhanced by inducing H_2S synthesis by using L-cysteine or other substrates. However, this approach could fail in condition where a down-regulation of the enzyme CBS and CSE occurs. Several studies have addressed these issues in preclinical setting (Kashfi and Olson 2013). The activity of garlic-

derived molecules, generally considered as H₂S releasers following metabolization, such as diallyl disulfide, diallyl sulfide, diallyl trisulfide, and diallyl tetrasulfide, has been profusely investigated (Jacob et al. 2008). Garlic and its bioactive component, the S-allyl cysteine, have been shown to restore erectile function in diabetic rats by preventing ROS formation through modulation of NADPH oxidase subunit expression (Yang et al. 2013). In particular, anti-inflammatory and anticancer effects have been demonstrated. However, no data, concerning their efficacy, on urogenital tract are available. The GYY4137 is actually the most attracting synthetic H₂S donor. It inhibits lipid accumulation exhibiting anti-atherosclerotic activity both in vitro and in vivo (Xu et al. 2014) and exerts anti-inflammatory (Li et al. 2013) and anticancer (Kashfi 2014) activity. Also in this case, there are no data available on GYY4137 effect in urogenital tract diseases.

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H₂S Is a Promoter of Angiogenesis: Identification of H₂S “Receptors” and Its Molecular Switches in Vascular Endothelial Cells

Bei-Bei Tao, Wen-Jie Cai, and Yi-Chun Zhu

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Abstract

Angiogenesis is a physiological process in organ development and also a compensatory response in ischemia. When ischemia occurs, oxygen sensors in vascular endothelial cells sense the decrease in oxygen, thus activating downstream signaling pathways to promote the proliferation, migration, and tube formation of the endothelial cells. The new vasculatures are formed by sprouting from preexisting vessels, in order to maintain oxygen homeostasis in ischemic tissues (Folkman and Shing 1992). Collateral circulation is sometimes established under chronic ischemic conditions such as chronic myocardial

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ischemia (Banai et al. 1994). However, naturally occurring angiogenesis is usually not sufficient to compensate for ischemia in ischemic tissues. Proangiogenic drugs may be useful to promote angiogenesis in these diseases.

Keywords

Angiogenesis • Myocardial ischemia • Endothelial cells • Hydrogen sulfide • Molecular switch

1 Introduction

Angiogenesis is a physiological process in organ development and also a compensatory response in ischemia. When ischemia occurs, oxygen sensors in vascular endothelial cells sense the decrease in oxygen, thus activate downstream signaling pathways to promote the proliferation, migration, and tube formation of the endothelial cells. The new vasculatures are formed by sprouting from preexisting vessels, in order to maintain oxygen homeostasis in ischemic tissues (Folkman and Shing 1992). Collateral circulation is sometimes established under chronic ischemic conditions such as chronic myocardial ischemia (Banai et al. 1994). However, naturally occurring angiogenesis is usually not sufficient to compensate for ischemia in ischemic tissues. Proangiogenic drugs may be useful to promote angiogenesis in these diseases.

Accumulating evidences have revealed some protective effects of hydrogen sulfide in the cardiovascular system. H₂S reduces blood pressure (Zhao et al. 2001) and ameliorates hypertension-induced myocardial remodeling in spontaneous hypertensive rats (Yan et al. 2004; Shi et al. 2007). H₂S provides protection in models of myocardial injury (Geng et al. 2004) and myocardial infarction (Chuah et al. 2007). However, the mechanisms underlying numerous biological effects of H₂S remain largely unknown. One of the most challenging questions in this new field is the identification of the “receptors” for H₂S to mediate its numerous biological effects. Here we review the studies reporting the proangiogenic role of H₂S and the recent work which reports the first H₂S “receptor” and its molecular switch identified in this proangiogenic model.

2 H₂S Promotes Angiogenesis

We report the proangiogenic role of *in vitro* and *in vivo* models of angiogenesis in 2007 (Cai et al. 2007). H₂S facilitates migration of vascular endothelial cells in both the wound healing model and the transwell model (Fig. 1A–D). H₂S also increases tube length and the number of branching points in an *in vitro* of tube formation (Fig. 1G–I). In addition, H₂S promotes angiogenesis in an *in vivo* model of Matrigel plug assay in mice which is evidenced by an increase in vascular density (Fig. 1L) and hemoglobin content (Fig. 1M) in the Matrigel plug. In 2009, Papapetropoulos

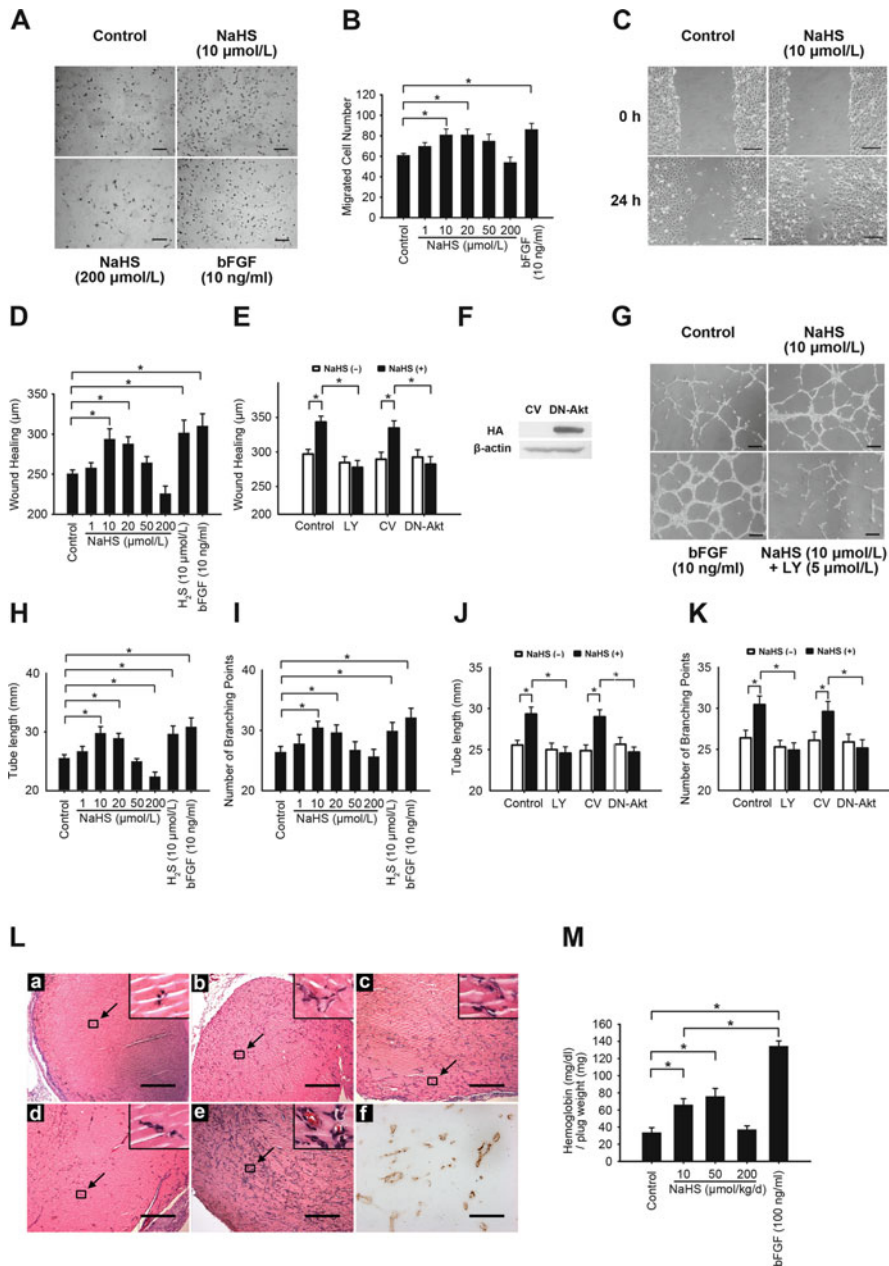


Fig. 1 H₂S promotes angiogenesis both in vitro and in vivo. (A and B) Cell migration was assessed by transwell migration assay. (C and D) Representative micrographs and statistical analysis of scratch wound healing assay of RF/6A endothelial cells. (E–K) NaHS-induced promotion of RF/6A endothelial cell migration was prevented by either LY 294002 (5 μmol/L) or transfection of the DN-Akt. The effects of H₂S on in vivo angiogenesis were assessed using

et al. further reported the proangiogenic role of endogenous H₂S in cystathionine γ -lyase (CSE) knockout mice. VEGF administration caused less angiogenesis in CSE knockout mice as compared with the wild-type mice. The authors suggested that endogenous H₂S has an important role in inducing angiogenesis (Papapetropoulos et al. 2009). This report also confirmed our previous results by using the chicken chorioallantoic membrane (CAM) model (Papapetropoulos et al. 2009). They found H₂S administration increased vascular length in chicken chorioallantoic membranes. In addition, the effects of H₂S in promoting the migration and tube formation of the vascular endothelial cells were also validated by Papapetropoulos et al. (2009).

3 The Proangiogenic Role of H₂S in Ischemic Disease Models

Though the proangiogenic role of H₂S had been reported by two independent groups using various *in vitro* and *in vivo* models, the role of H₂S in the angiogenesis in diseases was not examined. In 2010, we reported that chronic treatment with exogenous H₂S using an H₂S donor NaHS improved regional blood flow in ischemic tissues in a rat model of femoral artery ligation (Wang et al. 2010). H₂S significantly promoted collateral vessel growth in ischemic hind limbs of rats (Fig. 2A, B) and causes an increase in capillary density (Fig. 2D). In addition, regional tissue blood flow was also increased by H₂S treatment in ischemic tissue (Fig. 2C). These findings indicate that H₂S is a proangiogenic factor under both physiological and pathophysiological conditions. In the following years, the proangiogenic role of H₂S in various disease models has been reported by several independent groups. In 2012, Bir et al. reported a proangiogenic role of H₂S in a mouse model of hind limb ischemia (Bir et al. 2012). More recently in 2012, Qipshidze et al. examined the role of H₂S in chronic heart failure model of myocardial infarction induced by ligation of the left anterior descending coronary artery (Qipshidze et al. 2012). They found that the expression level of both H₂S-producing enzymes (CSE and CBS) was decreased in mice with myocardial infarction. Chronic treatment with NaHS for 1 month promoted angiogenesis in the myocardium using X-ray angiography and blood flow measurement with Doppler.

In addition to myocardial ischemia, H₂S also protects against cerebral ischemia by inducing angiogenesis in the brain (Hyunduk et al. 2014). In a model of transient focal cerebral ischemia/reperfusion induced by suturing endovascular internal carotid artery, H₂S administration improved significantly the neurological outcomes on the limb placing test and corner turn test without reducing the

Fig. 1 (continued) Matrigel plug assay in mice (**L** and **M**). (**L**) Representative photomicrographs of hematoxylin-eosin-stained Matrigel sections of mice treated with vehicle (**a**), various doses of NaHS (**b**, **c**, and **d** for 10, 50, and 200 $\mu\text{mol} \cdot \text{kg}^{-1} \cdot \text{day}^{-1}$ NaHS, respectively) and bFGF (**e**, 100 ng/ml in Matrigel). (**M**) Neovascularization in the Matrigel plugs was quantified by measuring hemoglobin content using the tetramethylbenzidine method

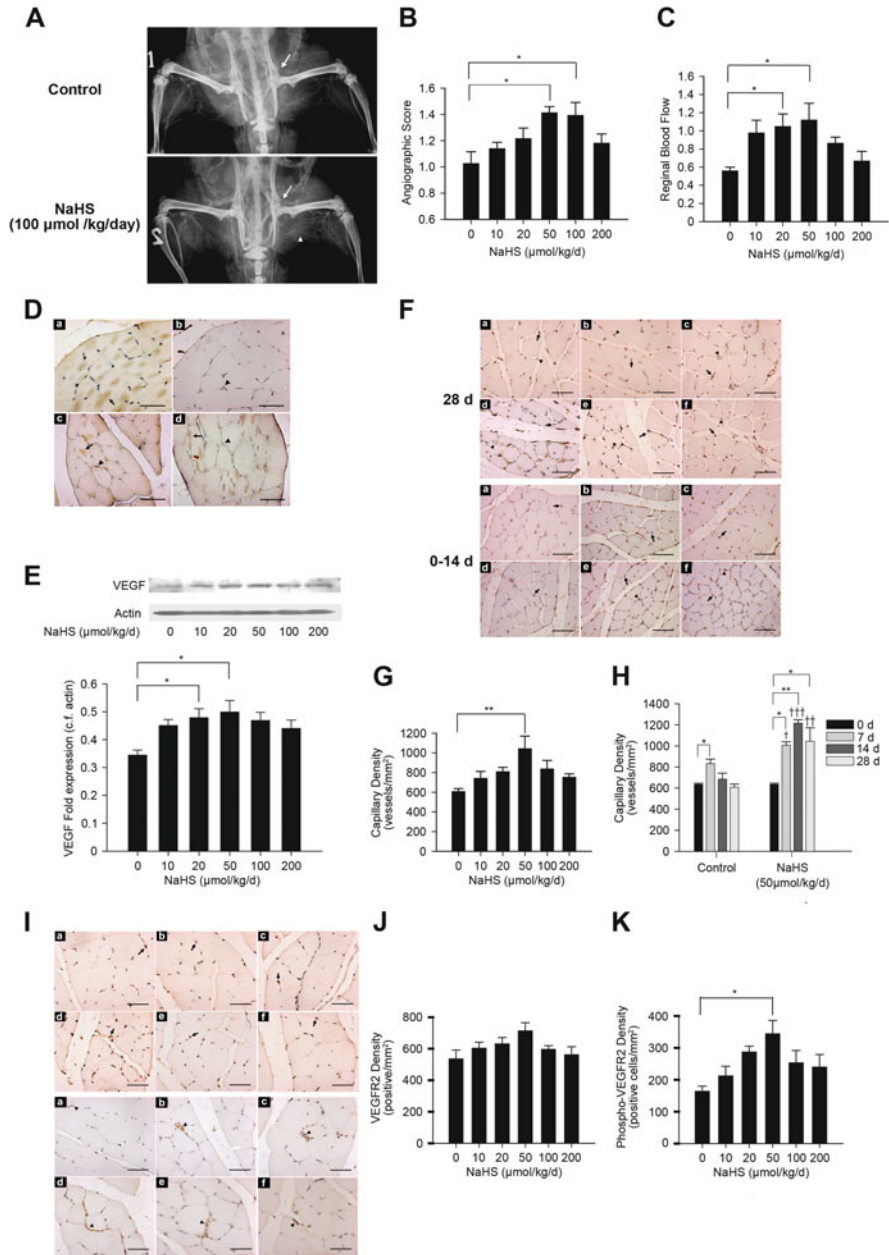


Fig. 2 NaHS treatment promoted collateral vessel formation and regional blood flow after femoral artery occlusion in the rat hindlimb ischemia model. (A) Representative postmortem angiograms obtained 4 weeks after surgery. *Arrow* denotes the site of ligation at the femoral artery. *Arrow head* indicates the typical “corkscrew” appearance of collateral vessels. (B) Quantitative analysis of collateral vessel development was performed by measuring the total length of the

infarction volume. In addition, H₂S enhanced the synthesis of endothelial cells in the ischemic region.

4 Signaling Pathways to Mediate the H₂S Effects

When H₂S was first reported as a proangiogenic factor, Cai et al. found that the proangiogenic effects of H₂S in vascular endothelial cells were dependent on the PI3K/Akt pathway (Fig. 1E, F, G, J, K) (Cai et al. 2007). However, the direct target molecule for H₂S remains unknown. It could be one of the signaling elements in this pathway or some elements in the upstream of the pathway in the vascular endothelial cells. This prompts us to explore for the “receptor” of H₂S in the subsequent studies.

On the other hand, the H₂S signals may be transduced with more complicated mechanisms where cell-cell interaction is involved. For example, the proangiogenic effects of H₂S in the hind limb ischemia model might involve an interaction between the vascular endothelial cells and the skeletal muscle cells. H₂S treatment caused an increase in VEGF expression in the skeletal muscle cells in the ischemic hind limb where collateral circulation was promoted by H₂S treatment (Fig. 2E) (Wang et al. 2010). An increase in the phosphorylation of Akt and VEGFR2 was observed in the vascular endothelial cells neighboring the skeletal muscle cells in the ischemic tissues (Fig. 2I, J, K). The data suggest that exogenous H₂S might stimulate the skeletal muscle cells to release VEGF which acts on VEGFR2 on the cell membrane of vascular endothelial cells to promote angiogenesis. In addition to VEGF, other proangiogenic factors are also involved in the proangiogenic effects of H₂S. It is reported that H₂S promotes biosynthesis of VEGF, Ang-1, and Ang-2 in

Fig. 2 (continued) contrast-opacified vessels. (C) Blood flow measured with microsphere assay. **F** (upper part) Representative micrographs showing the vascular endothelial cells (arrow) and capillary vessels (arrow head) in the gastrocnemius muscles 4 weeks after surgery stained with anti-rat CD34 antibody (a for the vehicle; b, c, d, e, and f for 10, 20, 50, 100, and 200 $\mu\text{mol} \cdot \text{kg}^{-1} \cdot \text{day}^{-1}$ NaHS groups, respectively). **F** (lower part) Representative micrographs showing the time-course study of CD34-positive cells. (a–c) Control group (a, 0 day; b, 7 days; c, 14 days); (d–f) 50 $\mu\text{mol} \cdot \text{kg}^{-1} \cdot \text{day}^{-1}$ NaHS group (d, 0 day; e, 7 days; f, 14 days). Bar = 100 μm . (G and H) Quantitative analysis of the capillary density. * $P < 0.05$, ** $P < 0.01$ vs. day 0; † $P < 0.05$, †† $P < 0.01$, ††† $P < 0.001$ vs. control (at the same time point) by ANOVA. (D and E) NaHS increased VEGF expression in the ischemic hindlimb muscles. (a and b) Two consecutive sections showing that VEGF expression (arrow) was mainly localized in the skeletal muscle cells but not the vascular endothelial cells (arrow head). (c and d) Double staining of VEGF (arrow) and the vascular endothelial cells (arrow head) further clarified the skeletal muscle cells expressed VEGF in the groups treated with 50 $\mu\text{mol} \cdot \text{kg}^{-1} \cdot \text{day}^{-1}$ NaHS administration (d) and the vehicle (c). Bar = 100 μm . I (a–f in the upper part) Representative micrographs of VEGFR2 (arrow) in the gastrocnemius muscles. (I) (a–f in the lower part) Representative micrographs of phospho-VEGFR2 in the gastrocnemius muscles. a for the vehicle; b, c, d, e, and f for 10, 20, 50, 100, and 200 $\mu\text{mol} \cdot \text{kg}^{-1} \cdot \text{day}^{-1}$ NaHS groups, respectively. (J and K) Quantitative analysis of VEGFR2 and phospho-VEGFR2 density, respectively

astrocytes, increases mouse brain endothelial cells (MBECs) in the *in vitro* OGD model, and increases VEGF and Ang-1 in ischemic brain (Hyunduk et al. 2014). The expression of VEGF, VEGFR1, and VEGFR2 is increased in ischemic heart with H₂S treatment. Whereas, some factors with anti-angiogenic property such as endostatin, angiostatin, and parstatin were decreased (Qipshidze et al. 2012). These findings support the idea that H₂S is also a proangiogenic factor in disease status and some paracrine mechanisms might be involved in *in vivo* models.

On the other hand, more signaling elements may be involved in addition to the PI3K/Akt pathway found earlier in single cells. For example, ERK and HIF-1 α might also be involved in mediating the H₂S effects (Hyunduk et al. 2014). Hyunduk et al. report that H₂S increases p-Akt, p-ERK, and HIF-1 α in cultured astrocytes and MBECs.

Some chemical inhibitors such as glibenclamide (K_{ATP} channel blocker) and p38 inhibitors have been shown to block the H₂S effects in inducing the migration of vascular endothelial cells (Papapetropoulos et al. 2009). These data all suggest that the H₂S signals are mediated by some signaling pathways in a single cell. However, the signaling mechanisms underlying the H₂S effects remain largely unknown.

On the other hand, other gaseous signaling molecules including nitric oxide (NO) and carbon monoxide (CO) have been suggested to interact with H₂S in their proangiogenic effects. Coletta et al. found that H₂S and NO were mutually dependent in promoting angiogenesis (Coletta et al. 2012). An inhibitor of NO biosynthesis, L-NAME, blocks the H₂S effects in promoting endothelial cell proliferation. H₂S-induced angiogenesis in aortic rings is abolished in cultured vessels isolated from eNOS knockout mice. In the *in vivo* Matrigel plug model, H₂S-induced angiogenesis is also abrogated completely in eNOS knockout mice. These results indicate that H₂S-induced angiogenesis is dependent on the NO pathway. Moreover, Bir et al. show that H₂S increases eNOS activity by stimulating the phosphorylation of Akt and thereby increases NO production (Bir et al. 2012). This may explain how H₂S can increase NO biosynthesis. Indeed, future works are required to clarify the signaling mechanisms underlying the H₂S effects in angiogenesis.

5 H₂S Donors Developed to Treat Ischemic Diseases

Kan et al. report that S-propargyl-cysteine (SPRC), a novel endogenous hydrogen sulfide modulator, promotes proliferation, migration, and *in vitro* tube formation of the vascular endothelial cells (Kan et al. 2014). This proangiogenic effect of SPRC is also validated in rat aortic ring angiogenic model and the *in vivo* Matrigel plug assay. SPRC also increases angiogenesis in the mouse hind limb ischemia model and the rat myocardial ischemia model. SPRC treatment significantly increased STAT3 phosphorylation, and the proangiogenic effect of SPRC is blunted by siRNA-mediated knockdown of STAT3. The data indicate the involvement of STAT3 in this H₂S effect. However, co-crystallization shows that STAT3 does not directly interact with SPRC. SPRC enhances the interaction between VEGFR2

and STAT3 and increases nuclear translocation of STAT3, leading to an activation of VEGF transcription.

Diallyl trisulfide, a stable H₂S donor, is also reported as a proangiogenic factor. It has been found to have therapeutic potential in heart failure and to increase vascular density in the heart (Polhemus et al. 2013). Chronic treatment with diallyl trisulfide for 12 weeks significantly increased VEGF expression and decreased angiotensin expression in the myocardium in a pressure overload model of mice. In addition, diallyl trisulfide increased eNOS phosphorylation and enhanced NO bioavailability in the heart.

6 VEGFR2 Is the “Receptor” for H₂S to Induce Angiogenesis

All the works reviewed above support the proangiogenic role of H₂S which is first reported by Cai et al. in 2007. Since then, numerous efforts have been endowed to clarify the mechanisms underlying the H₂S effects. A series of signaling elements such as VEGF, PI3K, Akt, NO, and STAT3 have been suggested to mediate the H₂S effects. However, all these elements may reflect some signaling event secondary to the initial H₂S signals. That means there must be some target molecules in the vascular endothelial cells to directly interact with H₂S and transduce the initial H₂S signals to downstream signaling pathways to induce the H₂S effects in the vascular endothelial cells. We name the direct target molecule for H₂S as the “receptor” for H₂S. A typical ligand has a conformation to match with the structure of its receptor. However, H₂S is too small a molecule to have a conformation sufficient to dock with its “receptors.” In fact, H₂S does cause significant biological effects in the vascular endothelial cells. This leads to a hypothesis that there is some “receptor” for H₂S in the vascular endothelial cells. This “receptor” must have a direct interaction with H₂S and initiate a cascade of signaling events leading to increased migration of the cells. With this idea, we searched for the H₂S “receptor” using the in vitro model of H₂S-induced migration of vascular endothelial cells. We focused on the PI3K/Akt pathway identified in the previous studies (Cai et al. 2007). Purified recombinant proteins of PI3K and Akt were reacted with H₂S in a cell-free system and the activity of these two kinases was not changed (Fig. 3b, c). Moreover, no chemical modification was found in these two kinases reacted with H₂S using tandem mass spectrometry. Since blockade of these two elements abolished the H₂S effects, we tried to search for the H₂S “receptor” in the pathways upstream of PI3K and Akt and finally found that VEGFR2 functioned as the direct target molecule for H₂S. H₂S directly increased the activity of recombinant VEGFR2 in a cell-free system (Fig. 3a). In vascular endothelial cells, blockade of VEGFR2 using either the chemical inhibitor SU5416 or siRNA-mediated knock-down both significantly prevented the proangiogenic effect of H₂S. Therefore, we believed that VEGFR2 functioned as a “receptor” for H₂S in the vascular endothelial cells. We proposed that there might be some chemical modifications in VEGFR2 induced by H₂S treatment. At glance of the data, we did not find any modifications in any of the free amino acid residues in VEGFR2 using tandem mass

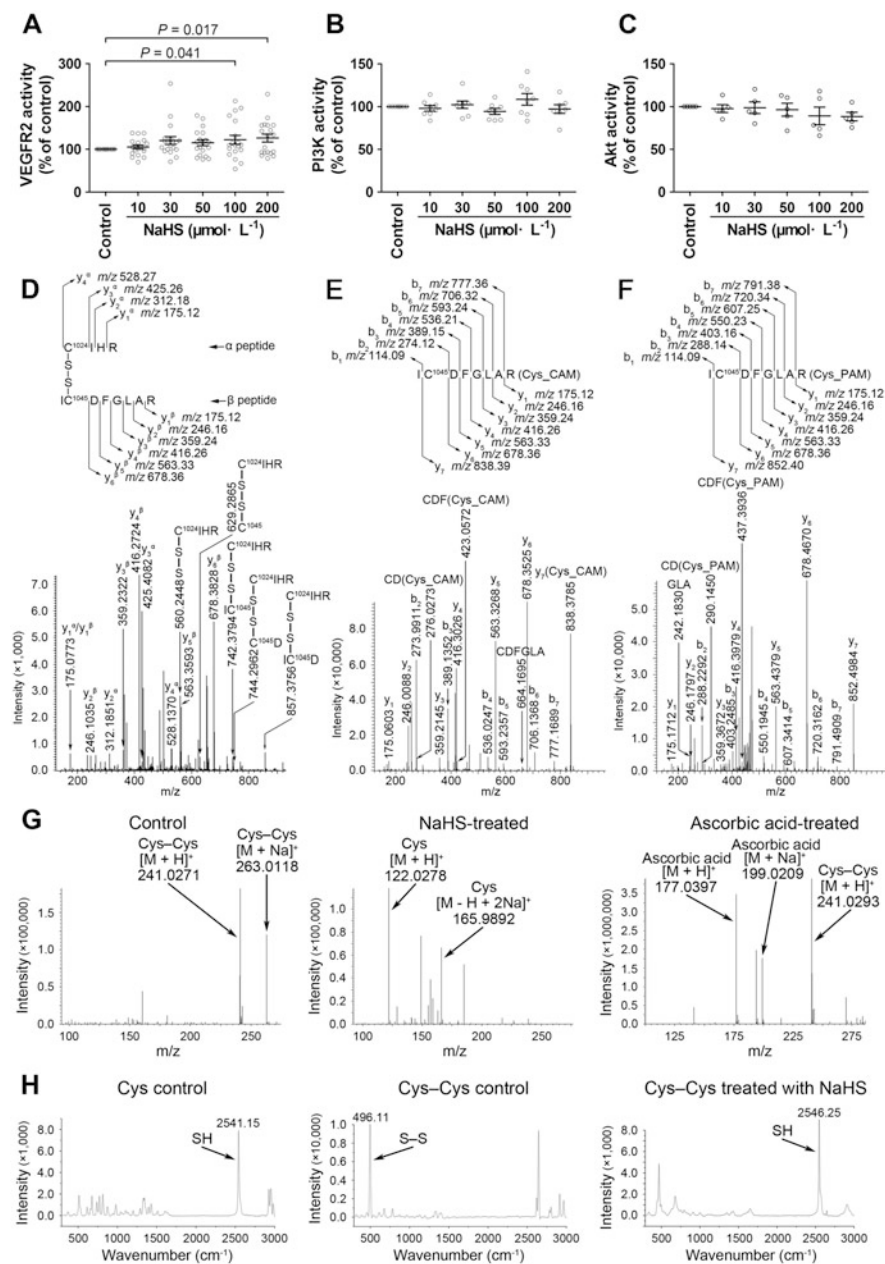
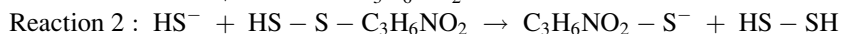
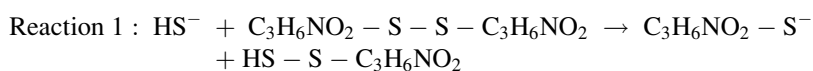


Fig. 3 (a-c) NaHS directly activated VEGFR2 without affecting the activity of PI3K and Akt in a cell-free system. (d) CID spectra of [M + 3H]³⁺ m/z 473.90 from a tryptic digest of VEGFR2 in the absence of NaHS showing an S-S bond between Cys1045 and Cys1024. (e) CID spectra of [M + 2H]²⁺ m/z 476.24 from a tryptic digest of VEGFR2 in the presence of DTT showing the β peptide containing Cys1045. (f) CID spectra of [M + 2H]²⁺ m/z 483.25 from a tryptic digest of VEGFR2 in the presence of NaHS showing the β peptide containing Cys1045. Cys_CAM,

spectrometry. However, a new disulfide bond between Cys1045 and Cys1024 was surprisingly discovered in the intracellular kinase core of VEGFR2, and this disulfide bond was cleaved by H₂S (Fig. 3d, e, f). More mass spectrometry and Raman spectroscopy experiments were performed to testify the effects of H₂S on each individual amino acid contained in protein molecules and the disulfide bonds contained in model chemicals. The results further validate our conclusion that H₂S does not modify any free amino acid residues in VEGFR2, however, cleaves the disulfide bonds (Fig. 3g, h). In order to further investigate the role of the Cys1045-Cys1024 disulfide bonds in VEGFR2, we mutated Cys1045 to alanine and acquired the mutant VEGFR2-C1045A. The activity of the recombinant VEGFR2-C1045A was significantly higher than the wild-type VEGFR2. The data suggest that Cys1045-Cys1024 is an inhibitory intrinsic motif in VEGFR2. Molecular dynamic calculations further showed that the breaking of Cys1045-Cys1024 disulfide bond restored the activity of VEGFR2 by shifting it to an active “Phe-in” conformation which is necessary for ATP binding (Fig. 4a, b). The data indicate that the Cys1045-Cys1024 disulfide bond actually serves as a molecular switch for H₂S to regulate the structure and function of its “receptor”, VEGFR2. The next question is how H₂S specifically targets this molecular switch? We further performed quantum chemical studies and mass spectrometry at various pH values and different time points. The data revealed that it was HS⁻ (the aqueous solution of H₂S is actually a mixture of H₂S and HS⁻), but not H₂S, that breaks the disulfide bond molecular switch (Fig. 4c, d, e, f). Quantum chemical calculations showed that HS⁻ specifically recognized the sulfur atoms of the disulfide bond and attacked one of the sulfur atoms of the disulfide bond. This was a nucleophilic attack where HS⁻ attacked the S–S bond via interaction with its frontier molecular orbitals: the highest occupied molecular orbital (HOMO) of the nucleophile HS⁻ and the lowest unoccupied molecular orbital (LUMO) of the electrophile under attack. Two steps of reactions and two HS⁻ anions are required to cleave one S–S bond (Tao et al. 2013):



With these data we conclude that H₂S/HS⁻ targets its “receptor” with a mechanism beyond the typical ligand-receptor docking. The targeting is based on an interaction of the electron orbitals of the nucleophile sulfur atom of H₂S/HS⁻ and the electrophile sulfur atom of the S–S bond. This is actually a new atomic biological mechanism based on the morphology and energy of the frontier molecular orbitals involved in the reaction.

Fig. 3 (continued) Carboxyamidomethyl cysteine; Cys_PAM, propionamide cysteine. (g and h) ESI-MS spectra and Raman bands showing hydrogen sulfide-induced cleavage of the S–S bond in the model chemical of Cys–Cys

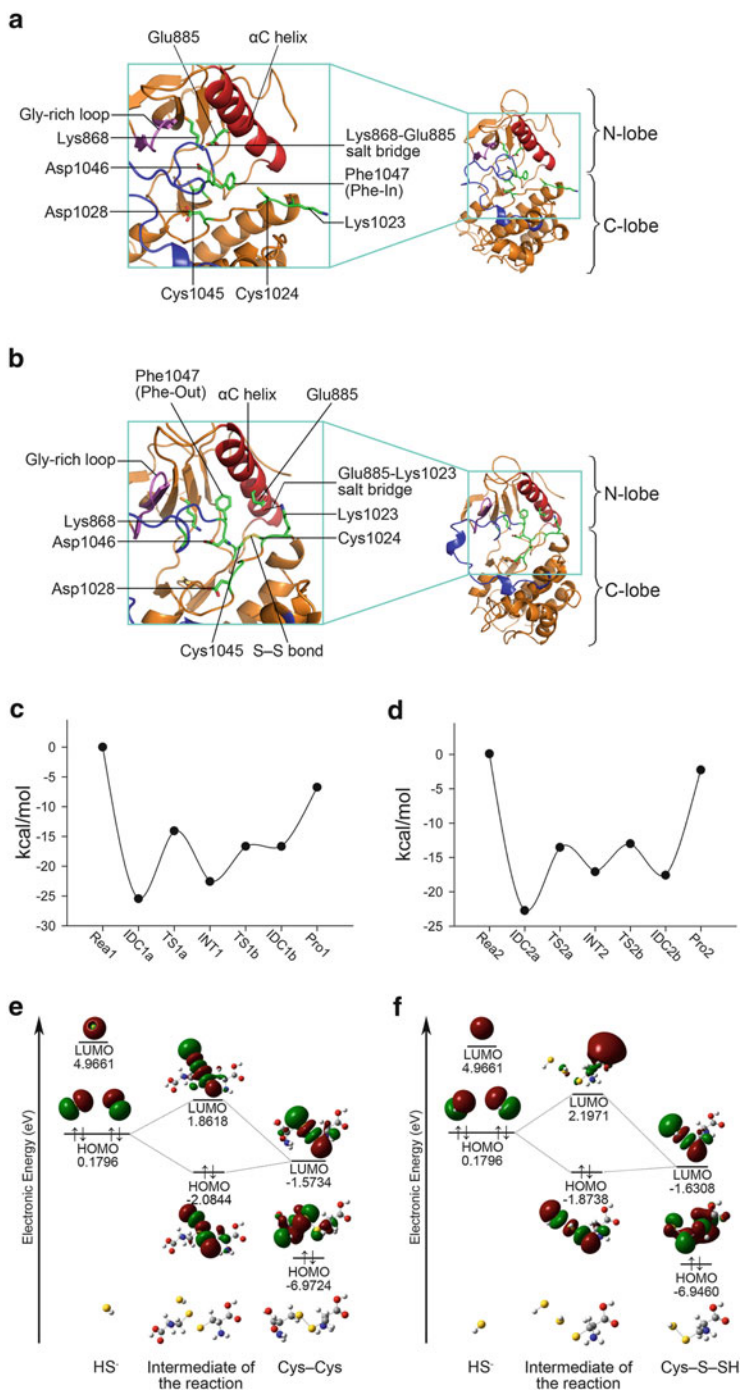


Fig. 4 (a and b) Molecular dynamics simulation showing conformation of the kinase core of VEGFR2. (a) There is a conserved salt bridge between Lys868 and Glu885 in VEGFR2 without the Cys1045-Cys1024 S-S bond. The DFG motif is in the “Phe-In” conformation. (b) The Lys868-

7 The Receptor Tyrosine Kinase Family Are Candidate “Receptors” for H₂S

Since the direct target of H₂S is actually the disulfide bond, it is reasonable that there might be other “receptors” for H₂S. If a protein contains a functional disulfide bond, this protein might be a candidate “receptor” for H₂S. It is known that VEGFR2 belongs to the family of receptor tyrosine where various members with different extracellular ligand-binding domain have a similar intracellular kinase domain. The Cys1045-Cys1024 molecular switch is located in the intracellular kinase domain of VEGFR2. H₂S may penetrate through cell membrane and directly activate the intracellular kinase domain without interacting with the extracellular ligand-binding domain of VEGFR2. This gives rise to an idea that other members of the receptor tyrosine kinase family may also contain such a molecular switch and thereby serving as potential “receptors” for H₂S. Interestingly, we have recently found that H₂S also directly activates EGFR (epidermal growth factor receptor) (Ge et al. 2014) and the insulin receptor (Xue et al. 2013), both of which belong to the receptor tyrosine kinase family (Fig. 5a, h). Some disulfide bond molecular switch was detected in EGFR which was labile to H₂S-induced cleavage (Fig. 5b, c). To find the precise location of the molecular switch in EGFR, mutation of some potential cysteine residue which is necessary to form a disulfide bond molecular switch was performed. The results showed that the Cys798 residue located in the intracellular kinase domain of EGFR is required to form the disulfide bond molecular switch for H₂S though the other cysteine residue required for the switch remains to be identified (Fig. 5d, e, f, g). Moreover, we have also shown that endogenous H₂S was required for VEGF to activate VEGFR2 in the vascular endothelial cells. This was evidenced by experiments using siRNA-mediated knockdown of CSE where the proangiogenic effect of VEGF was blunted in the vascular endothelial cells with CSE knockdown (Tao et al. 2013). VEGFR2 does not respond to its extracellular ligand without endogenous H₂S. The receptor tyrosine kinase family may be renamed as H₂S-dependent receptor tyrosine kinase family. In addition to the receptor tyrosine kinase family, some phosphatases, including protein tyrosine phosphatases (PTPs), the lipid phosphatase (PTEN), and low-molecular-weight PTPs, have functional disulfide bonds which are sensitive to redox regulation (Salmeen and Barford 2005). Those phosphatases all contain cysteine residues that are prone to form disulfide bonds or other oxidized forms such as a sulfonamide bond. Cho et al. found a redox regulation of disulfide

Fig. 4 (continued) Glu885 salt bridge is disrupted and the DFG motif rotates anticlockwise from the “Phe-In” conformation in VEGFR2 with the Cys1045-Cys1024 S–S bond. (c) Potential energy surface for the nucleophilic attack of HS[−] on Cys–S–S–Cys (reaction 1) (IDC1a, the ion dipole complex 1a; TS1a, transition state 1a; INT1, intermediate 1; TS1b, transition state 1b; IDC1b, ion dipole complex 1b). (d) Potential energy surface for the nucleophilic attack of HS[−] on Cys–S–SH (reaction 2) (IDC2a, the ion dipole complex 2a; TS2a, transition state 2a; INT2, intermediate 2; TS2b, transition state 2b; IDC2b, ion dipole complex 2b). (e and f) Quantum chemical mechanisms underlying hydrogen sulfide-induced breaking of the S–S bond with the model chemical Cys–Cys

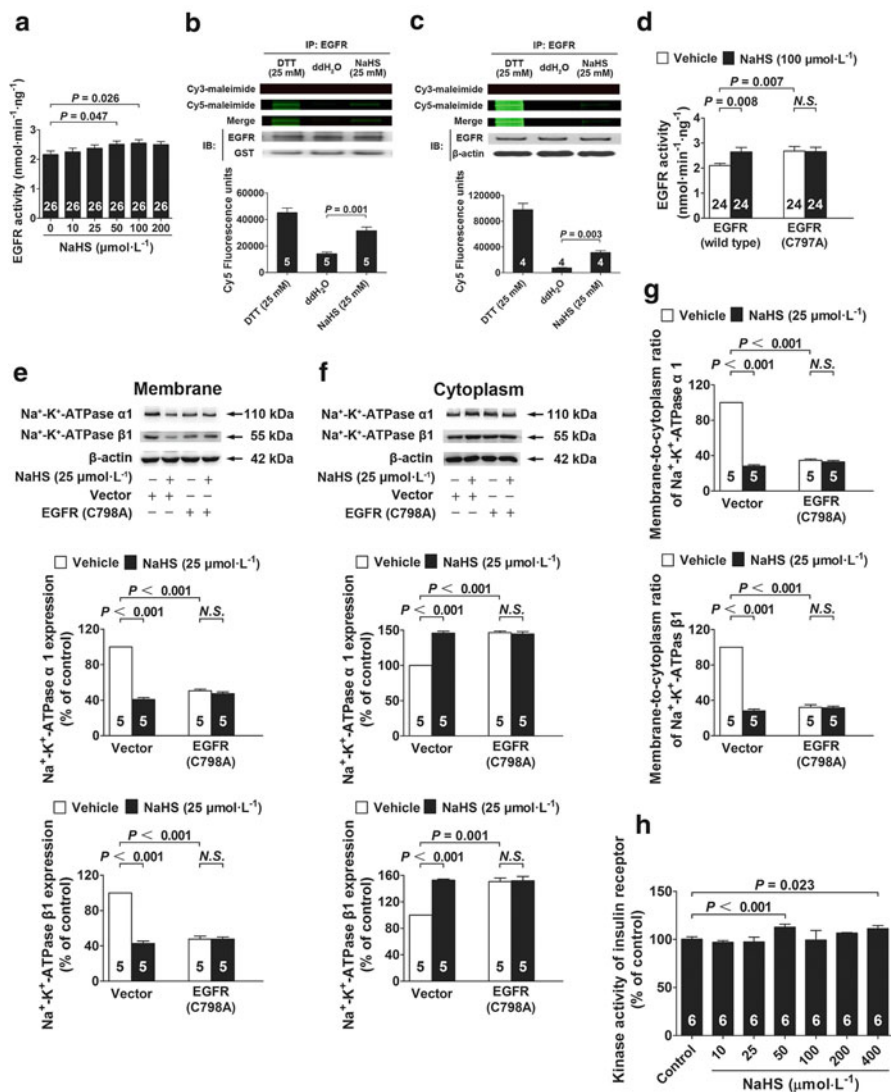


Fig. 5 H₂S directly activates EGFR (a) and breaks its disulfide bond 9 (b and c). NEM was used to cover all -SH groups of Cys residues in the intracellular kinase domain of recombinant EGFR proteins (b) or EGFR immunoprecipitated from cell lysate (c). Free thiols in EGFR were labeled firstly by Cy3-conjugated maleimide and then by Cy5-conjugated maleimide after DTT or NaHS treatment. (d-g) EGFR Cys797 (human) or Cys798 (rat) is required for H₂S to activate EGFR and induce Na⁺/K⁺-ATPase endocytosis. (d) Mutation at Cys797 in human recombinant EGFR proteins. (e-g) Rat renal tubular epithelial cells transfected with the vectors expressing the mutant EGFR (C798A) or control vectors. (h) H₂S directly increases activity of insulin receptor

bonds in PTEN (C71-C124), Cdc25A (C384-C430), and LMW-PTP (C12-C17) (Cho et al. 2004). Therefore, it is possible that these phosphatases might also function as “receptors” for H₂S.

Moreover, it is reported that a disulfide bond could be formed between two protein molecules to regulate their function. For example, a disulfide bond is formed between Cys32 in the active site of Trx-1 and Cys212 at the C-terminal of PTEN and this binding inactivates PTEN (Song et al. 2007; Meuillet et al. 2004). The data suggest that a protein complex may also function as potential “receptor” for H₂S with some disulfide bond between two protein molecules serving as molecular switch for H₂S.

8 Two Theories: Disulfide Bond Molecular Switch Versus Protein S-sulfhydration

Mustafa et al. proposed an S-sulfhydration theory to explain the interaction between H₂S and its target molecules (Mustafa et al. 2009). According to the S-sulfhydration theory, H₂S oxidizes the –SH group of the cysteine residues to yield an –S–SH group and thereby change the conformation and function of a protein molecule. Future works are required to clarify how H₂S could oxidize the –SH group and how such a modification at the cysteine residues would change the conformation and function of a protein molecule. Another interesting question is does H₂S-induced oxidization of the cysteine residues (S-sulfhydration) have selectivity for different cysteine residues in a protein molecule? In case of VEGFR2, there are 11 cysteine residues in the intracellular kinase domain. Does H₂S oxidize all the cysteine residues or some selective cysteine residues in VEGFR2 and how such a modification would activate this kinase? We have examined this idea using mass spectrometry and did not find any S-sulfhydration in the kinase domain of VEGFR2 (Fig. 3f) (Tao et al. 2013). Actually H₂S did not cause chemical modification on any of the 20 amino acids that constitute protein molecules in a cell-free system (Tao et al. 2013). However, H₂S cleaved disulfide bonds in various model chemicals (Tao et al. 2013). It is possible that H₂S might be oxidized to sulfane sulfur (Bailey et al. 2014). It is the oxidants of H₂S to oxidize the free thiols of the cysteine residues in a protein molecule to yield S-sulfhydration of a protein (Greiner et al. 2013). However, this is actually the effects of the oxidants of H₂S. Indeed, it remains to be clarified if the H₂S effects are mediated by the oxidants of H₂S but not H₂S itself.

A more interesting field about the role of element sulfur is now arising. We believe that element sulfur can be transformed to a reductive form like H₂S or an oxidative form like sulfane sulfur depending on the microenvironment in the body. For instance, H₂S can be easily oxidized under normoxic conditions but will remain in a reductive form during hypoxia. The role of endogenously generated H₂S may be tissue/cell type specific.

All these hypotheses including the disulfide bond molecular switch proposed by us remain to be validated in future studies. This will surely prompts the field of H₂S biology to a new landmark.

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Hydrogen Sulfide and Platelets: A Possible Role in Thrombosis

Michael Emerson

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Abstract

Platelets are circulating blood elements with key roles in haemostasis and thrombosis. Platelets are activated by a range of stimuli including exposed subendothelial components. Haemostasis also depends upon the effects of inhibitory substances, including the gasotransmitter nitric oxide whose effects on platelets are well documented. Evidence is also emerging to suggest that H₂S is generated enzymatically by platelets and can impact their function. Exposure of platelets to H₂S from slow-release compounds inhibits aggregation and exerted anti-thrombotic effects in vivo. The mechanisms by which H₂S impacts platelet function and the importance of interactions between H₂S and other gasotransmitters remain unclear. H₂S is therefore emerging as a potentially important regulator of platelet activation and thrombosis. Further study is required to evaluate its importance as a regulator of platelet physiology and associated pathological conditions such as myocardial infarction and stroke.

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Keywords

Cardiovascular • Haemostasis • Hydrogen sulfide • Nitric oxide • Platelet • Thrombosis

1 Platelet Development and Structure

Platelets are anucleate circulating blood elements derived from pluripotent megakaryocytes in bone marrow. Platelets are 2–3 μm in diameter and circulate for 5–9 days before undergoing phagocytosis in the spleen and liver.

Platelet structure and morphology are maintained by the cytoskeleton which consists of three discrete parts, the spectrin-rich membrane skeleton, which includes the open canalicular system and maintains membrane integrity. The other cytoskeletal components are the actin-rich cytoskeleton, which generates contractile forces, and the microtubule ring, which maintains the discoid shape of the resting platelet. Platelets are also distinctly rich in intracellular vesicles which, upon activation, release a range of mediators that drive adhesion, aggregation and immune responses. Platelets also contain a specialised endoplasmic reticulum called the dense tubular system which stores calcium. A distinct feature of the platelet external membrane is the presence of a range of receptors which upon occupation drive the process of platelet activation. Platelets express receptors for subendothelial proteins such as collagen; the product of the coagulation cascade, thrombin; substances released from granules such as ADP and products of enzyme activity such as thromboxane A_2 derived from cyclooxygenase. The basic platelet intracellular structure is shown in Fig. 1.

2 Platelet Activation and Role in Haemostasis

The primary role of platelets is to maintain vascular integrity and cease blood loss following injury. Platelets circulate in a quiescent state, but upon vascular injury and exposure of blood to the subendothelial matrix, platelet surface receptors bind to exposed collagen and other macromolecules leading to adhesion of platelets to the vessel wall. Adhesion initiates a cascade of signalling events culminating in calcium release. The rise in intracellular calcium leads to rearrangement of the cytoskeleton and extension of filopodia so that the platelet changes from a spherical disc to a stellate shape with an increased surface area. In addition, intracellular granules release their contents to propagate the platelet activation process by activating adjacent platelets. Platelet aggregation, the linking of adjacent platelets by fibrinogen, follows calcium-mediated activation of the surface integrin $\alpha\text{IIb}\beta\text{III}$ leading to the formation of a platelet plug which seals the breached vessel.

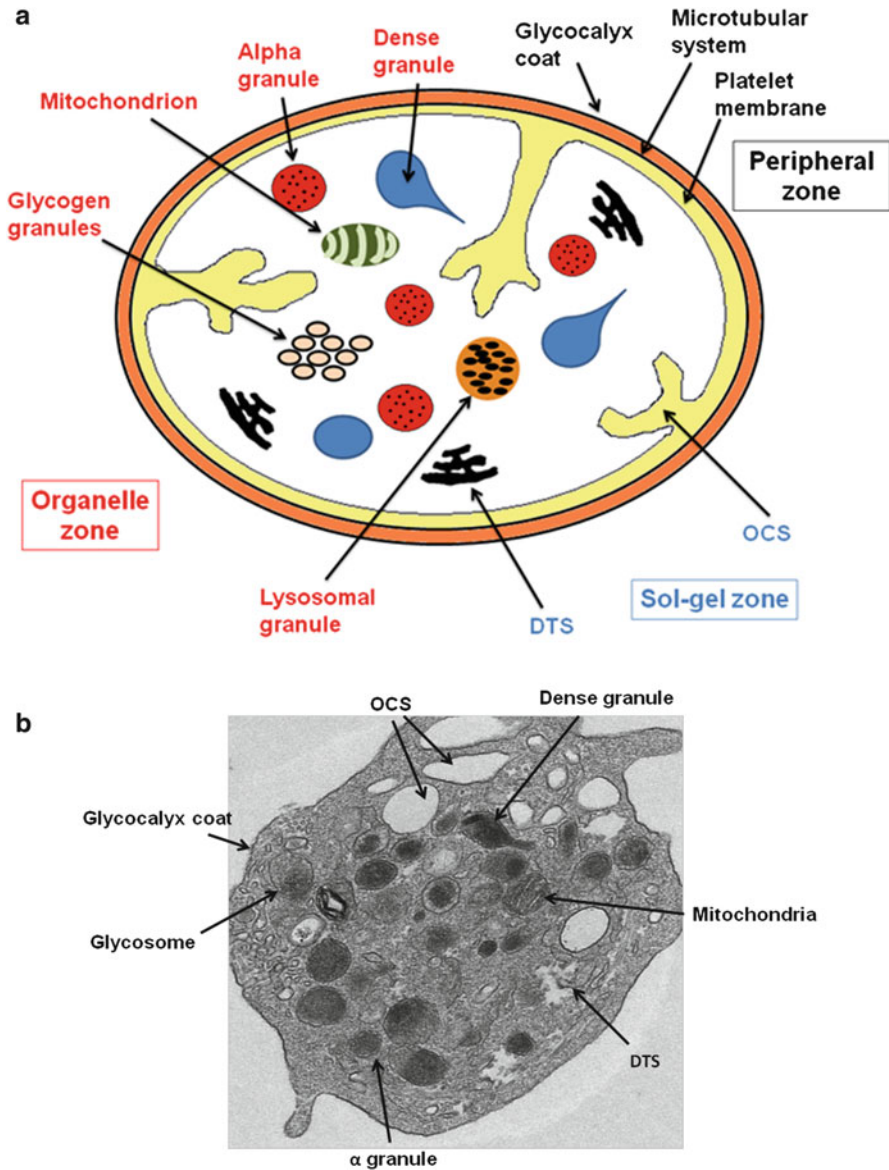


Fig. 1 Platelet intracellular morphology. (a) Basic diagram demonstrating the structure and organisation of platelets. DTS-dense tubular system; OCS-open canalicular system. (b) Electron microscopy image of a platelet from author's laboratory (Emerson, unpublished)

3 Platelets and Thrombosis

Thrombosis is characterised by the formation of a deposit of blood constituents (the thrombus) in a blood vessel leading to partial or full occlusion and results in events including myocardial infarction or stroke. Arterial thrombosis is a high-shear, platelet-dependent process that occurs when blood is exposed to sites of arterial injury such as atherosclerotic plaque rupture. Surgical devices used in revascularisation can also provide a surface that initiates thrombosis. Despite the use of anti-thrombotic drugs, such as aspirin, which targets platelet cyclooxygenase, arterial thrombosis and resulting conditions such as myocardial infarction, unstable angina and stroke remain the most common end points in atherosclerotic disease. In summary, inappropriate activation of platelets is a key driver of common cardiovascular diseases, and platelets are a major therapeutic target in the prevention and treatment of conditions such as myocardial infarction. Limitations in the efficacy of platelet-targeted antithrombotics are due to drug resistance in a high proportion of individuals and incidences of bleeding associated with use of antithrombotics (Schroeder et al. 2006).

4 Nitric Oxide and Platelets

Of the main gasotransmitters, the best characterised in terms of impact upon platelet function is nitric oxide (NO). NO is a major negative regulator of platelet function and acts to counterbalance the positive drivers of platelet activation during both haemostatic and thrombotic platelet responses. NO impacts a range of platelet activation processes including shape change, secretion and calcium signalling. NO acts in platelets predominantly via activation of soluble guanylyl cyclase and subsequent PKG activation resulting in interaction with a wide range of target proteins which explains its diverse inhibitory activity. For review, see Naseem and Roberts (2011).

NO generated by endothelial nitric oxide synthase (eNOS) within endothelial cells is released into surrounding tissues such as vascular smooth muscle and the blood. NO activity on vascular smooth muscle cells induces vasodilation (Huang et al. 1995) and so critically regulates vessel tone and blood pressure. NO has been reported to reduce leukocyte adherence to the vessel wall (Kubes et al. 1991; Lefer et al. 1999), which potentially accounts for its protective effect against atherogenesis (Kuhlencordt et al. 2001). NO can inhibit platelet aggregation *in vitro* in whole blood (Yoshimoto et al. 1999) and *in vivo* (Emerson et al. 1999; Freedman et al. 1999; Moore et al. 2010).

Determining the role of endogenous NO and eNOS in the regulation of platelets and thrombosis has been somewhat controversial, with conflicting reports published. A lack of thrombotic phenotype in eNOS^{-/-} mice has been reported by a number of groups (Dayal et al. 2006; Ozuyaman et al. 2005), whereas others contradict these reports by describing both enhanced and blunted thrombotic responses (Freedman et al. 1999; Heeringa et al. 2000; Iafrazi et al. 2005). Some

clarity in the field eventually emerged when real-time monitoring of the duration of the platelet aggregation response *in vivo* showed that loss of eNOS led to an effect primarily on the duration rather than the amplitude of the platelet aggregation response (Moore et al. 2010).

A second area of controversy has surrounded the issue of the expression and functional relevance of eNOS within platelets. Researchers have reported the presence of the L-arginine/NO pathway in platelets (Freedman et al. 1997; Radomski et al. 1990) and NO release from platelets *in vitro* (Malinski et al. 1993; Zhou et al. 1995). Furthermore, the hypercoagulability seen in eNOS^{-/-} mice was restored by the transfusion of wild-type platelets suggesting that platelets generated NO from eNOS (Freedman et al. 1999). eNOS protein and mRNA expression have been reported in platelets (Aytekin et al. 2012; Berkels et al. 1997; Mehta et al. 1995; Patel et al. 2006); however, the presence of eNOS in platelets is a subject of fierce debate. Contrary to the above studies, other research groups have reported the lack of eNOS mRNA and protein expression in platelets (Gambaryan et al. 2008; Ozuyaman et al. 2005; Tymvios et al. 2009). In fact, Gambaryan et al. (Gambaryan et al. 2008) have shown that some commercially available eNOS antibodies apparently detect eNOS expression in eNOS^{-/-} mice. Gambaryan et al. (2008) emphasise the need to have appropriate positive and negative controls when detecting protein expression, a method adopted by Tymvios et al. (Tymvios et al. 2009), who showed a lack of eNOS protein in human and mouse platelets and perhaps more importantly, the lack of a functional role of platelet NOS in regulating aggregation *in vitro* regardless of any argument regarding expression of NOS protein.

5 Hydrogen Sulfide and Platelets

The study of the biological relevance of H₂S in the regulation of platelets and thrombosis is in its infancy, and there is currently no consensus on the regulation of platelets by H₂S. Gaining an understanding of the biological roles of NO has been dependent upon studies with NO donors and NO synthase inhibitors. The roles of H₂S are, in contrast, unclear due to limited biological tools with which to explore its effects. As improved tools such as slow-release H₂S donors that mimic physiological exposure to H₂S and selective inhibitors of H₂S generating enzymes are developed and applied, the roles of H₂S in regulating platelets are likely to become more clear.

H₂S generating salts have been shown to inhibit platelet aggregation induced by a range of agonists including ADP, collagen, adrenaline, arachidonic acid and thrombin (Zagli et al. 2007). It is tempting therefore to speculate an inhibitory role of H₂S similar to that established for NO. These data cannot, however, be used to infer an inhibitory role of H₂S physiologically since the concentrations used (100 μM to 10 mM) were considerably higher than those likely to be found physiologically. A mechanism by which H₂S may inhibit platelets was not proposed so that a lack of clarity remains. A more recent study similarly found H₂S

generating salts in the μM to mM range to inhibit collagen-induced human platelet aggregation (Zhong et al. 2014). Inhibition of aggregation was associated with reduced secretion, impaired mobilisation of calcium from intracellular stores and reduction of a range of collagen-induced signalling events including PLC γ 2, Akt and MAPK phosphorylation. In this study, H_2S did not affect platelet viability, suggesting an inhibitory effect mediated via suppression of key signalling events downstream of collagen activation rather than a toxicologic effect on the platelet.

In contrast to the above studies, H_2S generating salts at lower concentrations (0.1–100 μM) have been reported to enhance human platelet aggregation induced by the thrombin receptor activator peptide (TRAP-6) and to contribute to the enhanced platelet aggregation occurring during hyperhomocysteinemia (d'Emmanuele di Villa Bianca R et al. 2013). A full causative mechanism was not described to explain the ability of H_2S to enhance platelet aggregation although the arachidonic acid cascade was shown to be implicated. The studies reported to date therefore report both inhibitory and stimulatory effects of H_2S -generating salts upon platelet aggregation responses. The reports cannot be described as contradictory due to differences in the concentrations of H_2S -generating salts used in the studies and the varying choices of platelet agonists. Nonetheless, further study is warranted to clarify the role of H_2S in regulating platelet activation and aggregation.

Given the slow manner of synthesis of H_2S from enzymes (Li et al. 2008), experiments with rapidly and transiently generating sulfide salts such as those described above may be of limited value in determining the physiological role of H_2S in regulating platelets. A limited study with the slow-release compound GYY4137 showed an inhibitory effect of H_2S upon human platelet aggregation at equivalent H_2S concentrations in the nM range (Ilkan et al. 2013). In addition, GYY4137 reduced activation-induced platelet adhesion molecule expression and attenuated the morphological signs of platelet activation (Grambow et al. 2013). *In vivo*, GYY4137 prolonged venular thrombus formation and tail-vein bleeding (Grambow et al. 2013) suggesting anti-thrombotic roles that may be in part platelet mediated. Further study is warranted, however, in order to clarify any physiological role of H_2S in the regulation of platelets at physiologically relevant concentrations.

As well as determining the functional roles of H_2S in regulating platelets, it is important to consider whether H_2S -generating enzymes are expressed by platelets and whether H_2S is generated endogenously by platelets. H_2S is synthesised by CBS and CSE in vascular endothelial cells (Shibuya et al. 2009; Whiteman et al. 2011) and has been shown to act as a physiologic vasodilator and regulator of blood pressure (Yang et al. 2008). Human platelets have been reported to possess both CBS and CSE by western blotting and to generate detectable amounts of H_2S from *L*-cysteine (d'Emmanuele di Villa Bianca R et al. 2013). In contrast, we have detected robust CBS expression by western blotting but very little or no CSE protein in human platelet lysates using three commercially available antibodies (unpublished data). There, therefore, appears to be a current consensus that platelets generate H_2S endogenously although the enzymatic source remains in dispute. In addition, application of *L*-cysteine, the substrate of H_2S generating enzymes, to

platelets has been shown to modulate their function (d'Emmanuele di Villa Bianca R et al. 2013) suggesting functional relevance of endogenous H₂S in the regulation of platelet activation and potentially thrombosis.

The fact that platelets express H₂S generating enzymes and generate H₂S suggests that further study of the role of H₂S from both exogenous and endogenous sources should be pursued. In addition, it is important to highlight that platelets are highly sensitive to gasotransmitters released from the vasculature. In the NO field, the major source of enzymatic NO impacting platelets is that generated by the vascular endothelium rather than the platelet itself (Tymvios et al. 2009). H₂S from the vasculature should therefore be investigated as a modulator, not only of vascular tone and blood pressure but as a regulator of platelet function and thrombosis. This will require *in vivo* studies, including work with models of thrombosis and *in vivo* platelet aggregation.

6 H₂S and Thrombosis/Disease

H₂S from GYY4137 has been investigated in a mouse model of microvascular thrombus formation following vascular injury (Grambow et al. 2013). In this chapter, H₂S delayed venular thrombus formation suggesting a potential anti-thrombotic effect. The current goal of anti-thrombotic research is to develop compounds with anti-thrombotic efficacy but limited impact upon haemostatic function. In this way, effective anti-thrombotic therapy can be achieved without the bleeding complications associated with current therapeutics such as aspirin. In this study, GYY4137 increased tail bleeding time threefold. Since tail bleeding is used as an index of haemostatic function, further study is required to evaluate the comparative efficacy of H₂S releasing compounds relative to established therapeutics.

In addition to exerting direct effects on thrombus generation, H₂S has also been suggested to exert antiatherogenic effects in studies involving the development of atherosclerotic lesions in ApoE knock-out mice (Wang et al. 2009). A number of mechanisms have been suggested by which H₂S may attenuate atherogenesis including inhibition of superoxide formation (Muzaffar et al. 2008) and modification of the atherogenic modification of LDL by hypochlorite (Laggner et al. 2007). If confirmed in human studies, H₂S may prove to be an important driver not only of platelet function and thrombus formation but a determinant of the atherogenic process driving platelet dysfunction during atherothrombotic diseases.

7 Current State of Play and Future Directions

Although limited in terms of quantity and depth of study, there is a reasonable body of evidence suggesting that H₂S is both generated by platelets and impacts platelet function. The enzymatic source of H₂S in platelets remains unclear but given that these issues remain unresolved in the NO field, this is not a surprising state of

affairs. Greater clarity of the role of H₂S in regulating platelet function and the mechanisms of action remain unresolved but are likely to become more clear as recently developed pharmacological tools are applied to the field.

One area of major interest to the platelet field is the interaction between NO and H₂S. Recently, NO and H₂S have been shown to interact in the regulation of angiogenesis, whereby H₂S amplifies NO signalling by inhibiting downstream phosphodiesterase (PDE) (Bucci et al. 2010; Coletta et al. 2012). Platelets are rich in PDE, and the importance of NO in the regulation of platelets as well as the emerging importance of H₂S as a modulator of platelet function suggests that interaction of gasotransmitters such as NO in the regulation of platelets and thrombosis should be considered. Evidence of the likely importance of the interaction between H₂S and NO recently emerged when the anti-thrombotic effect of H₂S was shown to be mediated in part by upregulation of NOS in a mouse model of thrombosis (Kram et al. 2013).

Currently, gasotransmitters are not direct targets for anti-thrombotic therapeutics, but their importance is highlighted not only by studies linking both NO and H₂S with platelet function and thrombosis but by their potential interactions with existing therapeutics, particularly prescribed antithrombotics. This is highlighted by the demonstration that blockade of the platelet P2Y₁₂ receptor, a standard therapeutic intervention, enhances the inhibitory effects of NO upon platelet function (Kirkby et al. 2013). Thus, it is essential to comprehensively analyse the effects of H₂S upon platelets, thrombosis and its interaction with anti-thrombotic reagents in order to fully understand the roles of H₂S in haemostasis and thrombosis.

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Part IV

Inflammation and Inflammatory Disease

H₂S and Inflammation: An Overview

Madhav Bhatia

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Abstract

Inflammation is a response to traumatic, infectious, post-ischemic, toxic, or autoimmune injury. However, uncontrolled inflammation can lead to disease, and inflammation is now believed to be responsible for several disease conditions. Research in our laboratory has shown that hydrogen sulfide (H₂S) acts as a novel mediator of inflammation. At present, work in several research

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groups worldwide is focused on determining the role of H₂S in inflammation. H₂S has been implicated in different inflammatory conditions. Most of this research involved working with animal models of disease and in vitro systems. Recent research, however, points to a role of H₂S in clinical inflammatory disease as well. This chapter describes our current understanding of the role of H₂S in inflammation.

Keywords

Hydrogen sulfide • Inflammation • Systemic inflammatory response syndrome • Leukocytes

1 Introduction

Inflammation is a response to traumatic, infectious, post-ischemic, toxic, or autoimmune injury. It is a highly orchestrated, tissue-based process, characterized by “rubor” (redness), “calor” (heat), “dolor” (pain), and “tumor” (swelling). Inflammation is a normal response to injury and a useful physiological event. However, uncontrolled inflammation can lead to disease. Recent research has shown a key role of hydrogen sulfide (H₂S) in inflammation. This has been possible with the use of experimental approaches, such as cell culture systems and in vivo disease models. At the time of writing this chapter, evidence has started emerging on the role of H₂S in clinical inflammatory disease.

In this chapter, our current understanding of the role of H₂S in inflammation is discussed.

2 Inflammation

Currently, inflammation is an important research question in systems biology in the academia, as well as a multibillion dollar market for the pharmaceutical industry. In a disease in which primary pathogenic events are unknown, control of inflammation is the next best option.

In response to the initial insult/injury, leukocytes are activated. A key component of the inflammatory process is the trafficking of inflammatory cells to the site of injury/infection. Cytokine/receptor interactions on the surface of these cells lead to the expression of gene products that bring about the inflammatory response. However, uncontrolled production of inflammatory products is injurious to host cells and, in some cases, can lead to cancer. Therefore, endogenous mechanisms have evolved to limit the production of inflammatory molecules and permit the resolution of the inflammatory response. An understanding of these mechanisms is important because defects in the pathway may contribute to inflammatory disorders, and the pathway itself may present targets for novel anti-inflammatory therapeutic strategies (Bhatia and Moochhala 2004; Bhatia 2010, 2012; Hegde and Bhatia 2011; Nathan 2002; Mazumder et al. 2010).

3 H₂S and Inflammation

Over the years, various studies have indicated a role of H₂S in the inflammatory process. In mammals, including humans, H₂S is produced by the action of the enzymes, cystathionine- β -synthase (CBS), cystathionine- γ -lyase (CSE), cysteine aminotransferase (CAT, EC 2.6.1.3), (followed by 3-mercaptopyruvate sulfurtransferase (MPST, EC 2.8.1.2), and cysteine lyase (EC 4.4.1.10), (Bhatia 2012; Li et al. 2011; Moore et al. 2003; Wang 2012). As the end product of CBS- and CSE-catalyzed cysteine metabolism, H₂S exerts a negative feedback effect on the activity of these enzymes (Bhatia 2012; Moore et al. 2003). Endogenous H₂S synthesized by CSE has been shown to be primarily responsible for its inflammatory action.

Recent work in our laboratory and others has shown a key role of H₂S as a mediator of inflammation in different clinical conditions. Current understanding of the role of H₂S in different inflammatory conditions is summarized in Table 1.

4 Disease Conditions with Role of H₂S in Inflammation

4.1 Acute Pancreatitis

Acute pancreatitis is a common clinical condition, the incidence of which has been increasing worldwide over recent years (Bhatia et al. 2000, 2005c; Bhatia 2012). For example, in the United States alone, acute pancreatitis is the most common reason for hospitalization (274,119 discharges in the year 2009, a 30 % increase over 2000) amongst all gastrointestinal diseases (Peery et al. 2012). It also inflicts a heavy economic burden; the direct cost in the United States alone in the year 2009 was US\$2,599,686,000 (Peery et al. 2012). Acute pancreatitis was the cause of death in 3065 cases and a contributing cause in an additional 5500 deaths in the year 2009 (Peery et al. 2012).

Most cases of acute pancreatitis are secondary to biliary disease or excess alcohol consumption. The exact mechanisms by which diverse etiological factors induce an attack are still unclear, but once the disease process is initiated, common inflammatory and repair pathways are invoked. There is a local inflammatory reaction at the site of injury, which if marked leads to systemic inflammatory response syndrome (SIRS), and it is this systemic response that is believed to be ultimately responsible for the majority of the morbidity and mortality (Bhatia et al. 2000, 2005c; Bhatia 2012). Lung injury, which is clinically manifested as acute respiratory distress syndrome (ARDS), is a major component of the multiple organ dysfunction syndrome (MODS) that results from SIRS in acute pancreatitis.

CBS and CSE, the two major H₂S synthesizing enzymes, are highly expressed in the pancreas. Endogenously produced H₂S has been shown as a mediator of inflammation in acute pancreatitis (Bhatia et al. 2005a). mRNA for CSE is expressed in mouse pancreas and that pancreas homogenates convert L-cysteine to H₂S *ex vivo*. Also, circulating levels of H₂S are increased in mice upon induction

Table 1 Role of H₂S in inflammatory disease

Disease model	Effect on inflammation	Reference
Cerulein-induced acute Pancreatitis in the mouse	Plasma H ₂ S is increased in inflammation. Treatment with the cystathione-gamma-lyase (CSE) inhibitor propargylglycine (PAG) protects against acute pancreatitis and associated lung injury	Bhatia et al. (2005a)
	Treatment with slow H ₂ S-releasing diclofenac protects mice against acute pancreatitis-associated lung injury	Bhatia et al. (2008a)
	Treatment with s-propargyl-cysteine (SPRC), a slow H ₂ S-releasing donor, protects against acute pancreatitis and associated lung injury	Sidhapuriwala et al. (2012)
	Knockout mice genetically deficient in CSE (CSE ^{-/-}) and protected against acute pancreatitis and associated lung injury	Ang et al. (2013)
Lipopolysaccharide (LPS)-induced endotoxemia in the mouse	Plasma H ₂ S levels, H ₂ S-synthesizing activity, and CSE expression are increased in inflammation. PAG treatment protects against inflammation	Li et al. (2005)
	Treatment with slow H ₂ S-releasing diclofenac protects mice against LPS-induced endotoxemia	Li et al. (2007)
	Treatment with morpholin-4-ium-4-methoxyphenyl(morpholino) phosphinodithioate (GY4137), a slow H ₂ S-releasing donor, protects mice against LPS-induced endotoxemia	Li et al. (2009)
Cecal ligation and puncture (CLP)-induced sepsis	Plasma H ₂ S levels, H ₂ S-synthesizing activity, and CSE expression are increased in inflammation. PAG treatment protects against sepsis. The H ₂ S donor sodium hydrogen sulfide (NaHS) further aggravates inflammation in sepsis	Zhang et al. (2006)
Carrageenan-induced hindpaw edema in the mouse	H ₂ S-synthesizing activity increased in inflammation. PAG treatment protects against inflammation in this model	Bhatia et al. (2005b)
	Treatment with slow H ₂ S-releasing diclofenac protects mice against carrageenan-induced hindpaw edema	Sidhapuriwala et al. (2007)
Burn injury-induced inflammation	Plasma H ₂ S levels, H ₂ S-synthesizing activity, and CSE expression are increased in inflammation. PAG treatment protects against burn injury-induced inflammation	Zhang et al. (2010)
Trinitrobenzene sulfonic acid-induced colitis	Slow H ₂ S-releasing mesalamine protects against inflammation in colitis	Fiorucci et al. (2007)
Direct administration of H ₂ S as NaHS to mouse	Direct administration of H ₂ S by intraperitoneal (i.p) administration of NaHS causes lung inflammation characterized by an increase in lung myeloperoxidase (MPO) activity and histological evidence of lung injury	Bhatia et al. (2006)

of acute pancreatitis. The conversion of L-cysteine to H₂S in pancreas homogenates is significantly reduced in mice pretreated with DL-propargylglycine (PAG) (Bhatia et al. 2005a). Furthermore, treatment of animals with PAG (either prophylactic or therapeutic) reduces the severity of pancreatitis as evidenced by a significant attenuation of hyperamylasemia, acinar cell injury/necrosis, and pancreatic myeloperoxidase (MPO) activity and by histological evidence of diminished pancreatic injury. Lung injury in severe acute pancreatitis is characterized by sequestration of neutrophils within the lung (i.e., increased lung MPO activity) and histological evidence of lung injury. Prophylactic or therapeutic administration of PAG additionally protected mice against acute pancreatitis-associated lung injury as evidenced by a significant attenuation of lung MPO activity and by histological evidence of diminished lung injury (alveolar thickening and leukocyte infiltration) (Bhatia et al. 2005a). The pro-inflammatory role of H₂S synthesized by CSE has been confirmed recently using knockout mice deficient in CSE (Ang et al. 2013). CSE knockout mice are protected against acute pancreatitis and associated lung injury, when compared to wild-type controls (Ang et al. 2013). These effects of CSE blockade/gene deletion suggested an important pro-inflammatory role of H₂S in regulating the severity of pancreatitis and associated lung injury and raised the possibility that H₂S may exert similar activity in other forms of inflammation. Following this study, an important role of CBS in the pathogenesis of acute pancreatitis and associated lung injury has been shown (Shanmugam et al. 2009). Acute pancreatitis was associated with increased plasma and tissue H₂S and ammonia (NH₃). Prophylactic or therapeutic administration of aminooxyacetate (AOA), a reversible inhibitor of CBS, directly inhibited CBS in the pancreas thereby reducing H₂S and NH₃ production, and protected against acute pancreatitis, showing a role of H₂S synthesized by CBS in inflammation in acute pancreatitis (Shanmugam et al. 2009).

4.2 Sepsis

Sepsis is defined as the presence of infectious organisms, such as bacteria, viruses, protozoa, or fungi, and/or their toxins in blood or tissue and the systemic response that follows. Sepsis leading to organ failure characterizes severe sepsis, and septic shock is defined by severe sepsis accompanied by hypotension unresponsive to fluid resuscitation. Severe sepsis and septic shock are one of the leading causes of mortality among intensive care unit and postoperative care patients (Bhatia 2012; Bhatia et al. 2009; Levy et al. 2010; Martin et al. 2003; Ramnath et al. 2006). Sepsis is a major health problem worldwide. The incidence of sepsis in North America, for example, is 3.0 per 1000 population, which transforms into an annual number of 750,000 cases, with 210,000 of them being fatal and a large socioeconomic burden (Bhatia 2012; Bhatia et al. 2009; Levy et al. 2010; Martin et al. 2003; Ramnath et al. 2006). The incidence of mortality due to sepsis is increasing, and the most likely causes are the increased incidence of resistant pathogens and the advances of medical and surgical procedures that save the lives of many patients but at the cost

of leaving them immunocompromized and in a state susceptible to death from severe sepsis and septic shock (Bhatia 2012; Bhatia et al. 2009; Levy et al. 2010; Martin et al. 2003; Ramnath et al. 2006).

H₂S has been shown to act as a mediator of inflammation in sepsis. In a clinically relevant mouse model of cecal ligation and puncture (CLP)-induced polymicrobial sepsis, liver CSE expression, H₂S synthesis, and plasma H₂S levels are significantly elevated (Zhang et al. 2006). Prophylactic, as well as therapeutic, administration of PAG significantly reduced sepsis-associated systemic inflammation, as evidenced by decreased MPO activity and histological changes in lung and liver and attenuated the mortality in CLP-induced sepsis (Zhang et al. 2006). On the other hand, administration of sodium hydrogen sulfide (NaHS), an H₂S donor, significantly aggravated sepsis-associated systemic inflammation (Zhang et al. 2006). Similar to CLP-induced sepsis, a pro-inflammatory action of H₂S has also been shown in lipopolysaccharide (LPS)-induced endotoxemia (Li et al. 2005; Collin et al. 2005). These studies show that H₂S plays a pro-inflammatory role in regulating the severity of sepsis and associated organ injury. H₂S has also been shown to modulate sinusoidal constriction in the liver and contribute to hepatic microcirculatory dysfunction during endotoxemia (Norris et al. 2013), and endogenous H₂S formation mediates the liver damage in endotoxemia in the rat (Yan et al. 2013).

4.3 Burn Injuries

Burn injuries are ranked among the leading causes of morbidity and mortality worldwide. In severe cases, burn injuries lead to SIRS, and associated MODS, which is a major contributor to death following burns (Bhatia 2012; Church et al. 2006; Endorf and Ahrenholz 2011). H₂S has been shown to act as a critical mediator of severe burn injury (to 25 % total body surface area full thickness burn)-induced inflammation in mice (Zhang et al. 2010). The result of this study shows that burn injury in mice resulted in a significant increase in plasma H₂S levels, liver and lung CSE mRNA expression, and liver H₂S-synthesizing activity. The enhanced H₂S/CSE pathway correlates with increased burn-associated systemic inflammation, as evidenced by increased MPO activity and histological evidence of lung and liver injury. There was protection against systemic inflammation and multiple organ damage by prophylactic or therapeutic treatment of PAG. Administration of NaHS at the same time of burn injury resulted in a further increase in MPO activity and more severe tissue injury in the lung and liver. These findings show that H₂S plays a key pro-inflammatory role in burn injury (Zhang et al. 2010).

4.4 Joint Inflammation/Arthritis

Joint inflammation/arthritis is a major health problem worldwide. In the United States, for example, back pain and arthritis (osteoarthritis, rheumatoid arthritis) are

amongst the most common and costly conditions, affecting more than 100 million individuals and costing more than US\$200 billion per year (Ma et al. 2014). The possible role of endogenous H₂S in the development of joint inflammation induced by intraplantar administration of carrageenan to the rat hindpaw has been investigated. An increase in H₂S synthesis in inflamed hindpaws was seen, suggesting a localized overproduction of H₂S during inflammation (Bhatia et al. 2005c). Pretreatment with PAG resulted in a dose-dependent inhibition of hindpaw edema as well as hindpaw MPO activity. These findings suggest that H₂S is an endogenous mediator of the development of hindpaw local inflammation. Recent reports in the literature in human patients point to a role of H₂S in rheumatoid arthritis, but there is a difference of opinion as to the pro- versus anti-inflammatory action of H₂S (Fox et al. 2012; Kloesch et al. 2012).

4.5 Colitis

Colitis, or inflammation of the colon, presents itself in several clinical forms, such as inflammatory bowel disease (IBD), ulcerative colitis (UC), and Crohn's disease (CD). Mucosal changes in colitis are characterized by ulcerative lesions accompanied by prominent inflammatory infiltrates in the bowel wall (Polytarchou et al. 2014). Mesalamine (5-aminosalicylic acid) is the first-line therapy for colitis. An H₂S-releasing derivative of mesalamine has been reported to protect against inflammation in trinitrobenzene sulfonic-induced colitis (Fiorucci et al. 2007) and against nociception in hapten-induced colitis (Coletta et al. 2012).

4.6 Chronic Obstructive Pulmonary Disease

Chronic obstructive pulmonary disease (COPD) is a common clinical condition characterized by progressive airflow obstruction that is only partly reversible, inflammation in the airways, and systemic effects or comorbidities. The main cause is tobacco smoking (Decramer et al. 2012). In patients with stable COPD, serum H₂S levels have been found to be significantly increased as compared to age-matched control subjects or those with acute exacerbation of COPD (AECOPD) (Chen et al. 2005). In this study, serum H₂S levels were negatively correlated with the severity of airway obstruction in patients with stable COPD, whereas they were positively correlated with the lung function in all patients with COPD and healthy controls. This study also reported that serum H₂S levels were lower in smokers than nonsmokers regardless of their health status (COPD or healthy controls). In addition, patients with AECOPD, whose serum H₂S levels were decreased, had greater neutrophil proportion but lower lymphocyte proportion in sputum than patients with stable COPD, suggesting a potential role of H₂S in regulating inflammatory response at different types or stages of COPD. This study (Chen et al. 2005) demonstrated that endogenous H₂S may participate in the development of airway obstruction in COPD and that the level of endogenous

H₂S may be correlated with the progression and severity of COPD. Also, there is a correlation between exhaled H₂S and exhaled nitric oxide (NO) in COPD (Sun et al. 2013).

5 Mechanisms of Action of H₂S in Inflammation

H₂S has been shown to contribute to inflammation via cytokines, chemokines, adhesion molecules (and leukocyte recruitment), and transient receptor potential vanilloid type 1 (TRPV1) and substance P.

5.1 Cytokines and Chemokines

In isolated pancreatic acinar cells stimulated by cerulein (an *in vitro* system that resembles acinar changes in acute pancreatitis), inhibition of H₂S formation by PAG has been shown to decrease mRNA expression and production of the chemokines monocyte chemoattractant protein (MCP)-1, macrophage inflammatory protein (MIP)-1 α , and MIP-2 (Tamizhselvi et al. 2007, 2008). Cerulein-induced acute pancreatitis is associated with a significant increase in mRNA for MCP-1, MIP-1 α , and MIP-2 in both the pancreas and lungs, suggesting that they are important early mediators in both local as well as distant inflammatory response (Tamizhselvi et al. 2008). Blockade of H₂S biosynthesis with PAG ameliorates the development of inflammatory process in cerulein-induced acute pancreatitis, acting through downregulation of chemokine expression (Tamizhselvi et al. 2008). Also, results indicate a key role of the phosphatidylinositol 3-kinase-protein kinase B pathway in relation to the action of H₂S on cerulein-induced cytokine production in isolated mouse pancreatic acinar cells (Tamizhselvi et al. 2009).

Both prophylactic and therapeutic administration of PAG significantly reduces the mRNA and protein levels of interleukin (IL)-1 β , IL-6, tumor necrosis factor (TNF)- α , MCP-1, and MIP-2 in lungs and liver, coupled with decreased nuclear translocation and activation of NF- κ B in lungs and liver following sepsis (Zhang et al. 2007a). In contrast, injection of NaHS significantly aggravates sepsis-associated systemic inflammation and increases nuclear factor (NF)- κ B activation. In addition, H₂S-induced lung inflammation is blocked by the NF- κ B inhibitor BAY 11-7082. Therefore, H₂S upregulates the production of pro-inflammatory mediators and exacerbates the systemic inflammation in sepsis through a mechanism involving NF- κ B activation (Zhang et al. 2007a).

In human monocyte cell line U937, treatment with H₂S donor NaHS results in significant increase in the mRNA expression and protein production of TNF- α , IL-1 β , and IL-6. This effect is mediated via NF- κ B and extracellular signal-related kinase (ERK) pathway (Zhi et al. 2007). Recent research, using mouse macrophage RAW264.7 cells, has shown that the activation of macrophages by lipopolysaccharide (LPS) results in higher levels of CSE mRNA and protein as well as the increased production of pro-inflammatory cytokines and chemokines IL-1 β ,

IL-6, TNF- α , and MCP-1. Silencing of the CSE gene by small interference ribonucleic acid (siRNA) results in decreased levels of pro-inflammatory cytokines (Badiei et al. 2013).

H₂S regulates inflammatory response by activating the extracellular signal-related kinase ERK pathway in polymicrobial sepsis (Zhang et al. 2008). Maximum phosphorylation of ERK1/2 and degradation of I κ B α in lung and liver is observed 4 h after CLP. Inhibition of H₂S formation by PAG significantly reduces the phosphorylation of ERK1/2 in lung and liver, coupled with decreased degradation of I κ B α and activation of NF- κ B. In contrast, injection of NaHS significantly enhances the activation of ERK1/2 in lung and liver, therefore leading to a further rise in tissue NF- κ B activity. In sepsis, pretreatment with PAG significantly reduces the production of cytokines and chemokines via NF- κ B and ERK1/2, whereas exogenous H₂S greatly increased it. In addition, pretreatment with PD98059, an inhibitor of MEK-1, significantly prevented NaHS from aggravating systemic inflammation in sepsis (Zhang et al. 2008). Binding site for NF- κ B has been shown on the CSE gene promoter and is critical for LPS-induced CSE expression (Wang et al. 2014). Therefore, H₂S may regulate systemic inflammatory response in sepsis via the NF- κ B-ERK pathway.

5.2 Adhesion Molecules and Leukocyte Recruitment

H₂S induces intercellular adhesion molecule (ICAM)-1 expression and neutrophil adhesion to cerulein-treated pancreatic acinar cells through nuclear factor (NF)- κ B and Src-family kinases (SFK) pathway (Tamizhselvi et al. 2010). H₂S activates SFKs in acinar cells, and inhibition of SFKs impairs H₂S-induced ICAM-1 expression secondary to the inhibition of NF- κ B activation. The effect of SFK inhibition on NF- κ B activation occurs together with I κ B α degradation. The results further demonstrate that neutrophil attachment onto H₂S-treated acinar cells is increased and that inhibition of SFK function inhibits H₂S-induced neutrophil attachment onto acinar cells. Taken together, these data indicate that H₂S engages SFKs in order to signal ICAM-1 expression by a mechanism involving induction of NF- κ B (Tamizhselvi et al. 2010).

Using intravital microscopy, it has been shown that in sepsis, prophylactic and therapeutic administration of PAG reduces leukocyte rolling and adherence significantly in mesenteric venules coupled with decreased mRNA and protein levels of adhesion molecules (ICAM-1, P-selectin, and E-selectin) in lung and liver. In contrast, administration of NaHS upregulates leukocyte rolling and attachment significantly, as well as tissue levels of adhesion molecules in sepsis. Conversely, in normal mice given NaHS to induce lung inflammation, NaHS treatment enhanced the level of adhesion molecules and neutrophil infiltration in lung. These alterations can be reversed by pretreatment with BAY 11-7082 (Zhang et al. 2007b). Therefore, H₂S acts as an important endogenous regulator of leukocyte activation and trafficking during an inflammatory response.

5.3 Transient Receptor Potential Vanilloid Type 1 and Substance P

Substance P is a mediator of inflammation and plays an important role in the pathogenesis of several inflammatory conditions. Intraperitoneal administration of the H₂S donor NaHS to mice causes an increase in circulating levels of substance P (Bhatia et al. 2006). NaHS, by itself, causes lung inflammation, as evidenced by a significant increase in lung MPO activity and histological evidence of lung injury. This is associated with a significant increase in lung levels of tumor necrosis factor (TNF)- α and interleukin (IL)-1 β . In preprotachykinin (PPT)-A^{-/-} mice, genetically deficient in substance P, H₂S does not cause any lung inflammation. Also, pretreatment of mice with CP-96345, an antagonist of the neurokinin-1 receptor (NK-1R), the receptor for substance P, protects mice against lung inflammation caused by H₂S. Depleting neuropeptide from sensory neurons by capsaicin significantly reduces the lung inflammation caused by H₂S. In addition, pretreatment of mice with capsazepine, an antagonist of the transient receptor potential vanilloid-1 (TRPV-1), protects mice against H₂S-induced lung inflammation. These results demonstrate an important role of substance P and neurogenic inflammation in H₂S-induced lung injury in mice (Bhatia et al. 2006).

In acute pancreatitis, PAG treatment significantly attenuates the increases in substance P concentrations in plasma, pancreas, and lung (Bhatia et al. 2008b). Moreover, administration of PAG significantly reduces PPT-A mRNA expression and NK-1R mRNA expression in both pancreas and lung when compared with cerulein-induced acute pancreatitis. This reduction in PPT-A mRNA expression and NK-1R mRNA expression is associated with a protection against acute pancreatitis and associated lung injury (Bhatia et al. 2008b). These results suggested that the pro-inflammatory effects of H₂S may be mediated by SP-NK-1R pathway in acute pancreatitis (Bhatia et al. 2008b). Furthermore, PPT-A deficiency and blockage of H₂S synthesis can regulate the toll-like receptor 4 (TLR4) pathway and subsequent innate immune response in acute pancreatitis, implying an interaction between SP/H₂S occurs via TLR4 and NF- κ B pathway. PPT-A gene deletion regulates H₂S-induced TLR4 signaling pathway in cerulein-treated pancreatic acinar cells, suggesting that in acute pancreatitis, H₂S may upregulate the TLR4 pathway and NF- κ B via substance P (Tamizhselvi et al. 2011).

In sepsis, PAG treatment significantly decreases the PPT-A gene expression and the production of substance P in lung, whereas administration of NaHS results in a further rise in the pulmonary level of substance P (Zhang et al. 2007c). PPT-A gene deletion and pretreatment with the NK-1R antagonist L703606 prevent H₂S from aggravating lung inflammation. In addition, septic mice genetically deficient in PPT-A gene or pretreated with L703606 do not exhibit further increase in lung permeability after injection of NaHS (Zhang et al. 2007c). These findings show that in sepsis, H₂S upregulates the generation of substance P that contributes to lung inflammation and lung injury mainly via activation of the NK-1R. Also, H₂S induces systemic inflammation and multiple organ damage characteristic of sepsis via transient receptor potential vanilloid type 1 (TRPV1)-mediated neurogenic

inflammation (Ang et al. 2010). Pretreatment with capsazepine, a TRPV1 antagonist, significantly attenuates systemic inflammation and multiple organ damage caused by sepsis. Moreover, capsazepine delays the onset of lethality and protects against sepsis-associated mortality. As expected, administration of NaHS exacerbates sepsis, but capsazepine reverses these deleterious effects. In the presence of PAG, capsazepine causes no significant changes to the PAG-mediated attenuation of systemic inflammation, multiple organ damage, and mortality. More importantly, capsazepine has no effect on endogenous generation of H₂S, suggesting that H₂S is located upstream of TRPV1 activation, and may play a critical role in regulating the production and release of sensory neuropeptides in sepsis. This study showed for the first time that H₂S induces systemic inflammation and multiple organ damage in sepsis via TRPV1-mediated neurogenic inflammation (Ang et al. 2010). Capsazepine treatment results in an attenuation of circulating and pulmonary levels of substance P in septic mice. Capsazepine also inhibits NaHS-augmented substance P production but has no effect on PAG-mediated abrogation of substance P levels in both plasma and lung. Capsazepine significantly reduces H₂S-induced inflammatory cytokines, chemokines, and adhesion molecules expression and protects against lung and liver dysfunction in sepsis. In the absence of H₂S, capsazepine caused no significant changes to the PAG-mediated attenuation of sepsis-associated systemic inflammatory response and multiple organ dysfunction. Capsazepine inhibits phosphorylation of ERK1/2 and IκBα, concurrent with suppression of NF-κB activation. Results in this study showed that H₂S regulates TRPV1-mediated neurogenic inflammation in polymicrobial sepsis through enhancement of SP production and activation of the ERK-NF-κB pathway (Ang et al. 2011a). Also, a recent study has shown that H₂S upregulates cyclooxygenase-2 (COX-2) and prostaglandin E metabolite (PGEM) in sepsis-evoked acute lung injury via TRPV-1 channel activation (Ang et al. 2011b).

6 H₂S Donors

NaHS and sodium sulfide (Na₂S) are the two “classical” H₂S donors that release the gas rapidly in solution. In most of the studies, they have been substantiated a pro-inflammatory action of H₂S. In recent years, slow/controlled release donors have emerged, and results using these compounds have been quite interesting. For example, S-diclofenac (ACS 15) is H₂S-releasing diclofenac, which comprises an H₂S-releasing dithiol-thione moiety attached by an ester linkage to diclofenac (a nonsteroidal anti-inflammatory drug—NSAID). The effect of treatment with the NSAID diclofenac and its H₂S-releasing derivative ACS15 on cerulein-induced acute pancreatitis and the associated lung injury has been investigated (Bhatia et al. 2008a). Although these two drugs do not have any significant effect on the local pancreatic injury, ACS15 affords significant protection against acute pancreatitis-associated lung injury (Bhatia et al. 2008a). ACS 15 also more effectively inhibits hindpaw swelling and neutrophil infiltration after carrageenan injection as compared to its parent NSAID (Sidhapuriwala et al. 2007). Furthermore, in a

rat model of LPS-induced endotoxemia, ACS 15 exhibits enhanced anti-inflammatory effect as compared to the parent drug (Li et al. 2007). Although these results suggest the potential for the use of controlled release H₂S donor compounds against inflammation, the protective action of ACS 15 in LPS-induced endotoxemia was associated with an inhibition of endogenous H₂S synthesis (Li et al. 2007). Therefore, protective actions of H₂S-releasing compounds may be caused by an inhibition of endogenous H₂S formation, possibly by a negative feedback mechanism caused by very low local levels of H₂S. Another slow H₂S-releasing drug is S-propargyl-cysteine (SPRC), a structural analog of S-allyl cysteine (SAC) with a common cysteine-containing structure. Pretreatment with SPRC has been shown to protect mice against acute pancreatitis and associated lung injury (Sidhapuriwala et al. 2012). Another slow H₂S-releasing compound is GYY4137 (morpholin-4-ium-4-methoxyphenyl(morpholino) phosphinodithioate), which has been reported to have an anti-inflammatory action (Li et al. 2009, 2013).

7 Conclusion

Figure 1 summarizes our current understanding of the mechanism of action of H₂S in inflammation. Several gaps in knowledge still remain on this subject. H₂S synthesis inhibitors, such as PAG, have proved useful in the proof of principle

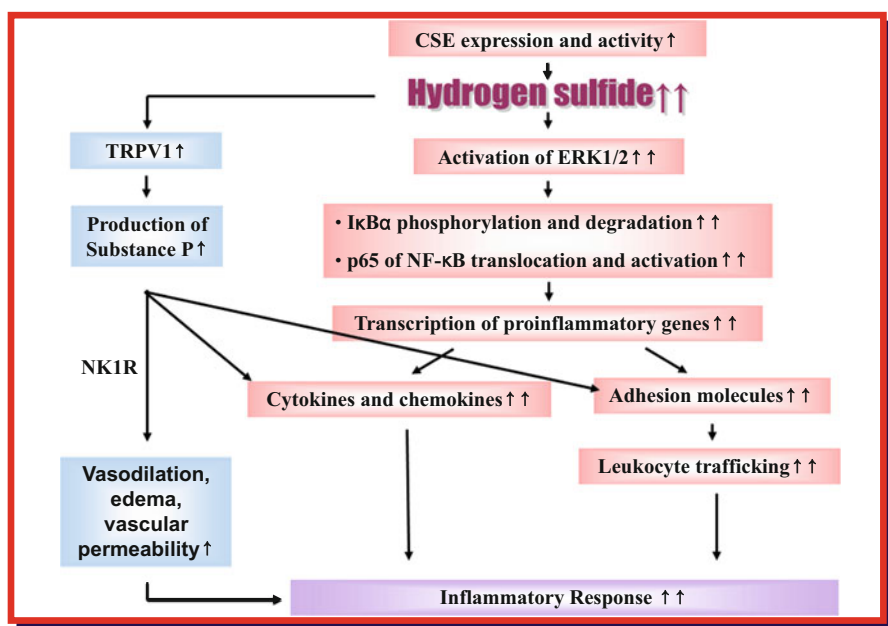


Fig. 1 A summary of our current understanding of the mechanisms of action of H₂S in inflammation

studies, but they are not entirely specific. Studies using CSE knockout mice and siRNA have contributed greatly to research in this area, but there is need to develop novel synthesis inhibitors that are more selective and have a better safety profile than the ones currently available. Also, following up on the progress made in basic/preclinical research, now is the time to investigate the role of H₂S in inflammation in clinical disease. As described in this article, some groups have already started working on this, and early results have been promising.

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Hydrogen Sulfide and Neuroinflammation

Kotaro Kida and Fumito Ichinose

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Abstract

The innate and adaptive immune system plays an important role in diverse forms of central nervous system (CNS) pathologies including neurodegenerative diseases and peripheral nerve injury. Evidence for an innate inflammatory response in Alzheimer's disease (AD) was described 20 years ago, and subsequent studies have documented roles of inflammation in Parkinson's disease (PD), amyotrophic lateral sclerosis (ALS), multiple sclerosis (MS), and a growing number of other CNS pathologies. Although inflammation may not be the initiating factor for neurodegenerative pathologies, experimental data suggests that persistent inflammatory responses involving microglia and astrocytes, as well as blood monocyte-derived macrophages, clearly contribute to disease progression.

High levels of hydrogen sulfide exert toxic effects to CNS. On the other hand, low and physiological levels of H₂S may have beneficial effects on number of tissues including CNS. For example, a number of studies have reported that H₂S exerts anti-inflammatory and anti-apoptotic effects in CNS. In this chapter, studies related to the role of H₂S in neuroinflammation and neurodegeneration

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will be reviewed and discussed. In particular, we will focus on the role of H₂S in neuroinflammation associated with PD.

Keywords

Neurodegenerative disease • Parkinson's disease • Microglia • Gliosis • Inflammatory cytokine

Parkinson's disease (PD) is one of the most common neurodegenerative diseases (Twelves et al. 2003). It is characterized by a slow and progressive degeneration of dopaminergic neurons in the substantia nigra (Hirsch et al. 1988). Although the etiology of PD is not fully understood, several mechanisms responsible for the neurodegeneration in PD have been suggested, including abnormal protein handling, oxidative stress, mitochondrial dysfunction, excitotoxicity, neuroinflammation, and apoptosis (Hirsch and Hunot 2009).

A number of animal models of PD, both toxin-induced and genetically engineered, have been created. While the toxin-induced PD models have provided considerable therapeutic insight into basal ganglia physiology and response to drug therapy, genetic models provide a powerful new set of molecular tools to study the etiology of PD (Melrose et al. 2006). Although none of the animal models accurately recapitulate the pathophysiological features of PD, 1-methyl-4-phenyl-1,2,3,6-tetrahydropyridine (MPTP) currently represents the most important and most frequently used parkinsonian toxin applied in animal models (Beal 2001; Przedborski et al. 2001; Schober 2004). MPTP is the only known dopaminergic neurotoxin capable of causing a clinical picture indistinguishable from idiopathic PD in humans (Langston et al. 1983). Moreover, MPTP produces a reliable and reproducible lesion of the nigrostriatal dopaminergic pathway after its systemic administration, which is often not the case for other neurotoxins (Bove et al. 2006).

Hydrogen sulfide appears to confer cytoprotection via multiple mechanisms including antioxidant and anti-apoptotic effects. For example, Kimura and colleagues demonstrated using a model of glutamate-induced oxidative stress that H₂S protects neurons from cell death by increasing the levels of the antioxidant, glutathione (Kimura and Kimura 2004). A recent study revealed that NaHS, an H₂S donor, protects PC12 cells from cytotoxicity and apoptosis induced by 1-methyl-4-phenylpyridinium ion (MPP⁺), the active metabolite of MPTP (Yin et al. 2009).

1 Effects of H₂S in Animal Models of PD

Neuroprotective effects of H₂S donors (i.e., NaHS and Na₂S) have been reported *in vitro* and *in vivo* in a variety of animal models (Kimura et al. 2010; Kimura and Kimura 2004). In particular, a recent report by Hu and colleagues demonstrated neuroprotective effects of NaHS in rat models of PD induced by 6-OHDA or rotenone (Hu et al. 2009). While these studies suggest neuroprotective effects of H₂S or related compounds, solution of NaHS has been found to include polysulfides

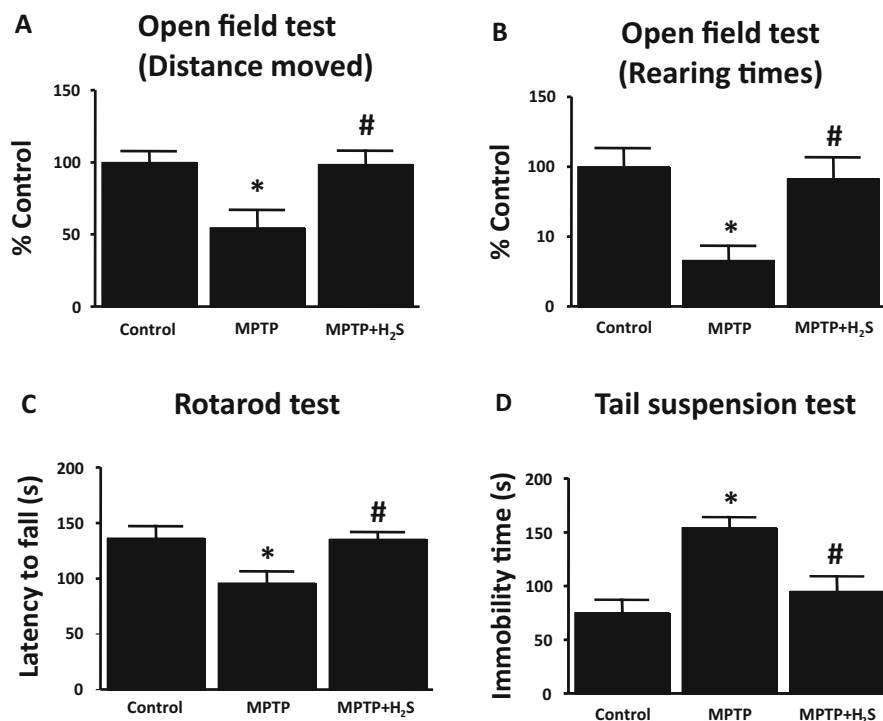


Fig. 1 Results of behavioral tests performed at 7 days after administration of MPTP or saline. Results of distance moved (a) and rearing times (b) in the open field test are shown as percent of the values in saline-treated control mice. (c) Latency to fall in the rotarod test is shown in seconds. (d) Immobility times in the tail-suspension test are shown in seconds. Control saline-treated mice, MPTP mice treated with MPTP that breathed air, MPTP + H₂S mice treated with MPTP that breathed H₂S. *N* = 7–9 in each group in all tests. **P* < 0.05 vs. control. #*P* < 0.05 vs. MPTP. Reproduced from Kida et al. *Antioxid Redox Signal* 2011 Jul 15;15(2):343-52 with permission

and elemental sulfur in addition to H₂S (Doeller et al. 2005). To examine the neuroprotective effects of authentic H₂S, we sought to examine effects of H₂S inhalation in the clinically relevant MPTP-induced PD model in mice (Kida et al. 2010). Inhalation of H₂S at 40 ppm for 8 h a day for 7 days prevented MPTP-induced movement disorder in mouse as measured by open field test, rotarod test, and tail-suspension test (Fig. 1). The neuroprotective effects of H₂S inhalation were associated with marked attenuation of the MPTP-induced loss of TH-containing neurons in substantia nigra and striatum (Fig. 2).

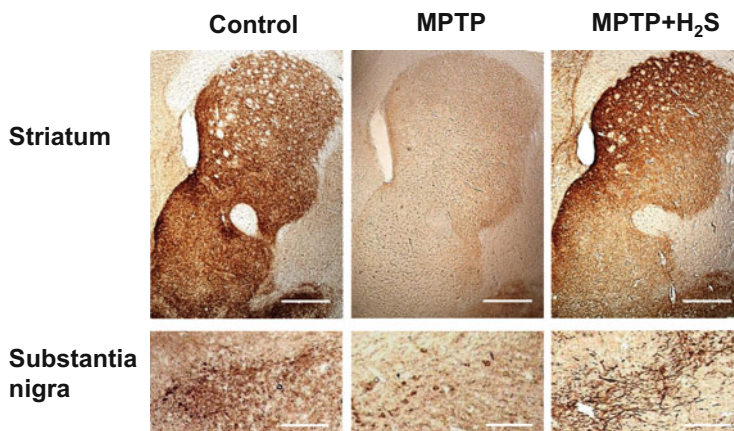


Fig. 2 Representative immunohistochemical staining of TH-positive neurons in mice 7 days after administration of MPTP or saline in striatum and substantia nigra. Reproduced from Kida et al. *Antioxid Redox Signal* 2011 Jul 15;15(2):343-52 with permission

2 Anti-apoptotic Effects of Inhaled H₂S

To determine the mechanisms responsible for the protective effects of H₂S inhalation after MPTP, we conducted extensive histological analysis of brain sections. Administration of MPTP increased the number of apoptotic cells in substantia nigra 1 day after MPTP administration as indicated by the presence of TUNEL-positive nuclei (Fig. 3a–d). In contrast, H₂S inhalation prevented the MPTP-induced apoptosis (Fig. 3e–h). Immunofluorescence double labeling demonstrated that the TUNEL-positive nuclei were found exclusively in substantia nigra especially in the area where TH-positive neurons reside (Fig. 3d, h, i). While majority of the TUNEL-positive nuclei did not overlay with TH-positive neurons (Fig. 3d, h, i, j), triple labeling with TUNEL, TH, and DAPI showed that a small number of TH-positive neurons contained a TUNEL-positive nucleus (Fig. 3j, k). These observations suggest that at least a part of TH-positive neurons died via apoptosis.

To further identify the cell types undergoing apoptosis in substantia nigra after MPTP administration, we stained brain sections with antibodies against GFAP (a marker of activated astrocytes) or Iba-1 (a marker of activated microglia) after TUNEL assay. Double labeling with TUNEL and GFAP revealed that GFAP-positive activated astrocytes did not overlay with any TUNEL-positive nuclei (Fig. 4a). On the other hand, double labeling with TUNEL and Iba-1 revealed that many TUNEL-positive nuclei were enveloped by Iba-1-positive activated microglia (Fig. 4b). In fact, triple labeling with TUNEL, Iba-1, and DAPI showed that majority of TUNEL-positive nuclei were distinct from nuclei of microglia (Fig. 4c), suggesting phagocytosis of apoptotic neurons by activated microglia. Taken together, these observations suggest that MPTP-induced apoptosis of

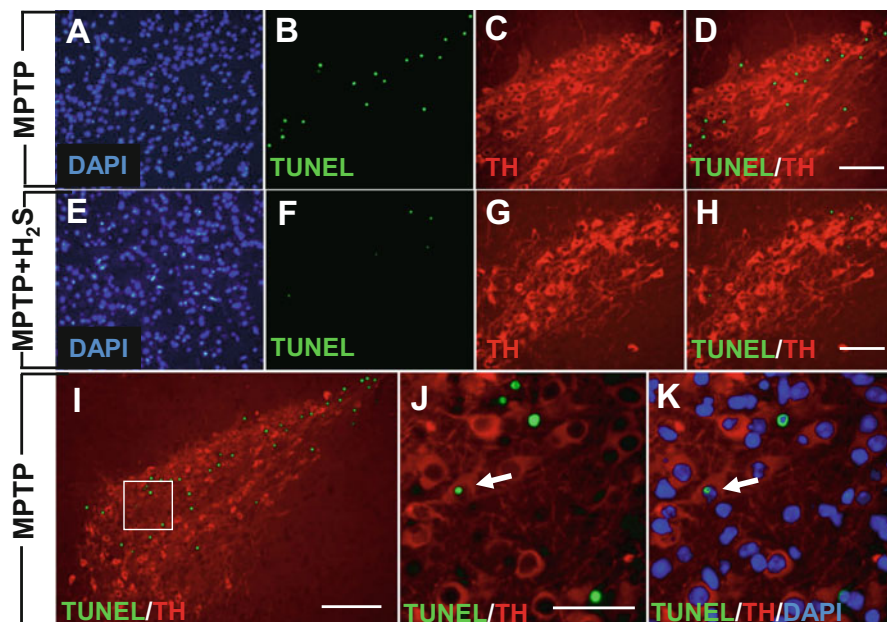


Fig. 3 Representative photomicrographs of brain sections containing substantia nigra obtained from mice 1 day after administration of MPTP without H₂S (MPTP, **a–d, i–k**) or MPTP with H₂S breathing (MPTP+H₂S, **e–h**). Sections were subjected to the TUNEL assay and stained with DAPI, anti-TH, anti-GFAP, or anti-Iba-1 antibody. Size bar = 100 μ m for (**a–h**), 200 μ m for (**i**), and 5 μ m for (**j**) and (**k**). *White arrows* in (**j**) and (**k**) indicate a TH-positive neuron containing TUNEL-positive nuclei. Reproduced from Kida et al. *Antioxid Redox Signal* 2011 Jul 15;15(2):343-52 with permission

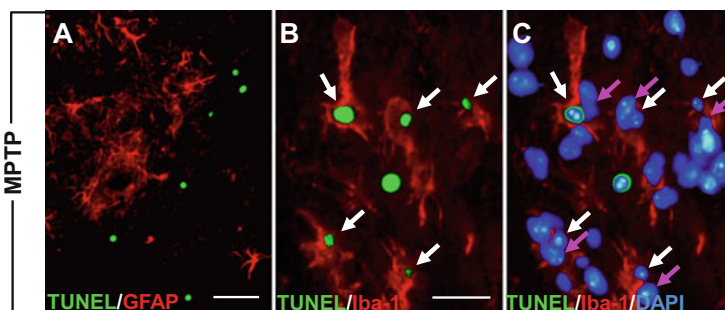


Fig. 4 Representative photomicrographs of brain sections 1 day after MPTP administration without H₂S breathing. (**a**) Double labeling with TUNEL (*green*) and GFAP (*red*). Size bar = 5 μ m. (**b**) Double labeling with TUNEL (*green*) and Iba-1 (*red*). (**c**) Triple labeling with TUNEL (*green*), Iba-1 (*red*), and DAPI (*blue*). Size bar = 2.5 μ m for (**b**) and (**c**). *White arrows* indicate TUNEL-positive nuclei and *pink arrows* indicate nuclei of microglia distinct from the TUNEL-positive nuclei. Reproduced from Kida et al. *Antioxid Redox Signal* 2011 Jul 15;15(2):343-52 with permission

TH-positive neurons in substantia nigra and apoptotic neurons were phagocytosed by activated microglia.

3 Effects of H₂S on Glial Activation

Immunofluorescence staining demonstrated few activated microglia and astrocytes in substantia nigra and striatum of saline-treated mice (Fig. 5a, d, g, j). In contrast, MPTP increased the number of activated microglia and astrocytes in substantia

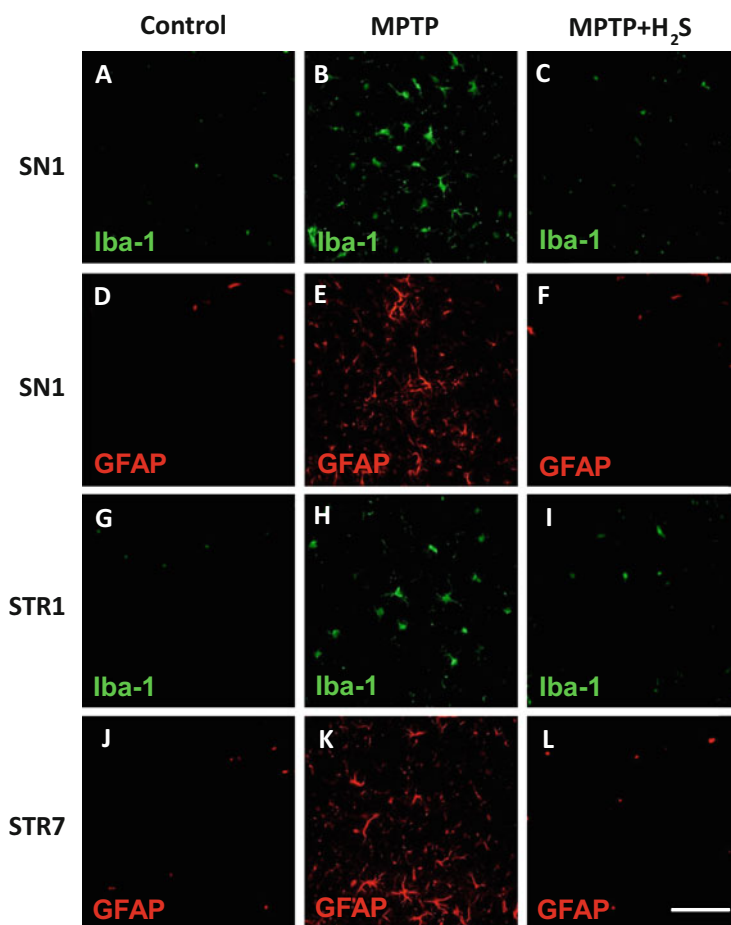


Fig. 5 Representative immunofluorescence staining for Iba-1 (microglia, *green*) and GFAP (astrocytes, *red*) in substantia nigra (SN) and striatum (STR) of saline-treated control mice (Control, **a, d, g, and j**), mice treated with MPTP that breathed air (MPTP, **b, e, h, and k**), and mice treated with MPTP that breathed H₂S (MPTP + H₂S, **c, f, i, and l**). These sections were obtained from mice 1 day (SN1 and STR1) or 7 days (STR7) after administration of MPTP or saline. Reproduced from Kida et al. *Antioxid Redox Signal* 2011 Jul 15;15(2):343-52 with permission

nigra on Day 1 (Fig. 5b, e). In striatum, the number of activated microglia increased on Day 1 (Fig. 5h) and the number of activated astrocytes increased on Day 7 after MPTP injection (Fig. 5k). Inhaled H₂S prevented the activation of both microglia and astrocytes in substantia nigra and striatum after MPTP administration (Fig. 5c, f, i, l).

4 Role of Endogenous H₂S in Parkinson's Disease

Parkin is an E3 ubiquitin ligase that ubiquitinates diverse substrates (Martin et al. 2011). Mutations in parkin that disrupt its catalytic activity are the most common cause of autosomal recessive form of PD, implying that loss of parkin is neurotoxic, while its enhancement is neuroprotective. Parkin may also participate in the pathophysiology of the common sporadic form of PD based on interactions with nitric oxide (NO). It has been reported that parkin is *S*-nitrosylated in brains of patients with PD (Chung et al. 2004; Yao et al. 2004). Nitrosylation of parkin inhibits its E3 ligase and neuroprotective activities, implying that the increased nitrosylation of parkin in PD is pathogenic. H₂S signals primarily by attaching to SH groups of cysteines in proteins, a process termed sulfhydrylation (Mustafa et al. 2009). Sulfhydrylation generally occurs on the same cysteines as nitrosylation so that the two processes may be reciprocal.

Vandiver and colleagues recently reported that parkin is physiologically sulfhydrylated and that, whereas nitrosylation inactivates parkin, sulfhydrylation stimulates its activity (Vandiver et al. 2013). These authors identified major declines of parkin sulfhydrylation in the corpus striatum of PD patients. Their experiments also showed cytoprotective actions of H₂S donors, which appear to reflect sulfhydrylation of parkin. Mutation of C95, a principal site of parkin sulfhydrylation, largely prevented the protective influences of H₂S donors indicating that the donors act via parkin sulfhydrylation to enhance its ubiquitination activity. In addition, H₂S donors were cytoprotective in parkin-related models of neurotoxicity. These studies suggest that diminished sulfhydrylation of parkin may be pathogenic in PD and selective H₂S donors may be therapeutic.

5 Conclusion

In the last several years, a number of studies reported that H₂S donors exert beneficial effects in rodent models of PD (Hu et al. 2010; Kida et al. 2010; Lu et al. 2012), and an H₂S donating variant of L-DOPA has shown promising effects in cellular models of PD (Lee et al. 2010). In addition, H₂S donors have been shown to exert potent anti-inflammatory effects in other forms of inflammatory pathologies (Li et al. 2009; Shirozu et al. 2013; Tokuda et al. 2012; Whiteman et al. 2009). While these beneficial effects had been speculated to reflect general antioxidant and anti-inflammatory actions of H₂S, numerous studies have failed to reveal a beneficial effect of antioxidants in PD. For this reason, recent findings by

Vandiver and colleagues are particularly notable because they may provide a molecular mechanism whereby H₂S therapy may prevent one of the initiating pathogenetic mechanisms of PD, as well as to mitigate propagation of the pathological process by chronic inflammation. These recent observations suggest a unique potential of H₂S-based approach to the treatment of PD and other CNS pathologies characterized with progressive neurodegeneration.

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Part V

H₂S and the Nervous System

Brain, Learning, and Memory: Role of H₂S in Neurodegenerative Diseases

B.V. Nagpure and Jin-Song Bian

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Abstract

For more than 300 years, the toxicity of hydrogen sulfide (H₂S) has been known to mankind. However, this point of view is changing as an increased interest was

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observed in H₂S biology in the last two decades. The scientific community has succeeded to unravel many important physiological and pathological effects of H₂S on mammalian body systems. Thus, H₂S is now referred to as a third endogenous gaseous mediator along with nitric oxide and carbon monoxide. Acting as a neuromodulator, H₂S facilitates long-term potentiation and regulates intracellular calcium levels, which are important processes in learning and memory. Aberrant endogenous production and metabolism of H₂S are implicated in pathogenesis of neurodegenerative diseases including Alzheimer's disease (AD) and Parkinson's disease (PD). Various H₂S donors have shown beneficial therapeutic effects in neurodegenerative disease models by targeting hallmark pathological events (e.g., amyloid- β production in AD and neuroinflammation in PD). The results obtained from many in vivo studies clearly show that H₂S not only prevents neuronal and synaptic deterioration but also improves deficits in memory, cognition, and learning. The anti-inflammatory, antioxidant, and anti-apoptotic effects of H₂S underlie its neuroprotective properties. In this chapter, we will overview the current understanding of H₂S in context of neurodegenerative diseases, with special emphasis on its corrective effects on impaired learning, memory, and cognition.

Keywords

Hydrogen sulfide • Neurodegeneration • Brain • Memory • Learning • Neuroinflammation

Abbreviation

3-MST	3-Mercaptopyruvate sulfurtransferase
A β	Amyloid- β
AD	Alzheimer's disease
APP	Amyloid precursor protein
CAT	Cysteine aminotransferase
CBS	Cystathionine- β -synthase
CNS	Central nervous system
CSE	Cystathionine- γ -lyase
H ₂ S	Hydrogen sulfide
K _{ATP}	ATP-sensitive potassium channel
LTP	Long-term potentiation
mitoK _{ATP}	Mitochondrial K _{ATP} channel
NaHS	Sodium hydrogen sulfide
NF- κ B	Nuclear factor kappa-light-chain-enhancer of activated B cells
NMDA	<i>N</i> -methyl-D-aspartic acid
NSAIDs	Non-steroidal anti-inflammatory drugs

PD	Parkinson's disease
ROS	Reactive oxygen species

1 Introduction

Alzheimer's disease (AD), Parkinson's disease (PD), and other neurodegenerative diseases share a number of common aspects—old-age onset, hereditary or sporadic appearance, and the aggregation and deposition of misfolded proteins. These diseases show overlapping spectrum of clinical signs and symptoms in their advance stages. In 2005, 24.2 million people worldwide had dementia, which is one of the most common sign of AD. The developed world i.e., North America and Western Europe showed the highest prevalence of dementia. It is estimated that these parts of the world along with China will be home to around 55 % of total affected population worldwide by 2040 (Reitz et al. 2011). The prevalence of dementia grows with age. The extensive Delphi consensus study done by Ferri et al. revealed the growth from 1 % in 60–64 years age group to about 30 % in those of >85 years age group (Ferri et al. 2005). The incidence rate of AD and other dementias also increase exponentially with the age, mirroring the prevalence rate. The worst affected is the seventh and eighth decade of life (Reitz et al. 2011). PD is the second most common neurodegenerative disorder, with incidence rate rising sharply after the age of 60. The prevalence of PD in people over age of 50 is feared to be doubled to approximately nine million by 2030 in the world's most populated and industrialized countries. Placing a great burden on caregivers, AD, PD, and other neurodegenerative diseases are costing dearly to today's society

In recent decades, the scientific community has witnessed the rise of a whole new class of gaseous biological mediators in mammalian cells. These are simple gas molecules, which are lipid soluble and hence freely membrane permeable reaching intracellular organelle. After the seminal discovery of physiological effects of nitric oxide (NO) on blood vessels, carbon monoxide (CO) and hydrogen sulfide (H₂S) have been recently recognized as two more “gasotransmitters.” The physiological role of H₂S was discovered by a Japanese group of scientists led by Kimura in 1996. In their pioneering study, the novel neuromodulator role of H₂S was transpired (Abe and Kimura 1996). Since then its possible roles in other body systems have been investigated worldwide. In mammalian central nervous system (CNS), its prominent effects include modulation of neurotransmission and long-term potentiation (LTP) (Abe and Kimura 1996) and induction of neuroprotection (Hu et al. 2011) from myriad of pathogenic agents. In mammalian cardiovascular system (CVS), its protective effects are deeply studied (Pan et al. 2006; Hu et al. 2008; Liu et al. 2012; Polhemus et al. 2014). The induction of relaxation (Lee et al. 2007; Yang et al. 2008) and constriction (Lim et al. 2008; Kohn et al. 2012) in various types of blood vessels is also documented (d'Emmanuele di Villa Bianca et al. 2011). The opposite effects of H₂S on systemic and localized inflammation have been observed in various mammalian tissues (Hegde and Bhatia

2011; Whiteman and Winyard 2011). In this book chapter, we will discuss the well-accepted and latest findings pertaining to neurophysiological and neuropharmacological effects of H₂S.

2 H₂S

2.1 Physical and Chemical Properties

At room temperature and ambient pressure, H₂S exists in a colorless gaseous form. The smell is pungent with distinctive rotten-egg odor. It is readily water soluble due to its weak acidic nature. Its solubility was measured to be 80 mM at 37 °C as equilibrium between H₂S, HS⁻, and S²⁻. The acid dissociation constant (pK_a) values of the first and second dissociation steps are recorded as 7.0 and >12.0, respectively (Vorobets et al. 2002; Kabil and Banerjee 2010; Mark et al. 2011). Thus, at physiological pH of 7.4 H₂S exists majorly as HS⁻ moiety along with minor presence of free H₂S in its dissociated form. The minute amounts of sulfide anions (S²⁻) can also be detected. Even with the advent of various methods of H₂S measurement, it is almost impossible to determine the active form of H₂S (H₂S, HS⁻, or S²⁻) present in the biological system. Hence the all-encompassing term of H₂S is now used to refer the total sulfide content present in the solution (i.e., H₂S + HS⁻ + S²⁻) (Zhao et al. 2014).

2.2 Biosynthesis in Mammalian CNS

In mammalian tissues, H₂S is biosynthesized from amino acid cysteine (Cys) and homocysteine (Hcy), which are recognized as the principle substrates for its endogenous production. They are acted upon by three different enzymes, namely cystathionine β-synthase (CBS), cystathionine γ-lyase (CSE), and 3-mercaptopyruvate sulfur transferase (3-MST) (Hu et al. 2011). Expressions of these enzymes are variable in different tissues. The study of this variation is important as the modulation of endogenous production of H₂S can be achieved by targeting each enzyme separately or concurrently.

CBS, a pyridoxal-5'-phosphate (PLP)-dependent enzyme, is abundantly expressed in brain and is the key source of H₂S there (Abe and Kimura 1996). The levels of CBS expression vary according to the stages of development. While the levels are low during embryonic brain, they increase notably from the late prenatal to the early postnatal period. After adulthood is reached, its level declines again (Enokido et al. 2005). Initially, the neuronal localization of CBS was demonstrated in major areas of brain including the hippocampus and the cerebellum (Robert et al. 2003). However, further study by Enokido et al. and Lee et al. revealed that H₂S is preferentially expressed in radial glia/astrocyte lineage of developing mouse brain (Enokido et al. 2005; Lee et al. 2009). In support of these findings, CBS has been found to be elevated in reactive astrocytes (Kimura and Kimura 2004). CBS initiates the *trans*-sulfuration pathway by catalyzing

β -replacement of serine by Hcy to generate cystathionine and water. Furthermore, serine replacement by Cys as a substrate results into production of cystathionine and H₂S. Besides above-mentioned *trans*-sulfuration pathway, CBS also catalyzes condensation reactions between two molecules of Cys and β -replacement of Cys by water to produce H₂S (Kabil and Banerjee 2014). The reaction replacing Hcy by Cys yields maximum generation (~96 %) of H₂S in vitro. Whereas the minor share of 1–3 % is contributed by condensation reactions between two molecules of Cys (Singh et al. 2009).

CSE is yet another PLP-dependent enzyme, which mediates a reaction between thiocysteine and a thiol compound R-SH to generate H₂S (Kimura 2011). The substrate thiocysteine is generated from L-cystine which in turn is produced by two L-cysteine molecules (Yamanishi and Tuboi 1981). Expression of CSE is rather widely distributed among peripheral tissues including liver, pancreas, uterus, and intestine (Kimura 2011). The activity of CSE in the human brain is 100 folds higher than mouse brain and it plays an important role in CNS physiology (Diwakar and Ravindranath 2007). CSE is a rate-limiting enzyme during the generation of Cys from methionine in *trans*-sulfuration pathway. The availability of Cys is an essential factor in glutathione (GSH) synthesis, which in turn is needed for mitochondrial function preservation to maintain redox homeostasis in cells. Thus CSE offers neuroprotection against oxidative stress by maintaining GSH and protein thiol homeostasis (Diwakar and Ravindranath 2007). Under physiologic conditions, CSE generates H₂S mainly from Cys. However under hyperhomocysteinemic (abnormally high level of homocysteine in the blood) conditions, Hcy (γ -elimination) becomes its major substrate taking over Cys. This alteration might be responsible for elevated production of H₂S in severe hyperhomocysteinemic conditions (Singh et al. 2009). Hence, it is probable that CSE becomes a major enzyme to produce H₂S in conditions like hyperhomocysteinemia (Hu et al. 2011), which incidentally is a common feature of neurodegenerative diseases including AD and PD.

The third enzyme, 3-mercaptopyruvate sulfotransferase (3-MST), was identified in the neurons. The research group detected the significant presence of H₂S in the brain homogenate preparation of CBS^{-/-} mice (Shibuya et al. 2009). Kimura further observed that 3-MST acts together with cysteine aminotransferase (CAT) to generate H₂S from Cys in the presence of α -ketoglutarate (Kimura et al. 2010). However, it is suggested that 3-MST is unable to produce H₂S in normal physiological conditions as they exert their activities at higher alkaline pH level (Shibuya et al. 2009). Furthermore, it requires endogenous reducing substances such as thioredoxin and dihydrolipoic acid (DHLA) for the production of H₂S (Kimura 2014). Aspartate can also act as a substrate for CAT, competitively binding to it and attenuating H₂S synthesis (Guo et al. 2012).

Recently, Shibuya et al. discovered the additional pathway for H₂S biosynthesis in mammalian cells. 3-MST along with D-amino acid oxidase (DAO) produces H₂S from D-cysteine by the interaction of mitochondria and peroxisomes. It was evident that this D-cysteine-dependent pathway operates predominantly in the cerebellum and the kidney. The protective effects of D-cysteine were observed against oxidative

stress in cerebellar neurons and against ischemia-reperfusion injury in the kidney (Shibuya and Kimura 2013).

2.3 Storage and Metabolism

Although endogenous H_2S can be synthesized and released immediately, the storage forms of H_2S are also known. Acid-labile sulfur is primarily contained in iron-sulfur center of mitochondrial enzymes and can release H_2S only in acidic pH of 5.4. Due to higher instability of iron-sulfur complexes, the release of H_2S is readily achieved. Bound sulfane sulfur, which is localized in cytoplasm, consists of divalent sulfur bond (e.g., persulfide form). It releases H_2S under reducing conditions of pH 8.4 (Ishigami et al. 2009). It is possible that H_2S produced by 3-MST/CAT enzymatic pathway is stored in the bound sulfane sulfur form. The decreased amount of bound sulfane sulfur has been detected in cells without 3-MST/CAT compared to the cells with 3-MST/CAT (Shibuya et al. 2009).

H_2S is catabolized in mammalian cells through various pathways. The major mechanism is through its mitochondrial oxidation in different tissues (Hildebrandt and Grieshaber 2008). In a reaction catalyzed by quinone oxidoreductase enzyme, H_2S is converted into persulfides. Persulfides are, in turn, oxidized in sulfite and thiosulfite. In physiological normoxic conditions, the thiosulfite is further metabolized into excretable form of sulfate. H_2S catabolism by quinone oxidoreductase enzyme seems to be universal in mammalian tissues, with possible exception of the brain (Mikami et al. 2011). H_2S can also be methylated to produce methane thiol by the action of enzyme thiol-S-methyltransferase. Non-mitochondrial heme proteins such as hemoglobin and myoglobin also catabolize intracellular H_2S by oxidation (Berzofsky et al. 1971; Stein and Bailey 2013). To a weaker extent, H_2S can also interact with reactive oxygen and nitrogen species. It is interesting to know that the presence of oxygen (O_2) is a very influential factor in deciding the fate of cellular H_2S as O_2 is capable of spontaneous oxidization of H_2S (van Kampen and Zijlstra 1983; Stein and Bailey 2013). The intracellular concentration of H_2S is kept firmly in low range, owing to the highly efficient nature of above-mentioned mechanisms.

3 Protective Roles of H_2S in CNS

The molecular basis of protective effects of H_2S in CNS during neurodegenerative diseases has received a considerable attention in recent years. This section summarizes the major protective effects of H_2S in CNS with brief discussion about underlying molecular mechanism (Fig. 1).

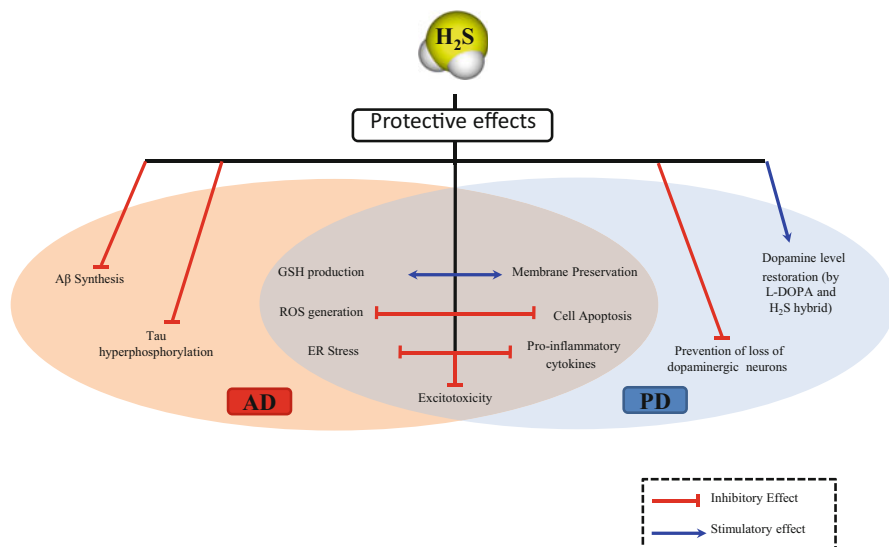


Fig. 1 The various targets of H₂S for its neuroprotective effects in the neurodegenerative diseases. The hallmark events in AD pathology i.e., A β generation and tau hyperphosphorylation are downregulated by H₂S. H₂S donors and hybrids produce positive therapeutic effects in PD pathology as well. Apart from reducing loss of neurons in substantia nigra of striatum, it also restores dopamine levels. H₂S elevates the intracellular levels of glutathione (GSH) while preserving membrane integrity, thus making neurons less susceptible to cellular injury. H₂S is also shown to prevent excessive cellular apoptosis, lower oxidative stress, reduce neuroinflammation by several pro-inflammatory cytokines, and prevent cells from excitotoxicity caused by aberrant neurotransmitter release from neurons and astrocytes

3.1 Anti-Inflammation

Neuroinflammation is strongly implicated in the pathologies of all major neurodegenerative diseases. The non-neuronal cells like microglia, astrocytes, and oligodendrocytes constitute the microenvironment under which normal neuronal functions are kept smoothly running. However, in case of any pathological or toxic insult, their over/chronic activation can initiate many unwanted and potentially neurotoxic cascades.

The pioneering work of Hu et al. revealed the anti-inflammatory properties of H₂S in lipopolysaccharide (LPS)-induced neuroinflammation in both primary cultured microglia and immortalized murine BV-2 microglial cells (Hu et al. 2007). In support of above findings, other studies also demonstrated the anti-inflammatory effects of H₂S-releasing NSAIDs in various neurodegenerative disease models. Although there is a lot to be done in order to completely reveal the underlying mechanisms of anti-inflammatory effect of H₂S, above-mentioned and other studies implicated inhibition of nuclear factor-kappa B (NF- κ B) and MAPK signaling cascades in the observed effects. Additionally, in an interesting study analyzing the neuroprotective effects of *S*-propargyl-cysteine (a novel H₂S-modulating agent),

its inhibitory effects on tumor necrosis factor (TNF)- α and TNF- α receptor 1 (TNFR1) were demonstrated (Gong et al. 2011). Earlier this year, Zhao et al. shed some light upon a previously unrecognized mechanism underlying H₂S suppression of neuroinflammation. They found that H₂S polarized microglia to an anti-inflammatory (M2) phenotype by activating calmodulin-dependent protein kinase kinase β (CaMKK β)-dependent AMP-activated protein kinase (AMPK) (Zhou et al. 2014). By suppressing neuroinflammation, H₂S imparts improvement in learning and spatial memory in various animal models of AD (Xuan et al. 2012; He et al. 2014).

3.2 Anti-Oxidation

The brain is highly prone to oxidative stress than other organs in mammalian body (Chance et al. 1979). Accounting for relatively small fraction of body weight (2 %), it is constantly supplied with large amount of total body oxygen consumption (20 %) and total body glucose (25 %) in order to maintain high metabolic turnover (Uttara et al. 2009). During the mitochondrial generation of ATP via oxidative pathway, huge amount of ROS and hydrogen peroxide (H₂O₂) are generated. Furthermore, the biochemical structure of neurons makes them particularly vulnerable to ROS. The double bonds present in abundant unsaturated fatty acids are easily susceptible to peroxidation (Butterfield et al. 2002). To worsen the conditions, brain is not equipped with high antioxidant activity. Additionally, the higher concentration of iron and vitamin C in various regions of brain leads to increased interaction between O₂ and metal ions, culminating in ROS generation (Floyd and Carney 1992).

Oxidative stress caused by overproduction of ROS is detrimental and one of the etiological factors of many neurodegenerative diseases. Kimura et al. found that H₂S protects primary neurons from oxidative glutamate toxicity (oxytosis) caused by glutamate. H₂S elevated antioxidant GSH levels by enhancing the activity of gamma-glutamylcysteine synthetase and upregulating cystine transport. The upregulation of gamma-glutamylcysteine synthetase activity facilitates the redistribution of GSH into mitochondria, thus protecting cells against oxidative stress damage (Kimura and Kimura 2004). Later, the same group also discovered that H₂S protects immortalized mouse hippocampal cells from oxytosis by activating ATP-dependent K⁺ (K_{ATP}) and Cl⁻ channels (Kimura et al. 2006). A study conducted by Lu et al. demonstrated that H₂S protects astrocytes via enhancing glutamate uptake function of glutamate transporter-1 and elevating GSH production. This phenomenon prevents excessive accumulation of glutamate in synaptic clefts protecting neurons from excitotoxicity (Lu et al. 2008). Besides these, H₂S has been show to downregulate peroxynitrite-mediated tyrosine nitration and inactivation of alpha1-antiproteinase inhibiting peroxynitrite-induced cytotoxicity, intracellular protein nitration, and protein oxidation in human neuroblastoma SH-SY5Y cells (Whiteman et al. 2004).

3.3 Anti-Apoptosis

Apart from antioxidant effects, H₂S is also known to possess anti-apoptotic properties conferring neuroprotection. Hu and colleagues discovered that H₂S inhibits apoptosis induced by rotenone (a toxin used to establish PD model) by preserving mitochondrial functions in human neuroblastoma cell line (SH-SY5Y). They observed that H₂S regulated the mitoK_{ATP} channel and thus impeded the apoptosis cascade (prevention of mitochondrial membrane potential (MMP) dissipation, cytochrome c release, and caspase-9/3 activation) (Hu et al. 2009). The anti-apoptotic effect was supported by other studies as well. Zhang et al. found out that H₂S attenuated neuronal injury induced by vascular dementia via inhibiting apoptosis in rats (Zhang et al. 2009). In yet another study, H₂S imparted the cytoprotective effect to PC12 cells against amyloid β (25–35)-induced apoptosis (Tang et al. 2008).

4 H₂S in Neurodegenerative Diseases

4.1 AD

AD is the most common neurodegenerative disease in today's society (Reitz et al. 2011). Classically, AD pathology is characterized by aggregation and deposition of A β plaques along with hyperphosphorylated tau in the brain (Goedert and Spillantini 2006).

More and more evidences are coming into light depicting a certain relevance of endogenous H₂S generation and AD pathology. In the very first study of its kind, Morrison et al. showed that concentration of S-adenosylmethionine, a CBS activator, was notably depleted in brains of AD patients (Morrison et al. 1996). It was later found out that any interference in *trans*-sulfuration pathway (please refer to Sect. 2.3 of this chapter) results in both elevation in total serum Hcy (Clarke et al. 1998) and depletion in H₂S production (Dwyer et al. 2004) in AD patients. It should also be noted that Hcy neurotoxicity itself inhibits CBS, reducing H₂S production (Tang et al. 2011). Furthermore, an extensive clinical study conducted by Liu et al. has proven the direct correlation between severity of AD and H₂S (Liu et al. 2008).

Recently, many studies have shown that H₂S affects amyloidogenesis in the brain. The downregulated generation of A β was observed in primary neuron culture (Zhu et al. 2014), SH-SY5Y neuroblastoma cell line (Nagpure and Bian 2014), and APP/PS1 transgenic mice (He et al. 2014). A β is synthesized by sequential enzymatic cleavage of amyloid precursor protein (APP). Besides suppressing the expression of APP (Nagpure and Bian 2014), H₂S has also been reported to inhibit the activities and expressions of the cleaving enzymes, BACE1 (Zhang et al. 2011) and γ -secretase (Nagpure and Bian 2014). In an interesting study investigating the effects of H₂S-rich Tabiano's spa-water on three experimental models of AD, Giuliani et al. found the lowered phosphorylation of tau protein at Thr181,

Ser396, and Ser202. In this particular study they also observed anti-amyloidogenic effects of H₂S in AD mouse model harboring human transgenes APPSwe, PS1M146V, and tauP301L (Giuliani et al. 2013). The therapeutic strategy of tau manipulation is of great interest. Many studies have revealed that heat-shock proteins (Hsps), the molecular chaperone families, are involved in prevention of tau aggregation and tau degradation (Thompson et al. 2012; Voss et al. 2012). As H₂S is known to elevate the expression of hsp90 (Xie et al. 2012), it is possible that H₂S could be used as a tau aggregation inhibitor to maintain intracellular microtubule infrastructure

H₂S targets multiple processes underlying AD pathology (Fig. 2) while exerting its neuroprotective effect. Hence H₂S can be of potential therapeutic value in treatment of AD.

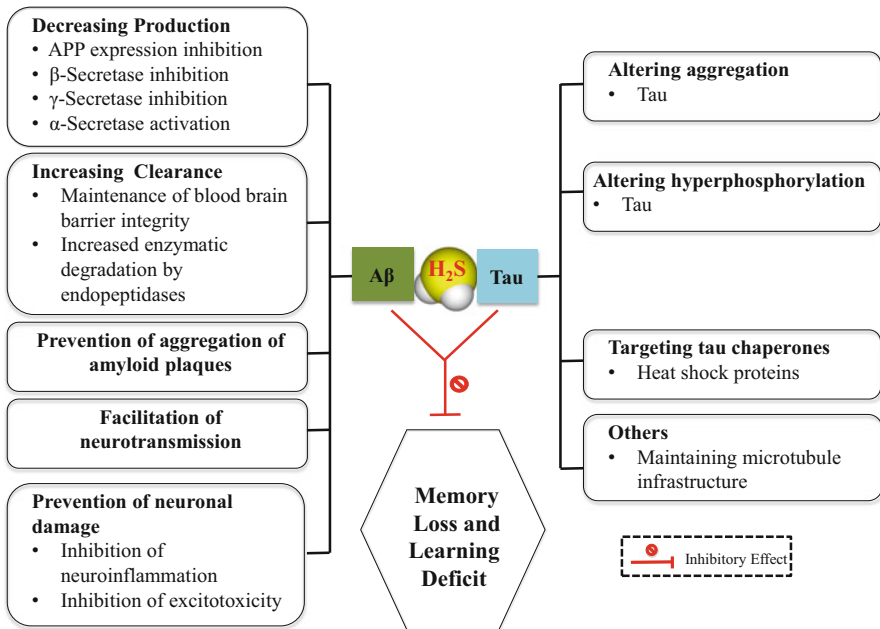


Fig. 2 Effect of H₂S on Aβ and tau aggregation and its potential role in prevention of memory loss and learning disabilities. H₂S inhibits aggregation of amyloid protein by decreasing the amyloid burden and enhancing its clearance. H₂S facilitates neurotransmission by acting as a neuro-modulator. It also inhibits the production of several pro-inflammatory cytokines, maintaining the microenvironment in which neurons can function properly. By preventing hyperphosphorylation of tau proteins and inhibiting tau chaperons (Hsps), H₂S can inhibit its aggregation into neurofibrillary tangles and preserve intracellular microtubule infrastructure

4.2 Vascular Dementia

Vascular dementia (VD), a heterogeneous group of brain disorders in which cognitive impairment is attributable to cerebrovascular pathologies, is responsible for at least 20 % of cases of dementia, being second only to AD (Gorelick et al. 2011; Iadecola 2013). Zhang et al. suggest that H₂S could protect the brain against VD injury induced by cerebral ischemia reperfusion through inhibiting the apoptosis in the hippocampus. They found that NaHS-treated rats had a greater ratio of Bcl-2 (anti-apoptotic) over Bax (pro-apoptotic) with increased Bcl-2 expression and decreased Bax expression in the hippocampus (Zhang et al. 2009). It is generally believed that inflammation (Malaguarnera et al. 2006; Liu et al. 2007), oxidative stress (Liu et al. 2007), and vascular factors (Brown et al. 2007; Stephan and Brayne 2008) play important roles in the VD pathology. As discussed earlier in this chapter, H₂S possesses potent anti-inflammatory (Hu et al. 2007) and anti-oxidative action (Kimura and Kimura 2004). Hence, it appears that H₂S may protect against VD injury by targeting multiple signaling pathways and events.

4.3 PD

PD is the second most common neurodegenerative disease histopathologically characterized by progressive degeneration of dopaminergic neurons in substantia nigra of midbrain.

Many studies indicate that hyperhomocysteinemia is common in the patients of PD (O'Suilleabhain et al. 2004; Zoccolella et al. 2010). Various *in vivo* experiments in PD animal models detected reduced levels of H₂S in substantia nigra and striatum regions of the brain. These findings suggest that impaired endogenous production of H₂S has a substantial effect on pathogenesis and progression of PD. Furthermore, the administration of exogenous H₂S has shown protective effects against underlying pathologic mechanisms. The study conducted by Yin et al. demonstrated that NaHS, a fast H₂S donor, protected PC12 cells from cytotoxicity and apoptosis induced by MPP⁺, the active metabolite of MPTP. They found that H₂S inhibited the loss of MMP and the accumulation of intracellular ROS (Yin et al. 2009). Recently, Xie et al. evaluated the therapeutic effects of ACS 84, a H₂S-releasing L-DOPA derivative, in a 6-OHDA-induced PD model (Xie et al. 2013). In this study, they observed that ACS84 not only attenuated 6-OHDA-induced cell injury and ROS production but also downregulated antioxidant enzyme expression. These results bolster the initial findings of Calvert and colleagues (Calvert et al. 2009) about upregulation of endogenous antioxidants by H₂S via stimulation of nuclear-factor-E2-related (Nrf2)-dependent signaling pathway. Previously, it was shown that pretreatment with NaHS could suppress rotenone-induced cellular injury and apoptotic cell death in human neuroblastoma SH-SY5Y cells via regulation of mitoK_{ATP} channel/p38- and JNK-MAPK pathway (Hu et al. 2009). In yet another study using neurotoxin 6-OHDA to induce PD

model, H₂S was shown to inhibit cell injury by stimulating pro-survival PKC/PI3K/Akt pathway (Tiong et al. 2010).

These findings in vitro studies were supported by the observations done in animal models of PD. Hu et al. found out that the systemic administration of NaHS dramatically reversed the progression of movement dysfunction, loss of tyrosine-hydroxylase (TH)-positive neurons in the striatum, and the elevated malondialdehyde level in injured striatum caused by 6-OHDA or rotenone (Hu et al. 2010). Inhaled H₂S also prevented the MPTP-induced movement disorder and the degeneration of TH-containing neurons by upregulating heme oxygenase-1 and glutamate-cysteine ligase (Kida et al. 2011). The anti-inflammatory, antioxidant, and neuroprotective properties shown by new H₂S releasing hybrids are encouraging, thus making them ideal candidates for PD treatment. ACS 84, a well-known L-DOPA hybrid, has been effective in reducing the release of pro-inflammatory cytokines and NO from stimulated microglia and astrocytes (Lee et al. 2010). Besides relieving from inflammation, H₂S-releasing L-DOPA hybrids also restore the depleted dopamine levels by inhibiting mono-amine oxidase B activity (Sparatore et al. 2011).

These findings highlight the potential therapeutic benefit of H₂S in PD which can be achieved either by the administration of exogenous H₂S or the modulation of endogenous H₂S production.

5 Learning and Memory

The life-long thoughts, behavior, and actions of humans are mostly controlled by two processes occurring in CNS: learning and memory. The intrinsic cellular mechanisms are different for learning (procurement), consolidation, and evocation of memories (Kandel 2001). Notably, the modifications of synaptic connectivity, in terms of number and structures of synapses, are probably the underlying mechanisms for learning and memory. The ability of synaptic plasticity of neurons relies heavily on many intracellular secondary signaling pathways. Therefore, modulation of synthesis of proteins involved in activation and/or deactivation of signal transduction pathways in CNS plays a key role in the processes of learning and memory (Milner et al. 1998).

5.1 Endogenous H₂S Levels: Relation with Learning and Memory

As mentioned earlier, H₂S is generated endogenously in brain, including hippocampus. It is a small but an important region primarily associated with learning and memory functions. The intracellular concentrations of a CBS activator, S-adenosylmethionine (SAM), were found to be decreased in AD brains than those in the normal brains (Morrison et al. 1996). In an extensive clinical study conducted in many patients of AD and VD, the plasma H₂S levels were demonstrated to be significantly lower than the normal controls (Liu et al. 2008). In accordance, Eto

et al. showed that the levels of H₂S are drastically declined in the brains of AD patients compared with those of the age-matched normal individuals (Eto et al. 2002). A recent work characterized the levels of H₂S at various time points through the development and progression of AD in double-transgenic APP/PS1 mice. This research group observed the decreased levels of H₂S and protein expression of CBS in the cortex and hippocampus of 9- and 12-month-old AD mice (He et al. 2014). Thus, the plasma H₂S concentration is directly correlated with the severity of memory-related symptoms of AD

5.2 Effect of H₂S on Glutamatergic Neurotransmission

The interaction of H₂S with the glutamatergic neurotransmission is well established. Kimura et al. found that H₂S enhances the responses of NMDA receptors to the neurotransmitter glutamate in neurons (Abe and Kimura 1996). According to the previously published studies, H₂S protects brain cells from death due to elevated glutamate levels in synapses. The underlying mechanism is via enhancing glutamate uptake function by glutamate transporters (Lu et al. 2008). It is well known that H₂S attenuates the development of opioid dependence and alleviates heroin withdrawal symptoms in animals. The neural mechanism involved in these effects is closely related to inhibition of cAMP/PKA pathway (Jiang et al. 2012; Yang et al. 2013). Although this is the case, the modulation of the glutamatergic system can also be involved since presynaptic glutamate synaptic transmission by NMDA receptors is regulated directly by μ -opioid receptors (Garzon et al. 2012). Thus, these results signify the importance of H₂S in modulating adaptations in the brain glutamatergic system involved in synaptic plasticity.

5.3 Role of H₂S in Intracellular Calcium ([Ca²⁺]_i) Regulation

Many research groups have evaluated possible mechanisms by which H₂S modulates learning and memory processes. The effect of H₂S on [Ca²⁺]_i homeostasis deserves a special mention here. [Ca²⁺]_i is critical for normal neuron-glia communication and regulation of synaptic plasticity. It has been found that H₂S is capable of regulating [Ca²⁺]_i in all important brain cell types, namely neurons (Yong et al. 2010), microglia (Lee et al. 2006), and astrocytes (Nagai et al. 2004). LTP is considered as a major cellular mechanism underlying the learning and memory functions. Kimura et al. discovered that the physiological concentrations of H₂S selectively enhance NMDA receptor-mediated currents and facilitate the induction of hippocampal LTP (Abe and Kimura 1996). L-type calcium channels contribute to LTP and fear memory formation (Bauer et al. 2002). Furthermore, they are essential in both the acquisition and retrieval of long-term recognition memory (Seoane et al. 2009). H₂S activates L-type calcium channels and NMDA receptors on plasma membrane (Lee et al. 2006). Yong et al. discovered that the

action of H₂S on [Ca²⁺]_i can be suppressed by using inhibitors of PKA, phospholipase C (PLC), and protein kinase C (PKC), suggesting the role of PKA and PLC/PKC pathways in the regulatory effect of H₂S on [Ca²⁺]_i (Yong et al. 2010) and possibly on functions of learning and memory. Voltage-gated Ca_v2.1 (P/Q-type) Ca²⁺ channels are essential in modulation of processes such as neurotransmitter release and synaptic plasticity. It was found that context-associated memory retrieval was dependent on these particular Ca²⁺ channels (Chen et al. 2012). Mallmann et al. observed deficits in spatial learning and reference memory, reduced recognition memory in the forebrain-specific Ca_v2.1 knockout mice (Mallmann et al. 2013). Interestingly these channels are also believed to be involved in AD pathology as Aβ downregulates Ca_v2.1 channels, thus blocking Ca²⁺ current (Nimmrich et al. 2008; Nimmrich and Ebert 2009). Furthermore, Gangarossa suggested the potential functional involvement of these specific Ca²⁺ channels in many brain disorders as the physiological activity of these specific Ca²⁺ channels is required for affective and cognitive behaviors (Gangarossa et al. 2014). H₂S can modulate the activity of voltage-gated Ca²⁺ channels. A recent study conducted by Sekiguchi et al. demonstrated that the function of these channels is tonically enhanced by endogenous H₂S synthesized by CSE in HEK293 cells transfected with Cav3.2, and that exogenous H₂S is capable of enhancing Cav3.2 function when endogenous H₂S production by CSE is inhibited (Sekiguchi et al. 2014).

5.4 Neuroinflammation and Other Factors

Neuroinflammation involves elevated production of myriad of pro-inflammatory cytokines, ROS, and activation of apoptotic pathways. These events lead to neurodegeneration as they negatively impact the functions of neurons, which affect the processes of learning and memory. In various behavior studies, it has been seen that the H₂S reverses neuroinflammatory changes, ultimately repairing the cognitive and memory impairments.

In the initial studies done in rat models of neuroinflammation, exogenous administration of H₂S in the form of NaHS (Gong et al. 2010) or *S*-propargylcysteine (SPRC) (Gong et al. 2011) ameliorated LPS-induced cognitive impairment as evaluated in the Morris water maze test. Shortly afterwards, Tang et al. demonstrated that the disturbances in endogenous H₂S generation played a pivotal role in formaldehyde-induced deficits in memory and cognition in animals (Tang et al. 2013). These results were confirmation of the findings of other study in which the neurotoxic effects of Hcy were investigated. Elevated plasma level of Hcy is a known risk factor for AD. The spatial memory acquisition and spatial learning as evaluated by probe trial and hidden-platform acquisition tests, respectively, showed that Hcy decreases spatial learning ability and memory of rats. They also employed novel object recognition test to study short-term, declarative memory and attention. Hcy, by decreasing endogenous production of H₂S, significantly decreased the discrimination index of rats in novel object recognition test highlighting the unsuccessful retention of memory of familiar objects (Tang

et al. 2011). Furthermore, the intraperitoneal administration of H₂S attenuated the spatial memory impairment in A β rat model of AD (Xuan et al. 2012). In continuation of their previous study, Xuan et al. demonstrated that the application of exogenous H₂S resulted in improved spatial learning and memory acquisition in double-transgenic APP/PS1 mice, an established animal model of AD (He et al. 2014)

6 Toxicity of H₂S

Although H₂S is produced endogenously in the brain and plays important biological functions, one should bear in mind that long-term treatment of neurodegenerative diseases may potentially cause brain H₂S accumulation and therefore neurotoxicity. Previous studies have shown that the exposure-response curve of H₂S is steep, and thus concentration of inhaled gas is more important compared to the duration of exposure (Prior et al. 1988; Guidotti 1996). The approximate concentrations (exposure levels) of inhaled H₂S for the major toxicological effects are given in Fig. 3. The toxidrome (i.e., a set of symptoms and signs associated with a particular poison) of H₂S is often considered as one of the most unusual and reliable toxidromes (Wang 1989; Milby and Baselt 1999). It is characterized by the “knock-down” (acute central neurotoxicity), pulmonary edema, conjunctivitis, and odor perception followed by respiratory paralysis (Guidotti 2010). Acute toxicity leading to reversible unconsciousness caused by H₂S inhalation is called as “knockdown” (Guidotti 1996). Although knockdowns can be fatal in the cases of prolonged high-concentration exposure (about 500–1000 ppm), the transient exposure is often reversible and apparently complete functionally (Burnett et al. 1977). Pulmonary edema is a well-recognized effect of acute H₂S toxicity. As H₂S has relatively low solubility, it penetrates deeply into respiratory track, causing alveolar injury culminating in acute pulmonary edema (Guidotti 2010). The conjunctivitis caused by prolonged low-concentration exposure (about 20 ppm) (Lambert et al. 2006) is peculiarly associated with reversible chromatic distortion and visual changes. These symptoms are sometimes accompanied by blepharospasm and photophobia (Tansy et al. 1981; Milby and Baselt 1999). H₂S is an odorous gas at low concentration (0.01–0.3 ppm). As the concentration increases, however, the victims start to experience olfactory fatigue. It is a sensory adaptation where the victims get accustomed to strong odor. At around 100 ppm concentration, H₂S paralyzes the olfactory mechanism, preventing perception of any smell. This phenomenon removes the primary warning sign of H₂S exposure (Ronk and White 1985; Turner et al. 1990).

The primary mechanism underlying H₂S toxicity is the inhibition of mitochondrial respiratory chain by HS[−] (a hydrogen sulfide ion). HS[−] binds to ferric iron (Fe³⁺) of cytochrome oxidase C. It culminates into sulfmethemoglobinemia, lactic acidosis, and hypoxia (Sastre et al. 2013). H₂S is not completely oxidized by brain, making it most vulnerable for the toxic effects. Initially, this pattern seemed identical to that observed in case of cyanide (HCN) poisoning (Dorman et al. 2002).

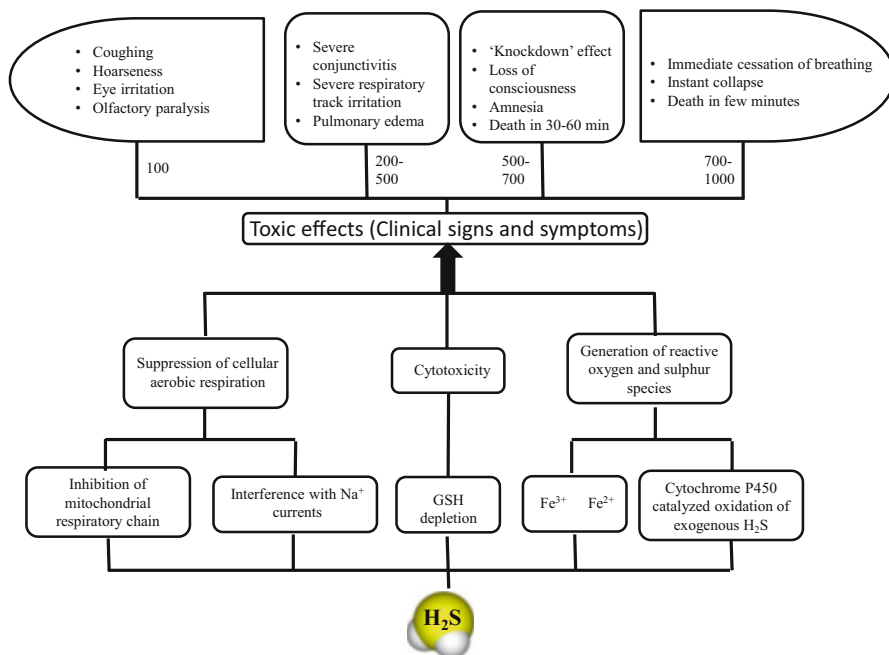


Fig. 3 Toxicity of H_2S . The spectrum of toxic effects of H_2S spans from mild irritation of eyes and respiratory tract to immediate deaths. The severity of symptoms depends on the exposure levels as depicted in the figure. The “toxidrome” of H_2S is often considered as fairly constant and reliable. The underlying molecular mechanisms for toxicity are varied. Along with suppression of cellular aerobic respiration, excessive generation of reactive molecules is proposed as primary mechanisms. The depletion of cytoprotective GSH also makes cells vulnerable for cellular injury

However, the anomalies began to appear when methemoglobin treatment did not yield effective results in cases of H_2S poisonings, despite its successful use in HCN poisoning (Truong et al. 2006). Hence, many other underlying mechanisms were proposed. Warenycia et al. demonstrated that H_2S interferes with sodium currents by abolishing sodium channel function, which might be responsible for depletion of cellular respiration in H_2S poisoning (Warenycia et al. 1989). H_2S was also found responsible for reduction of bound form of iron (ferric, Fe^{3+}) to free ferrous (Fe^{2+}) form. This along with cytochrome P450-catalyzed oxidation of the exogenous H_2S compound resulted into reactive sulfur and oxygen species generation. The concurrent depletion of GSH made neuronal cells more susceptible to neurotoxicity (Truong et al. 2006).

7 Concluding Remarks

The prevalence and morbidity of neurodegenerative diseases are increasing at rapid pace across the globe. Despite the investigations studying the therapeutic effects of H₂S are still in their infancy, plentiful evidence has proven a protective role for this gasotransmitter in the pathology of neurodegenerative diseases. However, our present knowledge on neuroprotection offered by H₂S mainly comes from cell and animal studies with the use of H₂S donors and inhibitors of endogenous H₂S. Whether the therapeutic effects of these donors and inhibitors in “bench” studies can be transferred to “bedside” clinical studies needs to be explored. Additionally, more information about drug safety and toxicity due to long-term H₂S-based therapeutic approaches is desirable. The development of H₂S-releasing drug with a sustained and controlled release is necessary due to non-physiological rapid generation of H₂S by most H₂S donors. Thorough understanding of these problems will enable us to study underlying mechanisms still deeper, and formulate H₂S-based therapeutic interventions to treat neurodegenerative diseases.

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H₂S and Pain: A Novel Aspect for Processing of Somatic, Visceral and Neuropathic Pain Signals

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Abstract

Hydrogen sulfide (H₂S) formed by multiple enzymes including cystathionine- γ -lyase (CSE) targets Ca_v3.2 T-type Ca²⁺ channels (T-channels) and transient receptor potential ankyrin-1 (TRPA1). Intraplantar and intracolonic administration of H₂S donors promotes somatic and visceral pain, respectively, via activation of Ca_v3.2 and TRPA1 in rats and/or mice. Injection of H₂S donors into the plantar tissues, pancreatic duct, colonic lumen, or bladder causes T-channel-dependent excitation of nociceptors, determined as phosphorylation of ERK or expression of Fos in the spinal dorsal horn. Electrophysiological studies demonstrate that exogenous and/or endogenous H₂S facilitates membrane currents through T-channels in NG108-15 cells and isolated mouse dorsal root ganglion

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(DRG) neurons that abundantly express $\text{Ca}_v3.2$ and also in $\text{Ca}_v3.2$ -transfected HEK293 cells. In mice with cerulein-induced pancreatitis and cyclophosphamide-induced cystitis, visceral pain and/or referred hyperalgesia are inhibited by CSE inhibitors and by pharmacological blockade or genetic silencing of $\text{Ca}_v3.2$, and CSE protein is upregulated in the pancreas and bladder. In rats with neuropathy induced by L5 spinal nerve cutting or by repeated administration of paclitaxel, an anticancer drug, the neuropathic hyperalgesia is reversed by inhibitors of CSE or T-channels and by silencing of $\text{Ca}_v3.2$. Upregulation of $\text{Ca}_v3.2$ protein in DRG is detectable in the former, but not in the latter, neuropathic pain models. Thus, H_2S appears to function as a nociceptive messenger by facilitating functions of $\text{Ca}_v3.2$ and TRPA1, and the enhanced function of the CSE/ H_2S / $\text{Ca}_v3.2$ pathway is considered to be involved in the pancreatitis- and cystitis-related pain and in neuropathic pain.

Keywords

H_2S • $\text{Ca}_v3.2$ T-type calcium channel • TRPA1 • Pain

1 Introduction

Hydrogen sulfide (H_2S), an endogenous gasotransmitter, is generated from L-cysteine mainly by cystathionine- β -synthase (CBS), cystathionine- γ -lyase (CSE), or 3-mercaptopyruvate sulfurtransferase (3-MST) along with cysteine aminotransferase (CAT) throughout the mammalian body (Kimura 2010) including the heart (Geng et al. 2004), blood vessels (Zhao et al. 2001), and central nervous system (CNS) (Warenycia et al. 1989). H_2S appears to play dual or complex roles in many tissues/organs. In the nervous system, there is evidence not only for the neurotoxicity of H_2S through activation of glutamate receptors including NMDA receptors (Cheung et al. 2007) and activation of the MEK/ERK pathway (Kurokawa et al. 2011) but also for the neuroprotection by H_2S via multiple mechanisms such as activation of ATP-sensitive K^+ (K_{ATP}) and Cl^- channels and elevation of intracellular glutathione levels (Kimura and Kimura 2004; Kimura et al. 2006). Endogenous H_2S is also involved in neuronal excitation via the Ca^{2+} /calmodulin-mediated pathway (Eto et al. 2002) and modification of long-term potentiation (Abe and Kimura 1996) via enhancement of the cAMP-induced NMDA receptor response (Kimura 2000).

H_2S appears to cause excitation of sensory nerves and play important roles in neurogenic inflammation, since NaHS, an H_2S donor, increases neuropeptide release in guinea-pig airways, an effect attenuated by capsaicin desensitization of C-fiber neurons and by the transient receptor potential vanilloid-1 (TRPV1) antagonist capsazepine. In addition, NaHS induces contraction of isolated airways, which is abolished by TRPV1 antagonism and pretreatment with the combination of tachykinin NK_1 and NK_2 receptor antagonists (Trevisani et al. 2005). It is consistent with the evidence that H_2S stimulates capsaicin-sensitive primary afferent nerve

terminals, from which tachykinins are released to produce concentration-dependent contractile responses by activating NK₁ and NK₂ receptors in the rat urinary bladder (Patacchini et al. 2004). Nonetheless, TRPV1 is not considered a direct target for H₂S, and Ca_v3.2 T-type Ca²⁺ channels and transient receptor potential ankyrin-1 (TRPA1) rather appear responsible for excitation of sensory neurons (Miyamoto et al. 2011; Andersson et al. 2012; Pozsgai et al. 2012; Sekiguchi and Kawabata 2013; Donatti et al. 2014). Sensory plasticity in nociception, known as sensitization, is thought to contribute significantly to the production of persistent, often pathological pain. Here, we review the role of H₂S in somatic, visceral, and neuropathic pain processing.

2 Targets of H₂S in Pain Processing

2.1 Ca_v3.2 T-Type Ca²⁺ Channels

Ca_v3.2 T-type Ca²⁺ channels (T-channels) are activated at near-resting membrane potential and play a crucial role in excitability of both central and peripheral neurons. T-channels are expressed in dorsal root ganglion (DRG) neurons including the peripheral and central axons/endings, suggesting the role of T-channels in sensory transmission (Fig. 1) (Nelson et al. 2005; Todorovic and Jevtovic-Todorovic 2011; Rose et al. 2013). Three isoforms of T-channels have been

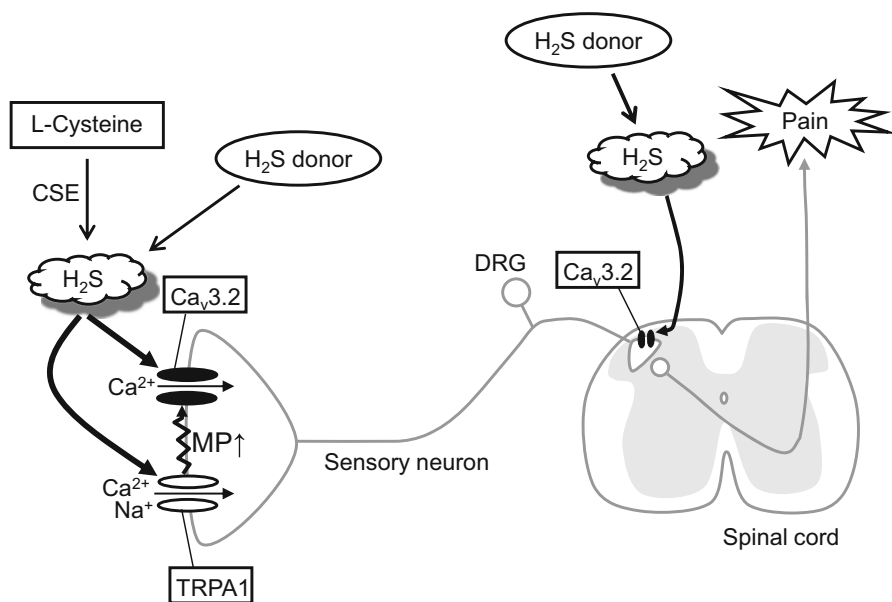


Fig. 1 Roles of H₂S in nociceptive processing at the peripheral and central terminals of the primary afferent neurons. *CSE* cystathionine- γ -lyase, *MP* membrane potential, *DRG* dorsal root ganglion

identified, i.e., $Ca_v3.1$, $Ca_v3.2$, and $Ca_v3.3$ (Perez-Reyes 2003). Studies using *in situ* hybridization have demonstrated that mRNA for $Ca_v3.2$ is the most abundant isoform of T-channels in peripheral sensory neurons (Todorovic and Jevtovic-Todorovic 2011). Bourinet and his coworkers have demonstrated thermal and mechanical analgesia in rats with molecular knockdown of $Ca_v3.2$ T-channels in DRG neurons (Bourinet et al. 2005). $Ca_v3.2$ T-channels are much more sensitive to inhibition by metals, such as zinc, copper, and nickel, than $Ca_v3.1$ or $Ca_v3.3$ (Todorovic and Jevtovic-Todorovic 2011; Sekiguchi and Kawabata 2013). It is now clear that a histidine residue at position 191 (His191) in the second extracellular loop of domain I of $Ca_v3.2$ is a critical determinant for the trace metal inhibition and is not conserved in $Ca_v3.1$ or $Ca_v3.3$ (Iftinca and Zamponi 2009). $Ca_v3.2$ T-channels appear to be tonically exposed to inhibition by Zn^{2+} in cultured cells and possibly in mammalian tissues under physiological conditions, and compounds that chemically interact with Zn^{2+} , such as L-cysteine and H_2S , selectively enhance the channel functions of $Ca_v3.2$ among the three T-channel isoforms (Nelson et al. 2007a; Matsunami et al. 2011; Sekiguchi and Kawabata 2013). Moreover, L-ascorbic acid, a reducing compound, at physiologically relevant concentrations selectively inhibits native T-currents in the DRG and thalamus and diminishes T-channel-dependent burst firing in reticular thalamic cells. This highly subtype-specific effect was achieved via metal-catalyzed oxidation of critical metal-binding and $Ca_v3.2$ -unique His191 residue (Nelson et al. 2007b). NaHS, an H_2S donor, facilitates membrane currents through T-channels in NG108-15 cells that naturally express $Ca_v3.2$ (Kawabata et al. 2007; Nagasawa et al. 2009) and in isolated mouse DRG neurons that abundantly express $Ca_v3.2$ (Matsunami et al. 2009). In $Ca_v3.2$ -transfected HEK293 cells, T-currents are not altered by H_2S donors, NaHS at 1.5 mM and Na_2S at 0.1 mM, but strongly suppressed by DL-propargylglycine (PPG), an inhibitor of CSE, an H_2S -forming enzyme. Further, in the presence of the CSE inhibitor, the H_2S donors, NaHS at 1.5 mM and Na_2S at 0.1–0.3 mM, enhance the decreased $Ca_v3.2$ T-currents, and CSE is abundantly expressed in both $Ca_v3.2$ -transfected and background HEK293 cells (Sekiguchi et al. 2014). Collectively, endogenous and exogenous H_2S appears to positively regulate the functions of $Ca_v3.2$ T-channels (Fig. 1). An independent group has reported that NaHS at low (0.1–1 mM) and high concentrations (3 mM) suppresses and enhances T-currents by 25–30 %, respectively, in $Ca_v3.2$ -transfected HEK293 cells or randomly chosen rat DRG cells (Elies et al. 2014), whereas we have never detected the suppressive effect of NaHS or Na_2S even at low concentrations on T-currents in our $Ca_v3.2$ -transfected HEK293 cells (Sekiguchi et al. 2014), NG108-15 cells (Kawabata et al. 2007), or small mouse DRG neurons (less than 30 μm in cell body diameter) (Matsunami et al. 2009), known to be rich in T-type Ca^{2+} channels (Todorovic and Jevtovic-Todorovic 2011). Particularly, Na_2S at 0.1–0.3 mM rather augments T-currents in $Ca_v3.2$ -transfected HEK293 cells pretreated with the CSE inhibitor (Sekiguchi et al. 2014), as mentioned above, being inconsistent to the suppressive effect of NaHS at equivalent concentrations in their report (Elies et al. 2014). The discrepancy has yet to be interpreted, although the effect of H_2S on $Ca_v3.2$ channels *in vitro* and *in vivo* might be dependent on the abundance

of H₂S-forming enzymes including CSE, the formation or contamination of polysulfides, and the concentrations of Zn²⁺ and ascorbic acid that negatively regulate Ca_v3.2 functions (Todorovic and Jevtovic-Todorovic 2011; Kimura et al. 2013; Sekiguchi and Kawabata 2013; Sekiguchi et al. 2014). Particularly, levels of Zn²⁺ would have great impact on the Ca_v3.2 channel activity and its regulation by H₂S, since Zn²⁺ binds to not only His191 of Ca_v3.2 (Todorovic and Jevtovic-Todorovic 2011; Sekiguchi and Kawabata 2013) but also H₂S itself (Munchberg et al. 2007). Given that Ca_v3.2 is involved in somatic and visceral nociceptive processing and in neuropathic pain (Todorovic and Jevtovic-Todorovic 2011; Francois et al. 2014), regulation of Ca_v3.2 by endogenous and exogenous H₂S is considered to have great impact on the pathophysiology of inflammatory and neuropathic pain (Sekiguchi et al. 2014). The T-channel suppression by NaHS at low doses, if any, might be overcome by TRPA1 activation by NaHS *in vivo*, as described below.

2.2 Transient Receptor Potential Ankyrin-1

Transient receptor potential ankyrin-1 (TRPA1), a member of the TRP channel family, is abundantly expressed in a subpopulation of unmyelinated nociceptors that also express TRPV1. It is activated by a diverse assortment of pungent or irritating reactive chemical compounds including those found in mustard oil (allylisothiocyanate), cinnamon oil (cinnamaldehyde), exhaust (acrolein), raw garlic, and onions (allicin). Moreover, TRPA1 has been put forth as a putative transducer of natural physical stimuli including both cold (<17 °C) and mechanical force (Stucky et al. 2009). It has been reported that TRPA1 is sensitized by proteinase-activated receptor-2 (PAR2) activation, contributing to inflammatory pain (Dai et al. 2007). We have provided evidence that pharmacological inhibition of TRPA1 prevents the spinal Fos expression following infusion of SLIGRL-NH₂, a PAR2-activating peptide, into the pancreatic duct, suggesting the pronociceptive role of TRPA1 as a downstream signal of PAR2 activation in the pancreatic nociceptors (Terada et al. 2013). Most interestingly, it has been reported that NaHS evokes a time- and concentration-dependent increase in cytosolic calcium concentration in TRPA1-transfected cells (Streng et al. 2008) and that NaHS induces calcium influx and inward currents in rat DRG cells, an effect inhibited by a TRPA1 antagonist (Miyamoto et al. 2011; Ogawa et al. 2012). The H₂S-induced TRPA1 activation appears to involve H₂S modulation of two cysteine residues located in the N-terminal intracellular domain of the TRPA1 (Ogawa et al. 2012). These findings suggest that TRPA1 in collaboration with Ca_v3.2 mediates H₂S-induced excitation of nociceptors, participating in pain processing (Fig. 1).

3 Somatic Pain

We first reported that intraplantar (i.pl.) administration of NaHS produces prompt hyperalgesia in rats, accompanied by expression of Fos protein, a marker for neuronal excitation, in the spinal dorsal horn, and that the H₂S-evoked hyperalgesia is blocked by ethosuximide and mibefradil, T-channel blockers (Kawabata et al. 2007). Another H₂S donor, Na₂S, also facilitates hyperalgesia by i.pl. administration, an effect completely blocked by pretreatment with NNC 55-0396 (NNC), a T-channel blocker (Sekiguchi et al. 2014). L-Cysteine, an endogenous source for H₂S, given i.pl., mimics the H₂S-induced hyperalgesia, an effect being abolished by PPG and β-cyanoalanine (BCA), inhibitors of CSE, in rats (Kawabata et al. 2007). The NaHS-induced hyperalgesia is also suppressed by pretreatment with i.pl. zinc chloride and/or ascorbic acid, known to selectively inhibit Ca_v3.2 among three T-channel isoforms, and by silencing of Ca_v3.2 T-channels in the sensory neurons by repeated intrathecal (i.t.) administration of antisense oligodeoxynucleotides (ODNs) targeting Ca_v3.2 T-channels in rats or mice (Maeda et al. 2009; Okubo et al. 2012). Interestingly, i.t. administration of NaHS also rapidly decreases nociceptive threshold in rats, as determined by the paw pressure method. The hyperalgesia caused by i.t. NaHS is also abolished by co-administration of mibefradil and pretreatment with i.t. zinc chloride and by silencing of Ca_v3.2 protein in the DRG (Maeda et al. 2009). These findings are consistent with the evidence that presynaptic Ca_v3.2 T-channel regulates spontaneous excitatory synaptic neurotransmitter release from the central terminal of nociceptors in the spinal dorsal horn, in addition to neuronal excitability in the peripheral endings of nociceptors (Jacus et al. 2012). Thus, positive regulation of Ca_v3.2 T-channel by H₂S is considered to have a great impact on nociceptive processing at both peripheral and central endings of the primary afferent neurons (Fig. 1). We have also demonstrated that the NaHS-induced hyperalgesia/allodynia in mice is significantly suppressed by AP18, a TRPA1 blocker, and by silencing of TRPA1 in the sensory neurons (Okubo et al. 2012), suggesting that TRPA1 and Ca_v3.2 mediate the pronociceptive effect of H₂S (Fig. 1). The involvement of TRPA1 in H₂S-induced hyperalgesia has been confirmed by independent studies (Andersson et al. 2012; Ogawa et al. 2012). It is likely that H₂S causes activation of TRPA1 followed by elevation of membrane potentials that activates Ca_v3.2 and then enhances the evoked Ca_v3.2 T-channel currents, leading to increased nociceptor excitability and hyperalgesia. Thus, both TRPA1 and Ca_v3.2 are considered to participate in the H₂S-induced hyperalgesia (Fig. 1).

We have shown that PPG and BCA, CSE inhibitors, partially block the hyperalgesia induced by i.pl. lipopolysaccharide (LPS), an effect being reversed by i.pl. NaHS (Kawabata et al. 2007). The suppression of LPS-induced hyperalgesia by CSE inhibitors might be associated with the prevention of neutrophil recruitment to the plantar tissue (Cunha et al. 2008). H₂S is also pronociceptive in the formalin model of persistent inflammatory pain in rats. Injection of formalin into the hindpaw evokes a dose-dependent increase in the concentration of H₂S in the paw tissue. Both nociceptive flinching and paw edema induced by formalin are

attenuated by pretreatment with PPG. Furthermore, PPG pretreatment attenuates the induction of c-Fos in spinal laminae I–II following injection of formalin (Lee et al. 2008). It has also been reported that pretreatment of mice with PPG reduces zymosan-induced articular hyperalgesia (Cunha et al. 2008). Recent evidence has shown that the inflammatory pain in temporomandibular joint by injection of complete Freund's adjuvant in rats is attenuated by subcutaneous administration of aminooxyacetic acid (AOAA), known as a CBS inhibitor (Miao et al. 2014), although the selectivity of AOAA for CBS is now questioned (Asimakopoulou et al. 2013). Thus, there is plenty of evidence that H₂S participates in inflammatory pain processing. Nevertheless, it is noteworthy that exogenously applied H₂S reveals antinociceptive activity in certain experimental conditions through activation of K_{ATP} channels or unknown mechanisms (Cunha et al. 2008).

4 Visceral Pain

4.1 Pancreatic Pain

Clinically, pancreatic pain is a serious problem for patients with pancreatitis or pancreatic cancer. In humans, pain elicited from the pancreas is often referred to the upper abdominal area and radiates to the back, and these skin areas are usually tender to touch. In pancreatic acinar cells isolated from mice, treatment with cerulein increases the levels of H₂S and CSE mRNA expression (Tamizhselvi et al. 2007). Furthermore, the activity and protein expression of CSE in the pancreatic tissues increase after the development of cerulein-induced pancreatitis in mice (Nishimura et al. 2009). Given the prevention by PPG, a CSE inhibitor, of cerulein-induced pancreatitis and referred hyperalgesia/allodynia (Bhatia et al. 2005; Nishimura et al. 2009), H₂S is considered to play a critical role in the development of acute pancreatitis and related pain. NaHS infused into the pancreatic duct causes the expression of Fos protein (Nishimura et al. 2009) and ERK phosphorylation (Fukushima et al. 2010), delayed and rapid markers for neuronal activation, respectively, in the superficial layers of spinal cord in anesthetized mice or rats, which are attenuated by i.p. administration of mibefradil, a T-channel blocker. Mibefradil also suppresses the referred hyperalgesia/allodynia in mice with cerulein-induced pancreatitis (Nishimura et al. 2009). Together, H₂S produced by upregulated CSE during the development of pancreatitis augments the function of T-channels, contributing to pancreatitis-related pain. As described above for somatic pain processing, TRPA1 in addition to T-channels appears to mediate H₂S-induced pancreatic nociception in naïve animals, because the TRPA1 inhibitor AP18 or the T-channel inhibitor NNC 55-0396 prevents spinal Fos expression following injection of NaHS into the pancreatic duct in mice without pancreatitis. In contrast, in mice with cerulein-induced pancreatitis, NNC 55-0396 or genetic silencing of Ca_v3.2 abolishes the referred hyperalgesia accompanying cerulein-induced pancreatitis, while AP18 or genetic TRPA1 silencing just facilitates the effect of NNC 55-0396, but had no significant effect on the pancreatitis-related pain

by itself (Terada et al. 2014). Given the downregulation of TRPA1, but not $\text{Ca}_v3.2$, in the pancreatic tissue following cerulein treatment (Terada et al. 2014), $\text{Ca}_v3.2$ is considered to primarily participate in the pancreatitis-related pain, while TRPA1 appears to play a secondary role in nociceptive signaling during pancreatitis.

4.2 Colonic Pain

In the colonic lumen, H_2S is also produced by sulfate-reducing bacteria and may be associated with the pathogenesis of inflammatory bowel disease (IBD) and colorectal cancer (Roediger et al. 1997; Ohge et al. 2003; Taniguchi et al. 2009). Nonetheless, there is evidence for a protective role of H_2S against IBD (Wallace et al. 2009). H_2S regulates the secretion of HCO_3^- in the duodenum (Ise et al. 2011) and Cl^- in the colon (Schicho et al. 2006), predicting possible involvement of H_2S in visceral nociception. Distrutti et al. (Distrutti et al. 2006) have described that i.p. injection of H_2S donors suppressed responses to colorectal distention by activating K_{ATP} channels. However, it has been reported that CBS in colonic sensory neurons is upregulated and contributes to visceral hypersensitivity in a rat model of irritable bowel syndrome (IBS) (Xu et al. 2009) and that upregulation of CBS expression by NF- κ B contributes to visceral hypersensitivity in rats (Li et al. 2012). We have demonstrated that intracolonic (i.col.) administration of NaHS induces visceral nociceptive behavior accompanied by referred hyperalgesia/allodynia and phosphorylation of ERK in the spinal cord in mice (Matsunami et al. 2009). The behavioral effects of i.col. NaHS are abolished by mibefradil and NNC 55-0396; T-channel blockers; and zinc chloride, known to selectively inhibit $\text{Ca}_v3.2$ among three isoforms of T-channels by genetic knockdown of $\text{Ca}_v3.2$ and also to be attenuated by AP18, a TRPA1 blocker, but not verapamil, an L-type Ca^{2+} channel blocker, or glibenclamide, a K_{ATP} channel blocker (Matsunami et al. 2009; Matsunami et al. 2011). Interestingly, $\text{Ca}_v3.2$ is constantly exposed to Zn^{2+} inhibition under physiological conditions, and zinc chelators applied exogenously enhance $\text{Ca}_v3.2$ functions by cancelling the Zn^{2+} inhibition (Nelson et al. 2007a). We have shown that two distinct Zn^{2+} chelators, N,N,N',N' -tetrakis (2-pyridylmethyl)-ethylenediamine (TPEN) and dipicolinic acid, when administered intracolonicly, mimics the NaHS-evoked visceral nociceptive behavior, referred abdominal hyperalgesia/allodynia, and phosphorylation of ERK in the spinal dorsal horn, which are inhibited by T-channel blockers (Matsunami et al. 2011). The contribution of $\text{Ca}_v3.2$ T-channels to colonic hypersensitivity has been confirmed by an independent group (Marger et al. 2011). Collectively, H_2S plays critical roles in colonic pain signaling through facilitation of $\text{Ca}_v3.2$ T-channel functions by cancelling the Zn^{2+} inhibition and also activation of TRPA1 channels. Our recent study has also shown that $\text{Ca}_v3.2$ -dependent excitation of capsaicin-sensitive sensory neurons by H_2S contributes to colonic mucosal cytoprotection in rats with 2,4,6-trinitrobenzenesulfonic acid (TNBS)-induced colitis (Matsunami et al. 2012a).

4.3 Bladder Pain

The most common visceral pain in the field of urology is bladder pain, a symptom of lower urinary tract diseases such as cystitis, cystolithiasis, and bladder cancer, which impairs the quality of life in patients. Interstitial cystitis or painful bladder syndrome is an idiopathic disease, presenting with bladder pain and urinary frequency or urgency (Kelada and Jones 2007). The bladder pain accompanying interstitial cystitis is regional and chronic and spreads over the lower pelvic and suprapubic area and mimics neuropathic pain (Theoharides et al. 2008). In isolated rat urinary bladder, H₂S stimulates capsaicin-sensitive primary afferent neurons and produces contractile responses (Patacchini et al. 2004). There is also evidence that intravesical NaHS produces changes in urodynamic parameters after chemical disruption of the urothelial barrier in rats (Streng et al. 2008). Intraperitoneal administration of cyclophosphamide induces cystitis in mice, accompanied by bladder pain-like nociceptive behavior and referred hyperalgesia in the region between the anus and urethral opening, which is now recognized as an animal model of human interstitial cystitis (Wantuch et al. 2007).

We have shown that inhibition of CSE by pre-administration of PPG prevents cyclophosphamide-induced bladder nociception, i.e., bladder pain-like nociceptive behavior and referred hyperalgesia, and that expression of CSE protein is markedly increased in the bladder tissue of mice with cyclophosphamide-induced cystitis (Matsunami et al. 2012b), as observed in the pancreatic tissue of mice with cerulein-induced pancreatitis (Nishimura et al. 2009). Post-administration of T-channel blockers, mibefradil and NNC 55-0396, and the knockdown of Ca_v3.2 T-channels by the antisense method suppress the cyclophosphamide-induced bladder nociception, while neither an L-type Ca²⁺ channel blocker, verapamil, nor TRPA1 blockers, AP18 and HC-030031, exert such effect (Matsunami et al. 2012b). The T-channel-dependent pronociceptive role of H₂S in bladder nociceptive processing is also supported by our evidence that intravesical administration of NaHS causes phosphorylation of ERK in the spinal cord, an effect prevented by pretreatment with the T-channel blocker NNC 55-0396 (Matsunami et al. 2012b).

5 Neuropathic Pain

Neuropathic pain is the consequence of damage to peripheral nerves or to the CNS and common in clinical practice and greatly impairs the quality of life of patients. The neuropathic syndromes include painful diabetic neuropathy, postherpetic neuralgia, phantom limb pain, complex regional pain syndrome (CRPS), etc. The molecular mechanisms underlying neuropathic pain involve abnormal functions and/or altered expression of voltage-gated ion channels such as sodium (McGowan et al. 2009), calcium (Boroujerdi et al. 2011), and potassium channels (Cao et al. 2010) in sensory neurons. There is evidence for upregulation of T-current density in small DRG cells isolated from rats with neuropathic pain due to chronic

constrictive injury (CCI) of the sciatic nerve (Jagodic et al. 2008) and that genetic silencing of $Ca_v3.2$ T-channels reverses CCI-induced neuropathic pain (Bourinet et al. 2005). We have shown that neuropathic hyperalgesia/allodynia caused by L5 spinal nerve cutting (L5SNC) is strongly suppressed by i.p. administration of CSE inhibitors and by i.pl. administration of T-channel blockers or genetic silencing of $Ca_v3.2$ T-channels (Takahashi et al. 2010). Given that the protein levels of $Ca_v3.2$ T-channels, but not CSE, are dramatically upregulated in the DRG of rats with L5SNC (Takahashi et al. 2010), our data suggest that the upregulated $Ca_v3.2$ T-channels in sensory neurons after spinal nerve injury are exposed to positive regulation by CSE-derived endogenous H_2S , contributing to the maintenance of neuropathic pain. H_2S is also considered to be involved in diabetes-associated peripheral neuropathy, since systemic administration of inhibitors of CSE or CBS and silencing of $Ca_v3.2$ T-channels in DRG reverse mechanical allodynia/hyperalgesia in streptozotocin (STZ)-induced diabetic neuropathy rats (Todorovic et al. 2001; Velasco-Xolalpa et al. 2013). It is also to be noted that T-channels, particularly of the $Ca_v3.2$ isoform, are upregulated and involved in the hyperalgesia/allodynia in the laboratory animals with the diabetic neuropathy (Messinger et al. 2009). Upregulation of T-channel current density has also been shown in DRG cells isolated from diabetic Bio-Bred/Worcester (BB/W) rats and from rats with STZ-induced diabetic neuropathy (Todorovic and Jevtovic-Todorovic 2011). Thus, the upregulation of $Ca_v3.2$ T-channels in the sensory neurons appears to cause diabetes-induced hyperexcitability of the nociceptors, contributing to the pathogenesis of diabetic neuropathy.

Many anticancer agents, such as vincristine, paclitaxel, and oxaliplatin, often cause painful peripheral neuropathies that limit their use in cancer therapy. It has been demonstrated that ethosuximide, a T-channel blocker, reverses paclitaxel- and vincristine-induced painful peripheral neuropathy in laboratory animals (Flatters and Bennett 2004). We have reported that paclitaxel-induced neuropathic hyperalgesia is reversed by i.pl. administration of NNC 55-0396 or mibefradil, systemic administration of PPG or BCA, CSE inhibitors, and genetic silencing of $Ca_v3.2$. Unlike L5SNC rats, neither $Ca_v3.2$ nor CSE are upregulated at protein levels in the DRG, spinal cord, or peripheral tissues including the hindpaws, whereas H_2S content in hindpaw tissues is elevated (Okubo et al. 2011). Together, $Ca_v3.2$, a target for H_2S formed by CSE, is a key molecule in the maintenance of distinct types of neuropathic pain.

6 Conclusion

H_2S plays important roles in processing of somatic, visceral, and neuropathic pain signals by targeting $Ca_v3.2$ T-channels and TRPA1. In this context, H_2S -generating enzymes, $Ca_v3.2$, and TRPA1 could be molecular targets for treatment of inflammatory and neuropathic pain including visceral pain. However, endogenous H_2S plays important protective roles in the cardiovascular and gastrointestinal systems, CNS, and so on. Therefore, we must consider the risk of side effects of CSE and

CBS inhibitors, especially in the case of their long-term use. Thus, blockade of Ca_v3.2 T-channels and/or TRPA1 would have greater advantage than inhibition of H₂S formation. In particular, the development of selective Ca_v3.2 inhibitors should impact on improvement of therapeutic strategy for neuropathic pain, although no isoform selective T-channel inhibitors are available at present.

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Part VI

H₂S, Angiogenesis and Cancer

Hydrogen Sulfide and Cancer

Mark R. Hellmich and Csaba Szabo

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Abstract

Recent studies revealed increased expression of various hydrogen sulfide (H₂S)-producing enzymes in cancer cells of various tissue types, and new roles of H₂S in the pathophysiology of cancer have emerged. This is particularly evident in cancers of the colon and ovaries, where the malignant cells both overexpress cystathionine- β -synthase (CBS) and produce increased amounts of H₂S, which enhances tumor growth and spread by (a) stimulating cellular bioenergetics, (b) activating proliferative, migratory, and invasive signaling pathways, and (c) enhancing tumor angiogenesis. Importantly, in preclinical models of these cancers, either pharmacological inhibition or genetic silencing of CBS was shown to be sufficient to suppress cancer cell bioenergetics in vitro, inhibit tumor growth and metastasis in vivo, and enhance the antitumor efficacy of frontline chemotherapeutic agents, providing a strong rationale for the

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development of CBS-targeted inhibitors as anticancer therapies. However, the observation that inhibition of H₂S biosynthesis exerts anticancer effects is contradicted by other studies showing that increasing H₂S with exogenous donors also exerts antitumor actions. Herein, we present a brief review of the scientific literature documenting the function of H₂S, H₂S donors, and transsulfuration enzymes in cancers from various tissue types, and propose that the paradoxical actions of H₂S can be resolved by considering the bell-shaped pharmacology of H₂S, whereby lower (endogenous) H₂S production tends to promote, while higher (generated from exogenously added H₂S donors) tends to inhibit cancer cell proliferation. Finally, we suggest areas for future investigations to expand our knowledge of this nascent field.

Keywords

Cancer • Hydrogen sulfide • Mitochondria • Bioenergetics • Angiogenesis • Proliferation • Signaling

1 Biological Roles of H₂S: A Brief Overview

Work over the last decade recognized the importance of endogenously produced H₂S in a variety of biological functions in the nervous, cardiovascular, and immune system, in health and disease (reviewed in: Szabo 2007; Kimura 2011; Whiteman et al. 2011; Wang 2012). The three principal enzymes involved in the physiological production of H₂S are CBS, CSE, and 3-mercaptopyruvate sulfurtransferase (3-MST). For the purposes of the current article, we focus on the biological effects of H₂S that are relevant for cancer biology; these include the regulation of vascular function (physiologically: vasorelaxation and stimulation of angiogenesis) (reviewed in: Wang et al. 2010; Szabo and Papapetropoulos 2011), regulation of cellular bioenergetics (physiologically: stimulation of mitochondrial electron transport and maintenance of cellular energetics) (reviewed in: Szabo et al. 2014; Módis et al. 2014a), and the regulation of intracellular signaling and cell death (physiologically, acting as a direct and indirect antioxidant and inhibiting oxidative damage and cell death in response to diverse stimuli) (reviewed in: Wang 2012; Kolluru et al. 2013). Appreciating the complexities of H₂S biology requires familiarity with its pharmacology, including the fact that it is a labile, diffusible gas, and that it has a bell-shaped (or biphasic) dose–response curve, whereby lower concentrations (or lower rates of production) can exert markedly different (often, opposing) effects compared to the effects of H₂S seen at higher concentrations (or higher rates of production) (reviewed in: Szabo et al. 2014).

2 Role of Endogenous H₂S Production in Colon Cancer

In 2013 we have compared human colon cancer specimens with patient-matched normal mucosa tissue and discovered that there is a selective upregulation of CBS in the cancer tissue, while the non-cancerous peritumor tissue has low CBS expression levels. The expression of the other two H₂S-producing enzymes, CSE and 3-MST, did not show an upregulation in the tumor tissue. When subsequently checking several colon adenocarcinoma-derived cell lines (HCT-116, HT-29, LoVo), we have also observed a selective upregulation of CBS, as compared to the non-malignant colonic epithelial cell line NCM356. We have also conducted cell fractionation studies to test the localization of CBS in colon cancer cells. While CBS is traditionally viewed as a cytosolic enzyme, it can also be translocated to the mitochondria (Teng et al. 2013; Szabo et al. 2014). Our results revealed that CBS in the HCT116 cancer cells is present both in the cytosol and in the mitochondria. As expected, homogenates of the patient-derived colon tumor specimens as well as homogenates of the colon cancer-derived cell lines showed increased rates of H₂S production, and this response was inhibited by the prototypical CBS inhibitor compound aminooxyacetic acid (AOAA) (Szabo et al. 2013).

Next, we have studied the functional role of CBS-derived H₂S in the control of colon cancer cell proliferation, migration, and invasion *in vitro*, by a combination of genetic (shRNA-mediated stable silencing of CBS in HCT116 cells or adenoviral overexpression of CBS in NCM356 cells) and pharmacologic (CBS inhibition by AOAA) approaches. Genetic silencing or pharmacological inhibition of CBS suppressed HCT116 cell proliferation, migration, and invasion (Szabo et al. 2013). We have subsequently also used *S*-adenosyl-L-methionine (SAM), an allosteric activator of CBS; this compound, at low concentrations, increased HCT116 cell proliferation (Módís et al. 2014b).

Part of the proliferative and pro-migratory effects of CBS-derived H₂S are likely to be due to stimulation of Akt/PI3K signaling, as prior studies have demonstrated that exogenous H₂S donors stimulate HCT116 cell migration via activation of these pathways (Szabo and Hellmich 2013). In addition, part of the effect of CBS-derived H₂S is due to mitochondrial, bioenergetic stimulatory effects. We have observed that both the silencing of CBS and the inhibition of CBS with AOAA suppressed HCT116 bioenergetic functions (including basal electron transport and a bioenergetic parameter known as “respiratory reserve capacity,” which is quantified as the increase in mitochondrial oxygen consumption in response to a mitochondrial uncoupling agent) (Szabo et al. 2013). CBS inhibition not only suppressed mitochondrial function but also glycolytic function in HCT116 cells (Szabo et al. 2013), an effect that may be attributable to the known stimulatory role of H₂S on the activity of GAPDH (Mustafa et al. 2009), an essential enzyme in the glycolytic pathway. Similar to the effects of the allosteric CBS activator SAM on proliferation, SAM, at low concentrations, increased HCT116 cell bioenergetic functions (Módís et al. 2014b).

Subsequent studies in nude mice bearing xenografts of either HCT116 cells or patient-derived tumor tissue (PDX) extended the findings into *in vivo* models.

Silencing of CBS expression and/or pharmacological inhibition of CBS with AOAA significantly reduced the growth rate of the tumor xenografts. While we conclude that part of the effects of CBS inhibition seen *in vivo* are likely related to intratumoral mechanisms (i.e. inhibition of cancer cell metabolism and signaling), part of the effect may also involve paracrine mechanisms in the tumor microenvironment, because CBS silencing or CBS inhibition suppressed the density and complexity of CD31-positive blood vessels within the tumor tissue (indicative of reduced tumor angiogenesis). Furthermore, in line with the role of H₂S as a local vasodilator, direct injection of AOAA into the tumor parenchyma reduced peritumor blood flow (Szabo et al. 2013). In addition to reducing primary tumor growth, inhibition of CBS with AOAA decreased the metastatic spread of HCT116 cell from the cecum to the liver (i.e., decrease the number of metastatic lesions per area) in an orthotopic xenograft model in nude mice, and AOAA synergizes with the anti-metastasis effects of oxaliplatin in the same model (Bohanon et al. 2014).

In HCT116 cells, silencing or pharmacological inhibition of CSE did not exert any effects on HCT116 proliferation, migration, or tumor growth *in vitro* or *in vivo* (Szabo et al. 2013). In contrast, in another human colon cancer cell line (SW480), high expression levels of CSE were observed; these levels were further increased by activation of the Wnt pathway in these cells. Moreover, pharmacological inhibition of CSE (with propargylglycine [PAG]) or genetic silencing of CSE attenuated cell proliferation *in vitro*. In addition, SW480 cells with CSE silencing tended to reduce tumor growth (significant reduction in tumor volume, but not in tumor weight) when injected into tumor-bearing nude mice (Fan et al. 2014).

The above findings are consistent with the conclusion that increased H₂S production (from CBS, but in other cell lines also from CSE) plays an essential role in colon cancer cell proliferation.

3 Role of Endogenous H₂S Production in Ovarian Cancer

Similar to colon cancer cells, CBS has been found to be overexpressed in primary epithelial ovarian cancer tissues, as well as in multiple ovarian cancer cell lines. When examining a collection of more than 200 patients' tissue microarrays constructed from primary epithelial ovarian cancers, Bhattacharyya and colleagues we found high expression of CBS in primary ovarian tumors, particularly in serous carcinoma, the most common histologic variant. Tumors that had serous histology and higher-grade cancers tended to contain higher levels of CBS. CBS expression was already strong in most of the early-stage (FIGO stages I and II) ovarian cancers studied (Bhattacharyya et al. 2013). In additional studies, quantitative RT-PCR and immunoblotting were used to compare the expression of CBS mRNA and protein levels in a variety of ovarian cancer cell lines, when compared to a control, non-malignant ovarian surface epithelial cell line (OSE). Most ovarian cancer cell lines studied showed high CBS expression (both at protein and mRNA level). CSE was not found to be overexpressed in ovarian cancers, but it was found in the normal ovarian epithelial cell line. Similar to the colon cancer study (see above),

CBS exhibited significant mitochondrial localization in A2780 cells (Bhattacharyya et al. 2013).

Next, Bhattacharyya and colleagues studied the functional role of CBS-derived H₂S in the control of ovarian cancer cell proliferation, migration, and invasion *in vitro*, by a combination of genetic (siRNA-mediated stable silencing of CBS in A2780, A2780/CP-70, OV202, and SKOV3 cells) and pharmacologic (CBS inhibition by AOAA) approaches. Downregulation or inhibition of CBS was found to inhibit cell proliferation *in vitro*, and AOAA treatment (especially at higher concentrations) also reduced cell viability. When studying the intracellular mechanisms responsible for these actions, Bhattacharyya and colleagues found that downregulation or inhibition of CBS reduces the intracellular content of the key antioxidant glutathione (GSH + GSSG) and triggers apoptotic cascades. This latter effect may well be the consequence of the intracellular antioxidant depletion after CBS inhibition/silencing. Another important consequence of the CBS silencing or CBS inhibition was an increase in cellular reactive oxygen species levels; this effect may well be secondary to antioxidant depletion (see above) or it may also be related to changes in mitochondrial function (see below). Finally, silencing CBS in A2780 cells also affected intracellular signaling pathways: Bhattacharyya and colleagues found that CBS silencing increases the expression of p53, while the expression of the RelA/p65 subunit of NF- κ B was decreased (Bhattacharyya et al. 2013).

Similar to the study in colon cancer cells, H₂S in ovarian cancer cells supports mitochondrial function and cellular bioenergetics. Bhattacharyya and colleagues found that silencing of CBS reduced mitochondrial oxygen consumption, and similar effects were seen when ovarian cancer cell lines were treated with the CBS inhibitor AOAA. Additional consequences of CBS silencing and/or CBS inhibition were (a) an increase in mitochondrial ROS production, (b) a decrease in intracellular NAD/NADH ratio, (c) a reduction in ATP synthesis, and an increase in ADP/ATP ratio (Bhattacharyya et al. 2013).

Subsequent studies in nude mice transplanted with A2780/CP-20 xenografts extended the findings into *in vivo* models. Silencing of CBS resulted in a significant approximately 40 % reduction in tumor weight and an even more marked (approximately 70 %) decrease in the number of tumor nodules. The reduction by CBS silencing of the proliferative capacity of the cancer cells was confirmed with Ki-67 staining. In addition (and similar to the colon cancer study discussed above), CBS silencing resulted in an inhibition of peritumor angiogenesis, as evidenced by a reduction of CD31 staining (Bhattacharyya et al. 2013).

Last, but not least, Bhattacharyya and colleagues also demonstrated that inhibition of CBS sensitizes the cancer cells to concomitant chemotherapy *in vitro* and *in vivo*. *In vitro*, CBS silencing of A2780 cells shifted the IC₅₀ of cisplatin from 13.1 to 7.9 mM. *In vivo*, the combination therapy of CBS siRNA and cisplatin produced a dramatic reduction in tumor weight and number of tumor nodules (both effects approximately 80–90 %), when compared to the group that received cisplatin alone (Bhattacharyya et al. 2013).

4 Lack of Functional Role of Endogenous H₂S Production in Melanoma

In contrast to colon cancer cells and ovarian cancer cells, Panza and colleagues found that CBS is not overexpressed in human nevi (compound, junctional, or dysplastic forms). Likewise, in samples of human primitive melanoma, CBS and 3-MST displayed a variable, but always very low level of expression. Furthermore, four distinct human melanoma cell lines (A375, Sk-Mel-5, Sk-Mel-28, and PES 43), when compared in normal human epidermal melanocytes (NHEM), showed no increase in CBS expression. In contrast, another H₂S-producing enzyme, CSE, exhibited significantly higher expression in all of the nevi, melanoma samples, and melanoma cell lines studied (Panza et al. 2015).

In order to evaluate whether the various H₂S-producing enzymes had a functional effect on the proliferation of human melanoma cell line, A375 cells were transiently transfected with siRNAs for either CBS, CSE, or 3-MST. Silencing of these enzymes did not affect melanoma cell proliferation (Panza et al. 2015). These data indicate that in melanoma (as opposed to colon cancer and ovarian cancer), the expression of H₂S-producing enzymes does not have an endogenous stimulatory role on cell proliferation.

5 CBS Silencing Accelerates the Development of Glioma

To assess the role of CBS in glioma tumorigenesis, Takano and colleagues established a subclone of U87-MG glioma cells with stable silencing of CBS. When subcutaneously or orthotopically injected into the flank of SCID mice, the subclone with CBS silencing exhibited a shorter latency period for tumor growth, as well as a more pronounced overall tumor growth rate. In addition to greater tumor volume, the CBS-silenced tumors exhibited increased depth of invasion, vascular density, cell proliferation, and more apoptosis (Takano et al. 2014). These findings unveil a role of CBS in glioma that is markedly different from its role in colon cancer or ovarian cancer. As far as the mechanisms responsible for the observed effects, *in vitro* studies by Takano and colleagues showed that glioma cells respond to CBS silencing with increased VEGF and ANGPTL4 levels, higher HIF2 α expression, and increased anchorage-independent cell growth (Takano et al. 2014).

6 Changes in H₂S-Producing Enzymes in Other Forms of Cancer

A limited number of studies are available that detect significant changes in various H₂S-producing enzymes in other forms of cancer. Perhaps the most detailed study was conducted by Guo and colleagues in human prostatic tissues and prostate cancer tissues. In normal prostatic tissue, CBS and CSE were both detected in the prostatic epithelium, while the periacinar stroma cells contained CSE, but not CBS.

LNCaP (an androgen-dependent prostate cancer cell line) exhibited marked CBS and CSE expression; this is in marked contrast with the low expression of CBS and CSE in RWPE-1 (a normal prostatic peripheral zone epithelial cell line). A lesser degree of CBS and CSE expression was seen in several other prostatic cancer lines. Both CBS and CSE were identified primarily cytoplasmic. Guo and colleagues did not investigate the effect of silencing or pharmacological modulation of CBS or CSE on the proliferation rate of LNCaP cells. However, they have tested dihydrotestosterone (DHT) on the expression of CBS and CSE and found that it increased the expression of both CBS and CSE in LNCaP cells (Guo et al. 2012). Other cancer cell types where the expression of CBS has been demonstrated include myeloma (De Vos et al. 2002), biliary track carcinoma (Hansel et al. 2003), and a significant portion of the cells contained in the NCI60 collection, with most consistent/highest increases seen in most breast cancers and most renal cancers present in the collection (Zhang et al. 2005). None of these studies have investigated the functional effect of modulation of H₂S synthesis in these cancer cell lines.

7 H₂S Donors and Cancer

There is an apparent paradox in the literature, since many reports (discussed in the previous sections) show that inhibition of H₂S biosynthesis exerts anticancer effects, while many studies show that H₂S donors of various types exert anticancer actions in vitro and in vivo (overviewed in: Hellmich et al. 2015). In addition, treatment of the recipients with high doses of the CBS/CSE substrate L-cysteine has been shown to increase the growth of melanoma (Panza et al. 2015). This paradox can be resolved by considering the bell-shaped pharmacology of H₂S, whereby lower (endogenous) H₂S production tends to promote, while higher (generated from exogenously added H₂S donors) tends to inhibit cancer cell proliferation (see for review: Hellmich et al. 2015).

8 Conclusions and Future Directions

Although limited in terms of quantity and mechanistic detail, there is a reasonable body of evidence suggesting that endogenous H₂S production is important for the growth and proliferation of at least two types of cancer: colon cancer and ovarian cancer. The data show that melanoma does not rely on endogenous H₂S production for its growth and proliferation, while in glioma, CBS silencing, in fact, accelerates tumor proliferation. It is reasonable to hypothesize that cancers from different organs and tissues will utilize different H₂S-associated pathways to regulate cell proliferation and survival mechanisms in a tumor cell-type dependent manner. The field of H₂S in cancer is a young one, where (as overviewed in Hellmich et al. 2015) additional work is necessary in many areas, including (a) studies to further delineate the mechanism of upregulation of H₂S-producing enzymes in cancer cells; (b) studies aimed to “place” the H₂S-related energetic mechanisms into the overall

scheme of tumor cell bioenergetics; (c) additional work on clinical samples, including biomarkers of H₂S production in patients with cancer; (d) studies aimed at “connecting” the H₂S pathway in cancer cells to various known pathways of cell proliferation and death/survival signaling; (e) further studies focusing on the interaction of tumor cell-derived H₂S with its microenvironment including cancer stem cells; (f) investigations not only into changes in the expression/activity of H₂S-producing enzymes, but also enzymes involved in its metabolism and degradation; (g) studies exploring potential interactions of the H₂S system with other gaseous mediators in cancer, such as nitric oxide and carbon monoxide; and (h) pharmacological and drug discovery studies into improved CBS inhibitors for the therapy of cancer, in forms of cancer where the preclinical data warrant such expansion.

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Role of H₂S Donors in Cancer Biology

Zheng-Wei Lee and Lih-Wen Deng

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Abstract

Hydrogen sulfide (H₂S) donors including organosulfur compounds (OSC), inorganic sulfide salts, and synthetic compounds are useful tools in studies to elucidate the effects of H₂S in cancer biology. Studies using such donors have shown the ability of H₂S to suppress tumor growth both in vitro and in vivo, with some of them suggesting the selectivity of its cytotoxic effects to cancer cells. In addition to promoting cancer cell death, H₂S donors were also found to inhibit cancer angiogenesis and metastasis. The underlying mechanisms for the anti-cancer activities of H₂S involve (1) cell signaling pathways, such as MAPK and STAT; (2) cell cycle regulation; (3) microRNAs regulation; and (4) cancer metabolism and pH regulation. Altogether, compiling evidences have demonstrated the great potential of using H₂S donors as anticancer agents. Nevertheless, the application and development of H₂S for therapy are still facing

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challenges as identification of molecular targets of H₂S awaits further investigation.

Keywords

Hydrogen sulfide donor • Organosulfur • GYY4137 • HS-NSAID • Cancer • Cell viability • Metastasis

Regarded as a hazardous gas with a pungent rotten egg smell, hydrogen sulfide (H₂S) has long been known to possess inhibitory effects on cellular mitochondrial activity and is toxic to central nervous and respiratory systems. Nevertheless, this gas molecule is found to be synthesized endogenously at least via three enzymatic pathways (cystathionine β-synthase, CBS; cystathionine γ-lyase, CSE; and 3-mercaptopyruvate sulfurtransferase, 3-MPST) in various mammalian tissues. Since then, there has been an upsurge of interest to study the role of H₂S in normal physiology and the pathophysiology of disease. H₂S has been reported to regulate various physiological processes including cardiovascular and neuronal functions as well as inflammatory responses. Yet, the roles of H₂S in cancer are less explored. This chapter gives an overview of the roles of H₂S in cancer biology, aiming to provide insights into utilizing H₂S donors as an anticancer agent.

1 Classification of H₂S Donors

The biological functions of H₂S have been mostly discovered through studies that made use of a variety of H₂S donors, either naturally existing or synthetically made. Classification of numerous H₂S donors and their respective bioactivities have been summarized in several reviews (Gu and Zhu 2011; Kashfi and Olson 2013; Song et al. 2014; Zhao et al. 2014b). Briefly, naturally occurring organosulfur compounds (OSCs) have been widely studied for their anticancer activities. Similarly, synthetic compounds such as anethole trithione (ADT-OH) derivatives and those that contain active P–S bonds, including Lawesson’s reagent and a slow-release H₂S donor, GYY4137, have also been used in cancer studies (Table 1). This chapter will describe in depth the bioactivities of these H₂S donors, particularly their effects on cancer cell viability as well as cancer angiogenesis and metastasis (Table 2). The underlying mechanisms and the associated signaling pathways responsible for the actions of H₂S in cancer will be summarized (Fig. 1). Lastly, some perspectives and future challenges pertaining to employment of H₂S as an anticancer agent will be discussed.

Table 1 Major classes of H₂S donors

Category	Example of donor(s)
Organosulfur compounds (OSCs)	Allicin
	Ajoene
	Allyl sulfides
	Diallyl sulfide (DAS), diallyl disulfide (DADS), diallyl trisulfide (DATS)
	Isothiocyanate
	Erucin, iberin, sulforaphane
Inorganic	Sulfide salts
	Calcium sulfide (CaS)
	Sodium hydrosulfide (NaHS)
	Sodium sulfide (Na ₂ S)
Synthetic	Anethole trithione (ADT-OH) derivatives
	ACS 5, ACS 48, ACS50, etc.
	HS-NSAIDs: HS-sulindac, HS-ibuprofen, HS-naproxen, HS-aspirin
	Cysteine-activated
	<i>N</i> -(Benzoylthio)benzamide
	Cysteine analogs
	<i>S</i> -propyl cysteine, <i>S</i> -allyl cysteine, <i>N</i> -acetyl cysteine, <i>S</i> -propargyl cysteine
	Lawesson's reagent
(<i>p</i> -Methoxyphenyl) morpholino-phosphinodithioic acid (GYY4137)	
	AP39 (mitochondria targeted)

Table 2 Effects of H₂S donors on cancer cell viability

Model	Results	Signaling/mechanism(s)	References
Organosulfurs			
<i>Aged garlic extracts (AGE)</i>			
Human patients with colorectal adenomas	Suppressed size and number of colon adenomas after 12 months of 2.4 mL/day treatment		Tanaka et al. (2006)
<i>S-allylmercapto-cysteine</i>			
Prostate cancer (PC3) orthotopic tumor mice	Inhibited primary tumor growth, reduced lung and adrenal metastasis	Upregulated E-cadherin adhesion molecule	Howard et al. (2007)

(continued)

Table 2 (continued)

Model	Results	Signaling/mechanism(s)	References
<i>Garlic oil</i>			
Pancreatic cancer (AsPC-1, MiaPaCa-2, PANC-1)	Inhibited cell proliferation, induced cell cycle arrest, and apoptosis		Lan et al. (2013)
<i>Fresh garlic extracts</i>			
Breast cancer (MCF7)	Altered cellular morphology, decreased cell growth	Deregulated E-cadherin, cytokeratin 8/18, and beta-catenin; downregulated cyclin D1, decreased ERK1 phosphorylation, and increased eIF2-alpha phosphorylation	Modem et al. (2012)
<i>Ajoene derivatives (p-methoxybenzyl-substituted end group)</i>			
Esophageal cancer (WHCO1)	Inhibited cell growth, induced G2/M cell cycle arrest	Induced apoptosis by caspase 3 activation	Kaschula et al. (2012)
<i>Allicin</i>			
Gastric cancer (SGC-7901)	Inhibited cell proliferation, induced G2/M cell cycle arrest, and apoptosis		Tao et al. (2014)
Colon cancer (HCT116, LS174T, HT29, Caco2)	Decreased cell growth, induced hypodiploid DNA, and apoptosis	Decreased Bcl-2 and increased Bax expression, increased cytochrome c release from mitochondria, induced translocation of Nrf2 into nucleus	Bat-Chen et al. (2010)
L5178Y lymphoma-bearing mice	Inhibited cell proliferation and induced apoptosis, increased survival rate of tumor-bearing mice	Increased caspase-3 activity	Padilla-Camberos et al. (2010)
<i>Tetra-arsenic tetra-sulfide (As4S4)</i>			
Retinoid acid (RA)-resistant human acute promyelocytic leukemic NB4-R1	Induced apoptosis	Decreased SET gene expression led to increased protein phosphatase 2 (PP2A) and decreased PML-RARA level	Liu et al. (2014)

(continued)

Table 2 (continued)

Model	Results	Signaling/mechanism(s)	References
<i>Isothiocyanate erucin (ER)</i>			
Prostate cancer (PC3)	ER decreased cell growth and proliferation	ER increased p21 expression and ERK1/2 phosphorylation	Melchini et al. (2013)
<i>Organosulfur compounds (OSCs)</i>			
Breast carcinoma (MCF7, Vcr-R)	Induced G2/M cell cycle arrest	Diallyltetrasulfide and dipropyltetrasulfide emerged as irreversible inhibitors of Cdc25 isoforms A and C	Viry et al. (2011)
<i>Garlic-derived organosulfurs</i>			
Colon cancer (HCT116)	Diallyltrisulfide and diallyltetrasulfide treatment reduced cell viability, induced cell cycle arrest, and apoptosis	Induction of apoptosis dependent on redox state of the cell, dispensable of p53; growth arrest and apoptosis associated with reduction of the level of cdc25c	Busch et al. (2010)
<i>Diallyl sulfide (DAS)</i>			
Colon cancer (Colo 320)	Induced G2/M cell cycle arrest and apoptosis	Increased ROS, upregulated NF-κB expression, promoted caspase-3 expression, suppressed ERK-2 activity	Sriram et al. (2008)
Human cervical cancer (HeLa)	Induced G0/G1 cell cycle arrest and sub-G1 apoptosis phase; DNA damage and fragmentation	Induced cytochrome c release and increased expression of pro-caspase-3 and pro-caspase-9	Wu et al. (2011)
<i>Diallyl disulfide (DADS)</i>			
Colon cancer (HCT116)	Induced G2/M cell cycle arrest and apoptosis	Increased ROS, increased expression level of cyclin B1	Song et al. (2009)
Neuroblastoma (SH-SY5Y)	Induced morphological changes and depolymerization of microtubules	Induced cytoskeleton oxidation, microfilaments reduction, and Tau dephosphorylation	Aquilano et al. (2010)

(continued)

Table 2 (continued)

Model	Results	Signaling/mechanism(s)	References
Colon cancer (HCT116)	Induced cell growth inhibition and G2/M arrest	Increased cyclin B1 expression but not affected by p53 silencing	Jo et al. (2008)
Prostate cancer (LNCaP)	Induced apoptosis and nuclear fragmentation		Gunadharini et al. (2006)
Lung carcinoma (A549)	DADS-induced G2/M cell cycle arrest and apoptosis	Increased ROS	Wu et al. (2005)
Colon cancer (HCT116, HCT15) and prostate cancer (PC3)	Inhibited cell proliferation	Induced p53 and nonsteroidal anti-inflammatory drug (NSAID)-activated gene (NAG1)	Bottone et al. (2002)
Gastric cancer (MGC-803)	Antitumor effect in vitro and in vivo	Upregulated miR-200b and miR-22 that target Wnt-1	Tang et al. (2013)
Gastric cancer (BGC823)	Inhibited cell growth, induced G2/M cell cycle arrest	Decreased Cdc25c and cyclin B1, increased p21, GADD45-alpha and p53; decreased Chk1 phosphorylation	Bo et al. (2014)
Leukemia (HL60)	Inhibited cell growth, induced apoptosis	Activated p38, reduced ERK activity	Tan et al. (2008)
Gastric cancer (MGC-803)	Inhibited proliferation, induced G2/M cell cycle arrest	Induced differentiation and decreased ERK1/2 phosphorylation	Ling et al. (2006)
Gastric cancer (MGC-803)	Inhibited cell growth, induced G2/M cell cycle arrest	Increased p38 phosphorylation, decreased Cdc25c expression	Yuan et al. (2004)
Colon cancer (HCT15)	Induced G2/M cell cycle arrest	Increased ERK phosphorylation	Knowles and Milner (2003)
<i>Diallyl trisulfide (DATS)</i>			
Osteosarcoma (U2OS, SaOS-2, MG-63)	Inhibited proliferation by triggering cell cycle arrest and apoptosis; suppressed wound-healing capacity, invasion and cancer media-induced HUVEC tube formation	Decreased Notch-1 expression; increased tumor-suppressive miR-34a, miR-143, miR-145, and miR-200b/c	Li et al. (2013)

(continued)

Table 2 (continued)

Model	Results	Signaling/mechanism(s)	References
Benzo(a)pyrene (BaP)-induced precancerous MCF10A	Inhibited cell proliferation, decreased G2/M cell cycle transition	Reduced BaP-induced peroxide formation and DNA strand breaks, thus preventing BaP-induced carcinogenesis	Nkrumah-Elie et al. (2012)
Basal cell carcinoma (BCC) and melanoma (A375)	Induced G2/M cell cycle arrest and apoptosis	Increased ROS, induced cytosolic Ca ²⁺ mobilization, decreased mitochondrial membrane potential	Wang et al. (2010)
Prostate cancer (LNCaP, C4-2, TRAMP-C1)	Inhibited cell proliferation	Decreased androgen receptor (AR) protein level and AR promoter activity	Stan and Singh (2009)
Transgenic adenocarcinoma of mouse prostate (TRAMP)	Inhibited progression to invasive carcinoma and pulmonary metastasis	Induced cyclin B1 and securin protein expression, suppressed protein expression of neuroendocrine marker synaptophysin	Singh et al. (2008)
Prostate cancer (PC3)	Oral gavage DATS retarded growth of PC3 xenografts	Induction of proapoptotic Bax and Bak, decreased VEGFR2 protein level	Xiao et al. (2006)
Gastric cancer (MGC803, SGC7901)	Decreased G0/G1 and increased G2/M phase cells	Upregulated p21 gene expression level	Ha et al. (2005)
Colon cancer (HCT15 and DLD-1)	Decreased G0/G1 and increased G2/M phase cells	Induced disruption of tubulin polymerization	Hosono et al. (2005)
Prostate cancer (PC3 and DU145)	Induced G2/M cell cycle arrest	Inhibited cyclin-dependent kinase 1 and hyperphosphorylation of Cdc25c	Herman-Antosiewicz and Singh (2005)
Inorganic and synthetic donors			
<i>NaHS</i>			
Prostate cancer (LNCaP)	Decreased proliferation of LNCaP and antiandrogen-resistant LNCaP-B cells	H ₂ S sulfhydrated androgen receptor, inhibiting AR binding to DNA and transactivation of its downstream genes	Zhao et al. (2014a)

(continued)

Table 2 (continued)

Model	Results	Signaling/mechanism(s)	References
TGF- β 1-treated MCF7	Decreased proliferation, G0/G1 phase cell cycle arrest, and apoptosis	Decreased TGF- β 1-induced p38 phosphorylation	Lv et al. (2014)
Colon epithelial (YAMC) and colon cancer (HT-29, SW1116, HCT116)	Inhibited proliferation, migration, and G1/S transition; increased LC3B autophagic vacuoles and acidic vesicular organelles	Increased AMPK phosphorylation and decreased mTOR and S6 kinase phosphorylation; induced p21 expression	Wu et al. (2012)
Systemic lupus erythematosus lymphocytes	Inhibited proliferation and S phase distribution of cell cycle	Decreased phytohemagglutinin (PHA)-induced CDK2 expression, AKT (S473) and GSK3b (S9) phosphorylation, increased p27 and p21 expression	Han et al. (2013)
Oral squamous carcinoma (Cal27, GNM, WSU-HN6)	Induced proliferation and accelerated cell cycle progression of synchronized cells	Increased Akt and ERK1/2 phosphorylation	Ma et al. (2014)
Colon cancer (HCT116, SW480)	Promoted cell proliferation, decreased G0/G1 phase, and increased S phase cells	Increased Akt and ERK phosphorylation; inhibited expression of p21, and reduced NO metabolite	Cai et al. (2010)
<i>GY4137</i>			
Hepatocellular carcinoma (HepG2, Bel7402)	Suppressed cell proliferation and tumor growth, inhibited G1/S cell cycle transition	Decreased Stat3 (Y705) activation; altered Bcl-2, cyclin D1, Mcl-1, survivin, VEGF, and HIF-1 α expression; increased cleaved caspase-9, caspase-3, and PARP	Lu et al. (2014)
HeLa, HCT116, HepG2, HL60, MCF7, MV4-11, U2OS	GY4137 but not NaHS decreased cancer cell survival; noncancer cells were less affected; reduced xenograft tumor growth	Induced apoptosis and partial G2/M cell cycle arrest	Lee et al. (2011)

(continued)

Table 2 (continued)

Model	Results	Signaling/mechanism(s)	References
Breast cancer (MCF7), liver cancer (HepG2)	Continual exposure but not single exposure to low concentration of NaHS decreased MCF7 and HepG2 viability; noncancer MCF10A and WI38 were less affected	Induced increased lactate production and impaired pH regulation; caused intracellular acidification	Lee et al. (2014)
<i>S-propargyl-cysteine (SPRC)</i>			
Gastric cancer (SGC-7901)	Decreased cell viability and suppressed proliferation and migration caused G1/S cell cycle arrest; reduced tumor weight and volumes	Increased p53 and Bax expression; increased CSE expression and tumoral CSE activity	Ma et al. (2011)
<i>HS-NSAIDs</i>			
Human colon, breast, pancreatic, prostate, and lung cancer and leukemia	Inhibited cell proliferation, caused G0/G1 cell cycle block	Induction of apoptosis	Chattopadhyay et al. (2012b)
<i>HS-aspirin (HS-ASA)</i>			
Leukemia (Jurkat)	Inhibited proliferation, induced apoptosis, G0/G1 cell cycle arrest	Downregulated B-catenin, cyclin D1, and c-Myc	Chattopadhyay et al. (2013)
Triple negative breast cancer (MDA-MB-231)	Suppressed cell growth, induced G0/G1 cell cycle arrest and apoptosis; reduced xenograft tumor volume and mass	Downregulated NF-κB, reduced thioredoxin reductase activity and increased ROS	Chattopadhyay et al. (2012a)
<i>NOSH-aspirin (NBS-1120)</i>			
Colon cancer (HT29)	Inhibited cell proliferation, induced apoptosis and G0/G1 cell cycle block; xenograft tumor volume reduction of 85 %		Chattopadhyay et al. (2012c)
<i>NOSH-naproxen (AVT-219) and NOSH-sulindac (AVT-18A)</i>			
Colon (HT-29), pancreas adenomatous (BxPC-3), and breast cancer (MCF7) and lymphocytic leukemia (Jurkat)	Inhibited proliferation, G0/G1 or G2/M block in cell cycle		Kodala et al. (2013)

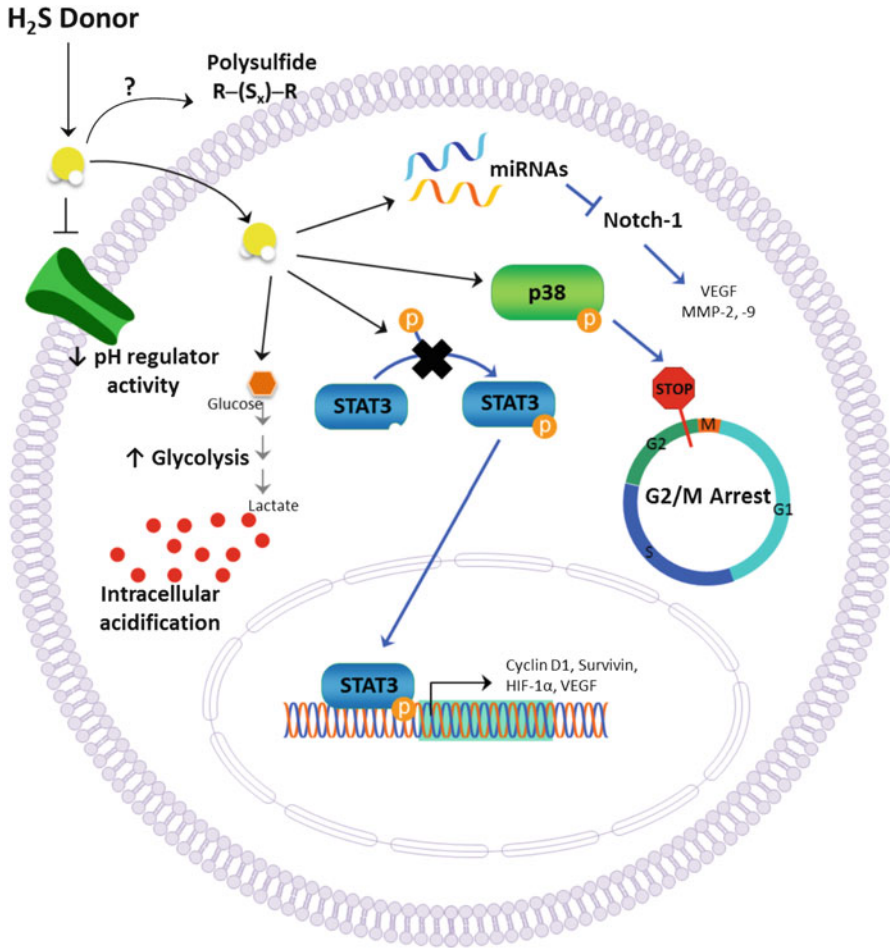


Fig. 1 H₂S-mediated pathways in cancer cells. H₂S potentially decreases STAT activation, increases p38 phosphorylation, induces tumor-suppressive miRNAs, impairs pH regulator activity, and increases glycolysis resulting in intracellular acidification in cancer cells

2 Effects of H₂S Donors on Cancer Cell Viability

2.1 Organosulfur Compounds (OSCs)

OSCs including allicin and ajoenes that are abundantly found in *Allium* vegetables, especially garlic, have been recognized to confer health benefits throughout recorded human history, dating back since ancient Egypt, Chinese, and Hindu civilizations (Block 1985). Consumption of garlic and other *Allium* vegetables is believe to reduce serum cholesterol and blood glucose levels (Sher et al. 2012) and to stimulate immune responses against microbial, viral, and pathogen infection

(Feng et al. 2012; Lau et al. 1991). In the mid-twentieth century, garlic extracts were shown to effectively inhibit growth of sarcoma 180 ascites tumor in CFW Swiss mice (Weisberger and Pensky 1957). Since then, increasing evidences for anticancer effects of OSCs from population-based case–control studies have suggested that greater intake of *Allium* vegetables reduces risks of developing cancer. For instance, a study carried out in Chinese population revealed that those who have the highest garlic intake experienced nearly 60 % reduction in gastric cancer occurrence as compared to those who consumed the least (You et al. 1989). In another double-blind, randomized clinical trial, Tanaka et al. reported that high dose treatment of aged garlic extracts significantly suppressed the progression of colorectal adenomas in patients (Tanaka et al. 2004, 2006). The effects of garlic or garlic oil in inhibiting cancer growth are also evidenced in several other cancer types including androgen-independent prostate cancer, pancreatic cancer, liver cancer, and breast cancer (Howard et al. 2007; Lan et al. 2013; Modem et al. 2012; Zhang et al. 2013).

Garlic is a complex mixture of different OSCs. The major forms of OSCs present in intact garlic are γ -glutamyl-*S*-alk(en)yl-L-cysteines and *S*-alk(en)yl-L-cysteine sulfoxides. Upon exposure to atmospheric water and oxygen, the cysteine sulfoxides rapidly get hydrolyzed and oxidized into several hundreds of thiosulfinates, of which 70–80 % is allicin. However, allicin is highly unstable and susceptible to further oxidation. In a rat liver perfusion model, 90 % of the allicin disappears from the tissue within the first 3 min while less than 1 % remains after 6 min. Decomposition of allicin produces the oil-soluble volatiles such as diallyl sulfides (DAS), diallyl disulfides (DADS), diallyl trisulfides (DATS), methyl allyl sulfides, and ajoene (Block 1985; Fenwick and Hanley 1985). Allicin and these volatile OSCs are the compounds that are responsible for the unpleasant odor of garlic.

Given the complexity of garlic chemistry, the observed biological activities mentioned above might be contributed by either one or collective effects of the OSC mixture. Therefore, to identify and characterize the OSC that is responsible for the anticancer activity, researchers have purified and isolated these individual compounds. Intriguingly, compiling evidences suggest that most of these OSCs possess anticancer activities even when they are singly administered. Allicin significantly inhibits cell proliferation and induces apoptosis in gastric cancer cell line SGC-7901 (Tao et al. 2014). Likewise, allicin effectively decreases cell proliferation of several colon cancer cell lines, including HCT-116, LS174T, HT-29, and Caco-2 (Bat-Chen et al. 2010). The inhibitory effect of allicin on cancer growth has also been demonstrated in an in vivo mouse model. Using L5178Y lymphoma-bearing mice, Padilla-Camberos and colleagues showed that administration of allicin significantly inhibited tumor growth. Additionally, normal splenocytes were able to tolerate with at least two times higher than the doses used, suggesting the low toxicity of allicin (Padilla-Camberos et al. 2010).

As mentioned previously, the half-life of allicin is very short. Therefore, the observed anticancer growth effects are unlikely due to the activity of allicin alone. Allyl sulfides produced from oxidative decomposition of allicin are comparably stable. In a study to evaluate the reactivity of different allyl sulfides in altering erythrocytes oxidative balance, tetrasulfides were found to be the most reactive

species. The reactivity decreased according to the order of tetra- > tri- > disulfides, while monosulfides exhibited no significant effect on the examining parameters (Munday et al. 2003). This suggests that the reactivity of the OSC is directly related to the number of sulfur atom present in the molecule. In fact, most of the studies used DADS or DATS rather than DAS.

DADS treatment alone was reported to be effective in inhibiting cancer cell growth, inducing apoptosis and causing partial G2/M cell cycle arrest in cancers of colon, pancreas, prostate, lung, brain, and stomach (Aquilano et al. 2010; Bo et al. 2014; Bottone et al. 2002; Gunadharini et al. 2006; Song et al. 2009; Wu et al. 2005). In particular, the anticancer proliferation effects were reported to be associated with an increase in reactive oxygen species (ROS) production. For instance, DADS was shown to induce cytoskeleton oxidation and microtubule depolymerization, thence morphological changes in neuroblastoma SH-SY5Y cells (Aquilano et al. 2010). However, these effects were reversed by administration of exogenous precursors of cellular antioxidants, *N*-acetylcysteine (NAC) or glutathione ester.

With an extra sulfur atom, DATS is thought to be more reactive than DADS. Nkrumah-Elie and colleagues reported that DATS effectively inhibited benzo(a) pyrene (BaP)-induced carcinogenesis in noncancer breast epithelial MCF10A (Nkrumah-Elie et al. 2012). BaP is a potent mutagen and carcinogen found in coal tar and cigarettes (Kaiserman and Rickert 1992). The author reported that either co-treatment or pretreatment of DATS on BaP-treated MCF10A significantly reduced cell proliferation. Furthermore, DATS decreased BaP-induced peroxide formation and DNA strand breaks, suggesting that DATS might be effective in preventing BaP-induced genomic instability and carcinogenesis. Concomitantly, anticancer activity of DATS has also been shown in other cancers, including colon, gastric, and prostate cancers (Ha et al. 2005; Herman-Antosiewicz and Singh 2005; Hosono et al. 2005). Not surprisingly, the involvement of ROS in DATS-mediated anticancer effects was evidenced by Busch's study that the apoptosis induced by DATS was able to be reversed by ascorbate or NAC (Busch et al. 2010). Moreover, DATS might selectively target cancer cells, as shown by Wang and colleagues that 25–100 μM of DATS significantly decreased viability of basal cell carcinoma BCC and melanoma A375 cells but had minimal effects on human keratinocytes HaCaT (Wang et al. 2010). Collectively, OSC (at least for DADS and DATS) exerts anticancer activity in cancers of various tissue origins.

Despite mounting studies reporting the anticancer activity of OSC or garlic derivatives, the contribution of H_2S remains elusive. Many other active side groups or moieties in the OSC molecules might play a part in the anticancer effects. By comparing the effects of DAS, DADS, DATS, diproylsulfide, dipropyl disulfide, and allyl mercaptan on the reduction of androgen receptor expression in prostate cancer cells, Stan et al. revealed that a subtle change in the structure and number of sulfur atom of OSC greatly affects their anticancer effects (Stan and Singh 2009). DATS with the three sulfur atoms flanking with allyl chains appeared to be the most effective OSC. The importance of these allyl side chains in determining OSC activities was evident as the allyl side chain produced from the breakage of S–S bond in DATS was able to oxidize the cysteine residue of tubulin protein, resulting

in the disruption of tubulin polymerization thence causing cell cycle arrest in colon cancer cell HCT15 (Hosono et al. 2005). Nevertheless, diallyl polysulfides present in garlic can also react with cysteine or glutathione to produce H₂S in human body, suggesting that H₂S might also be involved in the abovementioned bioactivities (Benavides et al. 2007; Jacob 2006).

2.2 Inorganic and Synthetic H₂S Donors

Inorganic salts of sulfide such as sodium hydrosulfide (NaHS) and sodium sulfide (Na₂S) are the simplest H₂S donors being utilized. In contrast to OSC or garlic derivatives, NaHS or Na₂S dissolves instantaneously in aqueous solution and produces only sulfide species. This simple chemistry allows one to easily correlate the observed bioactivities with H₂S. NaHS was shown to decrease breast cancer MCF7 and prostate cancer LNCaP cell proliferation, coupled with an increase in apoptosis (Lv et al. 2014; Zhao et al. 2014a). In contrast, NaHS stimulated proliferation and accelerated cell cycle progression of synchronized oral squamous carcinoma Cal27, GNM, and WSU-HN6 cell lines (Ma et al. 2014). This ambiguous role of NaHS has even been observed in the same colon cancer cell line, HCT116. Cai et al. reported that 200 μM of NaHS treatment induced HCT116 cell proliferation and promoted S phase progression, while Wu et al. observed that 1 mM NaHS inhibited cell proliferation and migration (Cai et al. 2010; Wu et al. 2012). It appears that cancer cells could tolerate H₂S derived from low micromolar concentrations of NaHS and get stimulated to proliferate faster. However, at a high concentration of NaHS exposure, pro-survival signaling was suppressed and cell death was triggered. In line with this hypothesis, HCT116 exposed to low micromolar concentration of NaHS increases Akt and ERK phosphorylation stimulating cell survival, while millimolar range of NaHS decreases mTOR and S6 kinase activity, leading to autophagy and apoptosis (Cai et al. 2010; Wu et al. 2012). These disparate results suggest that the anti- or pro-cancer effect of NaHS appears to depend on the concentration and duration of exposure.

One concern of utilizing sulfide salts is that it releases massive amounts of the gas almost instantaneously, likely leading the effect of the gas on cells to be acute and toxic. In addition, the amount released by NaHS or Na₂S does not mimic physiological levels. In view of this, researchers have been working on chemically synthesizing H₂S donor drugs with the ability to release H₂S at different rates. Synthetic H₂S donors such as (*p*-methoxyphenyl)morpholino-phosphinodithioic acid (GYY4137) were synthesized in order to achieve a prolonged and consistent H₂S release in culture as compared to the sulfide salts (Li et al. 2008). In a study done by Lee and colleagues, a wide range of cell lines were treated with either GYY4137 or NaHS (400 μM), and their viability was examined after 5 days of incubation (Lee et al. 2011). GYY4137 was able to reduce cell viability in various cancer cell lines including HeLa, HCT116, HepG2, HL60, MCF7, MV4-11, and U2OS but spare noncancer cell lines, IMR90 and WI38. In contrast, NaHS marginally affected the cancer cell viability. The differential effects of GYY4137 and NaHS are speculated to be resulted from their differential H₂S-release manner. As

opposed to NaHS which releases H₂S instantaneously, GYY4137 slowly releases H₂S and sustains over days in culture. This observation proposed that a prolonged and low exposure to H₂S might be the key to selective anticancer activity of H₂S. In a follow-up study, the authors demonstrated that the viability of cancer cells, exposed to low concentration (5–20 μM) of fresh NaHS solution every 2 h over 5 days, was significantly decreased (Lee et al. 2014). On the other hand, noncancer cells tolerated the same H₂S treatment regime. It is also worth mentioning that a high and continual H₂S exposure (>80 μM) killed both cancer and noncancer cells, likely due to the long known inhibitory effects of H₂S on mitochondrial activity (Savolainen 2010).

Besides GYY4137, a group of synthetic H₂S-releasing hybrid compounds was also shown to exhibit anticancer effects. Among these, nonsteroidal anti-inflammatory drugs (NSAIDs) appear to be good candidates. NSAIDs have been recorded to exhibit a certain extent of anticancer activity. Patients undergoing aspirin medication had significantly lower colorectal cancer incidence, corrected for age and gender (Kune et al. 1988). Analysis of data generated from 51 randomized trials also revealed that daily aspirin intake reduces cancer-associated death and incidence (Rothwell et al. 2012). However, long-term medication on NSAIDs may cause significant side effects, mainly gastrointestinal and renal damages (Harirforoosh and Jamali 2009; Sostres et al. 2013). Therefore, owing to the potential cytoprotective effects of H₂S (reviewed in other chapters), new hybrids of HS-NSAIDs were designed (Kashfi 2014).

Four HS-NSAIDs including HS-sulindac, HS-ibuprofen, HS-naproxen, and HS-aspirin (HS-ASA) were used to study their activities in 11 cancer cell lines derived from six tissue origins, including colon (HT29, HCT15, SW480), breast (MCF7, MDA-MB-231, SKBR3), leukemia (Jurkat), pancreas (BxPC3, MIAPaCa-2), prostate (LNCaP), and lung (A549) (Chattopadhyay et al. 2012b). All four HS-NSAIDs are effective in inhibiting the growth of the 11 cancer cell lines, whereas their respective NSAID treatment alone requires 38–1300 times higher concentrations to achieve the same extent of inhibitory effect. Among the four HS-NSAIDs, HS-ASA is the most potent hybrid even though ASA is the least potent NSAID in anticancer cell growth. The author speculates that the conjugation of ASA with HS moiety forms a hybrid which releases H₂S over a long period of time. Taken together, the authors conclude that HS-NSAID, as a H₂S donor, is a potential anticancer agent independent of cancer tissue types.

3 Underlying Mechanisms of H₂S Donors in Suppressing Cancer Cell Growth

Mitogen-Activated Protein Kinase (MAPK) Signaling and Cell Cycle Regulation

MAPK signaling plays a central role in regulating cell proliferation. The specific activation and deactivation of MAPK in response to H₂S donors have been documented. For instance, p38 was found to be activated but ERK activity was

reduced in DADS-induced apoptotic HL60 cells (Tan et al. 2008). Antagonists against p38 or ERK are able to decrease or enhance toxicity and apoptosis induced by DADS, respectively, suggesting direct involvement of these two MAPKs in mediating DADS anticancer activity. Consistently, DADS induces differentiation or G₂/M arrest in gastric cancer MGC803 cell line, and such activity is mediated by an increase in p38 activation and a decrease in phosphorylation of ERK (Ling et al. 2006; Yuan et al. 2004). However, in colon cancer HCT15 model, an elevated level of ERK phosphorylation has been shown to be an early signaling event that led to G₂/M arrest induced by DADS (Knowles and Milner 2003). All these studies converge to the fact that anticancer activity of DADS is p38 and more importantly ERK mediated.

STAT Pathway

Signal transducer and activator of transcription (STAT) pathway is implicated in many pathophysiologicals, including cancer. Specifically, constitutive activation of STAT3 was reported to be involved in maintaining progression of many malignancies and solid tumors (Siveen et al. 2014). Therefore, inhibition of STAT3 pathway in cancer potentially serves as a therapeutic strategy (Santos and Costa-Pereira 2011). It has been reported that GYY4137 suppressed hepatocellular carcinoma tumor growth via inhibition of STAT3 pathway both in vitro and in vivo (Lu et al. 2014). GYY4137 significantly decreased phospho-STAT3 (Y705) level in HepG2 and Bel7402 cells, leading to decreased expression of its downstream genes, including Bcl-2, cyclin D1, Mcl-1, survivin, vascular endothelial growth factor (VEGF), and hypoxia-inducible factor-1 α (HIF-1 α). Altered levels of these molecules thence contributed to the inhibitory effect of GYY4137 on hepatocellular carcinoma progression. Having known that STAT3 is widely activated in various types of cancer, the potential of using GYY4137 to deactivate STAT3 in other cancers is worth exploring.

miRNA Regulation

Several studies have reported that miRNA is involved in the bioactivity of H₂S donors. miR-200b and miR-22 were found to be upregulated in DADS-treated MGC803 gastric cancer cells (Tang et al. 2013). The authors further identified Wnt1 to be a target of these two miRNAs and reported a synergistic effect between miR-200b and miR-22 in the inhibition of gastric cancer growth in vitro and in vivo. Another study showed that DATS treatment on osteosarcoma U2OS, SaOS-2, and MG-63 cell lines increased expression of several tumor-suppressive miRNA such as miR-34a, miR-143, miR-145, and miR-200b/c (Li et al. 2013). Re-expression of miR-34a into the osteosarcoma cells resulted in the attenuation of Notch-1 activity and downregulation of its downstream genes such as VEGF and matrix metalloproteinase (MMP)-2 and matrix metalloproteinase (MMP)-9. Moreover, miR-200b/c was found to be downregulated in osteosarcoma. miR-200b/c is known to suppress cancer epithelial to mesenchymal transition. DATS treatment successfully re-expressed these miRNAs that in turn inhibited osteosarcoma growth. The role of miRNA in modulating anticancer activity of H₂S donors requires more future studies.

Cancer Metabolism and pH Regulation

A recent report by Lee and colleagues showed that a continual and low exposure to H₂S, either by one-time administration of GYY4137 or NaHS replaced every 2 h for over 5 days, significantly increased glycolysis causing overproduction of metabolic acid lactate (Lee et al. 2014). Simultaneously, such low and prolonged exposure of H₂S impaired the ability of cancer cells to regulate their own intracellular pH by inhibiting anion exchanger and sodium/proton exchanger activity. Combinatorial effects of increased glycolysis and defective pH regulation cause accumulation of metabolic acid, resulting in an uncontrolled intracellular acidification leading to cancer cell death (Lee et al. 2014). Neither increased metabolic acid production nor inhibition of pH regulator alone is sufficient to induce cancer cell death. Importantly, these mechanisms do not operate in normal cells, and hence, H₂S has the capability to kill cancer cells but spare normal cells. This study suggests a possibility of developing novel selective anticancer drugs based on the principle of H₂S donation and their effect on a fundamental difference in the energy-producing mechanisms of cancer, vis-a-vis, normal cells.

4 Effects of H₂S Donors on Cancer Angiogenesis and Metastasis

The vasodilatory role of H₂S has been well documented and so has the modulatory effects of different H₂S donors on endothelial cells (reviewed in other chapters). However, the effects of H₂S donors specifically on cancer angiogenesis and metastasis have been poorly studied. Nonetheless, a study by Thejass and Kuttan (2007) provided evidences that DAS exhibited anticancer angiogenesis property. B16F-10 melanoma cells were injected intradermally on the ventral skin of mouse and were treated with DAS for 5 consecutive days. After 9 days of tumor induction, the mice were sacrificed and ventral skin was dissected for quantification of tumor-directed capillaries. The DAS treatment reduced almost 67 % of the number of tumor-directed capillaries as compared to the control. The anticancer angiogenic effect of DAS was also evidenced using ex vivo rat aortic ring assay. Conditioned medium collected from DAS-pre-treated B16F-10 melanoma cells resulted in a more profound reduction of microvessel outgrowth as compared to the non-treated conditioned medium. This may imply that DAS treatment alters the release of cancer-associated angiogenic factors. One important factor identified in the study was VEGF as DAS treatment significantly decreased mRNA expression of VEGF in the melanoma cells.

With respect to cancer metastasis, NaHS treatment significantly reduced migration of transforming growth factor- β 1 (TGF- β 1)-stimulated MCF7 in Boyden chamber assay (Lv et al. 2014). TGF- β 1 potentiates epithelial-mesenchymal transition (EMT), a cellular process that allows cancer cells to acquire higher motility and invasiveness. The ability of NaHS to suppress TGF- β 1-induced cancer migration suggests that H₂S may be an anti-EMT agent. Inhibition of cancer invasion by NaHS was then identified partly due to the downregulation of Snail protein and

reduction of p38 phosphorylation. Besides, OSCs including DAS, DADS, and DATS were shown to inhibit migration and invasion of colon cancer Colo 205 (Lai et al. 2013). The number of OSC-treated cells which migrated into the lower part of the Boyden chamber was nearly threefold lesser than that of non-treated cells. The inhibitory effect was not only concentration dependent, but also enhanced with the number of sulfur atom of OSC. In the same study, the author reported that these OSCs significantly decreased expression level of proteins in MAPK pathway leading to inhibition of MMP-2, MMP-7, and MMP-9 activities. MMPs are proteinases expressed by cancer cells in order to degrade extracellular matrix, thereby facilitating their extravasion and migration to distant sites (Deryugina and Quigley 2006; Itoh and Nagase 2002). Downregulation of MMP by OSC underlies the antimetastatic property of the H₂S donor. In an in vivo transgenic adenocarcinoma of mouse prostate (TRAMP) mice model, DATS administration significantly suppressed incidence and pulmonary metastatic event of poorly differentiated prostate cancer (Singh et al. 2008). Dorsolateral prostate of TRAMP mice given DATS were harvested, and the percentage of the area occupied by poorly differentiated prostate cancer cells was quantitated. 1 or 2 mg treatment of DATS, thrice weekly, lowered the area of poorly differentiated prostate cancer by 34 and 41 %, respectively, as compared to non-treated control. Poorly differentiated cancer resembles high-grade cancer with high metastatic event. The inhibitory activity of DATS on prostate cancer development and metastasis is encouraging and more work is needed to further explore the anti-angiogenic and antimetastatic activities of other H₂S donors using various in vivo and in vitro cancer models.

5 Therapeutic Potential and Challenges of Utilizing H₂S Donors as Anticancer Agents

Organosulfur compounds (OSCs), GYY4137 and HS-NSAID have been shown by many independent groups to suppress the proliferation of various cancer cell lines but exhibit minimal effects on the noncancer cells. The lower toxicity of H₂S donors, as compared to other cytotoxic agents on noncancer cells, is of a great value. Wang's group showed that DATS was tolerated by normal keratinocytes, but it inhibited basal cell carcinoma and melanoma cell growth (Wang et al. 2010). Similarly, allicin spared normal splenocytes at more than double the dosages used to kill lymphoma cells (Padilla-Camberos et al. 2010). The slow-releasing H₂S donor, GYY4137, was also shown to be selective in inhibiting proliferation of various cancer cells lines including colon, liver, breast, and cervical cancers and leukemia (Lee et al. 2011). In vivo studies have demonstrated that these compounds were effective in reducing tumor volume and mass. At whole animal level, although not measured objectively, the authors reported that no detrimental effects on animal physiology were observed after H₂S donor treatment. These evidences clearly show that low toxicity chemotherapy is possibly achieved using H₂S donors. Furthermore, several studies observed increased ROS production upon OSC treatment, and

some clearly showed that ROS was required for the anticancer activity. Therefore, further examinations are needed to evaluate if the increase in ROS can exert detrimental effects to normal physiology.

One of the major challenges in developing H₂S donors as an anticancer agent is the identification of its molecular targets. Being a lipophilic gas, H₂S is cell permeable; therefore, it is difficult to estimate its distribution and to identify the exact site of its actions. Nevertheless, using octanol and hexane organic solvents as membrane bilayer mimics, Cuevasanta et al. demonstrated that H₂S is highly soluble at hydrophobic core and therefore likely to exhibit greater activity within lipid membrane layers (Cuevasanta et al. 2012). Given the dissociation constant of H₂S, about two-third of H₂S exists as HS⁻ at physiological pH. One possible modulation of protein structure and activity by H₂S is via a posttranslational modification, sulphydration, on the cysteine residue (Mustafa et al. 2009). HS⁻ can covalently link with sulfur atom of cysteine amino acid, forming S–SH bond. Another mechanism in which H₂S may modulate protein functions is via breaking of disulfide linkage between two cysteine residues. Tao et al. reported that Cys1045–1024 disulfide bond of VEGFR2 was labile and susceptible to H₂S-mediated reduction to form S–SH modification (Tao et al. 2013). Sulphydration has also been detected in glyceraldehyde 3-phosphate dehydrogenase (GAPDH) and actin and was shown to alter the protein activity (Paul and Snyder 2012). Based on these findings, it is reasonable to speculate that the effects of H₂S on cancer cell glycolysis and pH can arise from its sulphydration on membrane-bound proteins, such as the glucose transporters and pH regulators (Lee et al. 2014).

As mentioned previously, OSC in polysulfide form could react with reduced thiols present in the cells to produce H₂S (Benavides et al. 2007). However, given the high reactivity of these OSCs, and competition by other reducing agents in the cells, H₂S production from OSC is subtle. More importantly, sustainability of H₂S-release from these compounds is a major concern as the key to selective anticancer activity of H₂S appears to result from a low and continual exposure to H₂S (Lee et al. 2014). In this context, synthetic H₂S donors such as GYY4137 or HS-NSAID, which can slowly release H₂S over long period of time, would be useful.

Whether H₂S is the sulfur species that gives rise to the observed physiological responses remains debatable. Recently, Greiner et al. reported that H₂S being a sulfur-containing molecule with the lowest oxidation state is unlikely to be the active species that can lead to oxidative thiol modification on protein cysteine residue (Greiner et al. 2013). They analyzed the effect of protein thiol modification by several different H₂S donors and found the formation of polysulfide in aqueous solution, regardless of the donors. The formed polysulfide then readily modified protein inside the cells. Kimura also reported that H₂S-derived polysulfide was able to activate transient receptor potential ankyrin-1 (TRPA1) channels in rat astrocytes (Kimura et al. 2013). In this study, polysulfide was found to be more potent than its parental H₂S. Taken together, these studies suggest that polysulfide may be the molecule that mediates sulfide signaling. Hence, identification of the sulfide-signaling molecule may greatly enhance our understanding and knowledge in using H₂S donors in cancer therapy.

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Part VII

H₂S and Ageing

H₂S: A New Approach to Lifespan Enhancement and Healthy Ageing?

Bedoor Qabazard and Stephen R. Stürzenbaum

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Abstract

Ageing, a progressive structural and functional decline, is considered to be a major risk factor for virtually all ageing-associated pathologies and disabilities, including Alzheimer's disease, Parkinson's disease, stroke, diabetes, atherosclerosis and certain cancers. Biogerontology research has now been largely

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directed towards finding novel drug targets to decelerate the ageing process and attain healthy ageing in order to delay the onset of all ageing-related diseases. H₂S has been reported to exert vasodilatory, antioxidant, antiapoptotic and anti-inflammatory actions and has been shown to act as a signalling molecule, neuromodulator and cytoprotectant. Intriguingly, H₂S has been reported to regulate cell cycle and survival in healthy cells which suggests that it may regulate cell fate and hence the ageing process. This chapter sets out to provide an overview of the current knowledge regarding the involvement of H₂S in ageing, with a specific focus on the invertebrate model nematode *C. elegans*.

Keywords

H₂S • Ageing • Lifespan • *C. elegans* • Nematode

1 The Ageing Process: An Overview

Ageing, or senescence, is the progressive structural and functional organismal decline which emerges as an organism grows older (Golden and Melov 2007). Ageing is characterised by homeostatic imbalances and an impaired ability to respond to stress, thereby increasing the incidence of pathology and probability of death, the ultimate inevitable consequence of ageing. Importantly, ageing is the major risk factor for the development of age-associated diseases, such as cancer, neurodegenerative illnesses, cardiovascular pathologies, dementias and other structural and functional derangements.

1.1 Theories of Ageing: Why Do We Age?

Our understanding of the ageing process has expanded considerably over the last few decades, and it is now apparent that ageing is not driven by a single causative gene or a key defective body system but rather a complex, multidisciplinary event controlled by several interacting cellular, molecular and environmental factors (Weinert and Timiras 2003). Numerous evolutionary-, molecular-, cellular- and systemic-ageing theories are available. For example, ageing has been extensively studied in light of the evolutionary theory, with the most influential theories being the disposal soma theory (Kirkwood 1977) and antagonistic pleiotropy (Williams 1957). Other theories postulate that life expectancy can be influenced by neuro-endocrine control (Hammerman 1987), metabolic rate (Pearl 1928) or telomere length (Campisi et al. 2001).

1.2 Ageing Pathways: The Genetic Control

The fundamental discovery that mutations in single genes (e.g. in *age-1* or *daf-2*) can drastically increase the lifespan of the nematode *Caenorhabditis elegans* (Klass

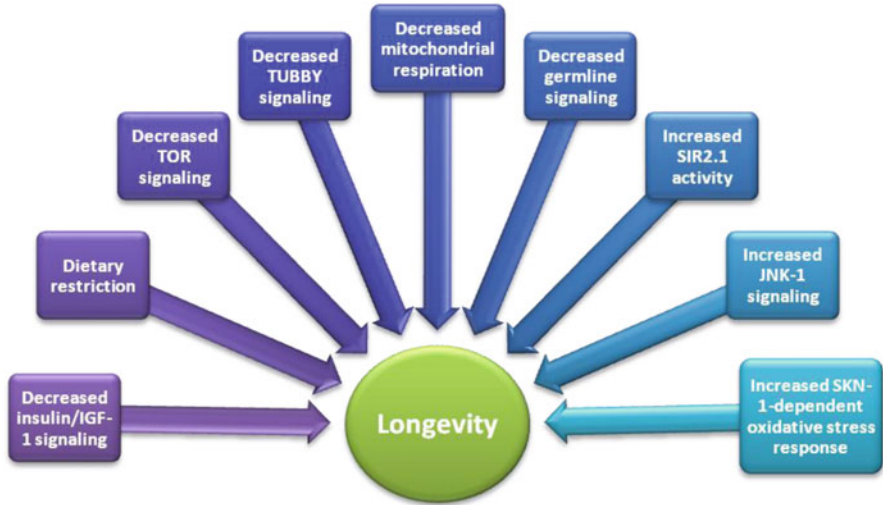


Fig. 1 Multiple evolutionarily conserved pathways are involved in the complex regulation of lifespan. Decreased or increased activity of the following signalling pathways and processes has been shown to modulate ageing (reviewed in Kenyon 2010)

1977; Friedman and Johnson 1988, 2002; Kenyon et al. 1993) supports the notion that ageing is part of a programmed biological route and subject to regulation by specific genetic pathways which confer plasticity to the rate of ageing. The process of lifespan regulation relies on evolutionarily conserved genetic pathways (Fig. 1), including the insulin/insulin growth factor-1 (IGF-1)-like signalling pathway (IIS), DAF-16/FOXO transcription factor, c-Jun N-terminal kinase (JNK) signalling, stress-response pathways, mitochondrial control, dietary restriction, SIR2 deacetylase activity, germ line signalling, sensory perception, TOR (target of rapamycin), TUBBY signalling as well as telomere length. Mutations in genes affecting these signalling pathways protect against stress, reduce disease susceptibility and dramatically extend the lifespan of organisms (Kenyon 2010).

Mutations in *daf-2* (an insulin receptor/IGF-1 receptor (Kimura et al. 1997)) or *age-1* (a phosphatidylinositol 3-kinase (PI3K) subunit (Morris et al. 1996)), key genes within the insulin/IGF-1 signalling (IIS) pathway, are known to modulate hormonal signalling and extend the lifespan of animals by more than 100 % (Kenyon 2010). The IIS signalling cascade is initiated after DAF-2 receptor activates the PI3K subunit AGE-1, which results in activation of Akt/PKB homolog PDK-1 (Paradis and Ruvkun 1998) and subsequently the phosphorylation and activation of AKT-1, AKT-2 and SGK-1 (Paradis and Ruvkun 1998). These kinases inactivate DAF-16, a Forkhead FOXO transcription factor homolog, by retaining it in the cytoplasm and preventing its nuclear translocation (Ogg et al. 1997). DAF-16 is the major downstream target of IIS that, when present in the nucleus, promotes life extension and stress resistance (Lin et al. 1997; Henderson and Johnson 2001; Landis and Murphy 2010). Thus, interventions that reduce the activity and

signalling of the upstream components of the IIS pathway will reduce the repressive signal on DAF-16 and hence enhance the protective pro-survival downstream of DAF-16 targets. In contrast, a null mutation of *daf-16* marginally shortens lifespan and antagonises lifespan extension caused by mutations in the upstream IIS components (e.g. *daf-2*, *age-1*, *akt1/2*, *aap-1*, *pdh-1* or *sgk-1*) (Ogg et al. 1997; Henderson and Johnson 2001). Microarray studies have identified several genes downstream of DAF-16, namely, stress-response genes (e.g. catalases, superoxide dismutases, glutathione *S*-transferases and metallothioneins), antimicrobial genes and chaperones (Lee et al. 2003; McElwee et al. 2003; Murphy et al. 2003). Another hormonal contribution to ageing regulation is believed to originate from the novel age-related gene *Klotho*, which encodes a cell-surface protein with an extracellular domain acting similar to a circulating hormone (Shiraki-Iida et al. 1998). Suppression of this gene accelerates ageing (Kuro-o et al. 1997), whilst upregulation slows ageing in mice (Kurosu et al. 2005) and *C. elegans* (Château et al. 2010). The underlying mechanism of the anti-ageing effect of *Klotho* is poorly understood but may involve IIS inhibition/DAF-16 activation and reduction of oxidative stress (Yamamoto et al. 2005; Château et al. 2010) and angiotensin II (AngII) suppression (Ishizaka et al. 2002).

Another evolutionarily conserved intervention influencing longevity is the caloric or dietary restriction (DR) in the absence of malnutrition (Mair and Dillin 2008). DR is the most efficient environmental intervention proven to delay ageing. An important nutrient-sensing component in this process is the Sirtuins family of NAD⁺-dependent protein deacetylases, which respond to metabolic changes in the cell (i.e. nutrient and energy availability and stress) and mediate the beneficial effects of DR on lifespan in yeast, worms and flies (Rogina and Helfand 2004; Wang et al. 2006). Overexpression of *sir-2.1*, the *C. elegans* ortholog of the yeast SIR2, increases lifespan by 15–50 % via a *daf-16*-dependent action (Tissenbaum and Guarente 2001), but more recent findings have challenged this effect (Burnett et al. 2011). The Sirtuins pathway has been shown to interact with the IIS pathway and directly deacetylate and activate DAF-16 (Berdichevsky et al. 2006). Pharmacological activation of Sirtuins has therefore been widely accepted as a potential DR mimetic approach to slow ageing, an example is the polyphenol drug resveratrol (Wood et al. 2004). Despite the evidence that resveratrol longevity and *sir-2.1* are interlinked in *C. elegans* (Viswanathan et al. 2005), Bass et al. (2007) did not observe lifespan extension with resveratrol in *Drosophila* in seven independent trials and reported only a slight increase in lifespan in wild-type and *sir-2.1* mutant *C. elegans* in some but not all trials (Bass et al. 2007).

Another nutrient- and amino acid-sensing pathway linked to DR is the TOR (target of rapamycin) pathway. TOR is a protein kinase that phosphorylates ribosomal S6 kinase and translation initiation factor 4E-binding protein 1 (4E BP1) in response to nutrients, thus stimulating growth and inhibiting autophagy when food is abundant (Kapahi and Zid 2004; Kenyon 2010). Mutations that reduce TOR signalling extend the lifespan of flies (Kapahi et al. 2004), yeast (Kaerberlein et al. 2005), mice (Harrison et al. 2009) and worms (Vellai et al. 2003) and increase resistance to environmental stress (Hansen et al. 2007). Since the lifespan extension

produced by TOR inhibition is not further increased by DR (Kapahi et al. 2004; Kaeberlein et al. 2005; Bjedov et al. 2010), TOR inhibition is thought to mimic the effects of DR (Kenyon 2010). Recently, the TOR inhibitor rapamycin was shown to return promising anti-ageing effects in mice (Wilkinson et al. 2012). Interestingly, TOR can be inhibited by other longevity-promoting energy sensors such as AMPK (AMP-activated protein kinase) (Inoki et al. 2003) or resveratrol (Baur et al. 2006; Dasgupta and Milbrandt 2007). Likewise, the antidiabetic drug metformin can extend mouse lifespan via AMPK activation (Anisimov et al. 2008). The role of AMPK in longevity is not clear, but there is evidence that its activation mimics DR and has been found to extend the lifespan of *C. elegans* (Apfeld et al. 2004; Greer et al. 2007a, b) and mice (Ingram et al. 2004; Anisimov et al. 2005) partly via FOXO/DAF-16 phosphorylation (Greer et al. 2007a, b).

The stress-induced MAP kinase, JNK, prolongs survival in worms (Oh et al. 2005) and flies (Wang et al. 2003). In *C. elegans*, *jnk-1* overexpression leads to a DAF-16-dependent increase in survival and improved resistance to heat and oxidative stress (Oh et al. 2005). JNK-1 interacts with IIS by directly phosphorylating/activating DAF-16, thereby inducing the expression of DAF-16 target genes which are crucial for longevity and stress tolerance. Another important stress-related pathway is SKN-1/Nrf2, which protects against oxidative stress by activating the conserved phase II detoxification system through constitutive and stress-inducible mechanisms (An and Blackwell 2003). Phase II enzymes mediate the synthesis of glutathione, scavenge free radicals and help to detoxify the reactive intermediates from the phase I reactions. Hence, SKN-1 induces the expression of genes involved in detoxification, cell repair and pathogen resistance and suppresses genes that decrease survival and stress tolerance (Oliveira et al. 2009). SKN-1 is expressed constitutively in ASI chemosensory neurons under normal conditions and induced in the intestinal nuclei (by reduced IIS signalling) in response to stress (An and Blackwell 2003; An et al. 2005; Tullet et al. 2008). Moreover, SKN-1 contributes to the longevity and stress resistance associated with IIS inhibition, and, likewise, DAF-16 is essential for the induction of some SKN-1 targets under reduced IIS signalling (Tullet et al. 2008), suggesting a possible crosstalk between the two pathways in regulating stress response and lifespan. In addition, an interaction between SKN-1 and p38 MAPK has also been suggested as p38 MAPK represses the inhibitory signal from glycogen synthase kinase-3 (GSK-3) on SKN-1 leading to intestinal SKN-1 expression and activation under oxidative stress (Inoue et al. 2005).

2 The Role of H₂S in Ageing and Age-Associated Diseases

2.1 H₂S: A Prehistoric Entity

H₂S is thought to have caused a mass extinction in the Permian-Triassic periods, some 250 million years ago (Kump et al. 2005; Ward 2006). Massive volcanic eruptions in Siberia and a cascade of environmental destructions resulted in a severe

depletion of oxygen levels in the oceans and atmosphere, which caused the majority of organisms to become extinct (Kump et al. 2005). However, certain species managed to survive these harsh habitats, thanks to their ability to utilise H₂S in the process of chemosynthesis (Lavu et al. 2011). For example, the anaerobic green sulfur bacteria (*Chlorobium*) respired, instead of oxygen, the sulfate dissolved in water and in doing so continued to generate more lethal H₂S gas (Wang 2012). Upon the subsequent environmental depletion of H₂S, primitive species gradually switched to use water instead of H₂S by means of photosynthesis (Lavu et al. 2011). Blackstone et al. (2005) found that very low levels of the toxic H₂S gas can reduce the oxygen demands and prolong survival in mice by inducing a hypometabolic hibernation-like or ‘suspended animation’ state that extend their lives.

2.2 Induced Hypothermia and Suspended Animation: From Animals to Humans

According to Blackstone et al. (2005), house mice (*Mus musculus*) exposed to low (non-toxic) doses of H₂S enter a reversible state of suspended animation-like hypothermia. In detail, H₂S exposure dose-dependently decreases the metabolic rate by 50 % with a 90 % reduction in oxygen demand. The core body temperature declines from 37 °C to about 15 °C, and the respiration falls from 120 to 10 breaths per minute. The mice survive this procedure for 6 h and show no negative health consequences afterwards (Blackstone et al. 2005). In this context it should be noted that mice do not naturally hibernate but can undergo a clinical torpor state during food deprivation. Further experiments uncovered that a pretreatment with H₂S primes the mice to tolerate and survive hypoxia. Mice exposed to 150 ppm H₂S for 20 min are more resistant to an oxygen challenge, possibly due to the reduction in metabolic rate and whole-body oxygen demand (Blackstone and Roth 2007). Furthermore, rats display a reduced degree of brain damage when exposed to H₂S for 48 h following an experimentally induced stroke (Florian et al. 2008). In these animals the expression of pro-inflammatory and pro-apoptotic genes is suppressed in the areas surrounding the infarct region, and the mice are characterised by superior memory and learning profiles. Likewise, H₂S inhalation (80 ppm) in mice (Faller et al. 2010) and NaSH infusion in rats (Aslami et al. 2010) protect against ventilator-induced lung injury.

Further studies with larger animals yielded conflicting results. For example, studies failed to induce hibernation or hypometabolism in pigs (Simon et al. 2008) and sheep (Haouzi et al. 2008), leading the authors to conclude that the effects seen in mice may not be achievable in larger mammals. One reason could be due to the difference in the ratio between surface area and body mass which may render large animals less responsive to the same relative dose of H₂S (Wang 2012). However, if H₂S-induced hibernation would be feasible in humans, it would promise potential therapeutic benefits regarding the emergency management of severely injured patients and in the conservation of donated organs during transplant surgery. Indeed, in 2010 it was announced that H₂S-induced hypothermia

had completed phase I clinical trials, but the trials were withdrawn and terminated for unknown reasons.

2.3 H₂S: Protective Effects That Promote Healthy Ageing

Researchers have recently explored the physiological importance of endogenously produced H₂S in cellular senescence and organism longevity and, hence, its contribution to ageing and age-related diseases. In addition to the effects of H₂S as an anti-inflammatory, antiapoptotic and antioxidant mediator, several lines of evidence support the notion that H₂S acts as a cytoprotective mediator in the ageing process. Plasma levels of H₂S in humans decrease with age (Chen et al. 2005). Similarly, the activities of CBS and CSE enzymes decline progressively in the livers of ageing mice with a corresponding decrease in cysteine and glutathione levels (Dröge et al. 2006). CSE gene and protein expression and enzyme activity are attenuated in the lenses of old rats, leading to cataract formation in vitro (Sastre et al. 2005). H₂S replacement promotes endothelial protection in hyperglycaemia due to the preservation of mitochondrial function (Suzuki et al. 2011). Likewise, the protective effect of calorie restriction in blood vessels of ageing rats has been attributed to the maintenance of normal H₂S levels (Predmore et al. 2010). Moreover, the suggested antioxidant role of H₂S seems to align well with the free radical theory of ageing and may contribute to its therapeutic potential in ageing-associated diseases, such as AD, PD, atherosclerosis and others. H₂S acts as a reducing agent that can scavenge other oxidative and nitrosative species (Whiteman et al. 2004, 2005), but additional mechanisms are believed to be responsible for the antioxidant role of H₂S. Importantly, H₂S can increase the intracellular GSH pool, the major cellular antioxidant, by stimulating γ -glutamylcysteine synthase and cysteine transport in neurons (Kimura and Kimura 2004) and glutamate uptake in astrocytes (Lu et al. 2008). NaSH confers protection to a murine hippocampal HT22 neuronal cell line from oxidative glutamate toxicity by opening K_{ATP} and Cl⁻ channels in addition to increasing GSH levels (Kimura et al. 2006). In agreement, the H₂S-releasing derivative of sildenafil, ACS6, inhibits superoxide formation and NADH oxidation in endothelial cells (Muzaffar et al. 2008). Moreover, the slow-releasing H₂S donor GYY4137 protects cultured chondrocytes and mesenchymal progenitor cells from oxidative and nitrate stress induced by SIN-1 and 4-HNE (Fox et al. 2012).

Molecular studies revealed that certain age-related genes are involved in the H₂S response, notably *Sirtuins* (Miller and Roth 2007) and *Klotho* (Zhang et al. 2013). H₂S also affects transcription factors, namely, Nrf-2 (nuclear factor erythroid 2-related factor 2), STAT3 (signal transducer and activator of transcription 3) and Hif-1 (hypoxia inducible factor 1). Nrf-2 regulates the expression of several cytoprotective and antioxidant enzymes (e.g. thioredoxin and heme oxygenase), and activation of this transcription factor contributes to the cardioprotective effect of H₂S (Calvert et al. 2009) and mediates H₂S responses in the nematode *C. elegans* (Miller et al. 2011). STAT3, the cell survival and proliferation regulator, is

activated by GYY4137 treatment in LPS-injected rats (Li et al. 2009). Lastly, enhanced Hif-1 activity and a genetic requirement for this factor were reported in *C. elegans* after H₂S exposure (Budde and Roth 2010; Miller et al. 2011). Recently, H₂S has been shown to prevent H₂O₂-induced senescence in human umbilical vein endothelial cells through SIRT1 activation (Suo et al. 2013) and protect against cellular ageing via *S*-sulfhydration of Keap1 and Nrf-2 activation in mouse embryonic fibroblasts (Yang et al. 2013). In SH-SY5Y neuronal cells, H₂S confers protection against D-galactose-induced cell injury by suppression of advanced glycation end-product formation and oxidative stress (Liu et al. 2013). H₂S has been associated with mitogen-activated protein kinase (MAPK); however, whilst most papers suggest an activation, others claim that H₂S is involved in the inhibition of the MAPK pathway. For instance, H₂S induces apoptosis of human aorta smooth muscle cells (Yang et al. 2004) and stimulates angiogenesis in cultured human umbilical vein endothelial cells (Papapetropoulos et al. 2009) via p38MAPK activation, whereas the antiapoptotic effect of H₂S in human polymorphonuclear leukocytes is likely due to the inhibition of p38MAPK (Rinaldi et al. 2006). In addition, H₂S activates and promotes the phosphorylation of ERK1/2 (extracellular signal-regulated kinase 1/2) in HEK-239 cells (Yang et al. 2004), human colon cancer cells (Cai et al. 2010) and rat vascular smooth muscle cells (Jeong et al. 2006). Moreover, akt activation/phosphorylation has also been suggested as a mechanism for H₂S-induced angiogenesis (Cai et al. 2007) and cardioprotection (Yong et al. 2008). JNK (c-Jun NH₂ terminal kinase) upregulation by NaSH increases rat intestinal epithelial cell proliferation (Deplancke and Gaskins 2003), whilst JNK downregulation by NaSH prevents apoptosis in a human neuroblastoma cell line (Hu et al. 2009).

S-Sulfhydration is a signalling mechanism for H₂S involving posttranslational modification of cellular proteins and enzymes (Mustafa et al. 2009). The mechanism entails covalent modification of reactive cysteine residues via the addition of one sulphur (from H₂S) to the thiol group of cysteine (-SH) to yield a hydro-persulfide molecule (-SSH). This process results in an increased activity of the modified protein bearing the -SSH moiety (Mustafa et al. 2009). H₂S can alter, via this process, the function of a wide range of cellular proteins and enzymes in multiple settings, most notably in cell survival and stress pathways. For example, the sulfhydration of cysteine residues can shield them from oxidation under oxidative stress conditions, thereby preserving protein function and preventing permanent damage (Paul and Snyder 2012). H₂S-mediated *S*-sulfhydration of p65 at cysteine-38 (Sen et al. 2012) and PTP1B phosphatase (Krishnan et al. 2011) inhibit the apoptotic effect of NF-κB and modulate ER stress responses, respectively. More recently, H₂S has been shown to regulate cellular redox signalling via electrophile *S*-sulfhydration (Nishida et al. 2012). Further studies are required to confirm and validate this mechanism and to what extent the various H₂S-mediated responses correlate with health and disease.

2.4 H₂S: A Novel Therapeutic Agent in Age-Associated Pathologies?

A considerable body of evidence suggests that H₂S acts as a protective mediator in the heart, blood vessels, brain and other body tissues and compartments by virtue of its possible antioxidant, antiapoptotic and anti-inflammatory properties. Although the role of H₂S in inflammation is controversial (Whiteman and Winyard 2011) with some studies advocating a pro-inflammatory effect of H₂S by activating leukocyte recruitment (Zhang and Bhatia 2008), numerous studies describe H₂S as an anti-inflammatory molecule. For example, H₂S was shown to counteract endothelial inflammation by inhibiting leukocyte adherence (Zanardo et al. 2006). Moreover, H₂S accelerates the resolution of inflammation by enhancing neutrophil survival through p38MAPK inhibition (Rinaldi et al. 2006). NaSH treatment reverses LPS (lipopolysaccharide)-induced iNOS expression and NO release (Oh et al. 2006; Muzaffar et al. 2008) and prevents homocysteine-induced toxicity and subsequent ROS formation in VSMCs (Yan et al. 2006). Likewise, exogenous H₂S attenuates oxidised low-density lipoprotein (oxLDL)-induced cytotoxicity in cultured HUVECs and was shown to protect against atherosclerosis (Laggner et al. 2007a, b; Jeney et al. 2009). The anti-atherosclerotic effect of H₂S is further supported by the reported inhibition of platelet aggregation (Zagli et al. 2007), inhibition of VSMCs proliferation (Du et al. 2004) and reduction of neointima formation of balloon-injured carotid arteries (Meng et al. 2007). More compellingly, H₂S appears to activate antiapoptotic signalling to protect against ischaemia/reperfusion (I/R) injury (Sivarajah et al. 2006; Jha et al. 2008). Activation of Nrf-2 (Calvert et al. 2009), Akt (Cai et al. 2007; Yong et al. 2008) and ERK1/2 and PI3K (Hu et al. 2008) signalling pathways and preservation of mitochondrial structure and function (Elrod et al. 2007) have all been suggested to contribute to H₂S-induced cardioprotection. H₂S may also mediate cardioprotection by influencing intracellular organelles such as the endoplasmic reticulum (ER), as evidenced by the suppression of cardiac myocyte death in hyperhomocysteinaemic rats (Wei et al. 2010).

Exogenous H₂S has been shown to protect against detrimental pro-inflammatory oxidants, such as peroxynitrite (ONOO⁻) (Whiteman et al. 2004), hydrogen peroxide (H₂O₂) (Muzaffar et al. 2008), hypochlorous acid (HOCl) (Whiteman et al. 2005) and β -amyloid (Tang et al. 2008). Given that H₂S is not very abundant suggests that the potential for scavenging activity is limited and/or that other mechanisms participate in the cytoprotective action of this gas. In agreement, H₂S protects the mouse HT22 neuronal cell line from peroxynitrite-induced oxidative stress by activating K_{ATP} and Cl⁻ channels and increasing GSH levels (Kimura and Kimura 2004; Kimura et al. 2006). In addition, H₂S-inhibited HOCl provokes cytotoxicity, protein oxidation and lipid peroxidation in the brain (Whiteman et al. 2005) and protects brain endothelial cells against methionine-induced oxidative stress (Tyagi et al. 2009). The interaction with cystic fibrosis transmembrane conductance regulator (CFTR) Cl⁻ channels may also be a mechanism for H₂S protection against oxidative stress (Lahousse et al. 2003). It has been

suggested that Cl^- channels play a critical role in maintaining normal cell survival as the reduction of Cl^- current causes cell death in hippocampal neurons, and indeed a downregulation of CFTR gene expression is observed in the brains of Alzheimer's disease (AD) patients (Lahousse et al. 2003). In fact, CBS has been stipulated to be a potential risk factor for the pathogenesis of AD (Beyer et al. 2004). H_2S levels are also reduced in rat models of Parkinson's disease (PD) and administering H_2S alleviates the condition (Hu et al. 2010). In animal models of PD, H_2S reduces the levels of $\text{TNF}\alpha$ and IL-6 (Lee et al. 2010); however, in Down syndrome more H_2S is produced (Kamoun et al. 2003). Interestingly, H_2S has been found to influence Cl^- conductance in the CNS leading to neuroprotection (Kimura et al. 2005). In support of this, H_2S activates CFTR Cl^- channels in HT22 neuronal cell lines leading to neuroprotection during oxytosis, which is reversed by the CFTR blockers NPPB and IAA-94 (Kimura et al. 2006). Moreover, H_2S activates $\text{Cl}^-/\text{HCO}_3^-$ transporters in smooth muscle cells (Lee et al. 2007). In addition to the interaction with ion channels in the CNS, H_2S can affect neurotransmitters, namely, GABA (the major inhibitory neurotransmitter) and glutamate (the major excitatory neurotransmitter). As such, H_2S ameliorates hippocampal damage associated with recurrent febrile seizures by upregulating the expression of GABA_B receptors (in Ca^{2+} -dependent manner) which in turn enhances GABA_B ergic inhibition (Han et al. 2005a). It can be therapeutically important during seizures/epilepsy to restore the disrupted balance between CNS excitation and inhibition (Han et al. 2005b). In contrast, H_2S seems to potentiate the effect of glutamate on the NMDA receptor with the subsequent toxicity and neuronal death due to Ca^{2+} overload (Gagliardi 2000; Garcia-Bereguain et al. 2008).

H_2S demonstrates a notable growth-suppressive effect on prostate cancer (Pei et al. 2011), breast cancer (Chattopadhyay et al. 2012a) and colon cancer (Chattopadhyay et al. 2012b) cell lines. Lee and colleagues (2011) suggest that the slow-releasing H_2S donor GYY4137 is a potential anticancer agent owing to its ability to kill seven different human cancer cell lines (HeLa, HCT-116, Hep G2, HL-60, MCF-7, MV4-11 and U2OS) in a concentration-dependent manner. Moreover, H_2S -releasing nonsteroidal anti-inflammatory drugs (NSAID) have been shown to constrain the growth of various human cancer cells by inhibiting cell proliferation, inducing apoptosis, and cause G_0/G_1 cell cycle block (Chattopadhyay et al. 2012c).

However, the precise role of H_2S in ageing and its contribution to age-related pathologies awaits further elucidation but promises to return new therapeutic leads to be added to the growing list of the beneficial effects of this gas.

2.5 H_2S -Mediated Lifespan Extension: Insights from *Caenorhabditis elegans*

Given that the monitoring of complete survival profiles in humans is challenging, model organisms such as the nematode *C. elegans* are valuable surrogates and arguably the premier animal model for lifespan studies and ageing research.

Miller and Roth (2007) observed that exposure to a small amount of H₂S (50 ppm, in a controlled atmosphere) extends the lifespan of *C. elegans* and increases thermotolerance without any major change in metabolic function. H₂S-treated worms exhibit a 70 % increase in survival rate compared to the untreated controls, and the authors suggest that the mechanism of lifespan extension is dependent on the SIR2 (the silent information regulator 2) pathway. However, it is noteworthy that the response to H₂S is not identical among different species. For example, whilst an exposure to H₂S does not affect the metabolic function in *C. elegans*, hypothermia and hypometabolism are observed in rodents; however, this may also be due to differences in the relative dose exposures and uptake dynamics. Nevertheless, most ageing pathways are conserved between man and worm, and thus *C. elegans* represents a great advantage in exploring the role of H₂S in the ageing process, which can subsequently be extrapolated to higher animals in order to unravel the riddles of longevity and allow a more targeted approach to combat ageing-associated diseases.

A recent study, which explored exogenous H₂S effects and endogenous H₂S metabolism in the nematode *C. elegans*, demonstrated that a slow-releasing H₂S donor (GYY4137) is able to extend the median survival by up to 23 % and increases tolerance towards oxidative and endoplasmic reticulum (ER) stress (Qabazard et al. 2013). These authors also found that *cysl-2*, a sulfhydrylase/cysteine synthase in *C. elegans*, is transcriptionally upregulated by GYY4137 treatment and that the deletion of *cysl-2* results in a significant reduction in lifespan which is partially recovered by the supplementation of GYY4137. Likewise, in a mammalian cell culture system, GYY4137 is able to protect bovine aortic endothelial cells (BAECs) from oxidative stress and (H₂O₂)-induced cell death (Qabazard et al. 2013). Taken together, this provides further support that H₂S exerts a protective function which is consistent with the longevity dividend theory (Olshansky et al. 2006).

Moreover, a related study revealed that the *C. elegans* genome encodes orthologues of the three H₂S synthesising enzymes, namely, CBS, CSE and 3-MST, and is capable of the endogenous synthesis of H₂S (Qabazard et al. 2014). Genetic deficiency of *mpst-1* (3-MST orthologue 1), but not *cth-2* (CSE orthologue), reduces the lifespan of *C. elegans*, an effect which is reversed in worms supplemented with GYY4137 (Qabazard et al. 2014). Treatment of *C. elegans* with GYY4137 increases the expression of several age-related, stress-response and antioxidant genes, increases the lifespan in short-lived *mev-1* mutants (which are characterised by an elevated level of oxidative stress) and protects wild-type *C. elegans* against paraquat poisoning (Qabazard et al. 2014). The authors conclude that the lifespan-prolonging and health-promoting effects of H₂S in *C. elegans* are likely due to the antioxidant action of this highly cell-permeable gas (Qabazard et al. 2014). Whether H₂S prolongs directly the lifespan and promotes healthy ageing in mammals is a key question which has yet to be fully explored.

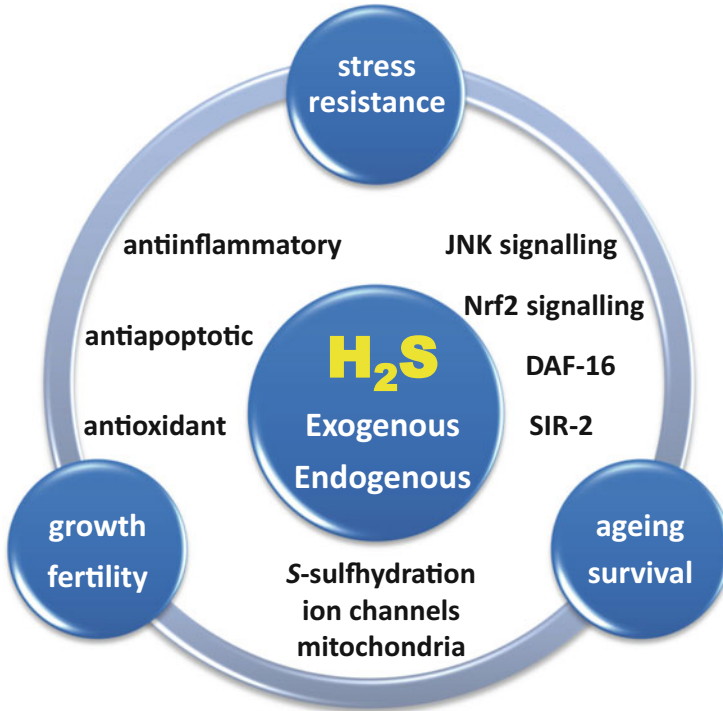


Fig. 2 Summary of H₂S contribution to ageing

3 Conclusions

Numerous findings suggest that endogenous H₂S plays a critical regulatory role in ageing and the cellular response to stress (Fig. 2), which in concert highlight that exogenous H₂S donors may prove valuable in the pharmacological intervention to delay the ageing process, the protection against age-associated diseases and ultimately the extension of a healthy lifespan.

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Part VIII

H₂S, Measurement and Possible Therapeutics?

Fluorescent Probes for H₂S Detection and Quantification

Wei Feng and Brian W. Dymock

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Abstract

Many diverse, sensitive and structurally novel fluorescent probes have recently been reported for H₂S detection. Quantification of H₂S requires a selective chemosensor which will react only with H₂S against a background of high concentrations of other thiols or reducing agents. Most published probes are able to quantify H₂S selectively in a simple in vitro system with the most

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sensitive probes able to detect H₂S at below 100 nM concentrations. A subset of probes also have utility in sensing H₂S in living cells, and there are now several with specific sub-cellular localization and a few cases of *in vivo* applications. Biologists studying H₂S now have a wide range of tools to assist them to aid further understanding of the role of H₂S in biology.

Keywords

Hydrogen sulfide • Fluorescent probe • Chemosensor

1 Introduction

Fluorescent probe molecules which can chemically detect or ‘sense’ H₂S are often called ‘chemosensors’ or ‘chemodosimeters’. Quantification of H₂S requires a very selective chemosensor which will react only with H₂S against a background of high concentrations of other thiols or reducing agents, as is found in a cell. Most published chemosensors are able to quantify H₂S in a simple *in vitro* system, but application in the cell is still very challenging in most cases. A wide range of diverse chemosensing probes have been reported with most being simple probe molecules that selectively react with H₂S to produce a fluorescent product that can be quantified using a standard laboratory fluorimeter. For application in a wide range of biological situations, a good probe molecule must fulfil a range of criteria:

- Rapidly react with H₂S under physiological conditions
- Selectively react with H₂S over high concentrations of biological thiols and anions
- Be sensitive to low concentrations of H₂S (high quantum yield)
- Linear response within physiological H₂S concentration range
- Be biocompatible with good permeability and intracellular stability
- Is not toxic and does not produce toxic products
- Be chemically stable for storage and longer-term studies
- Be straightforward to synthesise or be commercially available

With those objectives in mind, most authors have addressed several but rarely adequately solved all of these challenges. This chapter discusses published probes operating by a fluorescence readout and will critically appraise their performance against the aforementioned key criteria. The probes are organised into three categories: (1) probes which react with H₂S through a nucleophilic reaction, (2) probes which undergo selective reduction by H₂S and (3) probes which trap H₂S using a coordinated metal.

The probes discussed herein are especially selected from the more recent literature from the past 2–3 years, for earlier material readers are directed to other reviews (Lin and Chang 2012; Peng and Xian 2014; Yu et al. 2014).

2 Probes Which React with H₂S Through a Nucleophilic Reaction

Under physiological conditions, H₂S (pK_a c.6.9) dissociates to form HS⁻ as the major form (estimated at 72 % (Li and Lancaster 2013)), which is a very good nucleophile. A number of reports on H₂S fluorescent probes which are based on this nucleophilic feature have been published recently. When the H₂S amount is as small as endogenous levels, the side fluorescence turn-on can be comparable to that resulting from H₂S. As a non-substituted thiol, H₂S is different from mono-substituted biothiols, such as cysteine and glutathione (GSH), in that H₂S can undergo nucleophilic reaction twice instead of once for normal biothiols. It seems that probes that will turn on fluorescence only after two nucleophilic reactions can distinguish H₂S from biothiols. The general design of nucleophilic substitution–cyclisation-based fluorescent probes has been reviewed in 2014 by Xian as shown in Fig. 1 (Peng et al. 2014). H₂S reacts with the electrophilic component in probe **A** to deliver an intermediate **B1** which contains a free thiol. In the presence of another electrophilic centre, such as an ester group, the free thiol will undergo intramolecular cyclisation spontaneously to form product **P** and release the fluorophore which can be detected by fluorescence measurement. Probe **A** may also react with biothiols to form intermediate **B2**. However, the cyclisation will not happen due to the lack of a free SH in **B2**. Biothiols are also not reactive enough to undergo intermolecular attack on the ester group. As a result, the fluorophore cannot be released.

In 2011, He and co-workers reported the synthesis and application of bis-electrophilic centred probes SFP 1 (**1**) and SFP 2 (**3**) (Fig. 2) (Qian et al. 2011). 1,3,5-Triaryl-2-pyrazoline and 4,4-difluoro-1,3,5,7-tetramethyl-4-bora-3a,4a-diaza-*s*-indacene (BODIPY) were employed as fluorophores for each probe, respectively. Aldehyde groups of **1** and **3** react with HS⁻ to form an intermediate hemithioacetal group which then undergoes Michael addition to a

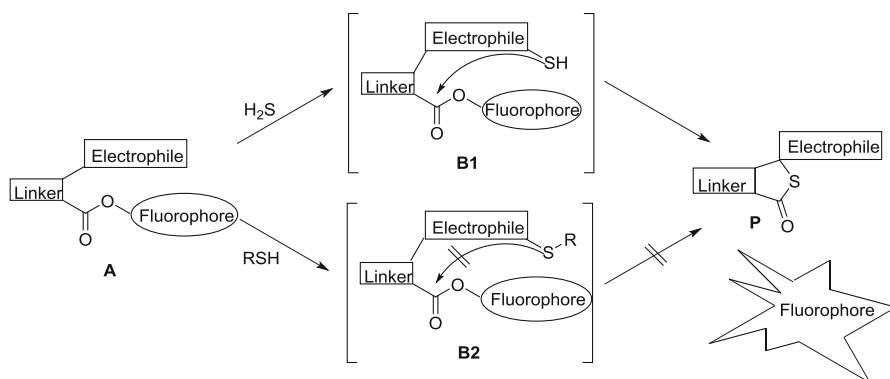


Fig. 1 General design of the nucleophilic substitution–cyclisation-based fluorescent probes for H₂S

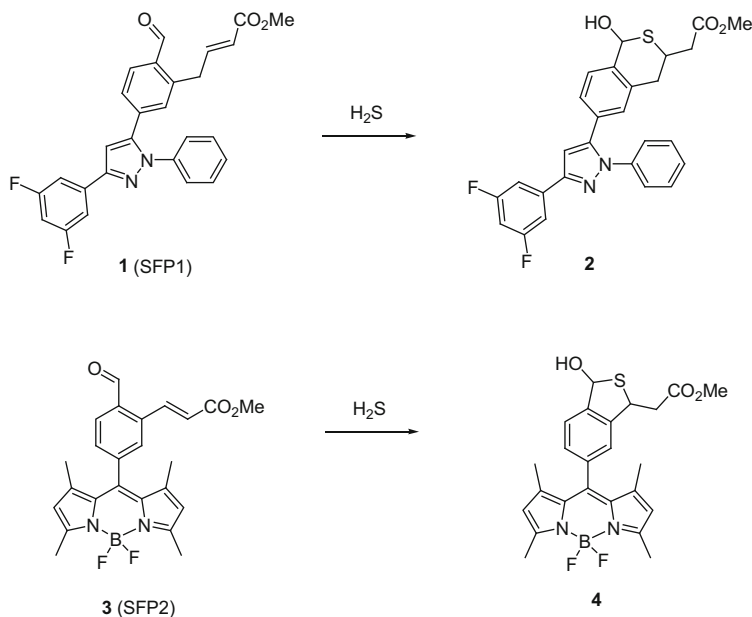
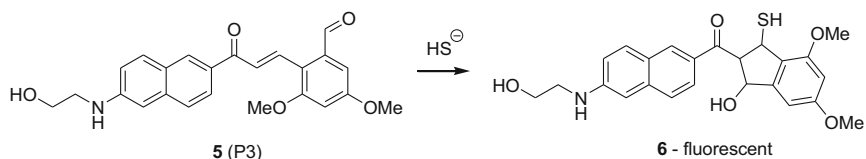


Fig. 2 Structures of probe SFP 1 and SFP 2 and their mechanisms

nearby acrylate ester give cyclised products **2** and **4**, respectively. Fluorescence response of these probes to sulphide ($50\ \mu\text{M}$) is about fourfold than that of biothiols ($1\ \text{mM}$). However, biothiols do turn on fluorescence, which is much higher than control experiments. Imaging of sulphide with these two probes in HeLa cells was successful. Images were obtained by incubating $10\ \mu\text{M}$ probes with Na_2S of different concentrations ranging from 10 to $200\ \mu\text{M}$. Furthermore, as well as addition of exogenous Na_2S , addition of GSH and cysteine can also lead to fluorescence turn-on in HeLa cells following increased production of H_2S . However, better sensitivity of H_2S detection in cells is required.

Also based on an aldehyde-Michael acceptor system, the recently reported probe P3 (**5**) was shown to form a five-membered carbocyclic product following a three-step mechanism that the authors propose as a revision to the generally conceived mechanism of initial Michael addition (Fig. 3) (Singha et al. 2015). Probe **5** was optimised using a computational approach which added two methoxy groups to the aromatic ring increasing the electrophilicity of the enone system as well as creating a steric hindrance to attack at the enone, hence favouring the smaller HS^- anion attack. **5** ($10\ \mu\text{M}$) reacts with Na_2S ($100\ \mu\text{M}$) in only 5 min (pH 7.4 in Hepes buffer). Very good selectivity over other biological thiols is achieved with cysteine ($200\ \mu\text{M}$) giving a 7 % response. However, with 10 % FBS added to the media, the selectivity over cysteine was improved indicating promise for cellular application. **5** was reported to have a LOD of only $50\ \text{nM}$ with S/N ratio of 3 making it one of the most sensitive probes published to date. In HeLa cells, **5** ($10\ \mu\text{M}$) was sensitive enough to detect endogenous levels of H_2S following a 30 min incubation,



Mechanism: (i) aldehyde addition of HS⁻, (ii) intramolecular Michael addition, (iii) aldol condensation

Fig. 3 Structure of probe P3 and the conversion to its fluorescent carbocyclic product via a three-step mechanism

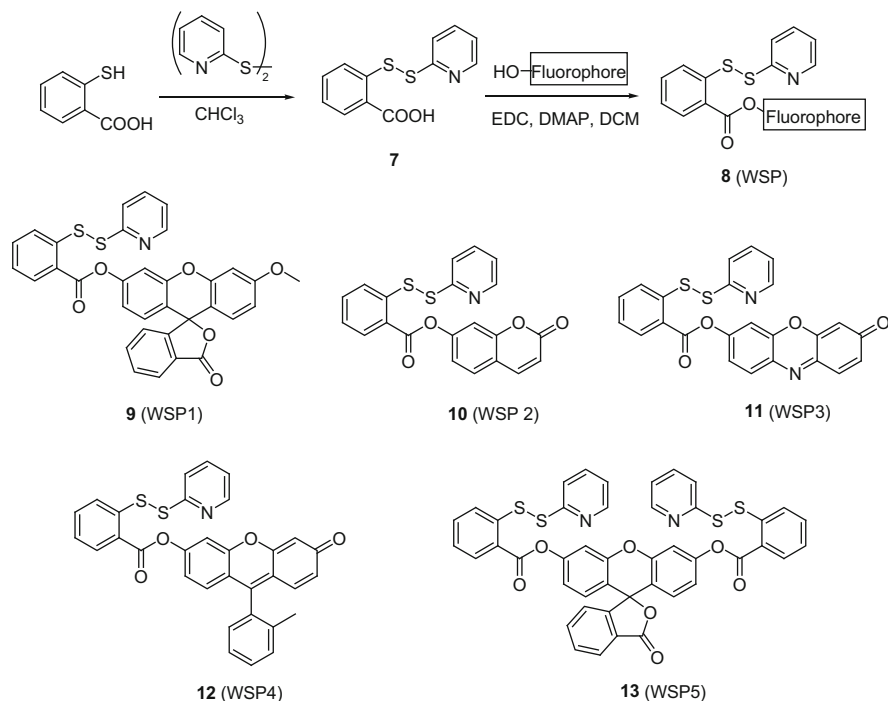


Fig. 4 Synthesis of WSP probes

further supported by a reduction in signal when H₂S synthesis inhibitors phorbol myristate acetate or propargyl glycine were added. With the addition of exogenous Na₂S (300 μM), a very strong fluorescence signal was obtained. Being a sensitive and selective probe, **5** demonstrates that other less selective probes have overestimated the levels of endogenously produced H₂S.

A series of WSP (**8**) probes were synthesised by Xian (Liu et al. 2011; Peng et al. 2014). Intermediate 2-(2-pyridinyldithio)-benzoic acid (**7**) was easily prepared in good yield. It was coupled to different fluorophores, such as methoxy fluorescein, 7-hydroxycoumarin, resorufin and 2-methyl Tokyo Green, through an ester formation reaction (Fig. 4). WSP 1 (**9**), WSP 2 (**10**), WSP 3 (**11**) and WSP

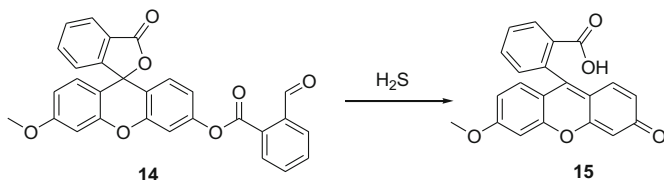


Fig. 5 Structure of probe **15** and its mechanism

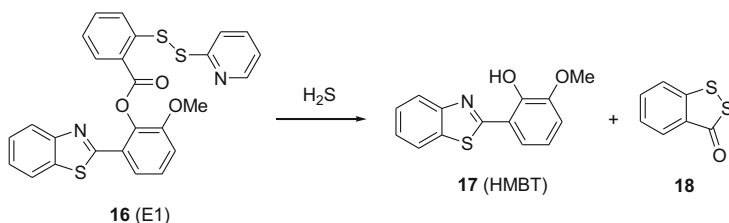


Fig. 6 Structure of probe E1 and its mechanism

5 (**13**) have low detection limits ranging from 47 to 79 nM, and WSP **4** (**12**) has a detection limit of 266 nM. A time period of 30 min is required for the fluorescence intensity to reach the maximum. However, it can be shortened to 5 min when a small amount of surfactant is present. All of these probes showed good selectivity of H_2S over other reactive sulphur species (200 μM) including cysteine, GSH, homocysteine, SO_3^{2-} and $\text{S}_2\text{O}_3^{2-}$. Good fluorescence images were obtained when **12** (30 μM) and **13** (50 μM) were incubated together with NaSH in HeLa cells, and no fluorescence was observed when no NaSH was added, indicating a good selectivity of H_2S over biothiols.

In 2014, Zhang and co-workers reported the synthesis and application of probe **14** (Fig. 5) (Liu et al. 2014b). It has a 2-formylbenzoate group which can react with HS^- to form a hemithioacetal. This probe is applicable for H_2S detection with H_2S concentrations ranging from 2.5 μM to 1000 μM . Confocal fluorescence imaging was achieved when probe **14** (5 μM) was incubated together with Na_2S (1 mM) in HeLa cells.

Probe E1 (**16**) was developed by Qian in 2012 (Xu et al. 2012). It is based on excited-state intramolecular proton transfer (ESIPT). 2-(20-Hydroxyphenyl) benzothiazole (HMBT) was chosen as the fluorophore, and the bis-electrophilic centres are disulphide and ester groups (Fig. 6). Following reaction of HS^- with the disulphide, a hydrogenpersulphide is formed which then attacks the ester forming the fluorescent HMBT (**17**) and by-product **18**. Probe **16** responds to H_2S within 2 min with a 30-fold fluorescence enhancement. The limit of detection can be as low as 120 nM. It is also applicable for H_2S imaging in HeLa cells, and the fluorescence can be significantly enhanced by addition of cetyltrimethylammonium bromide (CTAB). However, there is a disadvantage of these probes. Although biothiols, such as cysteine and GSH, cannot react to elicit fluorescence, they do

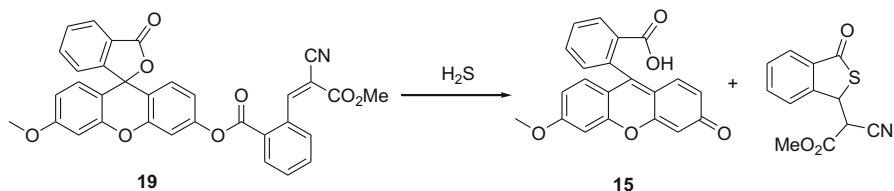


Fig. 7 Structure of probe **19** and its mechanism

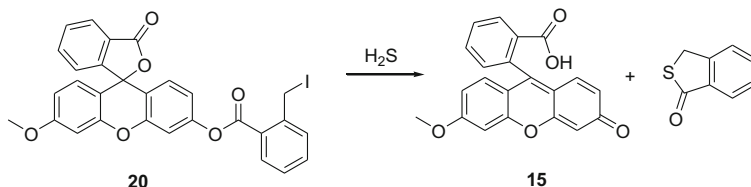


Fig. 8 Structure of probe **21** and its mechanism

react with the disulphide or aldehyde moieties, thus leading to the requirement of high probe loadings. Since H₂S and biothiols have different pK_a values in aqueous solution (approximately 6.9 for H₂S and 8.5 for thiols), a probe that has bis-electrophilic centres which are both reactive to H₂S but not to biothiols is possible.

In 2012, Xian developed probe **19** which has bis-electrophilic centres: a Michael acceptor and an ester group (Fig. 7) (Liu et al. 2012). Experiments showed that the Michael acceptor does not react with cysteine or GSH. Upon treatment with NaSH, probe **19** responds with an 11-fold fluorescence increase within 30 min, generating fluorescent product **15**. However, when **19** was used in COS-7 cells for H₂S imaging, only a weak fluorescence image was observed.

In 2013, Guo and co-workers reported the design and synthesis of probe **20**, which employs an alkyl iodide as the first electrophilic centre (Fig. 8) (Zhang et al. 2013). Experiments showed that cysteine or GSH (200 μM) does not react with **20** (20 μM) in PBS:MeCN (1:1, v/v), while reaction of **20** with H₂S is complete within 1 h. The capability of **20** (5 μM) to image H₂S (100 μM NaSH) in a biological system was demonstrated in COS-7 cells. Fluorescence intensity can be further enhanced by adding CTAB without any interference from biothiols.

Probes DSP 1 (**21**), DSP 2 (**22**) and DSP 3 (**23**) were also developed by Xian (Fig. 9) (Liu et al. 2014a). **23** gave a much stronger fluorescence response than **21** or **22**. This probe was designed for the detection of hydrogen polysulphide H₂S_n through aromatic substitution and cyclisation reactions. With **23**, H₂S_n can be detected with a very low LOD of 71 nM. When H₂S was mixed together with reactive oxygen species (ROS), fluorescence enhancement was also observed. However, **23** does not respond to H₂S itself.

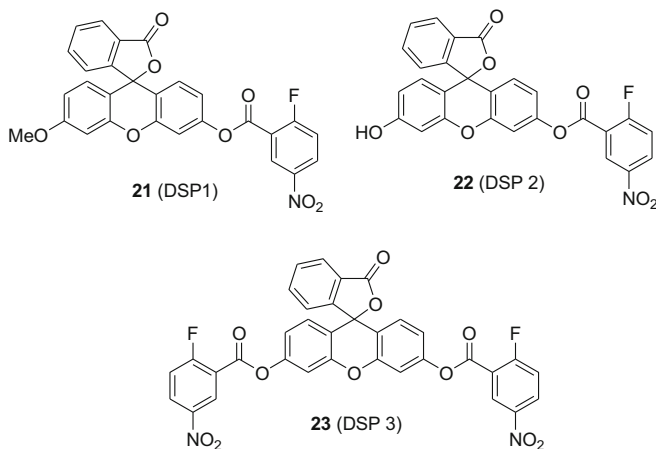


Fig. 9 Structures of probes DSP 1, DSP 2, DSP 3

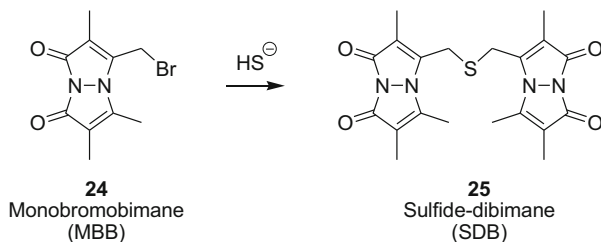


Fig. 10 Reaction of MBB with sulphide to produce fluorescent SDB which is accurately quantified by reversed-phase HPLC using a fluorescence detector

Probably the most sensitive detection system yet published is the protocol developed by Kevil for reaction of sulphide with monobromobimane (MBB, **24**) (Shen et al. 2011, 2012). Following alkylation with 2 MBB molecules, the resulting product, sulfide-bimane (SDB, **25**), is detected by reversed-phase HPLC with fluorescence detection (excitation 390 nm, emission 475 nm) (Fig. 10). This complex series of experiments requires many precautions to be taken, such as rigorous exclusion of oxygen, to achieve the claimed LOD of 5 nM for sulphide. Using this method, plasma levels of total H₂S were quantified at around 1.5 μM where free H₂S levels were in the range of 200 nM (Shen et al. 2012).

3 Probes Which Undergo Selective Reduction by H₂S

Certain functional groups can be reduced by H₂S to reveal a new functionality in a very rapid and selective manner. When the product function of this reductive process is part of a fluorescent molecule, the overall reaction sequence is an

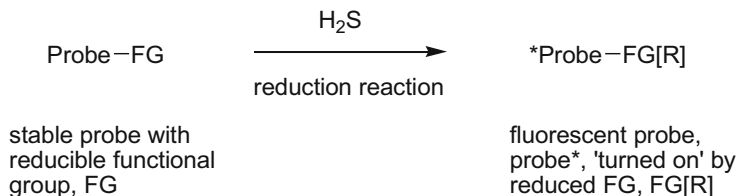


Fig. 11 Scheme showing reduction of a functional group (FG) with H₂S to give a fluorescent product

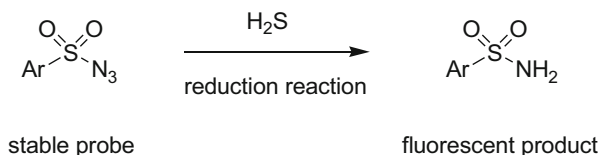


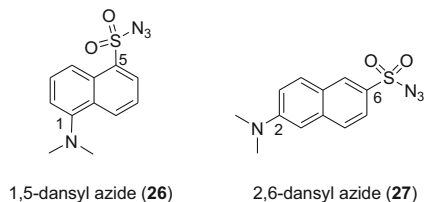
Fig. 12 General reduction of aromatic sulphonyl azides by H₂S gives fluorescent primary sulphonamide products

effective fluorescent turn-on signal upon reaction between the probe and H₂S (Fig. 11). These probe molecules are in most cases simpler structures than the other categories discussed herein, increasing the likelihood of easier access to the probe and better probe stability. The reduction of one functional group by H₂S is a well-known chemical reaction but rarely used in modern synthesis mostly due to the hazards of handling H₂S gas. However, for the sensitive detection of H₂S, this process is well suited. It can be viewed as a mild reductive process so only certain more susceptible functional groups would react rapidly and completely. One of the most popular groups in probe molecules is the azide (N₃). Usually azide is attached to a carbonyl, sulphonyl and/or aromatic system so that the resulting NH₂ product function is conjugated with the rest of the system, the overall product being a new fluorophore. For H₂S detection, azide directly attached to an aromatic system is by far the most studied. However, aromatic sulphonyl azide (ArSO₂N₃) undergoes rapid and complete reduction when exposed to H₂S in a wide range of media (Fig. 12) and will be discussed first.

3.1 Dansyl Sulfonyl Azides

Wang and co-workers have reported two sulphonyl azide probes based on the dansyl template. In this series, the first reported probe was 1,5-dansyl azide (**26**) (Fig. 13) (Peng et al. 2011). This probe has a very simple structure with a central naphthalene ring bearing only two substituents: a dimethylamino group and the reactive sulphonyl azide. **26** can be prepared easily in a single step from the sulphonyl chloride with sodium azide. The authors selected the sulphonyl azide group on the assumption that

Fig. 13 Dansylazide probes reported for specific use in H_2S quantification



the azide moiety would react rapidly with H_2S since it is activated by the electron withdrawing naphthylsulphonyl. This reaction does indeed take place with a lower detection limit of $1\ \mu\text{M}$ H_2S with signal:noise ratio of 3. Experiments were conducted in $20\ \text{mM}$ sodium phosphate buffer with $0.5\ \%$ tween-20. A 40-fold increase in fluorescence was recorded with $25\ \mu\text{M}$ sulphide (in the form of sodium sulphide, Na_2S). The response was seen to be linear at up to $100\ \mu\text{M}$ sulphide. Good selectivity was observed with over 18 anions studied at or above their physiological concentrations, and only HSO_3^- , $\text{S}_2\text{O}_4^{2-}$ and $\text{S}_2\text{O}_5^{2-}$ led to small responses. Thiols such as benzyl mercaptan did give stronger responses but still only one fifth of that of H_2S . **26** was also unreactive towards very high concentrations of glycine or lysine. In bovine serum, rapid and clear responses were obtained leading these authors to study the quantification of H_2S levels in mouse blood. This was found to be $31.9 \pm 9.4\ \mu\text{M}$; however, this level does not agree with other authors such as the work from the Kevil group who show in their sensitive method that plasma levels of free H_2S are well below $1\ \mu\text{M}$ (Shen et al. 2012). Although probe **26** appears to fulfil all the desirable criteria, there may be other reactions in blood which lead to an increase in signal. The same authors then published a variant of **26** with a different arrangement of the same two substituents. Probe **27**, 2,6-dansyl azide, is an isomer of **26**, but has a much higher quantum yield, 40-fold higher than **26** (Wang et al. 2014). This desirable property increases the sensitivity of **27** towards reduction by H_2S . Furthermore, the authors have shown that **27** does not require any added surfactant to boost the signal. Another benefit of **27** is its short and straightforward synthesis, well within the capabilities of most chemistry labs. When studied in only phosphate buffer (no added surfactant), **27** ($20\ \mu\text{M}$) detected H_2S from a sample of Na_2S ($10\ \mu\text{M}$) with a 100-fold increase in fluorescence within 1 h. The corresponding increase with **26** is reported as eightfold in phosphate buffer. Probe **27** shows a good overall anion selectivity as for **26**, including glutathione and cysteine at $100\ \mu\text{M}$. A linear response is seen in the reaction of **27** with H_2S at up to $10\ \mu\text{M}$ with a limit of detection of $1\ \mu\text{M}$ and S/N ratio of 3:1, in aqueous media. **27** gives a similar response in fetal bovine serum (FBS) indicating its promise in cellular applications.

3.2 Aromatic Azides

Direct attachment of azide to an aromatic group produces a fluorescent aniline product upon reduction by H_2S . Lippert et al. have reported two probes based on azide-caged rhodamines with either a carbamate (sulphide fluor-1, SF1, **28**) or urea

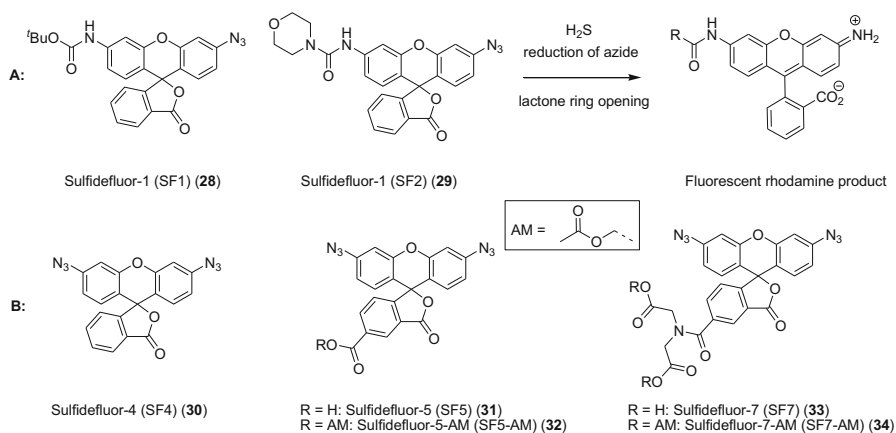
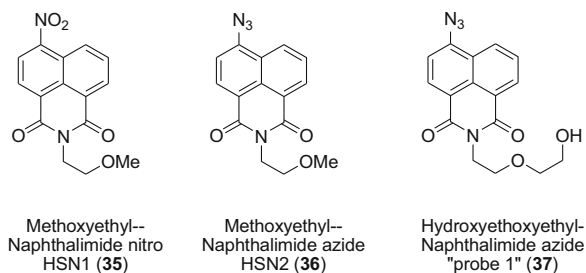


Fig. 14 Azide-caged rhodamine probes undergo lactone ring opening upon reduction by H₂S (**A**). Second-generation bis-azide rhodamines with intracellular hydrolysable ‘cell-trappable’ AM pro-drug (**B**)

(sulphide fluor-2, SF2, **29**) side chain (Fig. 14) (Lippert et al. 2011). **28** and **29** were synthesised in two steps from rhodamine 110, a commercially available starting material. These probes are described as ‘caged’ since the fluorescent rhodamine product is only revealed upon lactone ring opening triggered following azide reduction by H₂S. Following reaction of either probe (10 μM) with NaSH (100 μM) for 1 h in aqueous Hepes buffer at pH 7.4, excitation at 488 nm and emission at 525 nm produced a seven- to ninefold turn-on response. Under these conditions, the LOD was 5–10 μM. **29** was found to be preferred due to its higher selectivity for other reactive species such as glutathione (fivefold selective). In a live-cell imaging study in HEK293T cells, **28** (5 μM) was found to give a slightly stronger response to H₂S possibly due to its higher cLogP (1.98) compared to **29** (1.21). The Chang group have further developed their ‘SF’ probes with the report of bis-azides SF4 (**30**), SF5 (**31**), SF5-AM (**32**), SF7 (**33**) and SF7-AM (**34**) with enhanced sensitivity (Fig. 14) (Lin et al. 2013). In aqueous buffer (20 mM Hepes, pH 7.4), probes **30**, **31** and **33** (10 μM) produced a 40-, 4- and 20-fold increase in fluorescence intensity when incubated with NaSH (100 μM) for 1 h. From these probes, **30** displays the lowest LOD of 125 nM. These probes showed good selectivity against 100 μM of a range of reactive sulphur, nitrogen and oxygen species, 500 μM cysteine and 5 mM glutathione. In live HUVEC cells, **30**, **32** and **34** were compared with the first generation probe SF2 before and after replacing the cell media. Compounds with the ‘cell-trappable’ group, an acetoxymethyl (AM) ester, retained their brightness and penetrated the nucleus, whereas **29** and **30** did not. In addition **29** and **30** were only present in the cytosol. Bis-AM esters SF5-AM (**32**) and SF7-AM (**34**) underwent intracellular saponification to reveal a bis-carboxylic acid, with two negative charges, which was unable to exit the cell. In this way, the ‘AM’ probes are ‘trapped’ in the cell; hence, brightness is retained for

Fig. 15 Naphthalimide probes with methoxyethyl (HSN2) and hydroxyethoxyethyl ('probe 1') and side chains. Nitro derivative HSN1 also reduces to a fluorescent aniline in the presence of H₂S



a much longer period of time. SF7-AM was the most sensitive cell-trappable probe with signal still evident with only a 1 μM dose of NaSH.

Naphthalimide probes based on the Lucifer yellow family are attractive due to their synthetic accessibility and large Stokes shift. Three probes have been described in this family, two of them azides, with a methoxyethyl HSN2 (**36**) (Montoya and Pluth 2012) and hydroxyethoxyethyl (**37**) side chain (Guo et al. 2014; Montoya and Pluth 2012) (Fig. 15). A nitro derivative, HSN1 (**35**), capable of being reduced by H₂S to the fluorescent aniline, was also described, a rare example in probe structures. However, nitro **35** (5 μM) only experienced a 15-fold increase in fluorescence when exposed to 500 μM H₂S (50 mM PIPES, 100 mM KCl, pH 7.4) over 90 min, whereas with azide **36**, a 60-fold increase was observed in 45 min. Under these conditions, with the more sensitive **36**, the LOD is 1–5 μM . Furthermore, **36** is more selective: with 100 equiv. (500 μM) of reactive oxygen species and biological thiols, both probes are selective with S/N of about 7:1; however, with 10 mM of cysteine or glutathione, only **36** retains selectivity over a 60 min reaction time. In live HeLa cells incubated with both probes (5 μM) for 30 min, convincing increases in intracellular fluorescence were observed when H₂S (250 μM) was added. Interestingly, **36** gave a signal without any addition of H₂S, whereas nitro **35** did not despite being less selective towards biological thiols. Hydroxyethoxyethyl derivative ('probe 1', **37**), prepared in a two-step synthesis, when tested at a concentration of 10 μM (in 1:1 DMSO:20 mM Hepes buffer, pH 7.4) with 10 equiv. of Na₂S (100 μM), gave a 25-fold increase in fluorescence intensity following a 60 min reaction time (Guo et al. 2014). With **37** the LOD was 0.37 μM . Interestingly, **37** was very selective against cysteine (1 mM) and a wide range of other anions but less selective against sulphate (100 μM), iodide (1 mM) and chlorate (1 mM) with a fluorescence enhancement of about 9–12-fold in those cases. **37** was used to assess a series of water samples, including mineral, lake and tap water, for their H₂S content. Spiked samples with 200, 300 and 400 ppb of added Na₂S were detected in all the samples with good recoveries. Lake water had the highest H₂S content with 17 ppb. No cellular experiments were reported with **37**.

Tang has reported two diphenylethene-based probes, DPE-Az (**38**) and TPE-Az (**39**) (Fig. 16) which operate by a photostable process known as aggregate-induced emission (AIE) (Kwok et al. 2014), unlike other reported probes (Cai et al. 2014). Both probes become aggregates at 10 μM concentrations in a 2:8 mixture of DMSO:Hepes buffer. When 5 equiv. of NaSH are added to **39** (10 μM), after

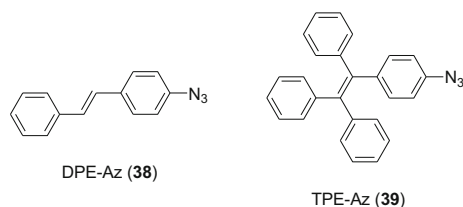


Fig. 16 Diphenylethene (DPE) and tetraphenylethene (TPE) probes based on aggregation-induced emission (AIE)

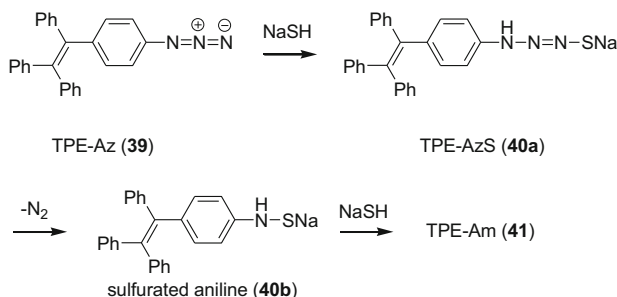


Fig. 17 Proposed mechanism of reaction of TPE-Az with NaSH explaining why more than 1 equiv. of H₂S is required before a signal can be detected

only a 5 min incubation, a 60-fold increase in emission at 493 nm was recorded (340 nm excitation). This reaction is complete after only 2 min, much faster than many other probes. Furthermore, the preferred probe, **39**, is selective for H₂S over other species such as cysteine (1 mM) and glutathione (5 mM). Importantly, **39** does not give a signal until *more than one equivalent* of H₂S has been added to the system. To explain this unusual phenomenon, the authors propose that one molecule of HS⁻ adds to the azide, to give TPE-AzS (**40a**), which then undergoes loss of nitrogen to form a sulfurated aniline (**40b**), which is then reduced to fluorescent TPE-Am (**41**) by a second equivalent of HS⁻ (Fig. 17). By varying the amount of the probe, the level where the signal starts will vary. In other words just by using different concentrations of **39**, the point at which a signal is obtained means that more than 1 equiv. of H₂S has been detected. In this way, **39** can directly quantify H₂S *without a separate calibration being required*. However, the authors only studied a lower concentration of 10 μM of **39** and did not report any studies in cells.

Han and co-workers reported a cyanine-based near-infrared fluorescence probe, Cy-N₃ (**42**), based on a strategy of internal charge transfer (ICT) (Fig. 18) (Yu et al. 2012). They reasoned that a change of substitution on a cyanine dye scaffold, upon reduction of azide to amine, would shift the absorption/emission wavelength. Indeed when **42** (10 μM) was treated with NaSH (100 μM) in Hepes (40 mM, pH 7.4) within 20 min, the NIR fluorescence maximum changed from 710 to 750 nm, following excitation at 625 nm. The increase in emission intensity

Fig. 18 Structure of near-infrared cyanine fluorescence probe, Cy-N₃, selective for H₂S

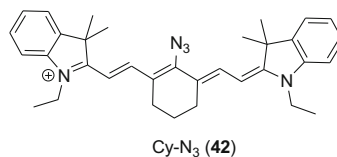
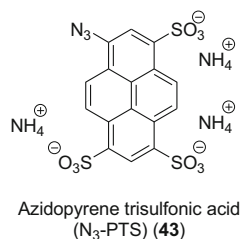


Fig. 19 Structure of an azidopyrene-based fluorescence probe selective for H₂S



was twofold with an excellent LOD of 80 nM. This represents one of the most sensitive fluorescence probes reported to date. In addition **42** was selective for 24 different anions and reactive oxygen, nitrogen and sulphur species, including cysteine and glutathione at 1 mM concentrations. Intensities from the most challenging species were still only 10 % of the sulphide response. This probe (10 μ M) was able to quantify H₂S from 0.08 to 100 μ M, with a linear response in both water and fetal bovine serum. Detection of H₂S released from a small molecule donor, ADT-OH (Li et al. 2007) (500 μ M), with **42** (10 μ M) was achieved in fetal bovine serum (40 mM Hepes, pH 7.4) at 37 °C. Release from ADT-OH at this concentration was equivalent to NaSH at 100 μ M. Moreover **42** was sensitive and selective enough to detect H₂S in cells. RAW264.7 cells (mouse macrophages) were incubated with 100 μ M H₂S then 10 μ M **42** for 20 min at 37 °C. Quantifiable increases in intracellular H₂S were clearly visualised. In summary **42** has been extensively validated as a sensitive, selective and cell-applicable fluorescence probe for the quantification of H₂S.

Highly water-soluble probe N₃-PTS (**43**), based on a pyrene-1,3,6-trisulfonic acid scaffold, was prepared in one step from known fluorescent dye APTS (Fig. 19) (Hartman and Dcona 2012). As a tris-ammonium salt, **43** has water solubility of >100 mM. Following addition of NaSH (50 μ M) to **43** (100 μ M), a sevenfold increase in fluorescence intensity (emission at 505 nm, excitation at 435 nm) was detected after 90 min of reaction in PBS buffer. When compared with NaSH (50 μ M), **43** gave at least a fivefold greater fluorescence intensity compared with a range of other reactive species, including cysteine and glutathione at 1 mM. A linear response is obtained when **43** (100 μ M) is reacted with NaSH from 2 to 100 μ M, suggesting the LOD is around 2 μ M or less. In fetal bovine serum, very little fluorescence quenching was observed and the concentration of NaSH required to increase the signal by twofold was 9 μ M. Generally the H₂S sensitivity in FBS was about 20 % that of PBS, and the authors suggested that new derivatives of **43**

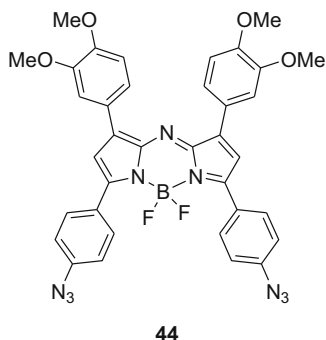


Fig. 20 Aza-BODIPY bis-aromatic azide dye

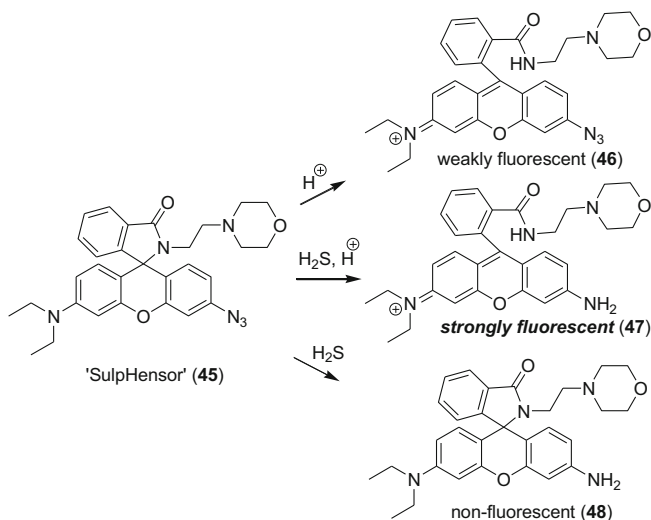


Fig. 21 Rhodamine probes modified with a spirolactam (acid sensitive) and an aromatic azide (H₂S sensitive) and a lysosome-targeting morpholine group

could increase the serum sensitivity. No cellular studies were carried out with this probe.

An aza-BODIPY dye (**44**) has been reported by Ramaiah and co-workers (Fig. 20) (Adarsh et al. 2014). Reaction of **44** (12 μM) in THF with 2 equiv. of Na₂S in water was complete in only 30 s and was selective for other ions; however, data on cysteine or glutathione were not reported. No data in cells was reported.

Yang and co-workers have devised a dual-mechanism acid-sensitive probe, 'SulpHensor' (**45**) targeted to acidic lysosomes (pH 4–6) in cells (Fig. 21) (Yang et al. 2014). **45** can react separately, and reversibly, with either acid or H₂S leading to separate products with little fluorescence (**46/48**). However, when both events

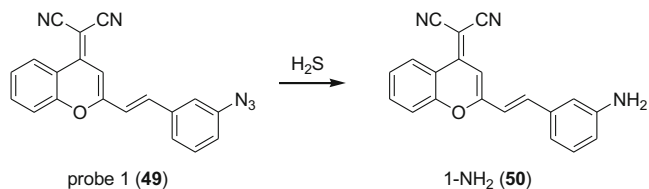


Fig. 22 Conjugated benzopyranazide probe **1** with two-photon absorbing properties undergoes reduction with H₂S to give fluorescent product 1-NH₂

happen, a strongly fluorescent product is formed (**47**). Hence in cells, **45** only visualises H₂S in acidic organelles. The rapid acid/base equilibration to give a nonfluorescent product at neutral pH was designed to lead to a clear selectivity within cells for the lysosomal compartment. **45** (5 μM) gave a 15-fold increase in fluorescence when reacted with NaSH (50 μM) at pH 4.5. With other reactive species, **45** was quite selective with the least being glutathione (5 mM) with an approximate 7:1 S/N ratio. In HeLa cells, **45** (5 μM), with a 30 min pre-incubation, was confirmed to detect H₂S (50 μM) selectively in lysosomes. Boosting endogenous production of H₂S with cysteine pretreatment in HeLa cells was confirmed with **45** demonstrating the specific useful applications of this probe.

Two-photon microscopy, as opposed to one-photon of high energy, utilises two photons of low energy to obtain the excited state of a pharmacophore, potentially allowing deeper tissue penetration of the probe (Kim and Cho 2009). Peng and co-workers developed two-photon probe **49**, which reacts with H₂S to give fluorescent 1-NH₂ (**50**) in the NIR wavelength range (Fig. 22) (Sun et al. 2013). Unlike other NIR probes, **49** was shown to be photostable when subjected to a 500 W lamp for 4 h. When a solution of **49** (5 μM) in 100 mM pH 7.4 PBS buffer, with 50 % DMSO, was reacted at 37 °C with NaSH (50 equiv.), a 65-fold fluorescence turn-on response was observed. **49** was generally selective for other anions with homocysteine being the least selective at about 20:1 S/N ratio. A pH range from 2.5 to 10 was found to be tolerable by this probe, and it could be used in serum (FBS) as well. Incubation of MCF-7 cells with **49** (5 μM) at 37 °C in PBS with NaSH (50 equiv.) for 60 min gave a strong signal with emission 575–630 nm. **49** was also tested in live animals, the first reported probe to have been studied in vivo. Using the skin-popping methodology, **49** (40 μM) and NaSH (25 equiv.) were administered to ICR mice. At the 4 h timepoint, a clear increase in signal was seen compared to treated groups indicating the in vivo applicability of the two-photon strategy.

3.3 Aliphatic Azides

In contrast to the aromatic azides, aliphatic azides are not directly conjugated to an extended chromophore system. However, they are still reduced efficiently to aliphatic amines which are then good nucleophiles for intramolecular processes

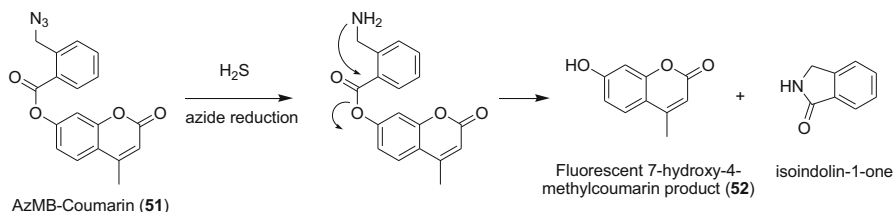


Fig. 23 Aliphatic azide in AzMB-coumarin is reduced by H₂S triggering an intramolecular cyclisation and ejection of fluorescent coumarin

which ultimately result in a fluorescent product. An elegant example is described by Wu et al. where probe molecule AzMB-coumarin (**51**) undergoes reduction followed by intramolecular attack on the nearby ester ejecting fluorescent coumarin (**52**) and the innocuous by-product isoindolin-1-one (Fig. 23) (Wu et al. 2012). The reaction is complete within 20–60 min either in acetonitrile or 20 % v/v sodium phosphate (pH 7.4, 100 mM) buffered acetonitrile. Control of pH is important since the hydrosulphide ion, HS⁻ (pK_a of H₂S is c.6.9), is present as the major species at pH 7.4. Using **51** (20 μM) and NaSH, following a 1 h incubation with excitation at 365 nm and emission at 450 nm, the LOD was 100 μM in aqueous acetonitrile and 10 μM in acetonitrile. **51** (100 μM) is selective over biological thiols such as glutathione and cysteine (1 mM) with an S/N ratio of about 6:1. In HeLa cells treated with **51** (1 mM) for 10 min then treated with or without NaSH (1 mM), a clear signal was seen with the treated cells visualised by fluorescence microscopy.

3.4 Probes Based on Reduction of a Nitro Group

Hydrogen sulphide can also reduce nitro groups to anilines; however, when the nitro is part of an aromatic phenol, a thiolytic process occurs, where the resulting aniline product is cleaved to reveal a fluorescent phenol. Using this principle, a three-channel-based fluorescent probe **53** (Fig. 24) was developed by Lin and co-workers in 2013 (Wang et al. 2013). This white light-emitting probe was constructed by connecting a blue fluorescent dye, 1*H*-phenanthro[9,10-*d*]imidazole, and an ESIPT (excited-state intramolecular proton transfer) component, 3-hydroxychromone. The reactive site was modified with a 2,4-dinitrophenyl group. When exposed to NaSH, **53** reacts through the dinitrophenol carbon ejecting fluorophore **54** responding with 3-, 6- and 16-fold emission enhancements at three channels around 440, 510 and 570 nm. **53** is stable over a pH range of 5.5–8.5 with a LOD of 10 μM at pH 7.4 in PBS:DMSO (v/v, 4/1) with 0.5 % tween-20. Selectivity of **53** (5 μM) for H₂S is good for a range of anions (S/N up to 20:1), wherein cysteine (1 mM) and glutathione (10 mM) responses were two- to threefold higher than for other anions (S/N 5:1). An MG63 cell line study indicated that **53** is not toxic at concentrations below 50 μM. Application of **53** (5 μM) for endogenous H₂S

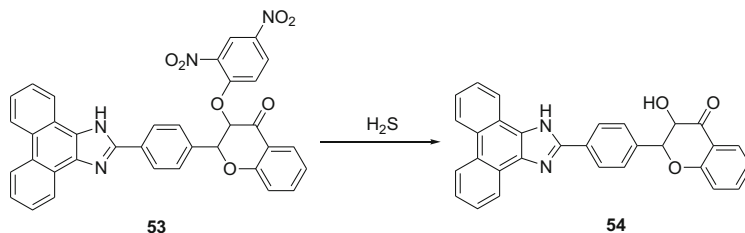


Fig. 24 Structure of probe **1** and its mechanism

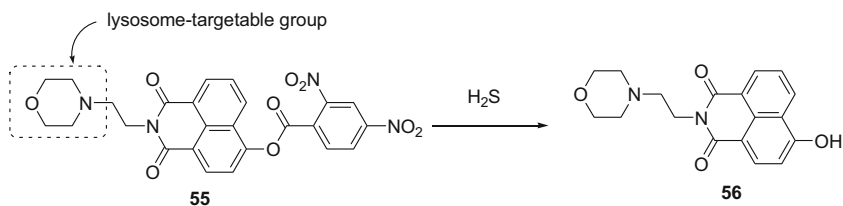


Fig. 25 Structure of probe **3** and its mechanism

(50 μM) detection was carried out in MG63 cells. Bright fluorescence images of blue, green and red colours were obtained, featuring the first three-channel sensor for hydrogen sulphide. However, incubation of **53** with cysteine and GSH also led to bright fluorescence images at three channels.

A 1,8-naphthalimide-derived fluorescent probe **55** (Fig. 25) for H_2S based on the thiolysis of a dinitrophenyl benzoate ester, to produce fluorescent **56**, was developed by Cui and co-workers in 2013 (Liu et al. 2013). **55** is applicable to H_2S detection over a pH range of 4–8 in bovine serum with a LOD of 0.48 μM . The response of 10 μM **55** to 1 mM NaHS was 12-fold of control and sixfold of that to 1 mM cysteine and GSH in less than 20 min. **55** was also shown to be suitable for H_2S detection in live cells with minimal toxicity. Because of the presence of a lysosome-targetable group, 4-(2-aminoethyl)morpholine, **55** is able to specifically localise in lysosomes at 5 μM and detect H_2S in lysosomes in live cells.

Lin and co-workers have created dinitrophenol ether attached to a BODIPY core (Fig. 26) (Cao et al. 2012). The resulting probe, NIR- H_2S (**57**) (5 μM), has an 18-fold turn-on response when reacted with NaSH (40 μM) in 50 mM PBS buffer with 3 mM cetyltrimethylammonium bromide and 10 % ethanol. **57** was selective with a range of other reactive species, the most challenging being cysteine (1 mM) and glutathione (10 mM), where the S/N was approximately ten- and sevenfold, respectively. In MCF-7 cells treated with **57** (5 μM) for 20 min then treated with NaSH (0–100 μM) for 10 min clear images were obtained showing H_2S in the cells, primarily in the mitochondria.

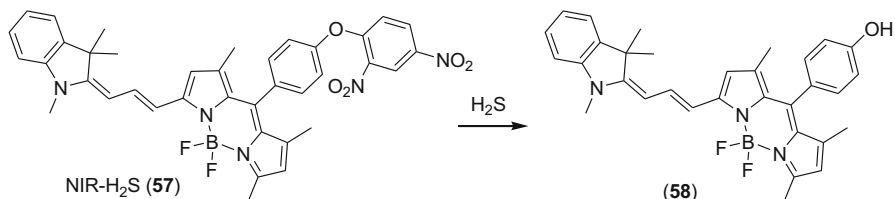


Fig. 26 Dinitrophenylether NIR-H₂S is reduced by H₂S triggering an intramolecular cyclisation and ejection of fluorescent coumarin

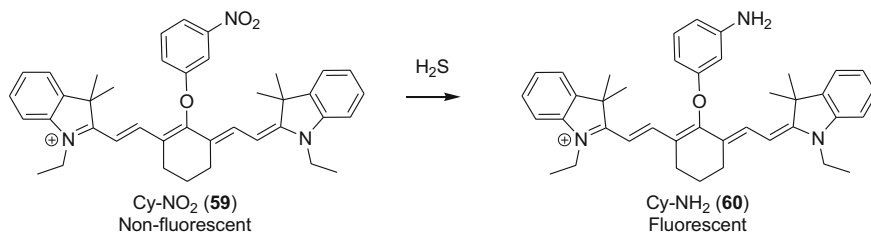


Fig. 27 Nitrophenylether-substituted cyanine dye Cy-NO₂ is reduced by H₂S to a fluorescent aniline product, Cy-NH₂

Earlier, a cyanine dye-based probe, **42** (Cy-N₃), was discussed (Fig. 18) where an aromatic azide was reduced by H₂S to give a fluorescent product. A nitro reduction design of a similar cyanine probe, Cy-NO₂ (**59**), was described by Chen and co-workers (Fig. 27) (Wang et al. 2012). Probe **59** was not affected by pH in the range 4.2–8.2 and operated well under physiological conditions in the NIR range (ex/em 755/789). Hence, **59** (10 μM) in 40 mM Hepes buffer at 37 °C gave a 12.7-fold fluorescence increase when treated with Na₂S (350 μM). Selectivity was also good with cysteine and glutathione being the most reactive, but still with a very good window. **59** was also effective in FBS which led on to studies in RAW264.7 cells. Cells were treated with **59** (10 μM) for 10 min, washed then treated with Na₂S (50–350 μM) then imaged by confocal fluorescence microscopy. A clear dose response was obtained with brighter images at higher concentrations of Na₂S. Further studies revealed that **59** is predominately located in the cytoplasm.

4 Probes Which Trap H₂S Using a Coordinated Metal

A fluorescein-based probe conjugated with an azamacrocyclic copper complex has been reported to specifically react with H₂S through the copper centre resulting in an increase in fluorescence (Fig. 28). Cyclen (1,4,7,10-tetraazacyclododecane,

Fig. 28 Fluorescein conjugated with an azamacrocyclic ‘cyclen’ copper complex. When H₂S binds the Cu²⁺, the copper is released resulting in an increase in fluorescence

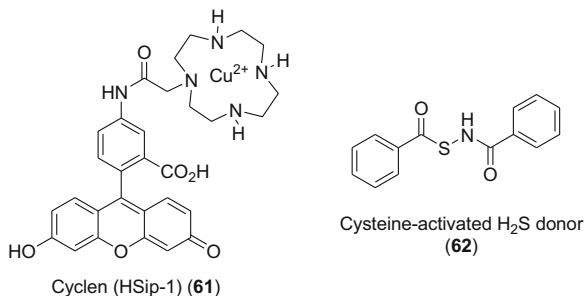
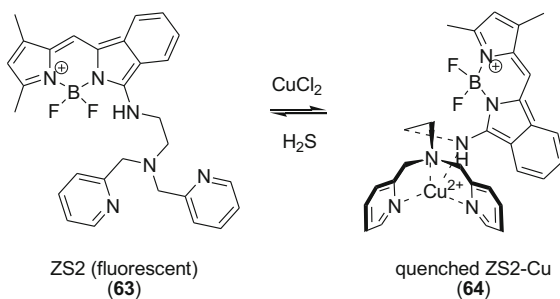


Fig. 29 *N*-Bis [2-pyridylmethyl] aminoethene attached to BODIPY (ZS2) forms a copper complex ‘quenched ZS2-Cu’. When H₂S binds the Cu²⁺, the copper is released resulting in an increase in fluorescence



HSip-1, **61**) was found to be the most sensitive and selective of four different macrocyclic ring sizes studied (Sasakura et al. 2011). At 10 μ M concentration, **61** gave a 50-fold turn-on response (ex/em 491/516 nm) when treated with NaSH (100 μ M) in 30 mM Hepes buffer, whereas with high concentration of GSH (10 mM), almost no increase in fluorescence was detected. When present together, response to NaSH was not affected by 10 mM glutathione. With added cysteine (1 mM), a >10-fold S/N ratio was achieved; all other tested anions were even more selective. **61** (1 μ M) was also tested in vitro with a cysteine-activated H₂S donor drug (50 μ M) (Zhao et al. 2011). In HeLa cells **61** was impermeable, so it had to be diacetylated, then it was able to image exogenously added H₂S (200 μ M).

Probes which undergo nucleophilic reaction with H₂S are often too slow to be useful in cells. For example, the reactivity of the aldehyde–acrylate reaction (see Fig. 2) takes 50 min to reach maximum fluorescence (Li et al. 2013). Another disadvantage is their poor water solubility requiring non-biocompatible solvents. Taking advantage of the strong affinity between sulphur and copper has produced probes such as **61** (Sasakura et al. 2011) and Cu-1 (Qu et al. 2013). A new probe, ZS2-Cu (**64**), has good water solubility (1.28 mg/ml) and membrane permeability and is highly sensitive (Fig. 29) (Li et al. 2014). BODIPY-based **63** is complexed with Cu²⁺ through a bis-pyridylethylenediamine group. BODIPY is membrane permeable and the bis-pyridyl feature provides good water solubility. In fact

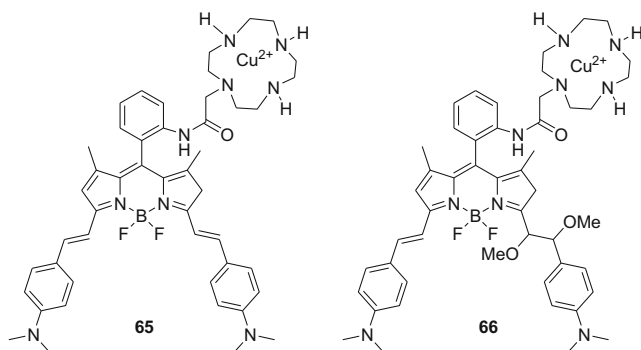


Fig. 30 Cyclen copper complexes, probe **7** and **8**, attached to BODIPY modified with substitutions to create an NIR-fluorescent probe upon complexation of Cu with S

experiments are carried out purely in aqueous solution without the need for an organic co-solvent. Fluorescence turn-off response of this probe was specific to copper; a range of 10 other cations did not quench the fluorescence of **63** even when mixed in the presence of copper. **64** exhibited over 30-fold selectivity towards H₂S over other anions and biothiols such as cysteine and glutathione and a 15–18-fold increase in fluorescence when NaSH was added in the presence of other anions or biothiols. When 2 equiv. of NaSH was added to **64** (10 μM), a 19-fold increase in fluorescence intensity was measured at 546 nm. The reaction occurs very rapidly with a lower limit of detection of 250 nM. In HeLa cells incubated with **64** (5 μM) for 15 min, weak fluorescence was observed, but when cells were washed and NaSH (100 μM) added, a strong increase in intracellular signal was observed confirming the good membrane permeability of **64**.

Among another series of BODIPY-based probes published recently appears to be a rare example of *in vivo* application of H₂S sensing using a fluorescence probe. Huang and co-workers have designed near IR (NIR) fluorescent probes **65** and **66** (Fig. 30) with excitation at 680/600 nm and emission at 680/765 nm, respectively (Wu et al. 2014). **65** was selective for reactive species with the lowest selectivity for 1 mM cysteine (about 10:1 S/N) and 10 mM glutathione (about 20:1 S/N). The LOD for **65** was 80 nM with a limit of quantitation (LOQ) determined at 0.27 μM. In RAW264.7 cells **65** (10 μM) was not cell penetrant and hence required liposomes as a carrier to achieve cell penetration. However, this was effective and led to an eightfold fluorescence enhancement upon exogenous application of Na₂S (200 μM). Probe **66**, also using the liposome formulation, was shown to be effective in MCF-7 cells and was able to detect H₂S evolved from diallyltrisulphide. In mice, **65** (40 μM) was studied using a skin-popping technique to administer the compound into the tissue near the skin. A dose of Na₂S (25 equiv.) was injected and detected with a sevenfold increase in fluorescence within 5 h.

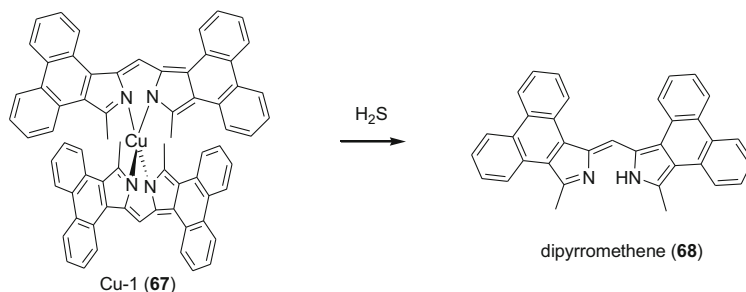


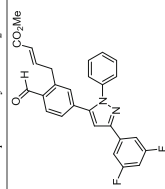
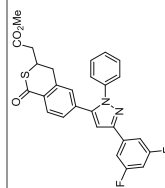
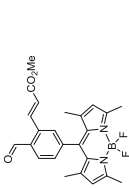
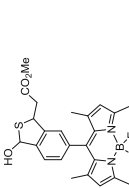
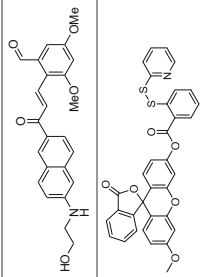
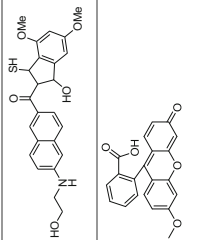
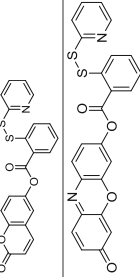
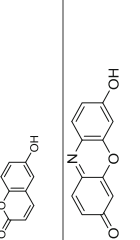
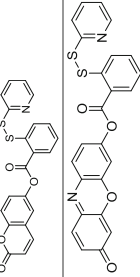
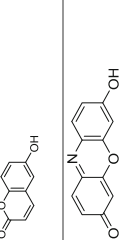
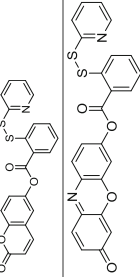
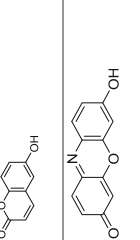
Fig. 31 Bis-phenanthrene-fused dipyrromethene complex, Cu-1, reacts with H₂S to give fluorescent product **1** with fluorescence in the NIR range

Another NIR probe based on the favourable copper-sulphide affinity, with potential for deep tissue application *in vivo*, was reported by Shen and co-workers (Fig. 31) (Qu et al. 2013). Dipyrromethene-based copper complex, Cu-1 (67), reacts instantly with H₂S (1.5 equiv.) in DMSO:water (1:1) solution to liberate fluorescent 68 with a 14-fold increase in fluorescence intensity. There is some interference when cations such as Zn²⁺ are added where the reaction rate slows down to about 5 h. However, good selectivity is seen with other reactive sulphur species with glutathione (10 equiv.) still giving an 8:1 S:N ratio. In HeLa cells, 67 (50 μM) treatment for 30 min with addition of NaSH (50 μM) gave bright images and indicated that the fluorescence product was located in the cytoplasm. In addition, 67 was found to be nontoxic up to 100 μM concentrations assessed by MTT assay.

5 Conclusions

There are now many diverse, sensitive fluorescent probes reported for H₂S detection and quantification with a particular acceleration in the past few years highlighting the upsurge in interest in H₂S biology in general. Table 1 summarises selected probes showing the range of detection limits (some down to low nanomolar), detection wavelengths, reaction times and conditions and application in a biological system. There are now many probes which have some utility in living cells and several with specific subcellular localisation. Biologists studying H₂S now have a wide range of tools to assist them which should lead to yet further progress in the understanding of the role of H₂S in biology and how to apply this knowledge therapeutically.

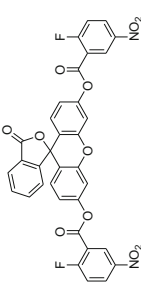
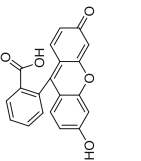

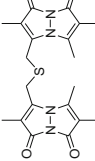

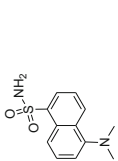
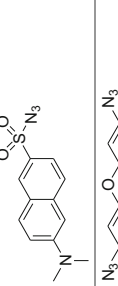
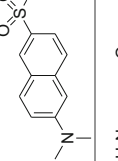
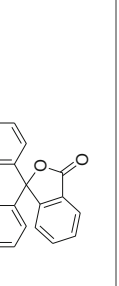
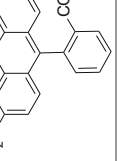
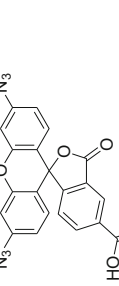
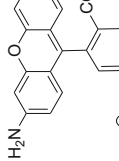
Table 1 Selected H₂S-sensitive fluorescent probes listed according to mechanism: probes which react nucleophilically with H₂S, probes which are reduced by H₂S and probes which involve metal coordination

Probe	Probe name	Probe structure	Fluorescent product	LOD (μM)	Wavelength (EX/EM, nm)	Reaction time (mins)	pH	Medium	Application in biosystem	References
		<i>Probes which react nucleophilically with H₂S</i>								
1	SFP1			<5	310/550	20	7.4	PBS	HeLa cells	Qian et al. (2011)
3	SFP2			<5	465/480–580	20	7.4	PBS	HeLa cells	Qian et al. (2011)
5	P3			0.05	375/505	5	7.4	Hepes, 10 % FBS	HeLa cells	Singha et al. (2015)
9	WSP1			0.06	476/516	60	7.4	10 mM PBS/MeCN (1:1) + Hexadecyltrimethylammonium bromide (CTAB)	COS7 cells	Liu et al. (2011), Peng et al. (2014)
10	WSP2			0.079	385/456	5	7.4	10 mM PBS/MeCN (1:1) + Hexadecyl trimethyl ammonium bromide (CTAB)	–	Peng et al. (2014)
11	WSP3			0.047	550/586	5	7.4	10 mM PBS/MeCN (1:1) + Hexadecyl trimethyl ammonium bromide (CTAB)	–	Peng et al. (2014)

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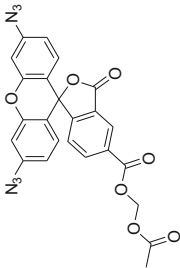
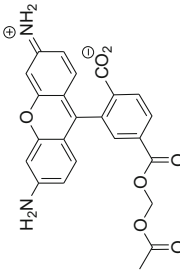
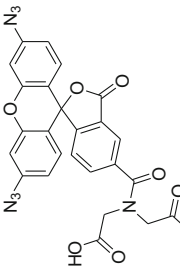
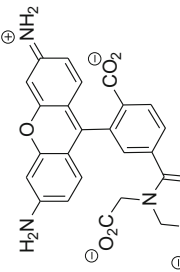
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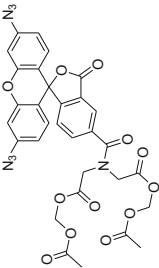
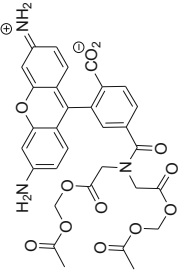
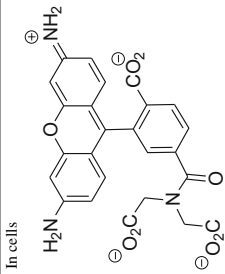
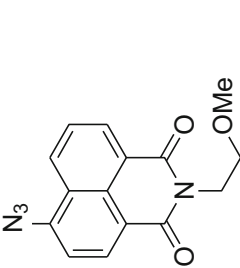
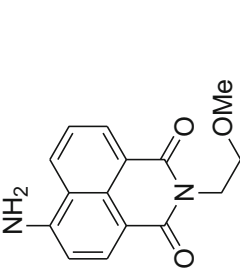
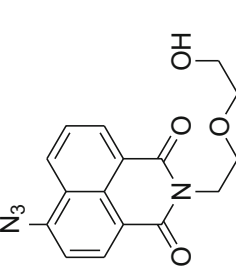
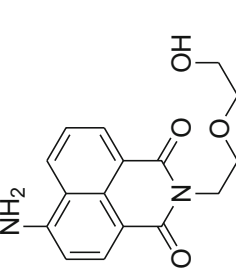
Probe	Probe name	Probe structure	Fluorescent product	LOD (μM)	Wavelength (EX/EM, nm)	Reaction time (mins)	pH	Medium	Application in biosystem	References
12	WSP4			0.266	512/531	5	7.4	10 mM PBS/MeCN (1:1) + Hexadecyl trimethyl ammonium bromide (CTAB)	HeLa cells	Peng et al. (2014)
13	WSP5			0.047	502/525	5	7.4	10 mM PBS/MeCN (1:1) + Hexadecyl trimethyl ammonium bromide (CTAB)	HeLa cells	Peng et al. (2014)
14	-			2.5	517	30	7.4	EtOH/PBS 1/1 (V/V)	HeLa cell	Liu et al. (2014b)
16	E1			0.12	295/487	5	7.4	Tris-HCl buffer	HeLa cells	Xu et al. (2012)
19	-			1	517	30	7.4	PBS	COS-7 cells	Liu et al. (2012)
20	-			0.1	517	40	7.4	PBS/MeCN (1:1, v/v)	COS7 cells	Zhang et al. (2013)

23	DSP3			0.07	517	10	7.4	PBS	HeLa cell	Liu et al. (2014a)
24	MBB			0.005	390/475	30	9.5	100 mM Tris-HCl buffer	Can measure plasma H ₂ S levels. Clinical trial on-going	Shen et al. (2011, 2012)
<i>Aromatic azide probes which are reduced by H₂S to produce fluorescent aniline products</i>										
26	1,5-Dns-Az			1	340/535	60	7.5	20 mM sodium phosphate buffer with 0.5 % tween-20	-	Peng et al. (2011)
27	2,6-Dns-Az			1 (6 μM in FBS)	325/450	60	7.4	PBS	-	Wang et al. (2014)
30	SF4			0.125	496/517	60	7.4	20 mM Hepes	-	Lin et al. (2013)
31	SF5			-	498/521	60	7.4	20 mM Hepes	-	Lin et al. (2013)

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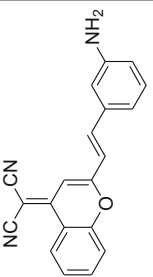
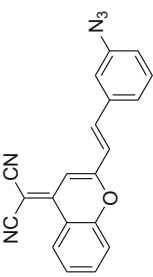
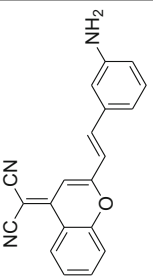
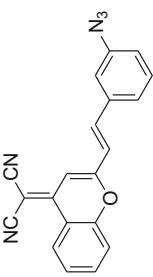
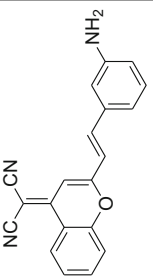
Probe	Probe name	Probe structure	Fluorescent product	LOD (μM)	Wavelength (EX/EM, nm)	Reaction time (mins)	pH	Medium	Application in biosystem	References
32	SF5-AM		 In cells	0.25	488/525	60	7.4	20 mM HEPES	HUVECs	Lin et al. (2013)
33	SF7			–	498/526	60	7.4	20 mM HEPES	–	Lin et al. (2013)

34	SF7-AM		  <p>In cells</p>	0.5	488/525	60	7.4	20 mM Hepes	HUVECs	Lin et al. (2013)
36	HSN2			1-5	435/490	45-60	7.4	50 mM PIPES, 100 mM KCl	HeLa cells	Montoya and Pluth (2012)
37				0.37	440/534	15	7.4	1:1 DMSO:20 mM Hepes	Tap, mineral and lake water. No cellular experiments	Guo et al. (2014)

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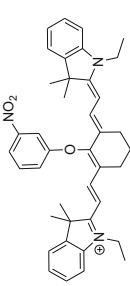
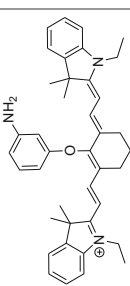
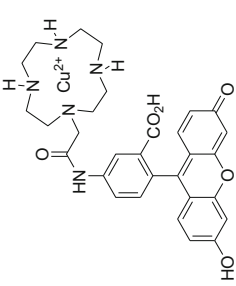
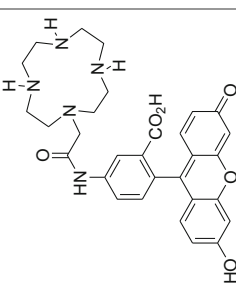
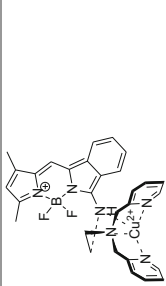
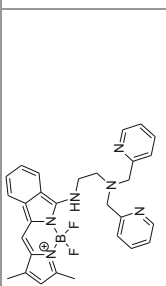
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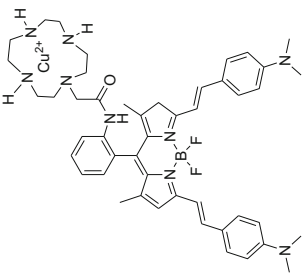
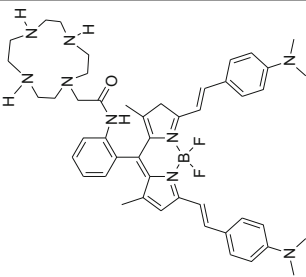
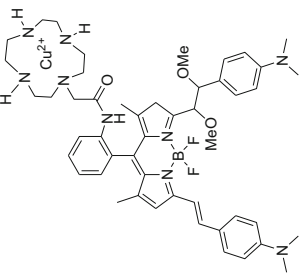
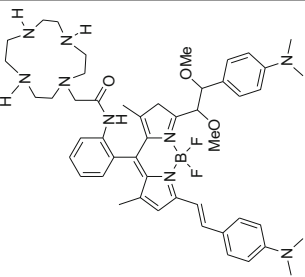
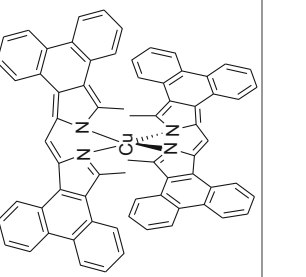
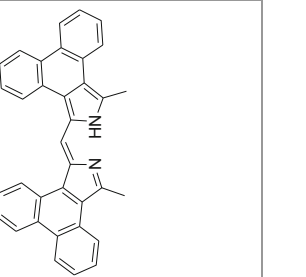
Probe	Probe name	Probe structure	Fluorescent product	LOD (μM)	Wavelength (EX/EM, nm)	Reaction time (mins)	pH	Medium	Application in biosystem	References
39	TPE-Az			–	340/493	5	7.4	Hepes:DMSO (4:1)	–	Kwok et al. (2014)
42	Cy-N ₃			0.08	625/750	20	7.4	40 mM Hepes	RAW264.7 macrophage cells	Yu et al. (2012)
43	N ₃ -PTS			2	435/505	90	7.4	PBS	–	Hartman and Deona (2012)
44	Aza-BODIPY			0.5	700/750	0.5	7.0	THF-water	–	Adarsh et al. (2014)
45	SulphSensor			0.5	530/555	60	4.0–6.0 (lysosome targeted)	Britton-Robinson buffer with 10 % DMF	HeLa cells Lysosome targeting	Yang et al. (2014)

49	Probe 1		3.05	520/670	60	7.4	100 mM PBS buffer in 50 % DMSO	MCF-7 cells	Sun et al. (2013)
51	AzMB-Coumarin		10 (pure MeCN)	365/450	60	7.4	20 % 100 mM sodium phosphate buffered acetonitrile	HeLa cells	Wu et al. (2012)
53			10	440 510 570	30	5.5-8.5	Buffer/tween	MCF63 cell lines	10
55			0.48	555	10	4-6	Bovine serum	MCF7 cells Lysosome targeting	8
57			0.05	650/708	<10	7.0	50 mM PBS buffer (pH 7.0) with 3 mM cetyltrimethylammonium bromide and 10 % ethanol	MCF-7 cells	Cao et al. (2012)

(continued)

Table 1 (continued)

Probe	Probe name	Probe structure	Fluorescent product	LOD (μM)	Wavelength (EX/EM, nm)	Reaction time (mins)	pH	Medium	Application in biosystem	References
59	Cy-NO ₂			<50	755/789	60	4.2–8.2	40 mM HEPES, pH 7.4, 37 °C	RAW264.7 cells. Probe located in the cytoplasm	Wang et al. (2012)
<i>Metal coordinating probes</i>										
61	HSip-1				491/516	<5	7.4	30 mM HEPES buffer	HeLa cells	
64	ZS2-Cu			0.25	480/546	15	7.4	PBS	HeLa cells	Li et al. (2014)

65	Probe 7			0.08 (LOQ 0.27)	680/765	15	7.4	10 mM PBS buffer with 0.5 % tween-20	RAW264.7 cells and in mice using skin-popping technique	Wu et al. (2014)
66	Probe 8			0.08	600/680	15	7.4	10 mM PBS buffer with 0.5 % tween-20	MCF-7 cells	Wu et al. (2014)
67	Cu-I			1	540/600	<1	7.0	1:1 DMSO:water	HeLa cells	Qu et al. (2013)

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The Pharmacological Effects of S-Propargyl-Cysteine, a Novel Endogenous H₂S-Producing Compound

Ya-Dan Wen and Yi-Zhun Zhu

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Abstract

S-propargyl-cysteine (SPRC), also named as ZYZ-802, is a structural analog of S-allylcysteine (SAC), the most abundant constituent of aged garlic extract. SPRC becomes a derivative of the amino acid cysteine, which contains sulfur atom, by changing allyl group in SAC to propargyl group in SPRC. Another

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analog of SPRC and SAC is *S*-propyl cysteine (SPC), which has propyl group instead in its cysteine structure. Drug formulation of SPRC has been investigated in the mixture of extenders, such as lactose, microcrystalline cellulose, and cross-linked povidone, showing good fluidity and scale-up production possibility. Controlled release formulation of SPRC (CR-SPRC) and leonurine-SPRC were invented and shown the decent pharmacological effects in heart failure and hypoxia injury, respectively. The pharmacological effects of SPRC have been shown that cardioprotection and proangiogenesis in several ischemic heart models, neuroprotection in Alzheimer's disease, proapoptosis in gastric cancer and anti-inflammation in acute pancreatitis. Moreover, CR-SPRC reduced infarct size and recovered partial cardiac function in heart failure rat model. Leonurine-SPRC protected hypoxic neonatal rat ventricular myocytes in much lower dose. Interestingly, since the propargyl group in SPRC has the stronger chemical bond in the cysteine structure than allyl group in SAC and propyl group in SPC, SPRC showed more extensive cardioprotection in ischemic rat hearts model compared to SAC and SPC. The mechanisms of pharmacological effects of SPRC have been unveiled that SPRC reduced Ca^{2+} accumulation, activated antioxidants, inhibited STAT3, decreased inflammatory cytokines, and elevated p53 and Bax. More pharmacological effects and mechanisms of SPRC will be discovered in atherosclerosis, hypertension, and other diseases.

Keywords

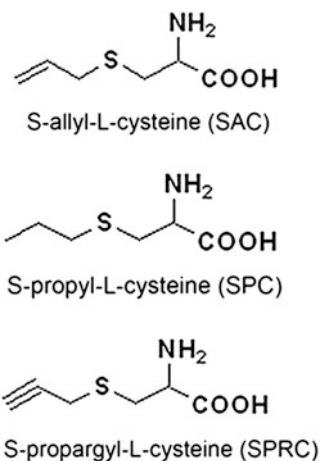
S-propargyl-cysteine • SPRC • S-allylcysteine • S-propyl cysteine • Garlic extraction • Hydrogen sulfide • ZYZ-802

1 General Introduction

Garlic is an ancient food consumed by human for thousands of years. This food is considered as a beneficial diet due to the characteristics of antioxidant, hepatoprotection, neuroprotection, and H_2S production (Butt et al. 2009). The major compounds containing sulfide in garlic are γ -glutamyl-*S*-allyl-L-cysteines and *S*-allyl-L-cysteine sulfoxides (alliin) (Amagase 2006). A major transformed chemical from γ -glutamyl-*S*-allyl-L-cysteines is *S*-allylcysteine (SAC), which is a sulfide amino acid presenting cellular (Gupta and Rao 2007), neural (Chauhan 2006), and cardiovascular protection (Shin et al. 2007). The structural analogs of SAC include *S*-propargyl-cysteine (SPRC) and *S*-propyl cysteine (SPC), which contain the same cysteine structure and only differ in the propargyl and allyl moiety, respectively, which were anticipated to have stronger pharmacological effects. Three cysteines all present similar H_2S effects through generating additional H_2S by being potential substrates of endogenous cysteine-metabolizing enzymes (Wang et al. 2009a), seen in Fig. 1.

SPRC is also named as ZYZ-802. Due to the cysteine structure, SPRC was considered as an endogenous H_2S -producing agent by providing the substrate for H_2S synthesis. In the H_2S generating pathway, there are three important synthases:

Fig. 1 The chemical structures of SAC, SPC, and SPRC



cystathionine β -synthase (CBS) and cystathionine γ -lyase (CSE) both with a cofactor of pyridoxal 5'-phosphate (vitamin B₆) in cytoplasm (Szabó 2007) and 3-mercaptopyruvate sulfurtransferase (3-MST) along with cysteine aminotransferase (CAT) (Shibuya et al. 2009) in third synthesis pathway of H₂S production in mitochondria. The three synthases have different priorities of expressions in different systems, which mean SPRC may play a role of endogenous H₂S-producing compound via CSE/H₂S in cardiovascular system or CBS/H₂S in neural system.

The pharmacological effects of this cysteine analog were firstly investigated in 2009 on adult rat hearts and neonatal cardiomyocytes (Wang et al. 2009a). From then on, SPRC showed its protective effects in cardiovascular disease, neurodegenerative disease, cancer, and inflammatory disease. Interestingly, SPRC showed stronger cardioprotective effects of decreasing animal mortality, reducing heart infarct zone, lowering the activities and levels of LDK and CK, and promoting cell viabilities through activate antioxidative effects, compared to SAC and SPC treatment (Wang et al. 2010). In this chapter, the chemical and pharmacological characteristics of SPRC will be introduced in the following lines.

2 Analogs of SPRC

SPRC is an analog of SAC, which is one of the major compounds in aged garlic extract. As an original cysteine chemical, SAC firstly investigated the effects on neural and cardiovascular system. It was found that SAC inhibited cellular A β -induced apoptosis and destabilized Alzheimer's A β fibrils in in vitro (Gupta and Rao 2007). In Alzheimer's transgenic mouse model with Swedish double mutation, SAC attenuates Alzheimer's cerebral amyloid, neural inflammation, and tau phosphorylation (Chauhan 2006). In rat ischemic brain, SAC showed neural protection through decreasing generation of toxic-free radical, lipid peroxidation, and cellular damage (Numagami and Ohnishi 2001). More studies were investigated on rat model of stroke-prone spontaneously hypertension (Kim

et al. 2006). After treatments of SAC, the incidence of stroke was decreased, behavioral syndromes were relieved, and mortality due to stroke was reduced (Kim et al. 2006). Intervention with SAC showed its cardioprotection by significant decrease of myocardial infarction zone and infarction mortality (Shin et al. 2007).

Another analog of SPRC is SPC, which contains the propyl group in its cysteine structure. Recently, the pharmacological effects of SPC were investigated in diabetes, hyperlipidemia, and oxidative stress injury to protect kidneys, hearts, brains, and livers. First, in diabetic mice, SPC showed renal protection by significantly decreasing plasma urea nitrogen concentration and increasing creatinine clearance. Meanwhile, SPC intake also lessened the renal levels of reactive oxygen species (ROS), nitric oxide (NO), and inflammatory cytokines, like interleukin (IL)-6, tumor necrosis factor- α (TNF- α), prostaglandin E₂, and activations of NF- κ B, p38, and ERK1/2 induced in diabetic mice (Mong and Yin 2012). Second, SPC reduced the plasma levels of triacylglycerol and cholesterol in hyperlipidemic mice through enhancing the activities of antioxidants and inhibiting the protein and gene expressions of lipogenic-related enzymes, such as malic enzyme, fatty acid synthase, and 3-hydroxy-3-methylglutaryl coenzyme A reductase, together with sterol regulatory element-binding protein (SREBP)-1c and SREBP-2 (Lin and Yin 2008). Third, the anti-inflammatory and antioxidative effects can be further found in ethanol-induced acute hepatotoxicity (Yan and Yin 2007) and PC 12 cells insulted by ischemia (Chen and Yin 2008) and β -amyloid (Tsai et al. 2010) for neurodegenerative disease studies. However, a comparative study among SAC, SPC, and SPRC showed that SPC had weaker performance of cardioprotection and antioxidative effects, compared to SPRC (Wang et al. 2010).

3 Synthesis and Characteristics

3.1 Synthesis

In 50 L of reaction kettle, 27.6 L of dH₂O, 2.4 kg of Na₂CO₃, and 2.65 kg of L-cysteine hydrochloride were added and mixed to a solution. Under reaction temperature 0–5 °C, 3.06 kg of propargyl bromide was dropwise added and mixed for 1 h. After adjusting pH to 6, solids were precipitated and collected by aspirator filter pump with water washing. The product was purified by recrystallization from an ethanol–water solution. After drying, 960 g of products were collected with 40 % yield. The final product was verified by ¹H nuclear magnetic resonance spectroscopy. The purity was 99.7 %, as measured by high-performance liquid chromatography.

3.2 Characteristics

The final product of SPRC was faint yellow color powder, with garlic odor and slightly bitter taste. SPRC can be dissolved in dH₂O, NaOH, and dilute HCl and undissolved in ethanol, acetone, ethyl acetate, and dimethyl sulfoxide.

3.3 Purity

Following “Chinese Pharmacopoeia,” inorganic impurity was tested that chloride <0.05 %, sulfate <0.01 %, ferric salts <0.003 %, heavy metals <0.001 %, arsenic salts <0.0001 %, and water <0.53 %.

The products of SPRC have weight loss on drying <0.4 % and weight increment by moisture absorption around 5 %. The only possible residual organic solvent is ethanol, <0.0013 %.

3.4 Stability

The contents of SPRC products were >96 %, after photostability testing, high-temperature test, high humidity test, long-term stability study within 6 months, and accelerated stability study, showing its stability to temperature, light, and humidity.

4 Metabolism and Pharmacokinetics

The SPRC distribution was measured through stable isotope-labeled technique (Zheng et al. 2012). Adult Sprague–Dawley rats were fed [³⁵S]PRC at a dose of 75 mg/kg. The tissue distribution of [³⁵S]PRC-derived radioactivity was shown widely. Kidney showed the highest [³⁵S]PRC-derived radioactivity, which exceeded that of plasma. On the other side, several targeted organs, like brain, heart, lung, and intestine, presented lower [³⁵S]PRC-derived radioactivity than that of plasma. The plasma protein binding of SPRC measured by 96-well equilibrium dialysis was low in human, rat, and dog samples. LC-MS/MS was used to analyze the excretion of SPRC in urine, bile, and feces, which is 2.18 ± 0.61 %, 0.77 ± 0.61 %, and undetectable, respectively. The major metabolite in rat biomatrices, which was identified by MRM information-dependent, acquisition-enhanced product ion (MRM-IDA-EPI) scans on API 4000 QTRAP system, was *N*-acetylation (Zheng et al. 2012). These pharmacokinetic properties of SPRC were observed similar in previous pharmacokinetic SAC study (Nagae et al. 1994; Krause et al. 2002).

5 Formulation

SPRC was investigated to prepare for capsulation. The preparation process started from sieving main drug through 100 mesh sieve and other accessories through 80 mesh sieves. A mixture of lactose, microcrystalline cellulose, and cross-linked povidone as extenders, together with magnesium stearate and silicon powder capsule as lubricants, was made into main drug. It was finished by filling capsules and packing. These capsules have good fluidity and are easy to large-scale production.

SPRC developed controlled release formulation of SPRC (CR-SPRC), generating more stable and sustainable gas of H₂S (Huang et al. 2013). CR-SPRC

was produced by the solid dispersion technique using Eudragit (R) RS30D (Rohm Pharma, Weiterstadt, Germany) as a carrier. In a round-bottomed flask, CR-SPRC was dissolved in Eudragit (R) RS30D with a constant magnetic stirring in midrange rotation speed and 80 °C of a thermostat-controlled water bath. Eudragit(R) RS30D was then removed. The precipitates were cooled down to -20 °C immediately. After heating at 50 °C for 4 h, coprecipitates of CR-SPRC were desiccated, pulverized, and finally sieved through 100 mesh. The collection of CR-SPRC can be dissolved in 0.5 % sodium carboxymethylcellulose (CMC-Na) for future use (Huang et al. 2013). The pharmacological effects of CR-SPRC has been investigated in heart failure rat model. For more information, please refer to Sect. 6.1.2.

Incorporation of two mutually pharmacological or biological active compounds into one chemical by conjugation may be a good drug design to enhance the drug effects. Because leonurine and SPRC have been reported the cardioprotective effects both in vitro and in vivo, the two entities were considered to be synthesized to leonurine-SPRC conjugate, which connected by a phenolic hydroxyl ester bond so that easy hydrolyzing and releasing bioactive leonurine and SPRC (Liu et al. 2010). The pharmacological effects of leonurine-SPRC conjugate were investigated in neonatal rat ventricular myocytes under hypoxia, which will be discussed in the last paragraph of Sect. 6.1.1.

6 Pharmacologic Effects

6.1 Cardiovascular Disease

6.1.1 Ischemia/Hypoxia Injury

The modulation on cardiovascular disease by SPRC was firstly investigated in myocardial infarction rat models and hypoxic cardiomyocytes cells (Wang et al. 2009b). SPRC treatments were found in association with obvious reduction of myocardial infarct size and improved cardiac function in in vivo (Wang et al. 2009b). Rescuing hypoxic cardiomyocytes by SPRC indicates its cardioprotective effects. These effects go in line with the increased CSE activity and H₂S plasma concentration, which means the cardioprotective effects of SPRC, an endogenous H₂S-producing modulator, is through CSE/H₂S pathway (Wang et al. 2009b).

Furthermore, daily i.p. injection of three cysteine analogs (SAC, SPC, and SPRC) for 7 days before and 2 days after coronary artery ligation of rat hearts attenuated significant heart damage against ischemic reperfusion through the activation of antioxidative enzymes, such as superoxide dismutase (SOD), catalase (CAT), and glutathione peroxidase (GPx), as well as elevation of glutathione (GSH) levels, taken together with inhibition of lipid peroxidation products, like malondialdehyde (MDA) content (Wang et al. 2009a). Meanwhile, CSE expressions and H₂S producing rate and contents were increased by SPRC treatments but abolished by CSE selective inhibitor, propargylglycine (PAG), indicating the protection of SPRC through CSE/H₂S pathway (Wang et al. 2009a). It was highlighted that the SPRC showed stronger heart protection

compared to SAC and SPC. This may be due to the stronger connection of propargyl group in SPRC cysteine structure instead of allyl and propyl in SAC and SPC, respectively.

More cardiac effects of SPRC were investigated in hypoxia/reoxygenation (H/R) model (Liang et al. 2014). Rat ventricular myocytes and left ventricular papillary muscles were isolated to endure hypoxic injury for 30 min and followed by reoxygenation solution for 30 min. Pretreatment of SPRC 30 min prior to hypoxia may attenuate the apoptosis of isolated papillary muscles induced by H/R injury and maintain the morphology of these muscle fibers. Hypercontracture of ventricular myocytes was significantly inhibited by SPRC pretreatments at 3 min of reoxygenation and closed to normal level at 30 min of reoxygenation. Meanwhile, SPRC attenuated Ca^{2+} accumulation, high upstroke velocity, and increase of resting $[\text{Ca}^{2+}]_i$ induced by hypoxia. Because Ca^{2+} is mainly released by ryanodine receptors (RyR) and uptaken by SERCA-2, SPRC showed preserving activities of RyR and SERCA-2 and enhancing Ca^{2+} handling ability. These cardiac protective effects of SPRC were blocked by CSE inhibitor, PAG (Liang et al. 2014).

Owing to the mutually cardioprotective effects by leonurine and SPRC, the newly synthesized compound, leonurine-SPRC, possessed strong cardiac cellular protections of hypoxic neonatal rat ventricular myocytes in very low molar concentration of 1/10 leonurine dose and 1/100 SPRC dose (Liu et al. 2010). The cardioprotective mechanisms of this leonurine-cysteine analog illuminated three aspects: antioxidative effects by increasing SOD and CAT activities and decreasing MDA and ROS levels, ultrastructural protections of myocytes, and regulation of apoptosis by inhibiting activation of caspase-3 and rising Bcl-2 expressions (Liu et al. 2011). These findings indicated that this leonurine-SPRC conjugate showed potentially multifunctional anti-ischemia in neonatal rat ventricular myocytes, providing an important clue for future design and synthesis of such mutually pharmacological drugs.

6.1.2 Heart Failure

For releasing H_2S more stable and sustainable, CR-SPRC was introduced to heart failure rat model (Huang et al. 2013). The reduced infarct size and recovered cardiac function were discovered through activation of antioxidant defenses by increasing SOD, CAT, and GSH and decreasing creatine kinase (CK) leakage (Huang et al. 2013). Meanwhile, more protective mechanisms of SPRC are through the prohibition of activation of caspases and elevation of ratio of Bcl-2/Bax, showing the antiapoptotic effects of SPRC. Compared to original SPRC, CR-SPRC showed better performance of cardioprotection. Due to its endogenous H_2S -producing characteristics, the inhibition of heart failure by SPRC may be mediated by CSE/ H_2S pathway.

6.1.3 Angiogenesis

A study designed for the therapy of myocardial reperfusion injury investigated the proangiogenic effects of SPRC to provide a novel treatment target of ischemia/reperfusion heart disease (Kan et al. 2014). First, SPRC was found promoting cell

proliferation, cellular adhesion, migration, and tube formation in human umbilical vein endothelial cells (HUVEC). Moreover, under both normal condition and ischemic injury, rat aortic ring and Matrigel plug-implanted mice found the promoted angiogenesis. In the mouse ischemic hind limb by ligation of the left femoral artery and the rat ischemic heart by ligation of coronary artery, angiogenesis was facilitated by SPRC to attenuate the ischemic injury. Interestingly, SPRC treatments can significantly induce STAT3 phosphorylation in HUVEC in a dose- and time-dependent manner, in which effects were abolished by WP1066, an inhibitor of STAT3. After transfecting STAT3 siRNA into HUVECs to silence the expression of STAT3, the effects of cellular adhesion, migration, and tube formation by SPRC were abolished due to the absence of STAT3 expression. More studies showed that SPRC may not directly interact with STAT3 by the method of co-crystallization, but may enhance the interaction between VEGFR2 and STAT3 by co-immunoprecipitation. The influence to STAT3 by SPRC additionally found the nuclear translocation of STAT3 and transcriptional activation of downstream promoters, especially the *Vegf* promoter. Therefore, angiogenesis may be strengthened by SPRC through STAT3-mediated mechanism, which provides a novel avenue to target on ischemic disease.

6.2 Neurodegenerative Disease

Due to the neuroinflammatory effects by producing various inflammatory mediators and neural toxic proteins to exacerbate amyloid- β ($A\beta$) and hyperphosphorylated tau in Alzheimer's disease (AD), the anti-inflammatory effects of SPRC showed neural protection on rat spatial learning and memory impairment induced by lipopolysaccharide (LPS) bilateral intracerebroventricular (i.c.v.) injection valued by Morris water maze test (Gong et al. 2011a). The accumulation of $A\beta$ protein precursor, a type I transmembrane protein, playing key function and metabolism in pathogenesis of AD, together with $A\beta_{1-40/42}$, was suppressed by pretreatments of SPRC of 40 and 80 mg/kg, showing the inhibited effects of SPRC on $A\beta$ generation. Additionally, the overwhelming evidences of activation of tumor necrosis factor (TNF), like TNF- α and TNF- α receptor 1 (TNFR1), were prohibited by pretreatments of SPRC of 40 and 80 mg/kg. Further evidences of upregulatory inflammatory cytokines like I κ B- α and activation of NF- κ B p65 phosphorylation were significantly decreased by SPRC treatments of a dose range of 20–80 mg/kg. These findings all indicate SPRC may be a potential anti-neuroinflammatory agent for AD.

More studies were investigated on rats not limited on previous neuroinflammatory injury by LPS, but directly given i.c.v. injection of 10 μ g of $A\beta_{25-35}$ to induce most common AD rat model (Gong et al. 2011b). Administration of SPRC of 40 and 80 mg/kg through intraperitoneal injection may prohibit cognitive impairment and neuronal ultrastructural damage in AD rats. The inhibition of pathogenesis of AD by SPRC may be associated with the decreased proteins and genes expressions of TNF- α and cyclooxygenase-2 in AD hippocampus, suppressed extracellular signal-regulated kinase (ERK 1/2), reduced I κ B- α degradation, and weakened activation of NF- κ B after pretreatments of SPRC with $A\beta$

neuronal injury. The recovery of AD syndromes like cognitive impairment, maintaining of neuronal ultrastructure in AD rats, and less expressions of neuroinflammatory cytokines may suggest the underlying mechanisms and evidences of neural protective effects of SPRC on AD.

6.3 Cancer

SPRC investigated its potential effects on SGC-7901 gastric cancer cells (Ma et al. 2011). Cell viability, proliferation, and migration of SGC-7901 were significantly suppressed by SPRC treatments. These gastric cancer cells found the cell cycle arrest at the G1/S phase by SPRC treatments.

Further studies were performed on nude mice that accepted injection of SGC-7901 cells (Ma et al. 2011). After 7 days post-implant, when the tumor size reached 0.3–0.5 cm, mice were given 50 and 100 mg/kg intraperitoneal injections of SPRC. Tumor volumes were significantly reduced with 50 and 100 mg/kg of SPRC with tumor growth inhibition of 40–75 %. The tumor growth suppression by SPRC was inhibited by the inhibitor of H₂S-generating enzyme CSE, PAG. This anticancer mechanism of SPRC may be related to elevation of p53 and Bax expressions in tumor and SGC-7901 gastric cancer cells, indicating that SPRC induces proapoptotic effect.

6.4 Inflammation

The regulation of inflammation by SPRC was investigated in an *in vivo* model of acute pancreatitis (AP) in mice (Sidhapuriwala et al. 2012). Hourly caerulein injections for 10 h were given to induce AP. Pre-administration of SPRC was given 12 or 3 h before induction of AP. Pancreatic and pulmonary injuries were observed through increased amylase plasma concentration, elevated myeloperoxidase activities, and inflammatory histological observation, like edema, destruction of histo-architecture of the acini, and infiltration of inflammatory cells, which were reduced by 3 h pretreatment of SPRC before AP induction. Additionally, augmentation of pro-inflammatory cytokines (IL-1 β and IL-6) and decline of anti-inflammatory cytokine IL-10 were significantly inhibited by 3 h pretreatment of SPRC before AP induction. 12 h SPRC pre-administration before AP induction did not show significant effects on lung and pancreatic inflammatory injury. Meanwhile, anti-inflammatory effects of SPRC were in consistence with significant changes of H₂S level. Therefore, SPRC pretreatment of 10 mg/kg prior to AP induction ameliorated lung and pancreatic inflammatory injury by suppression of inflammatory cell infiltration and modulation of pro-/anti-inflammatory cytokine profile in plasma.

More anti-inflammatory effects of SPRC were investigated in other systems, like cardiac system, vascular endothelium, and neurodegenerative disease. First, in H9c2 cardiac myocytes, LPS was introduced as an inflammatory stimuli and SPRC as a protective treatment (Pan et al. 2011). In this LPS-induced inflammatory

response, NF- κ B was activated through phosphorylation of NF- κ Bp65 and I κ B- α degradation, which were inhibited by SPRC, as well as phosphorylation of ERK 1/2 and ROS generation. Other inflammatory cytokines, like TNF- α , inducible nitric oxide synthase, and intercellular adhesion molecules-1, were increased by LPS injury, which were suppressed by SPRC. Moreover, SPRC leads to phosphorylation of Akt in concentration-dependent and time-dependent manners, which were abolished by phosphoinositide 3-kinase (PI3K) inhibitor LY 294002, indicating the activated effects of SPRC on PI3K/Akt signaling pathway. The mechanisms of impairing NF- κ B/I κ B- α and inhibition of inflammatory cytokines had association with increasing H₂S and CSE levels induced by SPRC, indicating the cardioprotective effects are through CSE/H₂S pathway.

Second, SPRC was introduced to assess its effects on HUVECs injured by an inflammatory cytokines, NF- κ B (Pan et al. 2012). In this cardiac inflammatory cellular model, SPRC pretreatments observed a significant dose-dependent inhibition on the increased adhesion of U937 monocytes/ HUVECs induced by TNF- α . This decrease by SPRC (5 and 10 μ M) may be through the mechanism of reducing ICAM-1 and VCAM-1 expressions which were raised by TNF- α . Meanwhile, the translocation and phosphorylation of NF- κ B p65 and I κ B- α degradation were stimulated by TNF- α but attenuated by SPRC (5 and 10 μ M), indicating that the activation of NF- κ B was inhibited by SPRC pretreatments. Additionally, findings of TNF- α -induced JNK1/2 activation prohibited by SPRC (5 and 10 μ M) indicated that the regulation of JNK1/2 signaling pathway is also involved in the anti-inflammatory mechanisms of SPRC.

Third, the anti-inflammatory effects on neurodegenerative disease have been discussed in Sect. 6.2.

7 Conclusion

SPRC, a cysteine analog, inspired by the structure and vulnerable characteristics of a garlic extraction SAC, was synthesized to a propargyl group instead of allyl within the same cysteine structure in order to form a more stable structure with stronger pharmacological effects. The stronger protective effects of SPRC were firstly found in rat hearts and neonatal cardiomyocytes, compared to SAC and SPC. More pharmacological effects of SPRC were shown in cardiovascular diseases (hypoxia, ischemic reperfusion, H/R injury, heart failure, and angiogenesis), neurodegenerative disease (AD), cancer (gastric cancer), and inflammatory disease (AP and inflammatory responses induced by LPS or TNF- α in myocytes, endothelial cells, and rats). Meanwhile, SPRC, the endogenous H₂S agent, can increase CSE and H₂S productions, indicating that the regulation of cellular Ca²⁺ and activation of NF- κ B, TNF signaling, redox status, ERK 1/2, apoptosis, and inflammatory cytokines are through CSE/H₂S pathway. However, the investigations on pharmacological effects of this novel cysteine analog were still limited. More studies will be carried out on atherosclerosis, hypertension, and other diseases in vivo, ex vivo, and in vitro to unveil its underlying mechanisms of these potential protections.

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Phosphinodithioate and Phosphoramidodithioate Hydrogen Sulfide Donors

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Abstract

Hydrogen sulfide is rapidly emerging as a key physiological mediator and potential therapeutic tool in numerous areas such as acute and chronic inflammation, neurodegenerative and cardiovascular disease, diabetes, obesity and cancer. However, the vast majority of the published studies have employed crude sulfide salts such as sodium hydrosulfide (NaSH) and sodium sulfide (Na_2S) as H_2S “donors” to generate H_2S . Although these salts are cheap, readily available and easy to use, H_2S generated from them occurs as an instantaneous and pH-dependent dissociation, whereas endogenous H_2S synthesis from the enzymes cystathionine γ -lyase, cystathionine- β -synthase and 3-mercaptopyruvate sulfurtransferase is a slow and sustained process. Furthermore, sulfide salts are frequently used at concentrations (e.g. 100 μM to 10 mM) far in excess of the levels of H_2S reported in vivo (nM to low μM). For the therapeutic potential of H_2S is to be properly harnessed, pharmacological agents which generate H_2S in a physiological manner and deliver physiologically relevant concentrations are needed. The phosphorodithioate GYY4137 has been proposed as “slow-release” H_2S donors and has shown promising efficacy in cellular and animal model diseases such as hypertension, sepsis, atherosclerosis, neonatal lung injury and cancer. However, H_2S generation from GYY4137 is inefficient necessitating its use at high concentrations/doses. However, structural modification of the phosphorodithioate core has led to compounds (e.g. AP67 and AP105) with accelerated rates of H_2S generation and enhanced biological activity. In this review, the therapeutic potential and limitations of GYY4137 and related phosphorodithioate derivatives are discussed.

Keywords

GYY4137 • AP67 • AP105 • Slow-release H_2S donor • cGMP

1 Introduction

Recent research into the regulation of endogenously synthesised hydrogen sulfide (H_2S) from the enzymes cystathionine γ -lyase (CSE), cystathionine- β -synthase (CBS) and 3-mercaptopyruvate sulfurtransferase (3-MST) has revealed a startling array of physiological and pathophysiological processes that are regulated by H_2S including learning and memory, regulation of vascular tone, ion channel regulation, cytoprotection, acute and chronic inflammation, metabolism, allergy and cell signalling, etc. The fact that there are three regulated cellular processes to synthesise this gaseous molecule highlights its importance, and key findings in this area of research are presented elsewhere in this book. The rapid advances in molecular biology have given researchers in this area highly selective tools to silence CSE, CBS and 3-MST from cells and tissues as well as genetic knock-out animals devoid of these enzymes. Whilst potent and selective inhibitors of CSE, CBS and 3-MST are still not available and the use of commonly used inhibitors should be viewed

with caution (Asimakopoulou et al. 2013; Whiteman et al. 2011), molecular techniques have graphically illustrated the detrimental effects “H₂S removal” has to cells, tissues and animals. Studies in animal models and human disease have revealed that endogenous H₂S synthesis is perturbed in ageing (Qabazard et al. 2014; Yang et al. 2013), cardiovascular disease (Calvert et al. 2009, 2010; Minamishima et al. 2009; Yang et al. 2008), pre-eclampsia (Wang et al. 2013), acute and chronic inflammation (Li et al. 2009, 2013), diabetes (Whiteman et al. 2010a) and respiratory pathologies such as asthma (Chung 2014) and chronic obstructive pulmonary disease (Chen et al. 2005; Saito et al. 2014; Kirkham et al. 2014). These observations have consistently led to the suggestion that strategies which increase H₂S bioavailability are viable therapeutic approaches to disease treatment. However, at present, there are very few molecules described in the literature that can achieve this. As such, this review will highlight the recent progress that has been made with phosphorodithioate-based H₂S “donors”, typified by the molecule GYY4137.

2 The Need for Slow-Release H₂S Donor Molecules

Although multiple studies have highlighted the diverse physiological role for H₂S and have emphasised the possible therapeutic opportunities for modulating H₂S bioavailability, it should be noted, and with some concern, that the bulk of the scientific literature is replete with the use of high concentrations (and doses) of crude sulfide salts such as sodium hydrosulfide (NaSH) or sodium sulfide (Na₂S) where they have been used as “fast-release H₂S donors”. Whilst these salts are undoubtedly convenient for use in the laboratory as one can readily generate solutions containing known quantities of H₂S, they are highly problematic. Although these problems have been reviewed in detail elsewhere (Whiteman et al. 2011; Whiteman and Winyard 2011), it is pertinent to briefly discuss the major flaws. Firstly, NaSH and Na₂S are simple salts and not H₂S donors. They undergo instantaneous and pH-dependent dissociation generating, not releasing, H₂S. In sharp contrast, enzymatic synthesis of H₂S from CSE, CBS and 3-MST is not instantaneous but a slow and sustained process (Whiteman et al. 2011; Li et al. 2013; Banerjee and Zou 2005; Kabil et al. 2011a, b). This obvious discrepancy in kinetics of H₂S generation from NaSH and Na₂S compared to CSE-, CBS- and 3-MST-derived H₂S should preclude their use for determining the physiological and biological properties of H₂S. Secondly, high concentrations, >100 μM and often 1–10 mM, are frequently required to elicit any biological response in cells or tissue, presumably because H₂S generated from these salts dissipates in seconds (Li et al. 2008; Whiteman et al. 2010b), rather than being synthesised slowly by CSE, CBS and 3-MST. Thirdly, commercial supplies of NaSH are often of very poor purity (typically stated as >60 % purity), and despite this information being clearly stated in the online product specification information and documents that will accompany NaSH, and any commercial research product, when it is delivered (e.g. Sigma-Aldrich (2014) and Caymen Chemicals (2014)), it continues to be

widely used. As such, sulfide salts are poor models for determining the role of endogenous H_2S in physiological systems or exploring the pharmacological and therapeutic potential of H_2S -donating molecules.

Although several detailed and excellent reviews have explicitly highlighted how Na_2S and NaSH should and should not be used (Whiteman et al. 2011; Yuan et al. 2015; Hughes et al. 2009; Olson 2013) in the laboratory (e.g. preparing and storing solutions, effects of pH and buffers, preventing loss of H_2S to the atmosphere, etc.), they do not address the problem that these salts generate H_2S in an instant and as a rapidly dissipating bolus. As such, there is great interest in the development of slow-release H_2S donor molecules which overcome the problems stated above.

3 Water-Soluble Slow-Release Hydrogen Sulfide Donors

The first water-soluble slow-release H_2S donor to be described was morpholin-4-ium 4 methoxyphenyl(morpholino) phosphinodithioate (GYY4137), developed by Moore and colleagues at the National University of Singapore (Li et al. 2008) (Fig. 1). GYY4137 offers considerable advantages over the use of sulfide salts such as NaSH or Na_2S since the rate of H_2S generation more readily mimics endogenously synthesised H_2S from enzymatic sources such as CSE, CBS and 3-MST, rather than delivering H_2S as a single concentrated bolus, e.g. the manner in which H_2S is generated from GYY4137 more accurately reflects endogenously produced H_2S . A second obvious advantage is that the final concentrations of H_2S generated from GYY4137 are likely more physiologically, pathophysiologically and therapeutically relevant than a bolus of concentrated and impure (Sigma-Aldrich 2014; Caymen Chemicals 2014) sulfide salt. Thirdly, the high solubility of GYY4137 in water allows concentrated stock solutions to be prepared and used immediately in the laboratory for use on cells, tissues and animals without the need for organic solvents (e.g. ethanol, DMSO, dimethylformamide) commonly required for other drugs. However, it should be noted with caution that H_2S generation will start upon contact with water, and the freezing and reuse of stock solutions should be avoided. This ease of use and highly desirable water solubility has no doubt contributed to its widespread use and application.

3.1 GYY4137: Cytoprotection

H_2S has been reported to exert various “antioxidant” effects and inhibit the detrimental effects of reactive oxygen and nitrogen species on cells and tissues. However, the concentrations of the H_2S (typically NaSH) were often far in excess of that detected in vivo. For example, NaSH (15–60 μM) inhibited peroxynitrite (Whiteman et al. 2004)- and hypochlorous acid (Whiteman et al. 2005)-induced cellular toxicity, protein and lipid oxidation in human SH-SY5Y neuroblastoma cells. Higher concentrations of NaSH (e.g. 400–500 μM) also protected neuronal,

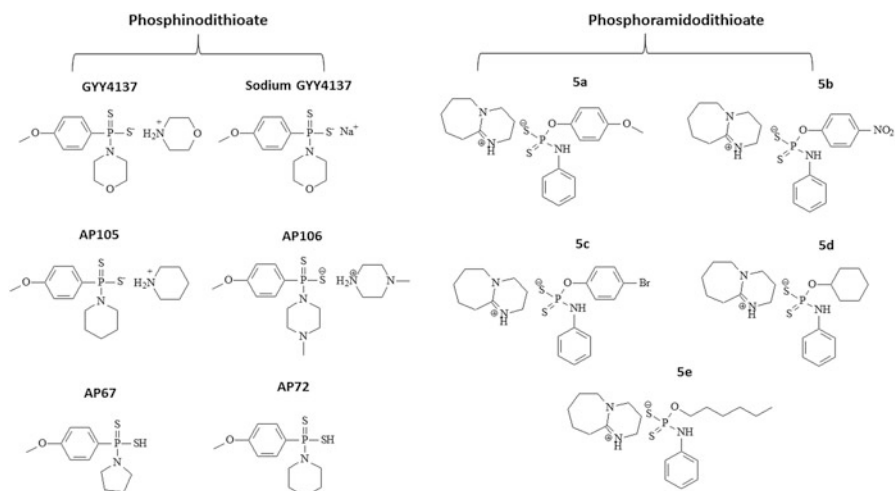


Fig. 1 Phosphinodithioate and phosphoramidodithioate H₂S donors. Examples of phosphinodithioates (GYY4137, sodium GYY4137, AP67, AP72, AP105 and AP106) and phosphoramidodithioates (5a–e) from the literature

microglial, myocardial and vascular cells from H₂O₂, lipid peroxides, chemical hypoxia and neurotoxins such as β -amyloid and MPP⁺ (Dong et al. 2012; Liu and Bian 2010; Xiao et al. 2012; Yin et al. 2009; Chen et al. 2010; Kimura et al. 2006; Kimura and Kimura 2004). Other studies have suggested NaSH (1 mM) may be cytoprotective by reducing lipid peroxides formed during the oxidation of LDL (Muellner et al. 2009). However, it is highly unlikely these concentrations are ever reached in the brain or vasculature (Olson 2013; Kolluru et al. 2011, 2013; Peter et al. 2013; Whitfield et al. 2008) or reached without inducing substantial toxicity and death. Based on the above in vitro observations, H₂S has been proposed as endogenous “antioxidant” effects of H₂S in vitro have been suggested (Guo et al. 2013). However, the low concentrations of H₂S in cells and tissues in vivo (nM to low μ M; Kolluru et al. 2013; Kolluru et al. 2011; Whitfield et al. 2008) and the very slow rate of reaction between sulfide and biologically relevant oxidants (e.g. peroxide, superoxide, peroxynitrite, etc.) strongly suggest direct antioxidant effects (e.g. “oxidant scavenging”) are unlikely (Carballal et al. 2013). As such, cytoprotection is probably achieved through more subtle processes, and it would be valuable to reassess some of the earlier observations of H₂S-mediated cytoprotection using donors which deliver realistic concentrations of H₂S in a physiologically relevant manner.

The slow-release H₂S donor GYY4137 has been shown to exert significant cytoprotective effects in a variety of cellular and animal models of human disease. For example, in human mesenchymal progenitor cells and human articular chondrocytes, GYY4137 (10–200 μ M) prevented cytotoxicity induced by the oxidants 4-hydroxynonenal, H₂O₂ and peroxynitrite (generated using SIN-1) (Fox et al. 2012). GYY4137 also preserved mitochondrial membrane potential ($\Delta\psi$ m)

and inhibited ATP loss as well as inhibited mitochondrial oxidant production in these cells, possibly through Akt- and ERK1/2-mediated signalling pathways. Activation of Akt was also observed in GYY4137 (100–500 μM)-treated human keratinocytes (Merighi et al. 2012). Furthermore, oxidative stress-induced cytotoxicity was potentiated by siRNA-mediated knockdown of CSE, further indicating a cytoprotective role for endogenous H_2S and the potential for synthetic H_2S donor molecules to overcome H_2S deficiency. More recently, these findings have been confirmed in human umbilical vein endothelial cells (HUVEC) exposed to H_2O_2 (Hine et al. 2015).

In other in vitro models, GYY4137 (100–200 μM) protected H9c2 myocardial cells from hyperglycaemia-induced toxicity via mitochondrial protection and activation of AMPK-/mTOR-dependent signalling (Wei et al. 2014). Mitochondrial protection induced by GYY4137 (100–500 μM) was also observed in oxLDL-treated RAW264.7 cells (Liu et al. 2013). Higher concentrations (100 mM) also protected human pulmonary arterial cells from hyperoxia, inhibited cellular and mitochondrial “ROS” production and reversed hyperoxia-induced alveolar cell growth arrest (Vadivel et al. 2014), although cellular H_2S levels from such an obviously high concentration of GYY4137 were not assessed in this study. In vivo cytoprotection was also observed in a hyperoxia-induced lung injury model in new born rat pups (Vadivel et al. 2014). In this study, intraperitoneal injection of GYY417 (37.75 mg/kg, i.p.) inhibited hyperoxia-induced alveolar and lung vascular cell growth arrest and pulmonary hypertension, reducing right ventricular hypertrophy and pulmonary arterial medial wall thickness and inhibiting pulmonary artery smooth muscle cell proliferation. When given therapeutically, GYY4137 rescued alveolarisation, activated Akt/PI3K signalling in hyperoxic lungs and restored mitochondrial $\Delta\psi\text{m}$ and lowered mitochondrial oxidant production in lung epithelial cells (Vadivel et al. 2014). However, it is not clear whether these effects were due to H_2S from GYY4137 since changes in blood, tissue or cellular H_2S before and after GYY4137 administration were not measured.

The above cytoprotective effects of slow-release H_2S donors have led the suggestion that manipulation of intracellular H_2S synthesis could hold the key to increasing life span (Qabazard et al. 2013, 2014; Vozdek et al. 1834; Budde and Roth 2011; Miller and Roth 2007). It has been known for some time that blood and tissue levels of “ H_2S ” decline with age (Zhang et al. 2013; Predmore et al. 2010), and up until recently (Li et al. 2013), there was no mechanism to replace the loss of this gas. Qabazard et al. (2014) were the first to investigate the effects of GYY4137 in a well-established model of whole animal ageing using *Caenorhabditis elegans*. GYY4137 (100–400 μM) significantly increased the life span, increasing both the growth rate and activity of the worms. GYY4137 also inhibited oxidative stress-induced *C. elegans* killing by paraquat presumably by inhibiting mitochondrial “ROS” generation (Qabazard et al. 2014) suggesting H_2S could also inhibit oxidative stress occurring during the “ageing process”. Subsequent studies (Hine et al. 2015) have confirmed these original observations, replicating them in *C. elegans*, yeast and the fruit flies *Drosophila melanogaster* where again, inhibition of ageing-induced oxidative stress by GYY4137 appeared to be a common

mechanisms across species. It is therefore tempting to speculate that strategies which promote endogenous H₂S synthesis and/or provide exogenous H₂S in a physiologically relevant manner could hold the key to healthful ageing and longevity. Although these studies clearly show GYY4137 to be cytoprotective in a variety of models, it is possible that compounds which generate H₂S more efficiently could offer greater selectivity and potency.

3.2 GYY4137: Anticancer Effects

Although there are no conclusive reports of H₂S directly causing cancer in humans, numerous clinical and epidemiological studies have evaluated in detail the relationship between commensal sulfate-reducing bacteria (e.g. H₂S synthesising), ulcerative colitis and colorectal cancer (Carbonero et al. 2012; Roediger et al. 1997; Huycke and Gaskins 2004; Medani et al. 2011). H₂S is thought to be directly toxic to colonic epithelium, initiating inflammation and tissue damage and subsequent cellular hyper-proliferation (Huycke and Gaskins 2004; Medani et al. 2011). Indeed, the addition of NaSH (1–5 mM) to cells from human colonic biopsies (Christl et al. 1996), intestinal epithelial cells (Deplancke and Gaskins 2003; Leschelle et al. 2005) and human HT29 adenocarcinoma cells (Leschelle et al. 2005) resulted in significant cellular proliferation. Additionally, NaSH (250 μM to 2 mM) induced genomic DNA damage HT29 cells via “free radical”-mediated processes in Attene-Ramos et al. (2006, 2007). More recently, Szabo et al. (2013) showed that in human colorectal carcinoma, CBS translocation to mitochondria and subsequent H₂S generation from CBS were a key mediator in driving the proliferation of carcinoma cells in vitro and in vivo via overstimulation of mitochondrial bioenergetics. Collectively, these studies strongly suggest that H₂S is pro-carcinogenic, at least with regard to colorectal carcinoma, and as such, inhibitors of CBS-mediated H₂S synthesis, or compounds which sequester or “scavenge” H₂S, could have therapeutic value (Szabo et al. 2013).

However, it is prudent to note that as with many aspects of the H₂S research field, diametrically opposite data towards cancer treatment are also presented. GYY4137 was further recently shown to exhibit prominent anticancer activity in vitro and in vivo (Lee et al. 2011). In this study, GYY4137 (100 μM to 1 mM) but not equivalent concentrations of NaSH induced substantial G₂/M phase cell cycle arrest and PARP-/caspase-9-dependent apoptotic cell death in seven disparate human cancer cell lines: cervical carcinoma (HeLa), colorectal carcinoma (HCT-116), breast adenocarcinoma (MCF-7), hepatocellular carcinoma (HepG2), osteosarcoma (U2OS), acute promyelocytic leukaemia (HL-60) and myelomonocytic leukaemia (MV4-11) cells. In sharp contrast, the same concentrations of GYY4137 had no effect on normal human lung fibroblasts (IMRO90 and WI-38) suggesting cell killing was cancer cell specific. The anticancer effects were not only limited to in vitro experiments. Both Lee et al. (2011) and then Lu et al. (2014) showed GYY4137 (100–300 mg/kg i.p.) inhibited tumour cell growth in HL-60, MV4-11 (Lee et al. 2011) and HepG2 xenografts (Lu et al. 2014)

in mice. Although the mechanisms for these clear observations have not been fully elucidated, it is known that GYY4137 induces intracellular acidification in cancer cell lines through increased glycolysis leading to enhanced lactate production and decreased anion exchanger and sodium/proton exchanger activity (Lee et al. 2014). Other mechanisms in HepG2 and Bel7402 hepatocellular carcinoma cells include suppression of STAT3 activation (Lu et al. 2014). One key aspect of these studies is that for the first time, a control compound incapable of generating H₂S was employed, albeit one structurally dissimilar to GYY4137, and shown to not have any suppressive effect on tumour cell proliferation in vitro or in vivo (Lee et al. 2011, 2014).

Since GYY4137 suppressed tumour cell growth in vitro and in vivo (Lee et al. 2011, 2014; Lu et al. 2014), but NaSH was either ineffective or promoted cell proliferation (Christl et al. 1996; Deplancke and Gaskins 2003; Leschelle et al. 2005), the cellular response to H₂S may be determined not only by the concentration of H₂S used, but by the manner in which cells, tissues and animals are exposed to H₂S, e.g. fluxes of low concentrations of H₂S from a slow-release H₂S donor (from GYY4137) vs. instant generation from a concentrated bolus of sulfide salts (such as NaSH) could be key to identifying novel therapeutic H₂S-based agents.

3.3 GYY4137: Anti-inflammatory Effects

When generated as a bolus of NaSH, H₂S has been shown to be both pro- and anti-inflammatory leading to considerable controversy in the literature (reviewed in Whiteman and Winyard (2011)). H₂S synthesis is induced in many cell types in vitro and in vivo by pro-inflammatory mediators such as endotoxin and cytokines, and tissue and elevated blood and tissue levels of H₂S are increased in animal models of acute and chronic inflammation, for example, haemorrhagic (Mok and Moore 2008) and septic shock (Li et al. 2005; Coletta and Szabo 2013), pancreatitis (Bhatia et al. 2005; Tamizhselvi et al. 2007) and arthritis (Li et al. 2013). In humans, increased blood levels of H₂S are detected in patients with endotoxic shock (Li et al. 2005) and either increased (Saito et al. 2013) or decreased (Wang et al. 2011a) in asthma. Increased (Chen et al. 2005) (or decreased (Kirkham et al. 2014)) sputum H₂S is observed in patients with chronic obstructive pulmonary disease (COPD). The ratio of blood/sputum H₂S levels has been proposed as a novel biomarker for airway disease (Chung 2014; Saito et al. 2013, 2014; Wang et al. 2011a). Similarly, synovial fluid H₂S levels are elevated relative to disease controls in patients with rheumatoid arthritis (Whiteman et al. 2010c) and inflammatory joint disease (Whiteman and Winyard 2011). Early studies generally showed inhibition of endogenous H₂S synthesis attenuated pro-inflammatory signalling in vitro and inflammation in vivo (reviewed in Whiteman and Winyard (2011)), although it should be noted that pharmacological inhibitors of CSE, CBS and 3-MST are non-specific necessitating their use at very high concentrations (reviewed in detail in Whiteman et al. (2011)) and the addition of NaSH to cultured

cells in vitro (Whiteman et al. 2010b; Zhang et al. 2008; Zhi et al. 2007; Kloesch et al. 2012a, b). Similarly, administration of NaSH to laboratory animals either induced systemic inflammation directly or exacerbated inflammation (Li et al. 2005; Bhatia et al. 2005, 2006; Tamizhselvi et al. 2007; Ang et al. 2010, 2011; Badiei et al. 2013).

Based on the findings above, it has been suggested by some investigators that endogenous and pharmacological H₂S (e.g. NaSH) is pro-inflammatory. However, when a slow-release H₂S donor such as GYY4137 is employed in cellular and in vivo models of inflammation, the opposite is observed suggesting the role of H₂S in inflammation needs to be thoroughly re-examined. For example, lipopolysaccharide (LPS)-induced NF- κ B activation, cytokine, \bullet NO and PGE₂ synthesis were exacerbated in murine RAW264.7 macrophages pretreated with NaSH (up to 200 μ M) but attenuated by GYY4137 (50–200 μ M) (Whiteman et al. 2010b). This attenuation was also observed when GYY4137 was added to RAW264.7 cells after LPS stimulation highlighting the therapeutic potential of slow-release H₂S donors. Similarly, GYY4137 (50–400 μ M) inhibited NF- κ B activation in oxLDL-treated RAW264.7 (Liu et al. 2013) and in human monocytic U937 cells and monocyte-derived macrophages infected with *Mycoplasma fermentans*, preventing MCP-1 production (2 mM) (Benedetti et al. 2014). GYY4137 (100–400 μ M) also inhibited NF- κ B activity in oxLDL (100 μ g/mL)-treated murine RAW264.7 macrophages resulting in lowered LOX-1 and iNOS expression and synthesis of macrophage-specific chemokines such as MIP-2 α and IP10 (Liu et al. 2013).

In articular chondrocytes isolated from patients with osteoarthritis (OA) (Burguera et al. 2014), GYY4137 also suppressed NF- κ B activity leading to lowered protein levels of the pro-inflammatory enzymes COX-2, prostaglandin E synthase, MMP-13 and iNOS leading to lowered synthesis of \bullet NO, PGE₂ and IL-6 suggesting H₂S donor compounds could be useful in inhibiting and/or reverse cartilage remodelling in degenerative joint diseases such as OA. Furthermore, in human articular chondrocytes and synovial fibroblasts (Li et al. 2013), GYY4137 (100–500 μ M) also inhibited LPS-induced synthesis of IL-6, TNF- α , IL-1 β , \bullet NO and PGE₂ as well as inhibited the activity of iNOS, COX-2 and TNF- α converting enzyme (TACE) suggesting additional anti-inflammatory effects of this compound in addition to NF- κ B inhibition.

The anti-inflammatory effects of GYY4137 have also been evaluated in in vivo. In Coxsackie virus B-infected rat cardiomyocytes, GYY4137 was recently shown to reduce cardiac damage and circulating TNF- α , IL-1 β and IL-6 levels in an NF- κ B-dependent manner (Wu et al. 2015). Similarly, in the LPS-induced sepsis model in rats, GYY4137 also inhibited the synthesis of these cytokines via suppression of NF- κ B activity, reduced tissue inflammation and elevated plasma levels of the anti-inflammatory chemokine, IL-10, although the mechanism for this is not clear (Li et al. 2009). The addition of GYY4137 to LPS-treated whole rat blood (Li et al. 2009) also resulted in the inhibition of TNF- α generation, possibly via inhibition of TACE (Li et al. 2013) and again highlighting the therapeutic potential for H₂S donors as anti-inflammatory compounds. Furthermore, GYY4137 (50 mg/kg i.p.) reversed LPS-induced hypotension. Although this may seem

counterintuitive since H₂S dilates blood vessels (see below) and sepsis markedly lowers systemic blood pressure and elevates blood and tissue levels of H₂S, the normalisation of blood pressure by GYY4137 is most likely due to inhibition of pro-inflammatory cytokines such as TNF- α and IL-1 β and other factors which drive the detrimental vascular in sepsis, rather than a dilating effect on blood vessels directly.

The therapeutic potential for slow-release H₂S donors has also been highlighted more recently in the rat complete Freund's adjuvant (CFA) arthritis model and in atherosclerotic apoE^{-/-} mice. In the CFA arthritis model (Li et al. 2013), CFA injection induced synovial H₂S synthesis fluid (SF), in agreement with *in vitro* studies using human joint cells (Li et al. 2013; Fox et al. 2012; Burguera et al. 2014) and observations with human SF from patients with various inflammatory arthritides (Whiteman and Winyard 2011; Whiteman et al. 2010c). GYY4137 (50 mg/kg *i.p.*) given 1 h prior to CFA resulted in increased tissue oedema although lowered neutrophil and macrophage infiltrate, presumably due to the vasodilatory effects of GYY4137 (Li et al. 2008). However, therapeutic administration of GYY4137 either 6 or 18 h post-CFA treatment resulted in decreased knee joint swelling, reduced SF neutrophil and macrophage accumulation and decreased SF levels of TNF- α , IL-1 β , IL-8 and IL-6 (Li et al. 2008). Furthermore, in atherosclerotic apoE^{-/-} mice, daily *i.p.* injection of GYY4137 (133 μ mol/kg/day) for 30 days resulted in the inhibition of aortic mitochondrial "ROS" production, LOX-1 expression and lowered cellular levels of IL-6, TNF- α and ICAM-1 (Liu et al. 2013). GYY4137 also partially restored endothelium-dependent vasorelaxation in aortic segments (Liu et al. 2013). Since loss of vascular H₂S occurs in atherosclerosis (Mani et al. 2013, 2014) and the onset of atherosclerosis is more rapid and aggressive in CSE^{-/-} mice compared to wild-type animals (Mani et al. 2013), these studies strongly suggest that slow-release H₂S donors could reverse (as well as prevent) the detrimental vascular remodelling associated with atherosclerosis, highlighting their therapeutic potential in vascular diseases. Although these studies consistently show GYY4137 exerted prominent anti-inflammatory activity *in vitro* and *in vivo* when used prophylactically and therapeutically, it is possible that slow-release H₂S donor compounds that are able to generate H₂S with greater efficiency than GYY4137 could offer improved potency and efficacy.

3.4 GYY4137: Vascular Effects

The effects of H₂S on vascular tissue are by far the most characterised. The role of H₂S as an "endogenous vasodilator" was first proposed by Zhao et al. (2001). The addition of NaSH or solutions of authentic H₂S gas to where more than 200 μ M H₂S (Zhao et al. 2001; Zhong et al. 2003; Ali et al. 2006) was required to relax precontracted rat aortic rings and lowered systemic blood pressure in rats via an endothelium and K_{ATP} channel-dependent mechanism, possibly through the S-sulphydration of critical cysteine residues in the K_{ATP} channel (Mustafa et al. 2011). Since then, numerous other studies have reported the vasorelaxant

effect of high concentrations of NaSH on other vascular tissues where effects on calcium channels, Na⁺/K⁺ ATPase, HCO₃⁻ and other K⁺ channels are activated or inhibited by H₂S (reviewed in Whiteman and Moore (2009) and Peers et al. (2012)) but often using near mM or mM concentrations of the salt (e.g. Al-Magableh and Hart (2011) and Hedegaard et al. 2014)). As a result of these studies, H₂S is often described as a “potent vasodilator” (Bhatia 2005; Pushpakumar et al. 2014; Ariyaratnam et al. 2013; Jang et al. 2014; Baragatti et al. 2013; Ghasemi et al. 2012) despite data clearly showing otherwise. However, the addition of GYY4137 at concentrations and incubation times which deliver nM to low μM H₂S also leads to vasodilation in vitro and in vivo (Table 1) suggesting blood vessels are acutely sensitive to small fluxes of H₂S when it is generated in a manner and a level comparable to enzymatic (e.g. via CSE, CBS and 3-MST). For example, in phenylephrine precontracted rat aorta, GYY4137 (10–100 μM) induced significant vessel relaxation, but >200 μM NaSH was required for comparable effects (Li et al. 2008). Although the dilatory effects of a bolus of NaSH (EC₇₀, 300 μM) were immediate, vessel relaxation in response to GYY4137 (EC₇₀, 200 μM) was slow and continued for up to 60 min. The effect of GYY4137 and NaSH was inhibited by glibenclamide and PNU37883A, indicating a requirement for K_{ATP} channel activation, and reduced by endothelium removal and pretreatment with L-NAME (NOS inhibitor) and ODQ (guanylate cyclase inhibitor), which inhibit the formation and vascular response to [•]NO, respectively. K_{ATP} channel dependence for vessel relaxation appears conserved as glibenclamide also prevented GYY4137-induced human internal thoracic artery relaxation (Webb et al. 2008) and dilation of human and rat term pregnant myometrium (Robinson and Wray 2012), although in this latter study, GYY4137 (1 mM) also inhibited L-type calcium channel activity suggesting a more complex mechanism of action. In agreement, bovine ciliary artery vessel dilation was observed with as little as 100 nM GYY4137 (EC₅₀ = 13.4 ± 1.9 μM) (Chitnis et al. 2013), and although this effect was inhibited by glibenclamide, it was potentiated by flurbiprofen suggesting prostaglandin-/prostacyclin-mediated effects of GYY4137 could also contribute, at least in part, to its mechanism of action. It is currently not known why the bovine ciliary artery appears to be exquisitely sensitive to H₂S.

In the original study (Li et al. 2008), the addition of 100 μM GYY4137 to rat aortic vascular smooth muscle cells did not result in cGMP accumulation suggesting guanylate cyclase and phosphodiesterase-independent effects of GYY4137 on vessel relaxation. Others have observed that higher concentrations of GYY4137 (300 μM) significantly increased cGMP in the same cells, albeit modestly, and whereas rat aorta relaxation by ≥100 μM NaSH was blocked by the PKG inhibitor DT-2, GYY4137-mediated vessel relaxation was independent of PKG (Salomone et al. 2014; Bucci et al. 2012). In contrast, much higher concentrations of GYY4137 (1 mM) reduced cGMP levels to below that of the controls in ophthalmic artery segments (Salomone et al. 2014) suggesting there could be tissue (as well as concentration)-dependent effects of the donor (and H₂S). These studies further highlight the differences between “instantly” generated H₂S and H₂S generated slowly and at low concentrations.

Table 1 Summary of in vitro and in vivo effects of GYY4137 and related compounds

Therapeutic area	Compound	Observation/comment
Cardiovascular	GYY4137	<ul style="list-style-type: none"> • K_{ATP} and endothelium-dependent dilation of rat aorta ex vivo (10–100 μM) • Inhibition of L-NAME-induced hypertension in rats in vivo (133 μM/kg i.v.) • Sustained blood pressure reduction in spontaneously hypertensive rats but minimal effect in normotensive WKY rats in vivo (133 μM/kg i.v.) • Decreased rat kidney perfusion pressure ex vivo • No acute or chronic toxicity observed <hr/> <ul style="list-style-type: none"> • Induced relaxation observed using 1–1000 μM GYY4137 with significant relaxation observed from 1 μM; K_{ATP} and L-type calcium channel-dependent effects <hr/> <ul style="list-style-type: none"> • Normalisation of mean blood pressure and increased foetal growth in pre-eclamptic mice (GYY4137, 0.25 mg/kg i.p.) (Wang et al. 2013) <hr/> <ul style="list-style-type: none"> • Relaxation of human airway smooth muscle via sarcolemmal K_{ATP} channels, although >10 mM was required (Fitzgerald et al. 2014) <hr/> <ul style="list-style-type: none"> • Variable effects on vascular smooth muscle cells • No cGMP accumulation with 100 μM GYY4137 (Li et al. 2008), increased with 300 μM (Bucci et al. 2012) or inhibited with 1 mM (Salomone et al. 2014) <hr/> <ul style="list-style-type: none"> • K_{ATP}- and prostaglandin-dependent dilation of bovine ciliary artery induced by 100 nM to 100 μM GYY4137 (EC50 = 13.4 \pm 1.9 μM) (Chitnis et al. 2013)
	AP67 and AP72	<ul style="list-style-type: none"> • K_{ATP}- and prostaglandin-dependent dilation of bovine ciliary artery induced by 1 nM to 10 μM AP67 (EC50 = 0.08 \pm 0.04 μM) and AP72 (EC50 = 4.3 \pm 0.09 μM) (Chitnis et al. 2015). Both donors were used in the free acid form (Fig. 1) • Restored endothelium-dependent vasorelaxation in apoE^{-/-} atherosclerotic mice
Cytoprotection	GYY4137	<ul style="list-style-type: none"> • GYY4137 (100–200 μM) inhibited 4-hydroxynonenal-, H₂O₂- and peroxynitrite-induced cytotoxicity in human articular chondrocytes and trabecular bone-derived mesenchymal progenitor cells • Preservation of mitochondrial $\Delta\psi$m and ATP levels • Activation of Akt signalling • No cytoprotection observed with equimolar concentrations of “decomposed” GYY4137 <hr/> <ul style="list-style-type: none"> • Activation of Akt in human keratinocytes by 100–500 μM GYY4137 (Merighi et al. 2012) <hr/> <ul style="list-style-type: none"> • Inhibition of H₂O₂-induced cytotoxicity in HUVEC by 100 μM GYY4137 (Hine et al. 2015)

(continued)

Table 1 (continued)

Therapeutic area	Compound	Observation/comment
		<ul style="list-style-type: none"> • Inhibition of oxLDL-induced cellular and mitochondrial toxicity in murine RAW264.7 macrophages by 100–500 M GYY4137 (Liu et al. 2013) • Inhibition of hyperglycaemia-induced cellular and mitochondrial toxicity in rat H9c2 myocardial cells by 100–200 μM GYY4137 (Wei et al. 2014) • Activation of AMPK-/mTOR-dependent signalling • Inhibition of mitochondrial “ROS” production and mitochondrial and cellular toxicity in hyperoxia exposed pulmonary artery smooth muscle cells and lung epithelial cells (100 mM) (Vadivel et al. 2014) • Inhibition of hyperoxia-induced alveolar cell growth arrest, pulmonary hypertension, right ventricular hypertrophy and pulmonary arterial medial wall thickness in vivo (GYY4137, 37.75 mg/kg i.p.) • Activation of Akt/PI3K signalling in hyperoxic lungs • Decreased age-dependent mitochondrial and cellular oxidant production in <i>C. elegans</i> (Qabazard et al. 2014; Hine et al. 2015) • Increased life span in <i>C. elegans</i> and <i>D. melanogaster</i> (Xiao et al. 2012; Peter et al. 2013)
	5a, 5b, 5c, 5d, 5e, GYY4137	<ul style="list-style-type: none"> • GYY4137 (200 μM) induced substantial cell killing to rat H9c2 myocardial cells (Putz et al. 1979) in contrast to other studies (Yin et al. 2009) but no toxicity observed with equimolar concentrations of 5a, 5b or NaSH. Marked toxicity observed with 5c (≥ 50 μM) • GYY4137 potentiated H₂O₂-induced cell death but cytoprotection observed with 5a and 5b (50–200 μM) (Putz et al. 1979) • GYY4137 caused nearly 100 % killing murine B16BL6 melanoma cells (100–200 μM). Some cell killing also observed with 5a and 5b
Cancer	GYY4137	<ul style="list-style-type: none"> • GYY4137 (100 μM to 1 mM) induced G2/M phase cycle arrest, PARP-/caspase-9-dependent cell death in seven disparate tumorigenic cell lines in vitro (Lee et al. 2011) • Cell killing through intracellular acidification and lactate accumulation and decreased anion exchanger and Na⁺/H⁺ exchanger activity • Inhibition of tumour cell growth in vivo in mouse HL-60 and MV4-11 xenografts (Lee et al. 2011) • No effect of NaSH or an inert analogue of GYY4137 (Lee et al. 2011) on cancer cell killing (Lee et al. 2014) • Inhibition of HepG2 and Bel7402 cell growth by suppression of STAT-3 activation (Lu et al. 2014) • Inhibition of HepG2 cell growth in mouse xenografts in vivo (Lu et al. 2014)

(continued)

Table 1 (continued)

Therapeutic area	Compound	Observation/comment
Inflammation	GYY4137	<ul style="list-style-type: none"> • Inhibition of LPS-induced IL-1β, TNF-α, PGE₂ and •NO synthesis in murine RAW264.7 macrophages by 50–500 μM GYY4137, effect conserved with prophylactic and therapeutic treatment. In contrast, NaSH potentiated cellular inflammation (Whiteman et al. 2010b) • Inhibition of cytokine synthesis in human synovial fibroblasts and articular chondrocytes (GYY4137 100–500 μM) (Qabazard et al. 2014) • Inhibition of NF-κB activation and activity in LPS- and oxLDL-treated (Whiteman et al. 2010b; Benedetti et al. 2014) RAW264.7 macrophages and <i>M. fermentans</i>-infected human monocytic U937 cells and monocytes by GYY4137 (100–400 μM) (Benedetti et al. 2014) • Inhibition of NF-κB activity, COX-2, prostaglandin E synthase, MMP-13 and iNOS expression and activity in human chondrocytes isolated from osteoarthritis patients (Burguera et al. 2014) • Inhibition of NF-κB activation, TNF-α, IL-1β and IL-6 synthesis in Cocksackie virus B-infected rat cardiomyocytes (Zhi et al. 2007) • Inhibition of <i>M. fermentans</i>-induced MCP-1 production in human monocytes (2 mM GYY4137) • Inhibition of oxLDL-induced LOX-1 and iNOS expression and synthesis of macrophage-specific chemokines (Chen et al. 2010) • Inhibition of iNOS, COX-2 and TACE activity (Li et al. 2013) • Inhibition of TNF-α synthesis in LPS-treated rat whole blood (Li et al. 2009) • GYY4137 normalised blood pressure, inhibited tissue inflammation and blood levels of IL-1β, TNF-α, IL-6 and increased IL-10 in LPS-induced sepsis in rats (Li et al. 2009) • Inhibition of serum-induced airway smooth muscle cell proliferation and IL-8 synthesis (100 μM) (Perry et al. 2011) • Therapeutic administration of GYY4137 (50 mg/kg i. p.) 6 or 18 h post-CFA treatment reduced knee joint inflammation and swelling; reduced synovial levels of TNF-α, IL-1β, IL-8 and IL-6; and inhibited neutrophil and macrophage synovial accumulation (Li et al. 2008) • Decreased blood levels of IL-6 and TNF-α and aortic expression of ICAM-1 synthesis and LOX-1 and mitochondrial “ROS” generation in apoE^{-/-} atherosclerotic mice with 30-day treatment of GYY4137 (133 μmol/kg/day i.p.) (Liu et al. 2013)

In the perfused rat kidney and heart, further differences between a bolus NaSH and GYY4137 are illustrated (Li et al. 2008). NaSH (10 nM) caused a transient fall in renal perfusion pressure, but higher doses (5 μM) caused a biphasic response, decreasing and then markedly increasing perfusion pressure, whereas no consistent effect was observed with GYY4137 (0.4–4.0 μM). Furthermore, in U46619, angiotensin II or noradrenaline-pre-constricted renal vessels, the addition of GYY4137 (100–500 μM) to the perfusion buffer caused concentration-dependent relaxation, an effect reversed when GYY4137 was removed or when angiotensin II was used in the presence of the K_{ATP} channel antagonist PNU3788A. In Langendorff-perfused rat heart, NaSH (100 μM) markedly reduced heart rate and left ventricular diastolic pressure. This effect was not observed with GYY4137 (50–200 μM) (Li et al. 2008). It is prudent to note that disparate effects of NaSH on heart rate have been reported: either no effect (Minamishima et al. 2009; Wang et al. 2011b; Shi et al. 2007; Alves et al. 2011; Derwall et al. 2010), tachycardia (Sodha et al. 2009) or bradycardia (Porokhya et al. 2012; Sikora et al. 2014). Although these observations have been made using different models and strains of animals, inconsistencies, such as these, question the reliability of sulfide salts to examine the physiological properties and pharmacological potential of H_2S .

The therapeutic potential for slow-release H_2S donors in vascular disease is also highlighted in Li et al. (2008). Although GYY4137 had negligible effect on mean arterial pressure (MAP) in normotensive WKY rats, spontaneously hypertensive rats, chronic dosing with GYY4137 (133 $\mu\text{mol/kg}$ i.v.) resulted in significant lowering of MAP within 2 days which persisted for 14 days. After withdrawal of GYY4137 treatment, MAP returned to basal levels but only after a further 14 days. In an additional model of hypertension, using L-NAME to inhibit the synthesis of endogenous $\bullet\text{NO}$, acute injection of GYY4137 (133 $\mu\text{mol/kg}$ i.v.) for 30 min before L-NAME treatment prevented L-NAME-induced increase in MAP. This preventative effect persisted for more than 60 min. In sharp contrast, a bolus of the sulfide salt NaSH (2.5 $\mu\text{mol/kg}$ i.v.) initially increased MAP above that of L-NAME treatment alone and failed to significantly decrease MAP, further reinforcing the marked differences between slow release of H_2S and an instantly generated bolus. More recently, Wang et al. showed pre-eclampsia in mice and humans led to the loss of vascular H_2S . Pharmacological inhibition or genetic silencing of CSE resulted in pre-eclampsia in mice, and “ H_2S supplementation” with GYY4137 (0.25 mg/kg i.p.) normalised mean blood pressure, increased foetal growth and lowered blood levels of soluble endoglin and soluble fms-like tyrosine kinase-1 (sFlt-1), the endogenous inhibitor of VEGF and placental growth factor (Wang et al. 2013).

3.5 GYY4137

GYY4137 is widely available from commercial suppliers, and its high solubility in water makes it a highly valuable laboratory tool. It is interesting to note that despite this high water solubility, commercial suppliers of GYY4137 state the compound

should be stored in organic solvent such as DMSO. We have found DMSO to be detrimental and accelerate GYY4137 decomposition and avoid storing the compound in organic solvents by preparing fresh solutions for each experiment immediately before use.

One major drawback of using GYY4137 is that due to inefficient hydrolysis, high concentrations are often required to elicit a biological response. Typically, concentrations in excess of 100 μM (and up to 100 mM) (Table 1) have been used. However, H_2S concentrations from GYY4137 will be considerably lower and should ideally be monitored to assess the effective concentration of H_2S . Similarly, the effects of a “decomposed” or “spent” donor control should be investigated to ensure any observed effects are due to “released” H_2S and not due to effects from the unhydrolysed parent compound or intermediates and/or hydrolysis products formed during GYY4137 hydrolysis and H_2S generation. These control experiments are generally not carried out (Table 1), but they are crucial since commercial GYY4137 (or GYY4137 prepared as described (Li et al. 2008)) is sold as a dichloromethane complex, with the dichloromethane being residual from initial synthesis and crystallisation. However, the molecular weights stated do not reflect the contribution of dichloromethane to the final mass of the molecule, and as such, previous publications with GYY4137 may contain erroneous, albeit lower, concentrations of the donor than reported. Additionally, dichloromethane is metabolised to carbon monoxide (CO) (Fagin et al. 1980; Mackison et al. 1981; Benzon et al. 1978; Baselt 1988; Barrowcliff and Knell 1979; Putz et al. 1979) an additional “gasotransmitter”. Since CO is reported to exert similar biological effects as H_2S (e.g. vasodilation, anti-inflammatory actions, modulation of cGMP signalling, etc.) (Peers et al. 2015; Peers and Lefer 2011; Nassour et al. 2015; Munoz-Sanchez and Chanez-Cardenas 2014), it is possible that some of the observed effects of GYY4137 could also be due, at least in part, to a contribution by CO. This may be especially important given the inefficient manner in H_2S generated from GYY4137 necessitates the use of high concentrations/doses of this compound to elicit pharmacological responses in cells, tissues and whole animals (see Table 1). An obvious solution to this problem is to employ control experiments with a time-expired (e.g. “spent” or “decomposed”) donor which would contain GYY4137 hydrolysis products (Alexander et al. 2014) as well as dichloromethane and morpholine to control for this possibility. However, these control experiments are not generally reported (see Table 1). An additional approach to circumvent this problem is to use structurally similar molecules devoid of residual solvent and organic counter ions.

3.6 GYY4137: Structural Modifications

Although the above studies consistently show GYY4137 exerted prominent anti-inflammatory, cytoprotective, anticancer and antihypertensive activity in vitro and in vivo, it is possible that slow-release H_2S donor compounds that are able to generate H_2S with greater efficiency than GYY4137 could offer improved efficacy.

Some structural modifications of GYY4137 such as *O*-substituted 1,8-diazabicyclo [5.4.0]undec-7-ene salts have been reported (Park et al. 2013), but they have not resulted in increased biological activity. For example, *O*-substituted GYY4137 derivatives such as compounds 5a–e (Fig. 1) showed no significant difference in H₂S release rates in physiological buffers or inside cells using rat H9c2 myocytes and the sulfide-specific fluorescent probe WSP-1 compared to GYY4137. Presumably, these structural changes did not improve H₂S release rates or efficiency, because of increased stability of the phosphorodithioate core by *O*-substitution (Park et al. 2013). Although, as yet, no mechanistic pathway appears to have been postulated, it would be anticipated that the generation of H₂S from the phosphoramidodithioates 5a–5e would follow a similar hydrolytic pathway to that elucidated for similar phosphorodithioates (Alexander et al. 2014). In contrast to previous studies using the same cell line (Wei et al. 2014), this report showed GYY4137 (200 μM) was cytotoxic, and although compounds 5a–c (Fig. 1) were markedly less toxic to H9c2 cells compared to GYY4137, they did not significantly increase the killing of mouse melanoma B16BL6 cells over that of GYY4137. In cytotoxicity assays using H9c2 myocytes, oxidative stress (using H₂O₂), modest (~10–20 %) but statistically significant inhibition of cytotoxicity was observed with 100–200 μM of 5a, 5b and 5c, but greater cytoprotection was observed with only 50 μM GYY4137, although it is intriguing to note that in this study, 200 μM GYY4137 potentiated H₂O₂-induced cell killing in sharp contrast to the results published in several studies elsewhere (Fox et al. 2012; Hine et al. 2015), including H9c2 myocytes (Wei et al. 2014). Since these compounds offered no significant advantage of GYY4137 in terms of biological activity and generation of H₂S, their physiological and/or pharmacological potential is not clear. However, these compounds have yet to be investigated in more complex *ex vivo* or *in vivo* models.

We have also modified the phosphorodithioate core of GYY4137 (Fig. 1) with the aim of improving H₂S generation and to have a selection of compounds which generate H₂S at different rates to use in the laboratory. This approach is analogous to the NONOates used extensively, and to great effect, to investigate the physiological role and therapeutic potential of •NO. Compounds with a range of different release characteristics could be used to more accurately model endogenously generated H₂S under basal conditions (e.g. using GYY4137) and induced H₂S synthesis (e.g. using AP67, AP105 and AP106) allowing for the examination of effects of H₂S concentration and H₂S generation rates on cells, tissues and animals. We have evaluated the biological activity of these compounds *in vitro* using cultured human synoviocytes and rat arterial smooth muscle cells and *ex vivo* using precontracted mouse aorta, and these studies clearly showed that the cellular and tissue effects of the H₂S donors were dependent upon the rates at which the compounds generate H₂S. For example, Fig. 2 compares the H₂S generation from equimolar donors in a physiological buffer (Fig. 2a) or in human synoviocytes (Fig. 2b) (e.g. AP67 > AP72 > AP105 > AP106 > GYY4137). Figure 3 shows that inhibition of LPS-evoked formation of •NO and PGE₂ in murine RAW264.7 macrophages and human synovial fibroblasts, respectively, was also dependent upon the rate of H₂S release (e.g. AP67 > AP72 > AP105 > AP106 = GYY4137).

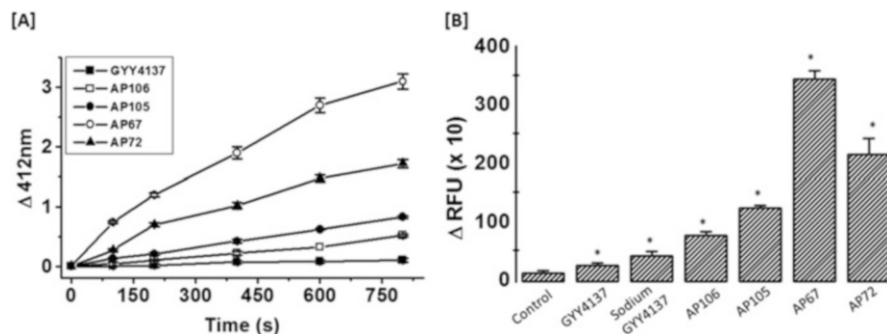


Fig. 2 H₂S generation from GYY4137 derivatives in the presence and absence of cultured cells. (a) H₂S generation in phosphate buffer saline was assessed using 5'-dithiobis-(2-nitrobenzoic) acid as described (Li et al. 2008). The final concentration of each H₂S donor was 500 μ M. (b) Intracellular H₂S was determined in human synovial fibroblasts cultured as described (Li et al. 2013) using the intracellular sulfide probe WSP-1 (10 μ M) (Szczyzny et al. 2014; Le Trionnaire et al. 2014). H₂S donors were added to cells at a final concentration of 100 μ M. Data are expressed as mean \pm SEM; *n* = 6

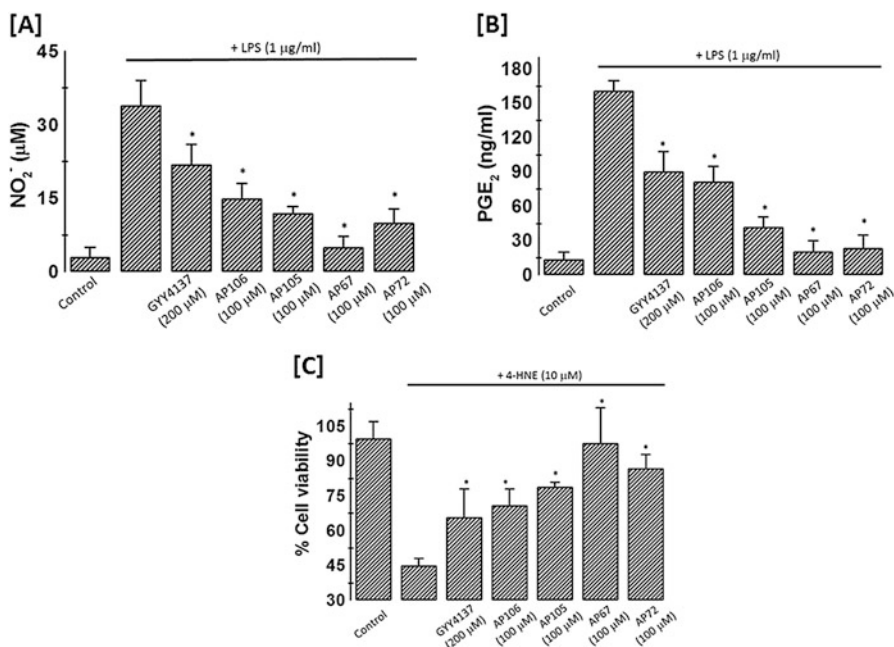


Fig. 3 H₂S release rate-dependent anti-inflammatory and cytoprotective activity of GYY4137 derivatives. Anti-inflammatory: RAW264.7 cells (a) and human synovial fibroblasts (b) were cultured as described (Li et al. 2013; Minamishima et al. 2009) and exposed to LPS (1 μ g/mL) in the presence or absence of H₂S donors (100 μ M) for 24 h. Levels of (a) nitrite (NO₂⁻, as index of \bullet NO synthesis) and (b) PGE₂ were measured in cell culture supernatant after 18 h by Griess assay and ELISA, respectively (Li et al. 2013; Whiteman et al. 2010b). Cytoprotection: human Jurkat T cells were cultured as described (Le Trionnaire et al. 2014) and exposed to 4-hydroxynonenal (10 μ M) in the presence or absence of 100 μ M H₂S donors. Cytotoxicity was assessed by trypan blue exclusion assay after 24 h. Data are expressed as mean \pm SEM; *n* = 6. **p* < 0.05, cf. LPS treatment (a, b) or 4-HNE-treated cells (c)

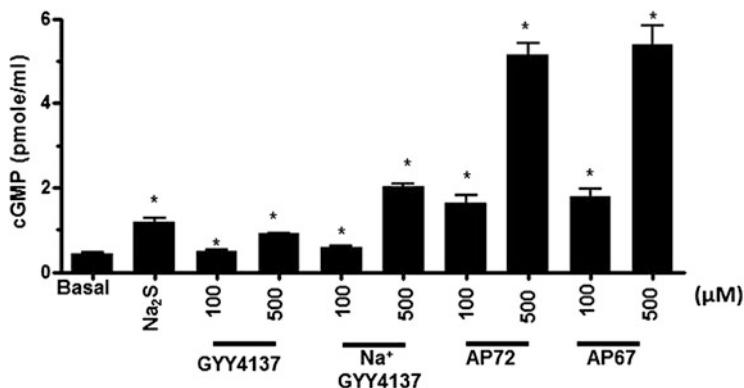


Fig. 4 H₂S released from donors enhances cGMP levels in rat aortic smooth muscle cells. Cells were isolated from 12–14-week-old male Wistar rats as described (Papapetropoulos et al. 1994). Cells between passages 2 and 5 were seeded on 24-well plates and after they reached confluence were incubated with the indicated H₂S donor at the concentrations stated in Hank's balanced salt solution for 5 min. Cellular cGMP was then extracted using 0.1 N HCl and quantified by ELISA (Assay Designs). Data are expressed as mean ± SEM; *n* = 4, **p* < 0.05, cf. basal cGMP levels

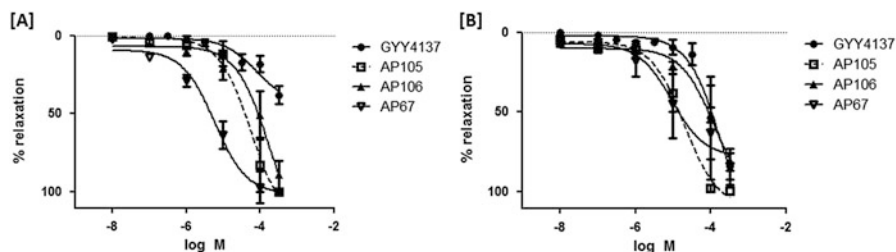


Fig. 5 H₂S release rate-dependent dilation of phenylephrine-precontracted mouse aorta. Mouse aortic rings were prepared as described in Bucci et al. (2012). (a) Endothelium-dependent vessel relaxation: EC₅₀ GYY4137, 80.9 μM; AP106, 167.4 μM; AP105, 57.5 μM; AP67, 5.9 μM. (b) Endothelium-independent vessel relaxation in denuded mouse aorta: EC₅₀ GYY4137, 186.1 μM; AP106, 120.2 μM, AP105 19.8 μM; AP67, 9.3 μM. Data are expressed as mean ± SD of three separate determinations

Additionally, the ability of the donors to induce cGMP accumulation in rat arterial smooth muscle cells was also dependent upon the rate of H₂S generation (Fig. 4). Intriguingly, the sodium salt of GYY4137 was more effective at stimulating cGMP formation, perhaps due to some inhibitory action of dichloromethane or morpholine in GYY4137. Furthermore, the H₂S release rate-dependent effects were also observed in phenylephrine-precontracted mouse aortic rings (Fig. 5), although this effect was not so clear in denuded aortic rings lacking endothelial cells suggesting the mechanisms for vessel relaxation are complex. However, the effects of the donors (e.g. H₂S release rate effects) were conserved in cells and in isolated blood vessels. The vasodilatory activity of AP67 and AP72 free acids (Fig. 1) has

also been shown to be more potent than GYY4137 in dilating precontracted bovine ciliary arteries *ex vivo* with markedly lower EC50 (AP67, $0.08 \pm 0.04 \mu\text{M}$; AP 72, $4.3 \pm 0.09 \mu\text{M}$; GYY4137, $13.4 \pm 1.9 \mu\text{M}$) (Chitnis et al. 2013, 2015) through K_{ATP} and prostaglandin-dependent mechanisms. The reasons for the differences in rate of H_2S release between GYY4137 and the other closely related phosphorodithioate donors such as AP67, AP105, etc., are not clear and are the subject of current and detailed chemical studies. However, since these molecules generate H_2S at different rates and elicit different effects on isolated cells and blood vessels, it is likely they will be useful and attractive additions to GYY4137 and alternatives to sulfide salts, for unravelling the complex physiology and pharmacology of H_2S where H_2S concentration and rate of H_2S generation need to be modulated and/or controlled.

3.7 Phosphorodithioate H_2S Donors: Summary

In summary, slow-release phosphorodithioate H_2S donors are powerful tools to study the physiological significance of H_2S in a wide range of cellular and animal models. By slowly generating fluxes of low concentrations of H_2S , they are more biologically relevant than using a bolus of instantly generated H_2S from often impure sulfide salts such as NaSH (and Na_2S) which are frequently used at supra-physiological concentrations/doses.

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Medicinal Chemistry: Insights into the Development of Novel H₂S Donors

Yu Zhao, Armando Pacheco, and Ming Xian

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Abstract

Hydrogen sulfide (H₂S) was traditionally considered as a toxic gas. However, recent studies have demonstrated H₂S is an endogenously generated gaseous signaling molecule (gasotransmitter) with importance on par with that of two other well-known endogenous gasotransmitters, nitric oxide (NO) and carbon monoxide (CO). Although H₂S's exact mechanisms of action are still under

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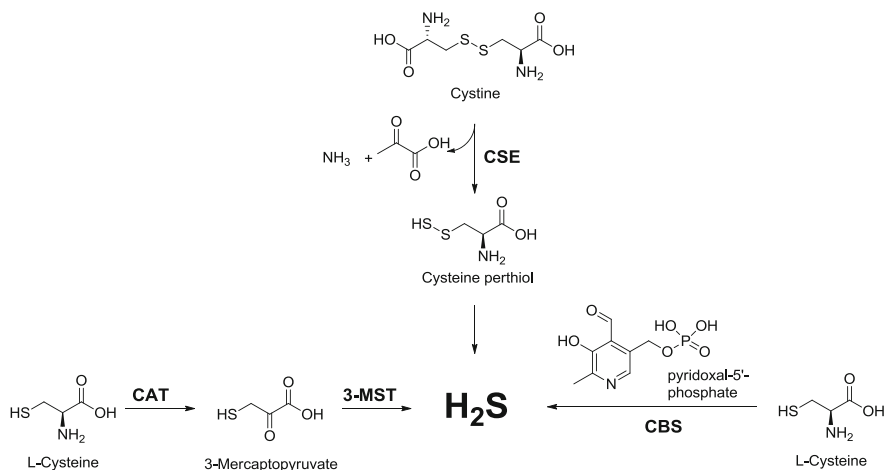
investigation, the production of endogenous H₂S and the exogenous administration of H₂S have been demonstrated to elicit a wide range of physiological responses including modulation of blood pressure and protection of ischemia reperfusion injury, exertion of anti-inflammatory effects, and reduction of metabolic rate. These results strongly suggest that modulation of H₂S levels could have potential therapeutic values. In this regard, synthetic H₂S-releasing agents (i.e., H₂S donors) are not only important research tools, but also potential therapeutic agents. This chapter summarizes the knowledge of currently available H₂S donors. Their preparation, H₂S releasing mechanisms, and biological applications are discussed.

Keywords

Hydrogen sulfide • H₂S donors • H₂S releasing • Synthesis

1 Introduction

Hydrogen sulfide (H₂S) is a colorless, flammable, and toxic gas which has been considered a poisonous air pollutant for centuries. It is well known for its characteristic odor of rotten eggs. Low concentrations of H₂S irritate the eyes, nose, throat, and respiratory system; high concentrations of H₂S lead to rapid loss of consciousness and death (Evans 1967; Li and Lancaster 2013; Lindenmann et al. 2010). Interestingly, this gaseous molecule has been recently classified as a member of the gasotransmitter family, sharing similarities with its congeners nitric oxide (NO), carbon monoxide (CO), and dioxygen (O₂) (Vandiver and Snyder 2012; Wang 2012; Li and Moore 2011; Szabó 2007; Kolluru et al. 2013; Fukuto et al. 2012; Olson et al. 2012; Olson 2012). The biosynthesis of H₂S in mammalian tissues has been attributed to enzymes, including cystathionine β-synthase (CBS), cystathionine γ-lyase (CSE), and 3-mercaptopyruvate sulfur-transferase (3-MST), in combination with cysteine amino transferase (CAT) (Scheme 1) (King 2013; Martelli et al. 2012a; Szabo 2012; Paul and Snyder 2012; Hu et al. 2011; Prabhakar 2012). It appears that CBS is predominant in the central nervous system. Brain homogenates produce H₂S in the presence of cysteine and pyridoxal-5'-phosphate. CSE is mainly found in the aorta, portal vein, and other vascular tissues. CSE converts cysteine disulfide (cystine) to cysteine perthiol, pyruvate, and ammonia. The resultant cysteine perthiol is highly reactive and unstable. Cleavage of the perthiol (S-SH) bond leads to H₂S generation. In comparison, 3-MST catalyzes H₂S formation in mitochondria (Whiteman et al. 2011). Kimura et al. demonstrated that together with cysteine aminotransferase (CAT), 3-MST was capable of producing H₂S from L-cysteine in the presence of α-ketoglutarate (Shibuya et al. 2009). Recently, it was also found that D-cysteine might be another substrate of 3-MST, as it can be converted to H₂S by 3-MST in the presence of D-amino acid oxidase (Shibuya et al. 2013). In addition to these enzymatic pathways, H₂S can also be produced by some “labile pools” of sulfide (such as persulfide, acid labile, and alkaline labile pools).



Scheme 1 Biosynthesis of H₂S

The production of H₂S in living systems is believed to have physiological reasons. In 1996, Abe and Kimura first revealed that endogenous H₂S acted as a neuromodulator in the brain (Abe and Kimura 1996). In their studies, CBS was found to be highly expressed in the hippocampus and cerebellum. They proved that high concentrations of H₂S (>320 μM) inhibited synaptic transmission in the hippocampus, and physiological concentrations of H₂S facilitated hippocampal long-term potentiation and enhanced *N*-methyl-D-aspartate receptor-mediated responses, which is the evidence that H₂S is involved in associative learning. Following this work, a number of studies have revealed many other biological activities of H₂S, including anti-inflammation, vasodilation, cardioprotection, anti-cancer, etc. (Köhn et al. 2012; Jackson-Weaver et al. 2013; Ariyaratnam et al. 2013; Predmore et al. 2012; King and Lefler 2011; Calvert et al. 2010a; Kimura et al. 2010; Kimura and Kimura 2004; Wen et al. 2013; Sivarajah et al. 2009; Lee et al. 2014). Associated with these discoveries, many biological reactions of H₂S have been disclosed. The oxidation state of the sulfur atom in H₂S is -2; therefore, it is a reductant species. H₂S is also a potent nucleophile under physiological pH. These properties make H₂S very reactive to many biological molecules. For example, H₂S can readily participate in 1 and 2 electron redox reactions. 1e-oxidation should form a highly reactive HS[•] radical, while the biological significance of this process is not well understood. H₂S can also proceed 2e redox reactions with many biologically important reactive oxygen species (ROS) such as hydrogen peroxide (H₂O₂), superoxide (O₂⁻), and peroxynitrite (ONOO⁻) (Filipovic et al. 2012a; Carballal et al. 2011). Because of these reactions, H₂S has been suggested to be a potent oxidant scavenger in vivo. Another interesting reaction is that H₂S can react with *S*-nitrosothiols to form thionitrous acid (HSNO), the smallest *S*-nitrosothiol, which may serve as a cell-permeable nitrosylating agent (Filipovic et al. 2012b). Recently, protein *S*-sulfhydration, i.e., generating *S*-SH groups on cysteine residues, has been suggested to be an important

oxidative post-translational modification mediated by H₂S. This reaction provides a possible way by which H₂S alters the functions of a wide range of cellular proteins and enzymes. However, the detailed mechanism(s) is still under investigation. It is well accepted that H₂S cannot directly react with protein cysteines (–SH) to form –S-SH adducts. It is possible that H₂S reacts with cysteine-modified residues, such as disulfides (–S–S–), sulfenic acids (S-OH), nitrosothiols (–S-NO), etc., to form S-sulfhydrated products. It is also possible that protein cysteines directly react with certain H₂S metabolites or precursors, such as sulfane sulfurs and hydrogen polysulfides, to yield S-SH products (Paulsen and Carroll 2013; Pan and Carroll 2013; Zhang et al. 2014; Mustafa et al. 2009; Gadalla and Snyder 2010; Sen et al. 2012; Krishnan et al. 2011; Vandiver et al. 2013; Paul and Snyder 2012; Yang et al. 2013). Nevertheless, more studies are needed to further understand this process, and reliable methods for the detection of S-sulfhydration would be the key.

Coordination chemistry of H₂S with transition metals has been well studied. Due to its soft character, H₂S binds tightly to coppers, and this property has been used in the development of fluorescent probes (Sasakura et al. 2011). Copper ion-mediated activation of odorant receptors via sulfur-copper coordination was found to contribute to human's high sensitivity to the smell of volatile sulfur compounds even at very low concentrations (Duan et al. 2012). From the toxicity point of view, H₂S interacts with cytochrome c oxidase (CcO) to inhibit mitochondrial respiration (involving reduction and tight sulfide coordination to copper center), which accounts for its toxicity to mammals. On the other hand, the interesting property of H₂S to induce a suspended-animation-like state (Blackstone et al. 2005) is believed to associate with a reversible inhibition of CcO via fine-tuned sulfide coordination chemistry that involves reaction with both the copper and iron centers (Collman et al. 2009).

To understand biological functions of H₂S and develop H₂S-related therapy, compounds that can release H₂S are often needed. Such compounds, known as H₂S donors, are used to mimic H₂S production under physiological conditions or disease states. In the past several years, the development of novel H₂S donors has been a rapidly growing field. Several types of donors have been reported (Martelli et al. 2012b; Kashfi and Olson 2013; Zhao et al. 2014a, b; Caliendo et al. 2010). These donors release H₂S under different mechanisms. It should be noted that the functions of H₂S reported in literature sometimes appear inconsistent. For example, in addition to anti-inflammatory effects, H₂S has also been claimed as a pro-inflammatory agent (Li et al. 2005; Tamizhselvi et al. 2008). These disparate results may be due to the different H₂S donors used in research. H₂S-releasing capabilities of the donors are quite different, therefore leading to very different results. Moreover, the by-products formed after H₂S release are different, and many times the effects of by-products were overlooked. Because of these reasons, the selection of suitable H₂S donors is crucial. Herein, we provide a review on currently available H₂S donors. H₂S-releasing profiles of selected compounds will be summarized, and their representative biological activities will be discussed.

2 H₂S Donors

2.1 Sulfide Salts

Inorganic sulfide salts such as sodium sulfide (Na₂S) and sodium hydrogen sulfide (NaHS) have been widely used as the equivalents of H₂S in research. A number of H₂S-regulated biological functions have been identified by using these sulfide salts. For example, Parkinson's disease (PD) is characterized by a progressive loss of dopaminergic neurons in the substantia nigra and depletion of the neurotransmitter dopamine in the striatum (Moore et al. 2005). The Bian group demonstrated that systemic administration of H₂S (using NaHS) could attenuate behavioral symptoms and dopaminergic neuronal degeneration in 6-hydroxydopamine (6-OHDA)-induced PD models (Hu et al. 2010). Using such models, they found that endogenous H₂S production was diminished during the development of PD, and the results suggested that administration of H₂S may have therapeutic benefits for Parkinson's disease. Similarly, H₂S was also found to be useful for Alzheimer disease (AD). H₂S biosynthesis in the brain is decreased in AD patients (Kamoun 2004). Giuliani and co-workers treated AD rats with different concentrations of NaHS (0.25, 0.5 and 1.0 mg/kg) and then studied the over time ability of AD rats to learn and remember. The results showed that AD rats receiving NaHS exhibited an improvement in learning and memory performance compared to the control group (Giuliani et al. 2013).

In addition to protective effects in nervous systems, H₂S has also shown protections in other tissues and organs. By employing Na₂S as the H₂S donor, Lefer et al. have proved short- and long-term cardioprotective activities of H₂S in a mouse model of myocardial ischemia injury (MI/R) (Calvert et al. 2009, 2010b). The evaluation of extent of myocardial infarction showed that mice receiving Na₂S exhibited a significant decrease in the infarct size. Na₂S also induced the reduction of oxidative stress and apoptosis following MI/R injury. Long-term treatment of Na₂S (7 days) resulted in a decrease in left ventricle dilatation, a decrease in cardiac hypertrophy, and an improvement in cardiac function. These findings provided strong evidence of the cardioprotective effects of H₂S therapy. Moreover, both anti-inflammatory and pro-inflammatory effects of H₂S have been reported when using sulfide salts as H₂S donors. Andruski and co-workers demonstrated that leukocyte-regulated inflammation of the knee joint was greatly inhibited after Na₂S injection (Andruski et al. 2008). In a separate study, Chen and co-workers proved that the administration of NaHS into the lungs in an asthmatic rat model attenuated inflammation (Chen et al. 2009). By using a smoke inhalation-induced lung injury model, Esechie et al. claimed Na₂S exhibited protective actions against severe lung injury (Esechie et al. 2008). All of these results suggest that H₂S (using NaHS or Na₂S) is a potent anti-inflammatory agent. On the other hand, pro-inflammatory effects of H₂S were also observed. For example, Li and co-workers found that H₂S exhibited pro-inflammatory activity in the animal model of endotoxic shock (Li et al. 2005). In addition, plasma H₂S concentrations, tissue H₂S synthesizing ability, and CSE expression were all found enhanced in

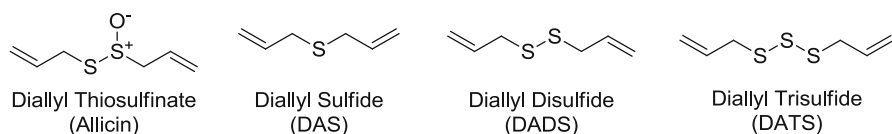
several animal models of inflammation (Bhatia et al. 2005; Mok and Moore 2008; Mok et al. 2004). These findings indicate the possible pro-inflammatory effects of H₂S.

Being effective H₂S donors, sulfide salts have the advantage of boosting H₂S concentration fast. In addition, they are “clean” donors, as no by-products would form after H₂S release. However, a major problem of this type of donors is that H₂S release is uncontrollable. Sulfide salts release H₂S immediately and completely as soon as they are dissolved in aqueous buffers. As a gaseous molecule, H₂S can be quickly lost from open stock solutions due to volatilization. Olson et al. found that the *t*_{1/2} of sulfide loss was about 5 min (DeLeon et al. 2012). Because of this property, modifications made between the time that a sulfide solution is prepared and the time that the biological effect is measured could dramatically affect the results. Unfortunately, this technique problem is difficult to solve and therefore often neglected. It may be the reason for some disparate conclusions.

It should also be noted that substantial polysulfide contaminations were detected in sulfide solutions, immediately upon sulfide salt crystal dissolution in water (Nagy et al. 2014). In fact, some commercially available sulfide crystals are yellow, indicating heterogeneous air oxidation on the surface. Polysulfides also form easily in sulfide solutions by metal ion-catalyzed air oxidation. It is very difficult and sometimes practically impossible to avoid sulfide oxidation. Therefore, appropriate control experiments should be carried out to determine whether the activity is from sulfide or from polysulfides.

2.2 Garlic-Derived Sulfur-Containing Compounds

Garlic is rich in sulfur-containing compounds and may be an active “H₂S” pool. The proven medical benefits of garlic (such as cardioprotective effects) have been recently revealed (at least in part) due to its inherent H₂S-releasing capability. Allicin (diallyl thiosulfinate) is the best characterized compound in garlic, and in aqueous solutions, it decomposes to a number of reactive sulfur-containing compounds including diallyl sulfide (DAS), diallyl disulfide (DADS), and diallyl trisulfide (DATS) (Scheme 2) (Amagase 2006). It was shown that glutathione (GSH)-rich human red blood cells (RBCs) could trigger H₂S release from garlic extract, as well as from DADS and DATS (Benavides et al. 2007). The fact that infusion of DADS in rats increased exhaled H₂S provides additional support for H₂S production from garlic in vivo (Insko et al. 2009). H₂S generation from these compounds is believed to be thiol dependent. The reaction should be initiated by the nucleophilic substitution of thiols (like GSH) at the α-carbon of the allyl



Scheme 2 Representative garlic-derived sulfur compounds

substituent, forming a perthiol intermediate, which then undergoes a thiol/disulfide exchange to release H₂S (Benavides et al. 2007).

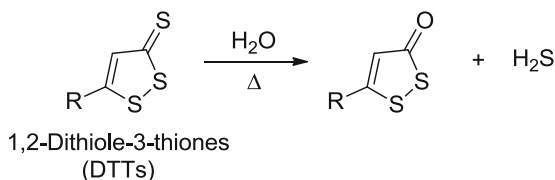
It is known that H₂S can induce vasodilation and increased garlic consumption in certain populations is associated with lower incidence of hypertension. The protective effects of garlic against renal, hepatic, cardiac, cerebral, and pulmonary ischemia/reperfusion injury are also well known (Sener et al. 2007). Garlic-derived compounds such as DADS and DATS can attenuate the deleterious effects of oxidized LDL on nitric oxide production (Lei et al. 2010) and attenuate myocardial ischemia/reperfusion in mice (Predmore et al. 2010). DADS is also known to inhibit 3-hydroxy-3-methylglutaryl-CoA and thus may be a potential antihyperlipidemic agent (Rai et al. 2009). It is also reported that oral administration of raw garlic homogenate in diabetic rats increased insulin sensitivity and reduced metabolic complications along with oxidative stress (Padiya et al. 2011). All these studies have revealed potential therapeutic applications of garlic-derived sulfur compounds.

2.3 1,2-Dithiole-3-Thiones

1,2-Dithiole-3-thiones (DTTs) have been well recognized as a H₂S-releasing core in the development of novel donors. However, the mechanisms of H₂S release from this type of molecules are still unclear. H₂S can be generated when heating DTTs to 120 °C in a DMSO-phosphate buffer mixture (Scheme 3), indicating that DTTs may be considered as hydrolysis-based H₂S donors (Qandil 2012). It was found that H₂S release from DTTs is a result of metabolic events, and enzymes such as esterases may be involved (Li et al. 2007). Nevertheless, this structure has been often used in making H₂S donor hybrid drugs. The pharmacological profiles of some lead candidates in the areas of inflammation, the cardiovascular system, urology, and neurodegenerative diseases have been reported. The hybrids often demonstrate significant improvements in activity and side effects, suggesting a useful pharmacological role of H₂S (Sparatore et al. 2011).

Nonsteroidal anti-inflammatory drugs (NSAIDs), such as aspirin and ibuprofen, have been widely used to treat inflammation. The usage of these drugs, however, is always accompanied with severe side effects (i.e., gastrointestinal ulceration and stomach bleeding) in alimentary systems. Recent studies revealed these unpleasant effects induced by NSAIDs could be significantly reduced by coupling DTT molecules, such as 5-(4-hydroxyphenyl)-3H-1,2-dithiole-3-thione (ADT-OH), with the parent NSAIDs (Caliendo et al. 2010). For example, Sparatore et al. synthesized a H₂S-aspirin hybrid (ACS14, *S*-aspirin) and compared the gastric

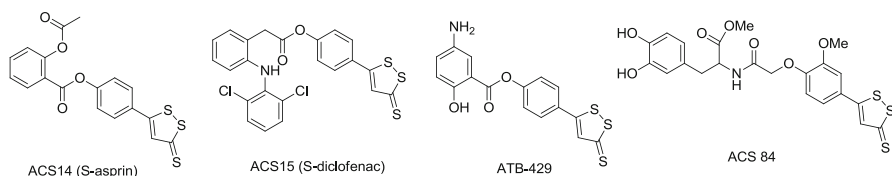
Scheme 3 H₂S release from DTTs



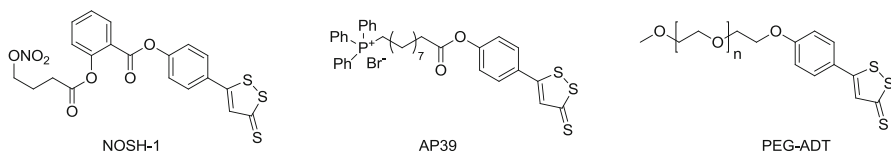
damages caused by ACS14 and aspirin in rats (Sparatore et al. 2009). They found that oral administration of aspirin to rats for 7 days led to extensive hemorrhagic lesions in the stomach. In comparison, rats receiving an equivalent dose of ACS14 did not exhibit such gastric damage. They also measured H₂S levels in plasma before and after ACS14 administration. The results showed that drug administration resulted in threefold increase of plasma H₂S concentration. These findings suggest that the advantages of ACS14 in gastric protection are likely related to the enhanced H₂S formation. ACS14 has also been shown to have strong protective effects against pathological cardiovascular alterations induced by buthionine sulfoximine (BSO), a GSH-synthase inhibitor (Rossoni et al. 2010). In this study, 7-day BSO treatment increased rats' systolic blood pressure and lowered their heart rates. Oral ACS14 treatment attenuated BSO-induced hypertensive effects. In comparison, aspirin failed to decrease the hypertension. BSO treatment resulted in a 77 % reduction of GSH levels, but ACS14 significantly reserved GSH levels (closed to control values). Moreover, BSO rats receiving ACS14 rather than aspirin displayed enhanced myocardial functions during myocardial reperfusion injury.

H₂S-releasing diclofenac (ACS15, *S*-diclofenac) has also been well studied. This hybrid drug showed improved anti-inflammatory activity as compared to diclofenac in several models. ACS15 significantly reduced lipopolysaccharide-induced infiltration of neutrophils into the lung and liver and did not stimulate leukocyte adherence, as had been observed with diclofenac (Li et al. 2007; Wallace et al. 2007). Another hybrid drug from mesalamine (ATB-429) has been well characterized in animal models of Crohn's disease and ulcerative colitis. In pre-clinical studies ATB-429 was found to be more effective than mesalamine in treating the visceral pain associated with inflammatory bowel disease (Scheme 4) (Fiorucci et al. 2007; Distrutti et al. 2006).

In addition to NSAIDs, DTTs have been conjugated with other pharmaceutical molecules to develop hybrids. For example, L-DOPA is used in the clinical treatment for PD and dopamine-responsive dystonia. Recent studies indicated that some side effects (such as the development of dyskinesia) of long-term L-DOPA treatment were due to the loss of neurons induced by oxidative stress (Fahn 1991; Rascol et al. 2000). Since H₂S is believed to diminish oxidative stress, L-DOPA coupled with a H₂S-releasing moiety might show improved bioactivity. Based on this idea, McGeer et al. synthesized a library of DOPA-DTT conjugates and investigated their antioxidant effects (Lee et al. 2010). The hybrid compounds (such as ACS84) were found to release H₂S when interacting with mitochondria.



Scheme 4 Representative H₂S-NSAIDs



Scheme 5 The structures of NOSH-1, AP39, and PEG-ADT

Further investigation also showed that the hybrids inhibited the dopamine-metabolizing enzyme MAO B, which could help to restore the PD-depleted dopamine levels in the brain.

The idea of hybrid drugs was not new. In fact, it had been applied to another gaseous transmitter, NO, and a number of NO-NSAIDs have been well studied (Kashfi 2009). Since H₂S and NO share many similar actions, it was postulated that the new hybrids that incorporated both H₂S and NO donors would be more potent than those incorporating either one alone. As such, some NO-H₂S-releasing hybrids (such as NOSH-1) were prepared and tested (Scheme 5) (Kodela et al. 2012; Chattopadhyay et al. 2012; Kashfi 2014). These compounds indeed can induce higher NO and H₂S concentrations in vivo. They showed very strong activities in inhibiting the growth of cancer cell lines, as well as good anti-inflammatory activities using the carrageenan rat paw edema model.

DTT has been coupled with a phosphonium salt to develop mitochondria-targeted H₂S donors. One of such donors, AP39, showed promising activities (Scheme 5) (Le Trionnaire et al. 2014; Szczesny et al. 2014). H₂S release from AP39 in mitochondria was confirmed by several fluorescent probes. Compared to standard and non-mitochondria-targeted donors such as GYY4137 and ADT-OH, AP39 significantly inhibited the oxidative stress-induced toxicity. It also attenuated the loss of cellular bioenergetics during oxidative stress and protected against oxidative mitochondrial DNA damage. These results suggest that mitochondria targeting is important.

Very recently, DTT-based polymeric H₂S donors have been reported (Scheme 5) (Hasegawa and van del Vlies 2014). In this work DTT was conjugated to poly(ethylene glycol) polymers. In murine macrophages the resultant polymers (PEG-ADT) showed slower and lower H₂S release as compared to DTT alone, presumably due to steric hindrance of PEG. The polymer conjugate was also found to be less toxic than the parent donor molecule. This effect was attributed to the colocalization of the PEG-ADT conjugate within the endolysosomes without escaping to the cytoplasm. This type of polymeric donors may have some unique applications.

It should be noted that although DTTs have been claimed as H₂S donors, to what extent their effects are actually related to H₂S is still not fully elucidated. DTTs can significantly elevate in vivo GSH levels and are suggested to be the inducers of enzymes involved in the maintenance of GSH balance (Zhang and Munday 2008). It is possible that the elevated GSH contributes much more than H₂S in those pharmaceutical benefits. Moreover, it is known that DTTs can participate in other

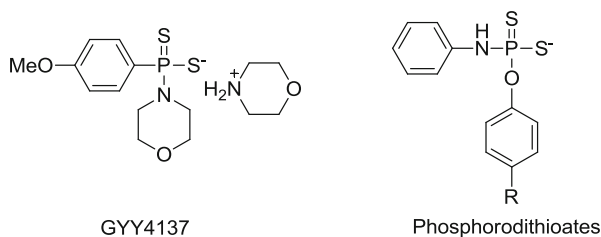
biological reactions such as electrophilic reactions (Talalay et al. 1988) and conversion of molecular oxygen to reactive oxygen radicals (Kim and Gates 1997). These possibilities should also be taken into consideration.

2.4 GYY4137 and Phosphorodithioate-Based Donors

In 2008, Moore and co-workers reported GYY4137, a Lawesson's reagent derivative, as a water-soluble H₂S donor. Since then, this compound has been widely used in the field (Scheme 6) (Li et al. 2008). GYY4137 releases H₂S very slowly and hydrolysis is believed to be the mechanism. In vitro experiments showed that 4.17 ± 0.5 nmol/25 min of H₂S was released from 1 mM of GYY4137. H₂S release from this donor is pH dependent with greater release at acidic pH (3.0) and less release at neutral or basic pH (7.4 and 8.5). Some recent studies also suggested that thiols may trigger H₂S release from GYY4137 (Martelli et al. 2013).

After the administration of GYY4137 to rats, H₂S concentrations in plasma could reach the maximal level after 30 min and maintain that level elevated over the 3-h experimental period. These results indicated that GYY4137, compared to NaHS, released H₂S in a much slower manner. Cytotoxicity test using rat vascular smooth muscle cells showed that GYY4137 did not cause significant cell damage under doses up to 100 μ M for up to 3 days. In contrast, NaHS (at similar doses and time course) promoted apoptotic cell death of smooth muscle cells (Baskar et al. 2007; Yang et al. 2004). This difference may be explained by different H₂S-releasing rates of these two donors. The slow release of H₂S is believed to be a major advantage of GYY4137.

The vasodilation effects caused by GYY4137 and sulfide salts were also compared. NaHS was found to initiate a fast and reversible (less than 30 s) relaxation of aortic rings. However, GYY4137 exhibited a much slower (~10 min) vasodilation. In a different study, Fitzgerald and co-workers demonstrated the single-cell level relaxation effects of GYY4137 on airway smooth muscle (ASM) (Fitzgerald et al. 2014). In this experiment, H₂S release from GYY4137 in culture media was confirmed by a methylene blue (MB) assay. A maximum of 275 μ M of H₂S was observed from 10 mM of GYY4137, and the generated H₂S was sustained in culture for 24 h. Compared to Na₂S, which induced acute relaxation of isolated ASM cells, GYY4137 led to a sustained relaxation presumably due to slow release.



Scheme 6 Phosphorodithioate-based donors

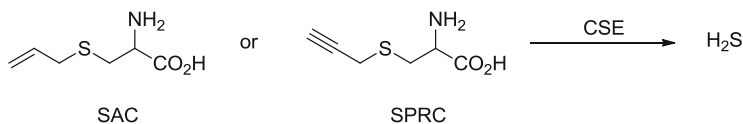
Other H₂S-related biological actions have also been observed for GYY4137. A recent study by Lee and co-workers claimed that GYY4137 exhibited anticancer effects both in vitro and in vivo (Lee et al. 2011). In this study, the MB assay indicated an extremely slow H₂S release (over 7 days) from GYY4137 in culture medium. After incubating several cancer cell lines with GYY4137, a significant reduction of cell proliferation was observed, indicating the potent anticancer effects of GYY4137. In comparison, the sulfur-lacking control compound, ZYJ1122, did not inhibit cell proliferation. Based on these findings, the authors claimed that the anticancer effects of GYY4137 were due to H₂S release.

The fixed structure and H₂S-releasing capability could limit the application of GYY4137 in different biological studies. By replacing the phosphorus-carbon bond in GYY4137 with phosphorus-oxygen bonds, Xian et al. reported a class of phosphorodithioate-based H₂S donors (Scheme 6) (Park et al. 2013). H₂S release from these donors was measured using a fluorescent probe. In general, these compounds were slow-release donors like GYY4137, but some structural modifications could completely shut down H₂S release. Selected donors were tested in cell-based experiments and found to have protective effects against H₂O₂-induced cell damage.

Although these phosphorodithioate-based compounds, especially GYY4137, have been widely used as H₂S donors, one should be careful when attributing their activities to H₂S. As have been demonstrated, these compounds released H₂S very slowly and at very low levels. Therefore, the dominated species in the testing systems should be the parent compounds, which might be bioactive themselves. In addition, their by-products (after H₂S release) have not been well identified, and the by-products' activities are unclear. Therefore, careful control experiments should be conducted when using these compounds as H₂S donors.

2.5 S-Allylcysteine and S-Propargyl-Cysteine

S-Allylcysteine (SAC) is a sulfur-containing amino acid. It is a cysteine derivative and a substrate for CSE (Scheme 7) (Kim et al. 2006). SAC has shown some interesting activities as an anti-oxidative (Herrera-Mundo et al. 2006; Numagami and Ohnishi 2001), anti-hepatotoxic (Hsu et al. 2006; Nakagawa et al. 1989), and anticancer agent (Chu et al. 2007; Welch et al. 1992). Padmanabhan and co-workers demonstrated that SAC exhibited cardioprotection in myocardial infarction (MI) by decreasing lipid peroxide products (Padmanabhan and Prince 2006). Zhu et al. also found that SAC upregulated the expression of CSE and led to enhanced H₂S



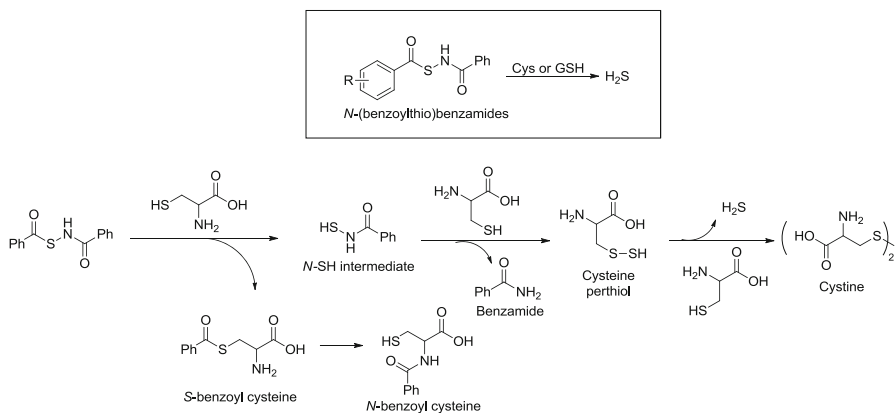
Scheme 7 H₂S release from SAC and SPRC

generation in the myocardium and plasma. The increased H_2S production then mediated the cardioprotection during acute MI (Chuah et al. 2007). It was found that rats receiving SAC prior to MI injury displayed a significant enhancement of plasma H_2S concentration. SAC-treated rats had smaller infarct size as compared to vehicle-treated ones, indicating SAC may be beneficial to the heart. *S*-Propargyl-cysteine (SPRC) is a structural analog of SAC (Scheme 7). Since SAC had been reported to be cardioprotective, SPRC was expected to show similar effects. Indeed, the MB assay revealed that rats receiving SPRC showed enhanced plasma H_2S concentrations. CSE activity was also increased. In addition, SPRC-treated MI rats were found to have smaller infarct sizes when compared to the vehicle group (Wang et al. 2009). These findings indicated that SPRC could be another protective molecule in MI, and the effects of SPRC are probably mediated by the CSE/ H_2S pathway. Other effects of SPRC, such as anticancer and anti-inflammation, have also been reported (Ma et al. 2011; Sidhapuriwala et al. 2012).

2.6 Thiol-Activated H_2S Donors

2.6.1 *N*-Mercapto-Based Donors

Xian et al. reported the first thiol-activated H_2S donors in 2011 (Scheme 8) (Zhao et al. 2011). Their design was based on the structure of *N*-mercapto (*N*-SH) compounds. It was expected that *N*-SH bonds are unstable and can be easily broken to form a hydrosulfide anion, which is the major form of H_2S under physiological conditions. This property triggered their idea of developing controllable H_2S donors based on the *N*-SH template. In Xian's work, acyl groups were used to protect the –SH residue of *N*-SH. The resultant compounds, i.e., *N*-(benzoylthio)benzamides, were found to be stable in aqueous buffers. Only in the presence of thiols (cysteine or glutathione) H_2S generation was observed. Structural-activity relationship of a series of such donors was studied in PBS buffers. It was found that H_2S release from



Scheme 8 H_2S release from *N*-SH-based donors

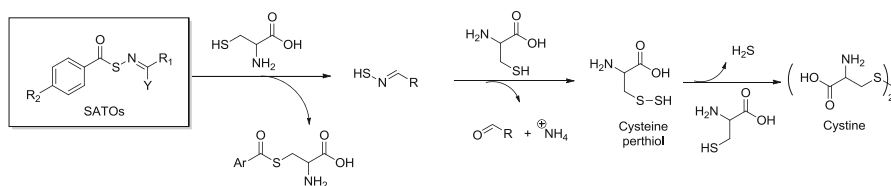
these compounds can be regulated by structural modifications. Electron-withdrawing groups caused faster H₂S generation, while electron-donating groups slowed down H₂S release. These donors were also evaluated in plasma and similar H₂S release profiles were observed. The mechanism of H₂S release from this type of donors was also studied. From reaction products isolated, the following mechanism was proposed: a thioester exchange between the donor and cysteine initiated the reaction. The resultant *S*-benzoyl cysteine was reactive and underwent a fast *S* to *N* acyl transfer to form a more stable *N*-benzoyl cysteine. Meanwhile, the free *N*-SH intermediate could react with another molecule of cysteine to form benzamide and cysteine perthiol, which further reacted with cysteine to produce cysteine disulfide (cystine) and release H₂S.

2.6.2 *S*-Aroylthiooximes

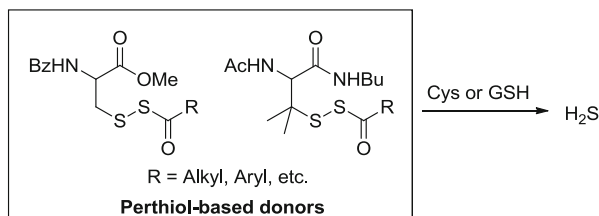
Matson and co-workers reported another type of *N*-SH-based H₂S donors based on *S*-aroylthiooxime (SATO) structures (Scheme 9) (Foster et al. 2014). These compounds were prepared by reacting substituted *S*-aroylthiohydroxylamines (SATHAs) with ketones or aldehydes. SATOs were found to be relatively stable in aqueous solutions under physiological pH. Hydrolysis of these compounds could be controlled by tuning the steric and electronic factors. H₂S release was triggered by thiols, and the half-life of H₂S release could be varied between 8 and 82 min by changing the substituents on the SATHA ring. The H₂S release mechanism from these donors was found to be similar to that of *N*-(benzoylthio)benzamides. The reaction was also initiated by a reversible thiol exchange between the SATO and cysteine to give the arylidenethiooxime along with *S*-benzoyl cysteine. Reaction of another molecule of cysteine with the arylidenethiooxime should produce cysteine perthiol, aldehyde, and ammonia. Finally, cysteine perthiol could react with cysteine to form H₂S and cystine.

2.6.3 Perthiol-Based Donors

As cysteine perthiol was found to be the key intermediate for H₂S generation from *N*-mercapto-based donors, perthiols were expected to be useful in the design of H₂S donors. In fact, cysteine perthiol is a “natural” H₂S donor, as it is involved in H₂S biosynthesis catalyzed by CSE (Caliendo et al. 2010; Stipanuk and Beck 1982). Therefore, perthiol-based donors may mimic H₂S bioproduction. With this idea in mind, a library of perthiol-based donors were reported by Xian and co-workers in 2013 (Scheme 10) (Zhao et al. 2013). These compounds were prepared as cysteine



Scheme 9 H₂S release from STAOS



Scheme 10 Perthiol-based H_2S donors

and penicillamine derivatives. Acyl groups were again used to protect the $-\text{SH}$ residue, so these donors were expected to be triggered by thiols. When H_2S -releasing profiles of these donors were evaluated in aqueous buffers (in the presence of thiols), cysteine-based donors were found to release very small amounts of H_2S (less than 10 % donor-to- H_2S conversion). This was due to an unwanted disulfide cleavage of the donors by thiols, leading to no H_2S production. In contrast, penicillamine-based donors showed high efficiency in H_2S generation (up to 80 % donor-to- H_2S conversion). Apparently, the two adjacent methyl groups prevented the cleavage of the disulfide bonds by thiols. Therefore, deacylation and subsequent H_2S generation dominated in the reaction, and H_2S production was enhanced. Mechanistic studies revealed that H_2S release from these donors followed a similar pathway as *N*-mercapto-based donors.

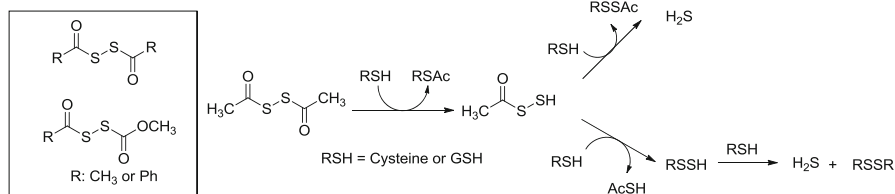
H_2S release from penicillamine-derived perthiol donors was demonstrated in cardiac myocytes (H9c2 cells) and in vivo. Cardioprotective effects of selected donors were also evaluated by a murine model of myocardial ischemia/reperfusion (MI/R) injury. Results showed that the administration of the donors to MI/R mice significantly reduced the infarct size, suggesting that these donors may have potential therapeutic benefits.

2.6.4 Dithioperoxyanhydrides

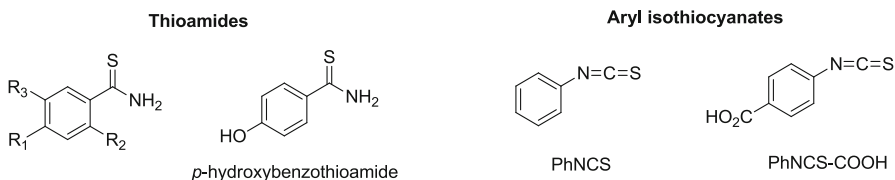
Dithioperoxyanhydrides have been reported as another type of thiol-activated H_2S donors (Roger et al. 2013). These compounds have some similarity with the perthiol-based donors, owing to the disulfide linkage in their structures. When treated with thiols, acylpersulfides were proposed to be formed as the key intermediate for H_2S production. As such, two possible pathways exist: (1) acylpersulfides could directly react with thiols to form H_2S and RSSAc ; (2) a new perthiol may be produced and in turn lead to H_2S and disulfide formation (Scheme 11). H_2S release from this type of donors was confirmed in both buffers and HeLa cell lysates. In addition, one donor, $\text{CH}_3\text{C}(\text{O})\text{SSC}(\text{O})\text{CH}_3$, was found to induce concentration-dependent vasorelaxation of pre-contracted rat aortic rings, presumably due to H_2S release.

2.6.5 Thioamide- and Aryl Isothiocyanate-Based Donors

In 2013, Calderone et al. reported that arylthioamides could be considered as a new class of thiol-activated H_2S donors (Scheme 12) (Martelli et al. 2013). However,



Scheme 11 Structures and H₂S release mechanism of dithioperoxyanhydrides

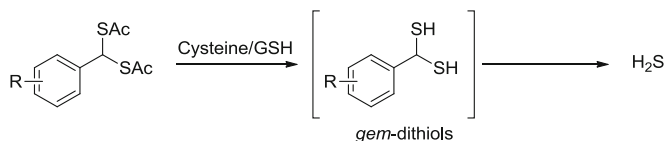


Scheme 12 Thioamide- and aryl isothiocyanate-based donors

these compounds seem to be weak and slow-release donors. Negligible amounts of H₂S were detected from the donors (1 mM) in the absence of thiols. Even when thiols were presented (4 mM cysteine or GSH), H₂S formation was low (~1 %). It is not clear how H₂S was formed, nor were the final products figured out. Nevertheless, these donors were tested for their effects on vasoconstriction. Pretreatment of aortic rings with a lead donor, *p*-hydroxybenzothioamide, significantly inhibited vasoconstriction induced by noradrenaline. Since the NaHS-treated group exhibited similar behaviors, the anti-vasoconstriction effects of *p*-hydroxybenzothioamide were proposed to be H₂S related. In addition, oral administration of *p*-hydroxybenzothioamide to rats caused a decrease of blood pressure, further demonstrating the protective actions of such donors in cardiovascular systems. It should be noted that in these experiments, neither cysteine nor glutathione was exogenously added. Endogenously presented biothiols seemed to be sufficient to promote H₂S release from the donors. Very recently, the same group reported that aryl isothiocyanate derivatives were another type of slow-release H₂S donors (Scheme 12) (Martelli et al. 2014). Thiol activation for these donors was also necessary. The lead compounds, such as PhNCS and PhNCS-COOH, showed vaso-relaxing effects on conductance and coronary arteries. However, detailed reactions behind these phenomena were unclear. It is likely that aryl isothiocyanates react with biomolecules (such as cysteine) to form thioamide derivatives, which then produce H₂S. Pharmacological effects of those possible products should also be studied.

2.6.6 *gem*-Dithiol-Based H₂S Donors

gem-Dithiols are known to be unstable in aqueous solutions, and H₂S is the decomposition product (Cairns et al. 1952; Berchtold et al. 1959). Based on this property, *gem*-dithiols were used by Xian and co-workers to develop thiol-activated



Scheme 13 H₂S release from acylated *gem*-dithiols

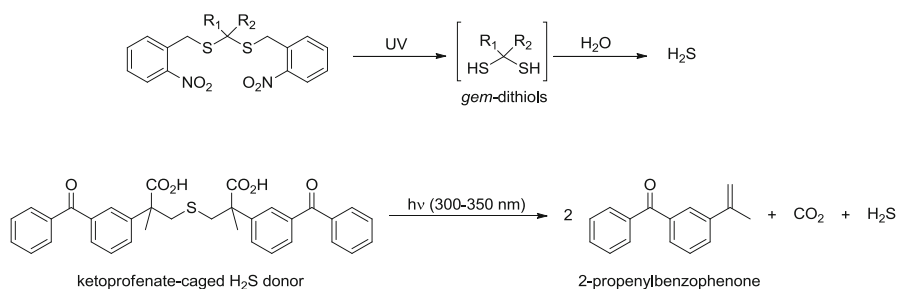
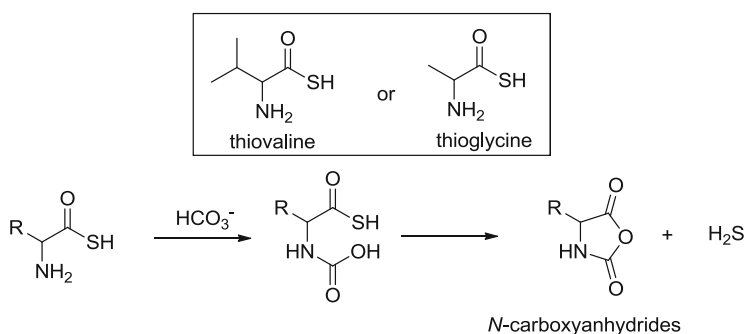
H₂S donors (Scheme 13) (Zhao et al. 2014a, b). In this study, *gem*-dithiols were stabilized by acylation and a series of such donors were synthesized. H₂S release from these donors was observed in the presence of cysteine or glutathione in PBS. Donor-induced H₂S release was also observed in HeLa cells (with endogenous thiols).

2.7 Photolabile H₂S Donors

gem-Dithiols were also used by Xian and co-workers to develop photolabile H₂S donors (Scheme 14) (Devarie-Baez et al. 2013). Their strategy was to introduce a photo-cleavable 2-nitrobenzyl group on *gem*-dithiols. The resultant compounds were expected to be sensitive to UV light. Indeed, such compounds were found to release H₂S only upon UV irradiation. This property was confirmed both in aqueous buffers and in HeLa cells. In another report, Nakagawa et al. utilized ketoprofenate as the photocage and developed another photosensitive donor (Scheme 14) (Fukushima et al. 2014). Their compounds can release H₂S with the elimination of 2 equivalents of 2-propenylbenzophenone and CO₂ upon irradiation at 300–350 nm. To evaluate the H₂S-releasing capability of the donor in complex biological systems, the donor was applied to fetal bovine serum. No H₂S was detected without irradiation. However, approximately 30 μM of H₂S was detected from 500 μM of the donor after irradiating for 10 min. Biological applications or pharmacological properties of these photo-inducible donors have not been reported.

2.8 Thioamino Acids

Thioamino acids, such as thioglycine and thiovaline, were reported to be H₂S donors in 2012 (Scheme 15) (Zhou et al. 2012). A unique property of these compounds is that they only release H₂S in the presence of bicarbonate. It was believed that bicarbonate reacted with the amine group of the donor first and then underwent a cyclization to form *N*-carboxyanhydride and produce H₂S. As such, thioamino acids may be effective H₂S donors given the high concentrations of bicarbonate in blood (~27 mM). In this work, both thioglycine and thiovaline were proven to enhance cGMP formation and promote vasorelaxation in mouse aortic rings.

**Scheme 14** Photolabile H₂S donors**Scheme 15** H₂S release from thioamino acids

3 Conclusions

Biomedical research on H₂S is a fast-growing field, and a number of H₂S donors have been developed in recent years. These compounds are not only useful research tools but also potential therapeutic agents. Current donors rely on different mechanisms to produce H₂S, and their H₂S production capabilities are also quite different. Therefore, it is impossible to define a universal best donor for all biological applications. One should carefully consider the donor's properties (such as how fast the release is, what the by-products are, cross-reactivity with other biomolecules, etc.) when choosing the donors for certain studies. The major problems associated with many widely used donors include the following: (1) H₂S release is uncontrollable, either too fast or too slow (such as sulfide salts or GYY4137, respectively); (2) H₂S release mechanism is still unclear, and the by-products after H₂S release are unidentified. Unless those by-products are identified and their biological activities are studied, it may be premature to claim the effects of the donors are due to H₂S. In addition, current efforts on "controllable donors" have been largely put on thiol-activation strategies. These donors (such as *N*-mercapto-/perthiol-based donors, dithioperoxyanhydrides, thioamides, and

thioisocyanates) deplete biological thiols to promote H₂S generation. Given the high endogenous concentrations of biothiols (such as GSH at mM levels), the anticipated active dose of these donors should be at low micromolar levels. Therefore, the consumption of free thiols by the donors may not cause significant changes in thiol redox balance. However, under some circumstance, free thiols' level might be low due to disulfide formation or binding to proteins. Caution should be taken when using these donors in such conditions, and careful control experiments are needed. Nevertheless, the development of H₂S donors is critical for understanding the biological functions of H₂S. We expect to see more interesting work arising in this field.

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