

Inflammation-associated S100 proteins: new mechanisms that regulate function

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Abstract This review focuses on new aspects of extracellular roles of the calgranulins. S100A8, S100A9 and S100A12 are constitutively expressed in neutrophils and induced in several cell types. The S100A8 and S100A9 genes are regulated by pro- and anti-inflammatory mediators and their functions may depend on cell type, mediators within a particular inflammatory milieu, receptors involved in their recognition and their post-translational modification. The S100A8 gene induction in macrophages is dependent on IL-10 and potentiated by immunosuppressive agents. S100A8 and S100A9 are oxidized by peroxide, hypochlorite and nitric oxide (NO). HOCl generates intrachain sulfinamide bonds; stronger oxidation promotes cross-linked forms that are seen in human atheroma. S100A8 is >200-fold more sensitive to oxidative cross-linking than low-density lipoprotein and may reduce oxidative damage. S100A8 and S100A9 can be S-nitrosylated. S100A8-SNO suppresses mast cell activation and inflammation in the microcirculation and may act as an NO transporter to regulate vessel tone in inflammatory lesions. S100A12 activates mast cells and is a monocyte and mast cell chemoattractant; a G-protein-coupled mechanism may be involved. Structure–function studies are discussed in relation to conservation and divergence of functions in S100A8. S100A12 induces cytokines in mast cells, but not monocytes/macrophages. It forms complexes with Zn²⁺ and, by chelating Zn²⁺, S100A12 significantly inhibits MMPs. Zn²⁺ in S100A12 complexes co-localize with

MMP-9 in foam cells in atheroma. In summary, S100A12 has pro-inflammatory properties that are likely to be stable in an oxidative environment, because it lacks Cys and Met residues. Conversely, S100A8 and S100A9 oxidation and S-nitrosylation may have important protective mechanisms in inflammation.

Keywords S100 · Calgranulins · S100A8 · S100A9 · S100A12 · Inflammation · Interleukin 10

Abbreviations

RA	Rheumatoid arthritis
IBD	Inflammatory bowel disease
ROS	Reactive oxygen species
NO	Nitric oxide
TLR	Toll-like receptor
mS100A8	Murine S100A8
mS100A9	Murine S100A9
TNF α	Tumor necrosis factor α
TGF β	Transforming growth factor β
IFN	Interferon
LPS	Lipopolysaccharide
IL	Interleukin
COX-2	Cyclo-oxygenase 2
cAMP	Cyclic adenosine monophosphate
MAP kinase	Mitogen-activated protein kinase
EC	Endothelial cells
FGF	Fibroblast growth factor
GC	Glucocorticoids
DEX	Dexamethasone
PPAR- γ	Peroxisome proliferator-activated receptor- γ
RAGE	Receptor for advanced glycation end products
NF κ B	Nuclear factor κ B

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EN-RAGE	Extracellular newly identified RAGE-binding protein
AGE	Advanced glycation end products
MCP-1	Monocyte chemotactic protein 1
NIF	Neutrophil immobilizing factor
NADPH	Nicotinamide adenine dinucleotide phosphate
NMR	Nuclear magnetic resonance
MPO	Myeloperoxidase
MMP	Matrix metalloproteinase

Introduction

Inflammation is a central component of acute responses to infection and wounding, and to numerous chronic human diseases including rheumatoid arthritis (RA), inflammatory bowel disease (IBD), atherosclerosis and allergic inflammation. Neutrophils and mast cells are important mediators of the acute innate immune response, but also contribute to the pathogenesis of chronic inflammatory diseases such as RA, asthma and other chronic lung disorders. Leukocyte recruitment to sites of infection/inflammation is mediated by generation of chemokines and other chemoattractants, and by cytokine stimulation of the vascular endothelium to facilitate leukocyte adhesion and extravasation. Initial activation of the innate immune system is usually self-limiting, but if this fails, chronicity ensues. In chronic inflammation, monocytes accumulate where they mature to macrophages that play an inflammatory and tissue destructive role. Activation of these cells by structures on pathogens known as pathogen-associated molecular patterns (PAMPs), through receptors such as the Toll-like receptors (TLRs) [reviewed in (Mogensen 2009)], modulates inflammatory processes via induction of inflammatory cytokines, reactive oxygen species (ROS) and nitric oxide (NO). However, macrophages are also important in resolution and interleukin (IL)-10 (which is produced by activated macrophages) and can induce anti-inflammatory effects via suppression of cytokine signaling 3 (SOCS3) (Gordon 2003).

In recent years, a subgroup of the S100 family (S100A12, S100A8 and S100A9) has been associated with acute/chronic inflammatory disorders (Foell et al. 2004a, b) and with various cancers (Salama et al. 2008). Collectively, these proteins are termed calgranulins, reflecting their calcium-binding properties and high expression in granulocytes. Despite intense research, the exact functions of these proteins remain enigmatic and disparate functional roles are proposed. This review will focus on the extracellular roles of these proteins in leukocyte migration and

chemotaxis, leukocyte activation, oxidant scavenging and their relevance in inflammatory processes.

Expression and gene regulation of calgranulins

The calgranulins are abundant neutrophil proteins; S100A8 and S100A9 comprise ~45% of the cytosol (Edgeworth et al. 1991). Constitutive expression is restricted to neutrophils, myeloid-derived dendritic cells, platelets, osteoclasts, hypertrophic chondrocytes and in trophoblasts in developing embryos. Myeloid cell expression is dependent on the stage of differentiation (Koike et al. 1992; Sellmayer et al. 1992). Monocytes constitutively express low levels of S100A8 and S100A9 (Hessian et al. 1993; Hsu et al. 2005), but this is reduced during their extravasation from blood; normal tissue macrophages do not express these proteins (Lagasse and Clerc 1988; Zwadlo et al. 1988). S100A8 and S100A9 are often strongly expressed in acute and chronic inflammatory lesions and one or both can be induced by appropriate stimulants in several cell types (see Table 1).

In addition to neutrophils, macrophages are often S100A8/A9 positive in inflamed tissues (Odink et al. 1987; Delabie et al. 1990; Goebeler et al. 1994; McCormick et al. 2005; Terasaki et al. 2007). Murine (m)S100A8 mRNA is induced in macrophages by TLR ligands, such as LPS (Hu et al. 1996) and dsRNA (Endoh et al. 2009), and by the cytokines, TNF α , transforming growth factor β (TGF- β) and interferon- γ (IFN γ) (Xu and Geczy 2000) and by IFN γ (Endoh et al. 2009). The stress response modulator norepinephrine also induces S100A8 and S100A9 in human monocytoïd cells, suggesting their regulation by stress (Suryono et al. 2006).

S100A8 induction by LPS and polyIC (a double-stranded RNA mimetic) is maximal at ~24-h post-stimulation and is strongly dependent on the earlier induction of the anti-inflammatory cytokine, IL-10. IL-10 alone has little effect, but is essential for, and potentiates, TLR ligand-induced responses (Xu et al. 2001; Endoh et al. 2009). S100A8 is not induced in stimulated macrophages from IL-10-null mice. We also demonstrated a crucial and convergent role for protein kinase R (PKR) in S100A8 induction by LPS and dsRNA, probably by virtue of its role in IL-10 induction (Xu et al. 2001; Endoh et al. 2009). TLR4, but not TLR3-mediated induction of mS100A8 mRNA, is also dependent on cyclo-oxygenase 2 (COX-2) via production of prostaglandin E₂ and cyclic AMP (cAMP) (Xu and Geczy 2000; Hsu et al. 2005). Stimulation of MAP kinases ERK1/2 and p38 pathways is required to generate responses to both agonists (Hsu et al. 2005; Endoh et al. 2009). Interestingly, although S100A8 and S100A9 are said to require co-expression for stability (Hobbs et al. 2003; Manitz et al. 2003), we find that murine S100A9

Table 1 Expression of S100A8 and S100A9 in various cell types

Cell type	Mouse		Human		Reference
	S100A8	S100A9	S100A8	S100A9	
Neutrophils	Constitutive	Constitutive	Constitutive	Constitutive	(Berntzen and Fagerhol 1988; Hessian et al. 1993)
Monocytes/macrophages	Inducible	Nil	Inducible	Inducible	(Hessian et al. 1993; Santhanagopalan et al. 1995; Hu et al. 1996; Xu and Geczy 2000; Xu et al. 2001; Hsu et al. 2005)
Myeloid dendritic cells	ND	ND	Constitutive	Constitutive	(Kumar et al. 2003)
Osteoclasts	Constitutive	Constitutive	Constitutive	Constitutive	(Zreiqat et al. 2007)
Microglia	ND	ND	Nil	Inducible	(Akiyama et al. 1994; Shepherd et al. 2006)
Eosinophils and basophils	ND	ND	Nil	Nil	(Hogg et al. 1989)
Lymphocytes	ND	ND	Nil	Nil	(Zwadlo et al. 1988)
Platelets	ND	ND	Constitutive	Constitutive	(Healy et al. 2006)
Microvascular endothelial cells	Inducible	Inducible	Inducible	Inducible	(Yen et al. 1997; McCormick et al. 2005)
Keratinocytes	Inducible	Nil	Inducible	Inducible	(Eversole et al. 1992; Eversole et al. 1993; Coleman and Stanley 1994; Grimbaldston et al. 2003; Eckert et al. 2004)
Fibroblasts	Inducible	Nil	ND	ND	(Rahimi et al. 2005)
Hypertrophic chondrocytes	Constitutive	Constitutive	Constitutive	Constitutive	(Zreiqat et al. 2007)
Trophoblasts	Transient	Nil	ND	ND	(Passey et al. 1999)

(mS100A9) is not induced in macrophages by any of these stimuli. In contrast, human S100A8 and S100A9 are generally co-induced in human monocytes and macrophages (Hsu et al. 2005; Endoh et al. 2009).

S100A8 and S100A9 induction in other cell types may be somewhat stimulus specific. In murine microvascular endothelial cells (ECs), mS100A8 is induced by LPS and IL-1 β , peaking 24 and 16-h post-stimulation, respectively, but not by TNF α or IFN γ . In contrast, mS100A9 is induced earlier by IL-1 β (maximal 8-h post-stimulation) and is also induced by TNF α in these cells (Yen et al. 1997). Co-stimulation with TNF α enhanced LPS responses, but not IL-1 β -mediated induction in vitro and in vivo. The S100s are not expressed in normal vessels, but mS100A8 is seen in microcapillaries in delayed-type hypersensitivity lesions in mice (Yen et al. 1997). Human S100A8 and S100A9 are expressed in neovessels, but not in large vessels or arteries in human atheroma (McCormick et al. 2005). ECs in lungs of mice bearing metastatic tumors also express these proteins, and vascular EC-derived growth factor and TNF α are implicated in their induction (Hiratsuka et al. 2006).

In murine fibroblasts, LPS, FGF-2 and IL-1 β , but not TNF α or IFN γ , induce mS100A8; induction by FGF-2 is markedly enhanced by heparin, but suppressed by TGF- β (Rahimi et al. 2005). Murine S100A9 was not induced with any stimuli tested (Rahimi et al. 2005). As for microvascular EC (Yen et al. 1997), the degree of cell confluence was important for expression in fibroblasts in vitro (Rahimi et al. 2005), indicating that cell-cell contacts may provide additional signals for optimal induction in these cell types.

S100A8/A9 is not found in normal skin, but the pattern of S100A8 expression in fibroblasts in rat dermal wounds suggests its involvement in fibroblast-to-myofibroblast differentiation at sites of inflammation and in repair/remodeling. In keeping with gene regulation studies, TGF- β mediates the later stages of wound healing when S100A8 levels decline (Rahimi et al. 2005). S100A8/A9 expression has been associated with the hyperproliferative responses typical of psoriasis (Gabielsen et al. 1986) and with wound healing (Thorey et al. 2001). We also found S100A8, but not S100A9 induction in UVA-irradiated skin of mice, and upregulation in a keratinocyte cell line by peroxide (Grimbaldeston et al. 2003). Both responses were suppressed by anti-oxidants, confirming ROS-mediated induction (Grimbaldeston et al. 2003). Cultured human keratinocytes also express S100A8 and S100A9 in response to TNF α , IFN γ (Mork et al. 2003) and IL-1 α (Hayashi et al. 2007). TGF- β and retinoic acid reduced induction, suggesting a relationship with keratinocyte differentiation (Hayashi et al. 2007). The late induction of these genes in keratinocytes also suggests that they are secondary-response gene products.

Our finding that S100A8 induction in macrophages is IL-10-dependent suggested that this protein may have anti-inflammatory properties. In keeping with this, we found that glucocorticoids (GC) directly upregulated S100A8 and S100A9 in human monocytes, and S100A8-positive macrophage numbers increased in rheumatoid synovium in RA patients treated with high-dose GC (Hsu et al. 2005). The synthetic GC, dexamethasone (DEX), strongly enhances

LPS-induced mS100A8 induction in murine macrophages (Hsu et al. 2005). Since the promoter region of the mS100A8 gene lacks a steroid-response element, and because anti-IL10 suppresses induction by DEX, this response is likely dependent on IL-10 induction by GC (Hsu et al. 2005). Increased levels of mS100A8 were also detected in the lungs of DEX-treated mice after intranasal administration of LPS (Bozinovski et al. 2005). GC also induced S100A8 in human dendritic cells (Kumar et al. 2003) and in a human keratinocyte cell line and potentiated its induction by IL-17 (Bai et al. 2007). However, DEX suppressed mS100A8 and mS100A9 expression associated with phorbol ester-induced skin inflammation in wild-type, but not in *c-fos*^{-/-} mice, suggesting that these are negatively regulated *c-fos/AP-1*-target genes in skin (Gebhardt et al. 2002). We also found that DEX potentiated S100A8 gene induction by LPS in primary murine fibroblasts and microvascular EC (Hsu et al. 2005). Because IL-10 and GC are anti-inflammatory and immunosuppressive [reviewed in (Couper et al. 2008) and (Wilckens 1995)], this pattern of gene expression indicates that S100A8 may be involved in the resolution phase of inflammation and may be more consistent with the proposed oxidant-scavenging role (Lim et al. 2009) than the pro-inflammatory role recently proposed via ligation of TLR4 (Vogl et al. 2007).

S100A12 is constitutively expressed in neutrophils at low levels (~5% of cytosolic protein) (Guignard et al. 1995), and expression patterns in myeloid cell lines (Vogl et al. 1999a, b) suggest that it is expressed later in myeloid cell differentiation than S100A8 and S100A9 (Robinson and Hogg 2000). PolyIC, LPS and TNF α induce S100A12 in monocytes (Yang et al. 2001; Endoh et al. 2009). Induction by TNF α is more rapid than for S100A8 and S100A9, with optimal S100A12 mRNA induction at 6–12 h (Yang et al. 2001). IL-6 also upregulates S100A12 expression in differentiated THP-1 macrophages and is dependent on the JAK-STAT pathway and de novo protein synthesis (Hasegawa et al. 2003). Pioglitazone, a peroxisome proliferator-activated receptor- γ (PPAR- γ) agonist, inhibited S100A12 mRNA induction in these cells (Hasegawa et al. 2003). Ligand-bound PPAR- γ represses signal-dependent transcription of many inflammatory cytokines through interactions with promoter-bound co-regulatory proteins (Pascual and Glass 2006). The rapid upregulation by pro-inflammatory stimuli and repression by PPAR γ agonists is consistent with a pro-inflammatory role for S100A12. In keeping with this, and in contrast to S100A8 and S100A9, we found that GC does not upregulate the S100A12 gene in activated human monocytes (K. Hsu, C. Geczy, unpublished data).

S100A12 is expressed by monocytes and/or macrophages in giant cell arteritis (Foell et al. 2004a, b), atherosclerotic lesions (Goyette et al. 2009) and macrophages

in asthmatic lung (Yang et al. 2007), rheumatoid synovium (Yang et al. 2001) and inflamed mucosa in IBD (Foell et al. 2003; Leach et al. 2007). S100A12 mRNA is detected in fetal bovine esophagus and skin (Hitomi et al. 1996), protein found in corneal fibroblasts (keratocytes) (Gottsch et al. 1995, 1999a, b), differentiating esophageal epithelial cells (Hitomi et al. 1998) and in human keratinocytes in psoriatic lesions, but not normal skin (Mirmohammadsadeh et al. 2000). IL1 α and TNF α increase S100A12 expression in bovine keratocytes (Gottsch et al. 1999a, b). Although circulating eosinophils do not express S100A12, eosinophils in the airways of asthmatic lung express S100A12 (Yang et al. 2007), indicating that it is also inducible in these cells and the mechanisms are currently being characterized by us.

Calgranulins and clinical conditions

There has been significant interest in realizing the clinical diagnostic potential of monitoring calgranulin concentrations in body fluids. Unlike acute-phase reactants primarily produced by the liver (e.g., C-reactive protein), these proteins are mostly expressed in cells in inflammatory lesions. Thus, calgranulin levels in the circulation are elevated in inflammatory conditions characterized by infiltration of neutrophils or activation of monocyte/macrophages or other stromal cells. Since leukocyte recruitment and activation is common to many inflammatory conditions, circulating levels may not be appropriate for diagnosis of any particular condition, although calgranulins are suggested to be non-specific markers of phagocyte activation (Foell et al. 2004a, b, 2007). On the other hand, samples derived directly from an inflammatory source (e.g., feces, sputum) may be more reliable markers of inflammatory disorders such as IBD (Foell et al. 2009) or eosinophilic asthma (Yang et al. 2007).

S100 proteins lack a signal peptide for secretion via the Golgi-mediated pathway and there is some debate regarding whether the high levels derived from inflammatory lesions are due to active secretion or passive release as a consequence of neutrophil necrosis. S100A8/A9 release correlates with loss of neutrophil viability (Voganatsi et al. 2001) and because these are short-lived, necrosis probably represents a significant extracellular source, particularly in situations such as acute inflammatory lesions or chronic conditions such as cystic fibrosis or RA, where neutrophil infiltration is significant. However, activation of protein kinase C by pro-inflammatory stimuli and elevation of [Ca²⁺]_i by contact with activated endothelium (Frosch et al. 2000) can also provoke S100A8/A9 release from phagocytes. LPS from several bacterial species (Kido et al. 2005) and chemoattractants such as C5a (Hetland et al.

1998) cause rapid release from neutrophils. Although mechanisms are unclear, secretion of S100A8 and S100A9 from monocytes and neutrophils is tubulin dependent (Rammes et al. 1997; Ryckman et al. 2004). Secretion following activation of neutrophils with monosodium urate crystals involves activation of CD11b, CD16, Src kinases, Syk and tubulin polymerization (Ryckman et al. 2004).

Secretion may also be cell-type specific. For example, S100A8 and S100A9 in activated microvascular ECs, fibroblasts and keratinocytes are mainly cell associated (Yen et al. 1997; Grimbaldston et al. 2003; Ravasi et al. 2004; Rahimi et al. 2005), implying that these proteins have predominantly intracellular roles in these cells. In contrast, stimulated monocytes and macrophages readily secrete the three calgranulins (Rammes et al. 1997; Xu and Geczy 2000; Xu et al. 2001; Hasegawa et al. 2003; Hsu et al. 2005; Endoh et al. 2009).

Extracellular functions

Calgranulins are implicated in numerous intracellular functions in phagocytes (reviewed in (Nacken et al. 2003; Roth et al. 2003; Gebhardt et al. 2006; Foell et al. 2007; Lim et al. 2009)), however this review will focus on some newer aspects of their extracellular roles. Tables 2 and 3 summarize some key extracellular functions proposed for human and murine S100A8, S100A9, the heterodimer (S100A8/A9) and for S100A12. However, there are several conflicting reports regarding their activities, making assessment of some functions difficult. Because the S100A8 and S100A9 genes are regulated by pro- and anti-inflammatory mediators, together with their emerging anti-inflammatory properties, the functions of these proteins may depend on several factors including the cell type and mediator involved in their induction in a particular inflammatory milieu, the receptors involved in their recognition and their post-translational modification. For this reason, we propose that they are mislabeled as 'pro-inflammatory' and in some circumstances we believe that their high expression levels may actually mediate host protection. For example, S100A8 and S100A9 expression patterns in incision wounds in skin lead to the suggestion that elevated levels may have a therapeutic effect against inflammation in wound healing (Wu and Davidson 2004), and their positive expression in skin may be crucial for the protection against bacterial infections (Champaiboon et al. 2009) and UVA-mediated oxidative damage (Grimbaldston et al. 2003). Interestingly, S100A8/A9 in stools from healthy newborns is elevated over 3–30 days and may be involved in the normal response to microbial infection (Baldassarre et al. 2007). Although S100A8/S100A9 has been associated with chondrocyte-mediated cartilage

destruction in RA (van Lent et al. 2007), anti-inflammatory properties of the injected proteins are indicated in rodent models of arthritis (Brun et al. 1995), autoimmune myocarditis (Otsuka et al. 2009) and endotoxin-mediated inflammation (Ikemoto et al. 2007). In addition, S100A9 has antinociceptive properties that are effective in various acute inflammatory pain models, and the use of a C-terminal peptide may represent a potential therapeutic (Paccola et al. 2008). Taken together with the pro-inflammatory properties described for these proteins, their ability to modulate inflammation should also be considered. Functions may be governed by the receptors that are activated by the calgranulins. In addition, post-translational modifications may occur in oxidative environments and other factors such as proteolytic cleavage by proteases released as a consequence of phagocyte activation (Greenlee et al. 2006), and zinc binding may also have important functional ramifications.

Endogenous ligands of TLR4?

Extracellular S100A8/S100A9 is reported to bind numerous receptors on cells. One of the most important may be the ability of murine S100A8 to bind TLR4; interaction between immobilized mS100A8 and recombinant TLR4 was demonstrated by surface plasmon resonance ($K_d \sim 11\text{--}25$ nM) (Vogl et al. 2007). Stimulation of murine bone marrow cells (a population normally comprising $\sim 30\%$ neutrophils, immature myeloid cells and haematopoietic stem cells) with mS100A8 activated nuclear factor κB (NF κB) via the MyD88-dependent pathway and secretion of TNF α . In contrast, mS100A9 or the mS100A8/A9 complex did not stimulate TNF α release, although the complex enhanced TNF α production in response to LPS (Vogl et al. 2007). Bone marrow cells from mS100A9 $^{-/-}$ mice are less responsive to LPS stimulation and the mice are resistant to LPS-induced septic shock (Vogl et al. 2007). The latter is not surprising, considering the fact that neutrophils from these mice do not migrate normally to various chemotactic stimuli (Manitz et al. 2003; Vogl et al. 2004; McNeill et al. 2007; Schnekenburger et al. 2008). The authors claim that mS100A9 $^{-/-}$ mice are also deficient in mS100A8 (essentially mS100A8 $^{-/-}$) because this is absent in circulating neutrophils (Hobbs et al. 2003; Manitz et al. 2003; Nacken et al. 2005) and thus implicate mS100A8/A9 in LPS-induced septic shock. However, mS100A8 was found in bone marrow cells in mS100A9 $^{-/-}$ mice (Manitz et al. 2003) and mS100A8 levels in serum, or from activated macrophages from these mice, which has not been reported.

In contrast to these findings, Ikemoto et al. (2007) showed that intraperitoneal injection of S100A8/A9 into

Table 2 Extracellular functions of human and murine S100A8, S100A9 and S100A8/A9

Protein	Suggested function		
Human S100A8	Integrin expression	Upregulates Mac-1 expression, induces high-affinity form and L-selectin shedding (Ryckman et al. 2003a, b), contradicted by reports showing that S100A8 has no effect, but that S100A9 is active (Newton and Hogg 1998; Anceriz et al. 2007)	
	Chemotaxis	Chemotactic for neutrophils (Ryckman et al. 2003a, b), contradicted by reports showing that S100A8 is not chemotactic for neutrophils or monocytes (Lackmann et al. 1993; Newton and Hogg 1998); chemotactic for periodontal ligament cells (active region is amino acids 21–45) (Nishimura et al. 1999) Fugetactic for neutrophils (oxidation sensitive) (Sroussi et al. 2006)	
	Oxidation	Protects from oxidation by scavenging oxidants (Harrison et al. 1999; Raftery et al. 2001; Raftery and Geczy 2002; Grimbaldston et al. 2003; McCormick et al. 2005), acts as an NO shuttle (Lim et al. 2008)	
	Cell metabolism	N-terminal peptide reduces glucose consumption/lactate production in myoblasts (Basso et al. 2006)	
	Human S100A9	Phagocyte migration	Induces high-affinity form of Mac-1, without L-selectin shedding or Mac-1 upregulation (Newton and Hogg 1998; Anceriz et al. 2007); another report shows upregulation of Mac-1 expression (Ryckman et al. 2003a, b)
Chemotaxis		Chemotactic for neutrophils (Ryckman et al. 2003a, b), contradicted by report showing that S100A9 is not chemotactic for neutrophils (Newton and Hogg 1998) C-terminal peptide has sequence identity with neutrophil immobilization factor (Goetzl and Austen 1972; Goetzl et al. 1973; Freemont et al. 1989), but function has not yet been validated with synthetic peptide Fugetactic for neutrophils (Sroussi et al. 2007)	
Inflammation-induced pain		Antinociceptive activity (Pagano et al. 2002; Pagano et al. 2006)	
Macrophage function		Decreases O ₂ ⁻ and H ₂ O ₂ release from <i>Mycobacterium bovis</i> -activated macrophages (S100A8 has no effect) (Aguiar-Pasesti et al. 1997)	
Atherosclerosis		Promotes dystrophic calcification in atheroma (McCormick et al. 2005)	
Fibrin formation/lysis		C-terminal peptide is homologous to high molecular weight kininogen; full-length protein inhibits fibrin formation (Hessian et al. 1995)	
Human S100A8/A9 heterocomplex		Phagocyte migration	Induces high-affinity form of Mac-1, L-selectin shedding and upregulates Mac-1 expression (Ryckman et al. 2003a, b), contradicted by reports showing that S100A8/A9 has no effect, but that S100A9 is active (Newton and Hogg 1998; Anceriz et al. 2007)
		Chemotaxis	Chemotactic for neutrophils (Ryckman et al. 2003a, b) and induces migration of prostate cancer cells (Hermani et al. 2006)
	Arachidonic acid transport	Intercellular transport of arachidonic acid via CD36-mediated S100A8/A9-arachidonic acid uptake (Kerkhoff et al. 2002)	
	Antimicrobial defense	Anti-microbial activity, particularly for <i>Candida albicans</i> (Steinbakk et al. 1990; Murthy et al. 1993; Sohnle et al. 2000)	
	Apoptosis	Induces apoptosis of numerous cell lines (Yui et al. 2002; Viemann et al. 2007)	
	MMP activity/expression	Inhibits MMP-1, -2, -3, -7, -8, -9, and -13 by sequestering Zn ²⁺ (Isaksen and Fagerhol 2001) and increases expression of MMP-2 in human gastric cancer cell line (Yong and Moon 2007)	
	Propagation of inflammation	Activation of NFκB in prostate cancer cells via ligation of RAGE (Hermani et al. 2006)	
	Anti-inflammatory	Injection of S100A8/A9 decreased serum levels of IL-6 and NO induced by LPS in rats (Ikemoto et al. 2007)	
	Amyloid deposits	Contributes to dystrophic calcification and amyloid deposits in aging prostate (Yanamandra et al. 2009), found in senile plaques of Alzheimers disease (Shepherd et al. 2006)	
Endothelial cell activation	Induces mild increases in transcription of pro-inflammatory cytokines and adhesion molecules (Viemann et al. 2005), potentiates RAGE-mediated pro-inflammatory cytokine and adhesion molecule production (Ehlermann et al. 2006) and impairs endothelial cell integrity (Viemann et al. 2007)		

Table 2 continued

Protein	Suggested function	
Murine S100A8	Chemotaxis	Chemotactic for neutrophils and monocytes; hinge region is the active domain (Lackmann et al. 1992; Cornish et al. 1996)
	Leukocyte recruitment	S100A8-SNO inhibits mast cell activation and suppresses leukocyte adhesion and extravasation in the microcirculation of mice
	Propagation of inflammation	Nuclear translocation of NF κ B and secretion of TNF α from bone marrow cells via TLR4 ligation; mS100A9 inhibits mS100A8-mediated activation (Vogl et al. 2007)
	Cartilage destruction	Enhances cartilage destruction in antigen-induced arthritis by upregulating MMP-2, -3, -9 and -13 expression in macrophages (van Lent et al. 2007)
	Oxidation	Protects from oxidation by scavenging oxidants (Harrison et al. 1999; Raftery et al. 2001; Raftery and Geczy 2002; Grimbaldston et al. 2003)
Murine S100A9	Inflammation-induced pain	Antinociceptive activity (Dale et al. 2004; Dale et al. 2006)
	Fibroblast proliferation	Promotes fibroblast proliferation (rat S100A9) (Shibata et al. 2004; Shibata et al. 2005)
	Macrophage function	C-terminal peptide inhibits macrophage spreading and phagocytosis of adherent peritoneal lavage cells (Pagano et al. 2005) Decreases O $_2^-$ and H $_2$ O $_2$ release from <i>Mycobacterium bovis</i> -activated macrophages (S100A8 has no effect) (Aguir-Pasesti et al. 1997)

Table 3 Extracellular functions of S100A12

Suggested function	Suggested mechanism	Reference
Induces		
IL-1 β and TNF α production in BV 2 microglial cells	Ligation of the receptor for advanced glycation end products (RAGE)	(Hofmann et al. 1999)
IL-2 secretion by PBMCs		
ICAM-1 and VCAM-1 on ECs		
Does not induce IL-1 β , TNF α , IL-6 or IL-8 in human monocytes/macrophages		(Goyette et al. 2009)
Inhibits filial growth and motility, filaricidal activity	Binding of nematode paramyosin	(Gottsch et al. 1999a, b; Akpek et al. 2002)
C-terminal peptide is antimicrobial and antifungal	Zn $^{2+}$ enhances activity	(Cole et al. 2001)
Chemotactic for monocytes and mast cells	Ligation of a G-protein-coupled receptor; hinge region is active	(Yang et al. 2001; Yan et al. 2008)
Enhances Mac-1 affinity and L-selectin shedding on neutrophils	Ligation of RAGE (assumed without evidence)	(Rouleau et al. 2003)
Stimulates mast cell degranulation and potentiates IgE-dependent mast cell activation	Ligation of an undetermined receptor	(Yang et al. 2007)
Inhibits MMP-2, -3 and -9	Ligation of Zn $^{2+}$	(Goyette et al. 2009)
Stimulates neurite outgrowth in rat hippocampal neurons	Ligation of RAGE and activation of phospholipase C, protein kinase C, Ca $^{2+}$ fluxes, calmodulin-dependent kinase II, and MAP kinase pathways	(Mikkelsen et al. 2001)

rats 1 or 3.5 h after LPS injection significantly reduced serum IL-6 and nitrite levels, whereas administration of anti-S100A8/A9 IgG 1 h after LPS injection slightly increased IL-6 levels. This study suggests that the S100 complex bound pro-inflammatory cytokines and also reduced NO levels. The latter would concur with our

studies showing that S100A8 and mS100A8 are readily S-nitrosylated on Cys $_{41/42}$ (Lim et al. 2008) by NO donors and these may have scavenged the NO generated in vivo.

Using recombinant human or murine S100A8, S100A9 or the complex, we cannot induce significant levels of pro-inflammatory cytokines, in the presence/absence of LPS, in

monocytes/macrophages from the relevant species (K. Hsu, C. Geczy, unpublished data). Taken together with the mechanisms regulating S100A8 gene expression that suggest an anti-inflammatory role, it is difficult to reconcile our results with a strictly pro-inflammatory role. The proteins used in the study showing that mS100A8 binds TLR4 were endotoxin tested, and heat treatment reduced activity, but polymyxin B (which binds to, and neutralizes LPS) did not, indicating that the activity was unlikely due to LPS contamination (Vogl et al. 2007). However, these recombinants were generated using a bacterial expression system, and contamination with other bacterial components, such as lipoproteins, is possible (Tsan and Gao 2004). Similar issues with contamination have been problematic for research characterizing endogenous ligands of TLR, such as heat shock proteins (Bausinger et al. 2002; Gao and Tsan 2003a, b; Reed et al. 2003) and HMGB1 (Tian et al. 2007). Clearly, more research is needed to clarify the role of TLR4 in mediating extracellular functions of S100A8.

EN-RAGE and the ‘S100/RAGE pro-inflammatory axis’

When S100A12 was found to bind the receptor for advanced glycation end products (RAGE), Hofmann et al. named it extracellular newly identified RAGE-binding protein (EN-RAGE) to signify the importance of this receptor to its function (Hofmann et al. 1999). These studies were supported by the observation that apoS100A12 binds RAGE in vitro with very low affinity ($K_d \sim 140 \mu\text{M}$), which increases $> 1,000$ fold when S100A12 is in the Ca^{2+} - (Xie et al. 2007) or Zn^{2+} -bound hexameric state (Moroz et al. 2009). The hypothesis that RAGE is the sole receptor mediating the pro-inflammatory functions of S100A12 has enjoyed wide support and is often referred to as the ‘RAGE/S100 pro-inflammatory axis’. This theory can be summarized as follows: RAGE ligation by S100A12 presents as a consequence of inflammatory conditions, triggers a signaling cascade in microvascular ECs, macrophages (inferred from results using BV-2, a microglial cell line) and lymphocytes (also cell lines), culminating in $\text{NF}\kappa\text{B}$ activation. This amplifies inflammation and, since the promoter region of the RAGE gene contains an $\text{NF}\kappa\text{B}$ binding site (Li and Schmidt 1997), RAGE itself is consequently upregulated. Thus, RAGE ligation initiates a feed-forward loop that could potentiate inflammation and some authors refer to RAGE as a ‘master switch’ in chronic inflammation (Schmidt et al. 2001; Bierhaus et al. 2005).

This theory set the scene for further studies by several groups investigating the role of RAGE in various functions provoked by numerous S100 proteins. Because of their structural similarities, S100s generally, and particularly the calgranulins, have been proposed to bind RAGE. This has

often led to the erroneous use of the term ‘S100/calgranulins’ to refer to any or all S100s, perhaps implying functional redundancy. However, although the S100 proteins are structurally related, differences in expression profiles and subtle differences in amino acid sequences within functional domains may underlie non-redundant functions. For example, gene deletion of S100A8 in the mouse results in rapid and synchronous embryo resorption by day 9.5 of development, indicating a necessary function in trophoblast invasion and/or maternal recognition (Passey et al. 1999), whereas mice carrying a deletion of the S100A9 gene are apparently normal, but have somewhat compromised neutrophil function (Hobbs et al. 2003; Manitz et al. 2003; Vogl et al. 2004; McNeill et al. 2007). Moreover, particular S100 proteins have very different extracellular functions that can be cell type and/or concentration dependent such as those reported for S100B, which has RAGE-dependent and RAGE-independent activities (Sorci et al. 2003, 2004; Bianchi et al. 2007, 2008). Taken together, these observations do not support a theory for a common receptor for all S100 proteins.

We showed that S100A12 activates mast cells in vitro and in vivo, but failed to detect RAGE mRNA/protein in these cells, indicating an alternate receptor (Yang et al. 2007). In keeping with this, we showed that the chemotactic activity of S100A12 for monocytes and mast cells is RAGE independent and likely mediated by a G-protein-coupled receptor (Yan et al. 2008). Importantly, and in marked contrast to the finding that S100A12 induces $\text{IL-1}\beta$ and $\text{TNF}\alpha$ in a murine microglial cell line stimulated with bovine S100A12 (Hofmann et al. 1999), we failed to reproduce these results with human S100A12 and primary human monocytes or macrophages (Goyette et al. 2009). This, together with the fact that neutralizing anti-S100A12 antibodies would be ineffective in murine models used to confirm its role in cell-mediated immunity and IBD because S100A12 is not expressed in the rodent genome (Ravasi et al. 2004), indicates that the central role for RAGE may need to be re-evaluated.

In addition to TLR4, other receptors such as heparin, heparan sulfate (Robinson et al. 2002), CD36 (Kerkhoff et al. 2001) and N-glycans (Srikrishna et al. 2001) are implicated in extracellular S100A8/A9 functions. Thus, some functions attributed to RAGE may depend on cell type. RAGE is implicated in proliferation of human estrogen receptor-negative breast cancer cells (Ghavami et al. 2008a, b) and in S100A8/A9-mediated cardiomyocyte dysfunction (Boyd et al. 2008), but not in activation of prostate cancer cells (Hermani et al. 2006) or monocytes (Sunahori et al. 2006). Results from different binding assays are somewhat contradictory. Because S100A8/A9 protected the VC1 domain of RAGE (recombinant) from protease digestion (Ghavami et al. 2008a, b), binding is implicated

although S100A8/A9 does not bind RAGE-transfected CHO cells (Robinson et al. 2002) or interact with RAGE in surface plasmon resonance studies (Vogl et al. 2007). In contrast, RAGE was co-immunoprecipitated with anti-S100A8 and anti-S100A9 from LPS-stimulated cardiac myocytes (Boyd et al. 2008) and colocalized with S100A8/A9 in prostate cancer cells (Hermani et al. 2006). Recent evidence indicates that the glycosylation status of RAGE is important. RAGE purified from various sources is glycosylated and a small proportion of molecules (~1–5%) contain epitopes recognized by an N-glycan-specific mAb (mAbGB3.1) (Turovskaya et al. 2008). S100A8/A9 bound a similarly small fraction (~1%) of RAGE molecules; binding was abolished by deglycosylation of RAGE and blocked by mAbGB3.1, indicating that these novel N-glycans mediate the interaction (Turovskaya et al. 2008). These results may explain some of the variability in previous studies since N-glycans may be present on other membrane proteins, and the extent of RAGE glycosylation may differ between cell types.

As proposed for HMGB1, it is possible that some RAGE-dependent extracellular calgranulin functions may depend on a co-stimulus to activate co-operative signaling pathways or enhance receptor affinity. For example, HMGB1 augments CpG-stimulated IFN α production in DCs, and this is dependent on RAGE and TLR9 (Tian et al. 2007). Stimulation of transfected HEK293 cells with HMGB1–CpG complexes induced association of RAGE and TLR9 indicating that HMGB1 and RAGE may aid in CpG presentation to TLR9 (Tian et al. 2007). Alternatively, functions may be dependent on pre-activation of cells to induce other necessary gene products. For example, in EC pre-treated with AGE-modified albumin for 72 h, S100A8/A9 upregulate IL-6, ICAM-1, VCAM-1 and MCP-1, whereas only MCP-1 expression was enhanced in untreated cells (Ehlermann et al. 2006). AGE pre-treatment also upregulates RAGE expression in these cells and the authors assume that S100A8/A9 functions through this receptor (Ehlermann et al. 2006), though another putative RAGE ligand, S100B (Leclerc et al. 2007), did not have similar effects (Ehlermann et al. 2006). Another interpretation is that AGE pre-treatment provided a second signal necessary for S100A8/A9-mediated upregulation of pro-inflammatory genes in EC.

Alternatively, RAGE may act as a composite receptor. That is, RAGE may need to physically associate (perhaps directly bound or in lipid rafts) with another receptor/receptors to mediate certain functions. For example, RAGE and CD18/CD11b interact both in cis (on the same cell) and trans (between two cells) and interestingly HMGB1 requires both CD18/CD11b and RAGE to initiate NF κ B transcriptional activity in neutrophils (Orlova et al. 2007). β 2 integrins interact with other cell surface

receptors in cis, including Fc γ RIIA (CD32), urokinase-type plasminogen activator receptor (CD87), CD14 and PDGF β receptor (Petty et al. 2002), and are implicated as signaling partners for some of these (Petty et al. 1997; Todd and Petty 1997). Therefore, it is possible that CD18/CD11b and RAGE act as a composite receptor wherein RAGE binds the ligand and CD18/CD11b transduces the signal. Alternatively, RAGE and CD18/CD11b-initiated signals may both be required in close proximity, for example within lipid rafts. Such a composite receptor mechanism may explain why S100B requires the extracellular domain of RAGE, but not RAGE signaling, to stimulate NO production in BV-2 microglia (Adami et al. 2004) and similar mechanisms could mediate some of the responses of calgranulins.

Post-translational modifications of S100A8 and S100A9 by oxidants

S100A8 and S100A9 are exceptionally abundant in neutrophils and upregulated in activated macrophages, cell types that produce high levels of ROS. Myeloperoxidase (MPO) expressed in phagocyte granules is released following activation and generates the powerful oxidant, HOCl from H₂O₂ and Cl⁻ (Klebanoff 2005). Large amounts (>100 μ M) can be produced from 10⁶ activated neutrophils within 2 h (Test and Weiss 1986). Activated macrophages also express high levels of MPO at chronic inflammatory sites such as in atheroma (Nicholls and Hazen 2005; Lau and Baldus 2006). Murine and human S100A8 are exceptionally sensitive to oxidation of their single, conserved Cys residue (Harrison et al. 1999). Mild oxidation with Cu²⁺ or low levels of H₂O₂ generate disulfide-linked dimers and activated neutrophils rapidly convert it to this form (Harrison et al. 1999). Intra- (between Lys_{34/35} and Cys₄₁) and intermolecular (between Lys₆, Lys₇₆, Lys₈₃ Lys₈₇ and Cys₄₁) sulfinamide bonds are generated with as little as 1:1 M equivalents of hypochlorite (HOCl) in vitro (Raftery and Geczy 2002; McCormick et al. 2005) and from PMA-stimulated neutrophils (Raftery et al. 2001). Covalent sulfinamide bonds are formed between thiol groups in Cys residues and the ϵ -amine group of Lys residues and, unlike disulfide bonds, cannot be broken by normal cellular reduction systems or the commonly used chemical reductant dithiothreitol (DTT) (Raftery et al. 2001). Some methionine residues in both proteins are also oxidized to methionine sulfoxides (Harrison et al. 1999). These processes are more fully covered in a recent review (Lim et al. 2009).

S100A8 and S100A9 are abundant at inflammatory sites and their sensitivity to modification by ROS/RNS may be important in protecting oxidant-sensitive proteins when oxidants are generated following phagocyte activation. In the context of atherosclerosis, oxidation by HOCl

promotes aggregation of low-density lipoproteins (LDL), which are taken up more readily by macrophages than the unmodified form, facilitating formation of pro-atherogenic foam cells (Hazell and Stocker 1993). Lys and Met residues in apolipoprotein B are the main targets in LDL and these mediate aggregation, whereas lipid peroxidation is a quantitatively minor product (Hazell et al. 1994). An unsuspected oxidation pathway involving oxidative cross-linking of Cys and Lys residues was proposed to contribute to the pathogenesis of atherosclerosis (Fu et al. 2002). Appreciable amounts of aggregated LDL are generated with molar ratios HOCl to LDL, of 300:1, some 300-fold more than required for oxidative cross-linking of S100A8, which is induced by equimolar concentrations of HOCl (Raftery and Geczy 2002; McCormick et al. 2005). This exquisite sensitivity indicates that S100A8, and to a lesser extent S100A9, may protect against excessive oxidation in inflammatory conditions where activated neutrophils/macrophages express high levels of MPO, a property that may protect the host against severe tissue damage. DTT-stable cross-linked complexes (dimeric S100A8₂, S100A9₂, S100A8/A9 and S100A8/A9₂) were found in extracts from human atherosclerotic lesions (McCormick et al. 2005), indicating that this could be the case.

Leukocyte migration

Transmigration to inflammatory sites involves the process of rolling of leukocytes on activated endothelial cell surfaces mediated by binding of endothelial selectins to glycoprotein and glycolipid ligands on leukocytes (Sperandio 2006). Leukocyte integrins bind vascular cell adhesion molecules, thereby mediating tight adherence and immobilizing leukocytes, and allow subsequent transmigration

through the endothelial layer toward chemotactic signals (Petri and Bixel 2006; Petri et al. 2008).

mS100A8 was the first S100 family member described as having potent chemotactic activity for murine neutrophils and monocytes and was initially named chemotactic protein 10 kDa (CP10) to reflect its function and molecular mass (Lackmann et al. 1992). Since then, several S100 proteins have been implicated in modulating migration of various cell types (Table 4). One feature is their potency; mS100A8 has maximal chemotactic activity for neutrophils and monocytes at 10^{-11} to 10^{-13} M and, in contrast to other chemoattractants (e.g. fMLP, IL-8), does not cause a Ca^{2+} flux or induce changes in integrin or L-selectin expression (Cornish et al. 1996). It causes profound shape changes in monocytes and may mediate initial events in diapedesis that are independent of changes in integrin expression (Cornish et al. 1996). These features are similar to those described for TGF- β , which is also considered to have anti-inflammatory functions (see (Cornish et al. 1996)).

Injection of mS100A8 into murine foot pads stimulates leukocyte recruitment (Lackmann et al. 1993) with kinetics typical of delayed-type hypersensitivity reactions elicited by injection of antigen into a sensitized animal, and with early recruitment of neutrophils followed by monocytes over 24 h (Devery et al. 1994; Lau et al. 1995). The mechanism through which mS100A8 exerts this function is unclear, but is possibly via a pertussis toxin-sensitive, G-protein-coupled receptor (Cornish et al. 1996).

Similar to mS100A8, S100A12 may regulate leukocyte trafficking by acting as a chemoattractant and influencing expression of adhesion molecules. It upregulates CD11b integrin expression and L-selectin shedding on neutrophils, which enhances their adhesion to ECs (Rouleau et al.

Table 4 Chemotactic activities of S100 proteins

S100	Responsive cells	Optimal concentration (M)	Reference
Bovine S100A2	Guinea pig eosinophils	10^{-10}	(Komada et al. 1996)
Human S100A7	Human CD4 ⁺ T cells, monocytes and neutrophils	10^{-11} to 10^{-8}	(Jinquan et al. 1996; Wolf et al. 2008)
Human S100A15	Human monocytes and neutrophils	10^{-9}	(Wolf et al. 2008)
Murine S100A8	Murine neutrophils and monocytes	10^{-13} to 10^{-11}	(Lackmann et al. 1992; Geczy 1996; Ryckman et al. 2003a, b)
Murine S100A8 ₄₂₋₅₅	Murine neutrophils and monocytes	10^{-11} to 10^{-10}	(Lackmann et al. 1993; Geczy 1996)
Human S100A8, -A9 and -A8/A9	Neutrophils	10^{-12} to 10^{-9}	(Ryckman et al. 2003a, b)
Human S100A8 ₂₁₋₄₅	Human periodontal ligament cells	10^{-12}	(Nishimura et al. 1999)
Human S100A12	Human monocytes, neutrophils, mast cells and THP-1 cells	10^{-12} to 10^{-8}	(Yang et al. 2001; Rouleau et al. 2003; Yan et al. 2008)
Human S100A12 ₃₈₋₅₃	Human monocytes and mast cells, murine mast cells	10^{-12} to 10^{-9}	(Yan et al. 2008)
Bovine S100B	Murine SMCs	Not determined	(Sakaguchi et al. 2003)

2003). S100A12 may also induce ICAM-1 and VCAM-1 expression on ECs, thereby ensuring the tight adhesion required for extravasation (Hofmann et al. 1999). S100A12 is chemotactic in vitro for monocytes (Miranda et al. 2001) and mast cells (Yan et al. 2008) and has weak activity for neutrophils, but not lymphocytes (Yang et al. 2001), with optimal activities of about 10^{-9} to 10^{-12} M. Because of the functional similarities in leukocyte trafficking, we suggest that S100A12 may represent the human homolog of mS100A8 (Ravasi et al. 2004).

The chemotactic properties of human S100A8 and S100A9 are less clear. Ryckman et al. demonstrated their chemotactic activity for human neutrophils (Ryckman et al. 2003a, b), although other studies (Newton and Hogg 1998) found no activity. Ryckman et al. (2003a, b) suggest that because the human proteins were active in the picomolar range, and inactivated by HOCl treatment, differences could be due to oxidation. At the concentration of HOCl used (100 μ M), S100A8 forms intermolecular sulfinamide-linked complexes (Harrison et al. 1999). We found no activity of disulfide-bonded mS100A8 dimers (Harrison et al. 1999), or of cross-linked complexes, although very mild oxidation with HOCl stabilized its function (Raftery et al. 2001). The hinge region of mS100A8 is responsible for its chemotactic activity (see below). The structure of human S100A8 is shown in Fig. 1 and the Cys and Lys residues highlighted are conserved in mS100A8. Figure 1 shows that Cys₄₁ lies adjacent to the hinge region (colored red) and on opposite sides of the dimer. Disulfide bond formation may sterically hinder access of the hinge to its receptor, whereas the Lys_{34,35} are in closer proximity to Cys₄₁ and intrachain sulfinamide bond formation may not disrupt access to the hinge and would stabilize the molecule.

Others report a fugetactic (cell repulsion) activity of S100A8 and S100A9 for human neutrophils (Sroussi et al. 2006, 2007), which in S100A9 was inhibited by oxidation of Met₆₃ and Met₈₃ (Sroussi et al. 2007). Thus, it is clear that oxidation can alter S100A8 and S100A9 functions and may regulate their role in leukocyte recruitment. Moreover, these studies stress the requirements for appropriate storage and testing of these proteins for potential oxidative modifications before functional assays are performed. In contrast to S100A8 and S100A9, the lack of Cys and Met residues in S100A12 would make it more resistant to modification by oxidation and allow it to function in oxidative conditions typical of inflammatory sites.

S100A8 and S100A9 may also contribute to leukocyte trafficking by influencing integrin expression and affinity. One study showed that S100A9 induces the high-affinity epitope of the β_2 integrin CD11b on neutrophils, without causing shedding of L-selectin or enhanced CD11b expression; this activity was negatively regulated by formation of the heterodimer with S1008 (Newton and Hogg

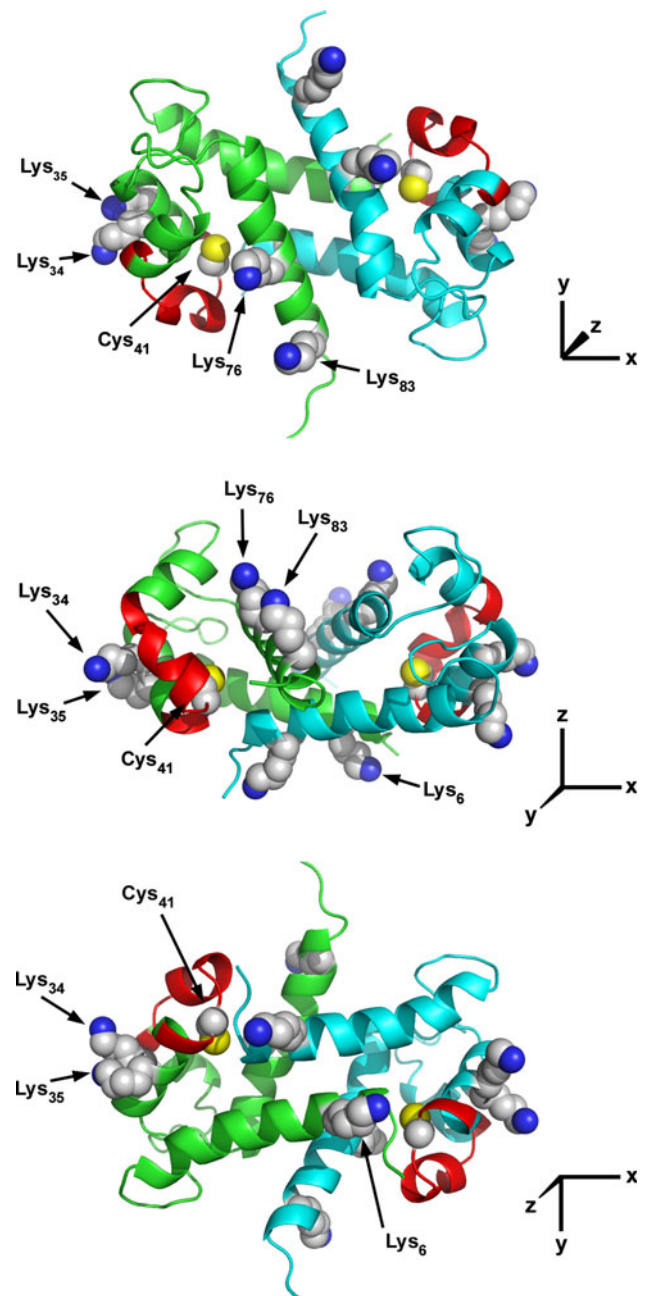


Fig. 1 Ribbon diagram structure of the S100A8 homodimer. The S100A8 homodimer is shown as a ribbon diagram with each subunit colored *green* or *blue* and the hinge regions colored *red*. Side chains forming inter- and intramolecular sulfinamide bonds are shown on both subunits, but labeled on only one. Images were produced with PyMOL and are based on the crystal structure of human S100A8 (PDB accession # 1MR8)

1998). However, another report (Ryckman et al. 2003a, b) showed that S100A8, S100A9 and S100A8/A9 cause L-selectin (CD62L) shedding and enhance CD11b expression on monocytes and neutrophils in addition to induction of its high-affinity epitope. Because of its ability to enhance CD11b affinity, S100A9 facilitates adhesion of neutrophils

to fibrinogen (Ryckman et al. 2003a, b) and fibronectin (Anceriz et al. 2007), and of monocytes to EC (Eue et al. 2000a, b) in vitro. Taken together, these data suggest that S100A9 modulates leukocyte adhesion to activated endothelium at sites of inflammation. However, this may be regulated by formation of the S100A8/S100A9 complex and, because the complex has been seen on endothelium in areas of contact between adherent leukocytes in vivo (Robinson et al. 2002), the significance of these observations is unclear.

S100A9^{-/-} neutrophils have reduced capacity to migrate through an EC monolayer in response to IL-8 or leukotriene B₄, and in contrast to wild-type cells, surface expression of CD11b was not increased by IL-8 in vitro (Manitz et al. 2003). These results, together with the reduced Ca²⁺ transients observed after stimulation of S100A9^{-/-} neutrophils with several chemokines (McNeill et al. 2007), indicate a role for S100A9 in response to chemotactic signals. It is proposed that S100A8/A9 regulates polymerization of the microtubule network in response to elevated [Ca²⁺]_i, an event antagonized by p38 MAP kinase-dependent phosphorylation of S100A9 (Vogl et al. 2004). S100A8 is the active subunit and phosphorylation of S100A9 may modulate this, and deficiencies in S100A9^{-/-} leukocyte migration may be due to dysregulation of cytoskeletal dynamics.

Some in vivo models also support a role for S100A8/A9 in leukocyte migration. In murine air pouch models, blockade of S100A8 and S100A9 by specific antibodies significantly suppressed neutrophil migration in response to LPS or monosodium urate crystals (Ryckman et al. 2003a, b; Vandal et al. 2003). Human S100A8/A9 injected into the murine air pouch recruits neutrophils, with maximal numbers 12-h post-injection, an effect independent of TLR-4 (Ryckman et al. 2003a, b). In S100A9-null mice, no differences in the accumulation or time course of leukocytes recruited in thioglycollate-induced peritonitis to the air pouch following LPS administration (Hobbs et al. 2003) were seen, whereas neutrophil infiltration into skin wounds (Vogl et al. 2004) and into the pancreas in a model of caerulein-induced pancreatitis (Schnekenburger et al. 2008) was reduced. Several responses of S100A9^{-/-} neutrophils appear to be stimulus dependent; for example, the oxidative burst generated by S100A9^{-/-} neutrophils stimulated with phorbol esters is similar to wild-type cells (Hobbs et al. 2003) whereas responses to ATP or platelet-activating factor are reduced (Nacken et al. 2005). It is proposed that S100A9 may regulate a specific signaling pathway, possibly diacylglycerol signaling (McNeill et al. 2007), and may explain why leukocyte influx is disrupted in some in vivo models while appearing normal in others.

The C-terminal region of S100A9 is identical to neutrophil immobilizing factor (NIF) (Freemont et al. 1989)

that inhibits neutrophil migration and chemotaxis in vitro (Goetzl and Austen 1972; Goetzl et al. 1973). The question of whether NIF is proteolytically cleaved from S100A9 was posed over 15 years ago (Freemont et al. 1989). Unpublished results from our laboratory indicate that NIF is efficiently produced when S100A9 is digested with α -chymotrypsin, and to a lesser extent with cathepsin G (L. Miranda, M. Raftery & C.Geczy, unpublished data). This suggests that granular contents comprising numerous proteases from activated neutrophils (Raptis and Pham 2005) could generate NIF and this may represent one means of limiting neutrophil influx into inflammatory lesions. A C-terminal domain peptide within this region of S100A9 (H92-H102) is implicated in regulation of neuropathic pain (Paccola et al. 2008) and inhibits spreading and mannose-dependent phagocytic activity of adherent peritoneal macrophages (Pagano et al. 2005), indicating a regulatory role for this peptide in macrophage function.

Mast cell regulation

S100A12 was recently identified as a novel mast cell activator (Yang et al. 2007). In addition to a potential role in allergic inflammation, this function may have important ramifications in infection and in chronic inflammation such as in RA, IBD and atherosclerosis, where S100A12 is expressed and mast cell activation is implicated in pathogenesis, although endogenous activators in these circumstances are unclear [reviewed in (Galli et al. 2005; Bischoff 2007)]. S100A12 provoked degranulation of murine and human mucosal and tissue mast cells and significantly potentiated IgE-mediated degranulation (Yang et al. 2007). This is an important observation considering the functional heterogeneity of mast cells from different species (murine and human) and tissues (connective tissue and mucosal types) (Gurish and Austen 2001; Bischoff 2007). In marked contrast to its inability to induce pro-inflammatory cytokines in human monocytes/macrophages, S100A12 induced IL-6, IL-8, MCP-1 and MIP-1 β from human cord blood-derived mast cells (Yang et al. 2007). Injection of S100A12 caused edema and leukocyte recruitment in mice; responses were not apparent in mast cell-deficient mice or wild-type mice injected with the mast cell stabilizer sodium cromoglycate. Leukocyte infiltration in mast cell-deficient mice was restored when these mice were reconstituted with bone marrow-derived mast cells from wild-type mice (Yang et al. 2007). Our recent studies show that S100A12 is also a mast cell chemoattractant in vitro and can sequester these cells when injected intraperitoneally (Yan et al. 2008). We propose that mast cells are a key target of S100A12 and their localized sequestration and

activation would generate chemokines, histamine and cytokines (e.g., TNF α) that promote activation of the vasculature, which is necessary for more sustained leukocyte recruitment, particularly of monocytes.

Importantly, potentiation of IgE-mediated mast cell activation indicates a primary role in allergic inflammation. S100A12 is expressed by macrophages in the vicinity of tryptase-positive mast cells and in eosinophils in the airways of lung tissue of patients with asthma, but not in the normal lung. It is elevated in the sputum of patients with eosinophil asthma (Yang et al. 2007) indicating its release.

S100A8 is also implicated in the regulation of mast cell activation by acting as an NO shuttle (Lim et al. 2008). By covalent coupling to cysteine residues in particular proteins, NO mediates numerous processes including vascular homeostasis, smooth muscle function, neuronal development, anti-microbial defense and regulation of transcription and apoptosis via S-nitrosylation (the covalent coupling of NO to Cys residues) (Jaffrey et al. 2001). Relatively few proteins are targets for S-nitrosylation and specificity of this modification may be conferred by a local hydrophobic environment and acidic and basic residues in proximity to the Cys, which are important for reducing the thiol pKa (Perez-Mato et al. 1999; Hess et al. 2005). We showed that S100A8 is readily S-nitrosylated on Cys₄₁ (Lim et al. 2008), which is conserved in human and murine S100A8, flanked by Glu₄₀ and Arg₄₆ residues, and lies within the hydrophobic pocket formed by the hinge and C-terminal regions. S100A8 is preferentially nitrosylated in the S100A8/A9 complex; S100A8-SNO was detected in normal human neutrophils and the levels increased in neutrophils treated with NO donors (Lim et al. 2008). The S100A8-SNO bond is relatively stable and able to trans-nitrosylate hemoglobin indicating that it may act as an NO shuttle. Importantly, S100A8-SNO inhibited mast cell degranulation stimulated by compound 48/80 in vitro, and leukocyte adhesion and transmigration triggered by mast cell activation in the mesenteric circulation of the rat (Lim et al. 2008). Since S-nitrosylated S100A8 can act as an NO donor, it may regulate other NO-dependent functions by trans-nitrosylating other NO targets. These properties indicate important functions of blood reflow in the microcirculation in resolving inflammatory lesions or in neovessels where S100A8 and S100A9 are expressed (McCormick et al. 2005).

Structure–function studies

S100s contain a low-affinity S100-specific N-terminal and a high affinity canonical C-terminal EF-hand Ca²⁺-binding domain (Gribenko and Makhatadze 1998; Heizmann and Cox 1998; Zimmer et al. 2003). The two EF hands are

encoded by separate exons with an intron invariably splitting the coding sequence at the ‘hinge’ region between the two Ca²⁺-binding motifs. Thus, an S100 can bind two Ca²⁺ ions, each with different affinities. This may be functionally important in second messenger roles, since the lower Ca²⁺ affinity of the N-terminal EF hand indicates that it may only be occupied when intracellular Ca²⁺ levels are elevated. Moreover, the Ca²⁺ concentration in the extracellular space is sufficiently high to ensure that S100s remain in the Ca²⁺-loaded state (Heizmann and Cox 1998), suggesting this as the functionally relevant form in those S100 proteins with extracellular roles.

Under physiological conditions, S100s exist as non-covalent antiparallel dimers held together by hydrophobic interactions, with hydrophobic residues within α -helices I and IV predominantly responsible for the dimerization interface. S100s typically form homodimers, although heterodimers, trimers and tetramers of S100A8/A9 (Hunter and Chazin 1998; Propper et al. 1999; Vogl et al. 2006) and hexamers of S100A12 (Xie et al. 2007; Moroz et al. 2009) are reported. The calcium-binding domains within the EF hands contribute to intermolecular stability of the S100A8/A9 complexes (Vogl et al. 1999a, b). Heterodimerization is independent of the extended C-terminal domain of S100A9 (Champaiboon et al. 2009). Recent studies mutating glutamic acid residues within the Ca²⁺-binding domains of S100A9 ablated resistance to bacterial invasion of a keratinocyte cell line over-expressing the S100A8/S100A9 complex. This directly confirmed the functional importance of EF-hand domains in resistance, by controlling the number of bound bacteria that could be internalized by the cells. A loss of the Ca²⁺-induced negative molecular surface to a positive face in these mutants is proposed to alter its binding properties to cytoskeletal components, such as negatively charged tubulin that is involved in bacterial invasion (Champaiboon et al. 2009).

Some functions of S100A8 and S100A9 are negatively regulated by, and others dependent on, heterocomplex formation (see Table 2), indicating that homo-/heterodimer formation may be an important regulatory mechanism. Since dimerization generates two binding sites on opposite sides of each monomer, S100 proteins could bridge target proteins. In the case of heterodimers, two different target proteins may be brought together in close proximity, a property that may facilitate the function of S100 proteins (Donato 2003). For example, this could conceivably be involved in the Ca²⁺-dependent assembly of enzyme complexes, such as the role proposed for S100A8/A9 in the assembly of a functional nicotinamide adenine dinucleotide phosphate (NADPH) oxidase complex in phagocytes (Kerkhoff et al. 2005).

Amino acid sequences within the EF-hand regions of S100 proteins are generally highly conserved, whereas

those within the hinge and C-terminal regions are the most divergent. Crystal and nuclear magnetic resonance (NMR) structures indicate that these regions form a hydrophobic pocket in the calcium-bound state, mediated by contraction of the C-terminal, high-affinity EF hand around the Ca^{2+} ion, which moves helix III into a position more perpendicular to helix IV and opens the hinge region. In contrast, the N-terminal EF hand undergoes very little Ca^{2+} -dependent conformational change (Smith and Shaw 1998; Heizmann et al. 2002; Zimmer et al. 2003). Structures of S100B, S100A10 and S100A11 bound to target peptides indicate that hydrophobic residues exposed in the hinge and C-terminus as a result of Ca^{2+} -ligation may mediate binding to target proteins (McClintock and Shaw 2003; Zimmer et al. 2003). The sequence conservation within the EF-hand regions indicates that this mechanism may be common to all S100s, whereas the sequence differences within the hinge and C-terminal regions could allow a diversity of target protein interactions that may determine the functions attributed to different family members (Kligman and Hilt 1988; Kube et al. 1992; Lackmann et al. 1993; Schafer and Heizmann 1996; Seemann et al. 1996; Heizmann et al. 2002; Ravasi et al. 2004; Yan et al. 2008).

Our earlier studies demonstrated that the chemotactic activity of mS100A8 for monocytes and neutrophils *in vitro* and *in vivo* was not found with human S100A8, which has only 57% amino acid identity, posing the question of whether the two proteins were orthologs (Lackmann et al. 1993) and whether there was equivalent chemotactic peptide in humans. We subsequently isolated a functional S100A8 homolog from human neutrophils, S100A12, which had potent monocyte chemotactic activity (Miranda et al. 2001; Yang et al. 2001) but failed to identify S100A12 in rodent genomes (Ravasi et al. 2004). The functional and sequence divergence suggested complex evolution of the S100 family in mammals. A synthetic peptide with sequence identical to the hinge region of mS100A8 (amino acids 42–55, mS100A8_{42–55}) mimicked the chemotactic activity of the full-length protein and was the first demonstration of an active hinge domain in the S100 protein family (Lackmann et al. 1993). However, synthetic mS100A8_{42–55} elicited only transient neutrophil recruitment *in vivo*, indicating other essential structural domains.

In contrast to the positive activity of the murine hinge domain, an equivalent peptide segment corresponding to the hinge sequence of hS100A8 had no chemotactic activity (Lackmann et al. 1993). Comparisons of the amino acid sequences of m/h S100A8 and S100A12 are given in Fig. 1. Using structural models of the three proteins, we found that the S100A12 and hS100A8 templates produced successful models for mS100A8, although the total amino acid identity between hS100A8 and mS100A8 was higher

than between S100A12 and mS100A8 (32%). Because of higher overall identity and the high surface hydrophobicity evident in the S100A12 model, the hS100A8 template was considered to be more accurate. However, the hinge regions of human and murine S100A8 have low homology, particularly in the C-terminal half. In the human S100A8 hinge, Gly₄₉ would allow a level of flexibility not afforded by Asp₄₉ in the mouse protein and consequently the murine hinge is unlikely to form the same kinked conformation described for hS100A8 (Ishikawa et al. 2000). Electrostatic potential maps indicated that the hinge regions of mS100A8 and S100A12 were structurally more similar. In contrast to the hinge regions of S100A12 and mS100A8, which are electro-negative, this region in hS100A8 is positively charged.

We next prepared a synthetic peptide based on the S100A12 hinge sequence, S100A12_{38–53}, and showed that, like the full-length protein, this was chemotactic for human THP-1 monocytoid cells and for human monocytes (Ravasi et al. 2004). Because S100A12 consistently provokes THP-1 cell migration in a biphasic manner with two prominent peaks at 10^{-9} and 10^{-12} M, two receptors of different affinities are implicated. S100A12_{38–53} was also chemotactic at these concentrations, although somewhat less potent. Using alanine substitutions, we showed that Lys₃₈, Leu₄₀, Asn₄₆, Ile₄₇, Asp₄₉, Lys₅₀ and Ile₅₃ were essential for monocyte chemotactic activity at 10^{-9} M, whereas Asn₄₂ and Ile₄₄ were important for activity at 10^{-12} M (Yan et al. 2008). In hydrophobic environments, the S100A12_{38–53} peptide adopts an α -helical conformation with the active hydrophobic residues on one side of the helix, suggesting that near the cell membrane (a hydrophobic environment), these residues may align to interact with a receptor (Yan et al. 2008). The hinge region of mS100A8 also has an α -helical structure in hydrophobic environments (Lazoura et al. 2000). As seen in Fig. 2, the sequence of S100A12's hinge region is more similar to the hinge region of mS100A8 than the human protein, and Asn₄₆ and Ile₄₇ are conserved at similar positions in mS100A8 but not human S100A8, which may explain why the mS100A8 hinge region is chemotactically active, whereas the human S100A8 hinge is not (Yan et al. 2008).

The hinge region of S100A12 also mimicks the stimulation of mast cell degranulation and chemotactic activities of full-length S100A12. Hydrophobic residues (Ile₄₄, Ile₄₇ and Ile₅₃) within this region are essential for monocyte chemotactic activity, mast cell activation and induction of edema *in vivo*. Mutation of Lys₃₈ or Lys₄₈ to Ala also reduced, but did not abolish edema caused by the S100A12 hinge region, indicating that these residues may contribute to the activity (Yan et al. 2008).

The redox-sensitive Cys residue in mS100A8 resides immediately before the hinge domain, but mutation to Ala

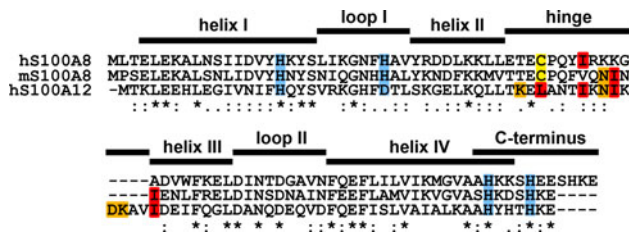


Fig. 2 Amino acid alignment of murine and human S100A8 and human S100A12. Positions of the helices, loops, hinge and C-terminal domains are indicated above the alignment. Residues involved in Zn^{2+} -binding are highlighted in blue. Hydrophilic (orange) and hydrophobic (red) residues implicated in the chemotactic activity of S100A12 (and where they are conserved in human and murine S100A8) are also highlighted. Asterisk indicates a conserved residue, colon conservative substitution, dot weakly conservative substitution. Sequences used were: human S100A8 (P05109), murine S100A8 (P27005) and S100A12 (P80511)

did not alter its chemotactic activity *in vitro* or *in vivo* (Harrison et al. 1999). In keeping with its non-essential role in chemotaxis, S100A12 has no Cys residues (Fig. 2). In this respect, the chemotactic activity of S100A12, in contrast to mS100A8 (in which disulfide bond formation causes loss of activity), would be resistant to covalent modification by oxidants. S100A12 also has no Met residues and, because a number of classical chemoattractants are susceptible to oxidative inactivation (Harrison et al. 1999), S100A12 may be important in propagating monocyte migration into chronic inflammatory lesions such as in RA (Yang et al. 2001) and atherosclerosis (Goyette et al. 2009) where ROS are continuously generated.

The S100 protein family is a good example that amplification occurs by gene duplication followed by divergence from the original duplicated gene. We propose that a duplication and divergence of S100A8 in humans, compared to rodents, has permitted the separation of two functions. In mice, S100A8 is abundant and acts as an anti-oxidant, whereas it is chemotactic within the picomolar range, but its chemotactic activity can be compromised by sensitivity to ROS (Harrison et al. 1999). Interestingly, rabbit S100A12 has a single Cys residue at position 30, whereas porcine and bovine S100A12 have no Cys residues. In human, and in addition to the numerous functions ascribed to the S100A8/A9 complex, S100A8 may function as an anti-oxidant, whereas S100A12 may chiefly act as a monocyte and mast cell chemoattractant and mast cell activator (Yang et al. 2001, 2007, 2008).

The absence of the S100A12 gene in rodents makes some studies using these in model systems difficult to interpret, particularly with neutralizing antibodies (Hofmann et al. 1999), and to relate particular functions exhibited by the human proteins to the calgranulins in these species. Moreover, the differences in expression patterns of murine and human S100A9 in activated monocytes/macrophages

(Hessian et al. 1993; Santhanagopalan et al. 1995; Hu et al. 1996; Xu and Geczy 2000; Xu et al. 2001; Hsu et al. 2005) indicate important functional differences and question the roles of the S100A8/A9 complex in chronic inflammation in mice. Moreover, there are other important considerations. For example, we (McCormick et al. 2005) and others (Eue et al. 2000a, b) found no S100A8 and little S100A9 in murine models of atheroma. Interestingly, and in keeping with a protective oxidant-scavenging role, activated murine macrophages produce little MPO, making the lesion less prone to HOCl oxidation. Thus, studies of these proteins in chronic inflammation in mice should be interpreted with caution.

Roles of zinc binding

Divalent cations such as Zn^{2+} and Cu^{2+} may regulate extracellular S100 functions, particularly because Zn^{2+} -binding causes structural changes that could influence ligand/receptor binding (Vogl et al. 2006; Moroz et al. 2009). The putative Zn^{2+} -binding sites are structurally conserved in the calgranulins (see Fig. 2). Ca^{2+} and Zn^{2+} , alone or together, induce heterotetramer formation of S100A8/A9 (Vogl et al. 2006) and oligomerization of S100A12, although in this case Zn^{2+} is more effective than Ca^{2+} (Goyette et al. 2009; Moroz et al. 2009). Evidence from chemical cross-linking, NMR, hydrophobic probe and intrinsic Tyr fluorescence experiments suggests that the structures of Zn^{2+} -, Ca^{2+} - and Ca^{2+}/Zn^{2+} -bound S100A12 complexes are subtly different (Goyette et al. 2009; Moroz et al. 2009). Zn^{2+} is required for S100A8/A9 to bind to a human microvascular EC line (Robinson et al. 2002) and more S100A12 is bound to the surface of human gastric carcinoma MKN74 cells in the presence of Zn^{2+} than in its absence (Moroz et al. 2009).

If the Zn^{2+} -bound forms of the calgranulins indeed have different functions, this may explain some of the contradictory reports in literature, particularly as different culture media have different levels of Zn^{2+} (Moroz et al. 2009), and this aspect warrants clarification. Furthermore, the physiological relevance of Zn^{2+} -binding has been unclear. Our demonstration of S100A12- Zn^{2+} complexes in human atheroma using an antibody that only recognizes Zn^{2+} -dependent S100A12 complexes is the first demonstration of Zn^{2+} -dependent calgranulin oligomers *in vivo* (Goyette et al. 2009). Since S100A12 inhibited matrix metalloproteinases (MMPs) by chelating Zn^{2+} , these complexes may form as a consequence of inhibiting the large amount of MMPs, particularly MMP-9 produced in human atheroma (Goyette et al. 2009). S100A8 and S100A9 can also suppress MMP activity in a similar manner (Isaksen and Fagerhol 2001), implicating these processes in the protection of vulnerable plaque from rupture.

Zn^{2+} is implicated in the apoptotic activity of S100A8/A9 (Yui et al. 1995, 1997, 2002), suggesting structural changes

that regulate function, although additional mechanisms probably involving receptor-mediated activation and generation of oxidants are also reported (Ghavami et al. 2004, 2008a, b). S100A8/A9 (known as calprotectin) also has broad-spectrum anti-microbial activity against numerous micro-organisms, particularly fungi, and can promote resistance to bacterial invasion (Steinbakk et al. 1990; Murthy et al. 1993; Sohnle et al. 2000; Champaiboon et al. 2009). Anti-microbial activity was proposed to be mediated by the ability of the complex to bind and control the levels of free Zn^{2+} , an essential microbial nutrient (Steinbakk et al. 1990; Sohnle et al. 2000; Lusitani et al. 2003), although this is still contentious, particularly because S100A8/A9 may precipitate when Zn^{2+} is in excess (Viemann et al. 2007). The crystal structure of the S100A8/S100A9 heterotetramer indicates burial of more hydrophobic residues, particularly in S100A9, leading to a dimer of heterodimers with major contributions from the N-terminal EF hands and the C-termini. This allows the emergence of two putative Zn^{2+} -binding sites at the subunit interface (Korndorfer et al. 2007). The C-terminal region of S100A9 contains three consecutive His residues and a His₉₁-x-x-His₉₅ motif that may mediate Zn^{2+} -binding. An essential role for S100A9 is implicated and studies using ⁶⁵Zn indicate that Zn^{2+} -depletion is an unlikely mechanism and favor the proposal that Zn^{2+} -dependent conformational changes may mediate activity (Murthy et al. 1993). In keeping with alternate/additional mechanisms, the anti-fungal activity of S100A8/A9 is abolished when Cys₄₂ of S100A8 or Met₆₃ or Met₈₃, but not Met₈₁, of S100A9 are mutated (Sroussi et al. 2009). Although the precise mechanisms need clarification, oxidation of these residues may contribute to their anti-microbial function.

Interestingly, S100A7 and the C-terminal domain of S100A12 have anti-microbial activity against Gram-negative bacteria (Glaser et al. 2005). These proteins do not have extended C-terminal domains containing three His residues, although both contain the His-x-x-x-His motifs. The anti-microbial activity of the C-terminal-derived S100A12 peptide is enhanced by Zn^{2+} (Cole et al. 2001), indicating that structural changes mediated by Zn^{2+} binding may modulate this activity. In contrast, Zn^{2+} does not mediate the antibacterial activity of S100A7 as it remains active in the presence of Zn^{2+} , and deletion of the C-terminus had little effect on activity (Lee and Eckert 2007).

Concluding remarks

The calgranulins are highly expressed in acute and chronic inflammatory conditions, although their exact contribution to the inflammatory process remains elusive. Although these proteins are commonly labeled as ‘pro-inflammatory’,

a comprehensive assessment of literature suggests that the calgranulins may play pleiotropic roles and this review attempts to highlight some of their often overlooked, protective functions. The mechanisms regulating the relative pro- or anti-inflammatory/protective roles of the calgranulins warrant more research. We propose that oxidative modifications, proteolytic cleavage to release active peptides, Zn^{2+} -binding and complex formation may be important factors in functional diversity. Receptor-mediated functions may be governed by glycosylation of the receptor and requirements for co-receptors and/or co-stimuli. A deeper understanding of the consequences of these modifications to calgranulin function and the receptors mediating these effects may explain some of the seemingly incongruent functions proposed for these proteins.

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