

# Biotechnology of Anoxygenic Phototrophic Bacteria

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**Abstract** Anoxygenic phototrophic bacteria are a diverse collection of organisms that are defined by their ability to grow using energy from light without evolving oxygen. The dominant groups are purple sulfur bacteria, purple nonsulfur bacteria, green sulfur bacteria, and green and red filamentous anoxygenic phototrophic bacteria. They represent several bacterial phyla but they all have bacteriochlorophylls and carotenoids and photochemical reaction centers which generate ATP and cellular reductants used for CO<sub>2</sub> fixation. They typically have an anaerobic lifestyle in the light, although some grow aerobically in the dark. Some of them oxidize inorganic sulfur compounds for light-dependent CO<sub>2</sub> fixation; this ability can be exploited for photobiological removal of hydrogen sulfide from wastewater and biogas. The anoxygenic phototrophic bacteria also perform bioremediation of recalcitrant dyes, pesticides, and heavy metals under anaerobic conditions. Finally, these organisms may be useful for overexpression of membrane proteins and photobiological production of H<sub>2</sub> and other valuable compounds.

**Keywords** Biogas, Bioremediation, Carotenoids, Green sulfur bacteria, Hydrogen sulfide, Membrane proteins, Photosynthetic bacteria, Purple bacteria

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## Abbreviations

BChl	Bacteriochlorophyll
$E_0'$	Standard reduction potential at pH 7 and 25 °C
EPS	Extracellular polymeric substances
FAP	Filamentous anoxygenic phototrophs
GSB	Green sulfur bacteria
PBR	Photobioreactor
PNSB	Purple nonsulfur bacteria
PSB	Purple sulfur bacteria

## 1 Introduction

Phototrophic organisms are defined by their ability to convert light energy into chemical energy in forms useful for growth and other metabolic processes [1–3]. Cyanobacteria and microalgae are the only phototrophic microorganisms that evolve  $O_2$  as a result of this process. This is because they have an enzyme complex capable of oxidizing water to oxygen ( $H_2O \rightarrow 2H^+ + \frac{1}{2}O_2 + 2e^-$ ) and donating the electrons to a type II photochemical reaction center. These organisms are therefore oxygenic (i.e., oxygen-evolving) phototrophs. Phototrophs that do not oxidize water may, for example, oxidize hydrogen sulfide ( $H_2S \rightarrow 2H^+ + S + 2e^-$ ) and donate the electrons to the photochemical reaction center in the organism. These organisms are therefore anoxygenic (i.e., not oxygen-evolving) phototrophs. In either case, these electron transfer reactions generate a transmembrane proton motive force used for ATP generation and other physiological purposes. The reductants generated by the reaction center are also used for biosynthetic purposes and cyclic electron transfer around the reaction centers to generate even more proton motive force.

The vast majority of anoxygenic phototrophic bacteria grow phototrophically only under anaerobic conditions [4]. Some may be capable of chemotrophic growth under aerobic conditions in the dark (typical of purple bacteria), but others are obligate anaerobic and are killed by  $O_2$  (typical of green sulfur bacteria; GSB). Anoxygenic phototrophic bacteria rarely form visibly large accumulations in natural environments because of their anaerobic lifestyle, but occasionally various

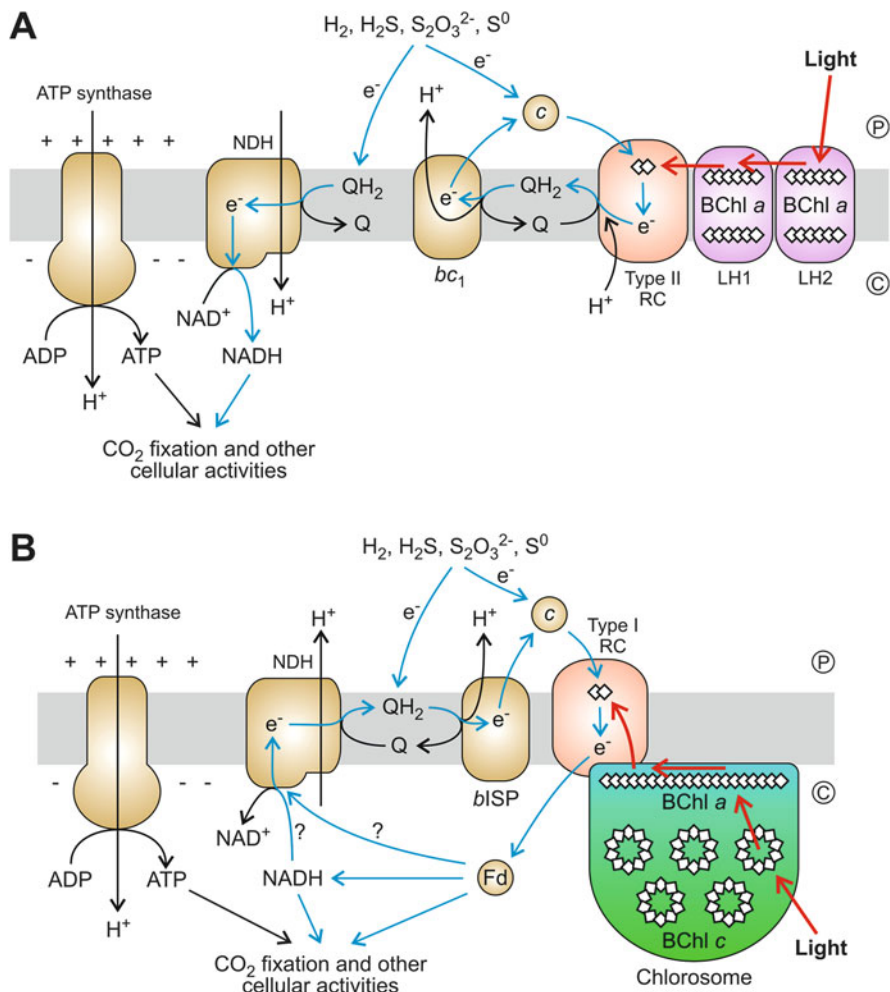
purple bacteria may bloom and color ponds, coastal areas, and wastewater reservoirs red, purple, or reddish-brown.

Anoxygenic phototrophic bacteria may not be as extensively used for biotechnological applications as chemotrophic microorganisms. However, given their diversity and the rising interest in biological solutions to societal challenges, more applications should be explored. For example, the extensive intracellular membrane systems in purple bacteria may have practical applications (Sect. 8). Another underexplored area is the potential for production of proteins and metabolites that are O<sub>2</sub> sensitive. In the following, a survey is made of biotechnological applications of the anoxygenic phototrophic bacteria. The bioremediating properties of oxygenic and anoxygenic photosynthetic bacteria were recently reviewed [5].

### ***1.1 Phototrophy and Photosynthesis***

There are only two principal sources of energy for living organisms: energy from light (phototrophy) or energy from chemical compounds (chemotrophy) [3]. Photosynthesis usually refers specifically to photoautotrophy, i.e., growth based on CO<sub>2</sub> fixation where the required energy is derived from light. Thus, all photosynthetic organisms are phototrophic but not all phototrophic organisms are photosynthetic. The principal cellular component that allows phototrophy is the photochemical reaction center [1]. This is a large enzyme complex that is always membrane-bound and has the ability to convert excitation energy obtained from light into chemical energy by creating a light-induced charge separation. The chemical energy is used for generation of strong reductants and ATP that are used in CO<sub>2</sub> fixation, biosynthetic pathways, and other cellular activities (Fig. 1). There are only two types of photochemical reaction centers: type I and type II. Type I (also called iron-sulfur type) has a relatively low redox potential and reduces soluble, cytoplasmic ferredoxins with  $E_0'$  values between  $-0.5$  and  $-0.6$  V. Type II (also called quinone-type) has a relatively high redox potential and reduces membrane-bound isoprenoid quinones with  $E_0'$  values between  $+0.1$  and  $-0.1$  V. The oxygenic phototrophs (cyanobacteria, algae, and plants) have both types of reaction centers (known as photosystems I and II) whereas the anoxygenic phototrophic bacteria have only either type I or type II. Light-harvesting pigment-protein antenna complexes are associated with the reaction centers to increase the amount of light energy harvested by the cells [1, 2]. The excitation energy in the antenna pigments is channeled to the photochemical reaction center in the order of picoseconds. There is a large diversity of evolutionary unrelated light-harvesting antenna complexes, but common to most of them is the presence of chlorophylls or bacteriochlorophylls and carotenoids. Most of the pigmentation in phototrophic bacteria (>99%) is found in these light-harvesting antenna complexes.

In principle, phototrophy can also be supported by alternative enzymes: the rhodopsin-like proteins [2, 3]. These membrane-bound enzymes absorb light energy and, as a result, transfer protons across the membrane, thereby conserving the light energy as a transmembrane electrochemical proton gradient. Rhodopsins



**Fig. 1** Simplified models of the photosynthetic machinery in (a) purple bacteria and (b) green sulfur bacteria (GSB) during photoautotrophic growth on inorganic sulfur compounds. Light is harvested by the peripheral antennae (LH1, LH2, or chlorosomes) and the excitation energy is transferred to the reaction center (RC). Light-driven cyclic electron transfer occurs when electrons are continuously transferred between the reaction center and the membrane-bound cytochrome  $bc_1/bISP$  in a light-dependent manner. This electron transfer generates a proton motive force, which fuels other physiological processes such as ATP synthesis. NADH is generated by reverse electron flow in purple bacteria under photoautotrophic growth conditions. If suitable organic substrates are available for photoheterotrophic growth in purple bacteria, NADH for cellular activities is instead obtained by oxidation of these organic substrates and the photochemical system may function exclusively in a cyclic mode to produce ATP. For details on oxidation of sulfur compounds ( $H_2S$ ,  $S_2O_3^{2-}$ , and  $S^0$ ) see Fig. 2. Pathways that carry electrons for cyclic electron transfer,  $CO_2$  fixation, and other cellular activities are shown with blue arrows. Transfer of light and excitation energy is shown with red arrows.  $bc_1$  cytochrome  $bc_1$  complex,  $BChl$  bacteriochlorophyll,  $bISP$  cytochrome  $b$  iron-sulfur protein,  $C$  (circled) cytoplasm,  $Fd$  ferredoxin,  $NDH$  NADH:quinone oxidoreductase,  $P$  (circled) periplasm,  $Q$  isoprenoid quinone (oxidized),  $RC$  photochemical reaction center. Modified from [1, 2]

do not transfer electrons and are not associated with large light-harvesting antennae as are photochemical reaction centers. Rhodopsins are well-known components of halophilic archaea where they support a phototrophic lifestyle under anaerobic conditions. Rhodopsin-like enzymes have recently been found in numerous very different marine microbes (archaea, bacteria, and eukaryotes) but the exact contribution to cell physiology is not always clear in these organisms. Phototrophy based on rhodopsins is not considered further here.

## 1.2 Types of Anoxygenic Phototrophic Bacteria

Purple bacteria belong to the phylum Proteobacteria and are characterized by having type II photochemical reaction centers and bacteriochlorophyll (BChl) *a* or BChl *b* as the major pigment [4, 6]. The light-harvesting (LH) antennae known as LH1 and LH2 are located in the cytoplasmic and intracytoplasmic membranes. As a group, the purple bacteria are very physiologically versatile and may grow photoautotrophically, photoheterotrophically, or chemotrophically in the dark by fermentation or by aerobic or anaerobic respiration. A subgroup of the purple bacteria, the purple sulfur bacteria (PSB), is characterized by a predominant photoautotrophic lifestyle where inorganic sulfur compounds such as sulfide serve as electron donors for photosynthesis. The other major subgroup of purple bacteria, known collectively as purple nonsulfur bacteria (PNSB), is characterized by a predominantly photoheterotrophic lifestyle where many organic compounds can be assimilated and these organisms are not as tolerant to sulfide as the PSB. PSB are Gammaproteobacteria whereas PNSB are Alphaproteobacteria and Betaproteobacteria.

GSB belong to the phylum Chlorobi and are characterized by having BChl *c*, *d*, or *e* organized into large light-harvesting organelles known as chlorosomes [4]. These organisms contain type I reaction centers and typically oxidize sulfur compounds for CO<sub>2</sub> fixation. Compared to the purple bacteria, the GSB have much more restricted growth requirements: they are strict anaerobic, obligate phototrophic, and obligate autotrophic.

Filamentous anoxygenic phototrophic (FAP) bacteria belong to the phylum Chloroflexi and are characterized by having BChl *a* organized into type II reaction centers and light-harvesting antennae similar to those found in purple bacteria [4]. There are two kinds of FAPs: the red FAPs have BChl *a* as the sole chlorophyll-like pigment, whereas the green FAPs, in addition to BChl *a*, also have BChl *c* or *d* organized in chlorosome structures similar to those found in GSB. The green FAPs are also known as green nonsulfur bacteria.

The only other groups of anoxygenic phototrophic bacteria known are the Heliobacteria, Chloracidobacteria, and the recently discovered Gemmatimonadiales [7]. There are no known archaea that contain photochemical reaction centers.

Cyanobacteria are oxygenic phototrophs [8]. However, a few cyanobacteria are capable of anoxygenic photosynthesis and an anaerobic phototrophic lifestyle

where the organisms consume hydrogen sulfide as electron donor for photosynthesis, very similar to the phototrophic GSB and PSB described above. Microalgae are also oxygenic phototrophs but are not known to grow phototrophically under anaerobic conditions. However, this does not exclude the possibility that some microalgae have light-independent lifestyles under anaerobic conditions or in the dark.

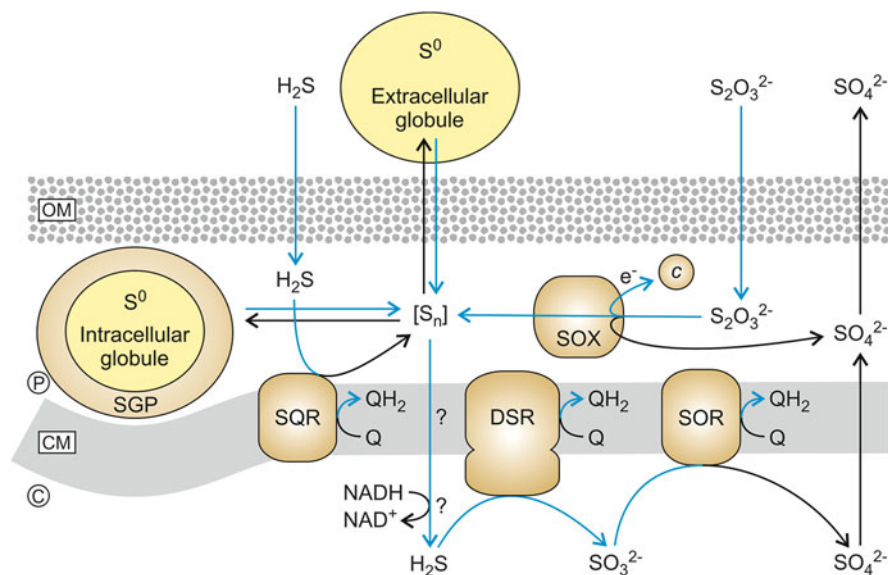
### ***1.3 Occurrence***

In general, anoxygenic phototrophic bacteria thrive where anaerobic conditions intersect with light [4]. In natural environments these conditions occur in lakes and sediments, often in narrow bands limited by the extent of light penetration. If sulfide is present, typically either PSB or GSB or both accumulate. Purple bacteria in general are physiologically versatile and thrive in many man-made polluted environments such as wastewater lagoons where light coincides with anaerobic or microaerobic conditions and high loads of organic compounds.

### ***1.4 Sulfur Metabolism and Sulfide Tolerance***

Reduced inorganic sulfur compounds are abundant in anaerobic environments because of the degradation of sulfur-containing organic materials and H<sub>2</sub>S production by sulfate-reducing bacteria. These compounds are oxidized by various chemotrophic bacteria (often collectively denoted “colorless sulfur bacteria”) under aerobic conditions or by phototrophic bacteria under anaerobic conditions. The metabolism of sulfur compound oxidation in phototrophic bacteria is complex and not fully elucidated (Fig. 2) [9]. Sulfide and thiosulfate are often incompletely oxidized to elemental sulfur if the sulfide and thiosulfate are supplied in excess. This elemental sulfur is deposited outside or inside the cells as sulfur globules with diameters ranging from very small up to 2 μm. Intracellular sulfur globules are found in PSB of the family *Chromatiaceae* and extracellular sulfur globules are found in GSB, PNSB, and most PSB of the family *Ectothiorhodospiraceae*. Upon depletion of sulfide and other electron sources, the sulfur globules are oxidized completely to sulfate.

Although sulfide is metabolized by most, if not all, phototrophic bacteria, the levels of tolerance vary significantly [9]. Sulfide concentrations above 15–30 mg/L H<sub>2</sub>S (0.5–1 mM) tend to inhibit the PNSB and favor the PSB and GSB. The highest sulfide concentrations tolerated by most PSB and GSB are about 100–150 mg/L H<sub>2</sub>S (about 3–5 mM), where the GSB typically are the most tolerant. In extreme cases, up to 375 mg/L H<sub>2</sub>S (11 mM) is tolerated.



**Fig. 2** Simplified model of the oxidative sulfur metabolism in phototrophic sulfur bacteria. Sulfur globules are shown in yellow. Intracellular sulfur globules in purple sulfur bacteria (PSB) are located in the periplasm and are associated with sulfur globule proteins (SGP). Pathways that carry electrons towards cellular electron carriers (isoprenoid quinones and cytochromes *c*) are shown in blue. *C* (circled) cytoplasm, *CM* (boxed) cytoplasmic membrane, *DSR* dissimilative sulfite reductase, *OM* (boxed) outer membrane, *P* (circled) periplasm, *Q* isoprenoid quinone (oxidized), *SGP* sulfur globule protein,  $[S_n]$  oligosulfide pool, *SOR* sulfite oxidoreductase (several types), *SOX* sulfur compound oxidizing system, *SQR* sulfide:quinone oxidoreductase. Modified from [9]

## 2 Removal of Hydrogen Sulfide from Wastewater Streams

Sulfide is a malodorous, corrosive, and toxic compound often present in liquid waste streams from domestic and industrial sources. The source is typically organically bound sulfur, which is liberated as sulfide or sulfate upon degradation, and inorganic sulfur (mostly sulfate), which is reduced to sulfide under anaerobic conditions by ubiquitous sulfate-reducing bacteria. Anoxygenic phototrophic bacteria offer an environmentally friendly biological approach to remove this sulfide [10]. Partial oxidation of H<sub>2</sub>S to elemental sulfur (S<sup>0</sup>) rather than complete oxidation to sulfate (SO<sub>4</sub><sup>2-</sup>) is desirable for a number of reasons. (1) Elemental sulfur can be physically removed from the waste stream by sedimentation of the sulfur granules. This alleviates downstream problems with sulfate (such as reduction to sulfide by sulfate-reducing bacteria). In addition, this biogenic elemental sulfur has commercial value as fertilizer and chemical feedstock. (2) Oxidation of sulfide to elemental sulfur generates less H<sup>+</sup> than oxidation of sulfide to sulfate. This minimizes corrosion and the requirement for pH control. (3) Conversion to elemental sulfur requires less light than conversion to sulfate; this maximizes the sulfide removal per light input. For these reasons, conditions in the photobioreactor

(PBR) have to be carefully controlled (e.g., flow rate, light intensity) to avoid complete sulfide oxidation and to maximize the output of elemental sulfur. Alternatively, naturally occurring or genetically engineered strains of phototrophic sulfur bacteria could be used which are deficient in sulfate formation and only oxidize sulfide to elemental sulfur, regardless of the sulfide load [11].

For sulfide removal from waste streams, GSB have certain advantages over other anoxygenic phototrophic bacteria [10]. (1) GSB deposit elemental sulfur extracellularly, which enables recovery of the sulfur by sedimentation (most PSB accumulate the elemental sulfur intracellularly). (2) GSB oxidize more sulfide per light input than purple bacteria, which is important for practical design and economic reasons. (3) GSB have higher tolerance for sulfide and higher affinity for sulfide uptake than purple bacteria. (4) Sulfide uptake in purple bacteria is diminished when organic nutrients are available. For these reasons, if a defined microbial culture is to be used for photobiological sulfide removal, a GSB culture may be beneficial. In one such study, a pilot-scale fixed-film continuous-flow PBR (21 mL) with the GSB *Chlorobium limicola* strain DSM 257 was designed and successfully optimized for sulfide removal [12]. At loading rates of 111–286 mg/L/h sulfide, about 100% of the influent sulfide was consumed and 92–95% recovered as elemental sulfur [12].

In another pilot-scale study in Brazil, up-flow anaerobic PBRs were tested with a continuous flow of domestic wastewater containing 1–6 mg/L sulfide [13]. These systems relied on naturally developing microbial communities under the influence of natural light. Sulfide removal efficiencies reached 90% and the effluent contained less than 0.5 mg/L sulfide, which is below the national sulfide discharge standard in Brazil of 1 mg/L. Most of the sulfide was oxidized to elemental sulfur ( $S^0$ ) although some sulfate ( $SO_4^{2-}$ ) was also formed. The sulfide was oxidized by a mixed microbial community which was shown by molecular techniques to contain GSB, PSB of the *Chromatiaceae* lineage, and green FAPs of the *Chloronema* lineage. Sulfide removal by chemotrophic denitrification was unlikely to occur because the influent and effluent wastewater contained very little nitrate and nitrite, and microbes capable of performing this process were not identified in the microbial community.

### 3 Removal of Hydrogen Sulfide from Gas Streams

Hydrogen sulfide is present in small amounts in biogas (typically 0.1–2%) and off-gases from wastewater treatment plants, for example. Besides being toxic and malodorous,  $H_2S$  in biogas causes corrosion and poisoning of the equipment using the biogas. Therefore, a number of chemical technologies are currently used to remove  $H_2S$  from biogas. A commercialized approach using chemotrophic sulfide-oxidizing bacteria in a fixed-film bioreactor under controlled oxygen conditions is also available [14]. Fixed-film or suspended-growth PBRs with anoxygenic phototrophic bacteria could be interesting alternatives for cost-effective  $H_2S$



removal because of their ability to operate for long periods of time without requiring a biomass separation step and their ability to operate under high and variable sulfide loadings [14]. Illumination and design of the PBR is a key concern for practical application because light is rapidly attenuated in dense cell suspensions [15]. Studies suggest that illumination using light-emitting diodes (LED) appears to be a very cost-effective approach if the LED emission wavelength is matched to the pigment absorption by the cells [16].

A variety of reactor designs have been investigated which allow the influent gas to exchange  $H_2S$  with an aqueous medium containing suspended or immobilized sulfide-oxidizing bacteria. In one study, a continuous-stirred PBR with *Chlorobium thiosulfatophilum* was used to remove sulfide from a synthetic gas stream containing 2.5%  $H_2S$  [17]. At its optimum operation, the 1.25-L PBR consumed 118 mg/h  $H_2S$  with negligible formation of sulfate corresponding to a conversion rate of sulfide to elemental sulfur per PBR volume of about 100 mg/h/L  $H_2S$ . Another study with a larger PBR working volume (11.9 L) used a flat-panel gas-lift PBR design with a PBR thickness of 10 cm and LED illumination [16]. In this system, the sulfide conversion to elemental sulfur was only about 20 mg/h/L  $H_2S$  but the energy expenditure for illumination per amount of sulfide removed was the lowest reported in the literature because of the use of LED for illumination.

## 4 Degradation of Recalcitrant Dyes and Pesticides

Azo dyes are the most important group of synthetic colorants and are generally recalcitrant to biodegradation because of their xenobiotic nature. Degradation of azo dyes in wastewater streams is most efficient under anaerobic conditions because many anaerobic bacteria apparently have a broad-range ability to reduce the azo bond [18]. Azo dyes are decolorized by many PNSB and this degradation is dependent on the enzyme azoreductase [19, 20]. Pure culture studies with different isolates of *Rhodospseudomonas palustris* have shown that this organism efficiently decolorizes various azo dyes in concentrations of around 1 g/L dye [20, 21]. In these studies, decolorization and at least partial degradation of azo dyes occurred only under anaerobic conditions and in the light. In another study, different PNSB strains isolated from various water sources (*Rhodobacter adriaticus*, *Rhodobacter blasticus*, *Rhodobacter capsulatus*, *Rhodovulum strictum*, and *R. palustris*) decolorized up to 96% of the tested azo dyes after only 2 days of illuminated and anaerobic incubation [22].

To circumvent problems associated with wastewater treatment using suspended cultures of bacteria, Wang and colleagues constructed a PBR to treat azo dye-contaminated wastewater [23]. The naturally developed biofilm contained anoxygenic phototrophic bacteria related to the *Rhodospseudomonas*, *Rhodomicrobium*, and *Chlorobium* lineages. This PBR allowed the removal of most of the organic load and up to 90% of the azo dyes. Over a 30-day run, the predominant phototrophic bacteria in the biofilm changed from purple bacteria to a

mixture of purple and GSB, and finally to mainly GSB. During this period the decolorization of azo dye increased from 60% to 90%, suggesting that GSB may also be capable of efficiently decolorizing azo dyes.

Purple bacteria belonging to the *Rhodospirillum* and *Rhodopseudomonas* genera have also been investigated for use in anaerobic biodegradation of halogenated aromatic pollutants such as 3-chlorobenzoate [5, 24].

## 5 Removal of Toxic Metals and Radioisotopes

Microorganisms – dead or alive, free or immobilized – can be used to remove toxic metals and radioisotopes from the environment [25]. Among anoxygenic phototrophic bacteria, this ability has especially been studied in PNSB [5].

The PNSB *Rhodobacter sphaeroides* bioaccumulates heavy metals including cadmium (Cd), nickel (Ni), and lead (Pb) [26–28], and metals with radioactive isotopes, cesium (Cs) and strontium (Sr) [29]. In one study, *R. sphaeroides* strain S accumulated  $\text{Cd}^{2+}$  with a higher efficiency under aerobic-dark conditions (93%) than under anaerobic-light conditions (50%) over 7 days at an initial  $\text{Cd}^{2+}$  concentration of 5 mg/L [27]. However, the ability to bioaccumulate metal ions is dependent on the strain of the organism under investigation. *R. sphaeroides* strain SSI is a spontaneous mutant of *R. sphaeroides* strain S with an increased ability to produce extracellular polymeric substances (EPS) on the cell surface [29]. The SSI strain has been shown to remove efficiently toxic metals including cadmium, uranium (U), cobalt (Co), mercury (Hg), and chromium (Cr). The removal of these metals has been attributed to the high amount of EPS produced on the surface of strain SSI because the strong negative charges of EPS efficiently adsorb many different kinds of metal ions. In one experimental setup with immobilized cells of strain SSI and 5 mg/L of each  $\text{Cs}^+$  and  $\text{Sr}^{2+}$ , about 100% of the cesium and 50% of the strontium were removed in 3 days. Other strains of *R. sphaeroides* and other PNSB have also been shown to remove efficiently (>90%) copper ( $\text{Cu}^{2+}$ ), zinc ( $\text{Zn}^{2+}$ ), and  $\text{Cd}^{2+}$  from contaminated shrimp pond water [30].

Zinc consumption by live *R. capsulatus* strain B10 cells has been reported at levels of 164 mg  $\text{Zn}^{2+}$  per gram of cell dry weight [31]. This is among the highest zinc biosorption capacities reported for any microorganism and this suggests that *R. capsulatus* could be useful for zinc bioremediation. The unusually high zinc biosorption capacity was again mainly attributed to the physicochemical properties of the EPS on the cell surface.

## 6 Photobiological Production of Hydrogen

Hydrogen ( $H_2$ ) has potential as an environmentally friendly fuel. Most, if not all, oxygenic as well as anoxygenic phototrophic microorganisms have the capacity to produce  $H_2$  but the physiological mechanisms for  $H_2$  production vary among these organisms [32]. Here, only  $H_2$  production in anoxygenic phototrophic bacteria is considered.

Although  $H_2$  production from phototrophic bacteria may not be economically viable by itself,  $H_2$  could be a valuable byproduct from other application of these organisms such as wastewater or biogas cleanup [33]. Photobiological  $H_2$  formation is catalyzed either by a hydrogenase ( $2H^+ + 2e^- \rightleftharpoons H_2$ ) or by nitrogenase as a byproduct from  $N_2$  fixation ( $8H^+ + N_2 + 8e^- \rightarrow H_2 + 2NH_3$ ). Most bacteria have the ability to re-uptake the produced  $H_2$  so this ability may have to be eliminated if the  $H_2$  production is to be useful.

In *Rhodospirillum rubrum* and other PNSB, photobiological  $H_2$  production is primarily caused by nitrogenase and therefore  $H_2$  production is induced under nitrogen limitation. Interestingly, if  $N_2$  is completely removed (for example, by using an inert gas), nitrogenase produces exclusively  $H_2$  and thus the  $H_2$  productivity is increased. In a nitrogen-limited batch culture of *R. rubrum*, a continuous production of 0.48 L/day  $H_2$  per gram cell dry weight was observed with lactate as electron donor [34]. *R. rubrum* and other purple bacteria also catalyze light-dependent  $H_2$  production from CO with a net reaction corresponding to the water-gas shift reaction ( $CO + H_2O \rightarrow H_2 + CO_2$ ) [35].

GSB also exhibit light- and nitrogenase-dependent  $H_2$  production. In these bacteria, inorganic sulfur compounds (sulfide, sulfur, and thiosulfate) are electron donors for photosynthesis and  $H_2$  production. Photobiological  $H_2$  production from organic compounds using these bacteria was demonstrated in a syntrophic co-culture of the GSB *Chlorobium vibrioforme* and the acetate-oxidizing, sulfur-reducing bacterium *Desulfuromonas acetoxidans* [36]. In this co-culture, the chemotrophic partner oxidized acetate and reduced sulfur to  $H_2S$ , and the phototrophic partner oxidized  $H_2S$  back to sulfur and produced  $H_2$  using nitrogenase. In this co-culture the productivity was 1.3 L/day  $H_2$  per gram cell dry weight with acetate as electron donor.

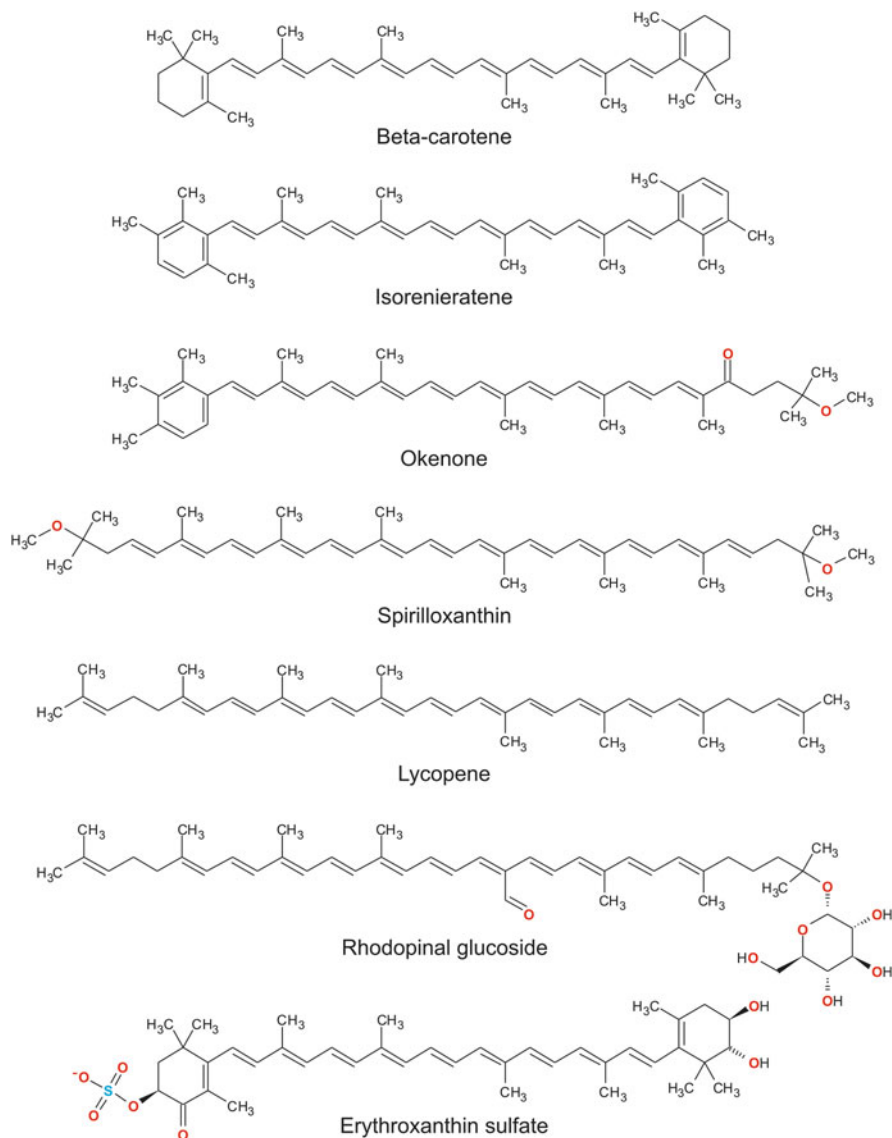
An alternative approach to photobiological  $H_2$  production in GSB might be engineered. In principle, the redox potentials of the type I photochemical reaction center ( $E_0'$  approx.  $-0.6$  V) and the soluble ferredoxins ( $E_0'$  approx.  $-0.5$  V) in GSB are low enough to allow reduction of  $H^+$  to  $H_2$  ( $E_0'$   $-0.42$  V). These strong cellular reductants are required for  $CO_2$  fixation by the reductive TCA cycle found in GSB. However, this reducing power might also be used for photobiological  $H_2$  production if a suitable hydrogenase could be inserted in these organisms that would accept electrons from the indigenous strong reductants. Then  $H_2$  production would be independent of  $N_2$  and nitrogenase.

## 7 Biosynthesis of Carotenoids and Other Terpenoids

Terpenoids (or isoprenoids) are compounds derived from one or more isoprene  $C_5$  units and constitute a very large range of natural compounds. A prominent group is the carotenoids ( $C_{40}$  compounds) found in all phototrophic organisms and some chemotrophic bacteria, archaea, and fungi. Carotenoids are yellow, orange, and red pigments with broad applications in the food, feed, nutraceutical, cosmetic, and pharmaceutical industries because of their vibrant colors and health-promoting activities [37]. Most commercialized carotenoids, such as beta-carotene and astaxanthin, are produced in microalgae although genetically modified *Escherichia coli* and yeasts that produce commercially valuable carotenoids are also available [38]. Anoxygenic phototrophic bacteria naturally produce a range of carotenoids (such as okenone and lycopene derivatives) which may have commercial interest [37, 39] (Fig. 3).

The most important role of carotenoids in phototrophic organisms is in protection from light [40]. The detrimental effects of light are much more serious under aerobic conditions than under anaerobic conditions because excitation of (bacterio) chlorophylls under aerobic conditions causes formation of reactive oxygen species (ROS). Mechanistically this occurs by reaction of excited triplet-state chlorophyll ( $^3Chl^*$ ) with ground-state molecular oxygen ( $O_2$ ), which results in formation of singlet oxygen ( $^1O_2$ ). Singlet oxygen is extremely reactive and detrimental to the cell. Carotenoids quench  $^3Chl^*$  and thereby prevent formation of singlet oxygen. This means carotenoids cannot be completely removed from phototrophic organisms growing under aerobic conditions. However, phototrophic bacteria growing under anaerobic conditions do not have this requirement and therefore carotenoids are not essential for anoxygenic phototrophic bacteria. This in turn means the carotenoids of these bacteria in principle are freely available for biosynthetic engineering.

Genetic engineering has been used to synthesize the commercially valuable carotenoid lycopene in the purple non-sulfur bacterium *R. rubrum* [41]. Here, the indigenous carotenoid biosynthetic pathway was interrupted by targeted gene inactivation and as a result the cells accumulated lycopene as the sole carotenoid in a content of 2 mg/g cell dry weight. Although this is not high compared to the yield obtained in genetically engineered *E. coli* (33 mg/g cell dry weight lycopene as the sole carotenoid [42]), additional engineering of *R. rubrum* could surely increase the yield. Using a similar approach, lycopene and zeta-carotene have also been produced as the sole carotenoid species in GSB by genetic manipulation of *Chlorobaculum tepidum* [43]. Because carotenoids are not required in anaerobic phototrophic bacteria, the flux of isoprene precursors to carotenoid biosynthesis in these organisms could be redirected to any isoprenoid compound such as valuable plant-type terpenoids [44].



**Fig. 3** Examples of carotenoids found in anoxygenic phototrophic bacteria. From [39] and Carotenoid Database (<http://carotenoiddb.jp/>)

## 8 Production of Functional Membrane Proteins

To study the structure and function of proteins, it is useful to overexpress the proteins in a foreign host organism to obtain amounts sufficient for experimentation. However, membrane proteins pose a challenge because they often denature in

the absence of a suitable membrane environment. For example, heterologous membrane proteins that are overexpressed in *E. coli* in a functional form are usually found in much lower titers than heterologous soluble proteins and tend to form inclusion bodies. To overcome this problem, the PNSB *Rba. sphaeroides* has been developed as a host for overexpression of functional membrane proteins [45]. This system takes advantage of the large content of intracellular membranes in purple bacteria which hold the membrane-bound antennae and enzymes of the photochemical machinery. In *Rba. sphaeroides* these internal membranes form intracytoplasmic membrane (ICM) vesicles that sequester newly synthesized foreign proteins and enable easy purification following cell lysis. Using this system, challenging membrane proteins in a functional form have recently been prepared and examined: human aquaporin 9 (hAQP9), human tight junction protein occludin (Occ), *Rba. sphaeroides* cellulose synthase enzyme complex (BcsAB), and *Rba. capsulatus* cytochrome  $c_y$  [46].

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