Review

The role of trace metals in photosynthetic electron transport in O_2 -evolving organisms

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Abstract

Iron is the quantitatively most important trace metal involved in thylakoid reactions of all oxygenic organisms since linear (= non-cyclic) electron flow from H_2O to NADP⁺ involves PS II (2–3 Fe), cytochrome b_6 -f (5 Fe), PS I (12 Fe), and ferredoxin (2 Fe); (replaceable by metal-free flavodoxin in certain cyanobacteria and algae under iron deficiency). Cytochrome c_6 (1 Fe) is the only redox catalyst linking the cytochrome b_6 -f complex to PS I in most algae; in many cyanobacteria and Chlorophyta cytochrome c_6 and the copper-containing plastocyanin are alternatives, with the availability of iron and copper regulating their relative expression, while higher plants only have plastocyanin. Iron, copper and zinc occur in enzymes that remove active oxygen species and that are in part bound to the thylakoid membrane. These enzymes are ascorbate peroxidase (Fe) and iron-(cyanobacteria, and most algae) and copper-zinc- (some algae; higher plants) superoxide dismutase. Iron-containing NAD(P)H-PQ oxidoreductase in thylakoids of cyanobacteria and many eukaryotes may be involved in cyclic electron transport around PS I and in chlororespiration. Manganese is second to iron in its quantitative role in the thylakoids, with four Mn (and 1 Ca) per PS II involved in O₂ evolution. The roles of the transition metals in redox catalysts can in broad terms be related to their redox chemistry and to their availability to organisms at the time when the pathways evolved. The quantitative roles of these trace metals varies genotypically (e.g. the greater need for iron in thylakoid reactions of cyanobacteria and rhodophytes than in other O₂-evolvers as a result of their lower PS II:PS I ratio) and phenotypically (e.g. as a result of variations in PS II:PS I ratio with the spectral quality of incident radiation).

Introduction

Trace metals play key roles in (non-cyclic) photosynthetic electron transport in the thylakoids of O₂-evolving organisms (Nicholls and Ferguson 1992; Kaneko et al. 1996; Merchant and Dreyfuss 1998). Mn is essential for O₂ evolution. Fe is involved in the PS II complex (cytochrome b_{559} and one additional Fe which is not in haem or part of a FeS complex), the cytochrome b_6 -f complex (cytochrome b_{563} , cytochrome f, Rieske non-haem iron-sulfur complex containing 2 Fe), cytochrome c_6 (an alternate to Cu-

containing plastocyanin), PS I complex (3 non-haem iron-sulfur complexes F_X , F_B and F_A , each containing 4 Fe) and ferredoxin (a protein with one non-haem iron-sulfur complex containing 2 Fe, an alternate to flavodoxin which contains no metals). Cu occurs in plastocyanin but, despite some claims, probably not in PS II. Ca^{2+} , at trace element levels, has a role in the thylakoid lumen in facilitating H_2O dehydrogenation and O_2 evolution (Hankamer et al. 1997).

In addition to these 'core' electron transport components involved in non-cyclic electron transport from H₂O to NADPH, there are also thylakoid compon-

ents which contain Fe, Cu and Zn that are involved in essential processes that are not part of the 'core' non-cyclic electron flow. These include the NAD(P)H dehydrogenase which may be involved in chlororespiration and cyclic electron flow around PS I; this complex contains non-haem Fe-sulfur centres. Additionally, the Mehler reaction involves the production of superoxide, which is converted to H₂O₂ using superoxide dismutase (containing Fe, Mn or Cu plus Zn when the whole range of organisms in which it occurs are considered); this H₂O₂ is ultimately converted to H₂O and O₂ using one of a number of enzymes. These generally involve Fe, such as the Fe-containing catalase (in a few 'primitive' cyanobacteria: Asada et al. 1993) or, with additional input of reductant from PS I, to H₂O using the Fe-containing ascorbate peroxidase (Asada 1992; Asada et al. 1993b). The thylakoidbound and stromal ascorbate peroxidases are produced by alternative gene splicing in Cucurbita moschata (Mano et al. 1997). In some algae, glutathione peroxidase, containing the non-metal Se, occurs; it is considered here because it 'spares' Fe in the alternate peroxidase, i.e. ascorbate peroxidase. The higher plant glutathione peroxidase which seems to be a lipid hydroperoxidase, does not contain Se (Stadtman 1996; Eshdat et al. 1997; Mullineaux et al. 1998). These metal-containing catalysts, which are involved in the Mehler peroxidase reaction and may also be regarded as means of detoxifying active oxygen species, are a part of the thylakoid membrane, although there is a parallel set of enzymes in the stroma that catalyze the same reactions. It is also significant that Couture et al. (1994) have reported a nuclear-encoded, thylakoid-expressed haemoglobin (containing Fe) in Chlamydomonas reinhardtii.

Our objectives in this review are:

- 1. To analyse the functional role of the metals in relation to the polypeptides, porphyrins, etc. in which they are found;
- 2. To assemble and interpret the adaptation-acclimation-related variations among taxa in the use of alternate catalysts containing different, alternative, metals (Cu or Fe) or a metal (Fe) *versus* no metal;
- 3. To describe and interpret the adaptation-acclimation-related variations in the ratios of metal-containing catalysis among taxa;
- 4. To use data discussed in 2 and 3. to indicate differences in the metal costs of photosynthesis at a given rate (mol O₂ or mol C per second;
- 5. To relate the discussion in 1.–4. to evolution in terms of the chemical appropriateness of the metal

to a given function and the evolutionary 'need' for that function at an earlier time when the availability of various metals was very different from the situation today, mainly as a result of a less oxidizing global environment earlier in the history of the biosphere.

We do not deal in detail with the involvement of trace metals in 'downstream' electron transport reactions associated with CO₂, NO₃⁻, NO₂⁻, N₂ and SO₄²⁻ assimilation which involve inputs, directly or indirectly, of photoproduced reductant and ATP.

For CO₂ assimilation, one trace metal requirement relates to the Zn in the carbonic anhydrase involved in CO₂ supply to Rubisco. Even when there is a functional CO₂ concentrating mechanism, there is usually some Rubisco oxygenase activity resulting in the production of phosphoglycolate and hence glycolate. Glycolate is metabolized in higher plants and some algae by Fe-containing catalase (Suzuki et al. 1991; Falkowski and Raven 1997; Iwamoto and Ikawa 1997). In other algae, glycolate is metabolized by glycolate dehydrogenase; here the reductant enters the respiratory chain which involves Fe and Cu (Raven 1988a, 1990; Suzuki et al. 1991; Falkowski and Raven 1997). Glyoxylate metabolism, insofar as this is known, involves a dehydrogenase step(s) which involves the respiratory chain, and hence Fe and Cu (Falkowski and Raven 1997). The alternative to glycolate metabolism is glycolate excretion (Falkowski and Raven 1997), which incurs no direct trace metal involvement (see H₂O₂ excretion: 2(c) below).

The trace metal requirements of the assimilation of NO_3^- and NO_2^- (cyanobacteria and eukaryotes) and NO_3^- , NO_2^- and N_2 (cyanobacteria only, at least among oxygenic organisms) have been dealt with in detail by Raven (1988a, 1990). The computed Fe and Mo requirement for growth at a given rate is significantly increased when NO_3^- , and especially N_2 , rather than NH_4^+ is the N source (Raven 1988a, 1990).

The functional roles of the metals

Linear electron flow with NADP⁺ as electron acceptor

Figure 1 shows the classic 'Z scheme' based on the concept of Hill and Bendall (1960), showing redox potential and emphasizing the places at which trace metals function. The quantities of trace metals indicated are for one protein (e.g. cytochrome c_6 or

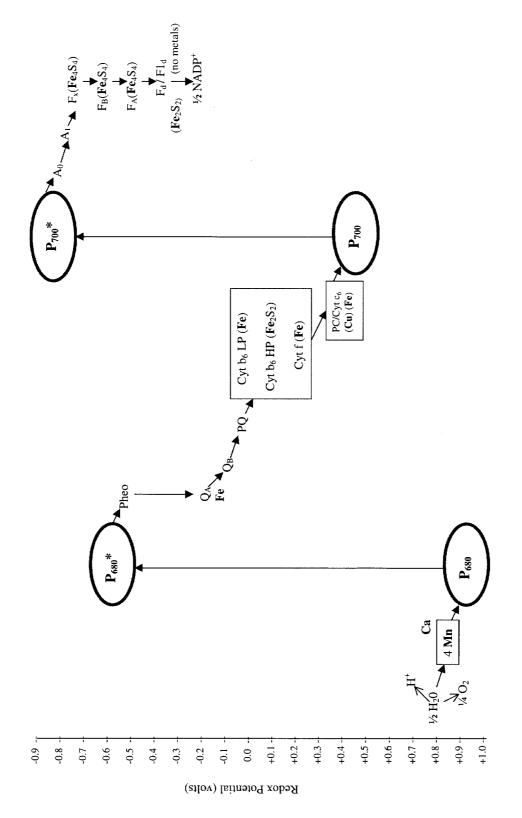


Figure 1. The 'Z scheme' of non-cyclic electron transport, showing the involvement of trace metals in various redox components in relation to their standard redox potential (modified from Falkowski and Raven 1997).

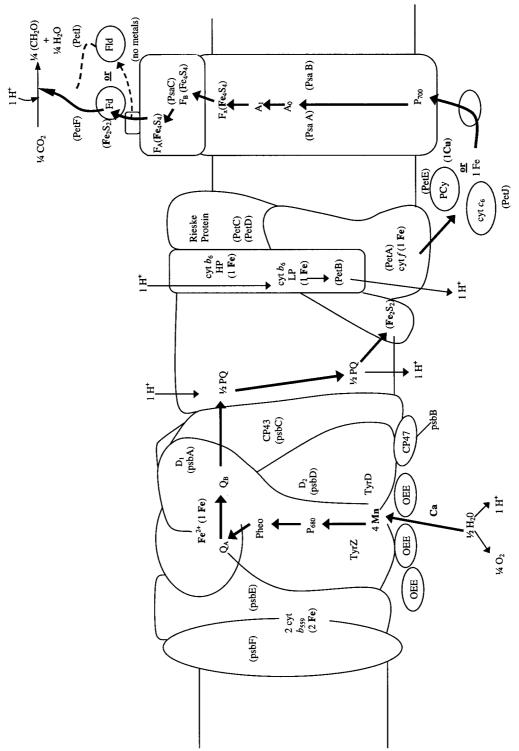


Figure 2. The spatial arrangement in the thylakoid membrane of the components of non-cyclic electron transport, showing the involvement of trace metals (modified from Falkowski and Raven, 1997, incorporating the likely arrangement of the b_6 — b_6 complex based on the crystal structure of the (larger) bovine b_{-c_1} complex from Iwata et al. 1998 and Smith 1998).

plastocyanin) or one complex (e.g. the cytochrome b_6 -f complex).

Figure 2 indicates the spatial (as opposed to the energetic, Figure 1) disposition of the various trace metal-containing catalysts of linear electron transport. Also indicated are the stoichiometries of electron flow to vectorial H^+ fluxes; with a proton-motive Q cycle the H^+/e^- ratio is 3 which, with a H^+/ATP ratio of 4, produces 0.75 ATP per electron transferred from H_2O to NADP⁺ (Falkowski and Raven 1997).

Table 1 gives molecular data on the proteins that bind trace metals in the linear electron transport chain, and indicates how the metals are bound. The data in Table 1 show that, with a 1:1:1:1:1 stoichiometry of PS II:cyt b_6 -f:cyt c_6 : PS I:ferredoxin, there are 23 Fe, 4 Mn and no Cu per 'linear electron transport chain', while with cytochrome c₆ replaced by plastocyanin and ferredoxin replaced by flavodoxin, the 'linear electron transport chain' with a 1:1:1:1:1 stoichiometry of PS II:cyt b₆-f:plastocyanin:PS I:flavodoxin has 20 Fe, 4 Mn and 1 Cu. The Fe content of these idealized linear electron transport chains assumes 2 cytochrome b_{559} , each with 1 Fe, per PS II; if, as is very possible there is only 1 cytochrome b_{559} per PS II, the number of Fe per linear electron transport chain here, and subsequently in this paper, should be reduced by 1. Table 2 lists the Fe content of the various complexes.

Cytochrome f, cytochrome c_6 , the iron sulphur centres of the Rieske protein and the electron acceptor complex of PS 1, plastocyanin, ferredoxin and flavodoxin have a straightforward role in the linear electron transfer pathway.

The two haems of the cytochrome b_6 provide the transmembrane electron transfer path for the Q cycle mechanism in the cytochrome b_6 -f complex.

Cytochrome b_{559} is an essential component of PS II, deletion of the genes for the polypeptide results in failure to assemble the complex. Under various experimental conditions, it may be either oxidised or reduced by the PS II reaction centre, it also has two redox potential forms. However its *in vivo* function is unknown. A role in protection against photoinhibition, possibly by providing a path for cyclic electron transfer around PS II, seems most likely (Whitmarsh et al. 1994; Prosil et al. 1996). There may also be a role for this cytochrome in cyclic electron flow round PS I (Miyake et al. 1995).

The function of the non-haem iron in PS II is also uncertain. Unlike the homologous Fe in the purple bacterial reaction centre it can undergo oxidation and

reduction. It is structurally placed between the primary and secondary quinone electron acceptors but is not thought to act as an electron carrier between them as its redox potential, +400 mV, is too high for oxidation by the quinone (Diner and Petrouleas 1988), electron flow only occurring when it is in the Fe²⁺ form. Electron flow is also dependent on the presence of bicarbonate as a ligand to the Fe. Removal of the bicarbonate or replacement by formate inhibits electron transfer and may contribute to photoinhibition (Blubaugh and Govindjee 1988).

Mn is an essential component of the water oxidation complex. Four Mn atoms are organized in two interacting pairs in the form of μ -oxo bridged dimers (Yachandra et al. 1993, MacLachlan et al. 1994b; Riggs-Gelasco et al. 1996). Ca is also an essential component of the complex (Yocum 1991) possibly as an integral component of the Mn complex (MacLachlan et al. 1994a; Latimer et al. 1995). The Mn complex provides the redox system for the accumulation of the 4 oxidizing equivalents required for water oxidation, providing the gating mechanism between the single electron turnover of the reaction centre and the oxidation of water. Three of the five oxidation states of the water oxidizing complex are known to involve different Mn redox states (Messinger et al. 1997).

Not listed in Table 1 are two other c-type cytochromes which are associated with the photosynthetic apparatus of many cyanobacteria and algae, and which have not so far been shown to act in linear electron transport or, indeed, in any other essential redox reactions in the organisms in which it occurs (Kerfeld and Krogmann 1998).

One of these c-type cytochromes is cytochrome c_{549} LP, with a redox potential of -250 mV (Navarro et al. 1995). Cytochrome c_{549} LP has been shown to occur in cyanobacteria, red algae (Rhodophyta), diatoms (Heterokontophyta) and the ulvophycean Bryopsis (Chlorophyta) (Evans and Krogmann 1983). Although some reports indicate thylakoid membrane association of cytochrome c_{549} LP, with the molecule located toward the lumenal side of the membrane and essential to the structure of PS II in the organisms in which it occurs, in other cases (e.g. Synechocystis sp. strain 6803) this cytochrome appears to be entirely water-soluble (Navarro et al. 1995; Kerfeld and Krogmann 1998). If there is one molecule of cytochrome $c_{549}LP$ per PS II reaction centre, then each 'linear electron transport chain' has one more Fe than in otherwise similar organisms which lack cytochrome c_{549} LP (e.g. higher plants). Cytochrome c_{549} LP can

Table 1. Thylakoid proteins involved in electron transport which bind trace metals

Gene	Polypeptide	Mass of mature polypeptide	Function	Metals bound	References
psbA psbD	Hydrophobic D1 protein of PS II reaction centre Hydrophobic D2	38 021 39 418	Catalysts of primary photochemistry of PS II	1 Fe per heterodimer as neither haem nor FeS; 4 Mn	
	protein of PS II reaction centre				
psbE	Hydrophobic α -cyt b_{559}	4409	Photoprotection? (not a	1 Fe per heterodimer as	Hankamer et al. (1997);
psbF	Hydrophobic β-cyt b ₅₅₉	4195	component of linear electron transport)	haem; 1 or 2 <i>b</i> ₅₅₉ (1 or 2 Fe) per PS II reaction centre	Hoganson and Babcock (1997); Krieger and Rutherford (1997); Rhee et al. (1998)
psbO	33 kDa extrinsic protein	26 539	Stabilizes Mn cluster bound to D1/D2; C1 ⁻ binding	Ca ²⁺ binding	
psbP	23 kDa extrinsic protein	20 210	Cl ⁻ binding		
psbQ	16 kDa extrinsic protein	16 523	Cl ⁻ binding		
psaA	Hydrophobic 83.2 kDa protein of PS I	83 200	Catalysts of primary photochemistry	4 Fe per heterodimer as F _x [4 Fe – 4S]; 1	Evans et al. (1981) Golbeck
psaB	Hydrophobic 82.5 kDa protein PS I	82 500	of Ps I	heterodimer per PS I, so 4 Fe per PS I	(1992); Evans and Nugent (1993); Berry et al (1997); Scheller et al. (1997); Schubert et al. (1997)
psaC	Hydrophilic stromal proteim pf PS I	8900	Transfer of electrons from F_x to ferredoxin or flavodoxin $psaC$; 1 $psaC$ per PS I, so 8 Fe per PS I	8 Fe per heterodimer as F_A [4 Fe – 4 S] and F_B [4 Fe – 4 S] per (1997); Schubert et al. (1997)	Evans and Nugent (1993); Scheller et al.

Table 1. Continued

Gene	Polypeptide	Mass of mature polypeptide	Function	Metals bound	References
PetF	Ferredoxin; hydrophilic stromal protein of PS I	11 000	Transfer of electrons from FA to ferredoxin-NADP oxidoreductase and thence to NADP+, to NO ₂ -reductase to ferredoxin-dependent glutamate synthase, to ferredoxin-dependent thioredoxin reductase, and to monodehydro-ascorbate	2 Fe as *[2 Fe – 2 S]	Evans and Nugent (1993); Miyake and Asada (1994); Scheller et al. (1997)
PetA	Cytochrome <i>f</i> component of cytochrome <i>b</i> ₆ - <i>f</i> –FeS complex; hydrophobic	32 038	Redox reactions; H ⁺ pumping	1 Fe per monomer bound as haem; 1 cyt f per b_6 - f FeS	
PetB	Cytochrome b_6 component of cytochrome b_6 - f FeS complex; hydrophobic	24 166	Redox reactions, H^+ pumping	1 Fe per monomer bound as haem; 2 cyt b_6 per b_6 - f comples;	Cramer et al. (1996 Hpe (1993)
PetC	Rieske non- haem Fe polypeptide; hydrophobic	19116	Redox reactions	2 Fe per complex 2 Fe as [2 Fe – 2S];	
PetJ	Cytochrome c_6 (alternative to plastocyanin); (hydrophilic protein in lumen)	10 000	Electron transfer from <i>b</i> ₆ - <i>f</i> -FeS complex to Ps I	1 Fe per monomer bound as Fe	
PetE	Plastocyanin (alternative to cytochrome c_6); (hydrophilic protein in lumen)	10 000	Electron transfer *from b ₆ -f-FeS complex to PS I	1 Cu	Falkowski and Raven (1997)

Table 2. Fe content of proteins or protein complexes involved in photosynthetic electron transport. From references in Tables 1 and 4, and in the text

Protein or protein complex	Number of Fe atoms per or protein complex
PS II	(2-)3
PS I	12
Cytochrome b ₆ -f	5
Cytochrome c_6	1
Ferredoxin	2
Catalase	4
Fe superoxide dismutase	1
Ascorbate peroxidase	1
NAD(P)H/PQ oxidoreductase	8-18

function *in vitro* as a catalyst of cyclic electron flow, via *its* rapid reduction by at least some isoforms of ferredoxin and its oxidation by plastocyanin (or, presumably, cytochrome c_6) (Navarro et al. 1995; Kerfeld and Krogmann 1998). Another suggested role of cytochrome $c_{549}\mathrm{LP}$ is in anoxia tolerance via an interaction with (soluble, cytosolic or stromal) hydrogenase (Kerfeld and Krogmann 1998; Merchant and Dreyfuss 1998). Neither of these roles seems to be particularly compatible with location of cytochrome $c_{549}\mathrm{LP}$ toward the lumenal side of the thylakoid membrane.

The other c-type cytochrome is cytochrome c_{552} (cyt M). This soluble cytochrome c has a redox potential similar to that of cytochrome c_6 and of plastocyanin, so that it could act as a third alternative to these two redox catalysis in certain cyanobacteria and algae (Kerfeld and Krogmann 1998; see 'Genetic and environmentally induced expression of alternate catalysts containing different metals or no metals' below). However, cyt M is present in much smaller amounts than cytochrome c_6 (or plastocyanin), which could constrain any obligatory role in linear (or cyclic) electron transport (Kerfeld and Krogmann 1998). Certainly the molecular genetic elimination of cytochrome M in a cyanobacterium has no observable phenotypic effect, albeit in cells that can express both cytochrome c_6 and plastocyanin (Kerfeld and Krogmann 1998; see also Manna and Vermaas 1997).

Table 3. Occurrence of plastocyanin and cytochrome c₆ (see Raven et al. 1989; Raven et al. 1990; Kerfeld and Krogmann 1998)

Taxon	Plastocyanin	Cytochrome c ₆	
Cyanobacteria	In most	In all?	
(including Chloroxybacteria?)			
Rhodophyta	Absent	Present	
Heterokontophyta			
Bacillariophyceae	Absent	Present	
Chrysophyceae	(Absent)	Present	
Fucophyceae	Absent	Present	
(= Phaeophyceae)			
Synurophyceae	(Absent)	Present	
Tribophyceae	Absent	Present	
(= Xanthophyceae)			
Haptophyta	(Absent?)	Present	
Cryptophyta	?	?	
Dinophyta	?	?	
Chlorophyta	In all?	In some	
Embryophyta	Present	Absent	
Euglenophyta	Absent	Present	
Chlorachniophyta	?	?	

Genetic and environmentally induced expression of alternate catalysts containing different metals or no metals

The major alternative catalysts in the linear electron transport pathway are the water-soluble proteins involved in the cytochrome b_6 -f complex to P_{700} electron transfer (plastocyanin or cytochrome c_6) and the F_A to ferredoxin-NADP⁺ reductase electron transfer (ferredoxin or flavodoxin). The former pair of redox carriers function on the lumenal side of the thylakoid membrane while the latter pair function on the stromal (eukaryotes) or cytosol (prokaryotes) side of the membrane.

Table 3 shows the taxonomic (phylogenetic) distribution of the occurrence of the members of the alternative pair plastocyanin/cytochrome c_6 . Although it has previously been thought that some cyanobac-

teria lack the capacity to make plastocyanin, more recent evidence, genetic and other, indicates a very widespread capacity for plastocyanin synthesis in cyanobacteria sensu lato, i.e. including chloroxybacteria or prochlorophytes (Arudchandran et al. 1994; Clarke and Campbell 1996; Kaneko et al. 1996). Among the green algae (Chlorophyta), plastocyanin seems to be universal, with the capacity to produce cytochrome c_6 in many cases. Plastocyanin seems to be present in all higher plants (Embryophyta, i.e. bryophytes and tracheophytes) and there is no evidence for cytochrome c_6 . The situation in the closest living relatives of the ancestors of the embryophytes, i.e. the class Charophyceae sensu lato, or Klebsomidiophyceae of the division Chlorophyta (van den Hoek et al. 1995), is unclear. All of the other groups of algae seem to lack plastocyanin and use only cytochrome c_6 . This is true of the organisms that obtained their plastids by secondary endosymbiosis of a green eukaryote, i.e. the Euglenophyceae (the situation in the Chlorarachniophyceae is not clear), the red algae which, presumably, like the green algae, have chloroplasts derived from endosymbiosis of a cyanobacterium (but see Stiller and Hall 1997), as well as the Heterokontophyta and, probably, Haptophyta which obtained their plastids by secondary endosymbiosis of red algae. Use of plastocyanin or cytochrome c_6 in other significant non-green algal taxa, which also obtained their plastids by secondary endosymbiosis (the Cryptophyta and Dinophyta), is not clear. Present evidence shows that goes both cytochrome c_6 and plastocyanin in eukaryotes are invariably encoded by nuclear genes.

A number of provisos are needed in considering the distribution among algae of cytochrome c_6 and plastocyanin.

- 1. The older literature was confounded by confusion of cytochrome c_6 with cytochrome f, a point which was elegantly clarified by Wood (1977; see Falkowski and Raven 1997).
- 2. The use of visible wavelength spectrometry (a broad redox-dependent band centred at ~ 580 nm) to characterize plastocyanin led to some 'false positives' in plastocyanin distribution, e.g. reports of its presence in red algae (see Raven et al. 1990; Falkowski and Raven 1997).
- 3. Nucleotide sequence data could be useful in showing whether the genetic information for only one, or for both, of the alternates is present in a given organism (e.g. Kaneko et al. 1996).
- 4. There is also the possibility that the gene for the cytochrome c_6 or plastocyanin expressed in the

plastid of organisms (hosts) whose plastids arose from secondary endosymbiosis may not have arrived in the host with the nucleus of the eukaryote ingested in the secondary endosymbiosis that gave the host its plastid, with transfer of the genes encoding plastocyanin or cytochrome c_6 to the host nucleus.

The alternative is that the cytochrome c_6 (or plastocyanin) genes could have been present in the host nucleus from a prior round of plastid acquisition by primary (prokaryotic) or secondary (eukaryotic) endosymbiosis, a suggestion made by Cavalier-Smith (1982, 1992). Although this may not be the simplest explanation of the data for cytochrome c_6 (or plastocyanin), it has relevance for 'explaining' odd juxtapositions of various carotenoids in different algal taxa (Raven 1987a). Although there is evidence that a mutant cyanobacterium (Synechocystis sp. PCC 6803) lacking both plastocyanin (as a result of Cu deprivation) and cytochrome c_6 (by mutation) still have functionality in electron transfer from the cytochrome b_6 -f complex to P_{700} culminating in photolithotrophic growth (Zhang et al. 1994), more recent work confirms that strain 6803 contains the genes for both cytochrome c_6 and plastocyanin (Kaneko et al. 1996) and that mutants of PCC 6803 with neither cytochrome c_6 nor plastocyanin are non-viable under conditions permitting phototrophic or chemoorganotrophic growth (Manna and Vermaas 1997). The Cu deficiency imposed on the Synechocystis sp. strain 6803 by Zhang et al. (1994) still permitted some expression of plastocyanin (and cytochrome oxidase) (Manna and Vermaas 1997). There is also the possibility that soluble cytochrome M, which is encoded in the genome of Synechocystis sp. strain 6803 (Kaneko et al. 1996) and which has a redox potential similar to that of cytochrome c_6 and of plastocyanin, could function in electron transfer from the cytochrome b_6 -f complex to P₇₀₀ (Kerfeld and Krogmann 1998). However, the level of expression of cytochrome M, at least in organisms that have normal expression of cytochrome c_6 and of plastocyanin, is much lower than that of these two redox proteins (Kerfeld and Krogmann 1998).

Plastocyanin and cytochrome c_6 have closely similar functional characteristics in the transfer of electrons between the cytochrome b_6 -f complex and P_{700} . They also have similar molecular masses, so in terms of evolutionary optimality there is little reason to use one rather than the other. Much physiological evidence, now supported by molecular data on the control of transcription and translation, has accumulated on

the relative expression of plastocyanin and of cytochrome c_6 as a function of Cu and Fe availability, the Cu-containing plastocyanin being expressed more than the Fe-containing cytochrome c_6 when Cu availability is high but Fe availability is low, and vice versa for higher Fe availability than Cu availability (Falkowski and Raven 1997). Plastocyanin represents a larger fraction of the total known cellular component containing Cu than does cytochrome c_6 as a fraction of the total known cellular components containing Fe, especially in organisms lacking Cu–Zn superoxide dismutase (i.e. almost all algae, the Charophyceae sensu lato and peridinin-containing dinoflagellates being exceptions) the relative sparing effect on Cu use by expression of cytochrome c_6 rather than plastocyanin is greater than that on Fe use by the expression of plastocyanin rather than cytochrome c_6 .

Less information is available on the occurrence of the other major alternative pair of catalysts, the iron-containing ferredoxin and the metal-free flavo-protein flavodoxin. It is believed (Raven et al. 1989) that the higher plants (bryophytes and tracheophytes) are unable to express flavodoxin and always rely on ferredoxin. Among the cyanobacteria and the algae, the alternative expression of ferredoxin and flavodoxin is common. However, by contrast with the cytochrome c_6 -plastocyanin pair, there is only one reported case of the occurrence of flavodoxin to the exclusion of ferredoxin, i.e. the occurrence of flavodoxin but not ferredoxin in the marine red macroalga *Chondrus crispus* (see Raven et al. 1990).

In those algae with the potential to express either ferredoxin or flavodoxin, the 'normal' electron carrier between the sequence A₀ (a form of chlorophyll a, the initial electron acceptor of PS I), A₁ (phylloquinone) and the iron-sulfur complexes F_X, F_B and F_A, and the ferredoxin-NADP oxidoreductase, is ferredoxin. However, with Fe deficiency there is increasing expression of flavodoxin. Indeed, the expression of flavodoxin as detected immunologically is used as a bioassay for Fe availability in the ocean (La Roche et al. 1996). Unlike the alternative pair cytochrome c_6 /plastocyanin, there is a significant difference between the catalytic effectiveness (on a molar basis) of ferredoxin and flavodoxin; ferredoxin has a higher specific reaction rate (mol electron transferred per mol protein per second) than does flavodoxin in their normal in situ place in non-cyclic electron transport. Although this may be a determinant of the timing of replacement of ferredoxin by flavodoxin during developing Fe deficiency in algae and cyanobacteria with

the potential to express either (or both) catalysts, La Roche et al. (1996) found that flavodoxin expression in the phytoplankton of the N.E. Pacific began rather early in the development of Fe deficiency.

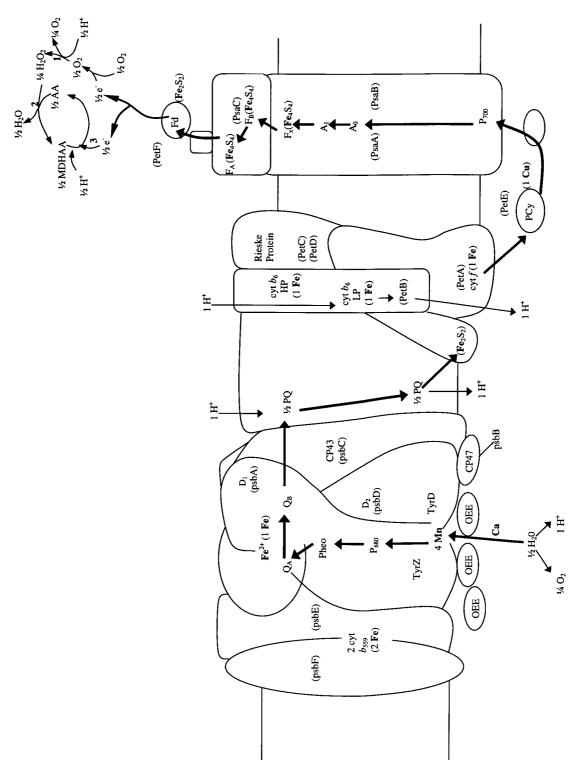
Linear electron flow with O_2 as electron acceptor: The Mehler peroxidase reaction

The overall reaction here is that of O_2 evolution by the water dehydrogenation reaction of the oxidizing side of PS II balanced by the reduction of O_2 to water by the reducing side of PS I. With 3 H⁺ pumped per electron transferred (and per 2 photons absorbed), the H⁺/ATP ratio of 4 gives 0.75 ATP produced per electron transferred and (with optimal excitation energy allocation to the two photosystems) 0.375 ATP produced per photon absorbed (Figure 3).

We now know (see 'Introduction', and Wiese et al. 1998) that the sequence of electron transport from the reducing side of PS I to O_2 involves superoxide and hydrogen peroxide as intermediates; this is termed the Mehler peroxidase reaction.

We have seen in the 'Linear electron flow with NADP⁺ as electron acceptor' section above that linear electron transfer to NADP⁺ with a 1:1:1:1:1 stoichiometry of trace metal-containing components has 20–23 Fe, 4 Mn and 0–1 Cu per electron transfer chain.

For the Mehler peroxidase reaction there is an additional need for trace metals as a result of the production of superoxide followed by its dismutation by a metal-containing superoxide dismutase (Fe, or Cu/Zn), and removal of the hydrogen peroxide produced by superoxide dismutase by the Fe-containing catalase or hydroperoxidase (Fe) in some cyanobacteria, glutathione peroxidase (Se) in some cyanobacteria and (eukaryotic) algae, and ascorbate peroxidase in most cyanobacteria and eukaryotic algae and all higher plants (Henry et al. 1976; Asada et al. 1980; Tel-Or et al. 1985; Shigeoka et al. 1987; Harrison et al. 1988; Price and Harrison 1988; de Jesus et al. 1989; Halliwell and Gutteridge 1989; Miyake et al. 1991; Shigeoka et al. 1991; Asada 1992; Asada et al. 1993b; Takeda et al. 1993; Kaneko et al. 1996; Stadtman 1996; Butow et al. 1997; Eshdat et al. 1997; Noctor and Foyer 1998; Obinger et al. 1998; Regelsberger et al. 1999) (Table 4). Not all of these enzymes are thylakoid-bound, but at least a fraction of cyanobacterial and plastidic superoxide dismutase and ascorbate peroxidase are thylakoid-associated (Asada et al. 1980; Mano et al. 1997).



dismutase (1) with 1 Cu and 1 Zn per molecule, ascorbate peroxidase (2) with 1 Fe per molecule and ferredoxin-monodehydroascorbate oxidoreductase (3) with no known trace metal requirement (Miyake and Asada 1992a,b; Miyake et al. 1993). A stromal pathway with the same trace metal electron, O₂ and H⁺ stoichiometry involves NADPH, NADPH-oxidized glutathione Figure 3. The Mehler peroxidase reactions at the reducing end of PS I in higher plants and charophycean algae. The reactions shown are for the thylakoid-associated enzymes superoxide oxidoreductase, and reduced glutathione-dehydroascorbate oxidoreductase.

Table 4. Turnover of metal-containing catalysts in different mechanisms of disposing of the O_2 - produced in the Mehler reaction by PS I

Organisms	Superoxide dismutase turnovers per 4 electrons from PS I	$ m H_2O_2$ -disposing enzyme turnovers per 4 electrons from PS I ^b
'Primitive' cyanobacteria	Two turnovers of Fe-SOD	Two turnovers of (Fe) catalase or hydroperoxidase
Many cyanobacteria; most eukaryotic algae (except Charophyceae and peridinin- containing Dinophyceae) (in absence of Se)	One turnover of Fe–SOD	One turnover of (Fe) ascorbate peroxidase ^a
Charophycean green algae; peridinin-containing Dinophyceae; higher plants	One turnover of Cu–Zn SOD	One turnover of (Fe) ascorbate peroxidase ^a
Chlamydomonas, many diatoms, grown with Se; some cyanobacteria	One turnover of Fe-SOD	One turnover of (Se) glutathione peroxidase ^a
(Aquatic plants?)	(Two turnovers of Fe–SOD (algae), Cu–Zn SOD (higher plants?)	No enzyme used; H ₂ O ₂ efflux?)

^aOne turnover of ascorbate on glutathione peroxidase is based on the overall stoichiometry for the two electron reaction in which one ascorbate (or two GSH) is converted to one dehydroascorbate (or one GSSG). While this is appropriate for glutathione peroxidase, the ascorbate peroxidase reaction converts two ascorbate to two monodehydroascorbate. In the case of the stromal enzyme the two monodehydroascorbate molecules dismute to one ascorbate and one dehydroascorbate (which is reduced in a two electron reaction by GSH to regenerate ascorbate), while for the thylakoid-associated enzyme the two monodehydroascorbate molecules are reduced by ferredoxin to regenerate two ascorbate (see Miyake and Asada 1994).

^bReferences: Henry et al. (1976); Asada et al. (1980); Price et al. (1987); Price and Harrison (1988); Harrison et al. (1988); de Jesus et al. (1989); Halliwell and Gutteridge (1989); Miyake et al. (1991); Shigeoka et al. (1991); Asada (1992); Miyake and Asada (1992a.b); Asada et al. (1993b); Miyake and Asada (1993); Takeda et al. (1993); Ogawa et al. (1995); Kaneko et al. (1996); Mano et al. (1997); Stadtman (1996); Butow et al. (1997); Eshdat et al. (1997); Regelsberger et al. (1999).

Attempts to compute the minimum requirement of Fe, Cu, Zn and Se downstream of ferredoxin in the various forms of the Mehler reaction relative to those of linear electron transport requires knowledge of the flux of electrons through the various reactions relative to those through the linear pathway, and of the specific reaction rate of the enzymes involved.

Dealing first with the stoichiometry issue, the simplest process is that found in 'primitive' cyanobacteria wherein (Fe) superoxide dismutase activity is followed by that of catalase. Here two turnovers

of Fe superoxide dismutase and of Fe-containing catalase are needed for every four electrons which eventually end up in 2 H₂O (Table 4). This 'primitive' cyanobacterial state can be produced in higher plants; thus, expression of *Escherichia coli* catalase in chloroplasts of *Nicotiana tabacum* greatly reduces the expression of ascorbate peroxidase (Shikanai et al. 1998b). In other cyanobacteria and eukaryotic algae other than the Charophyceae in the Division Chlorophyta and the peridinin-containing dinoflagellates, the H₂O₂ produced by (Fe) superoxide dismutase is fur-

ther metabolized by Fe-containing ascorbate peroxidase. However, in this case, only half as much H₂O₂ is generated per electron flowing out of PS I as in the catalase case mentioned above. This is because only two of these electrons pass from the reducing end of PS I to H₂O via O₂. and H₂O₂: the other pair of electrons is routed via ferredoxin, NADPH and glutathione to reduce back to ascorbate the dehydroascorbate generated by ascorbate peroxidase in the case of soluble (stromal) ascorbate peroxidase, and via ferredoxin to reduce back to ascorbate the monodehydroascorbate generated by the thylakoid-bound ascorbate peroxidase (Table 4; Figure 3). For charophycean green algae, higher plants and peridinin-containing dinoflagellates the situation is as described for the ascorbate peroxidase-containing cyanobacteria and non-charophycean algae, except that the superoxide dismutase involved is the Cu–Zn type rather than the Fe form. A further stoichiometric variation is in those eukaryotic algae (Chlamydomonas; some diatoms) expressing glutathione peroxidase, where a pair of electrons is transferred via one turnover each of Fe superoxide dismutase and of Se-containing glutathione peroxidase (Table 4).

The final possibility was suggested by Collén (1994) and Collén and Pedersén (1996), who pointed out that the high membrane permeability of H₂O₂ makes H₂O₂ efflux an option for disposing of this intermediate in the Mehler peroxidase reaction in aquatic organisms with a large extracellular sink for H₂O₂ (see Patterson and Myers 1973) (Table 4). Collén (1994) and Collén and Pedersén (1996) found a significant efflux of H₂O₂ from the green marine (ulvophycean) macroalga *Ulva lactuca* relative to the (presumed) rate of H₂O₂ production in the Mehler reaction in the light, bearing in mind that the Ulvophyceae have glycolate dehydrogenase rather than the H₂O₂-generating glycolate oxidase which deals with the small glycolate production rate found in those ulvophyceans (e.g. Ulva) with a CO2 concentrating mechanism (Suzuki et al. 1991; Raven 1997c).

A diffusive efflux of $\rm H_2O_2$ in algae is rendered more plausible by the smaller inhibition of photosynthesis by (added) $\rm H_2O_2$ in those algae tested than in higher land plants (Takeda et al. 1995; Korb and Raven, unpublished data). In *Chara*, a close freshwater relative of higher plants, photosynthetic $\rm CO_2$ fixation appears to be more sensitive to $\rm H_2O_2$ than in other algae (Korb and Raven, unpublished data). The high permeability of phospholipid bilayers to $\rm H_2O_2$ (Takahashi and Asada 1983; Mathai and Sitava-

man 1994), augmented by $\rm H_2O_2$ permeation through aquaporins (Henzler and Steudle 1998) also renders a diffusive efflux of $\rm H_2O_2$ in algae more plausible, although aquaporins may not occur in all plasmalemmas, or at all in plastid envelopes (Raven 1982, 1995). The 'basal' $\rm H_2O_2$ permeability is equivalent to a water layer of 200 μ m. This diffusion boundary layer thickness is similar to what may occur around many aquatic macrophytes in poorly stirred environments (Raven 1984).

The efflux of all of the H₂O₂ produced in the Mehler reaction would obviate the need for H₂O₂metabolizing enzymes such as catalase, ascorbate peroxidase or glutathione peroxidase, provided that the H₂O₂ levels were reduced to values similar to those achieved by these enzymes. Such an absence of H₂O₂metabolizing enzymes would mean two turnovers of Fe superoxide dismutase per four electrons from PS I used in producing 2 H₂O₂. However, all O₂evolvers have activities of at least one of the H₂O₂metabolizing enzymes, so the efflux possibility suggested by Collén (1994) apparently does not obviate the need for enzymic H₂O₂ disposal. Furthermore, the rates of H₂O₂ efflux in Enteromorpha intestinalis and Codium fragile in the light (and the dark) are much lower (Korb and Raven, unpublished) than the rate measured in *Ulva lactuca* (a very close relative of Enteromorpha) by Collén (1994) and Collén and Pedersén (1996). The conclusion about diffusive H₂O₂ efflux is that it is not a universal and major means of disposing of H₂O₂ produced in the Mehler reaction in aquatic organisms, and hence of possibly economizing on metal use.

The very low membrane permeability of O_2 .—, and its very high reactivity (Takahashi and Asada 1983) mean that there is no possibility of replacing superoxide dismutase by the efflux of O_2 .— in aquatic organisms.

Quantification of minimal metal requirements for the Mehler (peroxidase) reaction requires knowledge of the specific reaction rate (on a catalytic metal basis) of the enzymes involved at the *in vivo* substrate concentrations. Specific reaction rate is not a simple function of the (sole) substrate concentration for two of the enzymes considered here, i.e. superoxide dismutase and catalase, since these dismutation reactions involve two ${\rm O_2}^{--}$ (and 2 H⁺) and two ${\rm H_2O_2}$ respectively.

Assuming a maximum specific reaction rate of $400~\rm s^{-1}$ for our idealised linear redox chain with a 1:1:1:1:1 stoichiometry at light saturation (i.e. $400~\rm electrons$ transferred per redox chain per second: see

Raven 1990), then we can use published specific reaction rates of the enzymes to compute how much of the enzyme is needed to consume the active oxygen species derived from the reaction of ferredoxin, etc., with O₂. Superoxide dismutase has a specific reaction rate of 10^6 s⁻¹ (mol O₂ ·- mol⁻¹ enzyme s⁻¹, i.e. mol electron mol^{-1} enzyme s^{-1}), so that only $400/10^6$ or 4.10^{-4} mol SOD would be needed per redox chain if the steady-state O2.- level were sufficient to saturate the enzyme ($\sim 1 \text{ mol m}^{-3}$): Schomburg and Stephan (1994). However, a higher ratio of superoxide dismutase to redox chain would be needed if the steady-state O_2 – concentration at the active centre of superoxide dismutase is lower, and especially if diffusion of O₂ - from each of 5000 redox chains to a superoxide dismutase molecule leads to high concentrations of O_2 at the site of O_2 generation, which might cause more damage than would otherwise be the case; with 1 Fe, or 1 Cu + 1 Zn per superoxide dismutase molecule, there must be at least 4×10^{-4} mol Fe, or 4×10^{-4} mol Cu + Zn per mol redox chain. These computed SOD requirements apply when catalase is the agent removing H₂O₂; if peroxidase is used, only half as much superoxide dismutase, and hence Fe or Cu + Zn is needed per linear redox chain (Tables 4 and 5). The volume-based quantities of Cu-Zn superoxide dismutase in chloroplasts (Asada and Takahashi 1987), together with the data in Table 4.1 of Lawlor (1993), yields values of up to 0.5 mol superoxide dismutase per mol electron transport chain, i.e. 10^3 times the computed minimum value. Recent data show even higher values of 1.0 mol superoxide dismutase per mol PS I (or per mol electron transport chain): Ogawa et al. (1995). This apparent excess capacity (see Table 5) is related to maintaining steady-state O₂⁻⁻ concentrations at far below the concentration needed to saturate superoxide dismutase (Asada and Takahashi 1987). However, transgenic experiments have shown that SOD activity in the chloroplast may restrict the rate of the Mehler (peroxidase) reaction under some conditions (Arisi et al. 1998).

One of the enzymes disposing of $\rm H_2O_2$, i.e. catalase, has a very high specific reaction rate at substrate ($\rm H_2O_2$) saturation, i.e. $16\,000~\rm s^{-1}$ (Schomburg et al. (1994). In view of the complex kinetics of catalase, it is difficult to assess the rate at the 'normal' in vivo $\rm H_2O_2$ concentration of perhaps 0.1 mol m⁻³ $\rm H_2O_2$ but would be at most one-tenth of the cited $\rm H_2O_2$ -saturated rate or $1600~\rm s^{-1}$. In terms of the redox reactions of $\rm H_2O_2$ and of linear electron flow, this corresponds to an electron flux of 3200 mol electron

 mol^{-1} catalase s^{-1} , so that the minimum catalase content needed is 400/3200 or 0.125 mol catalase per mol non-cyclic redox chain; with 4 Fe per molecule this means 0.5 mol Fe per mol redox chain (Table 5).

Ascorbate peroxidase has a H₂O₂ (and ascorbate) saturated specific reaction rate of 323 mol H₂O₂ mol^{-1} enzyme s^{-1} , or 646 mol electron mol^{-1} enzyme s^{-1} (Schomburg et al. 1994). Because two electrons pass through the linear redox chain per electron passing through H₂O₂ into ascorbate peroxidase, the minimum ascorbate peroxidase needed is 400/1252 or 0.32 mol enzyme (0.32 mol Fe) per mol linear redox chain (Table 5). More enzyme is needed if the H₂O₂ level is less than the $0.15 \, \text{mol m}^{-3} \, \text{H}_2\text{O}_2$ needed to saturate the enzyme (Table 5). Data in Asada and Takahashi (1987) on the quantity of ascorbate peroxidase per unit chloroplast volume and data in Table 4.1 of Lawlor (1993) give values of 0.02-0.1 mol ascorbate peroxidase per mol linear redox chain, i.e. rather less than the value computed above. However, more recent data (Miyake et al. 1993) show that there is about 1 mol ascorbate peroxidase (half thylakoid-bound) per mol linear redox chain, i.e. in excess of what is required, and consistent with a lower steady-state concentration of H2O2 in chloroplasts than is needed to saturate the enzyme.

For glutathione peroxidase the reported specific reaction rate is 3100 mol H₂O₂ mol⁻¹ enzyme s⁻¹ at substrate (H₂O₂) saturation (Schomburg et al. 1994). Because the affinity for at least organic hydroperoxidases can be high (K_m 0.006 mol m⁻³) this specific rate may also apply to the reaction with in vivo concentrations of H₂O₂ in chloroplasts. An in vivo rate of 3100 mol H₂O₂ mol⁻¹ enzyme s⁻¹ means a rate of 6200 mol electron mol⁻¹ enzyme s⁻¹. As the stoichiometry of electron flow through glutathione peroxidase to that through the non-cyclic redox chain in the Mehler peroxidase reaction is, as for ascorbate peroxidase, 1:2, the minimum required ratio of ascorbate peroxidase to linear redox chains is 400/12,400 or 0.032 mol enzyme per mol redox chain. With 2 Se per molecule this corresponds to 0.064 mol Se per mol redox chain (Table 5).

These computations of the minimum requirement for trace metals in the Mehler catalase and Mehler peroxidase reactions are summarised in Table 5. To put these values on a whole thylakoid basis it is necessary to know what fraction of the electron flow through the linear redox chain can flow through the O_2 —consuming and H_2O_2 -consuming reactions of the Mehler catalase and Mehler peroxidase reactions.

Table 5. Computed (in text) ratio of the *minimum* content of enzyme (and their trace metals) dealing with active O species per idealized non-cyclic redox chain for the 'Mehler catalase' reaction and for the 'Mehler peroxidase' reaction using ascorbate or glutathione peroxidase

Reaction	Enzyme	Computed minimum (and observed) content per non-cyclic redox chain	Conputed minimum (and observed) trace metal content per non- cyclic redox chain
'Mehler catalase' in 'ancestral'	Fe superoxide dismutase	4.10^{-4}	4.10 ⁻⁴ (Fe)
Catalase	Catalase or		
	hydroperoxidase	0.125	0.5 (Fe)
'Mehler ascorbate peroxidase' in most	Fe superoxide dismutase	2.10^{-4}	2.10 ⁻⁴ (Fe)
cyanobacteria and algae other than Charophyceae and peridinin-containing Dinophyceae	Ascorbate peroxidase	0.32	0.32 (Fe)
'Mehler glutathione peroxidase' in some	Fe superoxide dismutase	2.10^{-4}	2.10 ⁻⁴ (Fe)
algae	glutathione peroxidase	0.032	0.064 (Se)
'Mehler ascorbate peroxidase' in Charophyceae;	Cu, Zn superoxide dismutase	2.10 ⁻⁴ (1.0)	2.10 ⁻⁴ (Cu) (1.0 (Cu)) 2.10 ⁻⁴ (Zn) (1.0 (Zn))
peridinin-containing Dinophyceae and higher plants	Ascorbate peroxidase	0.32 (0.5)	0.32 (Fe)

Where data are available (Miyake et al. 1993; Ogawa et al. 1995), the observed contents are given in brackets beside the computed values.

In some cyanobacteria and chlorophyte microalgae, the O2 uptake and evolution rate in the Mehler reaction in the induction phase upon illumination after a dark period is equal to the CO₂ fixation and O₂ evolution rates in subsequent steady-state photosynthesis, i.e. the electron flow through the reactions involving active oxygen species can equal the maximum capacity of linear electron flow to CO2 (Radmer and Kok 1976; Radmer, Kok and Ollinger, 1978; Radmer and Ollinger 1980; 1981; Kana 1990, 1992, 1993). Data from higher plants do not generally show such a high potential for the Mehler peroxidase reaction relative to linear electron flow to other acceptors (Badger 1985). Thus, in at least some algae, and cyanobacteria, the ratios cited in Table 5 can be taken as referring to the whole linear electron transport catalytic capacity, while in other organisms it could refer to a smaller fraction (Osmond and Grace 1995; Biehler and Fock

1996; Cheeseman et al. 1997) of the overall catalytic capacity of the linear redox chain.

With 3 Fe per PS II, 5 Fe per cyt b_6 -f, 12 Fe per PS II, 2 Fe per ferredoxin and 1 Fe per ascorbate per-oxidase, and a 1:1:1:1:1 ratio of these 5 catalysts, the Fe requirement of the Mehler peroxidase reaction is 23 Fe per electron transport chain. This ratio relates to a higher plant, where there is the additional requirement of 2 Cu (1 Cu for plastocyanin, 1 Cu for Cu–Zn superoxide dismutase, both present at 1 molecule per electron transport chain). In algae with cytochrome c_6 rather than plastocyanin, and Fe–SOD rather than Cu–Zn SOD, the Fe requirement is 25 Fe and 0 Cu per electron transport chain. The Fe requirement is reduced by 2 if flavodoxin replaces ferredoxin.

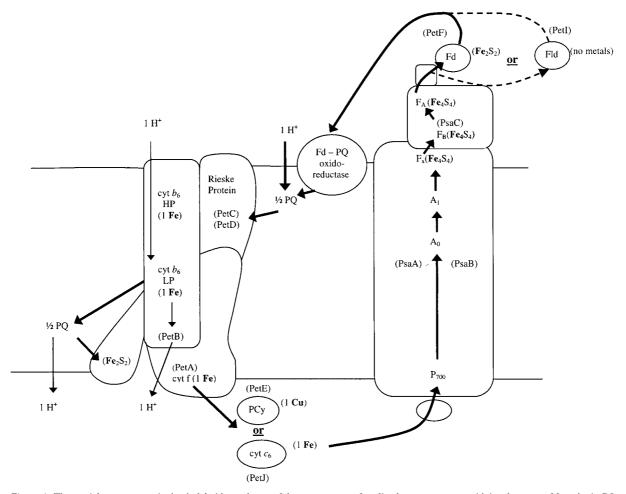


Figure 4. The spatial arrangement in the thylakoid membrane of the components of cyclic electron transport with involvement of ferredoxin-PQ oxidoreductase, showing involvement of trace metals.

Cyclic electron transport, chlororespiration and the role of NAD(P)H dehydrogenase

Cyclic electron transport around PS I (Figures 4 and 5), with associated H⁺ pumping and ADP phosphorylation, can be demonstrated *in vivo*, in chloroplasts *in vitro* and (with supplementation of redox catalysts) in isolated thylakoids (Raven 1976a,b; Herbert et al. 1990; Heber and Walker 1992; Bendall and Manusso 1995). The extent to which cyclic electron transport and photophosphorylation occur in parallel with the non-cyclic or Mehler peroxidase variants of linear electron transport in the same thylakoid membrane is unclear (Bendall and Manusso 1995). Also unclear is the maximum capacity for cyclic electron transport under optimal redox conditions and at light saturation, although the available data suggest a value

of 10% or less of the capacity for linear-chain electron transport except in specialized cells such as the N₂-fixing heterocysts of certain filamentous cyanobacteria and the bundle sheath cells of NADPme-C₄ flowering plants (Raven 1976a,b; Cha and Mauzerall 1992; Fork and Herbert 1993; Asada et al. 1993a; Bendall and Manusso 1995; Malkin and Canaani 1994).

Whatever the pathway(s) of cyclic electron transport around PS I in vivo there seems to be a role for A_0 , A_1 , F_X , F_B , F_A and ferredoxin or flavodoxin at the reducing end of the pathway, and for PQ, the cytochrome b_6 -f complex and cytochrome c_6 or plastocyanin at the oxidizing end of the pathway (Bendall and Manusso 1995; Miyake et al. 1995; Scheller 1996). Even this incomplete pathway, with no link between ferredoxin or flavodoxin and PQ, has a very substantial requirement for trace metals. Thus, with a 1:1:1:1 stoichiometry of

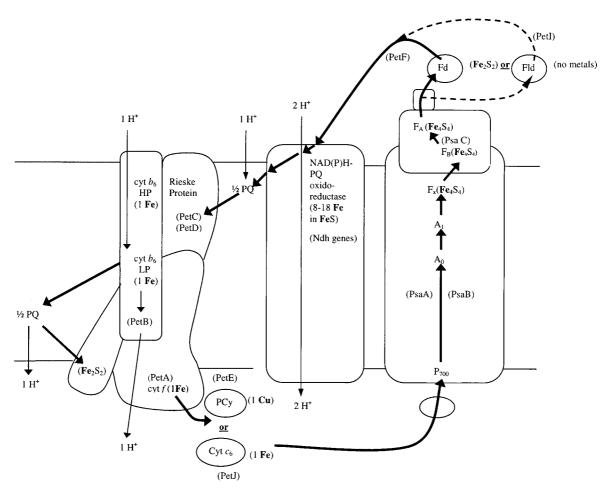


Figure 5. The spatial arrangement in the thylakoid membrane of the components of cyclic electron transport with involvement of NAD(P)H-PQ oxidoreductase, showing involvement of trace metals.

cytochrome b_6 -f, cytochrome c_6 , PS I and ferredoxin there is a requirement for 20 Fe, no Mn and no Cu for each cyclic electron transport chain, while with a 1:1:1:1:1 stoichiometry of cytochrome b_6 -f, plastocyanin, PS I and flavodoxin there are 17 Fe, no Mn and 1 Cu in each cyclic electron transport chain. These trace metal requirements apply to those pathways of cyclic electron transport that do not contain Fe for the ferredoxin (or flavodoxin) to PQ segment of cyclic electron transport, e.g. a 'direct' ferredoxin-PQ oxidoreductase (Bendall and Manusso 1995) or some sequence involving the psaE gene product (Yu et al. 1993; Zhao et al. 1993).

A significantly greater Fe requirement would follow from involvement of NAD(P)Hdh (NAD(P)H: PQ oxidoreductase) in cyclic electron flow (Mi et al. 1992a.b; Bendall and Manusso 1995; Scheller 1996; Endo et al. 1997a,b, 1998; Teicher and Scheller 1998). Depending on the assumed Fe content of the NAD(P)H-PQ oxidoreductase (8–18 Fe per complex: Siedow 1995; Brandt 1997), a cyclic electron pathway could have from 25 Fe (1:1:1:11:1 stoichiometry of NAD(P)H-PQ oxidoreductase (8 Fe):cytochrome b_6 -f complex: plastocyanin:PS I:flavodoxin) to 38 Fe (1:1:1:1:1 stoichiometry of NAD(P)H-PQ oxidoreductase (18 Fe), cytochrome b_6 -f complex:cytochrome c_6 : PS I:ferredoxin). Before discussing these data in the context of the iron requirement, and photon cost, of cyclic photophosphorylation, we shall briefly consider a second possible function of NAD(P)-PQ oxidoreductase, i.e. chlororespiration.

Chlororespiration (Bennoun 1982; Godde 1982; Bruce et al. 1983; Bennoun 1983; Caron et al. 1986; Klöck et al. 1989; Klöck et al. 1989; Scherer 1990; Wilhelm and Duval 1990; Avelange and Rebeille 1991; Peltier and Schmidt 1991; Walker 1992; Groom et al. 1993; Ting and Owens, 1993; Bennoun 1994; Schmetterer 1994; Buchel and Garab 1995; Endo and Asada 1996; Casper-Lindley and Björkman 1997; Buchel et al. 1998; Feild et al. 1998) involves the oxidation of carbohydrate within chloroplasts in darkness or in low light, ultimately using O₂ as the terminal electron acceptor, and appears to function in generating ATP in the dark and in low light.

The pathway of chlororespiration is unclear. Studies on mutants of Chlamydomonas reinhardtii (Bennoun 1993) show that chlororespiration does not obligatorily involve the Rieske Fe-S protein or the cytochrome f components of the cytochrome b_6 -fcomplex, plastocyanin, or the CF₀-CF₁ ATP synthetase (see also Feild et al. 1998, for work on higher plants). Belkhodja et al. (1998) found more expression of chlororespiratory (dark PQ reduction) in Fe-deficient leaves of higher plants than of controls, a result which would not be predicted from the involvement of the Fe-rich NAD(P)H-PQ oxidoredustase or cytochrome b_6 -f complex. Evidence on the nature of the terminal oxidases, and hence of their trace metal content, does not yield a clear picture (Bennoun 1982; Buchel and Garab 1995; Goyal and Tolbert 1996; Wiekkowski and Bojko 1997; Feild et al. 1998). Indeed, Rich et al. (1998; see also Bennoun 1994), having taken great care to minimize contamination from other organelles, conclude that higher plant chloroplasts lack NAD(P)H-PQ oxidoreductase and plastoquinol oxidase, and that they lack cytochrome c or alternative oxidase.

Requirement for cyclic photophosphorylation

The requirement for cyclic photophosphorylation is apparently variable among different photosynthetic cells; on a unit chromophore basis the requirement is probably highest in heterocysts of cyanobacteria and bundle sheath cells of C₄ NADPme plants (see Bendall and Manusso 1995). In other photosynthetic cells the *in vivo* capacity for cyclic photophosphorylation is apparently lower, i.e. only about one-tenth that of linear photophosphorylation (Raven 1976a). Although this restricted capacity may relate to the capacity of

ATP-consuming reactions in vivo under conditions that isolate cyclic photophosphorylation as the ATP source, it is worth considering possible Fe-economies of limiting the capacity of the ferredoxin/NAD(P)H to PQ (i.e. cyclic-specific and chlororespiration-specific) part of the pathway by restricting the quantity of the catalysts of this redox pathway. The in vivo specific reaction rate of mitochondrial NADH-UQ oxidoreductase is high relative to that of subsequent redox complexes in the mitochondrial electron transport chain (Siedow 1995; Whitehouse and Moore 1995) and bearing in mind the low demand for the products of NAD(P)H-PQ oxidoreductase in cyclic redox reactions, considerable Fe economies could occur if the NAD(P)H-PQ oxidoreductase content was very low. However, there must be a lower limit on the number of NAD(P)H-PQ oxidoreductase complexes per unit light-harvesting pigment if the high photon yield of cyclic photophosphorylation in vivo (Raven 1976a) is to be explained. This is because there must be limits on excitation energy migration from the totality of pigments associated with PS I reaction centres to the few PS I reaction centres associated (albeit via the membrane-dissociable ferredoxin or flavodoxin) with the restricted number of NAD(P)H-PQ oxidoreductase complexes, which then interact (via the mobile PQH_2/PQ pool) with the cytochrome b_6 -f complex and thence (via membrane-dissociable cytochrome c_6 or plastocyanin) to the oxidizing end of the same few PS I reaction centres. Of course, similar quantitative arguments (complexes per unit chlorophyll) apply to any non-Fe-requiring means of electron transfer from ferredoxin or flavodoxin to PQ, but without the implications for Fe cost.

Now that we have considered the two putative client reactions for a functional NAD(P)H-PQ oxidoreductase, we can consider the evidence that this Fe-costly redox complex is involved in these reactions, and the consequences of its involvement relative to other, relatively uncharacterized, redox reactions.

The occurrence of NAD(P)H-PQ oxidoreductase genes

The genes encoding this complex are found in the genome of cyanobacteria (Kaneko et al. 1996) and (with some exceptions) in the plastid genome of 'green plants', i.e. the higher plants or embryophytes, and the green algae *sensu lato* including not only the Chlorophyta but also the euglenoids and chlororachniophytes, which obtained plastids by (second-

ary) endosymbiosis of unicellular members of the green algae *sensu stricto* (Bhattacharya and Medlin 1998). Organisms within the green plants that lack the plastid-located NAD(P)H-PQ (*ndh*) genes include some flowering plant parasites, some *Pinus* spp., two *Chlamydomonas* spp. and *Euglena gracilis* (de Pamphilis and Palmer 1990; Haberhauser and Zetschke 1994; Wakagusi et al. 1994).

However, there are no known occurrences of *ndh* genes in the plastid genome of red algae, nor of those algae (e.g. diatoms, brown algae, cryptophytes, haptophytes) which obtained their plastids by (secondary) endosymbiosis of unicellular members of the red algae (Reith 1995). Thus, any plastid-expressed NAD(P)H-PQ oxidoreductases in these organisms lacking *ndh* genes in their plastid genome must have these genes for this redox complex in their nuclei; although this is not impossible, there is apparently no evidence that it occurs (Reith 1995; Martin et al. 1998).

In summary, plastid-located *ndh* genes are found in many but not all of the O₂-evolving eukaryotes tested. Although a nuclear location of these genes in organisms lacking the plastidial genes is possible, there is no apparent evidence that this is the case, despite the occurrence of cyclic electron transport and photophosphorylation (Raven 1976a,b, 1984; Cha and Mauzerall 1992; Fork and Herbert 1993; Malkin and Canaani 1994; Bendall and Manusso 1995) and chlororespiration (Caron et al. 1986; Büchel and Garab 1995) in organisms both with and without known NAD(P)H-PQ oxidoreductase genes.

The expression of NAD(P)H-PQ oxidoreductase genes

The NAD(P)H-PQ oxidoreductase genes are expressed in cyanobacteria where they function in normal dark respiration as well, apparently, as cyclic electron flow and photophosphorylation (see below, and Scherer 1990; Mi et al. 1992a, b, 1995; Schmetterer 1994; Tanaki et al. 1997). In eukaryotes it is clear that steps must be taken to prevent interference from the mitochondrially expressed NADH-UQ oxidoreductase in the immunological or functional assay of plastidial NAD(P)H-PQ oxidoreductase (Rich et al. 1998; Teicher and Scheller 1998). Thus, while there is evidence for expression of NAD(P)H-PQ oxidoreductase in thylakoids of green algae sensu stricto (Godde 1982) and of higher plants (Berger et al. 1993; Guedeney et al. 1996; Kubicki et al. 1996; Endo et al. 1997a,b; Wiekkowski and Bojko 1997; Burrows et al.

1998; Sazanov et al. 1998a,b; Shikanai et al. 1998a; Teicher and Scheller 1998), it is important to bear in mind the need for rigorous exclusion of mitochondrial contamination.

Quantitation of expression has been less frequently attempted. There are semi-quantitative indications that, as expected if there is a predominant role of cyclic electron flow in bundle sheath cells of NADPme-C₄ plants and NAD(P)H-PQ oxidoreductase is involved in this cyclic electron flow, there is greater expression of NAD(P)H-PQ oxidoreductase in bundle sheath chloroplasts than in mesophyll chloroplasts of the NADPme-C₄ plant Sorghum bicolor (Kubicki et al. 1996). More quantitative estimates include those of Sazanov et al. (1998a) who found one NAD(P)H-PQ oxidoreductase complex per 100 (linear) 'electron transport chain' in the C₃ Pisum sativum chloroplasts, and Burrows et al. (1998) who found one NAD(P)H-PQ oxidoreductase per 70 PS II in the C₃ Nicotiana tabaccum. As there are 22 Fe per linear electron transport chain with 3 Fe in PS II, 5 Fe in the cytochrome b_6 -f complex, 12 Fe in PS I and 2 Fe in ferredoxin, and 8-18 Fe per NAD(P)H-PQ oxidoreductase (see above), there is 86-275 times as much Fe in the components of linear electron transport as in NAD(P)H-PQ oxidoreductase.

The low level of expression of NAD(P)H-PQ oxidoreductase relative to that of the components of the linear electron transport pathway in eukaryotes imposes constraints on the capacity of electron flow through the NAD(P)H-PQ oxidoreductase relative to that of linear electron transport. One feature that may help to offset the low level of expression of NAD(P)H-PQ oxidoreductase is that it might, by analogy with the respiratory NAD(P)H-UQ oxidoreductase have a high specific reaction rate relative to other photosynthetic redox agents; NAD(P)H-UQ oxidoreductase has a higher specific reaction rate than do subsequent respiratory redox catalysts (Whitehouse and Moore 1995). This permits the same steady state electron flux during NADH oxidation by O2 to occur through the NAD(P)H-UQ oxidoreductase as occurs through the more abundant cytochrome b- c_1 complex and the even more abundant cytochrome oxidase (Siedow 1995; Whitehouse and Moore 1995).

However, the inequality of abundance is much greater in the case of the plastid NAD(P)H-PQ oxidoreductase. Even if the maximum electron transport rate through the NAD(P)H-PQ oxidoreductase could approach a value that is 70 (*Nicotiana*) – 100 (*Pisum*) times that through linear electron trans-

port, there might be constraints on the rate at which electrons could be delivered from the NAD(P)H-PQ oxidoreductase to the cytochrome b_6 -f complex viaPQ/PQH₂, despite the known lateral transport properties of PQ/PQH₂ in the thylakoid membrane, from PS II in appressed to cytochrome b_6 -f and PS I in non-appressed regions of eukaryotic green plant thylakoids. In fact NAD(P)H-PQ oxidoreductase occurs with PS I and cytochrome b_6 -f in non-appressed thylakoids of Nicotiana, thus diminishing the extent of transport of PQ/PQH2 that would be needed. Nevertheless, the low level of expression and the likelihood that the NAD(P)H-PQ oxidoreductase can only catalyze a specific rate of electron flow which is a few-fold greater than the specific flux through the linear electron transport chain means that the level of expression of NAD(P)H-PQ oxidoreductase would only permit a light-saturated electron flux through the NAD(P)H-PQ oxidoreductase at 5-10% of the linear electron flux. Although the flux-limiting steps in chlororespiration are not known, it is of interest that the chlororespiratory rate in Helianthus annus is only 1/300 that of linear electron transport at light saturation (Feild et al.

Spatial constraints may apply to the light-limited rate of electron transfer through NAD(P)H-PQ oxidoreductase if it is functional in cyclic electron transport. Here the argument involves the number of PS I complexes with which each of the rare NAD(P)H-PQ oxidoreductase complexes can inteact. If this number is less than the overall ratio of PS I to NAD(P)H-PQ oxidoreductase, then a maximal photon yield of cyclic electron transport demands that excitation energy is transferred efficiently from PS I units that are not in redox contact with NAD(P)H-PQ oxidoreductase to those that are in such contact. Here, the argument does not concern the maximum specific reaction rate of NAD(P)H-PQ oxidoreductase; these quantum yield considerations are all within the ceiling imposed by this maximum specific reaction rate. We shall return to this photon yield question again under 'Fe requirements for cyclic, and Mahler reaction electron transport and H⁺ pumping: Fe and photon costs' section below.

Evidence for functioning of NAD(P)H-PQ oxidoreductase in cyclic electron flow and chlororespiration

A powerful tool in assessing the role of NAD(P)H-PQ oxidoreductase in cyclic electron flow and chlororespiration, and in metabolism more generally, is to inactivate one or more of the *ndh* genes that encode parts of the NAD(P)H-PQ oxidoreductase complex. Generally, deletions or inactivations have no phenotypic effect under optimal growth conditions.

The absence of a phenotypic effect of NAD(P)H-PQ oxidoreductase deletion under optimal growth conditions has been reported for cyanobacteria (Schluchler et al. 1993; Yu et al. 1993; Zhao et al. 1993; Howitt et al. 1996; Klugehammer et al. 1997; Sültemeyer et al. 1997; Ohkawa et al. 1998) and for C₃ flowering plants (Rochaix 1997; Burrows et al. 1998; Sazanov et al. 1998a,b; Shikanai et al. 1998a; Nixon and Malinga 1999; Roldán 1999). These results indicate that if any NAD(P)H-PQ oxidoreductase-requiring process, such as cyclic electron flow and photophosphorylation, and chlororespiration, are needed for growth under optimal conditions then there must be a bypass (or bypasses) of the step involving NAD(P)H-PQ oxidoreductase, which permits the function to be fulfilled to an extent that does not interfere with normal growth and development.

However, phenotypic effects of preventing NAD(P)H expression have been found under suboptimal growth conditions in cyanobacteria. Synechocystis sp. strain PCC 7002 had a reduced rate of photolithotrophic growth at low (but not high) photon flux density, and was unable to grow photoorganotrophically on glycerol when PS II activity was inhibited by DCMU (Schluchler et al. 1993). These two effects can be attributed to decreases in cyclic photophosphorylation and/or chlororespiration, and cyclic photophosphorylation, respectively. These effects were mimicked when the psaE gene product (a component of PS I with no direct redox capabilities) was absent, either on its own or in combination with a lack of ndhF (NAD(P)H-PQ oxidoreductase (Yu et al. 1993; Zhao et al. 1993).

The other set of sub-optimal growth conditions in which the absence of NAD(P)H-PQ oxidoreductase manifests itself is in response to low inorganic C supply, i.e. when CO₂ concentrating mechanisms (CCMs) are maximally expressed (Klugehammer et al. 1997; Sültemeyer et al. 1997; Ogawa et al. 1985; Ogawa 1991; Ohkawa et al. 1998). Here, it seems to be the

HCO₃⁻-transporting rather than the CO₂-transporting variant of CCM seems to be affected by the absence of NAD(P)H-PQ oxidoreductase (Sültemeyer et al. 1997). Although these findings seem to be consistent with the operation of cyclic electron flow via NAD(P)H-PQ oxidoreductase in this light-dependent HCO₃⁻ transport, evidence from P₇₀₀ redox kinetics indicates that NAD(P)H-PQ oxidoreductase is *not* essential for cyclic electron flow in *Synechocystis* sp. strain 7002 (Sültemeyer et al. 1997). Similar results were found for mutants lacking *psaE* as for those lacking *ndhF* (i.e. lacking NAD(P)H-PQ oxidoreductase) (Sültemeyer et al. 1997), a somewhat different conclusion to that arrived at for the same strain by Schluchler et al. (1993), Yu et al. (1993) and Zhao et al. (1993).

No such direct effect on phenotype has been investigated for higher plants lacking NAD(P)H-PQ oxidoreductases under sub-optimal conditions. However, Sazanov et al. (1998b) showed a reduced rate of PQ photoreduction in heat-stressed *Nicotiana tabacum* lacking NAD(P)H-PQ oxidoreductase relative to wild type.

The absence of plastid-encoded NAD(P)H-PQ oxidoreductase is the norm for a number of green eukaryotic algae and higher plants and for all other eukaryotic O₂-evolvers, despite the evidence for cyclic electron transport and for chlororespiration in these organisms (see 'Cyclic electron transport, chlororespiration and the role of NAD(P)H dehydrogenase' and 'The occurrence of NAD(P)H-PQ oxidoreductase gene' sections above).

Other evidence relating to the function of NAD(P)H-PQ oxidoreductase in cyclic electron transport and in chlororespiration concerns the sensitivity of these processes to inhibitors that are allegedly specific to NAD(P)H-PQ (or UQ) oxidoreductases such as rotenone and amytal, and the phenotypic oxidoreductase nucleotide specificity of the complex (Bendall and Manusso 1995; Miyake et al. 1995; Scheller 1996; Darrouzet et al. 1998). The inhibitor evidence is somewhat ambivalent. As to nucleotide specificity, many reports indicate complete, or substantial, selectivity for NADH rather than NADPH. This permits a role in chlororespiratory use of glycolytic NADH, but not of NADPH from the oxidative pentose phosphate pathway or from PS I unless a transhydrogenase is present; the occurrence of such an enzyme in plastids and cyanobacteria is equivocal at best (Schluchler et al. 1993; Bendall and Manusso 1995). However, NAD(P)H-PQ oxidoreductase from Synechocystis PCC 6803 was NADPH-specific and had ferredoxin-NADP+ reductase activity (Matsuo et al. 1998). Similar results were found for Teicher and Scheller (1998) for the *Hordeum vulgare* and by Funk et al. (1999) for the *Zea mays* thylakoid enzymes.

The conclusion is that the NAD(P)H-PQ oxidore-ductase may well be a component of a cyclic electron transport pathway, but not necessarily the only pathway of cyclic electron transport in a given organism. Of the alternative pathways from ferredoxin to PQ a commonly suggested, but poorly characterized, possibility is a ferredoxin-PQ oxidoreductase (Bendall and Manusso 1995); this possible metal (Fe?) involvement is unclear. It is also not known if all variants of chlororespiration involve NAD(P)H-PQ oxidoreductase.

Fe requirements for cyclic, and Mehler reaction electron transport and H⁺ pumping: Fe and photon costs

We have seen in 'Cyclic electron transport, chlororespiration and the role of NAD(P)H dehydrogenase' section above that the Fe content per 'idealized' cyclic electron transport chain containing NAD(P)H-PQ oxidoreductase is 27-37 Fe for the case of plastocyanin rather than cytochrome c_6 , and ferredoxin rather than flavodoxin, depending on the assumptions about the Fe content of NAD(P)H-PQ oxidoreductase, while in the absence of NAD(P)H-PQ oxidoreductase the value is 19 Fe. The Fe cost of ATP synthesis by this pathway depends on the H⁺/e⁻ ratio and the H⁺/ATP ratio that are assumed. The H⁺/e⁻ ratio of cyclic electron transport is generally thought to be 2. This accords with the 'Q-cycle' operation of the cytochrome b_6 -f complex, but has no role for H⁺ pumping by the NAD(P)H-PQ oxidoreductase which, by analogy with the mitochondrial NADH-UQ oxidoreductase, would have a H⁺/e⁻ ratio of 2 (Siedow 1995; Whitehouse and Moore 1995; Brandt 1997). An H⁺/e⁻ ratio of 4 (2 from NAD(P)H-PQ oxidoreductase, 2 from cytochrome b_6 -f complex) is consistent with early data summarized by Heath (1972) and more recent work of Heber et al. (1995a). With an H⁺/ATP ratio of 4 (Falkowski and Raven 1997), an H⁺/e⁻ ratio of 4 would give, with 1 electron cycled per photon contributing excitation energy to PS I, one ATP per photon absorbed by pigments contributing to light harvesting by PS I reaction centres effective in cyclic electron flow (see 'The expression of NAD(P)H-PQ oxidoreductase genes' section) above. In vivo estimates from the 1970s of the photon yield of cyclic photophosphorylation also gave val-

ues of 1 ATP per photon absorbed by PS I (Raven 1976a,b), although at that time the interpretation was in terms of an H⁺/e⁻ ratio of 2 and an H⁺/ATP ratio of 2. The absence of the NAD(P)H-PQ oxidoreductase H⁺ pump (or its equivalent) would, on current views of the H⁺/ATP, give only 0.5 ATP per photon absorbed by pigments related to PS I reaction centres effective in cyclic electron flow. These computed photon yields of ATP production by cyclic photophosphorylation may be compared with the values for the Mehler peroxidase reaction where 2 photon are absorbed and used by PS II and PS I to move 1 electron from H₂O to O_2 and pump 3 H⁺ and hence produce 0.75 ATP. The net result is that 0.75 ATP is made with 2 photons or 3/8 (0.375) ATP per photon absorbed. This value is lower than the *in vivo* estimates of 1 ATP per photon from the 1970s for cyclic photophosphorylation based on total photons absorbed by PS I using irradiation at 700 nm (see above). However, it is closer to values predicted from current perceptions of the H⁺/ATP ratio (4) and an H⁺/e⁻ ratio of 2 with (potentially) all PS I units participating and an H⁺/e⁻ ratio of 4. Some PS I units are presumably not involved in cyclic electron transport involving NAD(P)H-PQ oxidoreductase due to lack of connection to a (scarce) NAD(P)H-PQ oxidoreductase (see 'The expression of NAD(P)H-PQ oxidoreductase genes' section above). The predicted photon yield of cyclic photophosphorylation would become even more comparable to that of ATP supply from the Mehler peroxidase reaction (0.375 ATP per photon absorbed; see below) if 400-700 nm photons absorbed by pigments that cannot transfer excitation energy to PS I reaction centres, are taken into account (see Raven 1976a,b).

The Fe cost of ATP synthesis by the cyclic photophosphorylation, with or without involvement of NAD(P)H-PQ oxidoreductase can be approached as follows. If we assume a maximum in vivo specific reaction rate of PS I of 400 mol electron per mol PS I per second in linear electron flow (based on the compilation in Raven 1990), and a capacity for cyclic electron flow of 10% of the linear rate, i.e. 40 mol electron per mol PS I per second, the maximum rate of electron transport with NAD(P)H-PQ oxidoreductase operative is then 1.08–1.48 mol electron per mol Fe in the cyclic redox pathway per second for the range of Fe contents cited in 'The occurrence of NAD(P)H-PQ oxidoreductose genes' section above. This calculation is for a pathway using plastocyanin rather than cytochrome c_6 and ferredoxin rather than flavodoxin. This means that, with an ATP/e⁻ ratio of 1, then 1.08-1.48 mol ATP produced per mol Fe in the cyclic redox pathway per second (Table 6). For a cyclic pathway lacking NAD(P)H-PQ oxidoreductase, the Fe content for the plastocyanin/ferredoxin alternative is 19 Fe per cyclic redox chain, and the ATP/e⁻ ratio is 0.5. The rate of cyclic electron transport per Fe is then 2.1 mol electron per mol Fe per second and the rate of ATP synthesis is 1.05 mol ATP per mol Fe per second (Table 6).

The computations of Fe costs show that, even with the highest assumed Fe content of the NAD(P)H-PQ oxidoreductase, the rate of ATP synthesis is, for a given electron flow rate, very similar on a mol Fe basis for the NAD(P)H-PQ oxidoreductase-containing redox sequence as for those that lack that complex and catalyse electron flow from ferredoxin (or flavodoxin) to PQ with a redox system that lacks Fe (or any other metal) and that does not pump H⁺. In the context of limiting factors other than Fe in the natural environment the NAD(P)H-PQ oxidoreductase-containing cyclic redox pathway can have a photon yield of ATP production which is twice that of a cyclic redox pathway lacking NAD(P)H-PQ oxidoreductase (see Raven 1988a, 1990 and Sunda and Huntsman 1997 for consideration of Fe and light colimitation).

We now compare the mol ATP produced per mol Fe in the redox chain per second for pseudocyclic photophosphorylation (Mehler peroxidase reaction) with cyclic photophosphorylation with and without NAD(P)H-PQ oxidoreductase, as the source of ATP without a net reduction of oxidant. The Mehler peroxidase pathway ('Linear electron flow with O2 as electron acceptor: the Mahler peroxidase reaction' section above) with plastocyanin rather than cytochrome c_6 and ferredoxin rather than flavodoxin has a minimum Fe, Cu and Zn content of 23 Fe, 1 Cu and 1 Zn (including the Fe, Cu and Zn needed for enzymes of O₂. and H₂O₂ metabolism) (Table 5) for a linear chain with all complexes at equal concentration. With 3H⁺/e⁻ in the Mehler peroxidase reaction and 4 H⁺/ATP, linear electron flow of 40 mol electron per mol complex per second (the same as that assumed for cyclic electron flow) means that 1.36 ATP synthesized per mol Fe in the redox chain per second (Table 6). This is within the range of rates computed for cyclic electron flow using NAD(P)H-PQ oxidoreductase with the highest and lowest likely Fe contents of the NAD(P)H-PQ oxidoreductase complex, and higher than the rate for a cyclic redox sequence lacking NAD(P)H-PQ oxidoreductase (Table 6). The photon cost of ATP synthesis in the Mehler peroxidase reaction, assuming equal photon allocation to PS I and PS II, is 0.375 ATP

Table 6. Computed (in text) Fe costs of ATP synthesis in cyclic and O_2 -linked linear (Mehler peroxidase) electron transport pathway with electron flow at 40 mol electron per mol complex per second

Reaction sequence	Mol ATP per mol Fe in redox sequence per second
Cyclic electron flow with NAD(P)H-PQ oxidoreductase, assuming 18 Fe per NAD(P)H-PQ oxidoreductase complex	1.08
Cyclic electron flow with NAD(P)H-PQ oxidoreductase, assuming 8 Fe per with NAD (P)H-PQ oxidoreductase complex	1.4
Cyclic electron flow without NAD(P)H-PQ oxidoreductase, i.e. assuming a non Fecontaining, non H ⁺ -pumping 'ferredoxin-PQ oxidoreductase'	1.05
O ₂ -linked linear (Mehler peroxidase) electron flow, including the observed Fe content of the ascorbate peroxidase involved in this reaction sequence	1.30

All pathways assumed to have equal quantities of all redox complexes, and to use plastocyanin rather than cytochrome c_6 and ferredoxin rather than flavodoxin. This 'plastocyanin' variant of the pathways is most commonly associated with the non Fe-containing Cu–Zn superoxide dismutase in the Mehler peroxidase reaction

per photon absorbed (2 photons absorbed per electron transported; 3 H⁺ per electron transported; 4 H⁺ per ATP: Falkowski and Raven 1997).

Three further considerations must be borne in mind when comparing the Fe requirements for ATP synthesis by cyclic electron flow and by the Mehler peroxidase reaction. One is that components of the Mehler peroxidase reaction have the additional role of disposing of active oxygen species other than those generated by PS I (Noctor and Foyer 1998). A second consideration is that the cyclic electron transport may be more effective in acidifying the thylakoid lumen in high light, thereby increasing non-photochemical quenching of excitation energy, than is the case for the Mehler peroxidase electron transport (Heber and Walker 1992; cf. Osmond and Grace 1995). A further role for coupled cyclic electron around PS I is in the regulation of PS II (Heber et al. 1995b). The final consideration is that the NAD(P)H:PQ oxidoreductase may be essential for chlororespiration as well as cyclic electron transport.

Genetic and environmentally driven variations in the ratio of trace metal-containing catalysts in the thylakoid membrane

Background

It is now clear that the major trace metal-containing complexes in the thylakoid membrane are not generally present in equal quantities in a given genotype grown under a given set of conditions, that these ratios are susceptible to environmental (growth condition) influences, and that there are substantial genetic (large-scale taxonomic or phylogenetic) variations in the range of ratios of these components.

Genetically determined variations in trace metal-containing catalysts

Most data are available for variations in the ratio of PS II to PS I as a function of genetic and environmental factors, with rather fewer data on the ratio of cytochrome b_6 -f complex, and even fewer data on the ratios plastocyanin/cytochrome c_6 and on ferredoxin/flavodoxin to the photosystems. The

best established genetic variation in content of metalcontaining catalysts is for the ratio of PS II to PS I. The highest ratio of PS II to PS I is found in cyanobacteria sensu stricto (i.e. excluding chloroxybacteria) and Rhodophyta (Cunningham et al. 1989; Malkin et al. 1990; Rögner et al. 1990; Murakami et al. 1997; Falkowski et al. 1998 cf. Greene and Gerard 1990; Kübler and Davison 1993). The ratio in these organisms is almost always less than 1, is commonly 1:3-1:5, is sometimes as low as 1:9, or is even 1:25 for the non-heterocystous N2-fixing marine cyanobacterium Trichodesmium spp. Higher ratios of PS II to PS I, usually more than 1.0 and as high as 4.0, are found for the chloroxybacteria, Chlorophyta and their Embryophyta (higher plant) derivatives and the Euglenophyta as well as for the Heterokontophyta (diatoms, brown algae), Dinophyta and Haptophyta; (Falkowski et al. 1981; Dubinsky et al. 1986; Falkowski et al. 1986; Friedmann and Alberte 1986; Chow and Anderson 1987; Smith and Melis 1987; Sukenik et al. 1987; Guenther et al. 1988; Raven 1988a; Falkowski et al. 1989; Lee and Whitmarsh 1989; Wilhelm et al. 1989; Burger-Wiersma and Matthijs 1990; Chow et al. 1990; Raven 1990; Bullerjahn and Post 1993; Trissl and Wilhelm 1993; Iglesias-Prieto and Trenck 1994; Partensky et al. 1997; cf. Caron et al. 1988). No data were found for the Cryptophyta or Chlorarachniophyta. These variations in the ratio of two major trace metal-containing complexes mean that, on a total chromophore basis, the Fe content in the two photosystems is greatest for cyanobacteria and Rhodophyta with a high ratio of the (high Fe) PS I to the (low-Fe) PS II, while the Fe content of the two photosystems is smallest for Heterokontophyta and such Chlorophyta as Mantoniella (Wilhelm et al. 1989) with a high ratio of PS II to PS I. By contrast, the Mn (in PS II only) content per unit total chromophore may be lower in cyanobacteria and Rhodophyta which have a lower PS II:PS I ratio than in organisms with higher PS II:PS I ratios.

The variations in PS II:PS I given above apparently represent genetically determined differences among major taxa (Divisions). Within these higher taxa there are genetically determined differences relating to the habitat in which the organisms normally grow. Thus, while the typical range of PS II/PS I ratios in sunadapted terrestrial higher plants (1.7–1.8) overlaps with that of shade-adapted terrestrial higher plants (1.19–1.88) (Chow et al. 1990a), the lowest value for the shade-adapted species is lower than that for the sun-adapted species. This mimics the acclimat-

ory effect seen when sun-adapted plants are grown at low PFDs, e.g. PS II/PS I in *Pisum sativum* is 1.79 when grown at high PFD and is 1.27 when grown at low PFD (Chow et al. 1990a). We shall see later ('Environmental (acclimatory) variations in trace metal-containing catalysts' section below) that this acclimatory effect of low growth PFD in decreasing PS II/PS I is not universal, so that in some eukaryotic algae growth PFD does not alter PS II/PS I, while in others PS II/PS I increases as the growth of PFD is decreased.

Environmental (acclimatory) variations in trace metal-containing catalysts

The PS II:PS I ratio, and the PS II:total chromophore and PS I:total chromophore ratios, vary with the photon flux density for growth and the nitrogen and iron supply for growth. Fe deficiency lowers the thylakoid protein content, and the activation state (but not the quantity) of Rubisco (Evans 1996). Berges et al. (1996) showed that N deficiency increased the PS II:PS I ratio in the marine green alga *Dunaliella tertiolecta* from 1.0 in N-replete cells to 2.4 in N-starved cells, and in the marine diatom *Thalassiosira weissflogii*, from 0.45 in N-replete to 1.4 in N-starved cells.

More relevant to in vivo functioning in terms of the effectiveness with which organisms use the potentially limiting resources Fe and Mn, are the rates of CO2 fixation per unit Fe or Mn at light and CO₂ saturation. The PS II:PS I ratio differences, and more importantly the PS I and PS II content per unit chromophore, combined with data on photosynthetic rate per unit chromophore, indicate that the cyanobacteria and red algae should have the most Fe-costly CO₂ fixation (mol Fe needed to yield a fixation rate of one mol CO₂ per second), while diatoms, brown algae and the green alga Mantoniella should be the least Fe-costly (Raven 1988a, 1990; Wilhelm et al. 1989). This analysis ignores the Fe used in the cytochrome b_6 -f complex, cytochrome c_6 (if present), ferredoxin (if present), thylakoid NAD(P)H-PQ oxidoreductase (Endo et al. 1997a,b; Mi et al. 1997; Matsuo et al. 1997), thylakoid Fe-superoxide dismutase (Asada et al. 1980; Asada 1992; Asada et al. 1993b) and thylakoid ascorbate peroxidase (Asada 1992; Asada et al. 1993b). The Fe content on a unit light-harvesting chromophore basis in all of these thylakoid-associated components, except the cytochrome b_6 -f complex, cytochrome c_6 and ferredoxin, is poorly understood. Although the

Fe content per molecule of Fe superoxide dismutase and per molecule of ascorbate peroxidase is known, the number of thylakoid-associated molecules of these enzymes on a total chromophore basis is rather poorly characterized (Asada et al. 1980; Hewitt 1983; Asada 1992, 1993b). Price et al. (1998) have elegantly shown the lower limit of expression of the cytochrome b_6 -f complex compatible with the wild-type phenotype in *Nicotiana*.

Variations in the light supply for growth can cause acclimation of the PS I and PS II content of photosynthetic cells, and hence in the computed Fe cost of photosynthesis in terms of the high Fe PS I. In some cases, e.g. the marine green microalga Dunaliella tertiolecta, there is no significant effect of PFD for growth on the PS II:PS I ratio, although there is an almost 3-fold increase in the reaction centre content per cell in cells grown at low PFDs and a rather larger proportional increase in chlorophylls a and b per cell (Falkowski et al. 1981). A very different acclimation is apparent in the marine diatom Skeletonema costatum, where the PS II:PS I is more than twice as high for cells grown at the lowest PFD tested as in those grown at the highest PFD. There is an approximately 1.5-fold increase in PS II and in chlorophylls a and c per cell, and an approximately 1.5-fold decrease in PS I per cell (Falkowski et al. 1981). Neither alga shows a large change in light-saturated photosynthetic rate on per cell basis as a function of PFD for growth (Falkowski et al. 1981), so that the light-saturated photosynthesis rate per PS I decreased from 136 (high light grown) to 45 (low light grown) electrons per PS I per second in Dunaliella tertiolecta, and increased from 231 (high light grown) to 285 (low light grown) electrons per PS I per second in Skeletonema costatum.

Photosynthesis at low PFDs for these two algae grown under similar conditions has been measured by Falkowski and Owens (1980). At low PFDs for measurement the photosynthetic rate on a chlorophyll basis for *Dunaliella tertiolecta* is independent of the PFD for growth, so that the photosynthetic rate per PS I measured at 20 μ mol photon m⁻² s⁻¹ is about 5 electrons per PS I per second regardless of the PFD for growth (Falkowski and Owens 1980; Falkowski et al. 1981). *Skeletonema costatum* cells grown at low PFDs have a lower chlorophyll-based O₂ evolution rate at low PFDs for measurement. Photosynthetic rates measured at a PFD of 20 μ mol photon m⁻² s⁻¹ are 135 electrons per PS I per second for high light cells, and 100–111 electrons per PS I per second for

cells grown at low PFDs (Falkowski and Owens 1980; Falkowski et al. 1981).

These computations show that, although low-light-acclimated *Skeletonema* cells have a lower photosynthetic rate per PS I than cells acclimated to high light, the rates are much higher than in *Dunaliella* cells where there is little or no acclimation of light-limited rates of photosynthesis per PS I as a PFD for growth. The computed electron transfer rates per PS I can be directly translated to electron transfer rates per atom of Fe in PS I by division of 12.

The increase in PS II:PS I with acclimation to lower PFDs in *Skeletonema costatum* cannot be generalised to all diatoms. Although the PS II:PS I ratio is higher in *Phaeodactylum tricornutum* (Friedmann and Alberte 1986) and *Cylindrotheca fusiformis* (Smith and Melis 1988) acclimated to lower PFDs, *Thalassiosira weisflogii* has an essentially constant PS II:PS I regardless of the PFD for growth (Dubinsky et al. 1986). However, as in *Skeletonema costatum*, the PS I (and PS II) per unit chlorophyll decrease with lower PFDs for growth, as is also the case for the haptophyte *Isochrysis galbana* and the dinoflagellate *Prorocentrum micans* (Dubinsky et al. 1986).

For another heterokont, the giant kelp *Macrocystis* pyrifera (Smith and Melis 1987), photosynthetic laminae (blades) growing at the surface had a PS II:PS I ratio of 1.79, while those growing at 20 m depth had a PS II:PS I ratio of 2.22, with an increased ratio of chlorophyll to PS II and to PS I in the deeper-growing blades. This acclimation in terms of PS II:PS I for blades growing at lower PFDs, i.e. at greater depths, is in the same direction as in the heterokont (diatom) Skeletonema costatum discussed above. There are no comparable photosynthetic data to permit assessment of specific reaction rate of the photosystems, and hence Fe cost, at the growth PFDs. The change in PS II:PS I with depth in *Macrocystis* is unlikely to be a response to a changed ratio of radiation exciting PS II to that exciting PS I with depth because the radiation exciting PS I relative to PS II probably decreases with depth (Kirk 1994). Thus, the depth response of *Macrocystis*, when interpreted in spectral terms, is the opposite of the acclimation of photosystem ratio as a response to light quality in cyanobacteria (Fujita et al. 1987; Murakami et al. 1997) and higher plants (Chow et al. 1990a,b). The light quality effect in cyanobacteria and higher plants involves a higher PS II:PS I ratio when growth occurs with a predominance of radiation exciting Photosystem I, and vice versa.

Further variations in the effect of photon flux density for growth, independent of far-red enrichment in shade light for terrestrial plants in nature, are a *decrease* in PS II:PS I with a lower photon flux density for growth in the flowering plant *Pisum sativum* (Chow and Anderson 1987, 1990b; Lee and Whitmarsh 1989; Chow et al. 1990b), and the absence of any effect of photon flux density for growth on the PS II: PS I ratio in the unicellular red alga *Porphyridium cruentum* (Cunningham et al. 1989).

The variability in acclimatory effects of photon flux density on PS II:PS I ratio is also seen in the effects of photon flux density for growth on the PS II:cytochrome b_6 -f complex ratio (see Table 3 of Raven 1990; see also Wilhelm and Wild 1984a,b, for effects of photon flux density for growth of *Chlorella* on the PS I:cytochrome b_6 -f complex ratio, and Chow and Hope 1998 for the PS II:cytochrome b_6 -f complex:PS I in *Nicotiana tabacum*). The *direction* of change of PS II:cytochrome b_6 -f complex as a result of varied photon flux density for growth is not always in the same as that for the change in PS II:PS I ratio.

We have emphasized the role of the light climate on phenotypic (acclimatory) variations in the ratio of Fe-containing components of thylakoids. Another obviously important determinant of the production of Fe-containing catalysts of thylakoids is the supply of Fe (Evans 1996; Imsande 1998; Behrenfeld and Kolber 1999).

Greene et al. (1991) examined the effect of Fe sufficiency and Fe limitation on the content of PS I, PS II and the cytochrome b_6 -f complex of the diatom Phaeodactylum tricornutum. Fe deficiency reduced the content of PS I and PS II reaction centres, but not that of the cytochrome b_6 -f complex, on a chlorophyll a basis. As Fe deficiency very substantially reduced the chlorophyll a content per cell, the content of all three complexes (PS I, PS II, cytochrome b_6 -f) is greatly reduced on a cell basis. The reduction in the content per cell of the three complexes is greatest for PS I and least for the cytochrome b_6 -f complex with a change in the PS II:cytochrome b_6 -f complex:PS I ratio from 2.9:7.9:1 in Fe-replete cells to 4.2:19:1 in Fe-deficient cells. The greater decrease in PS I than of the other complexes accords with minimizing the Fe content of the thylakoids. Greene et al. (1991) using the Fe/complex values quoted by Raven (1990), calculated that about 40% of the cellular Fe in Fe-deficient Phaeodactylum tricornutum can be accounted for by PS I, PS II and the cytochrome b_6 -f complex. Hewitt (1983) cites 18.5% as the fraction of leaf Fe

in higher plants that can be accounted for in known photosynthetic and non-photosynthetic components.

Greene et al. (1991) found that the content of the Fe-containing PS I and PS II complexes on a chlorophyll a basis is decreased in Fe-deficient cells. This suggests that, predominantly, the reduced chlorophyll a content per cell in Fe-deficient organisms relates to the decreased content of Fe-containing pigmentprotein complexes, while the pigment-protein complexes which do not contain Fe are associated in a stoichiometric way. It is known that some steps in the chlorophyll synthesis pathway are Fe-dependent. Thus, Figure 9.1 of Marschner (1995) identifies an Fe requirement in the conversion of coproporphyrinogen to protoporphyrinogen, and in the conversion of Mg protoporphyrin to protochlorophyllide. We note that the Fe requirement in converting succinyl CoA plus glycine to δ -aminolevulinic acid shown in Figure 9.1 of Marschner (1995) is not relevant to chlorophyll synthesis since this invariably involves the glutamate pathway of δ -aminolevulinic acid synthesis: Falkowski and Raven (1997). However, the evidence of Greene et al. (1991) suggests that the Fe requirement for chlorophyll synthesis is less significant than a decreased content of Fe-containing photosynthetic apoproteins in causing the lower chlorophyll content of Fe-deficient cells.

Restricted Fe supply also interacts with other growth rate limiting resource inputs such as a low incident photon flux density. Kudo and Harrison (1997) measured the critical Fe value, i.e. the Fe content of cells that was just sufficient to achieve the maximum possible specific growth rate under a given set of growth conditions, as a function of the light supply in the cyanobacterium Synechococcus, and found a higher Fe requirement for low light-grown than high light-grown cells. This agrees with suggestions of Raven (1998a, 1990) that low light adapted (or acclimated) cells, which usually have more pigment per cell and often have fewer reaction centre complexes per unit chlorophyll (but sometimes more per cell) than high light adapted or acclimated cells, have lower achieved rates of electron transport through Fe-containing redox catalysts under their low-light growth conditions than do high light adapted (or acclimated) cells under their high light growth conditions. Therefore cells growing at low light need more Fe per unit C fixed per second than do cells growing in high light (Raven 1990).

One possibility for offsetting this greater Fe cost of C fixation during photosynthesis at low photon

flux densities is a reduction in cell size, and thus in the package effect. The package effect (Kirk 1994) involves a lower specific absorption coefficient of pigment, and hence a lower efficiency of light absorption per unit pigment, within suspended particles relative to that in a monodisperse solution at the same concentration of pigment (mol m^{-3}). This effect increases, for a given concentration of pigment per unit volume of particle, with increasing size of the particle (subthylakoid particles; whole thylakoids; chloroplasts; cells; multicellular organs). The package effect means that the rate of photon absorption per pigment molecule is, for a given incident photon flux density, greater for structures with a smaller package effect. For a given ratio of pigment to thylakoid Fe (or Mn), the greater efficiency of photon absorption and hence the increased rate of excitation energy transfer to reaction centres and of electron flow through Fe- (and Mn-) containing catalysts, should be greater in cells with a smaller package effect (Raven 1986, 1987b, 1997c, 1998a, 1999). Thus, a smaller package effect can not only lead to an increased rate of photosynthesis per unit pigment at low photon flux densities, but can also increase the rate of photosynthesis per unit Fe (or Mn) (Raven 1998a, 1999). The work of Sunda and Huntsman (1997) and Van Leeuwe et al. (1997) are examples to which this package effect argument can be applied in relation to photosynthetic rates of marine phytoplankton cells when Fe and light are limiting growth rates.

Work in which the ratio of the major ironcontaining complexes in thylakoids has been altered phenotypically in response to treatments other than Fe supply, i.e growth under putatively Fe-sufficient growth conditions, includes photon flux density (400– 700 nm) for growth (e.g. Falkowski et al. 1981; Dubinsky et al. 1986; Wilhelm et al. 1989; Burger-Wiersma and Matthijs 1990; Greene and Gerard 1990), the spectral composition of the radiation supplied for growth (e.g. Fujita et al. 1987; Chow et al. 1990b; Murakami et al. 1997), the inorganic C supply (Manodori and Melis 1987; Murakami et al. 1997; cf. Li and Canvin 1998) and the external Na⁺ concentration (Murakami et al. 1997). Where variations in incident photon flux density of a given spectral composition influence the ratio of PS I to PS II, it is invariably to decrease the ratio at lower photon flux densities, although there is sometimes no phenotypic influence of the rate of photon supply for growth on the PS I:PS II ratio. For those spectral effects that relate predominantly to the fraction of incident light absorbed by

antenna pigments serving particular light reactions, it is found that light predominantly exciting PS I leads to an increased PS II:PS I ratio and *vice versa*. Restricted inorganic C supply and high Na⁺ decreased the PS II:PS I ratio in *Synechocystis* (Manodori and Melis 1987; Murakami et al. 1997).

These effects may be related to the influence of the various environmental factors on the redox state of PQ and/or the cytochrome b₆-f complex, a high PQH₂/PQ ratio decreasing expression of PS II and vice versa (see Hihara et al 1998; Allen 1993; Pfannschmidt et al. 1999). For changes in photon flux density, a tendency for reactions downstream of both photoreactions, CO₂ fixation and the Mehler reaction, to have a greater capacity to accept electrons than PS I and PS II have to supply them after a decrease in incident photon flux could lead to a lower PQH₂/PQ ratio and a smaller extent of reduction of the b_6 -f complex. This could cause differential gene expression such that the photoinhibition-sensitive PS II is expressed more at lower than at higher photon flux densities relative to the expression of PS I, with the possibility of an increase in the overall photon yield of photosynthesis. However, this sort of mechanism does not guarantee an increase in the overall photon yield and poses problems at high photon flux densities, where photoinhibition sensu lato can lower the PQH₂/PQ ratio and decrease the effective PS II:PS I ratio.

For spectral variations in light supply, the logical chain is less tortuous. Light that preferentially excites PS I causes a decrease in the PQH₂/PQ ratio and increases expression of PS II, balancing the excitation inputs to PS I and to PS II. The reverse occurs with light that preferentially excites PS II.

In the case of restricted inorganic C supply, there is expression (or increased expression) of inorganic carbon concentrating mechanisms dependent on cyclic electron transport, if not cyclic photophosphorylation (Klugehammer et al. 1997; Sültemeyer et al. 1997; Ohkawa et al. 1998; cf. Ogawa et al. 1985; Ogawa 1991). High Na⁺ concentrations increase the rate of the directly or indirectly ATP-powered active Na⁺ efflux (Murakami et al. 1997). In both cases, the argument an increased rate of ATP consumption would, if ATP is only available from non-cyclic photophosphorylation, lead to an increased NADPH/NADP⁺ ratio and hence to an increased PQH₂/PQ ratio. This increase in PQH₂/PQ not only leads (in most organisms) to a state transition increasing the ratio of excitation energy reaching PS I relative to PS II but also, if the increased PQH₂/PQ is long sustained (more

than minutes), to changed degrees of gene expression (Allen 1993). These longer-term effects yield an increased expression of PS I, with the likelihood of increased ATP supply by cyclic photophosphorylation. However, it may be difficult with such redox control of the capacity for cyclic photophosphorylation, to avoid over-reduction of the redox carriers of the cyclic pathway. This is due to reductant inputs from the reducing end of both PS II and PS I, unless cyclic photophosphorylation involves distinct PS I units having no redox contact with PS II. This would have important implications for flexibility of operation of thylakoid reactions.

Overall, the treatments that produce a high PQH₂/PQ would favor an increase in the high Fedemanding PS I relative to the low Fe-demanding PS II; the influence on the cytochrome b_6 -f complex, with its intermediate Fe requirement, is variable, sometimes tracking PS II and sometimes, more closely following PS I expression (Wilhelm et al. 1989; Murakami et al. 1997). It would be teleologically appropriate if the greater relative Fe demand engendered by a high PQH₂/PQ ratio resulted in enhanced Fe acquisition by cell surface Fe(III) reductases. This would require that Fe(III) supply to the cell surface would not be limiting.

These predictions of Fe costs of photosynthetically supported growth are at least qualitatively validated for Fe contents and growth rates of microalgae as a function of differences in combined N source (NH₄⁺ or NO₃⁻) (Raven 1988a; Rueter, 1993; Maldonado and Price 1996; Kudo and Harrison 1997; cf. Henley and Yin 1998). However, the Fe costs of growth as predicted by Raven (1988a) are greatly (10-fold) in excess of the values measured for an oceanic diatom (Thalassiosira oceanica) adapted to the low-Fe open ocean habitat, although they are closer to the values measured for a coastal diatom (Thalassiosira weisflogii) from a higher-Fe habitat (Sunda et al. 1991; Sunda and Huntsman 1995a). This difference between coastal and open ocean phytoplankton organisms is also seen for coccolithophorids and dinoflagellates, oceanic coccolithophorids being the most Fe-efficient of the organisms tested by Brand (1991). The least Fe-efficient organisms tested by Brand (1991) were coastal and other marine cyanobacteria (see Wilhelm et al. 1996; Muggli and Harrison 1997). The high Fe cost of growth of cyanobacteria (low PS II:PS I) relative to chlorophyll b-containing prasinophytes and chlorophyll c-containing raphidophytes, dinophytes, coccolithophorids (high PS II:PS I) agrees with predictions of Raven (1988a, 1990). However, the very

low Fe cost of growth in some oceanic eukaryotic phytoplankton organisms is a considerable challenge to our belief that we understand the Fe content of particular catalysts, their *in vivo* specific reaction rate, and the stoichiometry of their reaction products with C assimilation and growth (Raven 1988a, 1990).

Mn presents a less complex case than than Fe, and the achieved Mn cost of growth (Sunda and Huntsman 1986, 1998) is in reasonable agreement with prediction (Raven 1990; Sunda and Huntsman 1998). The prediction here is based on the use of Mn predominantly in PS II, with some involved in Mn superoxide dismutase in mitochondria of eukaryotes and in thylakoids of cyanobacteria (Okada et al. 1979).

For Cu and Zn, the other trace metals involved in thylakoid-associated and other electron transport reactions neither estimates of the metal cost of growth nor measurements of the actual metal cost of growth are available.

Cu has a major photosynthetic (thylakoid) role in plastocyanin. Plastocyanin is not encoded in the genome of all organisms, nor, if encoded, is it always expressed (Falkowski and Raven 1997); the ratio of Cu/Fe availability is a major determinant of expression (Falkowski and Raven 1997). The suggestion that there is a necessary occurrence of Cu in PS II (Droppa et al. 1984; Droppa and Horváth 1990), seems to result from nuclear contamination of PS II preparations (Avellano et al. 1994). A well substantiated role for Cu in the thylakoids of peridinin-containing dinoflagellates, Charophyceae sensu lato and Embryophyta is in thylakoid-associated Cu-Zn superoxide dismutase (Henry et al. 1976; Asada et al. 1980; de Jesus et al. 1989; Butow et al. 1997; cf. Asada et al. 1977). Roles of Cu elsewhere in photosynthetic organisms include stroma and cytosol Cu-Zn superoxide dismutase (in peridinin-containing dinoflagellates, Charophyceae sensu lato and Embryophyta) cytochrome oxidase, and certain other oxidases.

Zn has a role in thylakoids in Cu–Zn superoxide dismutase in the Dinophyta (or at least the 'normal' peridinin-containing representatives), Charophyceae sensu lato and Embryophyta (Henry et al. 1976; Asada et al. 1980; de Jesus et al. 1989; Butow et al. 1997; cf. Asada et al. 1977) and a non-electron transport role for any thylakoid-associated carbonic anhydrase (Karlson et al. 1996, 1998; Raven, 1997a,d; Stemler 1997), regardless of which of the three gene families of carbonic anhydrase might be involved because they all involve Zn (Hewett-Emmett and Tashian 1996). Non-thylakoid roles and locations for Zn in

photosynthetic organisms include Cu-Zn superoxide dismutase in stroma and cytosol (Charophyceae and Embryophyta) and thylakoid, stroma, cytosol and extracellular carbonic anhydrase required for inorganic carbon assimilation using diffusive CO2 entry and when active inorganic carbon transport occurs, as well as in 'Zn fingers' and some phosphatases and other enzymes (Vallee and Galdes 1984; Williams and Fraústo da Silva 1996; Raven 1997d; Karlson et al. 1998). It is of interest that Chlorella pyrenoidosa requires a much higher Zn concentration in the growth medium for photolithotrophy (0.77 mmol m^{-3}) than for chemoorganotrophy (77 nmol m^{-3}). This presumably reflects a very large Zn requirement for carbonic anhydrase in this alga (Eyster 1964) which lacks Cu–Zn superoxide dismutase. Although HCO₃⁻ has a protective role on the oxidising lumen side of PS II (Klimov et al. 1997; Hulsebosch et al. 1998), it is not known if this involves the intrathylakoid carbonic anhydrase (Karlson et al. 1998). Eyster (1964) also showed the expected (Raven 1988a) greater need for Fe (18 mmol m^{-3}) and Mn (0.1 mmol m^{-3}) in photolithotrophic than in chemoorganotrophic growth (Fe $1 \,\mu\mathrm{mol}\,\mathrm{m}^{-3}$ and Mn $1 \,\mu\mathrm{mol}\,\mathrm{m}^{-3}$, respectively). However, Tortel et al. (1996) showed a higher Fe:C ratio in marine chemoorganotrophic bacteria than in eukaryotic phytoplankton. This is contrary to the predictions of Raven (1988a) if the achieved C assimilation rates in the work of Tortel et al. (1996) are equal for the chemoorganotrophs and the photolithotrophs. The maximum specific growth rates of the chemoorganotrophic bacteria in Figure 1 of Tortel et al. (1996) are about five times those for eukaryotic phytoplankton at the same temperature (20 °C; Raven 1986). This means that the C assimilation rate per unit (catalytic) Fe in the chemoorganotrophic cells is higher than in the eukaryotic photolithotrophs, as predicted by Raven (1988a). Further work is needed with specific attention to the achieved in situ growth rate of chemoorganotrophic bacteria (Pakulski et al. 1996) and phytoplankton in the same, Fe-limited, conditions (Pakulski et al. 1996). The situation is further complicated by the high Fe costs of growth found for phagotrophic marine protozoa (Chase and Price 1997).

Evolution of trace metal requirements in thylakoid reactions

In addition to the chemical 'fitness' of a given trace metal to fulfill a given role in thylakoid redox reactions, a consideration of the evolution of such roles requires that we consider the requirement, as perceived by an inadequately informed human observer from, so frequently, several billion years distance for that role at a given time as well as the availability of trace elements at that time. It is also necessary to include any known or likely needs for a given trace metal in any other roles in the organism.

The main change in the environment of the earth that has influenced the availability of Fe, Mn, Cu and Zn over the last 4 billion years has been the increased O₂ level, and increased redox potential (biological scale, with H₂ electrode negative, O₂ electrode positive). Because photodissociation of H₂O vapor can only produce O_2 at $\leq 10^{-8}$ of the present atmospheric level, the accumulation of O₂ in the original CO₂/N₂/H₂O vapor atmosphere depends on the occurrence of O₂-producing photosynthesis. O₂ accumulation also depends on the oxidation of all of the primeval O2-consuming inorganic oxidants (mainly Fe^{2+} , S^{2-}) and on the sequestration of the organic C counterpart of the O₂ as products of photosynthesis so that it cannot recombine with the O₂ by biotic (respiratory) or abiotic means.

There is evidence for Rubisco-based photosynthetic CO_2 assimilation, the core of CO_2 fixation process in all extant O_2 -evolvers, as long as 3.85 billion years ago (Schidlowski 1988; Mojzis et al. 1996; Rosing 1999; cf. Eiler 1997), and there is fossil evidence for cyanobacteria, all of which today have the capacity for O_2 evolution, from 3.45 billion years ago. However, O_2 build-up *globally* to more than about 10^{-8} of the present atmospheric level did not occur until about 2 billion years ago, approximately coincidental with the origin of eukaryotes. If photosynthetic O_2 evolution indeed began as early as 3.45 billion years ago, then the absence of a global O_2 build-up until more than a billion years later could be related to the oxidation of Fe^{2+} to Fe^{3+} and S^{2-} to SO_4^{2-} .

The significance of this change from global anoxia before 2 billion years ago to (irregularly) increasing O₂ levels after 2 billion years ago for metal availability is as follows. Fe²⁺ and Mn²⁺ were relatively abundant in the anoxic ocean before 2 billion years ago. Cu was unavailable because it was present as cuprous ion (Cu⁺), which was precipitated as the very insoluble Cu₂S, and Zn (Zn²⁺) was very probably less available than it is in today's oxygenated ocean due to the high dissolved S²⁻ level (Williams and Frausto da Silva 1996; Raven 1997b,d, 1998d). This global scenario permits the 'luxurious' use of Fe and Mn in

the photosynthetic redox catalysts PS I and PS II, cytochrome b_6 -f complex, cytochrome c_6 , ferredoxin and NAD(P)H dehydrogenase. Perhaps this is not true in Fe- or MnSOD, in catalase or in ascorbate or glutathione peroxidases, because their substrates O_2^{-} and H₂O₂ would not have been readily available in early anoxic oceans (although FeSOD is found in some extant obligately anaerobic bacteria: Asada et al. 1980). However, this scenario would not favor the use of Cu in plastocyanin or of Cu and Zn in Cu-Zn superoxide dismutase, which in any case would not have a substrate in the absence of significant free O2. Oceanic dissolved Fe (Fe²⁺) could also have had a role in providing a UV screen for early biota near the ocean surface before significant atmospheric O₂ built up, permitting the production of an atmosphere O₃ screen for UV and paralleling the occurrence of very low UVabsorbing Fe³⁺ in the ocean (Cleaves and Miller 1998; cf. Mulkidjianian and Junge 1997).

This view has an early recruitment of Fe and Mn into the redox reactions of O2-evolving photosynthesis. Many of the Fe-containing components could have been derived from earlier Fe-containing catalysts in chemolithotrophs, phototrophs which could not evolve O2, or chemoorganotrophs using an electron acceptor for respiratory electron transport. Such an electron acceptor could be abiotically generated NO_2^- , derived from 2 $CO_2 + N_2 \rightarrow 2 CO_2 + 2 NO^-$, or N₂ + 2 H₂O \rightarrow 2 H₂ + 2 NO , energized by lightning or by meteorite impacts (Mancinelli and McKay 1988; Summers and Chang 1993; Yung and McElroy 1997; Raven and Yin 1998). Fe could have had earlier important involvement in the origin of life via continued production of the UV-labile NH₃ needed for amino-acid and nucleotide base synthesis from NO₂ derived from the NO generated by lightning or bolide impacts (Summers and Chang 1993; cf. Falkowski 1997). However, the need for NH₃ synthesis would be negated by an origin and early evolution of life using magmatic NH₃ at hydrothermal vents well out of the range of UV-B penetration (Russell and Hall 1997). Further possible involvement of Fe in the chemolithotrophic evolution of life has been proposed by Wächterhäuser and by Russell (summarized by Maden 1995; see Russell and Hall 1997), using FeS, and leading to FeS protein catalysts consistent with very early occurrence of chemolithotrophic cells (Pace 1997). A delayed recruitment of Cu and Zn into redox catalysts could have been a result of one or more of:

1. Unavailability of Cu and Zn in the early anoxic ocean;

- 2. The occurrence of Fe and Mn-containing catalysts that would perform the same function (Fe or Mn superoxide dismutases; cytochrome *c*₆ instead of plastocyanin); and
- 3. The absence of substrates for the catalysts (little or no ${\rm O_2}^-$ for superoxide dismutase or ${\rm H_2O_2}$ for catalase or ascorbate peroxidase when ${\rm O_2}$ was $\leq 10^{-8}$ of the present level).

A number of considerations suggest that this is an oversimplification of the sequence of metal use, or at least under-estimates the time for which Cu has functioned in redox reactions. One line of evidence comes not from photosynthesis per se but from respiration. Schäfer et al. (1996) point out that the Cu-containing cytochrome oxidase seems to have a time of origin, based on molecular phylogenetic considerations, which considerably predates the global occurrence of O_2 at above 10^{-8} of the present atmospheric level, i.e. it originated well before 2 billion years ago. This presents two O₂-related problems: one is the global absence of O2, the 'normal' substrate for electron donation by cytochrome oxidase; the other is the global absence of available Cu for synthesis of cytochrome oxidase in the global absence of O₂. A second problem with a later origin of Cu-dependent catalysts, this time directly related to O₂-evolving photosynthesis, is the phylogenetic distribution of plastocyanin. The widespread occurrence of plastocyanin in cyanobacteria indicates an origin early in the evolution of cyanobacteria (e.g. Kaneko et al. 1996). Of course, the distribution patterns of the cytochrome oxidase and plastocyanin could be explained by lateral gene transfer. This may help to explain the distribution of plastocyanin among extant cyanobacteria, but it is very unlikely to be the cause of the (apparently) monophyletic origin and subsequent phylogenetic distribution of cytochrome oxidase.

This 'Cu-anomaly' can be rationalized by the evolution of Cu-requiring catalysts in oxygenated microenvironments in a globally anoxic environment. There seems to be no such problem with the timing of the evolution of another photosynthesis-related, Zn-containing, redox enzyme, i.e. the Cu, Zn form of superoxide dismutase because it did not appear until after the origin of eukaryotes 2 billion years ago. Sporadic presence of the enzyme in prokaryotes could be explained by lateral gene transfer. This also fits with the use of Zn in zinc fingers in eukaryotes. However, there are arguments that Zn could have been involved earlier than 2 billion years ago in carbonic anhydrase, which is not a photosynthetic *redox* enzyme, although

Co could have substituted for it here (Morel et al. 1994; Sunda and Huntsman 1995b; Raven 1997d). The relatively low early availability of Zn relative to its present availability, and especially to the ready availability of Mg throughout the history of life, does not favor the early use of Zn in chlorophylls in place of mg. Although Zn occurs in bacteriochlorophyll in the acidophilic photosynthetic bacterium *Acidiphilium rubrum* (Wakao et al. 1996) Zn-chlorophyll has not been reported from O₂-evolvers.

Although not strictly a metal, Se is a trace element that is present as selenocysteine in the (H₂O₂-removing) Fe-containing glutathione peroxidase (Stadtman 1996), a significant peroxidase in some algae (Chlamydomonas; many diatoms) and perhaps some cyanobacteria (see 'Linear electron flow with O2 as electron acceptor: the Mehler peroxidose reaction' section above). It is of interest that glutathione peroxidase of higher plants, where it is a minor component relative to ascorbate peroxidase, has cysteine rather than selenocysteine and has very low maximum specific activity relative to the Secontaining mammalian enzymes (Eshdat et al. 1997). The use of selenocysteine rather than cysteine may to be a primitive feature related to anoxic or low O₂ habitats as -SeH is much more readily auto-oxidized by O₂ than is -SH (Lee and Berry 1996). However, the occurrence of Se-containing glutathione peroxidase in O₂-evolving organisms means that it can function in cellular environments with O2 concentrations in excess of those in an air-equilibrated solution (see Raven et al. 1994).

The high Fe (and Mn) requirements of thylakoid redox reactions, and hence the high Fe requirement for photolithotrophic growth, were probably a significant contribution to the natural selection pressures related to the evolution of the range of mechanisms for acquiring Fe from low-solubility Fe(III) complexes found in cyanobacteria, algae and higher plants in aerobic environments. These mechanisms involve surface acidification and reduction of Fe(III) to Fe(II), which is then taken up as Fe²⁺ by specific transporters in the plasmalemma (in eukaryotic primary producers other than grasses), or production of siderophores that chelate Fe(III), with uptake of the Fe(III) siderophore complex by specific transporters in the plasmalemma (cyanobacteria, grasses) (Raven 1988b; Hutchins 1995; Marschner 1995; Walls et al. 1995). The terrestrial higher plants thus have both the cyanobacterial (siderophores in grasses) and eukaryotic algal (Fe(III) reduction in a wide range of vascular plants) mechanisms. The Fe requirement for thylakoid redox reaction in higher plants is, of course, relatively immutable, subject to modification only by changes in the ratio of the complexes for the two Photosystems and other redox complexes in the thylakoid membrane. There is no substitution of ferredoxin by flavodoxin (slightly reducing the Fe requirement) or substituting plastocyanin by cytochrome c_6 (thereby slightly increasing the Fe requirement).

An enigmatic 'Fe-sparing' effect of V has been reported by Meisch and Beilig (1975) and Wilhelm and Wild (1984a), but not in higher plants (Hewitt 1983). This differential effect was attributed to an effect of V on a non-enzymatic transamination of δ -aminolevulinic acid in algae but not higher plants (Meisch et al. 1978; Hewitt 1983). However, it is now clear that the (enzymic) glutamate pathway of δ -aminolevulinic synthesis is ubiquitous for chlorophyll synthesis in O₂-evolvers (Falkowski and Raven 1997). There is no evidence that Fe is replaced by V in any Fe-containing catalyst of electron transport in photosynthesis, or that V was involved in the evolution of the Fe-containing catalysts of photosynthetic processes.

Conclusions

The involvement of trace metals in thylakoid reactions can in large part be explained by their redox chemistry and the availability during the evolution of the particular reaction sequence.

Although the quantitative role of trace metals in thylakoids is well understood for the major trace metal-containing complexes of linear electron transport from H₂O to NADP⁺ and thence to CO₂, there are many fewer data on minor components. These minor components include the NAD(P)H-PQ oxidoreductase, which may be involved in chlororespiration and in cyclic electron flow and photophosphorylation, and the superoxide dismutases and peroxidases associated with the Mehler peroxidase reaction.

Major variations in Fe costs of linear electron transport occur via phylogenetically determined and acclimatory changes in the content of major Fecontaining complexes (PS II, cytochrome b_6 -f, PS I). However, the acclimatory changes are not overtly driven by Fe availability but rather by changes in light quantity and quality. By contrast, variations in Fe availability alter the ferredoxin/flavodoxin ratio in some algae and cyanobacteria, while changes

in Cu and Fe availability alter the cytochrome c_6 /plastocyanin ratio in some green algae and cyanobacteria. These switches have a relatively minor impact on overall Fe costs of linear electron transport, but have a very large impact on the Cu costs of linear electron transport.

Note added in proof

Hippler et al. (1998) give a detailed review of the use of *Chlamydomonas* genetics in studying biogenetic pathways, including H_2 metabolism. Hoefnagel et al. (1998) consider the interdependence between chloroplasts and mitochondria in the light and the dark, including a critical consideration of chlororespiration. Finally, Stewart and Brudwig (1998) review the stoichiometry and functioning of cytochrome b_{559} of Photosystem II.

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