

Regulating the proton budget of higher plant photosynthesis

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Communicated by Clarence A. Ryan, Washington State University, Pullman, WA, May 12, 2005 (received for review February 7, 2005)

In higher plant chloroplasts, transthylakoid proton motive force serves both to drive the synthesis of ATP and to regulate light capture by the photosynthetic antenna to prevent photodamage. *In vivo* probes of the proton circuit in wild-type and a mutant strain of *Arabidopsis thaliana* show that regulation of light capture is modulated primarily by altering the resistance of proton efflux from the thylakoid lumen, whereas modulation of proton influx through cyclic electron flow around photosystem I is suggested to play a role in regulating the ATP/NADPH output ratio of the light reactions.

ATP synthase proton conductivity | cyclic electron flow | linear electron flow | energy-dependent nonphotochemical quenching | protein motive force

Photosynthesis converts light energy into chemical energy, ultimately powering the vast majority of our ecosystem (1). Higher plant photosynthesis is initiated through absorption of light by antennae complexes that funnel the energy to photosystem (PS) II and I. The photosystems operate in sequence with the plastoquinone pool, the cytochrome *b₆f* complex, and plastocyanin to oxidize H₂O and reduce NADP⁺ to NADPH in what is termed linear electron flow (LEF). LEF is coupled to proton translocation, establishing a transthylakoid electrochemical gradient of protons, termed the proton motive force (*pmf*) (2), comprised of electric field ($\Delta\psi$) and pH (ΔpH) gradients (3).

Dual Role of the *pmf*

The *pmf* plays two central roles in higher plant photosynthesis (4). First, *pmf* drives the normally endergonic synthesis of ATP through the CF₁-CF₀ ATP synthase (ATP synthase) (5). Both the ΔpH and $\Delta\psi$ components of *pmf* contribute to ATP synthesis in a thermodynamically, and probably kinetically, equivalent fashion (6). Second, *pmf* is a key signal for initiating photoprotection of the photosynthetic reaction centers through energy-dependent nonphotochemical quenching (q_E), a process that harmlessly dissipates excessively absorbed light energy as heat (7–10). Only the ΔpH component of *pmf*, through acidification of the lumen, is effective in initiating q_E by activating violaxanthin de-epoxidase, a lumen-localized enzyme that converts violaxanthin to antheraxanthin and zeaxanthin, and by protonating lumen-exposed residues of PsbS, a pigment-binding protein of the PS II antenna complex (11).

A Need for Flexibility in the Light Reactions

A major open question concerns how the light reactions achieve the flexibility required to meet regulatory needs and match downstream biochemical demands (12). In LEF to NADP⁺, the synthesis of ATP and the production of NADPH are coupled, producing a fixed ATP/NADPH output ratio. LEF alone is probably unable to satisfy the variable ATP/NADPH output ratios required to power the sum of the Calvin–Benson cycle (13, 14) and other metabolic processes (alternate electron and ATP sinks) that are variably engaged under different physiological conditions (12, 15, 16). Failure to match ATP/NADPH output with demand will lead to buildup of products and depletion of

substrates for the light reactions, leading to inhibition of the entire process.

The generation of *pmf* is likewise coupled to LEF, so it is clear that the sensitivity of antenna regulation (or q_E) must also be modulated in some way to avoid catastrophic failure of photoprotection (12, 15, 17–19). Longer-term acclimation of the q_E response can involve altering the sensitivity of the regulatory machinery to lumen pH by changing the xanthophyll pigment and/or PsbS levels (12, 20). However, dramatic changes in light intensity and/or CO₂ availability can occur over the seconds-to-hours time scale (8), requiring short-term adjustments. Indeed, it has been demonstrated that short-term alteration of CO₂ or O₂ levels can strongly modulate (by up to 6-fold) the sensitivity of q_E with respect to LEF (17, 18).

Two Types of Flexibility Mechanisms

Two general types of models have been proposed to account for the flexibility required to meet these changing demands (12). In “Type I” mechanisms, proton flux into the lumen is increased through alternate electron transfer pathways, especially cyclic electron flow associated with PSI (CEF1), a mechanism that returns electrons from PSI to the plastoquinone pool, thereby increasing the magnitude of the *pmf* relative to that generated by LEF alone (12). Other processes are also possible, for example, turnover of a plastid terminal oxidase (21, 22), but these processes would have to run at relatively high rates to significantly impact the overall ATP/NADPH balance. For C₃ vascular plants, CEF1 has been suggested to supply the relatively small fluxes (10–15% of that supplied by LEF) of protons required to balance ATP/NADPH output for the Calvin–Benson cycle and nitrogen assimilation (13, 14). It is a matter of intense debate (23, 24) as to whether CEF1 can run at sufficiently high rates to alter q_E responses by up to 6-fold, especially given the expected large ATP/NADPH imbalances such large fluxes would likely incur (12, 16).

In Type II mechanisms, lumen acidification with respect to LEF is adjusted without changing the relative flux of protons into the lumen, thus modulating q_E sensitivity without impacting ATP/NADPH output. This phenomenon is thought to be achieved by varying either the conductivity of the CF₁-CF₀ ATP synthase to proton efflux as measured by electrochromic shift (ECS) decay (g_{H^+}), i.e., the inverse of the resistance to proton efflux from the lumen or the relative fraction of *pmf* stored as ΔpH (12, 16–18, 24).

Probing the *pmf* to Gain Insight into the Flexibility Mechanisms

Recently, a series of *in vivo* probes of the *pmf* have been introduced (2, 3, 16, 25–28), allowing contributions from Types I and II flexibility mechanisms to be directly assessed. These

Abbreviations: CEF1, cyclic electron flow associated with PSI; ECS, electrochromic shift; LC, low CO₂ (50 ppm CO₂, 21% O₂); LEF, linear electron flow; PS, photosystem; *pmf*, proton motive force; *pmf*_{LEF}, *pmf* generated by LEF; q_E , energy-dependent nonphotochemical quenching.

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techniques are based on kinetic analyses of the ECS (26) of photosynthetic pigments, which yields absorbance changes proportional to changes in transthylakoid $\Delta\psi$ (29). Several useful parameters can be obtained from analysis of ECS decay kinetics during brief dark perturbations of the steady state, including estimates of the relative flux of protons through the ATP synthase (v_{H^+} , which at steady state equals flux of protons into the lumen), the magnitude of the light-induced pmf , the fraction of pmf stored as ΔpH and $\Delta\psi$, and g_{H^+} (3, 16–18, 25, 26, 28). Combined with standard chlorophyll a fluorescence assays, from which estimates of LEF can be obtained (30), one can calculate the pmf generated by LEF alone (i.e., $pmf_{LEF} = LEF/g_{H^+}$), a key parameter for estimating fractional changes in CEF1 turnover (17, 18).

Using these probes of the proton circuit, it was shown that in intact *Nicotiana tabacum* (tobacco) leaves, lowering atmospheric CO_2 from 372 to 0 ppm led to a ≈ 5 -fold increase in the dependence of q_E on LEF (17). The effect could be entirely accounted for by a proportional (i.e., 5-fold) decrease in g_{H^+} , so that even modest rates of LEF generated a substantial pmf and a robust q_E response (17, 18). A similar (≈ 6 -fold) change in q_E sensitivity was observed when both O_2 and CO_2 were lowered (to 1% and 50 ppm, respectively), but in this case, both changes in g_{H^+} and increased partitioning of pmf into ΔpH were invoked to explain the effect (18). In both cases, the ratio of v_{H^+}/LEF remained essentially constant (within noise levels), indicating that contributions from CEF1 to proton flux were either small or remained a relatively constant fraction of those from LEF, as previously found for tobacco (25). On the whole, these results support a large role for Type II mechanisms in modulating q_E sensitivity upon short term changes in CO_2/O_2 levels, but they do not rule out smaller contributions from Type I mechanisms in balancing ATP/NADPH output (12, 16, 28).

On the other hand, Munekage *et al.* (31, 32) recently presented partial characterization of a mutant strain of *Arabidopsis thaliana*, termed *pgr5* for proton gradient regulation, which showed two provocative phenotypes. First, nonphotochemical reduction of the plastoquinone pool, attributed to the key step in CEF1, was inhibited in *pgr5*. Second, q_E was severely diminished. It is reasonable to hypothesize that the loss of PGR5 blocks CEF1 and, thereby, abolishes a significant flux of protons needed to activate q_E (31, 32). Evidence for such a hypothesis would support a large role for Type I mechanisms in modulating q_E sensitivity (33) while arguing against Type II models (12, 17, 18). On the other hand, mutation of *pgr5* could indirectly affect q_E by disrupting downstream processes and modulating metabolic pool sizes (31, 32). Here, we present an experimental test for causal links between the loss of PGR5, steady-state proton flux, and the q_E response, allowing us to determine the relative roles of Type I and II flexibility responses.

Materials and Methods

Plant Strains and Growth Conditions. Wild-type *A. thaliana* (Wt-background strain *gl1*) (31) and *pgr5* plants were grown in tightly controlled chambers under a 16:8 photoperiod at an average of $\approx 70 \mu\text{mol photons m}^{-2}\text{s}^{-1}$ photosynthetically active radiation and at 23°C. These growth conditions stably reproduced the phenotypes seen previously over the entire experimental period (31). Wt (*gl1*) and *pgr5* seeds were a gift from T. Shikanai (Nara Institute of Science and Technology, Ikoma, Nara, Japan).

Spectroscopic Assays. Fully expanded leaves from ≈ 23 - to 26-day-old plants were used in spectroscopic assays. Room air (372 ppm $CO_2/21\% O_2$) or premixed gases from cylinders (i.e., 50 ppm $CO_2/21\% O_2$) were bubbled through water (for humidification) before entering the measuring chamber of the spectrophotometer. Leaves were clamped into the measuring chamber of a nonfocusing optics spectrophotometer/chlorophyll fluorometer,

specifically designed for use on leaves (17, 18, 34). Leaves were first exposed to 26–216 $\mu\text{mol photons m}^{-2}\text{s}^{-1}$ photosynthetically active radiation from a series of red light emitting diodes (maximum emission wavelength of 637 nm) to reach steady-state conditions (10 min). Further preillumination had little additional effect. After this actinic period, the steady-state (F_s) and light saturated (F_M') levels of chlorophyll a fluorescence yield were obtained (17, 18), from which estimates of the efficiency of PSII photochemistry (Φ_{II}) were calculated (30). Estimates of LEF were obtained by using Φ_{II} as in ref. 35. Analyses of the ECS decay kinetics upon perturbation of the steady state with an ≈ 300 -ms dark period were performed as described in refs. 17, 18, and 26. Previous assays showed linear correlations between estimates of LEF taken from fluorescence and absorbance measurements with our instruments (25–27, 34), suggesting that the spectroscopic techniques probed similarly responding populations of chloroplasts. Absorbance changes at 505, 520, and 535 nm were recorded in series, and those attributable to changes in ECS were deconvoluted from background signals according to the following equation (25, 26):

$$\Delta ECS = -\Delta I/I_{0(520)} - ((-\Delta I/I_{0(535)} + -\Delta I/I_{0(505)})/2). \quad [1]$$

An estimate of steady-state, light-induced pmf , termed ECS_t , was taken as the total amplitude of ECS decay from its steady-state level to its minimum quasi-stable level after ≈ 300 ms dark period (16–18). Relative estimates of the conductivity of the thylakoid membrane to protons (g_{H^+}), primarily attributable to the turnover of the ATP synthase, were obtained by taking the inverse of the time constant for ECS decay (τ_{ECS}) (16–18, 28). Relative estimates of the pmf attributable to proton flux from LEF, termed pmf_{LEF} , were calculated by using the following equation (16, 18, 28):

$$pmf_{LEF} = LEF/g_{H^+}. \quad [2]$$

This parameter estimates the light-driven proton flux through the ATP synthase based on the extent of LEF and the kinetic properties of the ATP synthase turnover, as reviewed in ref. 28.

Western Blot Analyses. Crude leaf extracts from Wt and *pgr5* were prepared as described in ref. 36. Flash-frozen tissue was ground in a mortar and pestle before resuspension in SDS/PAGE sample buffer. Ten micrograms of protein, as estimated by using the BCA Protein Assay Kit (Pierce), from each preparation was loaded onto an SDS/PAGE gel. Protein was transferred to poly(vinyl difluoride) membranes and probed with antibody directed against the β -subunit of the ATP synthase (a gift from Alice Barkan, University of Oregon, Eugene, OR). Immunoreactive bands were detected on radiographic film by using the SuperSignal West Pico Chemiluminescent Substrate kit (Pierce). Similar conclusions were reached when the gel was loaded with 10, 30, or 90 μg of protein, indicating that the assay was within the linear range of detection (data not shown).

Results and Discussion

Effects of Lowering CO_2 Levels and Loss of PGR5 on LEF and q_E Sensitivity. Fig. 1A shows plots of q_E as a function of LEF from 26–216 $\mu\text{mol photons m}^{-2}\text{s}^{-1}$ for the wild type (Wt, *gl1*) (31) under ambient air (372 ppm $CO_2/21\% O_2$) and two different treatments that lowered light saturated LEF by about the same extent. Low CO_2 air (LC; 50 ppm $CO_2/21\% O_2$) reduced light-saturated LEF in Wt by $\approx 30\%$, a typical response for *A. thaliana* (35). A similar lowering of light-saturated LEF was obtained by using *pgr5* under ambient air. These conditions were chosen to avoid significant photoinhibition, which appeared in *pgr5* at $>216 \mu\text{mol photons m}^{-2}\text{s}^{-1}$ as a decrease in LEF, as well

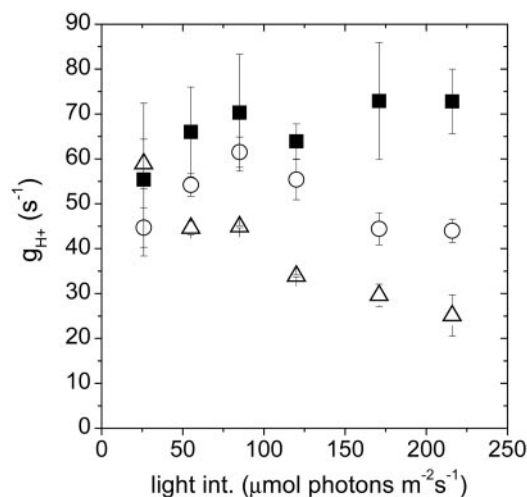


Fig. 3. The light intensity dependence of the proton conductivity of the ATP synthase (g_{H^+}). Estimates of g_{H^+} in Wt and *pgr5* from 26 to 216 $\mu\text{mol photons m}^{-2}\text{s}^{-1}$ were obtained by taking the inverse of the time constant for ECS decay during a 300-ms dark perturbation of steady-state conditions. Conditions and symbols are as in Fig. 1. Error bars represent SE for $n = 3-6$.

these results supports the validity of comparisons of the ECS-derived parameters between the two strains.

Differences in q_E Sensitivity Between Wt and *pgr5* Can Be Largely Attributed to Changes in g_{H^+} . The above flux estimates suggest differences in contributions to light-induced *pmf* from processes other than LEF, consistent with a difference in CEF1 engagement between Wt and *pgr5* (31, 32). However, the modest ($\approx 13\%$) decrease in ν_{H^+} in the absence of PGR5 was far too small to directly account for the corresponding 4- to 6-fold decrease in the q_E response at light-saturated LEF (Fig. 1A). In this regard, it was striking that the *pgr5* mutant exhibited lowered LEF without a corresponding increase in q_E sensitivity, in contrast to what was observed in the Wt upon lowering CO_2 (Fig. 1A).

Fig. 3 shows that g_{H^+} decreased in the Wt upon lowering CO_2 but remained similar or was substantially increased in *pgr5*, as is especially evident at the higher light intensities. Within the noise level, plots of q_E against pmf_{LEF} for Wt under the two CO_2 levels and *pgr5* overlapped (Fig. 4), indicating that, as was reported in refs. 17 and 18, changes in g_{H^+} could predominantly account for the differences in the q_E response. We thus conclude that in *pgr5* more facile proton efflux from the lumen through the ATP synthase, accompanied by decreases in LEF and probably CEF1, prevented the buildup of steady-state *pmf* and, thus, inhibited the q_E response.

In principle, g_{H^+} could be modulated by changing the specific activity of ATP synthase or its content in the thylakoids. Hence, a ≈ 2 -fold increase in the size of the ATP synthase pool could give rise to the observed ≈ 2 -fold increase (i.e., at higher light intensities) in g_{H^+} in *pgr5* (Fig. 3). However, ATP synthase content in Wt and *pgr5* was estimated by Western analyses and found to be essentially identical (Fig. 4 Inset). In addition, low light-induced activation of the ATP synthase by thioredoxin and leakage of the thylakoid membrane to protons were indistinguishable between Wt and *pgr5* (data not shown), essentially as seen for other C_3 plants (38). These data, taken together with the observed similarities in g_{H^+} at low light, lead us to conclude that the differences in g_{H^+} between Wt and *pgr5* were caused by alterations in steady-state substrate or affector concentrations (17).

The decrease in maximal LEF in *pgr5* is probably due to loss

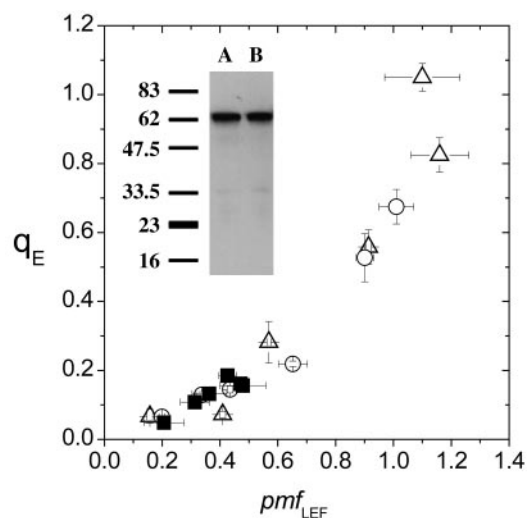


Fig. 4. The relationship between energy-dependent exciton quenching and the *pmf* generated solely by LEF. Estimates of energy-dependent quenching (q_E) and the *pmf* generated solely by LEF (i.e., pmf_{LEF}) were obtained as in Figs. 1 and 2, respectively. ATP synthase content in Wt (A) and *pgr5* (B) was estimated by Western blot analyses by using polyclonal serum directed against the β -subunit of the ATP synthase (Inset). Conditions and symbols are as in Fig. 1. Error bars represent SE for $n = 3-6$.

of PSI electron acceptors and a buildup of reduced intermediates (31, 32). A similar decrease in LEF was seen when CO_2 was lowered, but in contrast to the enhanced g_{H^+} that occurred in the absence of PGR5, such a decrease in LEF was accompanied by substantial decreases in g_{H^+} (Fig. 3), resulting in a net increase in both *pmf* and q_E . These results demonstrate an important role for ‘tuning’ the activity of the ATP synthase in the signal pathway that regulates light capture (39). Excessive turnover rates (i.e., large g_{H^+} values) will result in facile proton efflux, preventing buildup of *pmf* and diminishing the q_E response. On the other hand, inappropriate decreases in ATP synthase turnover rates can result in excessive buildup of *pmf*, over-acidifying the lumen and causing subsequent pH-induced degradation of the photosynthetic apparatus (4, 40).

From the above, we conclude that changes in CEF1 upon loss of PGR5 constitute a flux of protons approximately $<13\%$ of that from LEF, resulting in a commensurate decrease in ATP output. Because consumption of ATP and NADPH by the Calvin-Benson cycle is coupled, even a small ATP/NADPH imbalance could conceivably give rise to not only a buildup of ADP and $[\text{P}_i]$, but also a substantial reduction of NADP^+ , restricting the availability of PSI electron acceptors and, thereby, lowering LEF, as was observed in *pgr5* both here and in ref. 31.

Possible Causal Relation Between *Pgr5*⁻ and g_{H^+} . We proposed in ref. 17 that lowering CO_2 will lead to the buildup of phosphorylated metabolites in the stroma, depleting stromal $[\text{P}_i]$ below its K_M (≈ 1 mM) at the ATP synthase. This reaction will result in lowering of the effective g_{H^+} and subsequent increases in steady-state *pmf* and q_E . A small ATP/NADPH imbalance is expected to result from the absence of the PGR5-mediated CEF1. The deficit is obviously satisfied but only by substantially slower processes, e.g., alternative cyclic electron transfer processes or export of NADPH (12, 16). We thus expect in *pgr5* a buildup of stromal $[\text{P}_i]$ above its K_M at the ATP synthase, maintaining high g_{H^+} even when LEF is restricted. Thus, in this model, the loss of CEF1 in *pgr5* indirectly attenuates both steady-state *pmf* and q_E .

These results support a “division of labor” model for *pmf*

