Quantification of the electrochemical proton gradient and activation of ATP synthase in leaves

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ARTICLE INFO

Article history:
Received 4 February 2008
Received in revised form 31 March 2008
Accepted 4 April 2008
Available online 12 April 2008

Keywords:
ATP synthase
Electrochemical proton gradient
Membrane potential

ABSTRACT

We have developed a new method to quantify the transmembrane electrochemical proton gradient present in chloroplasts of dark-adapted leaves. When a leaf is illuminated by a short pulse of intense light, we observed that the light-induced membrane potential changes, measured by the difference of absorption (520 nm–546 nm), reach a maximum value (~190 mV) determined by ion leaks that occur above a threshold level of the electrochemical proton gradient. After the light-pulse, the decay of the membrane potential follows a multiphasic kinetics. A marked slowdown of the rate of membrane potential decay occurs ~100 ms after the light-pulse, which has been previously interpreted as reflecting the switch from an activated to an inactivated state of the ATP synthase (Junge, W., Rumberg, B. and Schröder, H., Eur. J. Biochem. 14 (1970) 575–581). This transition occurs at ~110 mV, thereby providing a second reference level. On this basis, we have estimated the ΔμH+ level that pre-exists in the dark. Depending upon the physiological state of the leaf, this level varies from 40 to 70 mV. In the dark, the ΔμH+ collapses upon addition of inhibitors of the respiratory chain, thus showing that it results from the hydrolysis of ATP of mitochondrial origin. Illumination of the leaf for a period longer than several seconds induces a long-lived ΔμH+ increase (~150 mV) that reflects the light-induced increase in ATP concentration. Following the illumination, ΔμH+ relaxes to its dark-adapted value according a multiphasic kinetics that is completed in more than 1 h. In mature leaf, the deactivation of the Benson–Calvin cycle follows similar kinetics as ΔμH+ decay, showing that its state of activation is mainly controlled by ATP concentration.

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1. Introduction

According to the chemiosmotic theory, ATP synthesis is driven by a transmembrane electrochemical proton gradient denoted as ΔμH+ [1,2]. In chloroplasts, ATP synthesis is promoted by a membrane-bound F0F1 ATP synthase that couples the transfer of H+ across the membrane to the synthesis of ATP. Conversely, it has been shown that, under aerobic conditions, this transmembrane potential has been ascribed to the hydrolysis of ATP of mitochondrial origin [6] evidencing the exchange of nucleotides between different cellular compartments, mitochondria, cytoplasm and chloroplasts. In this paper, we present new approaches based on the measurement of the membrane potential changes induced by a pulse of strong light to characterize the permanent ΔμH+ level in dark-adapted or pre-illuminated leaves.

Another issue addressed in this paper deals with the activation process of the ATP synthase in vivo. It is known that an electrochemical proton gradient is required to activate the ATP synthase in addition to supply energy for ATP synthesis [7,8]. The magnitude of this ΔμH+ is strongly decreased by the reduction of a disulfide bridge located in the γ subunit of the F1 complex [9–12]. In addition, Junge et al. evidenced, in isolated thylakoids, the necessity for a critical membrane potential to activate the ATP synthase [13,14]. These authors observed that after illumination by a series of flashes, the decay of the light-induced membrane potential is biphasic. They proposed that the break between the first phase (completed in ~100 ms) and the second one reflects a transition between a high and a low level for ATP synthase activity. Such a threshold potential for the activation of the ATP synthase has also been characterized in dark-adapted unicellular algae [15,16] and leaves [15,16]. Junesh and Gräber [12] and Strelow and Rumberg [17], have quantitatively analyzed the activation of ATP synthase in its oxidized or reduced form as a function of ΔpH. The half-maximum activation levels of the oxidized and reduced states of the ATP synthase are ΔpH = 2.2 and 3.4, respectively.

In this paper, we have analyzed the dependence of the activation state of the ATP synthase upon the membrane potential and ΔpH with dark-adapted or pre-illuminated leaves by measuring the kinetics of membrane potential decay following pulses of saturating light. On this basis, we propose a new method to estimate the ΔμH+ level that is in equilibrium with the phosphate potential in the dark.

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doi:10.1016/j.bbabio.2008.04.010
2. Materials and methods

Experiments were performed with young spinach leaves. When stated, the leaf was infiltrated under low pressure with 0.15 M sorbitol. During experiments, humid air was continuously flowed to the leaf.

Spectrophotometric and fluorescence measurements were performed using a new spectrophotometer (Biologic Science Instruments JTS10) in which absorption changes are converted to optical density changes (10-nm bandwidth) by light-emitting diodes (LED). The time resolution of the method is equal to the duration of the detecting flashes (12 µs). Continuous illumination was provided by LEDs peaking at 630 nm. Saturating actinic flashes were provided by a dye laser (630 nm) pumped by the second harmonic of a Nd:Yag laser (5-ns total duration). Membrane potential changes were measured by the difference of absorption (520 nm–546 nm). This difference eliminates the contributions of P700 and scattering changes. Owing to the low noise level of the method (ΔΔFI/10⁻³), a single experiment was performed at each wavelength. The extent of the field-indicating changes measured at 520 nm is proportional to the transmembrane electrical potential (reviewed in [18]). It has been explained on the grounds of a strong pre-existing polarization of the pigment probes. We have previously shown [19] that, in young spinach leaves, the concentration of PSII and PSI centers is about equal so that the absorption change measured 100 µs after a short saturating flash is proportional to the membrane potential increase associated with the transfer of two charges per photosynthetic chain. Membrane potential changes were calibrated with respect to the membrane potential increase induced by a single charge separation per photosynthetic chain (1 r. u.) that is typically equal to the threshold potential, ΔΔψleak that is determined by the extrapolation to time zero of the second phase of the decay kinetics (Fig. 1B, arrow) is independent of the light intensity (ΔΔψleak ~5.5 r. u.). The origin of the rapid ion transfers triggered above ΔΔψleak will be discussed in Section 3.7.

3.2. Determination of the membrane potential in dark-adapted leaves and light-induced activation of ATP synthase

As a working hypothesis, we consider that the ΔΔψleak level characterized in Section 3.1 provides a reference level for the transmembrane potential. In order to test the validity of this hypothesis, one must estimate the membrane potential that may pre-exist in the dark ΔΔψdark to which ΔΔψleak would be superimposed. The rate of charge recombination between PSII acceptor and donor measured in thylakoids in the presence of 3-(3,4-dichloro-phenyl)-1,1-dimethy lurea (DCMU) is known to increase with both the membrane potential and the proton gradient [23–25]. The kinetics of fluorescence decay induced by short flashes has been measured with dark-adapted Chlorella cells in the presence of DCMU [3]. The half time for the reoxidation of Qb is increased by a factor of 2 upon addition of tri-butylin, a specific inhibitor of ATP synthase. This was interpreted by assuming that a pool of ATP is present in the chloroplasts, the hydrolysis

![Fig. 1](https://example.com/fig1.png)

Fig. 1. Light-induced membrane potential changes for different light intensities. kiPSII – kiPSII ~7000, ~670 and ~330 s⁻¹ for curves 1–3, respectively. The leaf is dark-adapted for more than 2 h. A: Membrane potential increase measured during the light-pulse. B: Membrane potential decay measured from 10 to 20 ms (linear scale) after the light-pulse. Dashed line: Membrane potential decay after subtraction of phase 1; arrow: ΔΔψleak ~5.5 r. u. C: Membrane potential decay measured from 10 µs to 10 s (logarithmic scale).
of which induces a permanent electrochemical proton gradient. Bennoun [6] demonstrated that, in aerobic conditions, this pool of ATP is of mitochondrial origin. We performed similar experiments with leaves and measured the fluorescence decay following a 10-ns flash in the presence of 20 μM DCMU. To get fluorescence changes roughly proportional to $\Delta\mu$, we used non-saturating flashes hitting ~20% of PSII centers [3,26]. The half-times of the fluorescence decay were ~1.1 s in the control and ~2.5 s in the presence of a saturating concentration of uncouplers (2-μM nigericin +2-μM nonactin) that fully collapses $\Delta\mu_{\text{H}^+}$ pre-existing in the dark. This dependence of the rate of charge recombination on the addition of uncouplers shows that a membrane potential, likely associated with ATP hydrolysis, is present in dark-adapted leaves, as previously observed in algae [3]. To test the mitochondrial origin of the ATP pool accounting for this $\Delta\mu_{\text{H}^+}$, we repeated the above experiment in the presence of antimycin, an inhibitor of the cyt b/c complex of the respiratory chain. The half times of the fluorescence decay measured in the presence of 5-μM and 40-μM antimycin were ~2.06 and ~2.53 s, respectively. Thus, the addition of 40-μM antimycin, as well as saturating concentration of uncouplers fully collapses $\Delta\mu$ and $\Delta\mu_{\text{H}^+}$. It proves that, in the dark-adapted chloroplasts, ATP is of mitochondrial origin.

If indeed a $\Delta\mu$ component is present in the dark, the light-induced $\Delta\mu$ measured in Fig. 1 is superimposed to a “pedestal”, which can be collapsed by addition of antimycin. In Fig. 2, we measured the effect of antimycin on the kinetics of the membrane potential increase induced by a pulse of strong light ($k_{\text{PSI}}$ ~ $k_{\text{PSII}}$ ~ 5000 s $^{-1}$) given to a dark-adapted leaf. As shown in Fig. 2A, $\Delta\phi_{\text{max}}$ is larger in the presence of antimycin than in its absence. These differences in $\Delta\phi_{\text{max}}$ do not reflect changes in the rate of charge separation (see equal initial slopes for curves 1–3, Fig. 2A) but rather witness the decrease of the pedestal since we have shown that addition of antimycin collapses $\Delta\phi_{\text{dark}}$. Accordingly, we propose that $\Delta\phi_{\text{max}} = \Delta\phi_{\text{dark}} + \Delta\phi_{\text{max}}$. Moreover, as just discussed, in the presence of 40-μM antimycin (Fig. 2A, curve 4) we have $\Delta\phi_{\text{dark}} = 0$. In Fig. 2B, we have plotted curves 1–4 assuming equal $\Delta\phi_{\text{max}}$ Value. This allows the determination of $\Delta\phi_{\text{dark}}$ in the absence of antimycin, which is equal to 3.6 r. u. (dashed line 1) and suggests that the addition of increasing concentration of antimycin results in a decrease of $\Delta\phi_{\text{dark}}$ owing to a decrease of ATP concentration.

Previous reports showed that the membrane potential decay measured after the illumination of thylakoids [14] or leaf [16] by a series of short flashes displays a biphasic kinetics. The first phase completed in ~100 ms has been ascribed to proton transfer through ATP synthase that is associated with ATP synthesis and the slower phase, completed in ~1 s, to ions leaks through the membrane. The break between the two phases has been interpreted as reflecting the switch of the ATP synthase from an active to an inactive state that occurs below a threshold level of the membrane potential [14].

Fig. 2C shows the multiphasic decay of the membrane potential measured after the light is switched off. The first phase completed in ~15 ms we ascribe to rapid ion transfers through the membrane (Section 3.1) has not been observed in [15,16]. In these experiments, thylakoids or leaves were submitted to a series of short saturating flashes that induces a lower membrane potential increase than that induced by a pulse of saturating light as we used in Figs. 1 and 2. The lack of phase 1 in the decay kinetics reported in [15,16] is explained by the fact that the flash-induced membrane potential increase is lower than $\Delta\phi_{\text{leak}}$.

Decay kinetics shown Fig. 2C display a second phase (phase 2, completed in 100 ms) of equal amplitude for curves 1 to 4. In agreement with [14], we assume that this phase, is associated with proton transfer that leads to ATP synthesis. Interestingly, the extrapolation to time zero of phase 3, completed in 2 s, leads to an equal value of $\Delta\phi_{\text{thr}}$ level for curves 1–4 ($\Delta\phi_{\text{thr}}$ ~ 5.9 r. u.). Thus, we now define 3 membrane potential reference levels that do not depend upon antimycin concentration: $\Delta\phi_{\text{max}}$ ~ 11.6 r. u., $\Delta\phi_{\text{leak}}$ ~ 9.2 r. u. and $\Delta\phi_{\text{thr}}$ ~ 5.9 r. u.

In Fig. 2C, curves 1–3 (dashed lines), the permanent membrane potential $\Delta\phi_{\text{dark}}$ is lower than $\Delta\phi_{\text{max}}$ but not null. It shows that, below $\Delta\phi_{\text{thr}}$, ATP synthase remains able to slowly hydrolyze ATP and thus, able to generate a permanent membrane potential in contradiction with the conclusion drawn in [14]. Phase 3 is followed by phase 4, completed in 1 to 2 min, which we ascribe to the slow metabolic consumption of the ATP that has been synthesized during and after the pulse of saturating light. It should be kept in mind that under the conditions of the present experiments (i.e. dark-adapted leaves), the Benson–Calvin cycle is not activated so that the pool of ATP is not engaged into CO2 fixation. We propose that 5 μM-antimycin inhibits

![Fig. 2](image-url)
cyt b/c complexes but that the electron flow via complex I and alternative quinol-oxidases maintains a low level of ATP in the dark. Interestingly, the half time of phase 4 measured in the presence of 40-μM antimycin (~10 s), is significantly shorter than that measured in the presence of 2- and 5-μM antimycin (~22 s). This acceleration suggests that antimycin has a low uncoupling activity able to fully collapse Δω. Consistent with this hypothesis, the addition of saturating concentrations of other inhibitors of the respiratory chains as myxothiazol (inhibitor of the cyt b/c complex), or KCN (inhibitors of the cytochrome oxidase) induces similar effect than 5-μM antimycin (not shown).

The membrane potential change induced by a single-turnover flash that induces the transfer across the membrane of two charges per photosynthetic chain has been estimated to ~40 mV [18,27,28] (1 r. u. ~20 mV). In the following, we have used this value to calibrate membrane potential changes. In the presence of 40-μM antimycin (Fig. 2B, curve 4), we have Δφ̂max ~11.5 r. u. and Δφ̂leak ~9.6 r. u., which corresponds to ~230 and ~192 mV, respectively. These values are in reasonable agreement with the estimate of the maximum potential an artificial or biological membrane can sustain (~200 mV, [29,30]). Further, we have Δφ̂leak ~5.9×20 ~98 mV and Δφ̂dark ~3.7×20 ~74 mV (dashed line 1, Fig. 2B).

3.3. Equivalence of chemical and electrical potential for the activation of ATP synthase

In the following, we will attempt to address the question as to whether the threshold levels for ATP synthase activity and for ion leaks depend exclusively on Δφ or rather on the electrochemical proton gradient Δμ̂H+. We quote the chemical potential associated with Δμ̂H+ as Δμ̂H+=0.06 ΔpH and we have Δμ̂H+ = Δμ̂H+ + Δμ̂pH.

In Fig. 3A, the leaf has been infiltrated with 0.6-μM nigericin to rapidly convert ΔpH into Δω. Addition of nigericin induces a decrease in ΔΔμ̂H+, and thus in ΔΔμ̂pH, levels (not shown). Curve 1 shows the kinetics of the membrane potential decay measured after a 12-ms light-pulse given to a dark-adapted leaf. In a second experiment, the same leaf has been first illuminated by a 200-μs light-pulse that induces a Δω increase that decays in several minutes (t1/2 ~1 min). We assume that Δω is in thermodynamic equilibrium with the phosphate potential for time longer than 10 s. It implies that the Δω decay is associated with the metabolic consumption of the ATP synthesized during the 200-ms light-pulse. In Fig. 3A, the red arrow indicates the membrane potential measured 26 s after the 200-ms pulse (ΔΔω ~1.31). Curve 2 shows the membrane potential decay measured after a 12-ms light-pulse given 26 s after the 200-ms pulse. Fig. 3A insert shows that the initial rate of the membrane potential decay and the amplitude of phase 1 are lower with the pre-illuminated leaf (curve 2) than with the dark-adapted leaf (curve 1). It implies that the rate of the charge separation at the end of the 12-ms light-pulse is lower for curve 2 than for curve 1. We have previously shown [18] that a large fraction of the PQ pool is still reduced 26 s after the 200-ms light-pulse. Thus, one expects that, at end of the 12-ms pulse, PSII turnover and the rate of charge separation are slower for curve 2 than for curve 1, which explains the lower amplitude of phase 1. Beyond 15-ms dark, curves 1 and 2 are identical, as are curves 1–3 of Fig. 1B. Consequently, both ΔΔωleak (~8 r. u.) and ΔΔωphie (~4.35) are the same for curves 1 and 2, Fig. 3A. We have checked that for dark times longer than 26 s between the 12-ms and the 200-ms light-pulses, phase 2 remains identical to that measured with the dark-adapted material.

Fig. 3B shows the same experiment as Fig. 3A but performed in the absence of nigericin and in the presence of 3-μM antimycin. Antimycin concentration was adjusted to get the same ΔΔμ̂H+ level that measured in the presence of nigericin (Fig. 3A and B, curves 1). The membrane potential measured 26 s after the 200-ms pulse (ΔΔω ~0.67, Fig. 3B, red arrow) is about half than that measured in the presence of nigericin (~1.31), suggesting that the Δμ̂H+ component in the dark is not null. Moreover, contrary to what is observed in the presence of nigericin, curve 2 (pre-illuminated leaf) is shifted toward lower membrane potential values with respect to curve 1 (dark-adapted leaf). This downshift (0.64 r. u.) is consistent with the following hypotheses: 1) in the absence of nigericin, the hydrolysis of ATP generated by the 200-ms pulse induces the formation of a proton gradient superimposed to a membrane potential increase; 2) the threshold level for ATP synthase activity is set by the electrochemical proton gradient rather than by the sole membrane potential; and 3) in the presence of either nigericin or antimycin, the 200-ms light-pulse induces the synthesis of the same amount of ATP that is associated with the same increase in the electrochemical proton gradient. In the presence of nigericin, we have: ΔΔμ̂H+ = ΔΔω ~1.31 r. u. In the presence of antimycin, ΔΔμ̂H+ ~0.64 r. u. equal to the downshift between curves 1 and 2, Fig. 3B and of the membrane potential ΔΔω = ~0.67 r. u. measured 26 s...
after the 200-ms pulse is given. It leads to \( \Delta \mu_H^\text{PSII} \approx 0.64 \mu \text{V} \), a value equal to that measured in the presence of nigericin, in agreement with hypothesis 3. Since \( \Delta \mu_H^\text{Cytb} \) and \( \Delta \mu_H^\text{leak} \) undergo similar downshifts after the 200-ms light-pulse (Fig. 3B and insert), we conclude that the threshold levels for ion leaks and for ATP synthase activity are both controlled by the electrochemical proton gradient.

We have proposed that the long-lived \( \Delta \mu_H^\text{PSII} \) increase induced by the 200-ms light-pulse reflects the light-induced increase in ATP concentration. In Fig. 4, using the same procedure as in Fig. 3, we have measured \( \Delta \mu_H^\text{PSII} \) increase as a function of the duration of the light-pulse. As just discussed, \( \Delta \mu_H \) was measured 30 s after the illumination and \( \Delta \mu_H^\text{PSII} \) by measuring the downshifts of \( \Delta \mu_H^\text{Cytb} \) and \( \Delta \mu_H^\text{leak} \). In Fig. 4, curves 1–3 display \( \Delta \mu_H \), \( \Delta \mu_H^\text{PSII} \) and \( \Delta \mu_H^\text{leak} \), respectively, as a function of the light-pulse duration. \( \Delta \mu_H^\text{PSII} \) displays a multi phasic kinetics with a first phase completed in 40–50 ms, followed by a ~200 ms phase. This kinetics presents similarity with the multiphasic fluorescence increase induced by a high-light illumination of a dark-adapted leaf [21,31] denoted phases OJP [32,33]. The O and J phases, completed in ~40 ms [21,31–33], were associated with the reduction of most of the PSII electron acceptors whereas the IP phase, completed in ~200 ms, was ascribed to the reduction of PSI electron acceptors [22]. Thus, the kinetics of fluorescence and \( \Delta \mu_H^\text{PSII} \) increase both reflect the number of charges transferred through the membrane which is roughly proportional to the number of ATP synthesized.

### 3.4. Life-time of the light-induced electrochemical proton gradient

In Fig. 5, the decay of \( \Delta \mu_H^\text{PSII} \) induced by a 12-s illumination has been measured, using a similar procedure as in Fig. 2. In Fig. 5A, the black curve displays the kinetics of \( \Delta \mu_H^\text{PSII} \) increase induced by a 12-ms pulse given to a leaf dark-adapted for more than 3 h. Following a 12-s illumination, short pulses are given at dark times as indicated on Fig. 5. The duration of each of the light-pulses has been adjusted to reach the maximum level \( \Delta \mu_H^\text{PSII} \text{max} \). Owing to the small number of charges transferred and to the high-buffering capacitance of the lumen, we assume that, during each of these short light-pulses, \( \Delta \mu_H^\text{PSII} \) closely reflects \( \Delta \mu_H \). The first pulse is given after 45-s dark, a time sufficient to allow the full oxidation of the PSI acceptors and a partial oxidation of the PQ pool. Fig. 5B shows the kinetics of \( \Delta \mu_H^\text{PSII} \) decay measured after each pulse. Curves 1–8 display a first phase completed in ~15 ms (phase 1). As already discussed (Section 3.3), both the initial decay rate and the amplitude of phase 1 are lower for the shorter dark periods owing to the fact that the PQ pool is still partially reduced. Solid lines in Fig. 5B display the initial part of phase 2 after subtraction of phase 1. As in Figs. 1–3, \( \Delta \mu_H^\text{PSII} \text{max} \) level is estimated by extrapolation to time zero of phase 2.

The data shown in Fig. 2 evidenced that addition of antimycin induced a \( \Delta \mu_H^\text{PSII} \text{max} \) increase that reflects a decrease of the \( \Delta \mu_H^\text{NAP} \) pedestal. We thus assume that in Fig. 5B, the lower \( \Delta \mu_H^\text{PSII} \text{max} \) and \( \Delta \mu_H^\text{PSII} \text{leak} \) levels reflect a larger \( \Delta \mu_H^\text{PSII} \text{pedestal} \). In Fig. 5C, we have plotted for each of the probing pulses the kinetics of \( \Delta \mu_H^\text{PSII} \) decay after subtraction of phase 1 assuming equal value for \( \Delta \mu_H^\text{PSII} \text{pedestal} \) as determined in Fig. 5B. As the \( \Delta \mu_H^\text{PSII} \text{pedestal} \) is not known, we have plotted the difference \( \Delta \mu_H^\text{PSII} \text{pedestal} = \Delta \mu_H^\text{PSII} \text{dark} \). Dashed lines show the \( \Delta \mu_H^\text{PSII} \text{dark} \) level measured at the times each of the probing pulses is given. This level decreases when the dark time increases. We ascribe the light-induced \( \Delta \mu_H^\text{PSII} \) increase to the ATP synthesis induced by the efficient cyclic electron flow that operates during the 12-s illumination [19,34]. During the subsequent dark period, \( \Delta \mu_H^\text{PSII} \) and ATP concentration slowly decrease to their dark-adapted values. For dark times longer than 15-min, the amplitude and time-course of phase 2 stay constant while the amplitude of phase 3 (completed in ~4-s) increases with the time of dark adaptation. Thus, beyond 15-min dark adaptation, the electrical activity of the luminal membrane decreases.

### Fig. 5

Electrochemical proton gradient induced by a short light-pulse (kiPSII – kIleak ~5000 s⁻¹) given at various dark times after a 12-s illumination (kiPSII – kIleak ~400 s⁻¹). Dark times as indicated in panels A and C. A: \( \Delta \mu_H^\text{PSII} \) increase measured during the short light-pulse. B: \( \Delta \mu_H^\text{PSII} \) decay measured after the short light-pulse. Solid lines: phase 2 (see text). C: \( \Delta \mu_H^\text{PSII} = \Delta \mu_H^\text{PSII}\text{dark} \) decay kinetics (phases 2 and 3) measured after the short light-pulses plotted assuming the same value for \( \Delta \mu_H^\text{PSII}\text{dark} \). Dashed lines: \( \Delta \mu_H^\text{PSII} = \Delta \mu_H^\text{PSII}\text{dark} \) measured at the time the short light-pulses are given. Dotted lines: extrapolation to time zero of phase 3, giving the threshold level for ATPase activation (\( \Delta \mu_H^\text{PSII} = \Delta \mu_H^\text{PSII}\text{dark} \) ~2.4 r. u.).
time, the curves display equal threshold level for ATP synthase activation ($\Delta \mu_{H^+}^{ther} - \Delta \mu_{H^+}^{dark} - 2.4$). For dark times up to 15-min, $\Delta \mu_{H^+}^{ther} - \Delta \mu_{H^+}^{dark}$ measured at the time the probing pulses are given is larger than $\Delta \mu_{H^+}^{ther}$. Under these conditions, ATP synthase is already in its activated state at the time the light-pulse is given and most of the $\Delta \mu_{H^+}^{ther}$ decay is completed in ~100 ms (phase 2). At the time the first probing pulse is given (45 s after the 12-s illumination), we have $\Delta \mu_{H^+}^{ther} - \Delta \mu_{H^+}^{ther} = 1.5$ r. u. According to Fig. 2C, the absolute $\Delta \mu_{H^+}^{ther}$ level is ~5.9 r. u. Thus, in Fig. 5, the absolute $\Delta \mu_{H^+}^{ther}$ level measured 45 s after the 12-s illumination is $\Delta \mu_{H^+}^{ther} = 1.5 - 5.9 = -7.4$ r. u. = -150 mV.

In Fig. 6, $\Delta \mu_{H^+}^{ther} - \Delta \mu_{H^+}^{dark}$ decay has been measured as a function of the dark time following an illumination by giving a series of probing pulses as previously described (Fig. 5C, dashed lines). Curves 1–3 show the $\Delta \mu_{H^+}^{ther} - \Delta \mu_{H^+}^{dark}$ decay after 6-s, 30-s and 10-min illumination, respectively. The extrapolation to time zero of curves 1–2 is $\Delta \mu_{H^+}^{ther} - \Delta \mu_{H^+}^{dark} = 4.1$ r. u., a value close to that measured in Fig. 5C after 45-s dark (3.9 r. u.). The same value is obtained for times of illumination from 6-s to 2-min (not shown). In Fig. 6, curve 3, the time of illumination (10-min) is long enough to fully activate the Benson–Calvin cycle. During the first minute in the dark $\Delta \mu_{H^+}^{ther}$ is slightly lower than that measured in curves 1–2. It may reflect a lower ATP concentration owing to its fast consumption by the Benson–Calvin cycle. Curves 1–3 display a lag phase, the duration of which increases with the time of illumination. This lag phase can be tentatively ascribed to the buffering capacitance associated with the storage of high-energy phosphate bounds present in the numerous compounds, the synthesis of which are coupled to ATP hydrolysis including those involved in the Benson and Calvin cycle. The $\Delta \mu_{H^+}^{ther} - \Delta \mu_{H^+}^{dark}$ decay that follows the lag phase displays two phases completed in ~15 min and ~1 h. The amplitude of the slower phase increases as a function of the time of illumination to reach a maximum value after ~10 min (Fig. 6, curve 3, ~25% of the total amplitude). This phase could reflect a slow equilibration between the chloroplast and the cytoplasm compartments.

3.5. Control of the activity of the Benson–Calvin cycle by ATP

Three molecules of ATP and two molecules of NADPH supply the energy and reductive power required for the assimilation of one CO₂ molecule. On the other hand, the activity of several enzymes of the Benson–Calvin cycle are in their reduced (active) state mainly controlled by ATP concentration. The absence of a deactivation phase in a few min time range suggest that the redox-regulated enzymes of the Benson–Calvin cycle are in their reduced (active) state in dark-adapted and pre-illuminated leaves as well. In the presence of 10-μM myxothiazol, all the cell compartments are deprived in ATP and we observed that short periods of illumination (~30-s) induce no long-lived $\Delta \mu_{H^+}^{ther}$ increase (not shown). It suggests that in these conditions, ATP synthesized by the cyclic electron flow is rapidly consumed in the different cell compartments that lack ATP.

3.6. Phosphate potential and ATP concentration

If one assumes that the membrane is in thermodynamic equilibrium, we have:

$$\Delta \mu_{H^+} = 60/n \times (\log K + \log (R/P))$$

(1)

in which $R$ is equal to the [ATP]/[ADP] ratio, $n$ is the number of protons translocated through the ATP synthase per ATP synthesized and $\Delta \mu_{H^+}$ is expressed in mV. According to Turina et al. [35], the $H^+$/ATP ratio is $n = 4$. From the value of the Gibbs free energy for ATP synthesis [35,36] and assuming a stromal pH between 7 and 8, we estimate $\Delta \mu_{H^+} = 2 \times 10^5$. On this basis, we have:

$$\Delta \mu_{H^+} = 80 + 15 \times \log (R/P).$$

(2)
As computed from Fig. 5, $\Delta \alpha_{\text{H}}$ level measured 45 s after the 12-s illumination is $-150$ mV, that leads to $R/[P]-34000$. If the membrane is in thermodynamic equilibrium and assuming $[P]=1$ mM, it would correspond to a large $[\text{ATP}]/[\text{ADP}]$ ratio $(R \approx 45)$. It suggests that, when a leaf is illuminated for more than 5 s, most of the ADP has been converted to ATP. Generation of ATP is associated with an efficient cyclic process that is operating during the first min of illumination of a dark-adapted leaf [19, 34].

In a dark-adapted leaf (Fig. 2C, curve 1), $\Delta \alpha_{\text{H}}$ has been estimated to $\approx 74$ mV. Depending upon the physiological state of the above leaves, $\Delta \alpha_{\text{H}}$ increases (proton slip). A third hypothesis is to assume that phase 1 synthase, leading to its transitory inactivation.

A membrane potential level has reached its maximum value likely limited by the rate of the rotation of the electrical rotor $F_0$ still increasing (proton slip). A third hypothesis is to assume that phase 1 and phase 2 are both coupled to ATP synthesis. It would imply that above $\Delta \alpha_{\text{H}}$, ATP synthase activity suddenly increases and ATP synthase activity would be controlled by two threshold levels. This hypothesis appears quite unlikely.

3.7. Origin of rapid ion transfers associated with phase 1

As shown in Sections 1 and 2, rapid ion transfers are triggered above a membrane potential level $\Delta \alpha_{\text{H}}$ varying between 30 mM and 80 mM. For $\Delta \alpha_{\text{H}}$ limited to 50 mV, we have: $R/[P] = 10^{-2}$, which corresponds to $[\text{ATP}]/[\text{ADP}]=5 \times 10^{-5}$ assuming $[P]=5$ mM. One expects that, at such a low ATP concentration and under conditions in which ATP synthase is mainly inactivated, Eq. (2) is not satisfied owing to passive ion leaks. It suggests that, in dark-adapted leaves, ATP concentration is larger than that computed from Eq. (2).

3.8. Regulation of ATP synthase activity

Junesh and Gräber [12] and Strelow and Rumberg [17] have measured in vitro the $\Delta \alpha_{\text{H}}$ that induces the half activation of reduced ATP synthase. Half activation occurs at $\Delta \alpha_{\text{H}} \approx 190$ mV. The origin of these fast ion transfers is presently unknown. A possible hypothesis is that the opening of which would be triggered above a given membrane potential. As an alternative hypothesis, one may assume that, at the level of the $F_1$ complex has reached its maximum value likely limited by the rate of the rotation of the electrical rotor $F_0$ still increasing (proton slip). A third hypothesis is to assume that phase 1 and phase 2 are both coupled to ATP synthesis. It would imply that above $\Delta \alpha_{\text{H}}$, ATP synthase activity suddenly increases and ATP synthase activity would be controlled by two threshold levels.

Acknowledgement

The authors thank Fabrice Rappaport for the valuable discussions and great help in the preparation of this manuscript.

References


