

## Research



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# On the origin of a slowly reversible fluorescence decay component in the *Arabidopsis npq4* mutant

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Over-excitation of photosynthetic apparatus causing photoinhibition is counteracted by non-photochemical quenching (NPQ) of chlorophyll fluorescence, dissipating excess absorbed energy into heat. The PsbS protein plays a key role in this process, thus making the PsbS-less *npq4* mutant unable to carry out qE, the major and most rapid component of NPQ. It was proposed that *npq4* does perform qE-type quenching, although at lower rate than WT *Arabidopsis*. Here, we investigated the kinetics of NPQ in PsbS-depleted mutants of *Arabidopsis*. We show that red light was less effective than white light in decreasing maximal fluorescence in *npq4* mutants. Also, the kinetics of fluorescence dark recovery included a decay component, qM, exhibiting the same amplitude and half-life in both WT and *npq4* mutants. This component was uncoupler-sensitive and unaffected by photosystem II repair or mitochondrial ATP synthesis inhibitors. Targeted reverse genetic analysis showed that traits affecting composition of the photosynthetic apparatus, carotenoid biosynthesis and state transitions did not affect qM. This was depleted in the *npq4phot2* mutant which is impaired in chloroplast photorelocation, implying that fluorescence decay, previously described as a quenching component in *npq4* is, in fact, the result of decreased photon absorption caused by chloroplast relocation rather than a change in the activity of quenching reactions.

## 1. Introduction

Plants use light as an energy source for their metabolism: solar energy is absorbed, and excitons rapidly transferred to the photosynthetic reaction centres by the light-harvesting complexes (LHC) of each photosystem (PS). These bind a large array of chlorophylls (Chls) that are tightly connected to yield more than 80% quantum efficiency of photochemical reactions [1]. Under stable light conditions, plants optimize the efficiency of photosynthetic machinery and yet light intensity changes during the day and the rapid fluctuations imposed by shading in the canopy or by clouds, easily result in over-excitation. Thus, quantum efficiency needs to be rapidly downregulated to decrease the probability of Chl *a* triplet (<sup>3</sup>Chl\*) and singlet oxygen formation [2,3]. This is performed by photoprotection mechanisms including leaf and chloroplast avoidance movement, reactive oxygen species (ROS) scavenging, and quenching of triplet and singlet Chl excited states [4–6]. The ability of plants to modulate light utilization efficiency in fluctuating light is crucial for plant fitness [7]. A major role for prevention of over-excitation is played by a set of inducible mechanisms referred to as non-photochemical quenching (NPQ) [8] that are triggered by a feedback loop in which excess light (EL) induces lumenal acidification, detected by the thylakoid protein PsbS, which in turn triggers qE, the most rapid component of NPQ, leading to dissipation of excess energy with a half-life of 1–2 min [5,9]. As PsbS is not a Chl-binding protein [10], its effect on Chl fluorescence must be achieved through interaction with the antenna system binding the xanthophylls zeaxanthin (Zea) and lutein (Lut) [11], in the absence of which quenching does not occur [12].

Besides qE, NPQ includes a slowly relaxing component ( $\tau > 60$  min) qI, which is independent of lumenal pH and has been attributed to inactive PSII

centres produced by EL stress [13]. Additional quenching components with intermediate half-lives ( $\tau = 10\text{--}15$  min) were originally attributed to state 1–state 2 transitions [8], and more recently to Zea binding to the LHC proteins, hence the names qT or qZ [14,15].

Two types of mechanism have been proposed for activation of quenching by PsbS: the first proposes a direct interaction of PsbS with a neighbour antenna protein, either LHCII or a monomeric complex, which causes a conformational change activating quenching site(s) within the antenna subunit itself [16,17] or through the trapping of a Zea molecule at the PsbS–LHCII interface [11]. The first type of mechanism relies on the notion that LHC proteins exist in two conformations with different fluorescence lifetimes [18], whose interconversion is controlled by changes in protein–protein interactions in the membrane, which are promoted by activation of PsbS [19,20]. The recent report that *npq4* plants lacking PsbS are competent in quenching, although longer exposure to EL is required than in WT plants [21], supports the first hypothesis.

In this work, we have studied the properties of light-induced fluorescence decrease in the *npq4* mutant, which develops a slow fluorescence decay. A component, qM, with the same amplitude and half-life, could also be deconvoluted from the kinetics of WT plants, was uncoupler-sensitive and unaffected by treatments inhibiting PSII repair or mitochondrial ATP production. Also it was induced by white light but not by red light. A targeted reverse genetic analysis showed that the *npq4phot2* double mutant which was impaired in chloroplast avoidance, was devoid of qM. On this basis, we propose that the fluorescence decay previously described as a quenching component in *npq4* is, in fact, the result of decreased photon absorption caused by chloroplast relocation rather than by a change in the activity of quenching reactions. This finding supports a direct role of PsbS in triggering the quenching reactions.

## 2. Experimental procedures

### (a) Plant material

*Arabidopsis thaliana* T-DNA insertion mutants (Col-0) *npq1* (At1G08550) and *npq2* (At5G67030) were a kind gift of K.K. Niyogi (University of California at Berkeley). Mutant *lut2* (At5G57030) was obtained from the NASC collection, Salk line 005018. *koLhcb4*, *koLhcb5* and *koLhcb6* were obtained as described in [22,23]. Mutants *npq4* and *npq4chl1* were a kind gift of K. K. Niyogi, *stn7npq4* was provided by E.-M. Aro (University of Turku, Finland), and *phot2* by M. Wada (Kyushu University, Japan). Double mutants were obtained by crossing single mutant plants and selecting progeny either by pigment analysis, western blotting [22,23] or by the light-induced change in the green colour of leaf blades [24]. WT and mutant plants were grown on compost in a growth chamber for five weeks under controlled conditions (150  $\mu\text{mol photons m}^{-2} \text{s}^{-1}$ , 23°C, 8 L/16 D cycle, 70% relative humidity).

### (b) *In vivo* fluorescence and non-photochemical quenching measurements

NPQ of Chl fluorescence was measured on leaves at room temperature (RT; 23°C) with a PAM 101 fluorometer (Walz, Germany). NPQ was calculated according to Van Kooten & Snel [25]. When red actinic light was used, the light intensities for these experiments were chosen in order to produce the

same value of qL in all genotypes. When indicated, fluorescence was measured on detached leaves infiltrated with 150 mM sorbitol containing either 50  $\mu\text{M}$  nigericin [26], 100  $\mu\text{M}$  lincomycin [27] or 2  $\mu\text{M}$  myxothiazol [28].

### (c) Pigment analysis

Pigments were extracted from leaf discs using 85% acetone buffered with  $\text{Na}_2\text{CO}_3$ . Separation and quantification of pigments were performed by HPLC [29].

### (d) Measurement of chloroplast movement

Chloroplast avoidance response was induced in leaves by EL treatment. Before measurements, plants were adapted in darkness for 1 h, then detached leaves on wet paper were exposed to 400  $\mu\text{mol photons m}^{-2} \text{s}^{-1}$ , white light at RT for 1 h. Distribution of chloroplasts in the mesophyll cells was determined by light microscopy. To take micrographs, we introduced the solution (150 mM sorbitol  $\pm$  inhibitors) into intracellular spaces under weak negative pressure before EL treatment, and removed the upper epidermis from the leaves just before mounting the microscope slide.

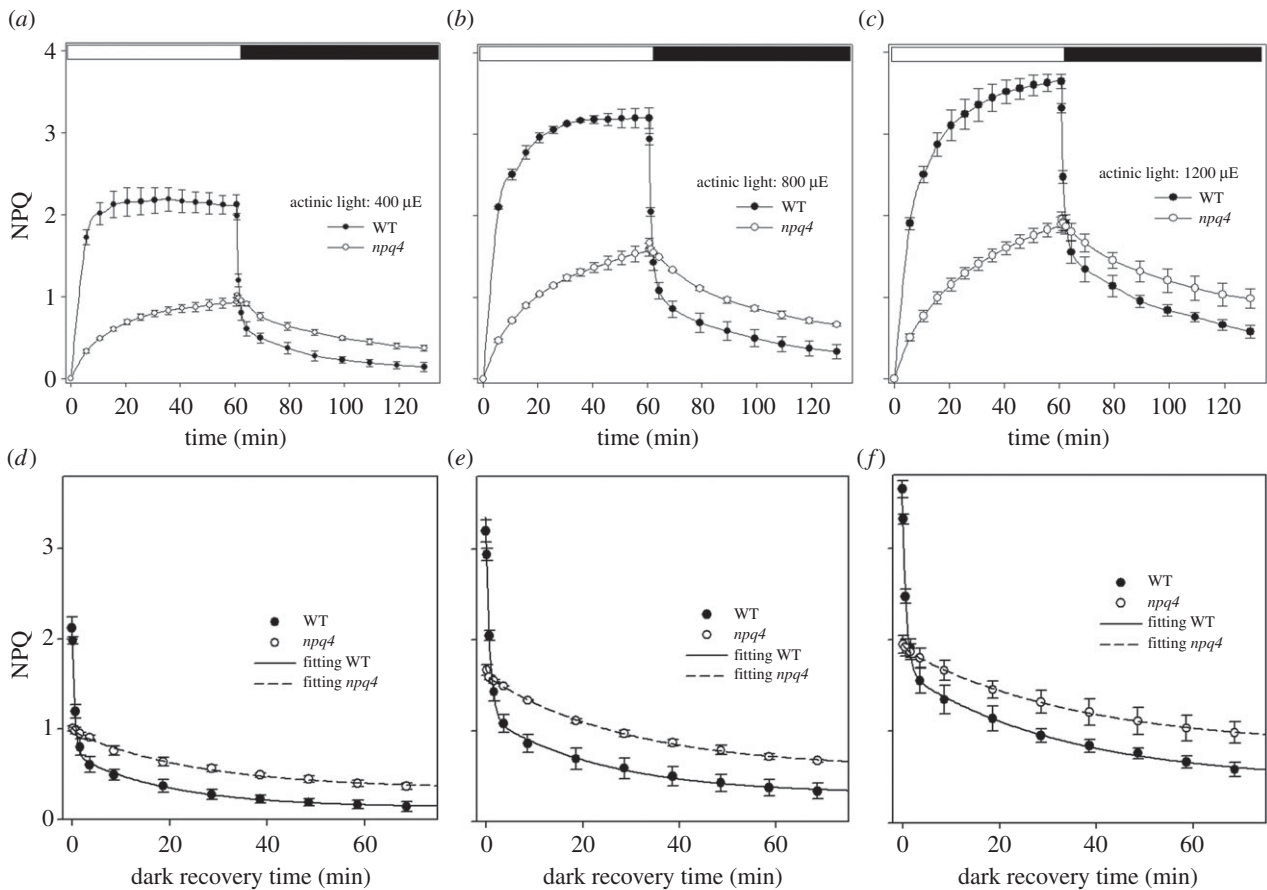
## 3. Results

### (a) Kinetics of non-photochemical quenching induction and relaxation in *npq4* versus wild-type leaves

Plants lacking PsbS are completely devoid of the fast quenching phase qE when illuminated for 8 min at 1200  $\mu\text{mol photons m}^{-2} \text{s}^{-1}$  (see electronic supplementary material, figure S1a), consistent with literature data [30]; the fluorescence quenching reached a maximum value four times lower than in WT plants. Dark relaxation of fluorescence quenching was clearly different between genotypes: *npq4* showed almost no relaxation, while WT recovered the most fluorescence quenching. Upon exposure to longer periods of actinic light, the quenching behaviour of *npq4* mutants showed a relative increase, reaching an NPQ value of 1.9, thus 45% with respect to WT plants (see electronic supplementary material, figure S1b). A major change was observed in the fluorescence recovery of *npq4* mutants in the dark, which showed a higher rate, although still slower than in WT plants. Thus, although the rapidly relaxing phase of the dominant qE component was missing in *npq4* mutants, a slowly reversible phase contributed to relaxation recovery, not only in WT plants but also in *npq4* mutants. Yet, the dark recovery was slower than in WT plants, yielding a higher qI.

The amplitude and kinetics of NPQ induction and relaxation were previously shown to be related to the activity of the xanthophyll cycle in EL [31]. We thus determined for Zea the extent of both EL-formation and dark-reconversion under the same experimental condition used to follow NPQ kinetics. Results in table S1 available in the electronic supplementary material show that the Zea content was the same in WT and *npq4* leaves for each condition. Both genotypes retained approximately 50% of the Zea produced upon in EL, upon 1 h dark recovery.

The NPQ kinetics of WT and *npq4* leaves were measured at different actinic intensities (figure 1a–c), showing similar behaviour of *npq4* leaves at all the light intensities used, including the rise in the light to half the amplitude of the WT leaves followed by a slow dark recovery.



**Figure 1.** Kinetic deconvolution of NPQ dark relaxation of WT and *npq4* leaves. The dark recovery of NPQ was measured in intact leaves of *Arabidopsis* WT and *npq4* mutants upon illumination for 60 min with different intensities of white actinic light: (a) 400, (b) 800, (c) 1200  $\mu\text{mol photons m}^{-2} \text{s}^{-1}$ . Symbols and error bars show mean  $\pm$  s.d. ( $n = 3$ ). Each dataset was fitted with a bi-exponential function  $\text{NPQ} = A_{qI} + A_{qE} e^{(-t/\tau_{qE})} + A_{qM} e^{(-t/\tau_{qM})}$  (d–f) and the kinetics of qE and qM relaxation were assessed in the different genotypes (table 1).

**Table 1.** Kinetics of NPQ dark relaxation in WT and *npq4* leaves. The dark-recovery of NPQ was measured in intact leaves of *Arabidopsis* WT and *npq4* mutants upon 60 min illumination with different actinic intensities (400, 800 and 1200  $\mu\text{mol photons m}^{-2} \text{s}^{-1}$ , white light, RT). Each dataset was fitted with a bi-exponential function  $\text{NPQ} = A_{qI} + A_{qE} e^{(-t/\tau_{qE})} + A_{qM} e^{(-t/\tau_{qM})}$  and the kinetics of qE and qM relaxation were assessed in the different genotypes by comparing amplitudes of parameter A, which describes the slope of exponential functions. Significantly different values (Student's *t*-test) with respect to WT are marked with asterisks.

actinic intensity ( $\mu\text{mol photons m}^{-2} \text{s}^{-1}$ )	genotype	$\tau_{qE}$ (min)	$A_{qE}$	$\tau_{qM}$ (min)	$A_{qM}$	$A_{qI}$
400	WT	$0.56 \pm 0.02$	$1.26 \pm 0.05$	$19.9 \pm 4.4$	$0.58 \pm 0.04$	$0.14 \pm 0.13$
	<i>npq4</i>	—	—	$25.7 \pm 3.9$	$0.63 \pm 0.03$	$0.39 \pm 0.07^*$
800	WT	$0.83 \pm 0.04$	$1.73 \pm 0.08$	$24.3 \pm 6.8$	$0.85 \pm 0.07$	$0.33 \pm 0.09$
	<i>npq4</i>	—	—	$29.3 \pm 2.2$	$1.02 \pm 0.03^*$	$0.67 \pm 0.02^*$
1200	WT	$0.81 \pm 0.04$	$1.65 \pm 0.08$	$33.3 \pm 8.4$	$1.20 \pm 0.10$	$0.58 \pm 0.08$
	<i>npq4</i>	—	—	$35.8 \pm 12.8$	$1.07 \pm 0.13$	$1.01 \pm 0.13^*$

The kinetic analysis of NPQ dark relaxation is frequently used as a tool for the characterization of quenching dynamics [15,32]. Thus, NPQ relaxation kinetics (figure 1d–f) were fitted with a bi-exponential decay function:  $\text{NPQ} = A_1 e^{(-t/\tau_1)} + A_2 e^{(-t/\tau_2)} + A_3$ . In WT plants, a fast decay component ( $\tau_1$ , half-life of 35–50 s) represented the dominant component of NPQ under all light conditions (around 60% of total quenching), and could reasonably be attributed to qE. The residual quenching at the end of dark recovery ( $A_3$ ) was due to processes that were essentially irreversible within the time range of the experiment ( $\tau > 60$  min), fitting with the

characteristics of the photo-inhibitory quenching, qI [33]. In *npq4* plants, the kinetics of NPQ relaxation were slowed down with respect to WT plants: a satisfactory fitting of NPQ dark relaxation could only be obtained with the introduction of an intermediate component with a decay rate intermediate between those of qE and qI ( $\tau_2$ , half-life of 20–35 min). This component also improved the fitting of WT curves where it contributes to about 30% of total NPQ (figure 1d–f and table 1). From now on this component will be referred to as qM. The rapid (qE) kinetic component was missing in *npq4* mutants under all experimental conditions explored in this work. According to the

**Table 2.** Kinetics of qM dark relaxation in *npq4* leaves. The kinetic components qM and qI were deconvolved from dark recovery of NPQ in *npq4* leaves. Dark-adapted leaves were exposed to white actinic light for 60 min at  $400 \mu\text{mol photons m}^{-2} \text{s}^{-1}$ , RT, following 60 min of dark recovery. (upper section) Before NPQ induction, leaves were vacuum-infiltrated with 150 mM sorbitol and either 50  $\mu\text{M}$  nigericin (uncoupler, collapsed the  $\Delta\text{pH}$  across the thylakoid membranes), 2  $\mu\text{M}$  myxothiazol (respiratory chain inhibitor) or 100  $\mu\text{M}$  lincomycin (chloroplast protein biosynthesis inhibitor). (middle section) The kinetic components qM and qI were measured in *npq4* double mutants lacking zeaxanthin (*npq4npq1*) or lacking lutein (*npq4lut2*). (lower section) Components qM and qI were measured in *npq4* double mutants depleted of Lhcb subunits CP26 and CP24 (*npq4koLhcb5/6*), CP29 and CP24 (*npq4koLhcb4/6*), lack the entire LHC (*npq4ch1*), or unable to activate state transition (*npq4stn7*) or chloroplast avoidance movement (*npq4phot2*). Each dataset was fitted with an exponential function  $\text{NPQ} = A_{\text{qI}} + A_{\text{qM}} e^{(-t/\tau_{\text{qM}})}$  and the kinetics of qM relaxation were assessed in the different samples by comparing amplitudes of parameters A. Significantly different values (Student's *t*-test) with respect to WT are marked with asterisks.

	$\tau_{\text{qM}}$ (min)	$A_{\text{qM}}$	$A_{\text{qI}}$
<i>npq4</i>	$28.4 \pm 4.5$	$0.63 \pm 0.03$	$0.35 \pm 0.03$
<i>npq4</i> + nigericin	—	—	$0.90 \pm 0.03^*$
<i>npq4</i> + myxothiazol	$34.7 \pm 13.9$	$0.52 \pm 0.08$	$0.40 \pm 0.09$
<i>npq4</i> + lincomycin	$37.3 \pm 6.3$	$0.58 \pm 0.04$	$0.39 \pm 0.04$
<i>npq4npq1</i>	$15.7 \pm 1.3^*$	$0.56 \pm 0.02$	$0.34 \pm 0.01$
<i>npq4lut2</i>	$27.7 \pm 3.1$	$0.80 \pm 0.03^*$	$0.33 \pm 0.03$
<i>npq4koLhcb5/6</i>	$22.8 \pm 3.4$	$0.49 \pm 0.03$	$0.28 \pm 0.03$
<i>npq4koLhcb4/6</i>	$20.7 \pm 3.5$	$0.49 \pm 0.03$	$0.29 \pm 0.03$
<i>npq4ch1</i>	$14.3 \pm 1.2^*$	$0.71 \pm 0.03^*$	$0.11 \pm 0.03^*$
<i>npq4stn7</i>	$14.2 \pm 1.9^*$	$0.55 \pm 0.04$	$0.37 \pm 0.06$
<i>npq4phot2</i>	$15.3 \pm 5.8^*$	$0.28 \pm 0.05^*$	$0.27 \pm 0.09^*$

former attributions, exponential parameters used for fitting of NPQ dark relaxation will be hereafter called  $A_{\text{qE}}$  and  $\tau_{\text{qE}}$  (qE amplitude and half-life, respectively),  $A_{\text{qM}}$  and  $\tau_{\text{qM}}$  (qM amplitude and half-life),  $A_{\text{qI}}$  (qI amplitude).

Interestingly, the amplitude and half-life of the intermediate phase qM did not significantly differ between WT and *npq4* plants at 400 and 1200  $\mu\text{mol photons m}^{-2} \text{s}^{-1}$ , while  $A_{\text{qM}}$  was only slightly higher in *npq4* than WT plants at 800  $\mu\text{mol photons m}^{-2} \text{s}^{-1}$  (table 1); therefore, these data suggest that the component qM accounts for a fluorescence decay process, common to WT and *npq4* plants, activated upon prolonged illumination and distinct from qE and qI.

Increasing the actinic light intensity caused a linear increase of the slowly relaxing component qI ( $A_{\text{qI}}$ , table 1), that can reasonably be attributed to PSII photoinhibitory processes in both genotypes. In *npq4*, the parameter  $F_v/F_m$  (PSII maximal quantum efficiencies) reached lower values than WT upon illumination at intensities higher than 400  $\mu\text{mol photons m}^{-2} \text{s}^{-1}$  (table S2 in the electronic supplementary material), thus showing that induction of PSII damage is more pronounced in *npq4* plants. However, the extent of PSII photoinhibition at 400  $\mu\text{mol photons m}^{-2} \text{s}^{-1}$  was nearly identical to WT, which indicates that this light treatment did not cause differential PSII damage in the two genotypes; moreover, qM amplitude and kinetics were similar in WT and *npq4* plants using this light regime (table 1), thus showing that the middle decay phase is not influenced by qE and is not related to PSII photoinhibition. Therefore, the molecular basis of the component qM was further investigated under the optimal NPQ induction conditions of 400  $\mu\text{mol photons m}^{-2} \text{s}^{-1}$ , 60 min.

### (b) Effect of $\Delta\text{pH}$ and D1 turnover on qM

Build-up of trans-thylakoid  $\Delta\text{pH}$  is the key event that triggers NPQ; indeed,  $\Delta\text{pH}$  collapses upon treatment with the

ionophore nigericin and the activation of thermal energy dissipation is prevented. *npq4* leaves vacuum-infiltrated with the ionophore still maintained a rise of NPQ in the light, whose amplitude was comparable to that of untreated *npq4* leaves (figure 2a); nevertheless, NPQ relaxation in the dark was missing, thus suggesting that the quenching was mainly because of photoinhibition of PSII reaction centres (table 2). A recent report [34] suggested that the slow phase of NPQ relaxation in the dark would reflect the consumption of ATP accumulated in the light phase. To test this possibility, amplitude and kinetics of qM in *npq4* leaves were measured upon inhibition of the respiratory chain with myxothiazol: upon this treatment, cytoplasm is depleted in ATP in the dark, therefore acting as a sink for chloroplastic ATP, and accelerating NPQ relaxation [34]. However, myxothiazol treatment did not affect either qM amplitude or its half-life (figure 2b and table 2), thus indicating that the middle phase of NPQ decay was not related to slow  $\Delta\text{pH}$  relaxation.

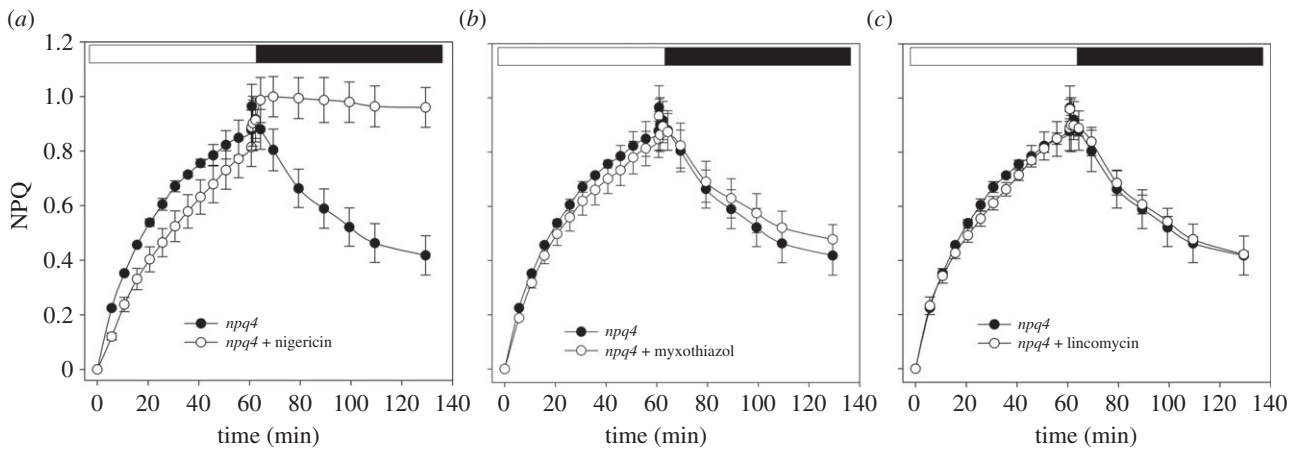
We further tested the possibility that turnover of D1 could account for qM; thus NPQ kinetics were measured on *npq4* leaves upon treatment with lincomycin, a plastid protein synthesis inhibitor. Results (figure 2c and table 2) show that lincomycin treatments failed to affect NPQ rise and decay in *npq4* leaves, indicating that turnover of the PSII reaction centre did not significantly contribute to qM.

To further investigate the molecular basis of qM, we used a targeted reverse genetic approach: we constructed a series of double and triple mutants combining *npq4* with mutations affecting mechanisms that are known to influence Chl fluorescence yield *in vivo*.

### (c) Role of xanthophyll composition

Induction and relaxation of NPQ were measured on *Arabidopsis npq4* mutants with altered xanthophyll composition (figure 3).





**Figure 2.** NPQ analysis of *npq4* leaves upon inhibition of trans-thylakoid  $\Delta pH$ , PSII repair mechanism or mitochondrial ATP production. Kinetics of NPQ induction and relaxation were measured in dark-adapted leaves, upon 60 min illumination at  $400 \mu\text{mol photons m}^{-2} \text{s}^{-1}$ , followed by a further 60 min of dark relaxation, in the absence or presence of  $50 \mu\text{M}$  nigericin (a),  $2 \mu\text{M}$  myxothiazol (b) or  $100 \mu\text{M}$  lincomycin (c). Symbols and error bars show mean  $\pm$  s.d. ( $n = 3$ ).

The *npq1* mutant lacks violaxanthin de-epoxidase activity and is thus unable to convert *Viola* into *Zea*; *qE* in *npq1* has approximately 40% amplitude with respect to WT [12], showing that *Zea* synthesis is needed for full expression of *qE*. A comparison of quenching dynamics showed that *qM* kinetics are very similar in *npq4* and *npq4npq1* plants (figure 3a): although *npq4* showed a somewhat more rapid NPQ rise within the first minutes of illumination, prolonged treatment lead to overlapping amplitude in both genotypes (table 2); likewise, *npq4npq1* plants showed a dark recovery which was initially faster than in *npq4*, while the extent of reversible NPQ was similar in the two genotypes at the end of dark period. The half-life of fluorescence dark recovery is lower in *npq4npq1* than in *npq4*, suggesting that residual *Zea* in the dark could account for the different kinetics in dark recovery.

Long-term NPQ measurements were performed on *lut2*, an *Arabidopsis* mutant devoid of lutein (figure 3b). *Lut*, together with *Zea*, affects quenching dynamics by modulating *qE* [35]. Prolonged illumination leads to overlapping NPQ traces in *npq4* and *npq4lut2*, and *qM* half-life and amplitude were very similar in both genotypes (table 2), implying that *qM* is not affected by *Lut* depletion. In conclusion, the analysis of NPQ dark relaxation kinetics in *npq4* double mutants indicates that *qM* dynamics were only slightly affected, if at all, by mutations in xanthophyll composition.

#### (d) Role of light-harvesting complexes and state 1–state 2 transitions

As NPQ depends on the antenna proteins [14,22,23,36], we evaluated the capacity of *npq4* mutants, devoid of specific LHC gene products, to modulate *qM*.

In mutants devoid of both *Lhcb5* and *Lhcb6* subunits (figure 3c and table 2), the amplitude and relaxation of NPQ were essentially the same as observed in *npq4* mutants. Similar results were obtained upon removal of both *Lhcb4* and *Lhcb6* (figure 3d and table 2), thus ruling out the possibility that minor antennae modulate *qM* amplitude and kinetics. These data are consistent with the behaviour of *npq4ch1* that lacks *Chlb*, and is thus devoid of all LHCs [37,38]; the slow phase of NPQ relaxation was found to be independent of LHC composition, indeed *qM* amplitude was similar in *npq4* and *npq4ch1* mutants (figure 3e and table 2). State transitions lead

to quenching of LHCII fluorescence by PSI [39] upon phosphorylation of LHCII, by STN7 kinase, driving its migration from PSII to PSI [40]; by using the mutant *npq4stn7*, we checked the possibility that state transition were involved in *qM*. The maximal amplitude of NPQ in *npq4stn7* was essentially the same as in *npq4* plants (figure 3f). The only difference was found in the kinetics of *qM* dark-recovery, which was faster in the double mutant than in the *npq4* mutant ( $\tau$  about 15 min in *npq4stn7* versus 29 min in *npq4*; table 2).

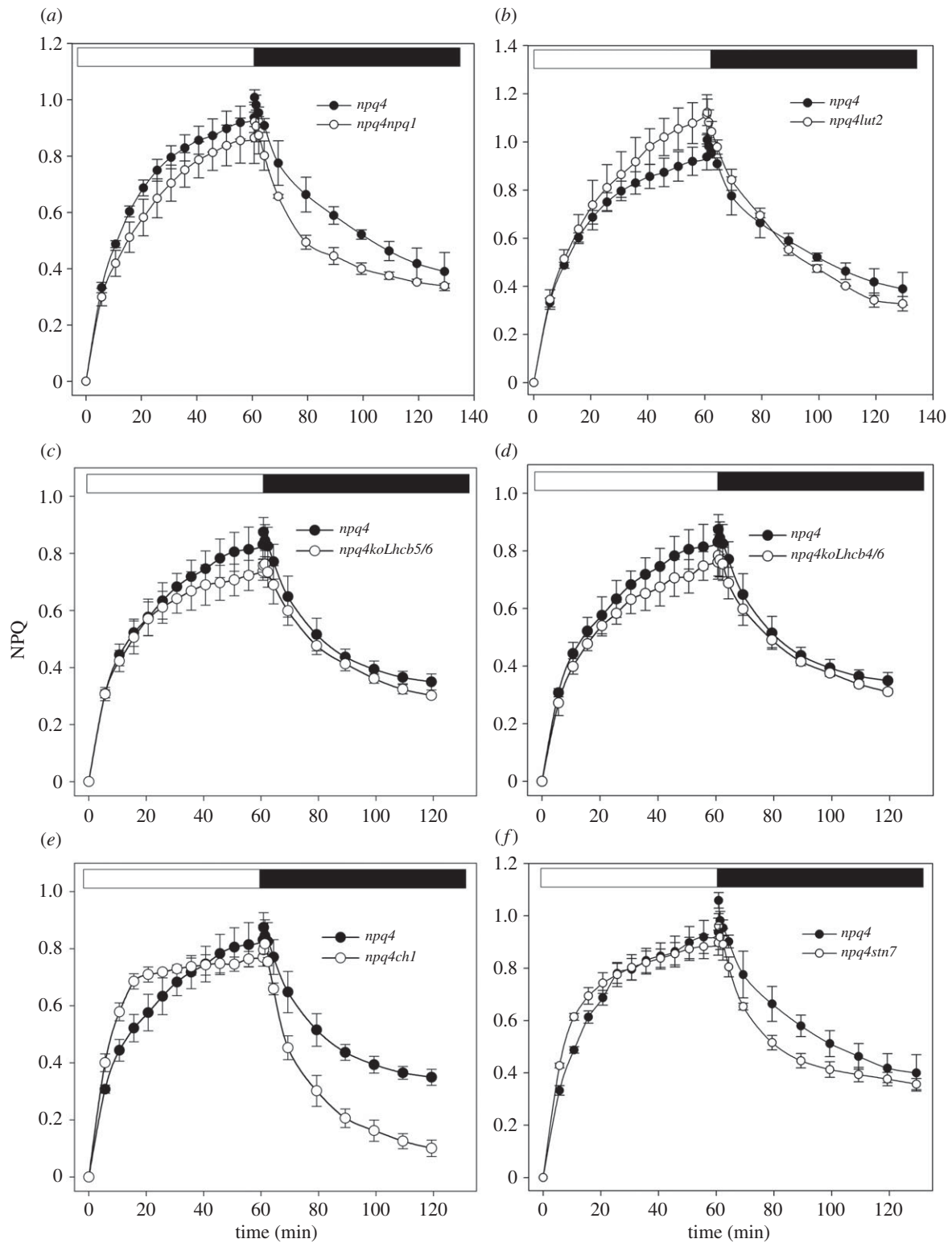
#### (e) Role of chloroplast photorelocation

Previous reports showed that light-induced chloroplast movements could affect Chl fluorescence emission [41,42]. We thus determined the *qM* relaxation kinetics in a double mutant *npq4phot2*, lacking the phototropin PHOT2 which activates the blue-light-dependent chloroplast avoidance movement [24].

Although two different components of NPQ dark-relaxation (*qM* and *qI*) were still detected in *npq4phot2*, the kinetics of the *qM* component were faster than in *npq4* (figure 4a and table 2), and the amplitude of the middle phase was decreased by 50% compared with *npq4*.

To further verify the hypothesis that phototropins were involved in the modulation of *qM*, we repeated NPQ measurements by using red light ( $600 < \lambda < 750 \text{ nm}$ ,  $350 \mu\text{mol photons m}^{-2} \text{s}^{-1}$ ,  $23^\circ\text{C}$ ) as actinic source. Results are shown in figure 4b: both *npq4* and *npq4phot2* matched the kinetics of *npq4phot2* in white light, implying that a specific fluorescence decay component is activated by actinic light  $\lambda < 600 \text{ nm}$  but not by illumination with red light, and affects the amplitude of *qM* specifically. This point was further studied through the analysis of Chl fluorescence parameters  $F'_m$  and  $F'_0$  during photosynthesis: we found that, upon illumination with white light at  $400 \mu\text{mol photons m}^{-2} \text{s}^{-1}$ , *npq4* leaves underwent a decrease in both  $F'_m$  and  $F'_0$  as the result of change in both quantum yield and leaf transmittance, respectively, while in *npq4phot2* only  $F'_m$  decreased upon irradiation (see electronic supplementary material, figure S2).

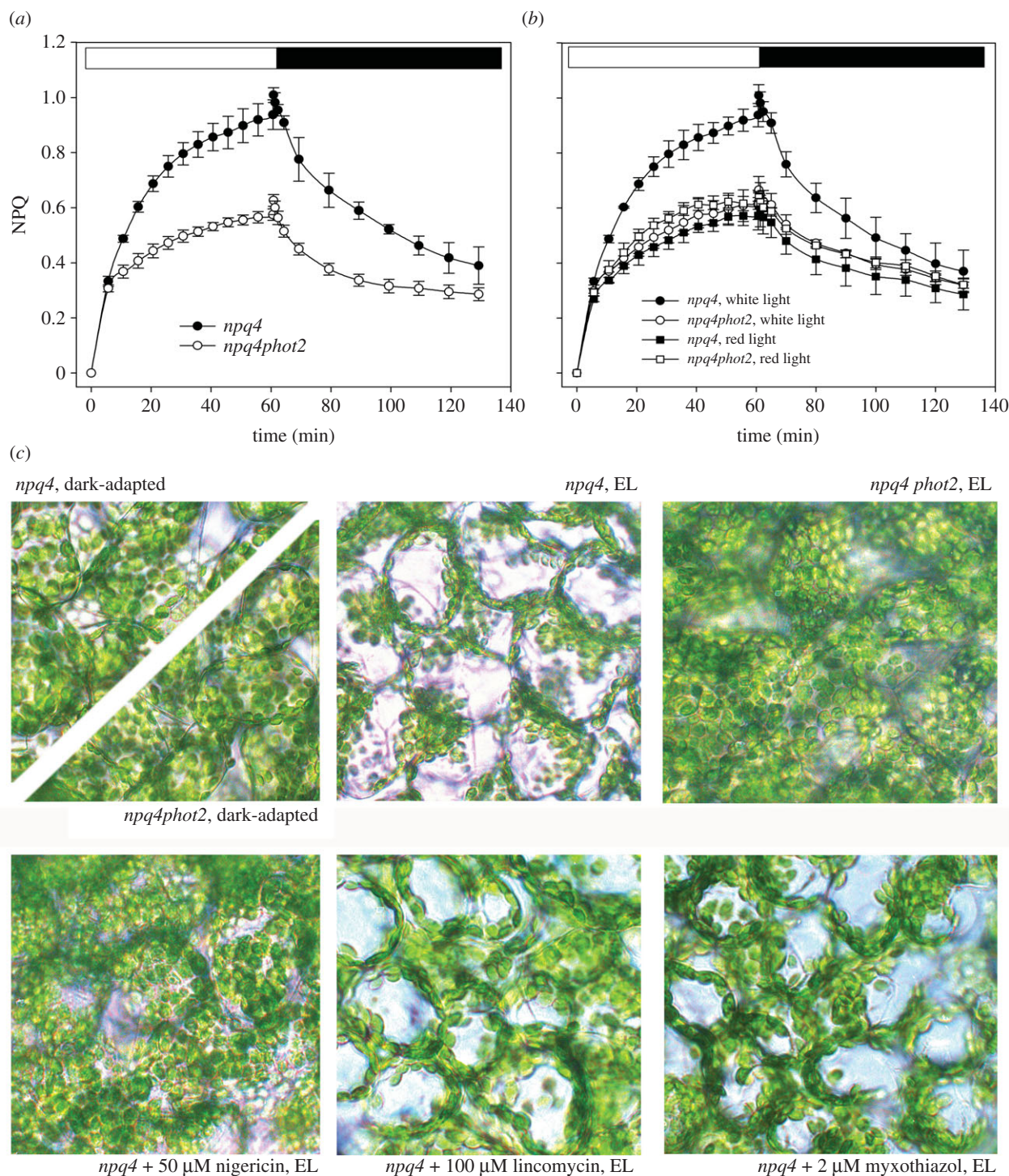
Thus, the chloroplast avoidance response is, among all the mechanisms examined, the only significant modulator of the concentration of Chl excited states independent of PsbS. To confirm the effect of the *phot2* mutation on the *npq4* background, we examined leaf cells by light microscopy (figure 4c). Upon irradiation with white light at  $400 \mu\text{mol photons m}^{-2} \text{s}^{-1}$ ,



**Figure 3.** NPQ analysis of *npq4* leaves with altered xanthophyll compositions or depleted of either PSII LHCs or state transition. NPQ induction and relaxation were measured in dark-adapted *npq4* plants lacking zeaxanthin (*npq4npq1* (a)) or lutein (*npq4lut2* (b)), or devoid of either minor Lhcb CP26 and CP24 (*koLhcb5/6* (c)), CP29 and CP24 (*koLhcb4/6* (d)) or the entire LHC owing to mutation (*ch1* (e)), or blocked in state transitions (*npq4stn7* (f)). NPQ kinetics were measured in dark-adapted leaves, upon 60 min illumination at  $400 \mu\text{mol photons m}^{-2} \text{s}^{-1}$ , followed by further 60 min of dark relaxation. Symbols and error bars show mean  $\pm$  s.d. ( $n = 3$ ).

*npq4* leaves showed chloroplasts located at the anticlinal cell walls, thus indicating that they had undergone avoidance movement. Instead, the chloroplasts of the mutant *npq4phot2* retained their preferential association with the periclinal walls, as in leaves adapted to darkness for 1 h. Treatment with nigericin blocks the chloroplast avoidance response (figure 4c), consistent with the complete depletion of qM after treatment with the ionophore (figure 2a).

In conclusion, the analyses of quenching relaxation dynamics on a number of mutants in the *npq4* genetic background, identified a kinetically intermediate component of fluorescence decay, distinct from either qE or qI, called qM, the triggering of which requires uniquely formation of a transmembrane proton gradient, but which is not related to xanthophyll or LHC composition, PSII turnover, consumption of ATP accumulated in the light phase or state transitions. Based on its relaxation



**Figure 4.** NPQ analysis of *npq4* leaves depleted of chloroplast avoidance movement. (a) PsbS-minus plants were crossed with *Arabidopsis* knock-out lines lacking the photoprotective mechanism of chloroplast avoidance movement (*phot2*). Kinetics of NPQ induction and relaxation of *npq4* and *npq4phot2* were measured in dark-adapted leaves, as described for figure 3. (b) NPQ kinetics were measured on *npq4* and *npq4phot2* leaves upon illumination with either white actinic light ( $400 \mu\text{mol photons m}^{-2} \text{s}^{-1}$ ) or red light ( $350 \mu\text{mol photons m}^{-2} \text{s}^{-1}$ ,  $600 < \lambda < 750 \text{ nm}$ ). Symbols and error bars show mean  $\pm$  s.d. ( $n = 3$ ). (c) Distribution of chloroplasts in the mesophyll cells of *npq4* and *npq4phot2* was determined by light microscopy. Leaves were dark-adapted for 1 h and then irradiated with white light at  $400 \mu\text{mol m}^{-2} \text{s}^{-1}$  for 1 h. Prior to light treatment, detached leaves were infiltrated with 150 mM sorbitol containing either 50  $\mu$ M nigericin, 100  $\mu$ M lincomycin or 2  $\mu$ M myxothiazol. (Online version in colour.)

kinetics, the mechanism of chloroplast photorelocation accounts for nearly 50% of qM amplitude.

## 4. Discussion

Here, we have investigated the light-induced decline of Chl fluorescence and its relaxation dynamics in the *npq4* mutant

of *Arabidopsis*, lacking the PsbS subunit essential for qE activity, in order to assess the basis for its residual light-induced fluorescence decline activity. Kinetic analysis of fluorescence dark recovery *in vivo* allowed an NPQ component to be identified which relaxes in the dark within the time range 16–25 min, intermediate between the fast qE component (1–2 min) and inhibitory quenching qI (more than 1 h). This component, qM, showed similar amplitude and



half-life in WT and *npq4* plants (figure 1 and table 1) and is uncoupler-sensitive (figure 2a). To search for the molecular basis of this process, the *npq4* genotype was crossed with others, which affected photosynthetic components and mechanisms known to alter the characteristics of light-induced Chl fluorescence changes, and analysed the fluorescence quenching in each.

### (a) The kinetic components of non-photochemical quenching dark relaxation in *npq4*

Analysis of NPQ dark relaxation of WT and *npq4* leaves identified three distinct kinetic components (figure 1 and table 1). The rapid phase (half-life 35–55 s) detected in WT but not in *npq4* leaves, can be safely assigned to qE [8]. The long-term relaxing component, whose half-life is longer than 60 min, can be assigned to photoinhibitory processes based on its amplitude dependence of photon fluency and increased incidence in *npq4* [13]. The third, intermediate kinetic component (half-life 20–35 min), can be detected with similar amplitude in both WT and *npq4* relaxation kinetics. Although the existence of a middle-phase kinetic component of NPQ has been reported previously [8,43], its physiological origin is still debated.

The middle phase component, qM, is saturated at moderate light intensity, maintaining the same amplitude at 800 and 1200  $\mu\text{mol photons m}^{-2} \text{s}^{-1}$  (table 1). This suggests that qM is not related to PSII photoinhibition, as the amplitude of a photoinhibitory component is expected to increase with irradiance; indeed, the component here defined as qI increases from 0.67 at 800  $\mu\text{mol photons m}^{-2} \text{s}^{-1}$  to 1.01 at 1200  $\mu\text{mol photons m}^{-2} \text{s}^{-1}$  in *npq4* (figure 1 and table 1). At all light regimes tested, the *npq4* leaves showed higher qI than WT leaves, consistent with the photoprotective role of qE in short-term exposure to EL [44].

We used an actinic intensity of 400  $\mu\text{mol photons m}^{-2} \text{s}^{-1}$  as at this irradiance qM is almost saturated and photoinhibition in *npq4* is as low as in the WT leaves (table S2 available in the electronic supplementary material). Lincomycin treatment failed to affect NPQ decay in *npq4* leaves, implying that at 400  $\mu\text{mol photons m}^{-2} \text{s}^{-1}$  turnover of D1 did not significantly contribute to qM.

A slow phase of NPQ dark relaxation was reported to depend on the hydrolysis of ATP accumulated during a light phase [34]. However, myxothiazol treatment did not affect qM in *npq4* (figure 2b and table 2), thus ruling out the possibility that the qM decay component was related to slow  $\Delta\text{pH}$  relaxation.

It is interesting to note that qM is sensitive to uncouplers: indeed, dark relaxation of fluorescence decline is prevented in leaves infiltrated with the ionophore nigericin. Besides the loss of qM, nigericin led to a strong increase in qI amplitude, implying that both qE and qM are crucial for PSII photoprotection in EL conditions (figure 2a). The intermediate component has been previously defined as qT or qZ for its possible dependence on state transitions or Zea biosynthesis, respectively [8,15,43]. However, data reported here show that blocking these processes with specific mutations does not interfere with qM as determined in the *npq4* strain. This mutant was chosen to avoid overlapping contribution of qE to the dark relaxation dynamics. In *npq4*, we could not detect qE type quenching ( $\tau < 5$  min) under any conditions, even upon 60 min of illumination (table 1).

### (b) Targeted reverse genetic analysis to identify the molecular basis for qM

NPQ kinetic of nigericin-treated leaves demonstrated that triggering of qM requires transmembrane proton gradient formation during the light phase. Among the effects of EL treatment is thylakoid lumen acidification and Zea synthesis, which are needed for full expression of qE in *Arabidopsis* [31]. Instead, the double mutant *npq4npq1*, which is depleted in both qE and Zea, showed the same kinetics and amplitude of qM as *npq4* (figure 3a). Therefore, unlike qE, Zea depletion did not prevent full expression of qM, which indeed reaches maximum value (although much lower than in WT) in both *npq4* and *npq4npq1* plants (table 2). The kinetics of Zea epoxidation in *npq4* do not fit with those of NPQ dark recovery, the former being far slower than the kinetic relaxation of qM [15]. We conclude that the xanthophyll cycle, one of the most efficient modulators of qE [14,31], does not affect the amplitude of qM but only slowed the dark relaxation rate (figure 3a). Similar considerations can be applied to Lut; indeed, the amplitude and kinetics of qM are essentially the same in both *npq4* and *npq4lut2* plants (figure 3b). qE is located in the antenna system, and an important role in NPQ was attributed to Lhcb4 and Lhcb5 [36,45]. Nevertheless, light-induced fluorescence decline was essentially the same in *npq4*, *npq4koLhcb4/6* and *npq4koLhcb5/6* leaves. Moreover, no significant differences were found when comparing *npq4* versus *npq4chl1* leaves depleted in all LHC proteins, including LHCI (table 2) [37], suggesting that these changes were not the result of qE type quenching. Finally, we examined the hypothesis of the involvement of state 1–state 2 transitions in PsbS-independent fluorescence decline; however, the similar behaviour of *npq4* and *npq4stn7* leaves (figure 3f) excludes this possibility.

### (c) Chloroplast avoidance response and qM

Among all the mutations introduced into the *npq4* genetic background, *phot2* was the only one that affected qM (figure 4a). Differences in *npq4* versus *npq4phot2* NPQ kinetics showed that the fluorescence recovery component is affected by chloroplast photorelocation. This process is mediated by the blue light receptors, phototropins. Consistently, we verified that the same effect on NPQ kinetics was obtained by using red light rather than white actinic light (figure 4b). Moreover, upon illumination with white light at 400  $\mu\text{mol photons m}^{-2} \text{s}^{-1}$ , *npq4* leaves underwent a decrease in both  $F'_m$  and  $F'_o$ , as expected for concomitant changes in PSII quantum yield and leaf absorption, respectively, while in *npq4phot2* leaves only  $F'_m$  was decreased upon irradiation (see the electronic supplementary material, figure S2). Thus, an avoidance response, causing chloroplast movement towards cell walls parallel to incident light, affects the apparent kinetics of NPQ, particularly the slower components. This is consistent with reports showing that light-induced chloroplast movements could affect Chl fluorescence emission [41], and with recent results which highlighted, in *phot2*, the lack of a fluorescence decay kinetic component, qM, intermediate between qE and qI [46]. The similar amplitude and half-life of the qM component in WT and *npq4* leaves (table 1), and its reduction in a mutant devoid of chloroplast photorelocation (table 2; see also [46]), strongly support the view that chloroplast relocation significantly influences the apparent kinetics of NPQ by decreasing the photon absorption of leaves, rather than changing the activity of quenching reactions.



It should be noted, however, that this effect arises from decreased photon absorption which gives a lower fluorescence yield, rather than from a genuine quenching process. In fact, during illumination, chloroplast relocation induces a change in the distribution of pigments within the cell, with the formation of localized chloroplast stacks along anticlinal cell walls. This effect reduces the overall optical density of the cell because of a 'sieve effect' resulting from the formation of highly transmitting paths across the periclinal cell surfaces while increasing the optical density beyond linearity in the vicinity of the anticlinal cell walls [47]. Thus, the fraction of excited, fluorescence emitting chloroplasts is decreased because of shading by neighbouring, ones. It is worth noting that the kinetics and timescale of qM formation and relaxation at 400  $\mu\text{mol photons m}^{-2} \text{s}^{-1}$  of white light (table 1) fits with that described for the chloroplast avoidance response under similar irradiance [48]. Light microscopy analysis (figure 4c) confirmed that movement of chloroplasts was inhibited in the presence of nigericin, consistent with the depletion in qM. The chloroplast avoidance response probably relies on the cytosolic  $\text{Ca}^{2+}$  signal for its activation [49]. Maintenance of a low cytosolic  $\text{Ca}^{2+}$  level requires an electrogenic pump which exploits, protonmotive force to actively extrude  $\text{Ca}^{2+}$  [50]. Phototropin signal transduction involves transient depolarization of the plasma membrane which, in turn, triggers cytosolic  $\text{Ca}^{2+}$  intake. Nigericin wrecks all the transmembrane electrochemical gradients, thus blocking several signal transduction events. The double effect of nigericin in collapsing the thylakoid pH gradient and in blocking chloroplast relocation (figure 4c) can easily lead to misinterpretation of qM as a slow qE response in the absence of PsbS.

Although chloroplast relocation is the major factor affecting the amplitude of qM in *npq4* (figure 4a,b), the fluorescence recovery kinetics of *npq4phot2* are not completely devoid of qM. The residual component accounts for about 18% of total reversible  $F'_m$  quenching in WT (tables 1 and 2) and reflects mechanism(s) sensitive to uncouplers (figure 2a) and yet distinct from the avoidance response, as it is still active in *npq4phot2*. Previous work led to different proposals for mechanisms leading to fluorescence recovery components with intermediate half-life between qE and qI. First, it was attributed to state 1–state 2 transitions [43]; second to PSII photoinhibition [51]; third to a slowly developing component of qE dependent on Zea [52]; fourth to light-induced dissociation of the complex Lhcb4-Lhcb6-LHCII-M [19]. Here, we

show that qM did not correlate with Zea accumulation nor was it related to qI. Although thylakoid membrane reorganization could well explain changes in chloroplast fluorescence yield since protein–protein interactions are responsible for nearly 50% of quenching [14,19], the need for PsbS to trigger domain reorganization [19,20] suggests this is not the source of residual qM. An interesting observation is that a substantial fraction of qM is retained in the absence of Lhcb in the *npq4chl1* mutant (figure 3e), although with a somewhat shorter half-life (table 2). This is consistent with the characteristics of Zea-independent NPQ localized in the PSII core complex [53].

Our results support the view that no qE occurs in *npq4* leaves within a wide range of actinic light intensities. Moreover light-induced fluorescence decline in *npq4* was always far lower than in WT plants, even upon 1 h of EL exposure (figure 1). Finally, the residual fluorescence decline in *npq4* leaves is due to avoidance of photon absorption, while quenching mechanisms can only be responsible for a minor component associated with the PSII core (figure 3e), rather than to reactions within the antenna system. Our results significantly differ from those of Johnson & Ruban [21], who reported that qE could be catalysed, although at a slower rate, in *npq4* plants. First, we found that the amplitude of fluorescence decline in *npq4* leaves did not match that observed in WT under any conditions, the fraction of reversible  $F'_m$  quenching in the absence of PsbS and triggered by lumen acidification being small (about 18%) (table 1). The photoprotective effect of fluorescence decline was consistently low, as shown from the higher amplitude of qI in *npq4* leaves under EL conditions (table 1). Second, we found that uncouple sensitivity is the result of the disruption of chloroplast relocation, also involving a proton gradient for signal transduction upon blue light activation of phototropins [49]. Overall, these results point to a crucial role of PsbS in the modulation of NPQ and show that sensing of trans-thylakoid  $\Delta\text{pH}$  by protonable residues in the LHC is not enough to induce WT levels of NPQ in the absence of PsbS. This conclusion is consistent with recent results showing that less than 0.5% of purified minor antennae underwent charge transfer quenching *in vitro*, whereas the fraction engaged in this process was more than 80 times higher in intact thylakoids with PsbS [36]. Thus, PsbS is indispensable for qE, within the trans-thylakoid  $\Delta\text{pH}$  that can be obtained by light treatment of leaves, rather than being only a modulator of the proton–antenna association constant, pK, of qE activation [54].

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