



## The role of the high potential form of the cytochrome *b559*: Study of *Thermosynechococcus elongatus* mutants



Fernando Guerrero<sup>a,b,1</sup>, Jorge L. Zurita<sup>a,b,2</sup>, Mercedes Roncel<sup>a</sup>, Diana Kirilovsky<sup>b</sup>, José M. Ortega<sup>a,\*</sup>

<sup>a</sup> Instituto de Bioquímica Vegetal y Fotosíntesis, Universidad de Sevilla-CSIC, Américo Vespucio 49, 41092 Seville, Spain

<sup>b</sup> Laboratoire de Bioénergétique Moléculaire et Photosynthèse, Institut de Biologie et de Technologies de Saclay (iBiTec-S), CEA Saclay, 91191 Gif-sur-Yvette cedex, France

### ARTICLE INFO

#### Article history:

Received 30 October 2013

Received in revised form 21 February 2014

Accepted 26 February 2014

Available online 5 March 2014

#### Keywords:

Cytochrome *b559*

High potential form

Photosystem II

Redox potentiometry

Site-directed mutagenesis

*Thermosynechococcus elongatus*

### ABSTRACT

Cytochrome *b559* is an essential component of the photosystem II reaction center in photosynthetic oxygen-evolving organisms, but its function still remains unclear. The use of photosystem II preparations from *Thermosynechococcus elongatus* of high integrity and activity allowed us to measure for the first time the influence of cytochrome *b559* mutations on its midpoint redox potential and on the reduction of the cytochrome *b559* by the plastoquinone pool (or  $Q_B$ ). In this work, five mutants having a mutation in the  $\alpha$ -subunit (I14A, I14S, R18S, I27A and I27T) and one in the  $\beta$ -subunit (F32Y) of cytochrome *b559* have been investigated. All the mutations led to a destabilization of the high potential form of the cytochrome *b559*. The midpoint redox potential of the high potential form was significantly altered in the  $\alpha$ R18S and  $\alpha$ I27T mutant strains. The  $\alpha$ R18S strain also showed a high sensitivity to photoinhibitory illumination and an altered oxidase activity. This was suggested by measurements of light induced oxidation and dark re-reduction of the cytochrome *b559* showing that under conditions of a non-functional water oxidation system, once the cytochrome is oxidized by P680<sup>+</sup>, the yield of its reduction by  $Q_B$  or the PQ pool was smaller and the kinetic slower in the  $\alpha$ R18S mutant than in the wild-type strain. Thus, the extremely positive redox potential of the high potential form of cytochrome *b559* could be necessary to ensure efficient oxidation of the PQ pool and to function as an electron reservoir replacing the water oxidation system when it is not operating.

© 2014 Elsevier B.V. All rights reserved.

### 1. Introduction

In higher plants, algae and cyanobacteria, the primary steps of photosynthetic light conversion take place in a large pigment–protein complex named photosystem II (PSII), which drives light-induced electron transfer from water to plastoquinone (PQ) with the concomitant production of molecular oxygen (for a review see [1]). The reaction center

(RC) of PSII is composed of an heterodimer formed by the D1 and D2 membrane proteins that bears all the cofactors needed for charge separation and stabilization: six chlorophylls (Chl) (including four of the Chl center, P680), two pheophytins, two plastoquinones  $Q_A$  and  $Q_B$ , one non-heme iron atom, one tetranuclear manganese cluster ( $Mn_4CaO_5$ ) and two redox active tyrosines. A heme protein, named cytochrome *b559* (Cyt *b559*), appears also tightly bound to RC in all photosynthetic oxygen-evolving organisms.

Cyt *b559* is one of the essential components of PSII RC, but its function although widely investigated, still remains unclear (for a review see [2,3]). The resolution of the 3-D structure of PSII has clearly established that Cyt *b559* is a heme-bridged protein heterodimer with two subunits ( $\alpha$  and  $\beta$ , encoded by *psbE* and *psbF* genes) of 9 and 4 kDa, respectively [4–7]. Each polypeptide chain forms a transmembrane  $\alpha$ -helix, the heme being located near the stromal side and perpendicular to the membrane plane. A bis-histidine axial coordination for the heme is largely supported by these structural studies.

Cyt *b559* is a *b*-type cytochrome with peculiar physicochemical properties. It is found in PSII preparations in three different states: a high potential form (HP) with a midpoint redox potential ( $E_m$ ) of  $\approx +400$  mV, another intermediate potential form (IP) with an  $E_m$  of  $\approx +200$  mV and a low potential form (LP) whose  $E_m$  ranges from  $+80$  and  $0$  mV depending on the organisms [3,8–12]. Many authors have proposed that the HP

**Abbreviations:** B band, thermoluminescence emission due to  $S_{2/3}Q_B^-$  charge recombination reaction; Chl, chlorophyll; Cm, chloramphenicol; CP43, a proximal antenna complex of photosystem II; Cyt, cytochrome;  $E_h$ , redox potential of the solution;  $E_m$ , midpoint redox potential; HP, IP and LP, high, intermediate and low potential forms; HTL, high temperature thermoluminescence; MES, 2-(N-morpholino) ethanesulfonic acid; OEC, oxygen evolving complex; P680, primary electron donor in PSII; PSII, photosystem II;  $Q_A$  and  $Q_B$ , the primary and secondary quinone acceptors of the reaction center of PSII; RC, reaction center;  $S_2$  and  $S_3$ , oxidized states of the manganese oxygen-evolving complex of PSII; Sm, streptomycin; Sp, spectinomycin; TL, thermoluminescence

\* Corresponding author. Tel.: +34 954489573; fax: +34 954460065.

E-mail addresses: [Fernando.guerrero@helsinki.fi](mailto:Fernando.guerrero@helsinki.fi) (F. Guerrero),

[jorge.zurita@juntadeandalucia.es](mailto:jorge.zurita@juntadeandalucia.es) (J.L. Zurita), [mroncel@us.es](mailto:mroncel@us.es) (M. Roncel), [kirilovsky@cea.fr](mailto:kirilovsky@cea.fr) (D. Kirilovsky), [ortega@us.es](mailto:ortega@us.es) (J.M. Ortega).

<sup>1</sup> Present address: Institute of Biotechnology, University of Helsinki, Viikinkaari 1, 00014 Helsinki, Finland.

<sup>2</sup> Present address: Fundación Progreso y Salud, Junta de Andalucía, Américo Vespucio 5, 41092 Seville, Spain.

form is the physiological and functional state of Cyt *b559* [3,10]. However, the HP form is labile and easily converted to the LP/IP forms by treatments that alter the membrane structure or during the process of PSII isolation [8,10]. In plants, algae and cyanobacteria, different kinds of PSII preparations with very variable percentages of the HP form, peaking at about 80% have been described [8,10,11,13,14].

A great number of hypotheses have been put forth in order to explain the function of Cyt *b559*. Most of them propose that this protein may participate in secondary electron transfer pathways protecting PSII against oxidative damage [15–18]. In these proposals, Cyt *b559* would donate electrons, via a  $\beta$ -carotene molecule, to reduce highly oxidizing chlorophyll radicals generated in PSII RC under donor-side photoinhibitory conditions [17,19,20]. On the other hand, Cyt *b559* may accept electrons from the acceptor side of PSII ( $Q_B$  or PQ) to prevent the formation of damaging singlet oxygen species under acceptor-side photoinhibitory conditions [8,21,22].

Different new roles as oxygen reductase, superoxide reductase, superoxide oxidase and plastoquinol oxidase have been recently proposed for Cyt *b559* [12,22–25]. The role as plastoquinol oxidase, taking electrons from the plastoquinone pool, has been supported by the existence of a third quinone,  $Q_C$ , different from the quinone sites,  $Q_A$  and  $Q_B$ , at 15 Å only from the heme of Cyt *b559* [7]. Some authors have postulated the existence of three quinones, besides  $Q_A$ , in PSII core complex preparations of *Thermosynechococcus* (*T.*) *elongatus* [26,27]. A function for  $Q_C$  associated with a secondary electron transfer along the Cyt *b559* in PSII has been proposed by Guskov et al. [7]. Therefore, a role for Cyt *b559* as part of a connecting channel between the plastoquinone pool and  $Q_B$  and  $Q_C$  sites would be possible.

Most of the site-directed mutagenesis studies of the  $\alpha$ - and  $\beta$ -subunits of Cyt *b559* have been carried out in *Chlamydomonas reinhardtii* and *Synechocystis* sp. PCC 6803. No stable PSII was assembled when the heme-coordinating histidines were replaced by leucines [28], neither when the histidine of the  $\alpha$ -subunit was replaced by tyrosine or methionine [29], or in deletion mutants of the  $\alpha$ -subunit [30] and of both  $\alpha$ - and  $\beta$ -subunits [31]. In fact, all these mutants were impaired in the functional PSII assembly and therefore not useful for determining the function of Cyt *b559* in PSII.

Recently, a set of site-directed mutants of Cyt *b559* in *Synechocystis* sp. PCC 6803 that grew photoautotrophically and accumulated stable PSII RC's has been constructed [14,32,33]. Mutants with changes on heme coordination (His22 of the  $\alpha$ - or  $\beta$ -subunit) and on charged residues of the cytoplasmic side of Cyt *b559* (R7E $\alpha$ , R17E $\alpha$  and R17L $\beta$ ) showed a higher sensitivity to photoinhibition than the WT strain. These results have led the authors to propose a functional role of Cyt *b559* in protection of PSII under photoinhibition conditions [14]. They also suggested that the electrostatic interactions between these arginine residues and the heme propionates of Cyt *b559* are essential in the maintenance of the structure and redox properties of the cytochrome. However, the *Synechocystis* sp. PCC 6803 strain used in this work seems to present an important limitation for the study of the Cyt *b559* role because the PSII preparations contained very low proportions of the HP form [14,33].

The thermophilic cyanobacterium *T. elongatus* has become in the last years a new model organism for photosynthesis researches, since genetic engineering techniques have been developed [34,35] and several 3-D structures of PSII have been solved from it [4,6,7,36]. In 2003, our group was already able to obtain PSII preparations from *T. elongatus*, 50% containing the HP form of cytochrome *b559* and 50% an IP form, both in their reduced state [11]. No LP form was present in these preparations. The HP form in *T. elongatus* showed the typical redox properties of the HP form in higher plants: (1) it has a highly oxidizing  $E_m$ , which is not affected by pH [11] and (2) it is partially converted into a form of lower potential (IP) by treatments such as washing with Tris or gentle heating [11]. Recently, PSII preparations with a higher degree of integrity and activity have been prepared in our groups using a new His-tagged CP43 *T. elongatus* strain ([37] and this work). This is essential

to study the properties of the Cyt *b559* HP form. The high rates of  $O_2$  evolution obtained in these preparations were good indicators of the integrity of the donor side of PSII ([37] and Table 1). The 1:1 ratio of the concentration of Cyt *b559* regarding Cyt *c550* [37] indicates that Cyt *c550* is not released from PSII during the isolation process, also a good indicator of donor side integrity. The most important characteristic of these isolated PSII complexes is that 85% of them presented Cyt *b559* in the reduced HP form with an  $E_m$  value of about +400 mV. Thus, our PSII preparations are an excellent material for the study of the effect of changes in the  $E_m$  values and on the percentage of HP/IP forms on the function of Cyt *b559*.

A series of site-directed mutants of Cyt *b559*, each carrying a single amino acid substitution, has been constructed in the thermophilic cyanobacterium *T. elongatus*, in order to modify the redox potential of the heme without altering the assembly properties of PSII. In this work we have analyzed in depth six of these mutant strains: five in the  $\alpha$ -subunit (I14A, I14S, R18S, I27A and I27T) and one in the  $\beta$ -subunit (F32Y). All mutants grew photoautotrophically and assembled functionally PSII RC. For the first time, the redox potential of site-directed mutants of Cyt *b559* has been determined. The  $E_m$  of the HP form of Cyt *b559* was significantly altered in two of the mutant strains:  $\alpha$ R18S and  $\alpha$ I27T. The  $\alpha$ R18S strain also showed a high sensitivity to photoinhibitory illumination. The results support the proposal that the HP form of Cyt *b559* may function in a secondary electron transfer cycle in order to protect PSII when the water oxidation system is impaired.

## 2. Material and methods

### 2.1. Construction of plasmids for the mutation of cytochrome *b559*

The genome region containing the genes *psbE*, *psbF*, *psbL* and *psbJ* (forming an operon) (Fig. 1) was amplified by PCR using the oligonucleotides *cytb559BssHIIa* and *cytb559BssHIIb* (Supplementary Table I). The amplified region which includes part of the *tr1540* gene encoding a serine–threonine kinase was digested by BssHII and cloned. The *tr1540* gene was interrupted by insertion of a 2.2 kb DNA fragment containing the *aadA* gene from Tn7, conferring resistance to spectinomycin (Sp) and streptomycin (Sm) [38], in the unique KpnI restriction site of the *tr1540* gene. Site-directed mutagenesis of the plasmid containing the interrupted *tr1540* gene was performed using the QuikChange XL site-directed mutagenesis kit of Stratagene as recommended by the manufacturer. The synthetic mutagenic oligonucleotides used to create the different point mutations in the *psbE* and *psbF* genes (encoding for the  $\alpha$  and  $\beta$ -subunits of the Cyt *b559*, respectively) are shown in Supplementary Table II. The mutagenic oligonucleotides also introduced a silent mutation creating or eliminating restriction enzyme sites for mutant screening.

### 2.2. Transformation of *T. elongatus* cells and genetic analysis of mutants

The construction of WT' strain of *T. elongatus*, containing a His-tagged CP43 for isolation of PSII complexes with high activity, was described in Guerrero et al. [37]. The plasmids containing the site-mutated *psbE* and *psbF* genes were introduced in WT' by electroporation according to Mühlenhoff and Chauvat [34] with slight modifications as described in [39]. To create the control strain WT'', a plasmid containing the *psbE* and *psbF* genes (without mutations) and the *tr1540* gene interrupted by the Sp/Sm resistance cassette was introduced into WT' cells.

After electroporation, cells were rapidly transferred to 2 ml of DTN-medium [34] and incubated for 48 h in a rotary incubator at 45 °C under low light conditions. Then, the cells in 0.1–0.2 ml aliquots were spread on plates containing chloramphenicol (Cm) (2  $\mu$ g ml<sup>-1</sup>) or Sp/Sm (12  $\mu$ g ml<sup>-1</sup>/6  $\mu$ g ml<sup>-1</sup>) and incubated at 45 °C, under dim light and a humidified atmosphere. Once green

Download English Version:

<https://daneshyari.com/en/article/10795714>

Download Persian Version:

<https://daneshyari.com/article/10795714>

[Daneshyari.com](https://daneshyari.com)