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The heterologous expression of a plastocyanin in the diatom Phaeodactylum tricornutum improves cell growth under irondeficient conditions

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Abstract

We have investigated if the heterologous expression of a functional green alga plastocyanin in the diatom Phaeodactylum tricornutum can improve photosynthetic activity and cell growth. Previous in vitro assays showed that a single-mutant of the plastocyanin from the green algae Chlamydomonas reinhardtii is effective in reducing P. tricornutum photosystem I. In this study, in vivo assays with P. tricornutum strains expressing this plastocyanin indicate that even the relatively low intracellular concentrations of holo-plastocyanin detected (\approx 4 μ M) are enough to promote an increased growth (up to 60%) under iron-deficient conditions as compared with the WT strain, measured as higher cell densities, content in pigments and active photosystem I, global photosynthetic rates per cell, and even cell volume. In addition, the presence of plastocyanin as an additional photosynthetic electron carrier seems to decrease the over-reduction of the plastoquinone pool. Consequently, it promotes an improvement in the maximum quantum yield of both photosystem II and I, together with a decrease in the acceptor side photoinhibition of photosystem II-also associated to a reduced oxidative stress-a decrease in the peroxidation of membrane lipids in the choroplast, and a lower degree of limitation on the donor side of photosystem I. Thus the heterologous plastocyanin appears to act as a functional electron carrier, alternative to the native cytochrome c_6 , under iron-limiting conditions.

1 | INTRODUCTION

Photosynthetic productivity, which is equally shared between terrestrial and oceanic systems, is due to a wide variety of organisms, from cyanobacteria to eukaryotic microalgae and plants (Field et al., 1998; Hall, 1976). Within the group of eukaryotic microalgae, diatoms constitute the dominant life form in oceanic phytoplankton, probably

Abbreviations: Cb_of , cytochrome b_of ; Cc_{550} , cytochrome c_{550} ; Cc_o , cytochrome c_o ; Cf, cytochrome f; Chl, chlorophyll; ET, electron transfer; Fd, ferredoxin; Fld, flavodoxin; HTL, high temperature thermoluminescence; P_{700} , photosystem I primary donor; Pc, plastocyanin; PQ, plastoquinone; PSl, photosystem I; PSII, photosystem II; PSII, standard thermoluminescence; PSII, half-life time; PSII, thermoluminescence; PSII, th

representing the largest group of biomass producers on earth (Bowler et al., 2010).

The oxygenic photosynthetic electron transfer (ET) chain is composed of three large multisubunit membrane-embedded complexes: photosystem II (PSII), cytochrome $b_{c}f$ (Cb $_{c}f$), and photosystem I (PSI), all of which contain iron cofactors and are connected by mobile electron carriers (Blankenship, 2014). In terrestrial plants, electrons from the Cb $_{c}f$ complex are transferred to PSI inside the thylakoidal lumen by plastocyanin (Pc), a small (ca. 100 amino acids) soluble Type-1 blue copper-protein. In contrast, electrons exiting PSI are delivered to ferredoxin (Fd), an iron–sulfur protein. However, in aquatic systems many photosynthetic microorganisms maintain alternative electron transporters couples (Hervás et al., 2003). Thus, in most cyanobacteria



and unicellular green algae, Pc is produced when copper is available, but under low-copper conditions Pc is replaced by cytochrome c_6 (Cc₆), a typical Class I c-type cytochrome (ca. 90 amino acids). In turn, the flavoprotein flavodoxin (Fld) replaces Fd under iron-limiting conditions (De la Rosa et al., 2006; Hervás et al., 2003; Sancho, 2006; Sétif, 2006). The functional and structural equivalence of both couples, Pc/Cc₆ and Fd/Fld, is well established (Bendall & Howe, 2016; Hervás et al., 1995, 2003; Hippler & Drepper, 2006).

The existence of redox alternative couples in photosynthesis can be understood as an adaptation to fluctuations in the bioavailability of iron in the aquatic environments. Many studies have revealed that iron limitation is the major factor controlling the primary productivity and growth of phytoplankton in vast oceanic areas (Boyd et al., 2007; Moore & Braucher, 2008). Massive oceanic iron fertilization experiments have also shown the appearance of algal blooms, in many cases dominated by diatoms, suggesting that these eukaryotic algae, in particular, are very sensitive to iron limitations (Morrissey & Bowler, 2012). Among other effects, iron starvation causes a severe decrease in the chlorophyll content and drastic alterations in the photosynthetic efficiency in diatoms, promoting a decrease in the content of both the iron-rich PSI and Cb_6f complexes and resulting in an increased PSII:PSI ratio (Allen et al., 2008; Lommer et al., 2012; Marchetti et al., 2012).

However, the occurrence of alternative couples is only partially maintained in the red lineage of photosynthetic organisms that diverged along evolution from the green lineage and that comprises, among others, red algae, cryptophytes, stramenopiles (that include diatoms), and haptophytes (Falkowski et al., 2004). Thus, whereas Fld is widely produced in the red lineage as an alternative to Fd under low iron conditions (Pierella Karlusich et al., 2014), Cc₆ is the only electron carrier from cytochrome f (Cf)-in the Cb₆f complex-to PSI in red algae and in the majority of organisms belonging to the red-plastid lineage, including diatoms, as confirmed by protein characterization, genome sequencing and transcriptomic analyzes (Akazaki et al., 2009; Bernal-Bayard et al., 2013; Blaby-Haas & Merchant, 2012; Bowler et al., 2010; Groussman et al., 2015). The absence of Pc as an alternative to Cc6 can thus represent an extra requirement of iron that affects the adaption of these organisms to iron limitations, as Cc6 replacement has been suggested to reduce iron requirements by up to 10% in open-ocean waters (Boyd et al., 2007; Guo et al., 2010). However, although petJ genes encoding Cc6 have been found in all the sequenced stramenopiles genomes, petE genes encoding Pc, closely related to those from green algae, have been also found in the genomes of several strains of oceanic diatom species. These include Thalassiosira oceanica and Fragilariopsis cylindrus, but also the haptophyte Emiliania huxleyi and the dinoflagellate Karenia brevis (Blaby-Haas & Merchant, 2012; Nosenko et al., 2006; Peers & Price, 2006). In addition, analyzes of microbial eukaryotic transcriptomes have detected transcripts encoding for putative Pcs in at least one species from each of the major diatom classes (Groussman et al., 2015). All of this has been attributed to the acquisition of petE Pc genes from green alga by horizontal gene transfer, as an adaptation to iron limitation (Groussman et al., 2015; Hippmann et al., 2017; Marchetti et al., 2012; Peers & Price, 2006). A particular strain of the open ocean diatom *T. oceanica* has been even described to express constitutively Pc as the main electron carrier from Cf to PSI, as the two Cc₆ genes of its genome are weakly expressed (Kong & Price, 2020; Lommer et al., 2012; Peers & Price, 2006). The presence of Pc in diatoms raises, however, some open questions, such as the mechanism regulating either its expression or a putative Cc₆/Pc alternation in function of the copper/iron levels (Hippmann et al., 2017; Lommer et al., 2012; Marchetti et al., 2012; Roncel et al., 2016). In addition, the existence of a still not found specific transporting system carrying copper into the lumen, to be incorporated into the Pc, is another unresolved question (Guo et al., 2010, 2015; Kong & Price, 2019; Levy et al., 2008).

Clear evidence for the production of a functional Pc protein has been only presented in the case of T. oceanica, where relatively small amounts per cell of the holoprotein were detected, purified and partially sequenced (Kong & Price, 2020; Peers & Price, 2006). However, there is a discordance between the theoretical molecular mass of the T. oceanica mature Pc (10,300 Da) and the apparent experimental mass found for the purified protein (18,200 Da) (Peers & Price, 2006). In any case, the Pc-translated sequence in diatoms is closely related to that from green algae, and thus the "red-type"-acquired Pc would show the features of a "green-type" Pc. In green algae (and plants) Pc. two well-defined functional areas can be described: (1) a hydrophobic ET area, determined by the accessibility to the solvent of the redox cofactor and (2) a strong negatively charged area for the interaction with complementary positive patches in Cf and PSI. Similar functional areas have been proved to be also present in green Cc6 (De la Rosa et al., 2006; Guss & Freeman, 1983; Hippler & Drepper, 2006; Ubbink et al., 1998). However, whereas Cc6 from red algae and diatoms still preserves the hydrophobic ET area, it shows a significant decrease in the negative character of the electrostatic area (Akazaki et al., 2009; Bernal-Bayard et al., 2013; Yamada et al., 2000), parallel with equivalent changes in the electrostatics features of PSI (Antoshvili et al., 2019; Bernal-Bayard et al., 2013, 2015) and Cf (unpublished data). Consequently, in vitro kinetic analyzes of the interaction of diatom PSI with green algae and plants Pcs, suggest that if an alternative Pc in diatoms could represent a favorable adaptation to iron limitation, its "green-type" character may, however, limit its ET efficiency. Remarkably, this efficiency increased in some green alga Pc mutants replacing negative charges for positive ones to approximate the electrostatics of diatom Cc₆ (Bernal-Bayard et al., 2015).

We have here investigated if the heterologous expression in the pennate model diatom *Phaeodactylum tricornutum* of a functional green alga Pc under iron-limiting conditions—as an alternative to Cc₆—can improve photosynthetic activity and cell growth. *P. tricornutum* is sensitive to iron limitation, and both lacks Pc and produces Cc₆ (Akazaki et al., 2009; Allen et al., 2008; Bernal-Bayard et al., 2013). For this purpose, we have carried out the construction of *P. tricornutum* strains expressing a mutant of the Pc from the green algae *Chlamydomonas reinhardtii*, that was previously shown to be more effective than the WT Pc in reducing diatom PSI in vitro (Bernal-Bayard et al., 2015).

2 | MATERIALS AND METHODS

2.1 | Cell cultures

Cells from the coastal pennate diatom Phaeodactylum tricornutum CCAP 1055/1 were grown in artificial seawater (ASW) medium (Goldman & McCarthy, 1978; McLachlan, 1964) in a rotatory shaker (50 rpm) at 20°C with regular transfer of the cells into fresh media. The cultures were illuminated by led white light (4500 K) lamps giving an intensity of 20 μmol m⁻² s⁻¹ (T8-150MWBL led lamps, Wellmax) following a light/dark cycle of 16/8 h. For the experiments to study the effects of iron deficiency, cells from 7 days old cultures were pelleted at 5000 g for 5 min and suspended and grown in fresh ASW medium under iron-replete (12 µM Fe) or iron-deficient (10 nM Fe plus 0.3 mg L^{-1} of copper) culture conditions, with transfer of the cells into the same fresh media after a week. Then, the cultures were grown for another 15 days. Except when indicated, most of the experiments were carried out using cultures with $\approx 6.5 \times 10^7$ cells ml⁻¹ and $\approx 2 \times 10^7$ cells ml⁻¹, respectively, for iron-replete and iron-deficient conditions.

2.2 | Plasmids construction

The nucleotide sequence of the WT Pc from the green alga-Chlamydomonas reinhardtii was retrieved from the GenBank-NCBI database. A synthetic Pc petE gene, with the E85K mutation (Bernal-Bayard et al., 2015), was designed with the codon usage of P. tricornutum (https://www.kazusa.or.jp/codon/). Moreover, the Fld promoter (induced in the absence of iron) and the transit peptide of Cc₆ (directing the protein to the thylakoid lumen) of P. tricornutum were added upstream to the Pc-translated sequence. However, as the native Cc6 gene of P. tricornutum has an intron (280 nucleotides) within the N-terminal plastid targeting sequence (Killian & Kroth, 2004), two constructions with alternative versions of the transit peptide of Cc6 were designed, with or without the sequence corresponding to the intron (Figure S1). The synthetic plasmids were generated by GeneCust (https://www.genecust.com/en/) by inserting the Pc petE E85K constructions into pBlueScript II SK (+) vector in EcoRV and Pstl sites. Constructions were then subcloned in the P. tricornutum transformation vector Isip1::YFP, kindly provided by the group of Prof. Chris Bowler (IBENS, Paris, France) (Kazamia et al., 2018), in the same restriction sites, to obtain the final pFld_PtCc6_PcE85K and pFld_PtVCc6_PcE85K plasmids (with and without the intron, respectively). Both plasmids were checked by sequencing (NZYTech), using the oligonucleotides pPisipF (5'-TCT CGTGTGGTCATGGGTTG-3') and pPisipR (5'-ACGACTTCTTCAA GTCCGCC-3'). To obtain antibodies against the Fld from P. tricornutum, the FLDA1 gene coding for Fld was first amplified by PCR from genomic DNA, using the iProof™ High-Fidelity DNA Polymerase (Bio-Rad) and adequate oligonucleotides. The PCR product was then cloned into the pET-28 b (+) vector (Novagen) to obtain the pET-28 b_Fld plasmid for expressing the His-tagged protein in E. coli. All plasmids were maintained in the *E. coli* DH5 α strain (Hanahan, 1983), isolated when required as described by Ish-Horowicz and Burke (1981) and quantified using a NanoDrop N-1000 spectrophotometer.

2.3 | Cell transformation by biolistic

P. tricornutum cells were transformed using the Biolistic PDS-1000/ He Particle Delivery System (Bio-Rad) as previously described (Apt et al., 1996; Falciatore et al., 1999). Approximately 5×10^7 cells, harvested at mid-logarithmic phase, were spread on ASW/1% (w/v) agar plates and incubated for 24 h. M17 tungsten microcarrier particles (1.1 µm diameter; Bio-Rad) were coated with plasmid DNA using CaCl₂ and spermidine as indicated by the supplier. Agar plates were located at level two within the chamber and a burst pressure of 1550 psi was used for the bombardment. The cells were cotransformed with the pPha-T1 vector containing zeocin resistance. also kindly provided by the group of Prof. Chris Bowler. As a control of the possible effects of the zeocin resistance, an experiment in which P. tricornutum was transformed only with the zeocin resistance vector was also carried out (B1 mutants). Afterward, the plates were incubated for 48 h in the dark before being spread onto new ASW agar plates containing 100 µg ml⁻¹ of zeocin (InvivoGen) to allow the selection of transformants. Plates were then maintained under illumination with a light/dark cycle of 16/8 h at 20°C as described above. Individual resistant colonies appearing 3-4 weeks after transformation were restreaked on new ASW agar plates containing zeocine. The presence of the Pc gene in zeocin-resistant colonies was checked by colony PCR, using the oligonucleotides pPisipF and pPisipR and the MyTag™ Polymerase (Bioline), according to manufacturer's instructions. The amplified products were visualized through DNA electrophoresis in 0.7% (w/v) agarose gel (Sambrook & Russell, 1989).

2.4 | Analytical methods

Growth cell parameters were obtained from at least 10 independent experiments. Cells were counted with a Neubauer-Improved hemocytometer (Marienfeld-Superior), according to the manufacturer's instructions. Cell specific growth rate (μ , day⁻¹) was calculated after 15 days of culture growth as: [μ = ln ($N_{\rm t}/N_{\rm 0}$) / Δ t] where $N_{\rm t}$ and $N_{\rm 0}$ are the final and initial cell concentration, respectively, and Δ t is the days of growth. Pigments extraction with acetone was carried out basically as described by Kong and Price (2020), but with an initial step of incubation at 70° C for 30 min. The total chlorophyll content in *P. tricornutum* cells was determined according to Arnon (1949). The chlorophyll a and c estimated content was determined as described by Jeffrey and Humphrey (1975), whereas carotenoid concentration was determined by the equation given by Strickland and Parsons (1972):

Chl
$$a \left(\mu g \, m l^{-1} \right) = 11.47 \times (A_{664} - A_{750}) - 0.40 \times (A_{630} - A_{759})$$



$$Chlc \left(\mu g \, ml^{-1} \right) = 24.34 \times \left(A_{630} - A_{750} \right) - 0.40 \times \left(A_{664} - A_{759} \right)$$

Carotenoid
$$\left(\mu g \, m I^{-1}\right) = 7.6 \times \left[\left(A_{480} - A_{750}\right) - 1.49 \times \left(A_{510} - A_{750}\right)\right]$$

For protein determination, *P. tricornutum* cells were pelleted and resuspended in $100-150 \,\mu$ l of lysis buffer [50 mM Tris–HCl, pH 7.5, buffer supplemented with 2% SDS, 10 mM EDTA and a tablet of the protease inhibitor cocktail (cOmpleteTM ULTRA Tablets, EASYpack, Roche)]. The cells were incubated at 37°C for 30 min with occasional vortexing. Soluble protein fractions were obtained by centrifugation at 15000 *g* for 30 min at 4°C, and the protein concentration was determined according to the Lowry method (Lowry et al., 1951).

The Cc₆ and Pc E85K content in P. tricornutum cells were determined following a modification of the method previously described (Bernal-Bayard et al., 2017). P. tricornutum cells were collected by centrifugation. Wet pellets (≈3 g) were weighed and resuspended in 20 ml of 20 mM Tris buffer, pH 7.5, supplemented with DNase and the proteases inhibitors PMSF, benzamidine and aminocaproic acid. The cells were then broken by three cycles of French press disruption (20,000 psi), followed by debris precipitation with 55% ammonium sulfate and further precipitation with 100% ammonium sulfate. Cc₆ and Pc content were determined in the final resuspended pellets by differential absorbance measurements between the fully reduced (sodium ascorbate) and fully oxidized (potassium ferricyanide) states at 552 nm (Cc6, reduced minus oxidized) or 600 nm (Pc, oxidized minus reduced), as previously described (Bernal-Bayard et al., 2015; Roncel et al., 2016). To reduce interferences from Cc₆ absorbance in the determination of Pc, the concentration of Cc6 was first determined and an equivalent concentration of purified Cc₆ was used as a baseline to measure Pc concentration more precisely. P. tricornutum Fld purification was carried out by FPLC (IBVF Chromatography Service) using a HisTrap Fast Flow Crude column (GE Healthcare Life Sciences) from the inclusion bodies of E. coli BL21 (DE3) cells transformed with pET-28 b Fld, after IPTG induction in the presence of 10 µM FMN. Purification then continued by following, with some modifications, protocols previously described (Purifying Challenging Proteins Handbook from GE Healthcare, and see also Rogl et al., 1998). MALDI-TOF mass spectrometry (IBVF Proteomic Service) was used to confirm the correct molecular weight of the purified protein.

2.5 | Western-blot analysis

For the immunodetection of *C. reinhardtii* Pc and *P. tricornutum* Cc_6 , cytochrome c_{550} (Cc_{550}) and Fld, polyclonal antibodies raised against those proteins were generated using standard procedures at the Animal Experimentation Facility (University of Seville, Spain), by subcutaneous injection of 1 mg of the purified proteins into a white New Zealand rabbit (Bernal-Bayard et al., 2013, 2015, 2017). Commercial antibodies against D1 and PsbO proteins were obtained from Agrisera (products no. ASO5 084A and ASO6 142–33, respectively),

whereas antibodies against PsaB were kindly provided by Dr. Anna Lindahl (IBVF, Sevilla, Spain).

Protein extracts (2.5–30 μg) were resolved on 15% (w/v) polyacrylamide gel electrophoresis, as previously described (Roncel et al., 2016), and transferred to a 0.2 μm nitrocellulose membrane included in the Trans-Blot® Turbo™ RTA Midi Nitrocellulose Transfer Kit through the Trans-Blot® Turbo™ Transfer System (Bio-Rad), following the manufacturer's instructions. The membrane was incubated overnight with a dilution (1:500 to 1:10000) of the selected primary antibody followed by 1 h incubation with a 1:10000 dilution of the secondary antibody (Goat Anti-Rabbit IgG [H + L]-HRP Conjugate, from Bio-Rad). Finally, the membrane was treated with the Immobilon Western Chemiluminescent HRP Substrate (Millipore) according to the manufacturer's instructions and visualized with the ChemiDoc™ Imaging System (Bio-Rad). Ponceau total protein stainings were used as loading controls in the immunoblottings.

2.6 | Microscopy

To monitor the Pc expression and localization in *P. tricornutum* transformed strains as well as cell morphology of the different strains, cells were observed and photographed using a Leica Microscope DM6000 B apparatus (IBVF Microscopy Service). The immunostaining and fluorescence microscopy were carried out following the protocol previously described in Szabo and Colman (2007), with some modifications. Cells were extended in a slide and fixed, incubated with rabbit polyclonal anti-Pc antibody, washed, incubated with anti-rabbit FITC secondary antibody (Sigma), washed again and mounted.

The morphology of WT and transformed cells of *P. tricornutum*, grown under iron-replete or iron-deficient conditions, was imaged in vivo also by fluorescence microscopy. Cell volumes and surfaces were calculated from images measurements of longitudinal and transversal cell dimensions from at least 30 cells, and treating *P. tricornutum* as a two cones model (Levy et al., 2008). In addition, the cell aspect ratio ([Major Axis]/[Minor Axis]) and circularity ($4\pi \times ([Area]/[Perimeter]^2))$ shape descriptors were measured by ImageJ Fiji (v. 1.46; Schindelin et al., 2012).

2.7 | Thermoluminescence analyzes

Thermoluminescence (TL) glow curves of *P. tricornutum* cell suspensions were obtained using a home-built apparatus designed by Dr. Jean-Marc Ducruet for luminescence detection from 1 to 80°C (standard thermoluminescence, STL) and from 10 to 160°C (high temperature thermoluminescence, HTL). A detailed description of the system can be obtained elsewhere (García-Calderón et al., 2019; Roncel et al., 2016). Temperature regulation, signal recording, and flash sequences were driven by a computer through a National Instrument DAQ-Pad1200 interface, using a specific acquisition program developed by Dr. Ducruet. An electrically insulated resistor heater (Thermocoax), powered by a variable (0 to 5 A) computer-driven power supply, was mounted below the chamber for temperature regulation. STL and HTL measurements were carried out as described in Roncel

et al. (2016). TL experiments were carried out in parallel cultures for each strain under iron-replete and iron-deficient conditions. TL parameters were obtained from three independent measurements. Most of the experiments were performed using cultures with a cellular concentration of $\approx 1.5 \times 10^8$ and $\approx 5 \times 10^7$ cells ml⁻¹ for STL and HTL, respectively.

2.8 Photosynthetic measurements

To study photosynthetic global activity, cells from 2.5 ml of the different cultures were collected by centrifugation and resuspended in fresh culture medium. Oxygen intake or evolution rates were then determined from three independent experiments by using a Clarktype oxygen electrode (Hansatech). Measurements were carried out at 20°C, both in the dark and under illumination (176 μmol m⁻² s⁻¹), to establish the net photosynthetic activity per cell.

The chlorophyll a fluorescence of PSII and the redox state of P₇₀₀ from intact cells were determined at room temperature using a pulseamplitude modulation fluorometer (DUAL-PAM-100, Walz). The photosynthetic parameters were obtained from five independent experiments, carried out basically as described in Roncel et al. (2016). Prior to measurements, cell suspensions with a cellular concentration of 1.1×10^8 cells ml⁻¹ were dark-adapted for 30 min. The maximum quantum yield of PSII (F_v/F_m) , the relative linear electron transport rates (rETR) and the effective PSII quantum yield [Y(II)] for each actinic light intensity were determined as previously described (Roncel et al., 2016).

The redox state of P₇₀₀ was monitored by following the changes in absorbance at 830 nm versus 875 nm in cells dark-adapted for 30 min prior to measurements. Most of the experiments were carried out using cultures with a cellular concentration of $\approx 5.5 \times 10^7$ and \approx 4.5 \times 10⁸ cells mL⁻¹ for iron-replete and iron-deficient cultures, respectively. The level of maximal P₇₀₀⁺ signal observed upon P₇₀₀ full oxidation, $P_{\rm m}$, was determined by pre-illumination of cell suspensions with FR light (730 nm) for 10 s and after that with a saturating pulse of red light (635 nm) at 10000 μ mol m⁻² s⁻¹ intensity and 0.2 s duration. Following P_m determinations, P₇₀₀⁺ reduction kinetic decays were recorded in darkness. The average of at least three individual traces from different cultures was taken. The half-life time ($t_{1/2}$) of the P₇₀₀⁺ absorption decay was determined by fitting the decays to a single exponential function. The quantum yields of PSI photochemistry, Y(I), donor side limitations, Y(ND), and acceptor side limitations, Y(NA) were determined as described in Roncel et al. (2016).

RESULTS 3

The E85K mutant Pc of C. reinhardtii can be heterologously expressed in the chloroplast of P. tricornutum

The expression of Pc in some oceanic species of diatoms has been proposed to constitute an alternative replacement of Cc6, similar to what occurs in cyanobacteria and green algae (Groussman et al., 2015; Hippmann et al., 2017; Lommer et al., 2012; Marchetti et al., 2012; Moore & Braucher, 2008; Peers & Price, 2006). However, there are no available data about the in vivo functional effects of expressing Pc in the thylakoidal lumen of diatoms under iron-limiting conditions. Thus, here we investigated the effects of the heterologous expression of a Pc in P. tricornutum, a diatom that naturally only produces Cc6. We have selected the E85K mutant of the Pc from the green alga C. reinhardtii to be expressed in P. tricornutum, as this mutant has previously shown to be the most effective Pc in reducing diatom PSI in vitro (Bernal-Bayard et al., 2015). The C. reinhardtii E85K Pc gene was placed under the control of the Fld promoter to ensure its expression in low iron conditions (Figure 1A), in which the levels of native Cc6 drastically decrease and become limiting for the photosynthetic activity (Roncel et al., 2016, and see below). The tripartite transit peptide of the native Cc6 was added to the Pc gene sequence to warrant the transit of the protein to the thylakoid lumen (Killian & Kroth, 2004). As the native Cc_6 gene of P. tricornutum has an intron (280 nucleotides) located within the region coding for the Nterminal plastid targeting sequence (Killian & Kroth, 2004), a version of the transit peptide of Cc6 without the sequence corresponding to the intron was also alternatively used (Figure S1).

The presence of the Pc gene in zeocin-resistant transformed colonies was checked by colony PCR and DNA electrophoresis in agarose gels (not shown). The nuclear transformation of P. tricornutum is random, which can result in a great variability in the expression of a protein in different clones. Consequently, several P. tricornutum strains with different levels of Pc expression were selected (clones #30, #45

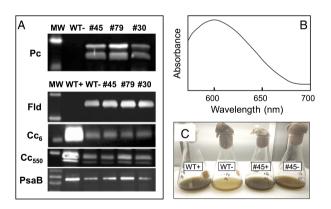


FIGURE 1 (A) Western blot analysis of different photosynthetic proteins in samples from P. tricornutum WT and #30, #45, and #79 mutant cells under iron-deficient conditions. MW, molecular weight standard; WT+ and WT-, control WT cells under iron-replete or irondeficient conditions, respectively; Cc₆, soluble cytochrome c₆; Cc₅₅₀, the extrinsic cytochrome c_{550} subunit of PSII; Fld, flavodoxin; Pc, E85K plastocyanin; PsaB, the membrane intrinsic subunit of PSI. Protein extracts: 30 μg for Pc; 20 μg for PsaB; 10 μg for Cc₆; 5 μg for Fld; 2.5 μ g for Cc₅₅₀. Membranes were incubated overnight with the following dilutions of the selected primary antibody: 1:500 anti-Pc; 1:1000 anti-Cc₆, Cc₅₅₀ and PsaB; 1:50000 anti-Fld. (B) Spectroscopic monitoring of holo-Pc (ferricvanide minus ascorbate) in cell extracts from mutant #45 under iron-deficient conditions. (C) Examples of WT and mutant #45 individual cultures under iron-replete (+) or irondeficient (-) conditions

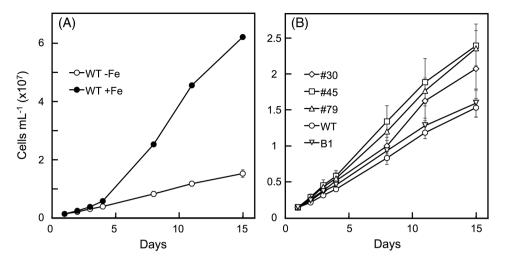


FIGURE 2 (A) Growth of P. tricornutum WT cultures under iron-replete (+Fe) or iron-deficient (-Fe) conditions. (B) Growth of WT, #30, #45, and #79 strains cultures under iron-deficient conditions. Mean values ± sD of measurements from 10 independent cultures are presented

and #79), as shown by the presence in western-blot analysis of a band corresponding to the expected size of Pc (10.2 kDa; Figure 1A). The following apparent progression in the levels of expression was observed: #30 < #45 < #79 (Figure 1A). Whereas mutants #30 and #45 were transformed with the construction that contained the intron in the plastid targeting gene sequence, mutant #79 lacked the intron (Figure S1), indicating that both constructions can be correctly processed. However, in the three mutant strains an additional band of an apparent higher size (12.6 kDa) was also observed (more abundant in the clone #79; Figure 1A). This band can be tentatively assigned to unprocessed Pc molecules still containing the Cc6 thylakoid transit peptide (24 extra amino acids; Killian & Kroth, 2004) and that would be considered as apo-Pc. Fluorescence microscopy monitoring of immunolabeled cells confirmed the detection of the heterologous Pc in the chloroplast, as the green fluorescence associated to the immunodetection of Pc merged with the red fluorescence of photosynthetic pigments (not shown). Finally, the presence of the Pc holoprotein (i.e., with the copper active cofactor) was directly detected by its absorption spectrum in cell extracts (Figure 1B). From the recorded differential spectra, analogous amounts of intracellular holo-Pc were estimated for mutants #45 (≈3.8 µM) and #79 (\approx 4.4 μ M) (Figure 1B), whereas the lower levels of Pc detected in the mutant #30 did not allow obtaining a reliable estimate of the holoprotein content. On the other hand, a similar amount of $\approx 3.6 \mu M$ of intracellular Cc6 was estimated for the WT and mutant strains under iron-deficient conditions according to specific spectral changes (Figure S2), which represents \approx 12% of the concentration detected under iron-replete conditions (\approx 30 μ M) (Bernal-Bayard et al., 2013).

3.2 | Relatively low intracellular concentrations of heterologous Pc are enough to promote an increased growth in *P. tricornutum* under iron-deficient conditions

It should be noted that, under iron-replete conditions, the transformed strains did not show significant differences with the WT strain (not shown), nor did the B1 strain (with the resistance to zeocin but

without Pc) in iron-deficient conditions (Figure 2B). However, the three selected transformed strains showed an increased cell growth under iron-deficient conditions compared with the WT (and B1) strain (Figures 1C and 2B). This increase was about 40% in the mutant #30, but up to 60% in the case of the #45 and #79 mutants (Figure 2B and Table 1), although the growth of the three transformed strains was still sensibly lower as compared with iron-replete conditions (Figure 2). Accordingly, the three transformed strains also showed higher values in the specific growth rate (µ, from 0.155 day⁻¹ in the WT to 0.175-0.185 day⁻¹ in the mutants), as well as in the total chlorophyll, chlorophyll a and c and carotenoid estimated contents per ml, as compared with WT cells under iron-deficient conditions (Table 1). Higher levels of pigments are still observed in the three mutants when chlorophyll content is normalized per cell, particularly in the mutants #45 and #79 (up to 40%; Table 1). Regarding global photosynthetic activity, the three transformed strains presented a significant enhancement in the net photosynthetic oxygen evolution normalized per cell compared with the WT strain, with the highest values observed again for the #45 and #79 clones (Table 1).

Fluorescence microscopy analysis showed changes in cell morphology in the three mutant strains (Figure S3 and Table 1). Thus, under iron-deficient conditions, WT fusiform cells showed to be more elongated than in the mutant strains (aspect ratio of 6.52, 4.89, 4.25, and 5.15 for WT, #30, #45, and #79 cells, respectively). Consequently, the mutant strains also appeared with an increased circularity, which actually resulted in an increased cell volume (Table 1) and chloroplasts with a thicker shape (Figure S3).

The changes in the levels of some photosynthetic proteins under low iron conditions were visualized by western-blot analysis. In addition of Fld (used as a control of iron deficiency), the studied proteins included soluble Cc_6 , PSII subunits (the intrinsic D1 subunit and the luminal extrinsic subunits cytochrome c_{550} –or Cc_{550} – and PsbO), as well as the intrinsic PsaB subunit of PSI. The deficiency of iron promotes a drastic decrease in the levels of the two luminal cytochromes, Cc_6 and Cc_{550} , more particularly in the case of Cc_6 , as previously described at less drastic iron limitations (Bernal-Bayard et al., 2017; Roncel et al., 2016) (Figure 1A). Western bands quantification allowed estimating a similar value of $\approx 10\%$ in the levels of Cc_6 and $\approx 30\%$ in

TABLE 1 General physiological and biochemical parameters of *P. tricornutum* transformed cells and cultures under iron-deficient conditions

Parameter (after 15 days of growth) ^a	WT	#30	#45	#79
Cells per ml	$(1.5 \pm 0.1) \times 10^7$	$(2.1 \pm 0.3) \times 10^7$	$(2.4 \pm 0.3) \times 10^7$	$(2.5 \pm 0.2) \times 10^7$
	$(100\%)^b$	(140%)	(159%)	(163%)
Specific growth rate, μ (day $^{-1}$)	0.155 ± 0.005	0.175 ± 0.008	0.183 ± 0.010	0.185 ± 0.013
	(100%)	(113%)	(118%)	(119%)
Total ChI per ml (µg ml ⁻¹)	4.19 ± 0.23	6.05 ± 0.06	8.13 ± 0.07	8.14 ± 0.06
	(100%)	(144%)	(194%)	(194%)
Chl a per ml (μ g ml ⁻¹)	2.97 ± 0.03	4.45 ± 0.09	6.12 ± 0.05	6.30 ± 0.02
	(100%)	(150%)	(206%)	(212%)
Chl c per ml ($\mu g m^{-1}$)	1.49 ± 0.04	2.24 ± 0.02	3.22 ± 0.04	3.14 ± 0.06
	(100%)	(150%)	(216%)	(211%)
Total carotenoids per ml ($\mu g \ ml^{-1}$)	2.04 ± 0.07	3.50 ± 0.15	4.81 ± 0.65	4.34 ± 0.41
	(100%)	(172%)	(236%)	(213%)
Total Chl per cell (pg cell ⁻¹)	0.27	0.32	0.38	0.37
	(100%)	(120%)	(140%)	(137%)
Total carotenoids per cell (pg cell ⁻¹)	0.13	0.16	0.20	0.18
	(100%)	(125%)	(150%)	(135%)
Cell circularity	0.33 ± 0.04	0.43 ± 0.07	0.44 ± 0.05	0.38 ± 0.05
	(100%)	(130%)	(133%)	(115%)
Cell volume (μm³)	40.2 ± 8.6	35.3 ± 9.5	55.8 ± 14.8	53.2 ± 10.9
	(100%)	(90%)	(139%)	(132%)
Cell surface (μm²)	85.1 ± 12.8	71.4 ± 13.6	92.8 ± 13.8	95.2 ± 11.4
	(100%)	(84%)	(110%)	(112%)
Net photosynthetic rate (nmol $O_2 h^{-1}$ per 10^6 cells)	37.2 ± 0.6	45.4 ± 2.5	78.5 ± 8.8	87.0 ± 1.6
	(100%)	(122%)	(213%)	(235%)

^aSee the Materials and methods section for more details.

Cc₅₅₀ in all the strains as compared with iron-replete conditions, in good agreement with the spectroscopic measurements (see above). Regarding other PSII components, the levels of PsbO and D1 were basically analogous in the WT and mutant strains under iron deficiency (Figure S4). However, in the case of the PsaB subunit of PSI, the decrease observed in the WT strain under iron-deficient conditions (to $\approx\!40\%$ of the iron-replete levels) seems to be somewhat recovered in the #45 and more clearly in the #79 mutant strains (Figure 1A). However, it is important to note that an increased level of PsaB does not necessarily imply an equivalent amount of active PSI, since other PSI iron-containing subunits are required to form functional PSI complexes (see below).

3.3 | P. tricornutum cells expressing Pc under low iron concentration show a lower decrease in PSII activity and a lower level of lipid peroxidation compared with WT cells

The effects of iron deficiency on PSII ET activity of *P. tricornutum* WT and mutant strains were investigated using the STL technique (Figure 3). Excitation of iron-deficient *P. tricornutum* WT cells with two flashes at 1°C induced the appearance of a TL glow curve with significant differences in comparison with the curves obtained in iron-replete cells: a very significant decrease in the total TL signal intensity of about 65% (Figure 3A). The decomposition analysis of these

emission curves allowed obtaining two components with similar $t_{\rm max}$ values of 14 and 28°C for both iron conditions. These two components can be assigned to the well-known B1 and B2 TL bands, originating from the recombination reactions of $S_3Q_B^-$ and $S_2Q_B^-$ charge pairs in PSII, respectively (Rutherford et al., 1984).

The TL glow curves obtained for the #79 (Figure 3A), #30 and #45 (not shown) mutant strains could be also decomposed in two TL components: B1 and B2 bands. The $t_{\rm max}$ values for these components were similar to WT in all strains and for the two iron conditions. The TL glow curves obtained in iron-deficient cells also showed a decrease in the total TL signal intensity with respect to iron-replete conditions (Figure 3). However, the comparison between conditions of iron sufficiency or limitation for each strain indicated a much lower decrease in intensity in the mutants compared with the WT strain: 39%, 56%, and 18% for #30, #45, and #79 mutant strains, respectively (Figure 3B).

The HTL technique was applied to detect lipid peroxidation in *P. tricornutum* WT and mutant cells, cultured in both iron-replete and iron-deficient conditions. Figure 4A shows that a broad HTL2 band with a maximum between 130–140°C was present in iron-deficient WT cells. This band was significantly lower in iron-replete cells. Thus, an important level of lipid peroxidation was observed in WT cells grown in iron-deficient conditions under the relatively low light intensity used. However, for the three mutant strains cultured under iron-deficient conditions, the level of lipid peroxidation was significantly lower (Figure 4B).

^bBelow, in parentheses, values referred to as the percentage of that of WT cells.

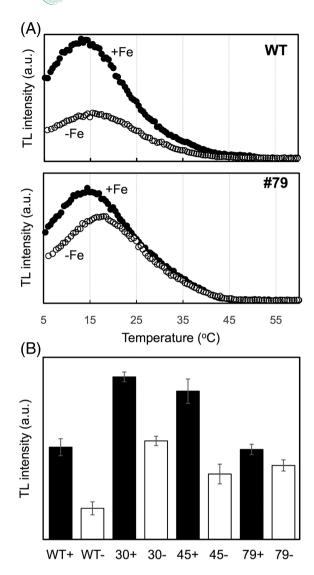


FIGURE 3 Effects of iron deficiency on the PSII activity of *P. tricornutum* cells measured by the STL technique. (A) Representative standard TL glow curves of the WT and #79 strains for iron-deficient (–Fe) and iron-replete (+Fe) conditions, as indicated. (B) Intensities of TL B1 and B2 bands for all the strains investigated under iron-replete (+) or iron-deficient (–) conditions, as indicated. Intensities were obtained from the component analysis of the curves of TL. Data in (B) represent mean values ± SD of three independent measurements

3.4 | Higher maximum quantum yields of PSII and relative electron transport rates are observed by fluorescence measurements in *P. tricornutum* cells expressing Pc under iron-deficient conditions

Differences in the photosynthetic activity of PSII between iron-replete and iron-deficient WT cells have been already described (Roncel et al., 2016). Thus, the effects of iron deficiency on PSII photochemistry in the mutants expressing Pc have been here investigated by chlorophyll a fluorescence measurements. The maximum quantum yield of PSII, $F_{\rm v}/F_{\rm m}$, was significantly decreased in iron-deficient WT cells ($F_{\rm v}/F_{\rm m}=0.331$) in comparison with iron-replete cultures ($F_{\rm v}/F_{\rm m}=0.331$)

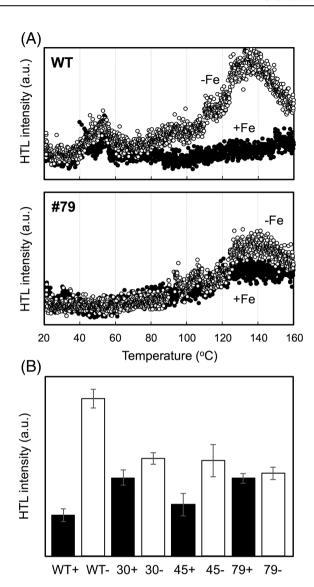


FIGURE 4 Effects of iron deficiency on the lipid peroxidation of *P. tricornutum* WT and mutant strains measured by the HTL technique. (A) Representative high-temperature TL glow curves (HTL2 band) of WT and #79 mutant strains for iron-replete (+Fe) or iron-deficient (-Fe) conditions, as indicated. (B) Intensities of HTL2 bands for all the strains investigated under iron-replete (+) or iron-deficient (-) conditions, as indicated. Intensities were obtained from the component analysis of the curves of HTL. Data in (B) represent mean values ± SD of three independent measurements

 $F_{\rm m}$ = 0.632) (Table 2). However, the three mutants showed a partial recovery (\approx 15–30%) of the $F_{\rm v}/F_{\rm m}$ values as compared to WT cells under conditions of iron deficiency, with higher values being obtained for the #45 (0.384) and #79 mutants (0.425) (Table 2).

It has been previously shown in cultures of *P. tricornutum* (Roncel et al., 2016) that the effective quantum yield of PSII, Y(II), is significantly lower in iron-deficient cultures as compared with iron-replete conditions at the range of irradiances used; iron-deficient cells being more sensitive to photoinhibition (Roncel et al., 2016). In our case, PSII light saturation curves carried out under iron deficiency have shown Y(II) values of a similar magnitude for the mutant strains compared to WT cells, although slightly higher values were

TABLE 2 Maximum quantum yield of PSII (F_V/F_m), photosynthetically active radiation intensity (PAR) for relative maximum electron transport rate (rETR_{max}), relative electron transport rate (rETR) at the maximum photosynthetically active radiation intensity (PAR_{max}), maximal P_{700}^{+} signal upon full oxidation ($P_{\rm m}$), and half-life time ($t_{1/2}$) of the ${P_{700}}^+$ re-reduction in iron-replete (+Fe) WT and iron-deficient (-Fe) WT cells and the #30, #45, and #79 mutants

Parameter	WT + Fe	WT-Fe	#30-Fe	#45-Fe	#79-Fe
F_{v}/F_{m}	0.632 ± 0.004	0.331 ± 0.014	0.381 ± 0.055	0.384 ± 0.036	0.425 ± 0.044
PAR for ETR _{max} (μ mol m ⁻² s ⁻¹) ^a	500	150	200	350	350
rETR (at PAR _{max}) ^b	28.32	3.90	5.65	7.27	7.67
P _m	0.683 ± 0.043 (–)	0.093 ± 0.011 (100%) ^c	0.110 ± 0.008 (118%) ^c	0.105 ± 0.003 (113%) ^c	0.112 ± 0.007 (120%) ^c
$t_{1/2}$ (ms)	4.45 ± 0.23	11.82 ± 0.93	5.35 ± 0.35	6.06 ± 0.37	5.84 ± 0.35

^aLight intensity for rETR_{max}.

observed in the mutants at lower light intensities (Figure S5). On the other hand, it has also been shown that P. tricornutum cells showed differences in the relative electron transport rate (rETR, i.e., the ratio between absorbed light quanta and transported electrons) under iron-replete and iron-deficient conditions (Roncel et al., 2016). In cells exposed to gradually increasing light intensities, the electron transport increases in parallel up to its limit capacity. Beyond this point, increasing light intensity induces photoinhibition, that is, a decrease of the rETR (Bailleul et al., 2015; Roncel et al., 2016). In this work, iron-deficient WT cells reached its maximum electron transport rate (rETR_{max}) at 150 μ mol m⁻² s⁻¹, showing inhibited rETR values at higher light intensities (about 30% of the maximum value at $\approx 1300 \, \mu \text{mol m}^{-2} \, \text{s}^{-1}$; Figure 5 and Table 2). In contrast, iron-replete WT cells have showed a rETR_{max} at $500 \mu mol m^{-2} s^{-1}$ (Table 2) and, under these conditions, the rETR value measured at the higher light intensity remained at about 85% of the maximum value (not shown). Therefore, these results confirmed a substantially higher sensitivity to light for P. tricornutum WT cells grown under low iron concentration (Roncel et al., 2016). Table 2 shows the higher values of photosynthetically active radiation (PAR) observed for the rETR_{max} for the three mutants (reached at 200-350 μ mol m⁻² s⁻¹), as compared to that obtained in the WT (150 μ mol m⁻² s⁻¹) under the same conditions. In addition, the three mutants showed higher rETR values at light intensities above 500 $\mu mol~m^{-2}~s^{-1}$ and up to $\approx 1300~\mu mol~m^{-2}~s^{-1}$, at which the rETR values still remained at about 50-65% of the maximum value. Higher values were again observed for the #45 and #79 mutants (Figure 5 and Table 2). The rETR values reported here under irondeficient conditions are similar to those previously described in P. tricornutum under a less severe iron-limitation (Roncel et al., 2016) and respiratory inhibition (Bailleul et al., 2015), but lower than those obtained in different culture and experimental conditions in response to CO₂ limitation (Shimakawa et al., 2017). In any case, our results suggest that under iron deficiency, the three mutant strains exhibit better light tolerance, confirming the specific effect of expressing Pc in preventing earlier photoinhibition. This is also in line with the observed faster growth rate of the three mutants.

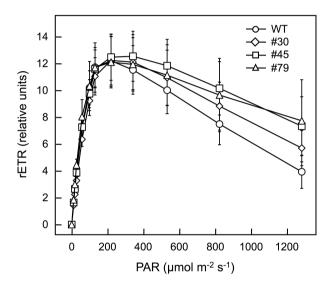


FIGURE 5 Relative linear electron transport rate (rETR) in the WT and #30, #45, and #79 mutant strains of P. tricornutum cells under iron deficiency, as indicated. rETR values were determined as a function of irradiance derived from steady-state light curves. Chlorophyll fluorescence was measured with a pulse-amplitude modulation fluorometer and rETR values were determined during stepwise increasing photosynthetically active radiation (PAR) from 0 up to 1292 μ mol m $^{-2}$ s $^{-1}$ light intensity. Data represent mean values ± SD of five independent measurements

The heterologous expression of Pc in P. tricornutum promotes an increase in the maximum quantum yield of PSI and faster P₇₀₀ reduction kinetics

The effects of Pc expression on PSI activity under iron deficiency were also investigated by measuring the P₇₀₀ redox state changes during illumination, as previously described (Roncel et al., 2016). In dark-adapted cultures, P₇₀₀ is reduced since the acceptor side of P₇₀₀, that is, the Calvin-Benson cycle and subsequent reactions, is deactivated. Under actinic light, P₇₀₀ is first oxidized and re-reduced by electrons coming from the plastoquinone (PQ) pool; thus, by applying saturating pulses, its ability to become oxidized and re-reduced can be determined (Roncel et al., 2016). Induction-recovery curves

^brETR values at 1292 μmol m⁻² s⁻¹ light intensity.

^cValues as percentage of that of WT-Fe cells.

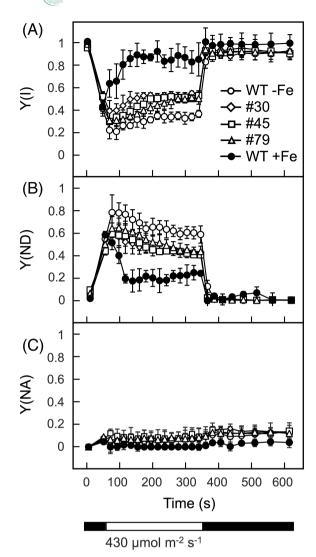


FIGURE 6 (A-C) PSI activity of WT (WT -Fe) and #30, #45, and #79 mutant cells of P. tricornutum under iron deficiency. Values of WT cells in iron-replete conditions (WT +Fe) are also included as a control. The redox state of the PSI reaction center, P₇₀₀, was monitored through the changes in absorbance at 830 nm versus 875 nm, measured with a pulse-amplitude modulation fluorometer. Cultures of the different strains were kept in the dark for 30 min prior to the measurements. After the initial determination of the maximal oxidation of P₇₀₀, actinic light was turned on at an intensity of 430 μmol m⁻² s⁻¹ and saturating pulses were applied every 20 s. after 5 min, the actinic light was switched off and measurements continued for another 5 min. Changes of (A) quantum yields of PSI, Y(I), (B) donor side limitations, Y(ND), and (C) acceptor side limitations, Y(NA), during the course of the induction curve are displayed. Data represent the mean values ± SD of five independent measurements. White and black bars below graphs indicate periods of illumination with actinic light and darkness, respectively

were performed in iron-replete and iron-deficient cultures showing that the calculated quantum yield of PSI photochemistry (Y(I); Figure 6A) increased in the three mutants compared to WT cells under the same conditions of iron deficiency. On the other hand, the high degree of donor side limitations, Y(ND), observed in iron-deficient WT cells, was slightly reversed in all the mutants (Figure 6B).

The higher Y(I) of the three mutants seems to indicate a recovery of the PSI activity missing in the WT cells as a result of the lack of availability of electron donors for PSI (Roncel et al., 2016). In contrast, similar acceptor-side limitations, Y(NA), were observed in all the cultures (Figure 6C). Thus, the presence of Pc in the three mutants seems to compensate for the deficiency in PSI activity by providing electrons to PSI in the light. To confirm this hypothesis, P_{700} reduction kinetics after saturating light pulses were recorded to obtain the maximal P_{700}^+ signal upon full oxidation (P_m) and the half-life time ($t_{1/2}$) parameters for P₇₀₀⁺ re-reduction (Table 2). After illumination with FR light and thereafter with a saturating light pulse, P₇₀₀ became oxidized and reached a maximal level of P₇₀₀⁺ (Figure S6). As expected, the P_m signal amplitude was significantly lower in WT iron-deficient cells than in iron-replete conditions (Table 2). However, under conditions of iron deficiency, the three mutants exhibited a slightly higher P_m signal compared to WT cells, thus indicating a somewhat higher amount of active P_{700} (Table 2). Following $P_{\rm m}$ determinations, P_{700}^{+} reduction kinetic decays were recorded in darkness and $t_{1/2}$ for the P_{700}^{+} absorption decay could be calculated (Figure S6 and Table 2). Thus, the value of $t_{1/2}$ in WT iron-deficient cultures was ≈ 3 times higher compared with iron-replete conditions (11.82 versus 4.45 ms; Table 2). However, under iron deficiency, $t_{1/2}$ values were \approx 2 times lower in the three mutants (≈5.8 ms) in comparison with WT cells (Table 2), thus indicating an acceleration of P₇₀₀⁺ re-reduction in the three mutants as compared with iron-deficient WT cells (Table 2).

4 | DISCUSSION

Adaptations of diatoms to natural oceanic low-iron environments include, among other effects, the induction of iron-uptake systems and the lowering of iron-containing photosynthetic proteins. In particular, a decrease in the levels of Fd (replaced by Fld) and the PSI and Cb₆f complexes has been reported, thus decreasing iron requirements (Lommer et al., 2012; Strzepek & Harrison, 2004). In addition, the presence of petE Pc genes in the genomes of some oceanic diatoms has been explained by their acquisition from green alga by horizontal gene transfer, as an adaptation to chronic iron limitations in these environments (iron concentration < 1 nM) (Moore & Braucher, 2008). Thus, Pc acquisition and expression in oceanic diatoms has been considered to represent an alternative (or even constitutive) substitution of Cc6, as it occurs in cyanobacteria and green algae (Groussman et al., 2015; Hippmann et al., 2017; Lommer et al., 2012; Marchetti et al., 2012; Moore & Braucher, 2008; Peers & Price, 2006). An open question is the mechanism regulating a putative Cc₆/Pc replacement in diatoms. In cyanobacteria and green algae, the copper levels determine the expression of one protein or another (Kropat et al., 2005; Merchant & Bogorad, 1986; Sandmann, 1986). In diatoms, however, both low copper and sufficient iron conditions have been reported to induce a decrease in Pc transcripts which, on the contrary, can be detected under iron-limiting conditions (Hippmann et al., 2017; Lommer et al., 2012; Marchetti et al., 2012; Rizkallah et al., 2020). Recently, a threefold decrease in the Pc protein content under copper deficiency has been reported in T. oceanica (Kong & Price, 2020). This

is in contrast to the drastic changes in Pc content in response to changes in copper concentration observed in cyanobacteria and green algae (Durán et al., 2005; Merchant & Bogorad, 1986; Sandmann, 1986). By its turn, our results indicate that Cc₆ levels in P. tricornutum depend on the availability of iron (Roncel et al., 2016), although iron uptake itself is connected to copper availability (Maldonado et al., 2006).

Although a Pc holoprotein was purified from *T. oceanica* (Peers & Price, 2006), this protein has neither been fully characterized nor cloned and expressed in *Escherichia coli*. Therefore, there are no functional data available for the interaction of the "red" partners (PSI or Cf) with the "green-type" Pc acquired by horizontal gene transfer. However, in vitro kinetic analysis of the interaction of *P. tricornutum* PSI with green algae and plants Pc indicated that diatom PSI is able to react with the very acidic Pc from these organisms, although with lower efficiency than with *P. tricornutum* Cc₆. Moreover, in the *C. reinhardtii* E85K Pc mutant designed to mimic the electrostatics of diatom Cc₆, this efficiency doubled as compared with the WT protein (Bernal-Bayard et al., 2015).

The estimated holo-Pc concentration per cell in T. oceanica under copper-sufficient conditions (≈3 µM) is about 10 times lower than that usually described for Cc6 under iron-sufficient conditions, including reported values in the coastal P. tricornutum diatom (≈28 μM) (Bernal-Bayard et al., 2013; Kong & Price, 2020; Peers & Price, 2006). However, Pc levels in T. oceanica are comparable to the Cc₆ content found under iron-deficient conditions in P. tricornutum (≈4-7 μM) (see above, and Roncel et al., 2016) and, interestingly, to the levels of heterologous holo-Pc here detected (≈4 μM). In this sense, although copper uptake systems have been described in T. oceanica (Kong & Price, 2019), the functional expression of Pc requires a specific copper-transporting system into the thylakoid that has not yet been found in red-type organisms. A copper-trafficking system could expect to include chaperones for copper traffic and, in particular, a Ptype ATPase transporter located in the thylakoid membrane, delivering copper to the lumen for incorporation into Pc (Blaby-Haas & Merchant, 2012; Guo et al., 2010, 2015; Kong & Price, 2019; Nouet et al., 2011). Thus, it is possible that, in the absence of specific mechanisms of copper import, the limitation in copper available in the lumen to be incorporated into Pc restrains the amount of a functional holoprotein to the levels described in T. oceanica and found in this work (Ho et al., 2003, Levy et al., 2008, Twining & Baines, 2013). However, our results indicate that even the relatively low concentrations of holo-Pc detected are enough to promote an increased growth (up to 60% in the case of the #45 and #79 mutants) under irondeficient conditions as compared with the WT strain (Figure 2). This increased growth can be measured as higher cell density, content in pigments (chlorophylls and carotenoids), global photosynthetic rates per cell and even cell volume (Table 1). Moreover, although the apparent partially recovered levels of psaB-more evident in mutant #79does not allow to directly inferring an equivalent increase of active PSI, a slightly higher PSI activity has been also observed in the mutants in PAM experiments. Previous studies have proposed that, in P. tricornutum, the iron-rich PSI is degraded in iron-deficient conditions, mobilizing this element for other metabolic activities as respiration (Allen et al., 2008). The presence of Pc should not modify the extent of PSI degradation, as it does not alter the cell metabolic demand for iron. Thus, the observed data suggest that the balance of PSI synthesis/degradation under iron-limiting conditions is perhaps affected not only by iron requirements, but also by limitations at the PSI donor side or the global redox stress status of the chloroplast. Finally, the Pc-expressing strains also show changes in cell morphology and chloroplasts with a thicker shape, which indeed reflects changes in the physiological and metabolic status of cells (Levy et al., 2008; Martin-Jézéquel & Tesson, 2013).

In addition, PSII activity of P. tricornutum cells was affected by low iron concentration in a different manner in the WT and mutant strains. The decreased PSII activity in iron-deficient conditions detected by TL experiments (Figure 3) may be attributed to a reduced amount of functional PSII complexes in cells (Msilini et al., 2011). The partial blocking of ET between PSII and PSI in cells cultured under iron-deficient conditions can generate the over-reduction of the PQ pool. Under iron-deficient conditions, P₇₀₀ cannot become reduced, possibly due to fewer available iron-containing ET proteins downstream of the PQ pool, such as the Cb₆f complex or the Cc₆ soluble donor (Allen et al., 2008; Bruce & Malkin, 1991; Greene et al., 1992; Roncel et al., 2016). In particular, in P. tricornutum cells grown under low iron conditions, the Cc_6 concentration is reduced to $\approx 10-12\%$ of the protein present in iron-replete cells, thus presumably disfavoring the PQ pool re-oxidation (Roncel et al., 2016). An over-reduction of the PQ pool may induce the appearance of the acceptor side photoinhibition process, which can decrease PSII activity. The existence of an alternative electron carrier in the mutant strains, the E85K Pc, could improve the ET to PSI, and partially eliminate the overreduction of the PO pool and the concomitant lack of activity of PSII. This hypothesis can explain the significant lower decrease of the TL intensity signal in the WT strain (Figure 3), as well as the higher maximum quantum yield of PSII obtained by fluorescence measurements in the mutants compared to the WT (Table 2). Higher values for rETR_{max} were also determined in the three mutants with respect to the WT (Figure 5, Table 2), which indicates that the aforementioned occurrence of the acceptor side photoinhibition process is delayed due to the presence of Pc.

In chloroplasts, the reducing site of PSI and the oxidizing side of PSII are the major generation sites of reactive oxygen species (Sgherri et al., 2018). However, recently it has been reported that superoxide can also be produced by Cb₆f and at the acceptor side of PSII (Roach & Krieger-Liszkay, 2014). On the other hand, the major source of singlet oxygen appears to be PSII (Krieger-Liszkay, 2005; Murata et al., 2007), the generation of this reactive oxygen species is enhanced as a consequence of the photoinhibition of the acceptor side in PSII. The formation of singlet oxygen can initiate the peroxidation of unsaturated lipids in membranes (Vavilin & Ducruet, 1998). The level of lipid peroxidation in photosynthetic membranes can be measured by the HTL technique (Roncel et al., 2007). Several luminescence high-temperature bands (HTL bands) have been observed without prior illumination at temperatures above 60°C (Ducruet & Vavilin, 1999; Roncel et al., 2007). A broad HTL band centered near 130°C (known as the HTL2 band) is generated because of the thermal



radiative decomposition of lipid peroxides that, in turn, leads to the formation of carbonyl groups in a triplet state followed by migration of excitation energy toward chlorophylls (Ducruet & Vavilin, 1999; Vavilin et al., 2002; Vavilin & Ducruet, 1998). The amplitude of this band has been well correlated with the accumulation of malondialdehyde, an indicator of lipid peroxidation in standard chemical tests (Vavilin et al., 2002; Vavilin & Ducruet, 1998).

Analysis of HTL2 bands of TL obtained in *P. tricornutum* cells (Figure 4) clearly showed a much higher level of lipid peroxidation in iron-deficient conditions in the WT strain. However, in the three mutant strains the level of peroxidation did not significantly increase under iron-restricted conditions. These results suggest a higher rate of generation of reactive singlet oxygen under iron-deficient conditions in the WT strain. As previously said, the low levels of Cc₆ probably induce the over-reduction of the PQ pool and the concomitant acceptor side photoinhibition in PSII, and thus singlet oxygen production. The existence of Pc in the mutant strains under iron-deficient conditions could partially eliminate the over-reduction of the PQ pool and the concomitant generation of singlet oxygen, thus decreasing oxidative stress and lipid peroxidation.

On the other hand, PSI activity is more sensitive to iron-limitation than PSII activity (Pushnik & Miller, 1989; Roncel et al., 2016). The significant decrease in the quantum yield of PSI, Y(I), observed in iron-deficient WT cells (Figure 6) can be ascribed to a deficiency of PSI donors, which causes that P₇₀₀ cannot become reduced (Allen et al., 2008, Bruce & Malkin, 1991, Greene et al., 1992, Roncel et al., 2016). An increase of Y(I) and a lower degree of limitation on the donor side of PSI, Y(ND), have been here observed in the three mutants (Figure 6A-B). The existence of Pc as an additional electron carrier in the mutant strains seems to improve ET to PSI and, consequently, to partially increase Y(I) as well as to decrease the over-reduction of the PQ pool and the concomitant donor-side limitation of PSI.

In summary, our results show that the presence of Pc as an additional electron carrier in *P. tricornutum* under iron-limiting conditions seems to decrease the over-reduction of the PQ pool. This consequently promotes an improvement in the maximum quantum yield of both PSII and PSI, together with a decrease in the acceptor side photoinhibition of PSII, oxidative stress and the peroxidation of lipids, and a lower degree of limitation on the donor side of PSI. As a result, the mutants expressing Pc in *P. tricornutum* exhibit an enhancement in cell growth under iron-limiting conditions compared to WT cells.

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AUTHOR CONTRIBUTIONS

José A. Navarro and Manuel Hervás conceived and designed the project. Carmen Castell and Pilar Bernal-Bayard carried out the molecular biology genetic work as well as cells transformation and selection. Carmen Castell, with the help of José A. Navarro and Manuel Hervás, carried out the physiological and biochemical characterization of the different strains here studied. Mercedes Roncel and Carmen Castell carried out PAM experiments and José M. Ortega and Carmen Castell the TL measurements. All the authors discussed the results and José A. Navarro wrote the manuscript with their collaboration. The manuscript was corrected, revised and approved by all authors.

DATA AVAILABILITY STATEMENT

The data that support the findings of this study are available from the corresponding author upon reasonable request

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SUPPORTING INFORMATION

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