

Characterization of cooperative bicarbonate uptake into chloroplast stroma in the green alga *Chlamydomonas reinhardtii*

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The supply of inorganic carbon (Ci; CO₂ and HCO₃⁻) is an environmental rate-limiting factor in aquatic photosynthetic organisms. To overcome the difficulty in acquiring Ci in limiting-CO₂ conditions, an active Ci uptake system called the CO₂-concentrating mechanism (CCM) is induced to increase CO₂ concentrations in the chloroplast stroma. An ATP-binding cassette transporter, HLA3, and a formate/nitrite transporter homolog, LCIA, are reported to be associated with HCO₃⁻ uptake [Wang and Spalding (2014) *Plant Physiol* 166(4): 2040–2050]. However, direct evidence of the route of HCO₃⁻ uptake from the outside of cells to the chloroplast stroma remains elusive owing to a lack of information on HLA3 localization and comparative analyses of the contribution of HLA3 and LCIA to the CCM. In this study, we revealed that HLA3 and LCIA are localized to the plasma membrane and chloroplast envelope, respectively. Insertion mutants of *HLA3* and/or *LCIA* showed decreased Ci affinities/accumulation, especially in alkaline conditions where HCO₃⁻ is the predominant form of Ci. HLA3 and LCIA formed protein complexes independently, and the absence of LCIA decreased *HLA3* mRNA accumulation, suggesting the presence of unidentified retrograde signals from the chloroplast to the nucleus to maintain *HLA3* mRNA expression. Furthermore, although single overexpression of HLA3 or LCIA in high CO₂ conditions did not affect Ci affinity, simultaneous overexpression of HLA3 with LCIA significantly increased Ci affinity/accumulation. These results highlight the HLA3/LCIA-driven cooperative uptake of HCO₃⁻ and a key role of LCIA in the maintenance of HLA3 stability as well as Ci affinity/accumulation in the CCM.

bicarbonate uptake | *Chlamydomonas* | chloroplast envelope | CO₂-concentrating mechanism | photosynthesis

Inorganic carbon (Ci; CO₂ and HCO₃⁻) transport is essential for a wide range of biological processes such as CO₂ metabolism, cellular pH homeostasis, and photosynthesis. Because HCO₃⁻ is not freely permeable to biological membranes, it must be transported across membranes by HCO₃⁻ transporters or channels. HCO₃⁻ transporters have been studied extensively in mammals and been found to cluster into solute carrier (SLC) 4 and SLC 26 families (1). In cyanobacteria, five types of Ci transporters have been identified (2), including three HCO₃⁻ transporters and two NAD(P)H dehydrogenase-dependent CO₂ uptake systems. In land plants, aquaporin-mediated CO₂ permeation has been suggested to play physiological roles in photosynthesis (3), and in a marine diatom, SLC4 family protein localized to the plasma membrane (PM) facilitates HCO₃⁻ uptake (4). However, no studies have validated the entire route of HCO₃⁻ transport from the outside of cells to the chloroplast stroma through the PM and chloroplast envelope (CE) in photosynthetic organisms.

Aquatic conditions are not well suited for efficient photosynthesis because the CO₂ diffusion rate is ~10,000-fold lower compared with that in atmospheric conditions (5). Therefore, aquatic photosynthetic organisms, including microalgae, are frequently exposed to limiting CO₂ stress. To acclimate to this stress, most microalgae possess a CO₂-concentrating mechanism (CCM) to accumulate CO₂ around the CO₂ fixation enzyme ribulose 1,

5-bisphosphate carboxylase/oxygenase (Rubisco) and to maintain adequate photosynthetic efficiency (6, 7).

The green alga *Chlamydomonas reinhardtii* has been used as a model organism for molecular and physiological studies of the CCM since it was first identified (8). A model of the CCM has been proposed based on the subcellular structure of *C. reinhardtii* (9, 10). Environmental Ci is transported to the chloroplast stroma by Ci transporters localized to the PM and CE. Carbonic anhydrase (CA) localized to the chloroplast stroma is predicted to contribute to the maintenance of the Ci pool, in the form of HCO₃⁻, by rapid conversion of CO₂ to HCO₃⁻, thereby preventing the loss of CO₂ by diffusion (11). It is known that tubule-like thylakoid membranes penetrate into the pyrenoid (12), a Rubisco-enriched structure in the chloroplast. HCO₃⁻ in the stroma is transported into the acidic thylakoid lumen by a putative channel or transporter localized to the thylakoid membrane, and HCO₃⁻ is rapidly converted to CO₂ by a constitutively expressed CA (13, 14). Then, CO₂ diffuses from the thylakoid lumen into the pyrenoid matrix and is fixed by Rubisco. It was also reported that *C. reinhardtii* acclimates to two distinct limiting CO₂ conditions, termed low CO₂ (LC; ~0.03–0.5% CO₂ or 7–70 μM CO₂) and very low CO₂ (VLC; <0.02% CO₂ or <7 μM CO₂) (15, 16), and different types of Ci uptake systems could function in the CCM in these separate conditions (16).

To identify CCM-associated components, several transcriptome analyses have been performed (17–22), and several genes encoding membrane proteins were focused on as candidate Ci transporter genes, including *LCII* (low CO₂ inducible gene 1) (23),

Significance

The entry of inorganic carbon (Ci; CO₂ and HCO₃⁻) into cells involves many biological processes in both animals and plants, and aquaporins as well as bicarbonate transporters play roles in Ci transport. Although transporting external HCO₃⁻ into the stroma through the chloroplast envelope is one of the rate-limiting factors for aquatic photosynthetic organisms, specific molecular components in this process have not yet been identified experimentally. Molecular identification of proteins essential for Ci uptake located in the chloroplast envelope and in the plasma membrane documented in this study helps in understanding how aquatic photosynthetic organisms developed machinery to acclimate to CO₂-limiting environment and to maintain adequate levels of photosynthesis for survival or growth.

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LCIA (low CO₂ inducible gene A) (19), and *HLA3* (high light activated 3) (24).

LCI1 is localized to the PM (25), and its expression is regulated by the MYB-transcription factor *LCR1* (low CO₂ stress response 1) (26). When *LCI1* was artificially expressed in HC conditions, the cells showed increases in the internal Ci pool, suggesting that *LCI1* is directly or indirectly associated with Ci uptake (25). *LCIA* (also known as *NAR1.2*) is a homolog of the nitrate transporter *NAR1* and belongs to the formate/nitrite transporter family (27). Although the expression of other *NAR1* family genes of *C. reinhardtii* is mainly regulated by nitrogen source, *LCIA* is specifically induced in LC conditions and is not under the control of nitrogen source (19). *LCIA* was predicted to localize to the CE (19), and this prediction was supported by indirect immunofluorescence assay evidence (16). Functional expression analysis using *Xenopus* oocytes showed transport activity of *LCIA* for both HCO₃⁻ and NO₂⁻ (27), and *LCIA* appears to be associated with HCO₃⁻ uptake in VLC conditions from analysis of an insertion mutant (16). *HLA3* is an ATP-binding cassette (ABC) transporter of the multidrug resistance-related protein subfamily, and its transcription is induced by high light and LC conditions (19, 24). Although *HLA3* is predicted to localize to the PM (10), no experimental data are available at present. Knockdown (KD) of *HLA3* mRNA expression resulted in modest decreases in photosynthesis affinity, but simultaneous KD of *LCIA* and *HLA3* mRNAs caused a dramatic decrease in growth rate, Ci uptake activity, and photosynthetic Ci affinity, especially in alkaline conditions, where HCO₃⁻ is the predominant form of Ci (28).

In this study, by use of indirect immunofluorescence assays and membrane fractionation, the subcellular localization of *HLA3* was elucidated. In addition, by analyses of the photosynthetic characteristics of *HLA3* and *LCIA* single insertion mutants, an *HLA3/LCIA* double insertion mutant, and overexpressing strains of *HLA3* and/or *LCIA*, we concluded that *HLA3* and *LCIA* are cooperatively associated with HCO₃⁻ uptake across the PM and CE, respectively.

Results

Accumulation of *HLA3* and *LCIA* in Very Low CO₂ Conditions. First, to define the acclimated states of limiting CO₂ conditions (LC or VLC) of cells grown in liquid culture, total Ci concentration in the culture medium at pH 7.0 was measured, and consequent CO₂ concentrations were calculated (Fig. 1A). CO₂ concentrations supplied with 0.04% CO₂ for 1, 2, 4, 6, and 12 h were estimated as 6.3, 3.1, 2.9, 1.9, and 1.8 μM, respectively, which correspond to the range for VLC (<7 μM CO₂) (16). Thus, we defined the limiting CO₂ conditions of liquid culture as VLC throughout this study. Next, the time course accumulation of *HLA3* and *LCIA* after VLC induction was examined (Fig. 1A). The accumulation of these proteins started within 1 h and reached their maximum levels within 4 h, as was the case for *LCI1* used as a control of VLC induction. The molecular masses of *HLA3* and *LCIA* were detected at sizes of ~133 and 27 kDa, respectively (Fig. S1A and *SI Results and Discussion*).

Subcellular Localization of *HLA3* and *LCIA*. To analyze the subcellular localization of *HLA3*, an indirect immunofluorescence assay was performed (Fig. 1B). Fluorescence signals from an anti-*HLA3* antibody were detected peripherally, suggesting the localization of *HLA3* to the PM. Fluorescence signals from an anti-*LCIA* antibody were detected as a single cup-shaped structure (Fig. 1B), as in the previous study (16). To further clarify the localization of *HLA3* and *LCIA* biochemically, protein samples from total cell, PM, and CE fractions were probed with antibodies against *HLA3*, *LCI1*, H⁺-ATPase, *LCIA*, and *CCP1* (Fig. 1C and *SI Results and Discussion*). *LCI1* and H⁺-ATPase were enriched in the PM fraction, consistent with the PM localization of these

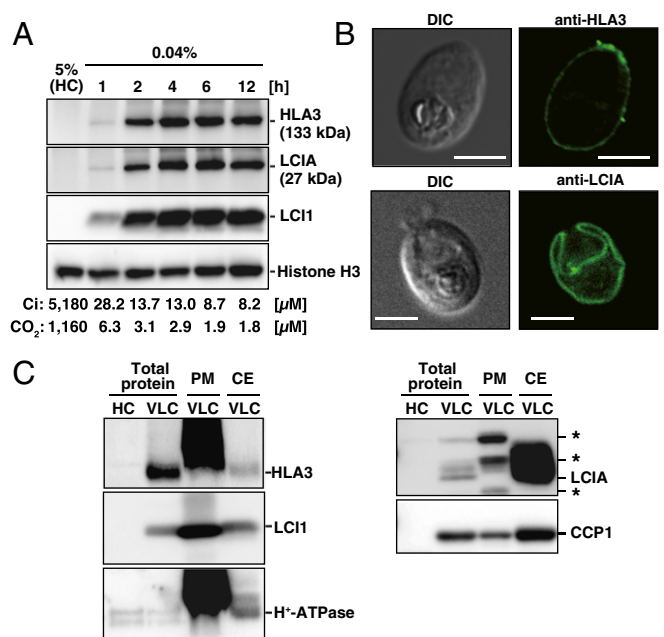


Fig. 1. Accumulation and subcellular localization of *HLA3* and *LCIA*. (A) Time-course of accumulation of *HLA3*, *LCIA*, and *LCI1* proteins in WT cells. For induction of limiting-CO₂ conditions, cells supplied with 5% CO₂ (high CO₂; HC) were centrifuged, suspended in new fresh medium, and cultured with 0.04% CO₂ for 1, 2, 4, 6, and 12 h. Histone H3 was used as a loading control. The total Ci concentrations and calculated CO₂ concentrations after each induction time are also indicated below the figures. Using an HCO₃⁻/CO₂ ratio of 4.47 at pH 7.0, CO₂ concentrations were calculated using the equation (pH = pK_a + log₁₀ [HCO₃⁻]/[CO₂]), where pK_a was an acid dissociation constant of 6.35. (B) Subcellular localization of *HLA3* and *LCIA* by an indirect immunofluorescence assay. WT cells were grown in very low CO₂ (VLC) for 12 h. DIC, differential interference contrast. (Scale bars, 5 μm.) (C) Immunoblot analysis in isolated plasma membrane (PM) and chloroplast envelope (CE) fractions with antibodies against *HLA3*, *LCI1*, H⁺-ATPase, *LCIA*, and *CCP1*. Asterisks indicate nonspecific bands.

proteins (25, 29). Similarly, a notable enrichment of *HLA3* was observed in the PM fraction. *LCIA* was highly enriched in the CE fraction, where CE protein *CCP1* (30) was also enriched. From these results, we concluded that *HLA3* and *LCIA* were localized to the PM and CE, respectively.

Isolation of an *HLA3* Insertion Mutant and Photosynthetic Characteristics. To evaluate the degree of contribution of *HLA3* to the CCM, we isolated an *HLA3* insertion mutant from our paramycin resistance gene-tagged mutant library by PCR-based screening, as described previously (31), and designated the strain *Hin-1* (Fig. S2A–C and *SI Results and Discussion*).

Next, the photosynthetic characteristics were evaluated by measuring the rates of Ci-dependent O₂ evolution of WT, *Hin-1*, and the complemented strain *Hin-1C* grown in VLC at different pH. K_{0.5} (Ci) values, the Ci concentration required for half maximal O₂-evolving activity, of WT and *Hin-1* were similar at pH 6.2 (ratio of HCO₃⁻:CO₂ = 0.7:1) and pH 7.8 (HCO₃⁻:CO₂ = 28:1), indicating that the difference in photosynthetic Ci affinity between WT and *Hin-1* was not significant (Fig. 2A). Because *HLA3* KD strains showed retarded growth rates at pH 9.0 (28) where the ratio of HCO₃⁻:CO₂ = 446:1 and HCO₃⁻ was the predominant form of Ci, we evaluated the changes in Ci affinity during acclimation to VLC at pH 9.0 in a time course analysis (Fig. 2B). Both WT and *Hin-1* showed a gradual decrease in K_{0.5} (Ci) during acclimation to VLC. However, although WT in VLC at 6 h showed almost the same Ci affinity compared with that at 12 h (241 ± 87 μM at 6 h and 290 ± 50 μM at 12 h), *Hin-1* still showed much

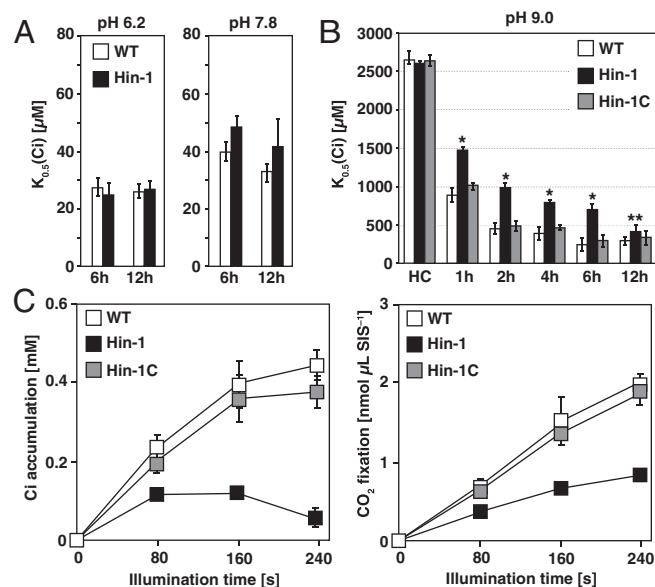


Fig. 2. Characterization of an *HLA3* insertion mutant. (A) Inorganic carbon (Ci) affinity of WT and *HLA3* insertion mutant (Hin-1) grown in very low CO₂ (VLC) for 6 or 12 h. Photosynthetic O₂-evolving activity was measured with different external Ci concentrations at pH 6.2 or 7.8, and the respective K_{0.5} (Ci) values, the Ci concentration required for half maximum O₂-evolving activity, were calculated. (B) Ci affinity of WT, Hin-1, and complemented Hin-1 (Hin-1C) grown in high CO₂ (HC) or VLC for 1, 2, 4, 6, and 12 h. O₂-evolving activity was measured at pH 9.0. **P* < 0.01 and ***P* < 0.05 by Student *t* test. (C) Accumulation and fixation of Ci in WT, Hin-1, and Hin-1C. Cells were grown in VLC for 6 h, and intracellular Ci accumulation (Left) and CO₂ fixation (Right) at pH 9.0 were measured using a silicone oil layer method. SIS, sorbitol impermeable space.

lower Ci affinity especially at 6 h ($691 \pm 143 \mu\text{M}$ at 6 h and $405 \pm 57 \mu\text{M}$ at 12 h), and the decreased Ci affinity was restored in Hin-1C ($296 \pm 78 \mu\text{M}$ at 6 h and $333 \pm 89 \mu\text{M}$ at 12 h). These results suggested that other Ci uptake systems could compensate for the absence of *HLA3* and contribute to the increase in Ci affinity at 12 h and that measuring photosynthetic characteristics at 6 h was appropriate for evaluating the contribution of *HLA3* to the CCM.

To evaluate the contribution of *HLA3* to actual Ci uptake activity, the accumulation and fixation of [¹⁴C]-labeled Ci in WT, Hin-1, and Hin-1C grown in VLC for 6 h were measured (Fig. 2C). Hin-1 showed significantly lower levels of Ci accumulation of 0.12 mM (0.57-fold of Hin-1C), 0.12 mM (0.32-fold), and 0.06 mM (0.17-fold) after 80, 160, and 240 s of illumination, respectively, and CO₂ fixation of 0.37 nmol·µL SIS⁻¹ (0.59-fold), 0.67 nmol·µL SIS⁻¹ (0.49-fold), and 0.83 nmol·µL SIS⁻¹ (0.44-fold), respectively, compared with that of Hin-1C. These results indicated that *HLA3* was a meaningful role in HCO₃⁻ uptake in VLC conditions.

Isolation of *LCIA* Insertion Mutants and Photosynthetic Characteristics.

A disruption mutant of *LCIA* has been characterized, and the contribution of *LCIA* to the CCM has been reported (16). To compare the degrees of the contributions of *HLA3* and *LCIA* to the CCM, we also isolated two *LCIA* insertion mutants (Fig. S2 D–G and SI Results and Discussion), designated as Ain (Ain-1 and Ain-2), and compared the photosynthetic characteristics with Hin-1. Interestingly, accumulation of *HLA3* was much lower in Ain compared with that in WT, and this decreased accumulation of *HLA3* was restored in the complemented strains Ain-1C and Ain-2C (Fig. 3A). This result was in sharp contrast to that of *LCI1* and *LCIB* (32), which were not affected by the impairment of the *LCIA* (Fig. 3A).

Next, the photosynthetic characteristics of Ain-1, Ain-2, Ain-1C, and Ain-2C were evaluated. As in the case of Hin-1, the K_{0.5} (Ci) of Ain-1 and Ain-2 was similar to WT at pH 6.2 (Fig. 3B). However, in contrast to Hin-1, the K_{0.5} (Ci) of Ain-1 ($57 \pm 2 \mu\text{M}$ at 6 h and $56 \pm 3 \mu\text{M}$ at 12 h) and Ain-2 ($57 \pm 1 \mu\text{M}$ at 6 h and $57 \pm 2 \mu\text{M}$ at 12 h) was significantly higher than that of WT ($40 \pm 3 \mu\text{M}$ at 6 h and $33 \pm 3 \mu\text{M}$ at 12 h), Ain-1C ($38 \pm 2 \mu\text{M}$ at 6 h and

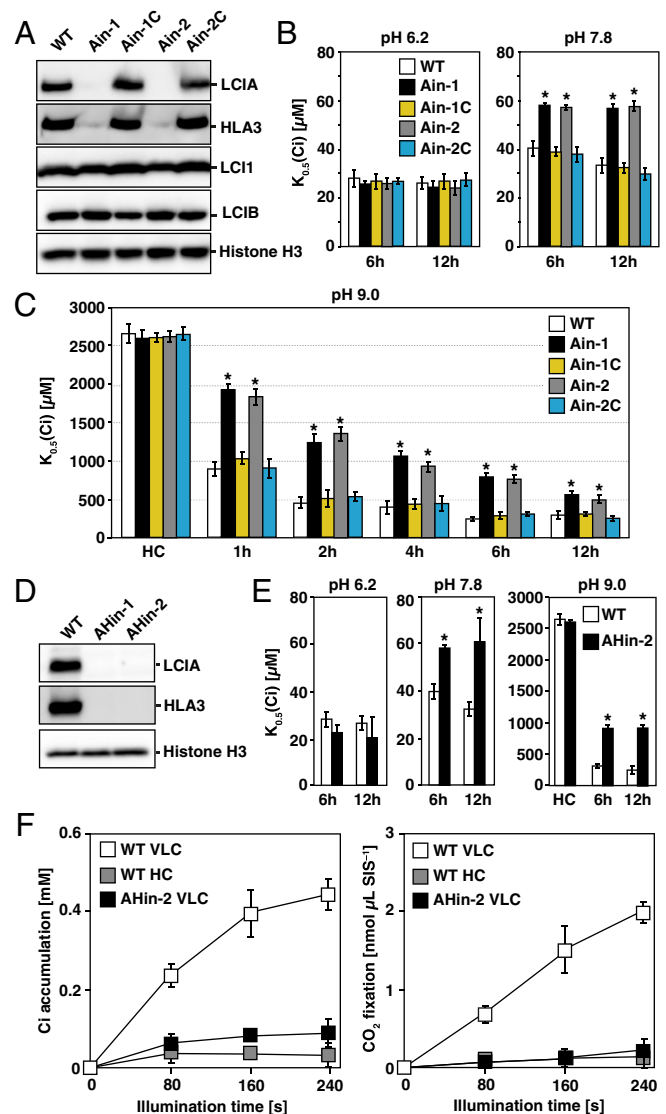


Fig. 3. Characterization of *LCIA* insertion mutants and an *LCIA/HLA3* double-insertion mutant. (A) Accumulation of *LCIA*, *HLA3*, *LCI1*, and *LCIB* in WT, *LCIA* insertion mutants (Ain-1 and Ain-2), and their complemented strains (Ain-1C and Ain-2C). Cells were grown in very low CO₂ (VLC) for 12 h. (B) Inorganic carbon (Ci) affinity of WT, Ain-1, Ain-2, Ain-1C, and Ain-2C grown in very low CO₂ (VLC) for 6 or 12 h. Photosynthetic O₂-evolving activity was measured with different external Ci concentrations at pH 6.2 or 7.8, and the respective K_{0.5} (Ci) values, the Ci concentration required for half maximum O₂-evolving activity, were calculated. **P* < 0.01. (C) Ci affinity of WT, Ain-1, Ain-2, Ain-1C, and Ain-2C grown in high CO₂ (HC) or VLC for 1, 2, 4, 6, and 12 h. O₂-evolving activity was measured at pH 9.0. **P* < 0.01. (D) Accumulation of *HLA3* and *LCIA* in WT and *LCIA/HLA3* double-insertion mutants (AHin-1 and AHin-2) grown in VLC for 12 h. (E) Ci affinity of WT and AHin-2 grown in HC or VLC for 6 or 12 h. O₂-evolving activity was measured at pH 6.2, 7.8, or 9.0. **P* < 0.01. (F) Accumulation and fixation of Ci in WT and AHin-2. Cells were grown in HC or VLC for 6 h, and intracellular Ci accumulation (Left) and CO₂ fixation (Right) were measured at pH 9.0. SIS, sorbitol impermeable space.

$32 \pm 2 \mu\text{M}$ at 12 h), and Ain-2C ($37 \pm 3 \mu\text{M}$ at 6 h and $29 \pm 2 \mu\text{M}$ at 12 h), even at pH 7.8 (Fig. 3B). At pH 9.0, although Ain also showed gradual decreases in $K_{0.5}$ (Ci) during acclimation to VLC, these cells always showed lower Ci affinity than Hin-1 (Fig. 2B), as well as WT and complemented strains (Fig. 3C). These results suggested a significant contribution of LCIA to increases in Ci affinity and to maintaining HLA3 stability in the CCM.

Isolation of LCIA/HLA3 Double-Insertion Mutants and Photosynthetic Characteristics. Because Ci affinity in VLC at 12 h was higher than that at 6 h in both *HLA3* and *LCIA* single mutants, either protein could partially complement each other to increase Ci affinity. Thus, we expected that *LCIA/HLA3* double-insertion mutants would show an additive decrease in Ci affinity compared with the single-insertion mutants. Thus, we isolated double-insertion mutants by crossing one of the Ain-2 progeny with Hin-1 and designated these as AHin (AHin-1 and AHin-2; Fig. 3D, Fig. S2 H–K, and SI Results and Discussion).

Next, the photosynthetic characteristics of AHin-2 were evaluated (Fig. 3E). As in the case of Hin-1 and Ain, the $K_{0.5}$ (Ci) of AHin-2 was similar to WT at pH 6.2. At pH 7.8, the $K_{0.5}$ (Ci) of AHin-2 ($58 \pm 2 \mu\text{M}$ at 6 h and $61 \pm 10 \mu\text{M}$ at 12 h) was significantly higher than that of WT, but it was similar to Ain. At pH 9.0, AHin-2 showed lower Ci affinity than both Hin-1 and Ain, and Ci affinity was not increased even at 12 h ($898 \pm 78 \mu\text{M}$ at 6 h and $901 \pm 94 \mu\text{M}$ at 12 h). Ci accumulation and fixation in AHin-2 grown in VLC at 6 h was also measured (Fig. 3F). After 80, 160, and 240 s of illumination, AHin-2 showed substantially decreased Ci accumulation of 0.05 mM (0.21-fold of WT and 0.41-fold of Hin-1), 0.06 mM (0.16-fold and 0.53-fold), and 0.06 mM (0.15-fold and 1.0-fold), respectively, and CO_2 fixation of 0.07 $\text{nmol}\cdot\mu\text{L}^{-1}\text{SIS}^{-1}$ (0.1-fold and 0.2-fold), 0.12 $\text{nmol}\cdot\mu\text{L}^{-1}\text{SIS}^{-1}$ (0.08-fold and 0.17-fold), and 0.14 $\text{nmol}\cdot\mu\text{L}^{-1}\text{SIS}^{-1}$ (0.07-fold and 0.17-fold), respectively, compared with that of WT and Hin-1.

Finally, the effect of absence of LCIA and/or HLA3 on cell growth was examined. Growth rates were measured in VLC at pH 8.4 (Fig. S2L) because there were no significant differences at pH 7.8, and none of the cell lines could grow at pH 9.0. The doubling time of WT was 7.2 h and that of Hin-1, Ain-1, Ain-2, and AHin-2 increased significantly to 7.6, 9.5, 9.3, and 12.7 h, respectively, reflecting the degree of decreased Ci affinity of each cell line. These results highlighted an additive decrease in Ci affinity/accumulation/growth rates of the double-insertion mutant compared with the *HLA3* or *LCIA* single-insertion mutants.

Isolation of LCIA and/or HLA3 Overexpressing Strains and Photosynthetic Characteristics. To demonstrate the physiological function of LCIA and HLA3 more directly, the photosynthetic characteristics of cells overexpressing LCIA and/or HLA3 were examined in HC conditions where other VLC-inducible proteins were not induced. For overexpression, two chimeric plasmids, pTY2b-LCIA and pTY2b-HLA3, were constructed (Fig. S3A). These plasmids allowed the induction of *LCIA* and *HLA3* transcripts by switching the nitrogen source from NH_4^+ to NO_3^- irrespective of the CO_2 conditions. In this study, we cultured the cells with four combinations of nitrogen sources in the medium and CO_2 concentrations, designated as HC- NH_4^+ , HC- NO_3^- , VLC- NH_4^+ , and VLC- NO_3^- .

First, we transformed WT cells with pTY2b-LCIA or pTY2b-HLA3 separately. The transformants showed accumulation of LCIA or HLA3 when grown in HC- NO_3^- conditions and were designated as Aox (Aox-1 and Aox-2) and Hox (Hox-1 and Hox-2), respectively (Fig. S3B and C and SI Results and Discussion). Next, by introducing pTY2b-HLA3 into Aox-1, we generated two independent transformants expressing LCIA and HLA3 simultaneously and designated these as AHox (AHox-1 and AHox-2; Fig. S3D). Accumulation of HLA3 in AHox-1 and AHox-2 was the same as that of VLC-grown WT. To isolate a strain overexpressing

both LCIA and HLA3 with greater abundance, the progeny of Aox-1 was crossed with Hox-1 and a strain designated as AHox-3 was obtained (Fig. S3E).

Next, to evaluate the effect of LCIA and/or HLA3 overexpression on the photosynthetic characteristics, rates of O_2 evolution at pH 6.2, 7.8, and 9.0 and Ci accumulation at pH 9.0 of these strains were measured. In Aox, there were no differences in Ci affinity at pH 7.8 and pH 9.0, as well as Ci accumulation compared with WT (Fig. 4A and Table S1–S3). In contrast, HC- NO_3^- -grown Hox showed a small but significant increase of Ci accumulation of 0.08 mM (2.5-fold of WT at 80 s), 0.07 mM (1.5-fold at 160 s), and 0.13 mM (2.4-fold at 240 s) in Hox-2, compared with that of HC- NO_3^- -grown WT, but the phenotype led to a slight increase in Ci affinity only at pH 9.0 in Hox-2 (Fig. 4B), suggesting that Ci in the cytosol transported by HLA3 could not efficiently enter the chloroplast stroma in the absence of LCIA. On the other hand, Ci affinity at pH 6.2 was increased in

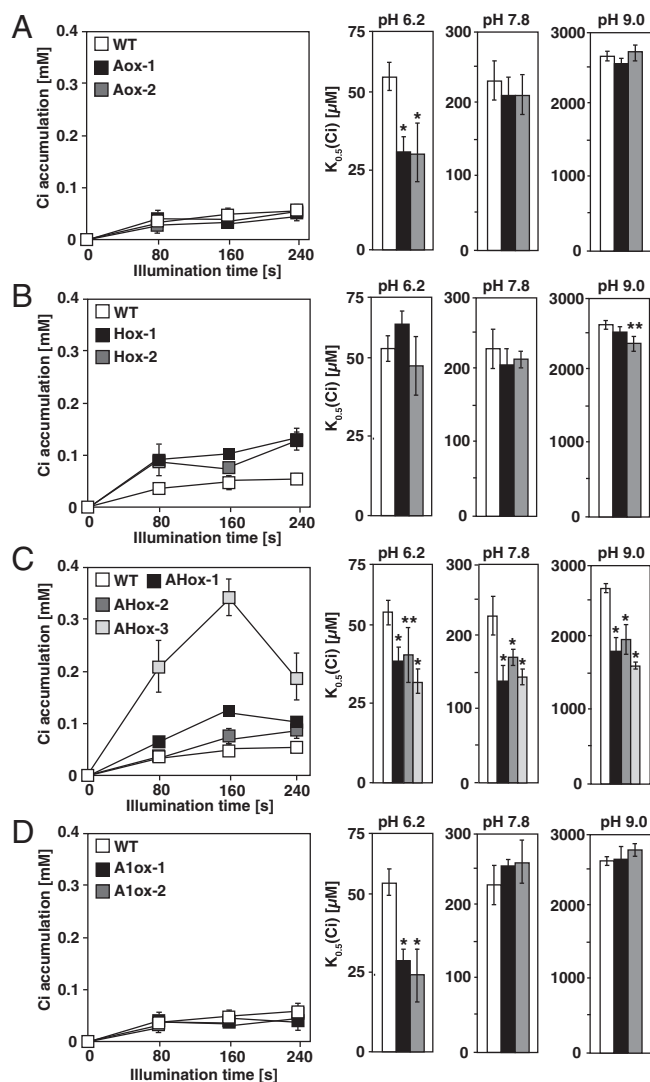


Fig. 4. Characterization of LCIA- and HLA3-overexpressing strains. Accumulation of inorganic carbon (Ci) (Left) and Ci affinity (Right) in WT and in strains overexpressing LCIA (A), HLA3 (B), LCIA/HLA3 (C), and LCIA/LCI1 (D). Cells were grown in high $\text{CO}_2\text{-NO}_3^-$ for 12 h, and Ci accumulation was measured at pH 9.0. For Ci affinity, O_2 -evolving activity was measured with different external Ci concentrations at pH 6.2, 7.8, or 9.0 and the respective $K_{0.5}$ (Ci) values, the Ci concentration required for half maximum O_2 -evolving activity, were calculated. * $P < 0.01$ and ** $P < 0.05$.

LCIA-overexpressing Aox (Fig. 4A) and AHox (Fig. 4C), but not in Hox (Fig. 4B).

In contrast to Aox and Hox, AHox showed a significant increase in Ci affinity and Ci accumulation compared with WT at alkaline conditions (Fig. 4C and Tables S2 and S3). In particular, HC-NO₃⁻-grown AHox-3 showed substantially increased Ci accumulation of 0.21 mM (6.3-fold of WT at 80 s), 0.34 mM (6.8-fold at 160 s), and 0.19 mM (3.6-fold at 240 s) compared with that of HC-NO₃⁻-grown WT. Consequently, the respective K_{0.5} (Ci) of AHox-1, AHox-2, and AHox-3 decreased to 141 ± 20 (0.61-fold of WT), 174 ± 20 (0.76-fold), and 147 ± 19 μM (0.64-fold) at pH 7.8 and to 1,821 ± 201 (0.68-fold of WT), 1,980 ± 198 (0.75-fold), and 1,626 ± 49 μM (0.61-fold) at pH 9.0. In HC-NH₄⁺ conditions at pH 7.8 where LCIA and HLA3 were not induced, the respective K_{0.5} (Ci) of 257 ± 28, 250 ± 30, and 262 ± 29 μM in AHox-1, AHox-2, and AHox-3 was not significantly different from that of 273 ± 31 μM in WT (Table S2). These results indicated that NO₃⁻-induced overexpression of LCIA and HLA3 could enhance HCO₃⁻ accumulation in the chloroplast stroma and increase Ci affinity.

Although PM-localized LCI1 could be associated with Ci uptake (25), the preferred Ci species of LCI1 remained elusive. To evaluate the degree of LCIA/HLA3-driven HCO₃⁻ uptake activity, we also isolated six transformants expressing LCIA with LCI1 by introducing pTY2b-LCI1 (Fig. S3A) into Aox-1 and designated two representatives as A1ox (A1ox-1 and A1ox-2; Fig. S3F). There were no differences in Ci accumulation and affinity in alkaline conditions compared with WT (Fig. 4D and Table S2 and S3), suggesting that LCI1 was not related to direct HCO₃⁻ uptake along with LCIA.

A Defect in LCIA Led to a Decrease in HLA3 Accumulation Caused by Suppression of HLA3 mRNA Accumulation. As described above, accumulation of HLA3 was much lower in Ain compared with that in WT (Fig. 3A). This result suggested two possibilities. First, HLA3 and LCIA undergo a physical interaction where the PM is associated with the CE and the absence of LCIA causes instability of HLA3. Second, the absence of LCIA causes the repression of HLA3 mRNA accumulation.

To examine the former possibility, the molecular masses of LCIA and HLA3 *in vivo* were estimated by Blue Native-PAGE. We expected that LCIA and HLA3 should be detected with the same molecular mass in nondenaturing conditions if these two proteins interact and form a complex. However, using 1.0% n-dodecyl β-D-maltoside (DDM) as a detergent, LCIA and HLA3 were detected with different sizes of ~240 and 580 kDa, respectively (Fig. 5A). We also estimated the molecular masses using different DDM concentrations (0.25%, 0.5%, 1.0%, or 2.0%) or using formaldehyde cross-linker, and LCIA and HLA3 were still detected at 240 and 580 kDa, respectively (Fig. S4A and B). Furthermore, LCIA and HLA3 could form respective complexes with the same molecular masses even in Aox, Hox, and AHox cells grown in HC-NO₃⁻ conditions (Fig. S4C). These results strongly suggested that LCIA and HLA3 did not interact physically *in vivo* and at least VLC-inducible proteins other than LCIA and HLA3 were not associated with the formation of the respective protein complexes.

For the latter possibility, HLA3 mRNA accumulation was evaluated by quantitative real-time PCR (Fig. 5B). The sequences of primers used are listed in Table S4. HLA3 mRNA levels were significantly reduced in Ain-1 and Ain-2 grown in VLC, but mRNA accumulation was restored in the complemented strains. In contrast, the mRNA levels of LCIA were not affected in Hin-1 (Fig. 5B), and those of other VLC-inducible genes *LCIB* and *LCII* were also largely unchanged in Ain-1 and Ain-2, as well as Hin-1 (Fig. S4D). These results suggested that LCIA localized to the CE could affect the mRNA

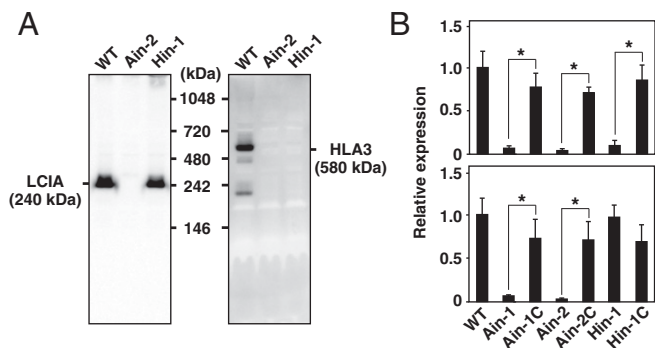


Fig. 5. Molecular masses of LCIA and HLA3 in nondenaturing conditions and effect of the absence of LCIA on HLA3 mRNA accumulation. (A) Molecular masses of LCIA and HLA3 in nondenaturing conditions. Total proteins were solubilized using 1.0% n-dodecyl β-D-maltoside and separated by blue-native PAGE. (B) Quantitative real-time PCR analyses of HLA3 (Upper) and LCIA (Lower) in WT, Ain-1, Ain-2, Ain-1C, Ain-2C, Hin-1, and Hin-1C. These cells were grown in very low CO₂ conditions for 4 h. Expression of each gene was normalized to *CBLP*. Data in all experiments indicate mean value ± SD from three biological replicates. **P* < 0.01.

expression level of HLA3 and subsequently caused a decrease in HLA3 protein accumulation.

Discussion

In this study, by characterizing the photosynthetic phenotype of LCIA and HLA3 insertion/overexpressing strains, it was revealed that HLA3 and LCIA are parts of the mechanism of HCO₃⁻ uptake through the PM and CE. These results elucidated a route of HCO₃⁻ uptake from the outside of cells to the chloroplast stroma by the cooperative function of HLA3 and LCIA.

Although LCIA could be associated with HCO₃⁻ uptake, the molecular mechanism remains elusive. LCIA is a homolog of formate transporter FocA and contains five amino acid residues (Fig. S2F) corresponding to those shown to form the pore of FocA (19, 33). FocA forms a symmetric pentamer that closely resembles the structure of aquaporin (33) and facilitates formate transport as a channel. Considering that LCIA was detected at 240 kDa in nondenaturing conditions (Fig. 5A), LCIA forms a protein complex as in the case of FocA. Furthermore, considering that the capacity for formate passage by FocA is increased by mutations of the aforementioned amino acids to smaller residues (33), examining the effect of similar mutations in LCIA could be helpful in elucidating the function of LCIA as a potential HCO₃⁻ channel. Relating to this hypothesis, a significant increase in Ci affinity at pH 6.2 was observed in LCIA-overexpressing strains (Fig. 4A, C, and D). Considering that external CO₂ at pH 6.2 should enter the cytoplasm continuously by passive influx, LCIA could function as a channel and cause an increase in the apparent Ci conductance with a minimal concentration gradient without waiting for a notable increase in Ci accumulation in the cytoplasm. In contrast, endogenous levels of HLA3 in HC conditions were not sufficient for Ci permeation toward the chloroplast stroma even with increased cytosolic Ci accumulation (Fig. 4B). These results suggested the functional importance of LCIA as a bottle neck step for increases in photosynthetic conductance across the CE.

By measuring the Ci accumulation and affinity of LCIA/LCI1-overexpressing strains and comparing the results with those of LCIA/HLA3-overexpressing strains, the degree of LCIA/HLA3-driven HCO₃⁻ uptake activity was evaluated (Fig. 4D). However, there were no differences in Ci accumulation and affinity at pH 9.0 compared with WT, suggesting that LCI1 was not related to the direct HCO₃⁻ uptake along with LCIA. Furthermore, although it was reported that Ci affinity was increased by the single overexpression of LCI1 at pH 7.8 (25), A1ox did not show a

significant increase in Ci affinity in the same pH conditions. This discrepancy could be caused by the difference in $K_{0.5}$ (Ci) values of the strains examined. For overexpressing LCI1 in the previous report, strain *lcr1* deficient in mRNA expression for at least three genes, *LCII*, *CAHI*, and *LCI6* (26), was used, and its $K_{0.5}$ (Ci) was $445 \pm 38 \mu\text{M}$ in HC conditions at pH 7.8 (25). In contrast, the $K_{0.5}$ (Ci) of strain C9 used as WT in this study was $230 \pm 27 \mu\text{M}$ in the same conditions, which was almost the same as $245 \pm 38 \mu\text{M}$ when LCI1 was overexpressed in *lcr1* (25). Thus, the effect of overexpressing LCI1 could be masked in A10x cells.

By means of *LCIA* insertion mutant analyses, it was shown that *LCIA* localized to the CE affected *HLA3* mRNA expression in the nucleus (Fig. 5B), which could throw new light on understanding the regulation of *LCIA* and *HLA3*. Considering that *LCIA* expression was not affected by the absence of *HLA3* (Fig. 5B), there may be unidentified retrograde signals from the chloroplast to the nucleus for maintaining *HLA3* mRNA expression. This possibility is supported by the recent study showing that transcript levels of *LCIA* and *HLA3* were simultaneously impaired in an HC-requiring mutant containing a disrupted *CAS* gene encoding a putative chloroplast calcium sensor protein and that other LC-inducible genes, such as *CAHI*, *LCII*, *LCIB*, and *LCIC*, were unaffected in the *CAS* mutant (34). Furthermore, this suggested that *LCIA* and *HLA3* could function cooperatively as part of the CCM and that *LCIA* has a key role in guaranteeing the maintenance of the HCO_3^- uptake system. Because *LCIA* and *HLA3* are conserved among aquatic algae, and owing to the structural

relationship of *LCIA* homologs with aquaporin (33), the *LCIA* and *HLA3* genes may have potential for genetic improvement of photosynthesis in land plants and algae.

Materials and Methods

C. reinhardtii strain C9 (photosynthetically WT strain available from the National Institute for Environmental Studies, Japan, as strain NIES-2235) was cultured in Tris-acetate-phosphate (TAP) medium for maintenance. For physiological experiments, cells were grown in liquid TAP medium for precultivation and diluted with modified high-salt medium [HSM (NH_4^+)] containing 9.35 mM NH_4Cl supplemented with 20 mM Mops (pH 7.0) to an OD_{730} of ~ 0.05 for photoautotrophic growth. To induce the expression of exogenous genes, cells grown in HSM (NH_4^+) medium for ~ 24 h to an OD_{730} of ~ 0.3 were collected by centrifugation and resuspended in fresh HSM (NO_3^-) containing 9.35 mM KNO_3 aerated with air enriched with 5% CO_2 (HC) or ordinary air containing 0.04% CO_2 (VLC). The culture conditions with combinations of medium and CO_2 concentrations are described as HC- NH_4^+ , HC- NO_3^- , VLC- NH_4^+ , and VLC- NO_3^- . For all culture conditions, cells were cultured at 25 °C with illumination at $80 \mu\text{mol photons}\cdot\text{m}^{-2}\cdot\text{s}^{-1}$.

Additional experimental procedures and methods are listed in the *SI Materials and Methods*.

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