



SHORT COMMUNICATION



## Requirement of ONE-HELIX PROTEIN 1 (OHP1) in early *Arabidopsis* seedling development and under high light intensity

Daniel Hey  and Bernhard Grimm 

Lebenswissenschaftliche Fakultät, Institut für Biologie, AG Pflanzenphysiologie, Humboldt-Universität zu Berlin, Berlin, Germany

### ABSTRACT

Plant ONE-HELIX PROTEINS (OHPs) are part of the light-harvesting complex superfamily whose members are involved in various processes related to sensing and capturing light as well as light protection. We recently showed the requirement of a functional OHP1-OHP2 heterodimer for efficient D1 synthesis. Interestingly, while the *ohp1* knockout mutant showed a strong defect in accumulation of the photosystem II and is hardly viable, virus-induced gene silencing of *OHP1* had no detectable impact on plant growth and performance under standard growth conditions. However, *in vivo* labeling assays with <sup>35</sup>S-methionine indicate a reduced D1 synthesis rate. Here, we show that VIGS-OHP1 plants are more susceptible towards elevated light intensities than control plants. This underlines an obligatory function of OHP1 for light acclimation.

### ARTICLE HISTORY

Received 16 October 2018  
Revised 7 November 2018  
Accepted 10 November 2018

### KEYWORDS

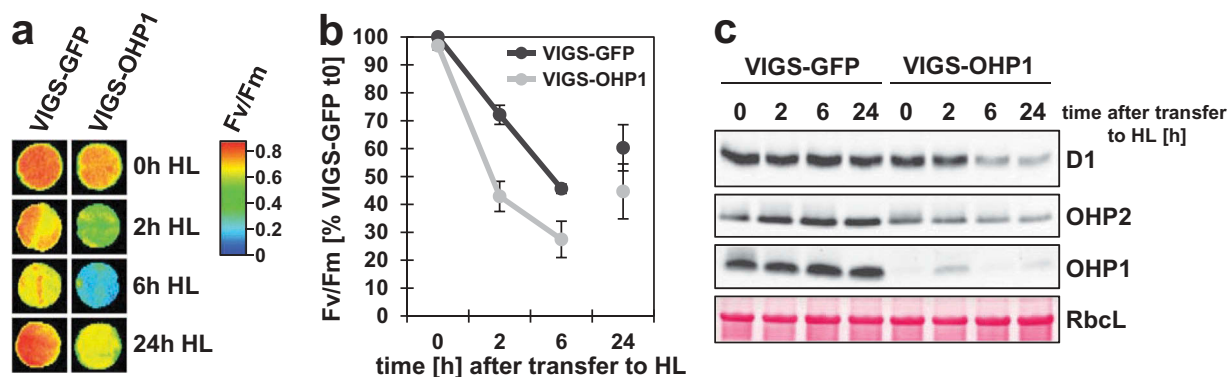
OHP1; one-helix protein; *Arabidopsis*; high light acclimation; photosynthesis and repair of photosynthetic complexes under light stress

Emerging insights into the function of the *Arabidopsis* one-helix proteins (OHPs) emphasize the important impact of OHPs on the accumulation of the photosystem II (PSII) core complex proteins<sup>1-3</sup> rather than a functional association of OHP2 with PSI, which was initially proposed.<sup>4</sup> However, a high light (HL)-induced increase in *OHP1* transcripts was reported<sup>2,5</sup> and OHP1 was postulated to be functionally correlated with the response to light stress. As it is currently assumed that both OHP variants act as OHP1-OHP2 heterodimers, their function depends on the mutual stability of both proteins.<sup>1</sup> Virus-induced gene silencing (VIGS) of *OHP2* phenotypically and biochemically resembled the *ohp2* T-DNA insertion mutant lines. They were characterized by an almost undetectably slow D1 synthesis rate, leading to a specific destabilization of PSII core proteins and PSII complexes.<sup>1</sup> In contrast, VIGS-OHP1 plants showed no phenotypical modification relative to the VIGS-GFP (green fluorescent protein) control seedlings under standard growth conditions as well as control-like levels of PSII proteins, although also in these plants the D1 synthesis rate was decreased.<sup>1</sup>

We continued to explore the impact of OHP1 on the D1 synthesis rate and stability and exposed the VIGS-plants to light stress conditions: Leaf discs of VIGS-plants were excised and exposed to HL (850  $\mu\text{mol photons s}^{-1} \text{m}^{-2}$ ) while floating on ultrapure water and being protected from the heat of the fluorescent lamps by a 5 cm water layer in a transparent container to prevent withering of the samples. Samples were harvested at the beginning as well as after 2, 6 and 24 hours of HL exposure and chlorophyll fluorescence was measured with a PAM imager after a 10 minute dark incubation (Figure 1(a)). Both VIGS-GFP and VIGS-OHP1 showed a similar Fv/Fm ratio at the beginning of the experiment, but in VIGS-OHP1 the Fv/Fm ratio decreased

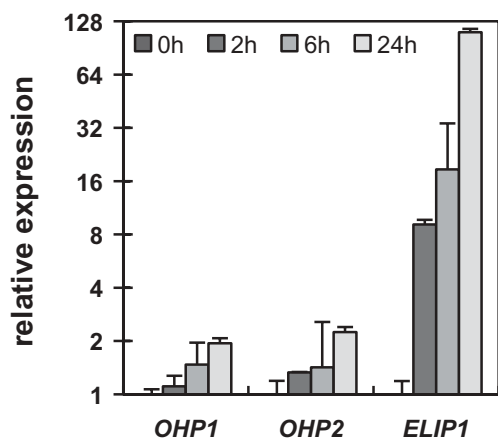
much faster upon HL exposure than in the VIGS-GFP control (Figure 1a, b)). After 24 hours of HL exposure the Fv/Fm ratio in both samples partially recovered, but still retained lower in VIGS-OHP1. To assess D1 levels, total proteins were separated by SDS-PAGE and analyzed by immunoblotting (Figure 1(c)). Whereas VIGS-GFP seedlings contained a stable amount of D1 during 24 hours of HL, D1 remarkably decreased after 6 hours of treatment in VIGS-OHP1 seedlings and remained at a low level until the end of the experiment. As reported earlier,<sup>1</sup> the OHP2 content decreased in VIGS-OHP1 leaf samples, while the content of OHP2 and OHP1 slightly increased in control seedlings during HL stress (Figure 1(c)).

The increased OHP1 content during the short-term HL exposure is in line with previous reports indicating a HL-dependent induction of *OHP1* expression.<sup>2,5</sup> When re-monitoring the expression of *OHP1* and *OHP2* by RT-qPCR, the VIGS-GFP leaf discs showed twice the amount of both transcripts after 24 hours of HL (Figure 2). We compared this result with other known HL-inducible genes, here for example *ELIP1*, which encodes EARLY LIGHT-INDUCIBLE PROTEIN 1, another member of the LHC protein family. However, the *ELIP1* transcripts accumulated more than 100-fold after 24 hours of HL exposure (Figure 2). In comparison, the twofold increased levels of the *OHP1* transcript seemed to be marginal. Thus, it remains open whether the observed increased OHP content (Figure 1(c)) corresponds to increasing transcript levels or is caused by HL-triggered accumulation or stabilization of OHPs. A decrease of the Fv/Fm ratio upon HL exposure, which indicates a lower maximum quantum efficiency of PSII, was determined for both lines, VIGS-GFP and VIGS-OHP1 (Figure 1(a, b)). Generally, a decreasing Fv/Fm ratio counts for stress exposure, such as HL, and is caused by damage or



**Figure 1.** VIGS-OHP1 plants are susceptible to high light intensities.

(a) Measurement of Fv/Fm ratios in leaf discs of VIGS-GFP and VIGS-OHP1 plants after exposure to high light ( $850 \mu\text{mol photons s}^{-1} \text{m}^{-2}$ ). False colors indicate Fv/Fm ratios and numerical values of the color code are indicated. One representative leaf disc for each time-point is shown. (b) Numeric representation of Fv/Fm ratios displayed in A. Data are normalized to Fv/Fm in VIGS-GFP at time-point  $t_0$ , which is set to 100%. Mean values and standard deviations of three biological replicates are shown. (c) Western blot analysis of OHP variants and D1 from leaf discs of VIGS-GFP and VIGS-OHP1 plants exposed to HL as described above. Ponceau S staining of the large subunit of RuBisCO (RbcL) is shown as the loading control.



**Figure 2.** Light-induced expression of OHP transcripts.

RT-qPCR analysis of *OHP1*, *OHP2* and *ELIP1* transcripts in VIGS-GFP plants during exposure to HL. Data were analyzed by the  $\Delta\Delta\text{Ct}$ -method and expression was normalized to the first time-point ( $t_0$ ). A mean value of three biological replicates is shown, error bars represent standard deviation.

inactivation of PSII, *i.e.* photoinhibition, or induced non-photochemical quenching.<sup>6</sup> The faster decreasing Fv/Fm ratios of VIGS-OHP1 compared to those of VIGS-GFP seedlings indicate a higher susceptibility of PSII towards excessive amounts of light in seedlings with OHP deficiency. *In-vivo* labeling assays with  $^{35}\text{S}$ -methionine unveiled a lower synthesis rate of D1 in VIGS-OHP1,<sup>1</sup> which is consequently also consistent with a strong decline of the D1 content after the HL periods (Figure 1(c)). In conclusion, impaired D1 recycling and/or lower stability correlate with the deficiency of OHP1 under HL conditions.

Besides the eukaryotic OHPs, the other members of the LHC superfamily possess one or two LHC-transmembrane helices. The cross-like configuration of these helices in the LHCPs and presumably also ELIPs contributes to the stability of the proteins.<sup>7</sup> By analogy, the two OHP variants are proposed to interact with each other to allow binding of

chlorophyll and/or precursors between the two transmembrane domains. It is proposed that the heterodimer formation is crucial for stability and function of both OHPs. Derived from the *ohp1* and the HL-exposed VIGS-OHP1 phenotype, OHP1 is predicted to be indispensable for effective functioning of the OHP1-OHP2 heterodimer during early seedling development and in adult plants under light stress. The early developmental stage is characterized by a rapid assembly of the photosynthetic complexes in a vulnerable environment, which resembles a light-sensitive state. Light stress conditions in adult plants require a high rate of the D1 repair cycle.<sup>8</sup> The complete set of (pigment supplying) auxiliary factors, including OHPs and HCF244<sup>9</sup> might be essential for acclimation of plants at both conditions. Thus, the seedling-lethal phenotype of the *ohp1* mutant and the stronger susceptibility of photosynthesis to HL-stress in the VIGS-OHP1 lines are explained with impaired action of the OHP heterodimer. Ultimately, the putative pigment-binding ability of the OHP-complex favors the hypothesis that the OHP1-OHP2 heterodimer formation is required to deliver chlorophyll for newly synthesized D1 proteins and to quench excitation energy of chlorophyll during a salvage pathway.<sup>10–13</sup>

## Materials and methods

### Plant material and growth conditions, vigs-assay

*Arabidopsis* wild-type plants (ecotype Col-0) were grown on soil in a growth chamber under long day conditions (16h photoperiod,  $100 \mu\text{mol photons s}^{-1} \text{m}^{-2}$ ,  $20^\circ\text{C}$ ). The VIGS-assay was performed with 12-day-old seedlings as previously described.<sup>1</sup>

### Light-stress assay

Leaf discs (7mm diameter) were excised from 5-week-old VIGS plants and incubated in HL ( $850 \mu\text{mol photons s}^{-1} \text{m}^{-2}$ ) for up to 24 hours floating on ultrapure water. To shield the discs from heat radiation of the lamps, a 5cm thick water basin was

placed on top of the samples. Samples were harvested at indicated time-points.

### Chlorophyll fluorescence measurements

Chlorophyll fluorescence kinetics of leaf discs were measured in a PAM-imager chamber (FluorCam 700MF, Photon Systems Instruments) after ten minutes of dark incubation. Fv/Fm ratios were normalized to time-point T<sub>0</sub> of VIGS-GFP and displayed in per cent.

### SDS-PAGE, Western blot

Total leaf protein extraction, SDS-PAGE, Western blotting and chemiluminescence detection was performed exactly as previously described.<sup>1</sup>

### RNA extraction, RT-qPCR

Total RNAs were extracted by a citric acid based extraction method<sup>14</sup> and cDNA-synthesis, reverse transcription and RT-qPCR was performed exactly as described before.<sup>1</sup> Data were analyzed by the  $\Delta\Delta C_t$ -method.<sup>15</sup>

### Abbreviations

GFP	green fluorescent protein
HL	high light
OHP	one-helix protein
PAM	pulse-amplitude modulation
PS	photosystem
RbcL	large subunit of Ribulose-1,5-bisphosphate carboxylase/oxygenase
VIGS	virus-induced gene silencing

### Funding

This work was supported by the German Research Foundation (Deutsche Forschungsgemeinschaft, DFG) to B.G. (Gr936 14-1, subproject of the DFG-Research Unit FOR2092).

### ORCID

Daniel Hey  <http://orcid.org/0000-0002-8749-8352>  
Bernhard Grimm  <http://orcid.org/0000-0002-9730-1074>

### References

- Hey D, Grimm B. ONE-HELIX PROTEIN 2 (OHP2) is required for the stability of OHP1 and assembly factor HCF244 and is functionally linked to PSII biogenesis. *Plant Physiol.* 2018;177:1453–1472. doi:10.1104/pp.18.00540.
- Myouga F, Takahashi K, Tanaka R, Nagata N, Kiss AZ, Funk C, Nomura Y, Nakagami H, Jansson S, Shinozaki K, et al. Stable accumulation of photosystem II requires ONE-HELIX PROTEIN1 (OHP1) of the light harvesting-like family. *Plant Physiol.* 2018;176:2277–2291. doi:10.1104/pp.17.01782.
- Beck J, Lohscheider JN, Albert S, Andersson U, Mendgen KW, Rojas-Stütz MC, Adamska I, Funck D. Small one-helix proteins are essential for photosynthesis in Arabidopsis. *Front Plant Sci.* 2017;8:7. doi:10.3389/fpls.2017.00007.
- Andersson U. Light stress-induced one-helix protein of the chlorophyll a/b-binding family associated with photosystem I. *Plant Physiol.* 2003;132:811–820. doi:10.1104/pp.102.019281.
- Jansson S, Andersson J, Kim SJ, Jackowski G. An Arabidopsis thaliana protein homologous to cyanobacterial high-light-inducible proteins. *Plant Mol Biol.* 2000;42(2):345–351.
- Murchie EH, Lawson T. Chlorophyll fluorescence analysis: a guide to good practice and understanding some new applications. *J Exp Bot.* 2013;64:3983–3998. doi:10.1093/jxb/ert208.
- Kühlbrandt W, Wang DN, Fujiyoshi Y. Atomic model of plant light-harvesting complex by electron crystallography. *Nature.* 1994;367:614–621. doi:10.1038/367614a0.
- Aro EM, McCaffery S, Anderson JM. Photoinhibition and D1 protein degradation in Peas acclimated to different growth irradiances. *Plant Physiol.* 1993;103:835–843. doi:10.1104/pp.103.3.835.
- Komenda J, Sobotka R. Cyanobacterial high-light-inducible proteins - protectors of chlorophyll-protein synthesis and assembly. *Biochim Biophys Acta* 2016;1857:288–295. doi:10.1016/j.bbabi.2015.08.011.
- Shukla MK, Llansola-Portoles MJ, Tichý M, Pascal AA, Robert B, Sobotka R. Binding of pigments to the cyanobacterial high-light-inducible protein HliC. *Photosyn Res.* 2018;137:29–39. doi:10.1007/s11120-017-0475-7.
- Staleva H, Komenda J, Shukla MK, Šlouf V, Kaňa R, Polívka T, Sobotka R. Mechanism of photoprotection in the cyanobacterial ancestor of plant antenna proteins. *Nat Chem Biol.* 2015;11:287–291. doi:10.1038/nchembio.1755.
- Lin YP, Endow JK, Dufour J, Zhu J, Inoue K. Analysis of an Arabidopsis heat-sensitive mutant reveals that chlorophyll synthase is involved in reutilization of chlorophyllide during chlorophyll turnover. *Plant J.* 2014;80:14–26. doi:10.1111/tpj.12655.
- Komenda J, Sobotka R, Nitschke W, Baymann F, Neehaul Y, Hellwig P, Richers S, Vonck J, Bott M, Hunte C. Cyanobacterial high-light-inducible proteins - protectors of chlorophyll-protein synthesis and assembly. *Biochim Biophys Acta.* 2016;1857:288–295. doi:10.1016/j.bbabi.2016.07.009.
- Onate-Sanchez L, Vicente-Carbajosa J, Stern AF, Byrd JR, Cherlin EJ, Wang Y, Yuan C, Nembhard I, Brush JE, Krumholz HM. DNA-free RNA isolation protocols for Arabidopsis thaliana, including seeds and siliques. *BMC Res Notes.* 2008;1:93. doi:10.1186/1756-0500-1-23.
- Pfaffl MW. A new mathematical model for relative quantification in real-time RT-PCR. *Nucl Acids Res.* 2001;29:e45. doi:10.1093/nar/29.9.e45.