

**Mechanism of fucoxanthin biosynthesis in *Phaeodactylum tricornutum* under different light intensities via FCP complex formation**

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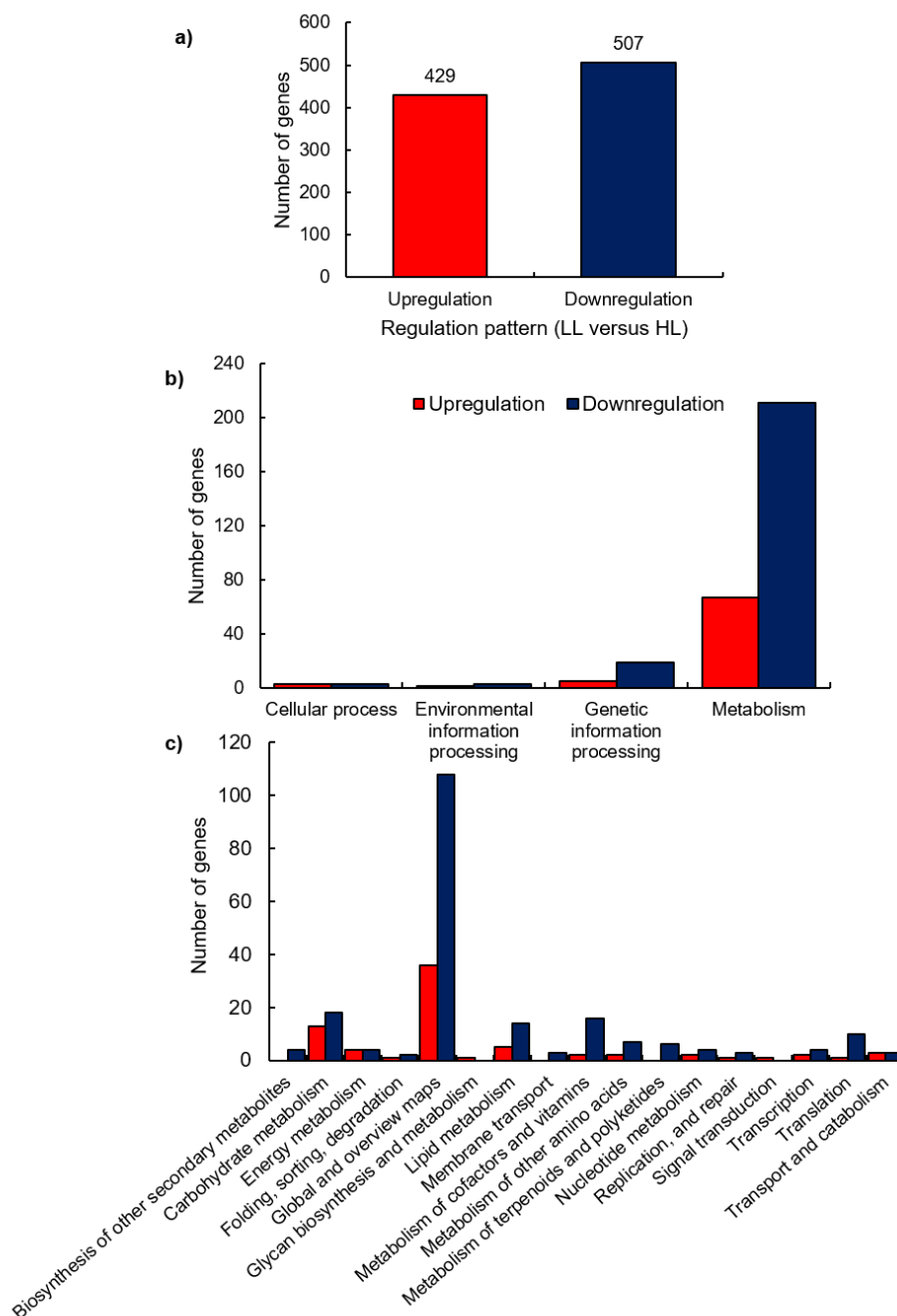
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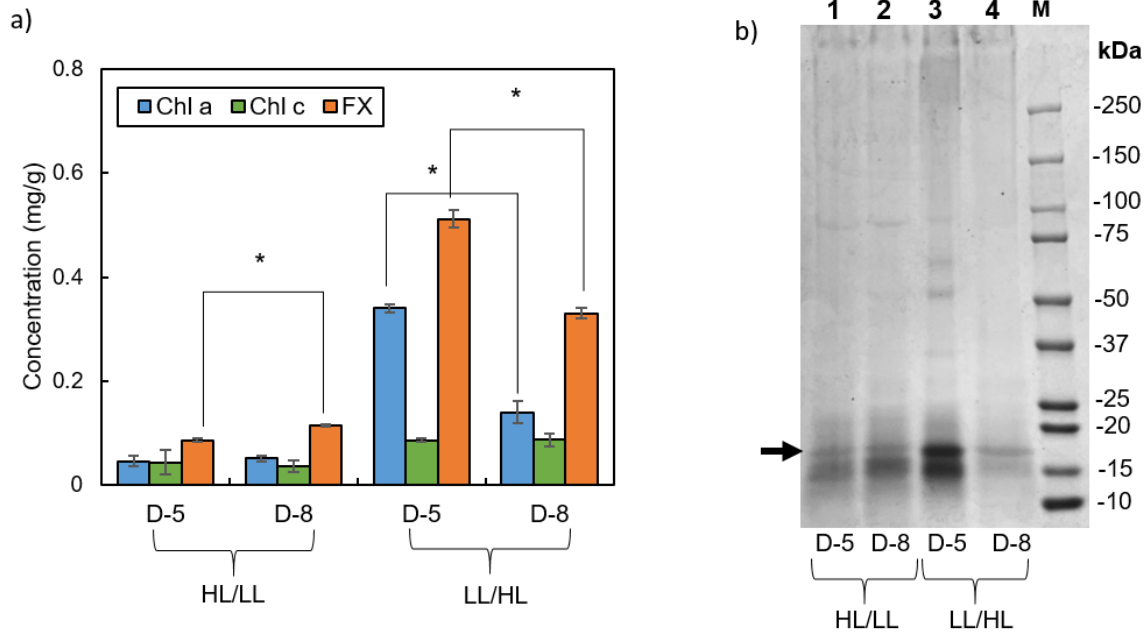
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**Supplementary Figure S1. Differentially expressed gene (DEG) analysis.** (a) Total DEGs downregulated and upregulated in low light (LL)-grown *P. tricornutum* on day 5 of culture; (b) Classification and (c) Sub-classification of KEGG pathways related to identified DEGs. Results were normalized to data obtained from high light (HL)-grown *P. tricornutum* on day 5 of culture.



**Supplementary Figure S2. Thylakoid membrane (TM) analysis.** (a) Photosynthetic pigments in TM fractions isolated from day 5 (D-5) and day 8 (D-8) *P. tricornutum* cultures; (b) Detection of the FCP antennae enrichment (black arrow) in TM fractions using SDS-PAGE. Light transition was the switch from high-to-low light (HL/LL) and from low-to-high light (LL/HL). Asterisk denotes significant differences analyzed using unpaired t-test, two-tailed  $p < 0.05$ . Chl a, chlorophyll a; Chl c, Chlorophyll c; FX, fucoxanthin; M, marker; Lane 1, day 5 culture (before HL/LL transition); Lane 2, day 8 culture (after 3 days of HL/LL transition); Lane 3, day 5 culture (before LL/HL transition); Lane 4, day 8 culture (after 3 days of LL/HL transition)

**Supplementary Table S1. Primers used for quantitative real-time polymerase chain reaction.**

H4, Histone H4; PSY, phytoene synthase; PDS, phytoene desaturase; VDL, violaxanthin de-epoxidase-like protein; LHCF, light harvesting chlorophyll a/c-fucoanthin binding proteins; Fw, forward; Rv, reverse.

| Annotation              |          | Primer sequence (5'–3')                       | Reference                 |
|-------------------------|----------|-----------------------------------------------|---------------------------|
| <b><i>PtH4</i></b>      | Fw<br>Rv | AGGTCCTTCGCGACAATATC<br>ACGGAATCACGAATGACGTT  | Siaut <i>et al</i> , 2007 |
| <b><i>PtPDS1</i></b>    | Fw<br>Rv | GTGGCAAACAATGCCCTACT<br>AGTGATCGACGGCTTTGAGT  |                           |
| <b><i>PtPDS2</i></b>    | Fw<br>Rv | GTCAGGAACACGTGACGGTA<br>GCATCGAGAGTTGTTCCGGT  |                           |
| <b><i>PtPSY</i></b>     | Fw<br>Rv | CCACGCCGAACATGCTTTAG<br>GACTTCTTGCACTTGTGCCG  | Yang and Wei, 2020        |
| <b><i>PtVDL1</i></b>    | Fw<br>Rv | GTTCTTCTCCCAGAGCGGTA<br>CGAATCCGTCGTACTTGACG  |                           |
| <b><i>PtLhcf3/4</i></b> | Fw<br>Rv | CGGTGACCAGGAGAAGTTTCG<br>ATCTCAGCGAAGGTCTTGCC |                           |
| <b><i>PtLhcf5</i></b>   | Fw<br>Rv | GACTTCCGGGAGACATCGAC<br>CGATCTGGGCAATACCAGCA  |                           |
| <b><i>PtLhcf8</i></b>   | Fw<br>Rv | AGGAAGTGGAAGGCAAGAGC<br>GTGGACCATGAGTGCGAGAA  |                           |