

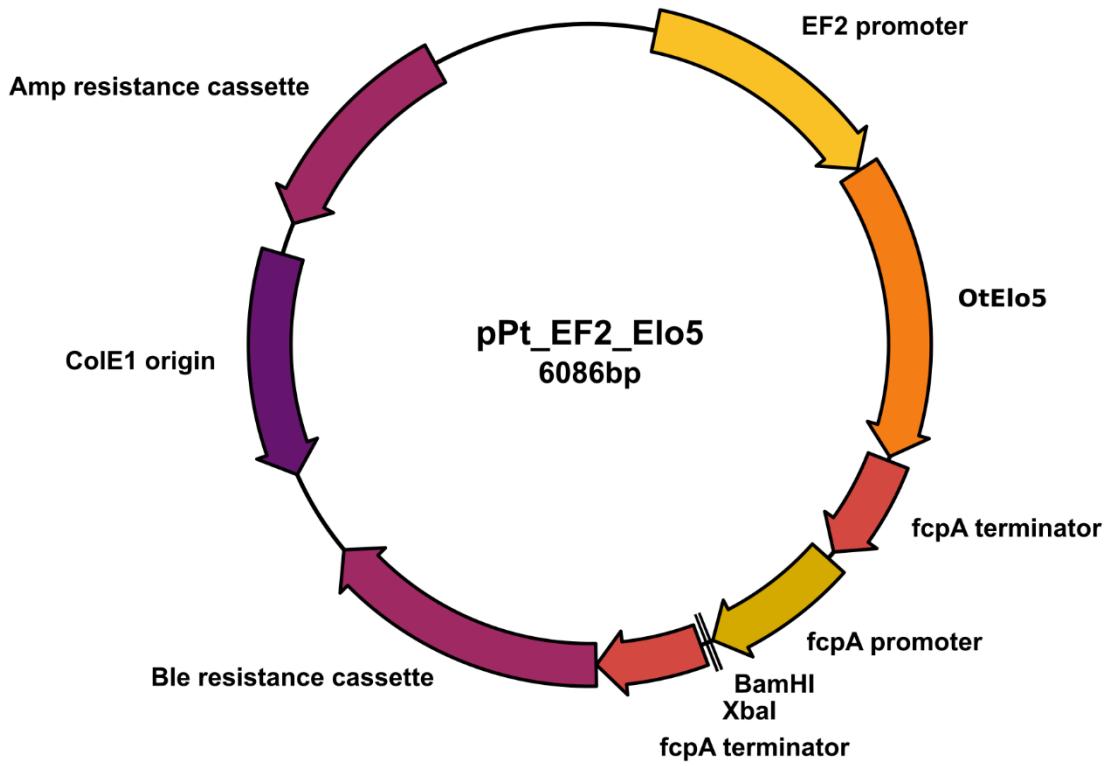
Supplementary Data Information

**Multifunctionilizing the marine diatom *Phaeodactylum tricornutum* for sustainable co-production
of omega-3 long chain polyunsaturated fatty acids and recombinant phytase**

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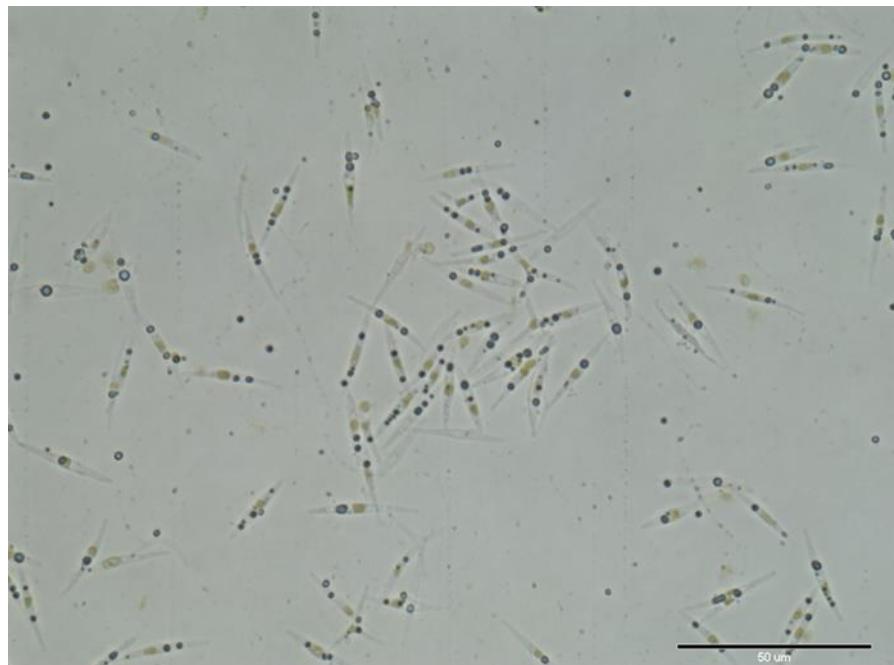
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Supplementary Figure S1. Plasmid map of the expression vector Pt_EF2_Elo5.

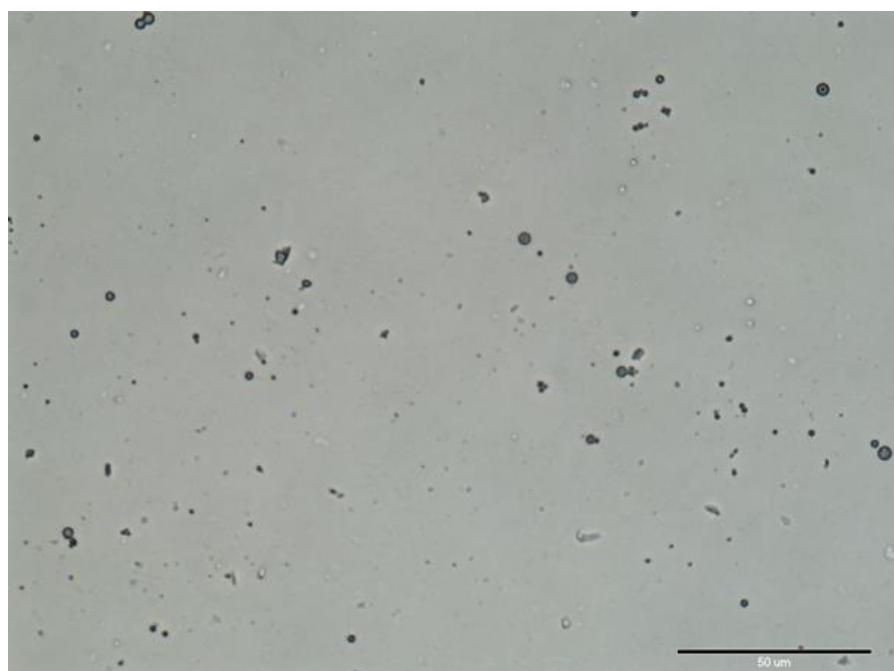


Supplementary Figure S2. Lysis of *P. tricornutum* cells by sonication. (a) *P. tricornutum* cells before sonication. (b) Visualization of cellular debris after sonication. Scale bar: 50 μ M.

a

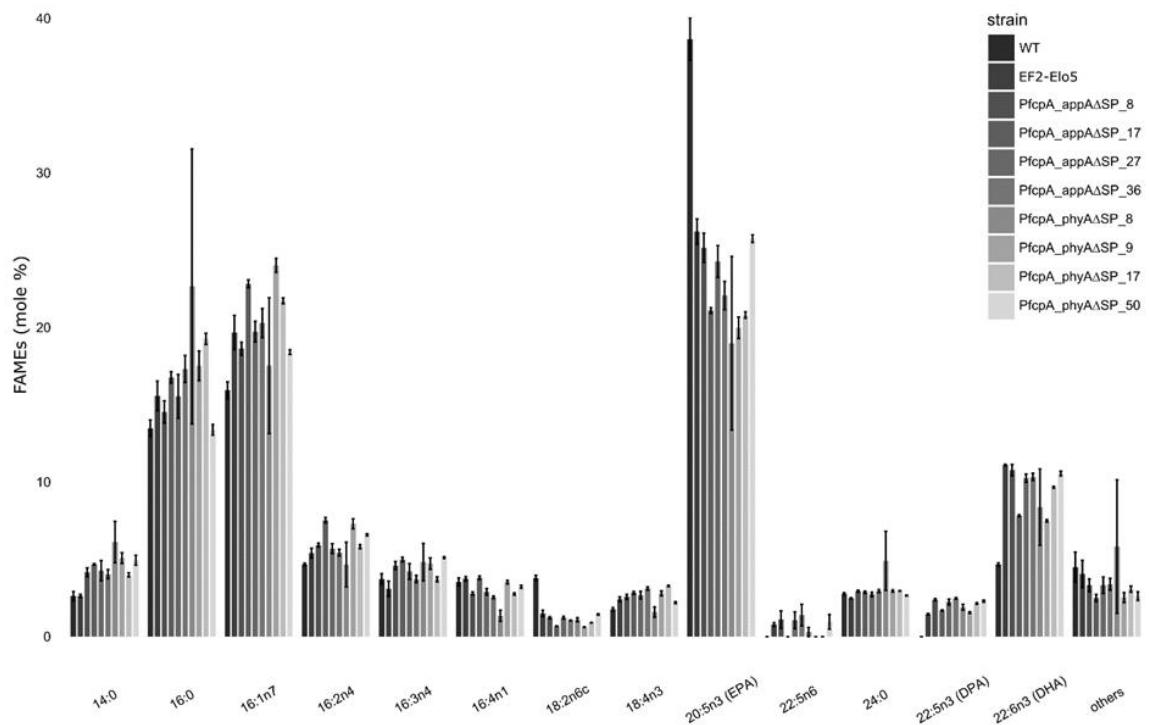


b

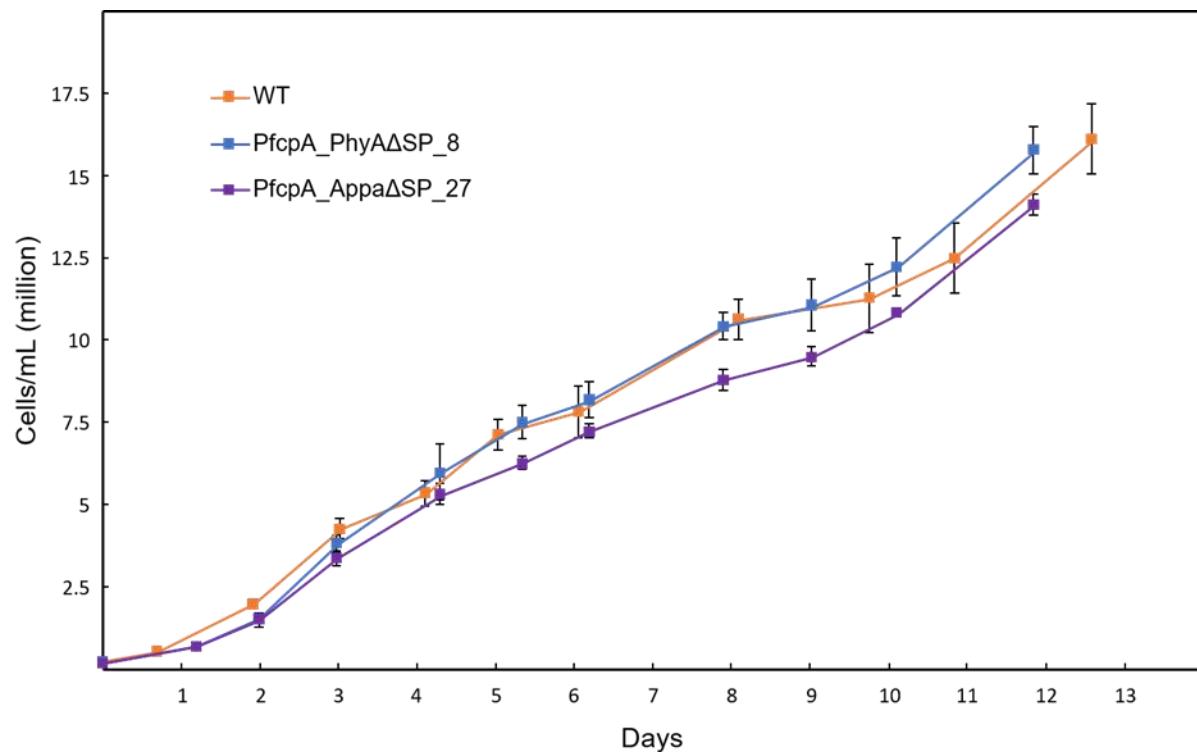


Supplementary Figure S3. Fatty acid composition of WT and transgenic strains of *P. tricornutum*.

Values are averages of measurements of at least three independent cultures, \pm standard error.



Supplementary Figure S4. Cellular growth of WT and transgenic *P. tricornutum* strains. Values are averages of measurements of three independent cultures, \pm standard error.



Supplementary Table 1. Fatty acid composition and phytase activities of WT and selected transgenic *P. tricornutum* strains. Values are averages of measurements of at least three independent cultures, \pm standard error; ND: not detected; FTU/mg soluble protein.

Construct	FTU/mg	EPA (mole %)	DPA (mole %)	DHA (mole %)
WT	3.13 \pm 0.36	38.64 \pm 1.35	ND	4.67 \pm 0.10
Pt_EF2_Elo5	ND	26.2 \pm 0.82	1.45 \pm 0.04	11.09 \pm 0.04
PfcpA_appA Δ SP_8	23.13 \pm 0.96	25.15 \pm 0.94	2.38 \pm 0.07	10.76 \pm 0.37
PfcpA_appA Δ SP_17	39.18 \pm 1.86	21.09 \pm 0.19	1.69 \pm 0.03	7.82 \pm 0.06
PfcpA_appA Δ SP_27	35.63 \pm 2.29	24.26 \pm 1.03	2.23 \pm 0.19	10.24 \pm 0.26
PfcpA_appA Δ SP_36	25.2 \pm 1.40	22.05 \pm 0.91	2.47 \pm 0.05	10.32 \pm 0.23
PfcpA_phyA Δ SP_8	40.2 \pm 1.22	18.97 \pm 5.62	1.9 \pm 0.19	8.37 \pm 2.47
PfcpA_phyA Δ SP_9	32.17 \pm 1.00	19.98 \pm 0.69	1.54 \pm 0.06	7.49 \pm 0.08
PfcpA_phyA Δ SP_17	31.59 \pm 0.61	20.8 \pm 0.20	2.14 \pm 0.05	9.65 \pm 0.05
PfcpA_phyA Δ SP_50	29.35 \pm 0.91	25.74 \pm 0.25	2.29 \pm 0.06	10.54 \pm 0.15

Supplementary Table 2. The total soluble protein content of WT and transgenic PfcpA_appAΔSP_27 and PfcpA_phyAΔSP_9 strains. Values are averages of measurements of at least three independent cultures, ± standard error.

	Cells /mL	Protein (mg /mL)	Protein (µg /mL) /10 ⁷ cells
WT	2.24 × 10 ⁷	2.48 ± 0.068	110 ± 3
PfcpA_phyAΔSP_8	2.06 × 10 ⁷	2.15 ± 0.102	104 ± 2
PfcpA_appAΔSP_27	2.28 × 10 ⁷	2.33 ± 0.125	102 ± 4

