

Version 3

Nov 06, 2019

Transient transformation of *Ostreococcus* species (OTTH595, RCC809 and RCC802) and *Bathycoccus* V.3

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Protist Research to Optimize Tools in Genetics (PROT-G)

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ABSTRACT

This protocol describes the preparation of cells and introduction of DNA into the cells by electroporation. For selection of stable transformants or measure of transient gene expression see related protocols.

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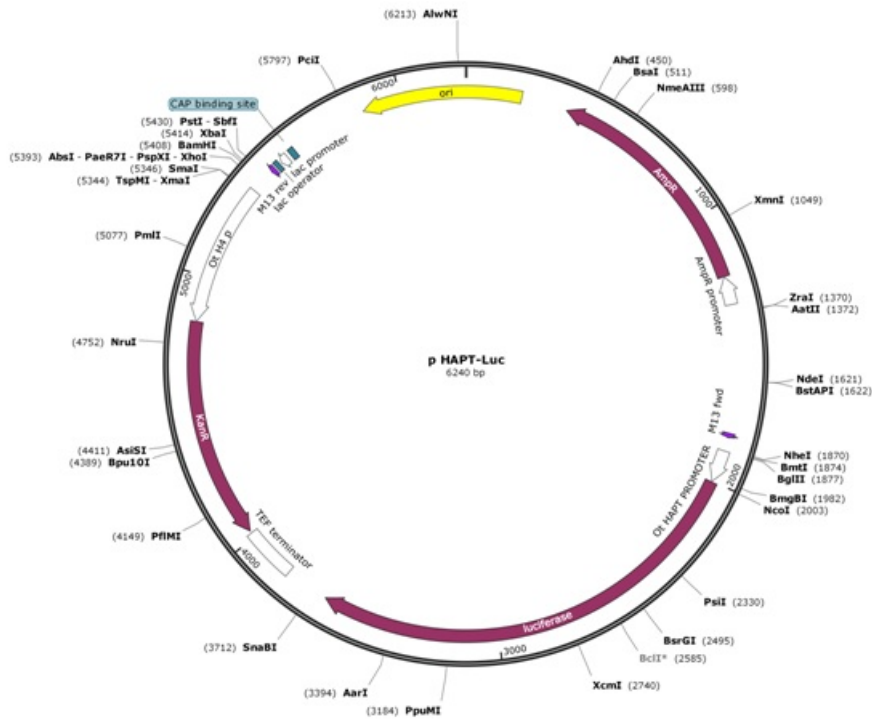
29524

MATERIALS TEXT

***Ostreococcus lucimarinus* (RCC802)**

pHAPT:Luc vector (Djouani Tahri et al., PLOS ONE 2011 <https://doi.org/10.1371/journal.pone.0028471>)

pHAPT:Luc map



PHAPT:Luc sequence

O. tauri pHAPT:Luc6240 bp ds-DNAcircular
 DEFINITIONsynthetic circular DNA

FEATURESLocation/Qualifiers

source1..6240

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/mol_type="other DNA"

CDScomplement(377..1237)

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/note="common sequencing primer, one of multiple similar variants"

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Validation of *O. lucimarinus* transgenic lines by PCR (Figure 4A)

Forward primer: 1GAGCGCAACGGTACCCGGGCGGTACCTGTGCG

Reverse primer: CTGGCGACGCTGACGGCGTACTTCACGT

***Bathycoccus prasinos* (RCC4222)**

pH4:KanMx pHAPT:Luc transgene (PCR product)

Construction of the transgene by PCR :

The transgene was generated by fusion PCR as described in Shevchuk et al. (2004) using DNA templates described below and oligonucleotides mapped on the sequence below.

Promoters of Histone H4 and high affinity phosphate (HAPT) transporters were amplified using (Fas1/Ras1) and (Fas4/Ras4) oligonucleotides respectively. KanMx and Luciferase sequences were amplified from pHAPT:luc plasmid template using oligonucleotides (Fas2/Ras2) and (Fas3/Ras3). Final PCR was done on amplified fragments using (Fas5/Ras5) oligonucleotides.

Oligonucleotide sequences:

Fas1: CCGGCTTCGTGATGCCTTGGATGTTGTCTC

Ras1: TCGAAACGTGAGTCTTTTCCTTACCcatTGTGTTTGATTTATAATGAGGTTTTCTT

Fas2: AAGAAAACCTCATTATAAATCAAACACAatgGGTAAGGAAAAGACTCACGTTTCGA

Ras2: CCAAGAAGGGCGGAAAGATCGCCGTGTAAGAAATACCGTCTATCATCGATGAATTCGA

Fas3: CGAATTCATCGATGATAGACGGTATTTCTTACACGGCGATCTTCCGCCCTTCTTGG

Ras3: AAGCATATATACATACACACATACAAAATGGAAGACGCCAAAAACATAAAGAAAGG

Fas4: CCTTCTTTATGTTTTGGCGTCTTCCATTTGTATGTGTGTGTATATATGCTT

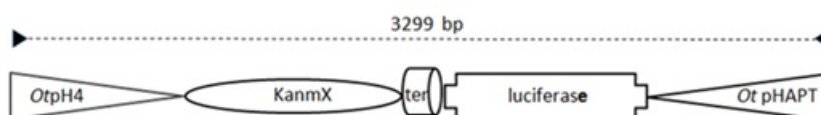
Ras4: AAGGTACACGACCAAACGCGCGTCGAC

Fas5: CCACCTTACCTCTGCCGGACATTGTGA

Ras5: GCGCGTAATATCTACGAGGTAGCACGAG

Note that transformants were checked by PCR using (to amplify Kanx

Map of the transgene:



Sequence of the transgene:

4020 bp ds-DNA linear

REFERENCE1(bases 1 to 4020)

AUTHORSFY Bouget and JC Lozano

FEATURES Location/Qualifiers

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1741 atcatctgcc cagatcgaa gtaagtgcg cagaaagtaa tatcatgctt caatcgtatg
1801 tgaatgctgg tcgtatact gctgtcatt cgatactaac gccgccatcc agtgcgaaa
1861 acgagctcga atcatcgat gatagacggg atttcttaca cggcgatctt tccgcccttc
1921 ttggccttta tgaggatctc tctgatttt cttgcgtcga gtttccggg aagaccttc
1981 ggtactcgt ccacaaacac aactcctccg cgcaactttt tcgcggttgt tacttgactg
2041 gcgacgtaat ccacgatctc ttttccgctc atcgctttc cgtgctccaa aacaacaacg
2101 gcggcgggaa gtcaccggc gtcacgctc ggaagacctg cgacacctgc gtcgaagatg
2161 ttgggtgtt ggagcaagat ggattccaat tcagcgggag ccacctgata gcctttgtac
2221 ttaatcagag acttcaggcg gtcaacgatg aagaagtgtt cgtctcgtc ccagtaagct
2281 atgtctccag aatgtagcca tccatcctg tcaatcaagg cgttggtcgc ttccggattg
2341 ttacataac cggacataat cataggacct ctcacacaca gttcgcctct ttgattaacg
2401 ccagcgttt tcccggtatc cagatccaca acctcgtt caaaaaatgg aacaacttta
2461 ccgaccgcgc cgggtttatc atccccctc ggtgtaatca gaatagctga ttagtctca
2521 gtgagccat atccttgcct gatactggc agatggaacc tcttggaac cgctccccg
2581 actccttag agaggggagc gccaccagaa gcaattcgt gtaaataga taaatcgtat
2641 ttgcaatca gagtgccttt ggcgaagaag gagaataggg ttggcaccag cagcgcact
2701 tgaatctgtt aatcctgaag gctcctcaga aacagctctt ctcaaatct atacattaag
2761 acgactcga atccacat caaataacc agttagtaa acattccaa accgtgatgg
2821 aatggaacaa cacttaaaat cgcagatcc ggaatgatt gattgcaaa aataggatc
2881 ctggcatgag agaattcac gcaggcagtt ctatgaggca gagcgacacc tttaggcaga
2941 ccagtagatc cagaggagtt catgatcgt gcaattgtc tctcctatc gaaggactc
3001 ggcacaaaat cgtattcatt aaaaccggga ggtagatgag atgtgacgaa cgtgtacatc
3061 gactgaaatc cctggaatc cgttttagaa tccatgataa taatttttg gatgattggg
3121 agctttttt gcacgttcaa aatttttgc aaccctttt tggaacgaa caccacggta
3181 ggctgcgaaa tgccatact gttgagcaat tcacgttcat tataaatgc gttcgcgggc
3241 gcaactgcaa ctccgataa taacgcgcc aacaccggca taagaattg aagagattt
3301 tcaactgata cgacgattct gtgatttga ttacgccat atcgtttcat agcttctgcc
3361 aaccgaacgg acatttcgaa gtactcagcg taagtgatg ccacctgat atgtcatct
3421 gtaaaagcaa ttgtccagg aaccagggcg tatctctca tagccttatg cagtgtctc
3481 ccagcgggic catctccag cggatagaat ggcgccgggc cttctttat gttttggcg
3541 tctccattt tgtatgtg tgtatgata tatgcttgg gaatatatgt tcacagaatg
3601 acgacttga aagcgcgtt gaattttaa acgaaaatc cgtgtggct gatattttt
3661 gcttttgc tttttcaac caccggatt ttgtttt ttcaaaaca cccaccgacc
3721 gtaaatgtg tgttcttg tttctgtg ggctgttc ttttaggga gggaggatg
3781 attcagagt aatatatt atgtgctcc agatctggt tatacagga gttggtgtg
3841 gcttttaac acacaaaata cgcctaacg cgaggaggcg tcttgaaac gtaaggtac
3901 tattactct gctacctg agatattacc gcgcgtaatt agaagctg ggagttgtg
3961 tegtgtct tgtatctc gacgcgctt tggctgtg cactttact gcgccgcgac
//

Validation of *B. prasinos* transgenic lines by PCR (Figure 4b)



Oligonucleotide sequences

1(Fas5) CCACCTTTACCTCTGCCGACATTGTGA

2TCGAAACGTGAGTCTTTTCCTTACC

3 TCGCCTCGACATCATCTGCCAGATGC

4 (Ras5)GCGCGTAATATCTACGAGGTAGCACGAG

Reference:

Nikolai A. Shevchuk, Anton V. Bryksin, Yevgeniya A. Nusinovich, Felipe C. Cabello, Margaret Sutherland, Stephan Ladisch /Nucleic Acids Research/, Volume 32, Issue 2, 16 January 2004, Page e19, <https://doi.org/10.1093/nar/gnh014>

Cell preparation

- 1) Starting from a culture of *Ostreococcus tauri*, RCC809 or *Bathycoccus* in stationary phase, inoculate cultures at 1 million cells/ml as determined by flow cytometry (Accuri C6 BD) in 200 ml plastic flasks in Artificial Seawater supplemented with Keller medium supplement (trace metals, vitamins, nitrate and Phosphate as described in Djouani Tahri et al., PLoS ONE 2011). For each transformation (including control), you should plan on using 50 ml de culture in exponential phase.
 - 2) Grow cells for 4 to 5 days depending on the light conditions, until they reach densities of 30 to 40.10⁶ cells/ml.
 - 3) Count cells by flow cytometry. Check by SyBR Green II staining that bacterial contamination is below 2%.
 - 4) Transfer lcultures to 50 ml Falcon tubes.
 - 5) Centrifuge at 8000g for 10 min at 4°C.
 - 6) Remove the supenatant, resuspend the cell pellet in 1 ml de sorbitol 1M (pH 7.5) in H₂O MQ, at 4°C.
 - 7) Transfére the cell suspension to 1.5 ml ependorf.
 - 8) Centrifuge at 8000g for 10 min at 4°C.
 - 9) Remove 900 µl of supernatant
 - 10) **Resuspend cells by gently pipeting.**

Electroporation of the transgene

- 2)
 - 1) Add 5µg of transgene DNA to cell suspension. Keep on ice for 5 minutes. The transgene consist of the high affinity phosphate promoter fused to the firefly luciferase (see Djouani Tahri et al., PloS one 2011).
 - 2) Transfer cells to a 2 mm electroporation cuvette (Biorad).
 - 3) Apply an electric field
For *Ostreococcus tauri* (OTTH595) : **capacitance**: 25µF, **resistance** 600 Ω, **voltage** 1.35KV.
For *Ostreococcus* sp RCC809 : **capacitance**: 25µF, **resistance** 600 Ω, **voltage** 1.4KV.
For *Bathycoccus* (RC4222) : **capacitance**: 25µF, **resistance** 600 Ω, **voltage** 1.5KV.
For *Ostreococcus lucimarinus* RCC802 : **capacitance**: 25µF, **resistance** 600 Ω, **voltage** 1.2KV.
 - 4) Add 1ml of fresh culture Medium to resuspend the cells.
 - 5) Add 40 ml of culture medium and transfer to a culture flask.
 - 6) Incubate at 20°C overnight in a light incubator.

At this stage, transient transgene expression can me measured or stable transformants can be selected (see relevant protocols).