

Nucleotide sequence of chloroplast CF1-ATPase ϵ -subunit and elongator tRNA^{Met} genes from *Arabidopsis thaliana*

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A lambda clone has been isolated from a genomic library of *Arabidopsis thaliana* as described (2). A 2.8 kb Eco R I fragment has been sequenced partially. The analysis of the sequence has shown that this fragment contains the chloroplast *atpB-atpE-trnM-trnV* gene cluster. Here, we present the nucleotide sequence of the *atpE* and *trnM* genes. The aminoacid sequence deduced from the *atpE* gene sequence from *A. thaliana* displays 87.9% and 89.4% homology with its chloroplast counterparts from tobacco (3) and spinach (4) respectively. The 3' end of the *atpB* gene overlaps the *atpE* gene by 4 bp. The elongator tRNA^{Met} gene shows only one nucleotide difference with its chloroplast counterpart from soybean and 5 nucleotide differences with its mitochondrial counterpart from *A. thaliana* (2).

<pre> GAAGGCTACCAACTTAAAGAGAGTAAACTGAAGAAATGACCTTAATCTTTGTACTGACTCCGAATCGAATTG atpB M T L N L C V L T P N R I V 80 120 160 TTGGGATTCAAGTAGAAGAAATCATTATCTACTAAATAGTGACAAATTGGTGTATTAGCAAACACGGCGCGATTG W D S E V E E I I L S T N S G Q I G V L A N H A P I 200 240 280 320 CCACAGCTGTTGATAGGTATTTGACAATACGCCCTGCTAACCAATGGTTAACGATGGCTCTGATGGCGGTTTGC A T A V D I G I L T I R L A N Q W L T M A L M G G F A 360 400 440 480 AGATAAGGCAATAATGAAATCACTATTTAGTAAATGATGCAGAGAAGAATAGTGACATTGATCCACAGAACGCTCAGCA R I G N N E I T I L V N D A E K N S D I D P Q E A Q Q 520 560 600 640 AACTCTGAAATAGCAGAACGCTAACTTGAGAAAAGCTGAGGTAAGAGCACAGACAATTGAGGCTAATCTAGCTCTCAGAC T L E I A E A N L R K A E G K R Q T I E A N L A L R 680 720 760 800 840 880 GAGCTGGCACCGAGTCGAGGCTCTAACAGCTTGTAACTAGCTGACGTGAAAAAGATGTATAAAAAT R A R T R V E A L N T I — 960 AGATCGAAAGCAGATAAAAACAATAAAATAAAAAGCTGTTACAAAAACTTATTAGACTACACCATGATCATGGTGT 1000 CTAATAAGTTATACCTACTATTGGATTTAACCAATGACTCTGCCGTATGAAAGCAATACTCTAACCACTGAGTTAACG atggatgataactcaaacttggtaactggactggaggacgcatacttcgttatgagattgtgactcaattca 1040 1080 AGTTTATTATCATTGTAAGAGAAGGAAAGGGATACCTATCACATCGATAGGTTATAAAATCCAATTATTCTAACGA tcca trnM 1120 1160 ATACCAATAAAACATGAGATCAAAGAGATAATGTTGATCATGTAATATTACCTTGACAAGAAATTCTACATGAT trnV 1200 1240 AAGATAAAATGTGTATCACAAACACAAGGGCTATAGCTCAGTTAGGTAGAGCACCTCGTTACACGTGCGCAAAGTTT 1280 1320 </pre>
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Fig.: Nucleotide sequence of *A.thaliana* chloroplast *atpE* and *trnM* genes. The 3' end of *atpB* gene and the 5' end of *trnV* gene are underlined. The non-transcribed strand of the *trnM* gene is shown in lower-case letters. The inverted repeat sequence located downstream of *trnM* gene is indicated by arrows.

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