

Nucleotide sequences of the *Rana catesbeiana* mitochondrial small (12S) and large (16S) ribosomal RNA genes

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The entire mitochondrial genome of *Rana catesbeiana* was cloned into a plasmid vector pBR322 at the unique Bam HI site (1) and the nucleotide sequences of the small (12S) and large (16S) rRNA genes were determined (Fig.1). The sequence of the 12 S rRNA gene showed 68% and 75% homologies, and that of the 16 S rRNA showed 69% and 74% homologies, to their counterparts of the human and *Xenopus laevis* mitochondrial genomes, respectively. Analysis of the contiguous genes revealed that the 5'- and 3'- ends of the 12 S rRNA gene were flanked by the tRNA genes for phenylalanine and valine, while those of the 16 S rRNA gene were flanked by the tRNA genes for valine and leucine (UUR), respectively (not shown).

A

CAAGGTTTG GTCCTAGCCT TATTATCAC TGTTTCTCAA CTTACACATC CAAGTATCAG CACACCCGTG AGAACGCCCT TTAATCTTA CCCAGATAAA	100
GGAGCTGTT TCAGGCCAA ACCCTCGCC CACACACCT AGCTCTACCA CACCCCTAGG GTAAATCAGC AGCTGAATTT TTTCGGTATC AGCGTCAGCT	200
TGACTCAGT AGAGAAAGA GGCGGCGTA ACCCGGTCG AGCGCGGCG GCTACAGCTT GGGGCTCGAG TTGATAGTCA TCGGGCTTAA CGCTGATTAA	300
AGATATGTTT AAGATTTATC TAACTGTTT ATGCTGTTT AAACCTAT CACACATC AGCTGTCCT AATTATTTAT CTGAAATACA	400
CGGGCTGCTG GAAGAACAC TGGATGAA TACGGGAA ACCCTGGCCG AACAGACCTT ATTAGGACCA ATTAGGCGCA GGAGGTTACG AGCAATGCTT	500
AAACCCCAA CGATTTGAG CGCTCCACC CGCTAGAGC AGCTGGTCTT ATTAACGATC ATTACGCGATA TACCGCCCA TTTCGCGGCT ATCAGCTGTT	600
ATACGCTTCA CGAACCTTA CGATGCAACG GCTTGGAGA CCTCTTAAAGA OCTAACACCTT CAACTACCA CGCTGACGTC TCGGCTTAA GAAATGAGT	700
ATACGCTTCA ATTCTTAAAT CTAGACAAA CGAACAGCA TATCAATTA TCTCATGA GTGGATTAA TTGATTTAA AAAATAGAG TGTTCTTTT	800
AACCGCGCTC TTGGACCGCT ACACACCCC CCTCACCGT TTGGATAGT TCTCACCGG TTCTCAACCC ACTTACATC TTGAGAGC CGAAGCTGTA	900
ACATGCTAAC TCTACTGAA ACTGCACTT GTTTATA	937

B

CCCCAAATC TAGCTACAC ACATCCCTAC GACCCCTTA CCCCTAGCCT CTTAACAAAT CATTAAACCA TTATAGTACA GGGGATCGAA AAATTTCTAA	100
GGCCCTCAGA CAAACTACCG CAAGGGAAAG ATGAAATAGA AATGAAATAA TTCTAAAGCC CAACTACCA GAGATTAACCT CTGGTACCTT TTGATCATGAT	200
GTCTAGCCAG TCTACTACAG CAAATTAAGA CTTTCCTTAACTTCCCGGAA ACCTAACATC AGCTCTGCTT GAAACCTCTT ATGGGACCAA CCCATCTCTG	300
CTTGCCTAAAGA GTGCGGAGAT CTCTTAACTG AGGTGTAAG CCTAACCGAC TTAGAGATAG CTGGTCTTGC AGGAAAGAC TTCTAGCTCT ACCTTAACT	400
CTCTCTTATA AACTAGAAA TCCCAACAGC TAAGAGCTAT TCAAAATAGG CACAGCTTAT TTGAAACAGG AAACAACTTC TAACACCGGG TGAAATTATAG	500
TAATCTCCTAT AAAGTGGGCC TAAACCCAGC CACCTTCTAA AACCCCTTAA AGCTTAACTA TAAATTACTA ATAAATCTCA AAATATTAAAT TAAACCTCTA	600
TTCCCTACTGA ACTTATTTAT ATCCCTATAT AACCAATTAT GTGAGACTA GTAACAGAA ATTGATTTT TTCTAAATCT AGTATAAAC CAAATGAC	700
CATCTCTGG TTGTTAACCC AAATCCAAA TCTATACCAA CATAACTAGA AACCCCTATA ACTACAAAGC TTAAACCTTAC ACTAGAACAT TCCAGAACG	800
ATTTAAACAG AAAGAGGAG CTCCGGCAAATT TTTCGGCTTCG CCTTGTTCACG CTCCTGATC AAACATAGA GTTCGACGCC GCCGAGTGC	900
AAAGTTCTACG GGGCCCGCTA CCTCTACCG CGGAAGCTAG CATATCTACT GTCTCTTAAAT AGTAGGCGT GTATCACCGA CCAAGGAGG GCTATAGCT	1000
CTCTCTTCTACG TCTCTCTACG ACTCTCTCTC CGCCGGAAAG AGCGGGATT ATTAAATAGG AGGAGAGAC CGCATGGCC TTAAATCTCA TCATACCTT	1100
CTTATCTACG TATCTACG CACAGAGCTG TCTCTCTCTC GTTTAGGGT GGCGGCTCCA CGGAGATATA CAAACCTTCA ATAACAAATG GCTAACAC	1200
CTTATCTACG AGATACACCTT CAACTACAT GTTAACTGTT CAACTACAT GTCATGCGA TGTGGGATAAC CAGCGGATTC ATGTCGAC	1300
TACTCTACG CGCCCTATCG ACAGCTACT GTTACGACCTC GATGGTGGAT CAGCTATAG TACTGCTGCA CGCCCTACTA ATGTCGAC	1400
TTAAACCCCT ACCTGATCTG AGTCTGACCC CGGATTAATCC AGCTGCTTTT CTATCTATAA AGTGGTCTCC CTAGTACGAA AGGACCGGGG CRACATGAC	1500
AAACCCCAT ATTCTCTCAC TATGAAATA ATCTTAATTG CGACAGACTT ATTATAGTACT TCTAAATCG AACAT	1585

Fig.1 Nucleotide sequences of *Rana catesbeiana* mitochondrial 12S (A) and 16S (B) genes. The sequence of the L-strand (5' to 3') is shown.

References

1. Fujii, H. et al. (1988) J. Biochem. 103, 474-481.