

# Nucleotide sequences of nine tRNA genes from *Micrococcus luteus*

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The nucleotide sequences of nine tRNA genes located in the different *Pst*I DNA fragments from *Micrococcus luteus* were determined (MLTR003-MLTR011). Each tRNA gene sequence is preceded by a putative promoter and followed by a probable terminator. The 3'-terminal CCA is coded for 4 out of 9 tRNA genes examined.

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- (1) MLTR003 Ala(GGC)<sup>1</sup>  
 CGATTTTGGTGCCCGCGGGGACCGTTACGCTGGTTCGAGTTGCCAGCGAGGGGAACGGGCGGAAACGCGCGTGAACATCGCGGGTACGCCACATCT 100  
GGGGTATGGCGCAGTTGGTAGCGCTCTGCATGGCATGCAGAAGGTCAGGGGTCGAATCCCTTACCTCCACCGCACGACAAGGCTCGAACCTTGC 200  
 Ala
- (2) MLTR004 Ala(GGC)<sup>2</sup>  
 GGCCGGTTTGGCGGTTGTCGCGAGGCGGGTATCTCTGTTTCAGGTGCTCCGGCCGGTCCGCCGAAACGGAGTCACCACCCACCTGGGGTATGGCGCAGT 100  
TGGTAGCGCTCTGCATGGCATGCAGAAGGTCAGGGGTCGAATCCCTTACCTCCACCGTCAGAAGGCCGGTCCGTCGGGACCGTCTGTCGTCTGAC 200  
 Ala
- (3) MLTR005 Arg(ACG)  
 GAGCGTCGATTTGCACTTCGCCGAGAGGTCCTGTAGTGTCTACGAGTTCGAGCCGCTCCGGCGGATCGGATGATGCGCCCATAGCTCAGCTGGATAGA 100  
GCGTCTGTCTACGGAACAGAAGGTCAGGGGTCGAATCCCTTGGGGCGCACGACCGGAAGGCCCGCTCTGCTGGAGCGGGGCTTCGTCATGTCGGG 200  
 Arg
- (4) MLTR006 Leu(GAG)  
 GACCATGGTTTGGAGCTAGCCGAAACCCGTCATCTTCTCGAGTTGCCGGAGCGGGACGGACCGGACAGGACCGGAAACGCGGTGGGCGACGGAGCC 100  
 CCTCCAGGGTCCACCGATGAGCGCGAGTGGCGGAATTGGTAGACCGCAGCGTTGAGGTCGCTGTTCGCAAGAGCTGGGGGTTCAAGTCCCCCTC 200  
GCGCACCGCTCGTCGAGCAGCGGGCCCGGTCGTCACGACCGGGGCCCTGCGTGTCTCCCGTCCGGGGTCCCCTCTAGAGTGGTGTCCGTCGACGGCA 300  
 Leu
- (5) MLTR007 Lys(CUU)  
 AGATTTGCACGGCGTCTGAAACCCCTGGTAGGGTTTCACCTCGTTGCACGGCGCACCGGAGAGATCCGAGAGGGCGTGGCCACCTGCACCTCTAGCTCA 100  
ACCGGTAGAGCATCTGACTCTTAATCAGCAGGTTCCGGGTTCCAGTCCCGGGGGTGCACCACACGAAGGCCCGCTCTCCACGGAGACGGGGCTTCCTC 200  
 Lys
- (6) MLTR008 Pro(UGG)  
 GTCGCGGCTTTTCGGGGCGCGAGACGGCGCGCTAGGATTCACAGTCTGCCTCGGGCAGTCCCTCGGGTGTGGGCGAGCCTCCACCGGTGAGCGCCGG 100  
 CGTCGGGACGACGGCGGGGTGTAGCTCAGCTTGGTAGAGCGCGCTTTGGGAGCGTGAGGCCGACAGGTTCAAATCCTGTACCCCGACTCGACGTCAA 200  
 GACCATCCACCACCACCAACAGGAGTCATCCGTGGTCAAGTCCACCGCAGAGAACCTCAGCCCGACCCGCTCAAGCTGACCGTCGAGGCGCCGTTTCGAG 300  
 Pro
- (7) MLTR009 Pro(CGG)  
 GCATCCCGGTTTGGCGGCAACCGCTGCGGCTGGTATAGTTCGCTCTCTGTTGGAATCAGGTCGAGACCTCAAACGGGGTGTGGCGCAGCTTGGTAGCGCC 100  
CGTCTTCCGGGACGACGAGGCGCGAGGTTCAAATCCTGTACCCCGACCAAGAGAGCCCGCGAATCGGGGGGGCTCTTCTCGTCTCCGGGTGCTCA 200  
 Pro
- (8) MLTR010 Thr(CGU)  
 GGCCGCGCGGTGGAATCGGGGCGCGAAACCGGTATGCTCGTCAGCCGTGGCCGTTGACGGTCACGTGCCTCCTTAGCTCAGCTGGCCAGAGCAGCTCCC 100  
TCGTAAAGAGCAGGTCGCGCGGTTTCGAATCCGGCAGGGGGCTCCCCCGGACCCCGCCGCTGATCAGCGCGGCTCTCTCGTCTCCGGGCGGCCCTCCGG 200  
 Thr
- (9) MLTR011 Trp(CCA)  
 GCCCGTTCGAGTCCCGCCCGCGATGGCGTATCCTGGATCTCCGGTCCCGGTTACGGTTCGATCCCGGAGGGTTCGGCCGGCCCGTCCGTCGGAAGG 100  
GTAGTGGCGCAATTGGTAGCGCAGCGGTTCCAAAACCGCAGGTTGCAGGTTTCGAGTCTGCTGCTGCTCGATGCCGGGCGCCGTCGCGGCATCGG 200  
 Trp

Figure 1. Putative promoter and terminator sequences are shown by double underlines and arrows, respectively.

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