



**Sequence (insert only)**

As compared to GenBank U00220:

1 mismatch

9 gaps

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GGCACTATGGGGTTTTCTGTACCTGTGACCGCCTCGATCCTCCCTTTATCCAGCCCTCACTCCTTCTCTAGGCGCC
GGAATTCGCCACCATGGGTGGCAAGTGGTCAAACGTAGTGTGAGTGGATGGCCTGCTGTAAGAGAAAGAAT
GAGGCGAGCTGAGCCAGCAGCAGAGGGAGTGGGAGCAGTATCTCGAGACCTGGAAAACATGGAGCAATTACA
AGTAGCAATACAGCAGCTACCAATGCTGCTTGTGCCTGGCTAGAAGCACAAGAGGAGGAAGAAGTGGGTTTTCCA
GTCAGACCTCAGGTACCTTTAAGACCAATGACTTACAAGGCAGCAGTAGATCTTAGCCACTTTTTAAAAGAAAAGG
GGGACTGGAAGGGTTAATTTACTCCCAAAAAGACAAGATATCCTTGATCTGTGGGTCTACCACACACAAGGCT
ACTCCCTGATTGGCAGAATAACACACCAGGGCCAGGAATCAGATATCCACTGACCTTTGGGTGGTGCTTCAAAT
AGTACCAGTTGAGCCAGAGAAGGTAGAAGAGGCCAATGAAGGAGAGAACAACCTGCTTGCTACACCCTATGAGCC
AGCATGGGATGGATGACCCAGAGAAAGAAGTATTAGTGTGGAAGTTTGACAGCAAACCTAGCATTTCATCATATGG
CCCGAGAGCTGCATCCGGAGTACTACAAAGACTGCTGAGAATTCGTTAACTCGAGGATCCGGCTGTGGAATGTGT
    
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GTCAGTTAGGGTGTGGAAAGTCCCCAGGCTCCCCAGCAGGCAGAAGTATGCAAAGCATGCATCTCAATTAGTCAG  
CAACCAGGTGTGGAAAGTCCCCAGGCTCCCCAGCAGGCAGAAGTATGCAAAGCATGCATCTCAATTAGTCAGCAA  
CCATAGTCCCGCCCCTAACTCCGCCCATCCCGCCCCTAACTCCGCCCAGTTCCGCCCATCTCCGCCCATGGCTGAC  
TAATTTTTTTTATTTATGCAGAGGCCGAGGCCGCCTCGGCCTCTGAGCTATTCCAGAGTAGTGAGGAGGCTTTTTTG  
GAGGCCTAGGCTTTTGCAAAGCTTGGGCTGCAGGTCGAGGCGATCTGATCAGAGACAGGATGAGATCGTTCGC  
ATGATGACCAGATGGATTGCAC