

1. US\_2006\_0217311\_A1\_3
2. US\_7374758\_B2\_15
3. US\_9669069\_B2\_1

1 ATGGTCAGCTACTGGGACACCGGGGTCCTGCTGTGCGCGCTGCTCAGCTGTCTGCTTCTCACAGGATCTAGTTCCGGAAG 80  
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2 TGATACCGGTAGACCTTTTCGTAGAGATGTACAGTGAAATCCCCGAAATTATACACATGACTGAAGGAAGGGAGCTCGTCA 160  
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1 TTCCCTGCCGGGTTACGTCACCTAACATCACTGTTACTTTAAAAAAGTTTTCCACTTGACACTTTGATCCCTGATGGAAAA 240  
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1 CACCAGGACTGGCTGAATGGCAAGGAGTACAAGTGCAAGGTCTCCAACAAAGCCCTCCCAGCCCCCATCGAGAAAACCAT 1040  
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1 TCAGCCTGACCTGCCTGGTCAAAGGCTTCTATCCCAGCGACATCGCCGTGGAGTGGGAGAGCAATGGGCAGCCGGAGAAC 1200  
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