

- 1. US_2006_0217311_A1_3
- 2. US_7374758_B2_15
- 3. US_9669069_B2_1

1 ATGGTCAGCTACTGGGACACCGGGGTCCTGCTGTGCGCGCTGCTCAGCTGTCTGCTTCTCACAGGATCTAGTTCGGGAAG 80
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1 TGTCTCCGGGTAAATGA 1377
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