

Query		ATA-TA-GC-AAGGACT-A-A-CCCC-T-ATACCTTCTGCATAATGAATTAAGTAACTAGAAAAT	
<u>NC_001807</u>	1823	ATA-TA-GC-AAGGACT-A-A-CCCC-T-ATACCTTCTGCATAATGAATTAAGTAACTAGAAAAT	1874
<u>NT_009237</u>	9317826	ATA-CA-GC-AAGGACT-A-A-TCCC-T-GTACCTTCTGCATAATGAATTAAGTAACTAGAAAAT	9317775
<u>NT_006713</u>	30541074	ATA-TA-GC-AAGGACT-A-A-CCCC-T-GTACCTTCTGCATAATGAATTAAGTAACTAGAAAAT	30541023
<u>NT_005612</u>	2832075	ATA-CA-GC-AAGGACT-A-A-CCCC-T-GTACCTTTTGCATAATGAATTAAGTAACTAGAAAAT	2832024
<u>NT_007914</u>	2998897	AAA-TA-GC-AAGGA-T-AGA-CCCT-T-ATACCTTCTGCGTAATGAATTAAGTAACTAGAAAAT	2998948
<u>NT_008413</u>	33647858	A-A-TA-GC-AAAAA-T-AGA-CCCT-T-ATACCTTCTGCATAATGAATTAAGTAACTAGAAAAT	33647908
<u>NT_008413</u>	5082622	A-A-GA--C-AA-G-----CCCT-T-ATACCTTCTGCATAAAGTATTAAGTAGAAAAT	5082665
<u>NT_024862</u>	339493	A-AGTA-GC-AAGGACA-A-A-CCCC-T-ATACCTTCTGCATAATGAATTAAGTAACTAGAAAAT	339544
<u>NT_016354</u>	80934604	A-A-AA-GC-AAAGAC--A-AGCCCT-T-ATACCTTCTGCATAATGTATTAAGTAACTAGAAAAT	80934554
<u>NT_016354</u>	41768304	A-A-AA-GC-AAAGAC--A-AGCTCT-T-ATACCTTCCGCATAATGTATTAAGTAACTAGAAAGT	41768354
<u>NT_011630</u>	2763299	A-A-AA-GC-AAAGACA-A-A-CCCT-T-ATATCTTCTGCATAATGTATTAAGTAACTAGAAAAT	2763249
<u>NT_004836</u>	2866222	A-A-AA-GC-AAAGA-TAA-A-GTCT-T-TTACCTTCTGCATAATGTACTAGCTAGAAAAT	2866172
<u>NT_011362</u>	20987791	A-A-AA-GC-AAAGAC--A-AGCCCT-T-ATACCTTCTGCATAATGCATTAAGTAACTAGAAAAT	20987741
<u>NT_022135</u>	6487931	A-A-AA-TC-AAAGAC--A-A-GCCC-T-ATACCTTTTGCATAATGTATTAACCAGAAAAT	6487980
<u>NT_022135</u>	29688448	AAA-AA-GC-AAAGA-T-A-A-ACCC-TTATACCTTCTGCATAATGTATTAAGTAACTAGAAAAT	29688397
<u>NT_022135</u>	32555841	AAA-AA-GC-AAAGACC-A-A-CCCT-T-ATACCTTCTGCATAATGTATTAAGTAACTAGAAAAT	32555790
<u>NT_005403</u>	62849509	AAA-AA-GC-AAAGAC--A-AGCCCT-T-ATACCTTCTGCATAATCTATTAAGTAACTAGAAAAT	62849458
<u>NT_023678</u>	902627	AAA--A-GC-AAAGACA-A-A-CCCT-T-ATACCTTCTGCATAATGTATTAACCAGAAAAT	902577
<u>NT_022517</u>	40234046	AAA--A-GC-AAAGAC--A-A-GCCC-TTATACCTTCTGCATAATGTATTAAGTAACTAGAAAAT	40234096
<u>NT_007995</u>	3190931	A-A-AA-GCAAAGCA-A-A-A-CCCT-T-ATACCTTCTGCATAATGTATTAAGTAACTAGAAAAT	3190981
<u>NT_008583</u>	5910010	A-A-AA-GC-AAAGACA-A-A-TCCT-T-GCATCTTTTGAATAATGAATTAAGTAACTAG-AAT	5910059
<u>NT_033899</u>	6843028	AAA--A-GC-AAAGACA-A-A-CCCT-T-ATACCTTCTGCATAATGTATTAAGTAACTAGAAAAT	6842978
<u>NT_010718</u>	19100337	AAA--A-GC-AAAGACA-A-A-CCCT-T-ATACCTTCTGCATAA-----CTAGAAAAT	19100379
<u>NT_011786</u>	26730286	A-A-AAGGC-AAAG-CT-A-ATCCCT-T-GTACCTTTTGCATAATGAATCAACTAG-CAT	26730336
<u>NT_008046</u>	17314257	AAA-AA-AC-AAAG-CT-T-A-CCCCTT-ATGCCTTTTGCATAATGAATTAAGTAACTAG-AAC	17314307