

NZ_LS483400.1
 NZ_CP026524
 NZ_CP011312

CTGCTTCCACCTGTTCTGGGGAATGAGTGGGATGTGCGTATCAATAGCGCCAGCGGTGA 60
 -AGCTTCCACCTGTTCCGGGGAATAGTGGGATGTGCGTATCAATAGCGCCAGCGGTGA 59
 CTGCTTCCACCTGTTCTGGGGAATGAGTGGGATGTGCGTATCAATAGCGCCAGCGGTGA 60
10.....20.....30.....40.....50.....60

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AATCTTGCCTTCCGACCGATAACCTCAGTTCCACACCGACGATGATGCGCAATGTTGT 120
 AATCTTGCCTTCCGACCGATAACTTCCGTACCCACCGACGATGATGCGCAATGTTGT 119
 AATCTTGCCTTCCGACCGATAACCTCAGTTCCACACCGACGATGATGCGCAATGTTGT 120
 0.....70.....80.....90.....100.....110.....120

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CCATAAATCGGGGTTCGGGCTTACCGAATCTTCAATGACCTGGCGCTTTTCAACAGCA 180
 CCATAAATCGGGGTTCGGGCTTACCGAATCTTCAATGACCTGGCGCTTTTCAACAGCA 179
 CCATAAATCGGGGTTCGGGCTTACCGAATCTTCAATGACCTGGCGCTTTTCAACAGCA 180
 20.....130.....140.....150.....160.....170.....180

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TATCAGCAGAGTAATAACCGGTGATGTCACGATAAATGGTGTTCGGTGATAACGAGGCTCG 240
 TATCAGCAGAGTAATAACCGGTGATGTCACGATAAATGGTGTTCGGTGATAACGAGGCTCG 239
 TATCAGCAGAGTAATAACCGGTGATGTCACGATAAATGGTGTTCGGTGATAACGAGGCTCG 240
 80.....190.....200.....210.....220.....230.....240

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GGACCTGATCTGCTCGACGCGCGGATTTTGACCCATGCCACCGCGG 287
 GGACCTGATCTGCTCGACGCGCGGATTTTGACCCATGCCACCGCGG 286
 GGACCTGATCTGCTCGACGCGCGGATTTTGACCCATGCCACCGCGG 286
 40.....250.....260.....270.....280.....