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>Alt
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>Ref 9:45172835-45173275 reversecomplement
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AACTTAAACACCCCTGCTCGGATAAAAATGAACCATGTCAGCCAGCATTTGGCTTTCAGCACGTGCTTCCACTC
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AAGCCTGGATGCCATCATGCAAGCTGCCGTCTGCCTGGTGAGACACATTTTTGACAGACATTTTAACAGGTTAC
ATTATCCATATTATGTGCTTAAGAGTTGTTTATAAATCATGTAAACAATGTTTTTGTGCTTCGTTGTACACAAT
GCAAAGAAACAATAGTTTACAGTGGGGGAAGTAAGTATTGAAAATGTCACCATTTTTCTCAGAAAACATTT


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Alt          1 AACCTTTCCTAATACAGGATTCGGTTCGTTTGTTGACAAAACACTTCTTCCATACACCGA
               |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Ref   45173275 AACCTTTCCTAATACAGGATTCGGTTCGTTTGTTGACAAAACACTTCTTCCATACACCGA

Alt         61 CACCAATGAAGCCAAACTTAAACACCCCTGCTCGGATAAAAATGAACCATGTCAGCCAGC
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Ref   45173215 CACCAATGAAGCCAAACTTAAACACCCCTGCTCGGATAAAAATGAACCATGTCAGCCAGC

Alt        121 ATTTGGCTTTCAGCACGTGCTTCCACTCACAGCAAATGGGAACGACTTCAAAGAATTGGT
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Ref   45173155 ATTTGGCTTTCAGCACGTGCTTCCACTCACAGCAAATGGGAACGACTTCAAAGAATTGGT

Alt        181 AAAGCAGCAGCATATATCTGGAAACCTGGACCCTCCAGAGGGAAGCCTGGATGCCATCAT
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Ref   45173095 AAAGCAGCAGCATATATCTGGAAACCTGGACCCTCCAGAGGGAAGCCTGGATGCCATCAT

Alt        241 GCAAGCTGCCGTCTGCCTGGCGTGACACATTTTTGACAGACATTTTAACAGGTTACATTA
               ||||||||||||||||||||| |||||||||||||||||||||||||||||||||||||||
Ref   45173035 GCAAGCTGCCGTCTGCCTGGTGAGACACATTTTTGACAGACATTTTAACAGGTTACATTA

Alt        301 TCCATACTATGTGCTTAAGAGTTGTTTATAAATCATGTAAACAATGTTTTTGTGCTTCAT
               |||||| ||||||||||||||||||||||||||||||||||||||||||||||||||| |
Ref   45172975 TCCATATTATGTGCTTAAGAGTTGTTTATAAATCATGTAAACAATGTTTTTGTGCTTCGT

Alt        361 TATACACAATGCAAAGAAACAATAGTTTACAATGGGGGAAGTAAGTATTGAAAATGTCAC
               | |||||||||||||||||||||||||||||||| |||||||||||||||||||||||||||
Ref   45172915 TGTACACAATGCAAAGAAACAATAGTTTACAGTGGGGGAAGTAAGTATTGAAAATGTCAC

Alt        421 CATTTTTATCAGAAAACATTT
               ||||||| |||||||||||||
Ref   45172855 CATTTTTCTCAGAAAACATTT


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>Alt
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AAGG


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Alt          1 AACCTTTCCTAATACAGGTTTCGGTTCGTTTGTTGACAAAACCCTTCTTCCATACACCGA
               |||||||||||||||||||| ||||||||||||||||||||||||| |||||||||||||
Ref   45173275 AACCTTTCCTAATACAGGATTCGGTTCGTTTGTTGACAAAACACTTCTTCCATACACCGA


Alt         61 CACCAATGAAGCCAAACTTAAACACCCCTGCTCGGATAAAAATGAACCATGTCAGCCAGC
               ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Ref   45173215 CACCAATGAAGCCAAACTTAAACACCCCTGCTCGGATAAAAATGAACCATGTCAGCCAGC


Alt        121 ATTTGGCTTTCAGCACGTGCTTCCACTCACAGCAAATGGTAACGACTTCAAAGAATTGGT
               |||||||||||||||||||||||||||||||||||||||| |||||||||||||||||||
Ref   45173155 ATTTGGCTTTCAGCACGTGCTTCCACTCACAGCAAATGGGAACGACTTCAAAGAATTGGT


Alt        181 AAAGCAGCAGCATATATCTGGAAACCTGAT-------GAGGGATGCCTGGATGCCATCAT
               |||||||||||||||||||||||||||||         |||||| |||||||||||||||
Ref   45173095 AAAGCAGCAGCATATATCTGGAAACCTGGACCCTCCAGAGGGAAGCCTGGATGCCATCAT


Alt        234 GCAAGCTGCCGTCTGCCTGGTGAGACACATTTTTGACAGACATTTTAACAGGTTACATTA
               ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Ref   45173035 GCAAGCTGCCGTCTGCCTGGTGAGACACATTTTTGACAGACATTTTAACAGGTTACATTA


Alt        294 TCCATATTATGTGCTTAAGAGTAGTTTATAAATCATGTAAACAATGTTTTTGTGCTTCGT
               ||||||||||||||||||||||| ||||||||||||||||||||||||||||||||||||
Ref   45172975 TCCATATTATGTGCTTAAGAGTTGTTTATAAATCATGTAAACAATGTTTTTGTGCTTCGT


Alt        354 TGTACACAATGCAAAGAAACAATAGTTTACGGTGGGGGAAGTAAATATTGAAAATGTCAC
               |||||||||||||||||||||||||||||| |||||||||||||| ||||||||||||||
Ref   45172915 TGTACACAATGCAAAGAAACAATAGTTTACAGTGGGGGAAGTAAGTATTGAAAATGTCAC


Alt        414 CATTTTTCTCAGAAAACTTTTTTAAAGG
               ||||||||||||||||| ||||||||||
Ref   45172855 CATTTTTCTCAGAAAACATTTTTAAAGG


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