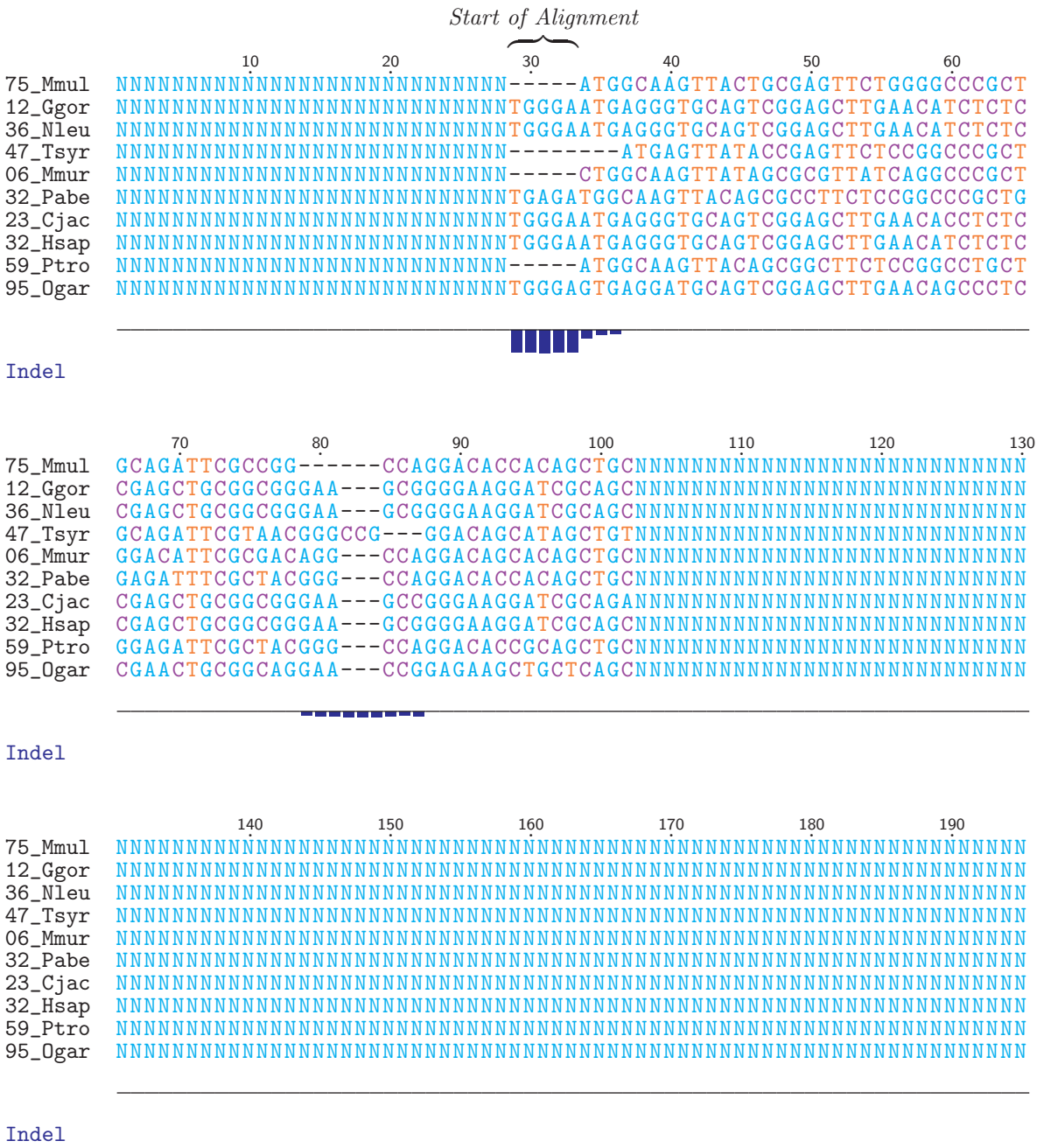
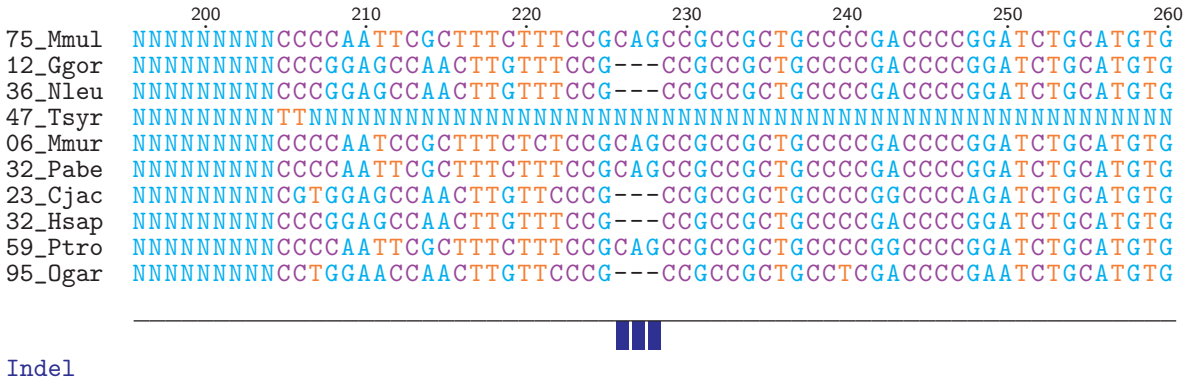
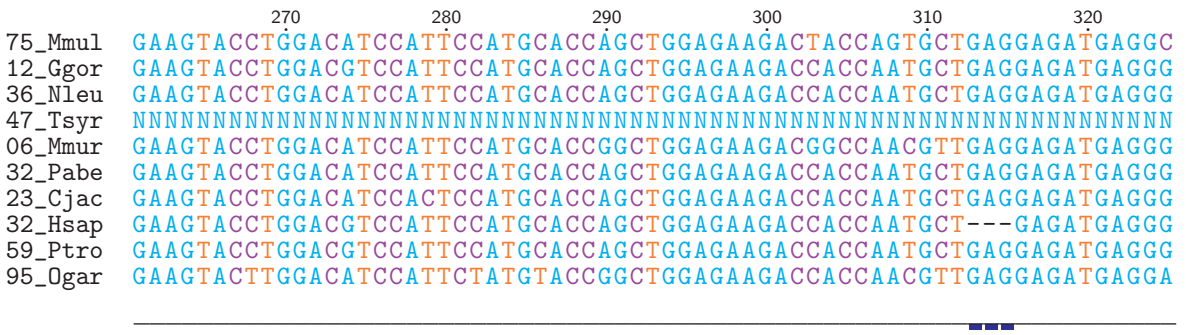
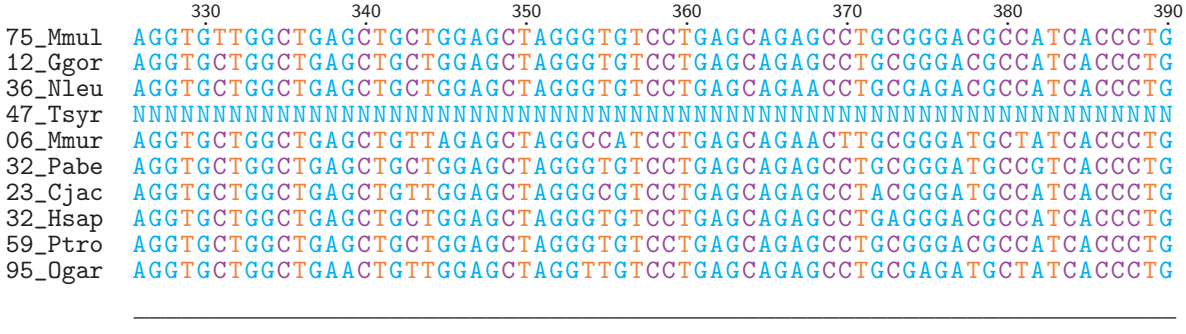

Start of Alignment
10
20
30
40
50
60
75_Mmul NNNNNNNNNNNNNNNNNNNNNNNNN-----ATGGCAAGTTACTGCGAGTTCTGGGGCCCGCT
12_Ggor NNNNNNNNNNNNNNNNNNNNNNNNNNNNTGGGAATGAGGGTGCAGTCGGAGCTTGAACATCTCTC
36_Nleu NNNNNNNNNNNNNNNNNNNNNNNNNNNNTGGGAATGAGGGTGCAGTCGGAGCTTGAACATCTCTC
47_Tsyr NNNNNNNNNNNNNNNNNNNNNNNNNNNN--------ATGAGTTATACCGAGTTCTCCGGCCCGCT
06_Mmur NNNNNNNNNNNNNNNNNNNNNNNNNNNN-----CTGGCAAGTTATAGCGCGTTATCAGGCCCGCT
32_Pabe NNNNNNNNNNNNNNNNNNNNNNNNNNNNTGAGATGGCAAGTTACAGCGCCTTCTCCGGCCCGCTG
23_Cjac NNNNNNNNNNNNNNNNNNNNNNNNNNNNTGGGAATGAGGGTGCAGTCGGAGCTTGAACACCTCTC
32_Hsap NNNNNNNNNNNNNNNNNNNNNNNNNNNNTGGGAATGAGGGTGCAGTCGGAGCTTGAACATCTCTC
59_Ptro NNNNNNNNNNNNNNNNNNNNNNNNNNNN-----ATGGCAAGTTACAGCGGCTTCTCCGGCCTGCT
95_Ogar NNNNNNNNNNNNNNNNNNNNNNNNNNNNTGGGAGTGAGGATGCAGTCGGAGCTTGAACAGCCCTC
Indel
70
80
90
100
110
120
130
75_Mmul GCAGATTCGCCGG------CCAGGACACCACAGCTGCNNNNNNNNNNNNNNNNNNNNNNNNNNN
12_Ggor CGAGCTGCGGCGGGAA---GCGGGGAAGGATCGCAGCNNNNNNNNNNNNNNNNNNNNNNNNNNN
36_Nleu CGAGCTGCGGCGGGAA---GCGGGGAAGGATCGCAGCNNNNNNNNNNNNNNNNNNNNNNNNNNN
47_Tsyr GCAGATTCGTAACGGGCCG---GGACAGCATAGCTGTNNNNNNNNNNNNNNNNNNNNNNNNNNN
06_Mmur GGACATTCGCGACAGG---CCAGGACAGCACAGCTGCNNNNNNNNNNNNNNNNNNNNNNNNNNN
32_Pabe GAGATTTCGCTACGGG---CCAGGACACCACAGCTGCNNNNNNNNNNNNNNNNNNNNNNNNNNN
23_Cjac CGAGCTGCGGCGGGAA---GCCGGGAAGGATCGCAGANNNNNNNNNNNNNNNNNNNNNNNNNNN
32_Hsap CGAGCTGCGGCGGGAA---GCGGGGAAGGATCGCAGCNNNNNNNNNNNNNNNNNNNNNNNNNNN
59_Ptro GGAGATTCGCTACGGG---CCAGGACACCGCAGCTGCNNNNNNNNNNNNNNNNNNNNNNNNNNN
95_Ogar CGAACTGCGGCAGGAA---CCGGAGAAGCTGCTCAGCNNNNNNNNNNNNNNNNNNNNNNNNNNN
Indel
140
150
160
170
180
190
75_Mmul NNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNN
12_Ggor NNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNN
36_Nleu NNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNN
47_Tsyr NNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNN
06_Mmur NNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNN
32_Pabe NNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNN
23_Cjac NNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNN
32_Hsap NNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNN
59_Ptro NNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNN
95_Ogar NNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNN
Indel



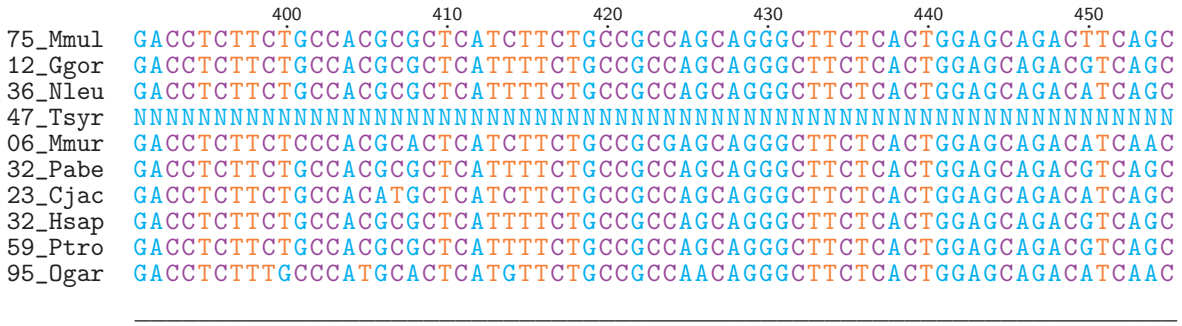
Indel



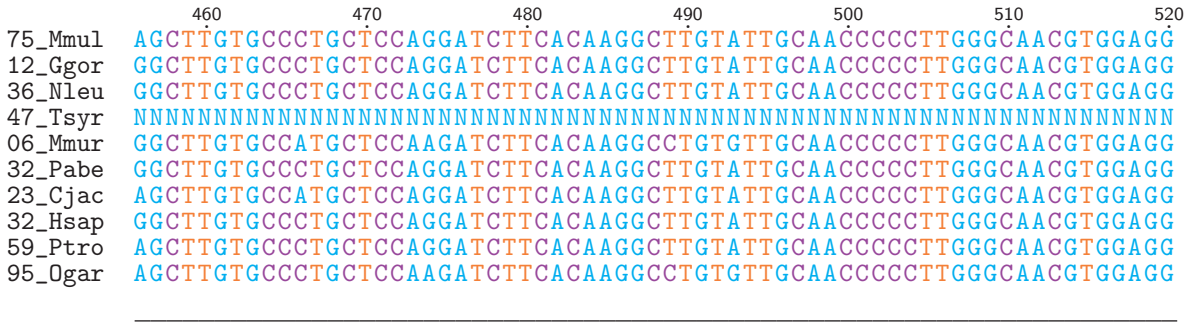
Indel



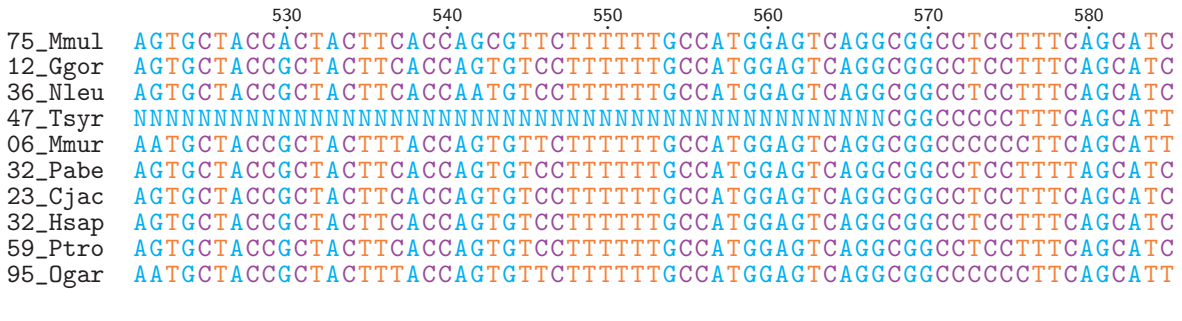
Indel



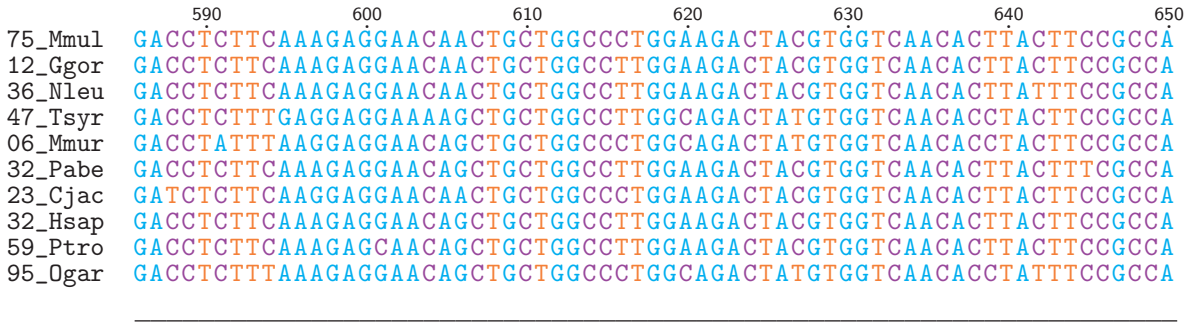
Indel



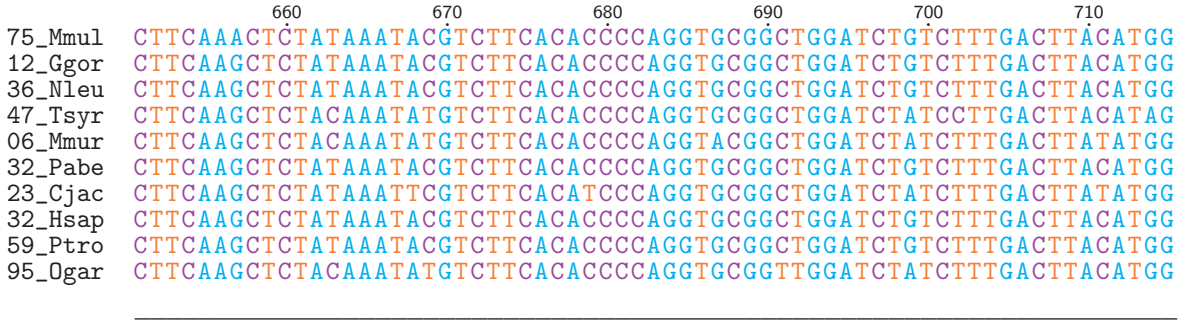
Indel



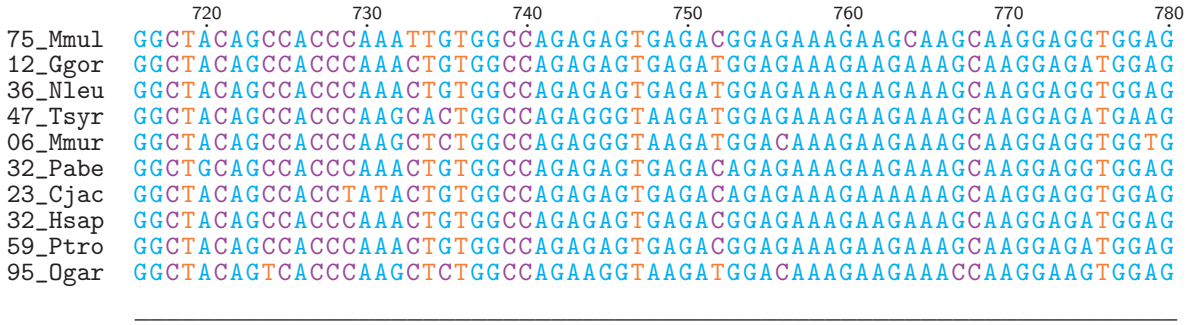
Indel



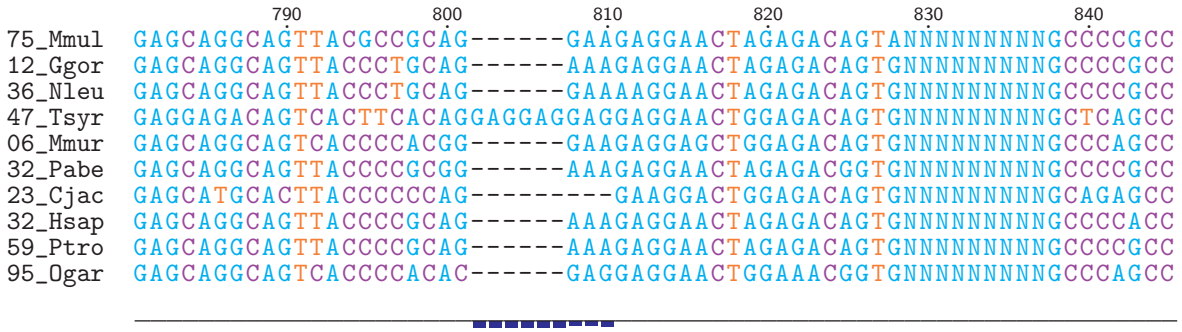
Indel



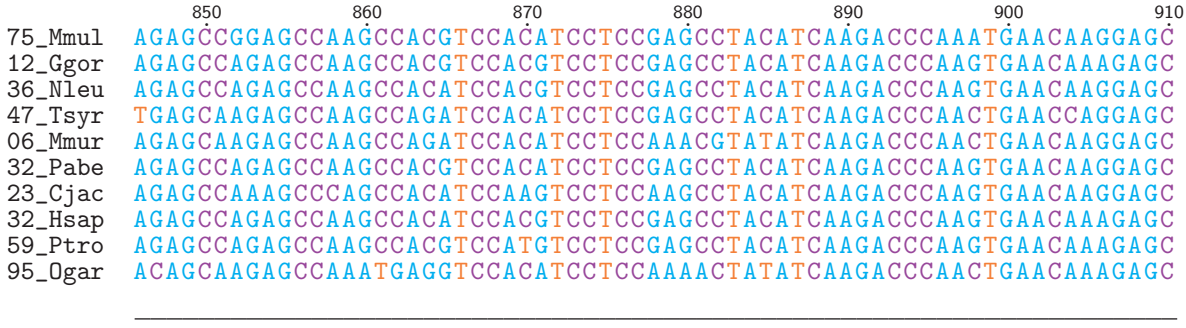
Indel



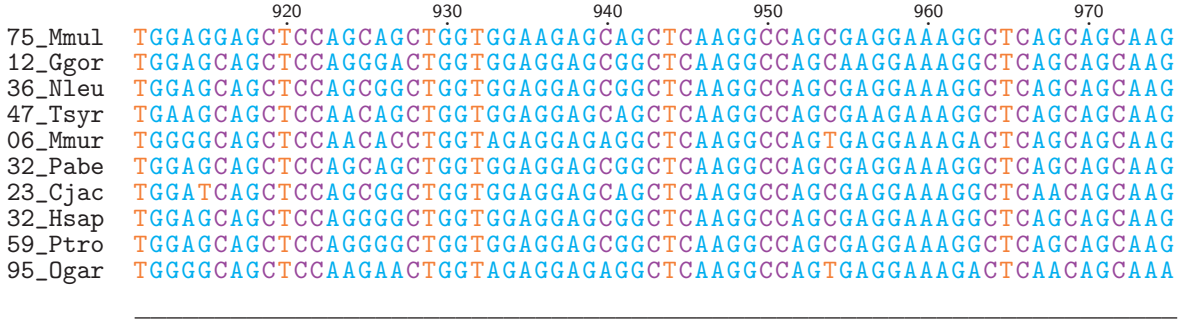
Indel



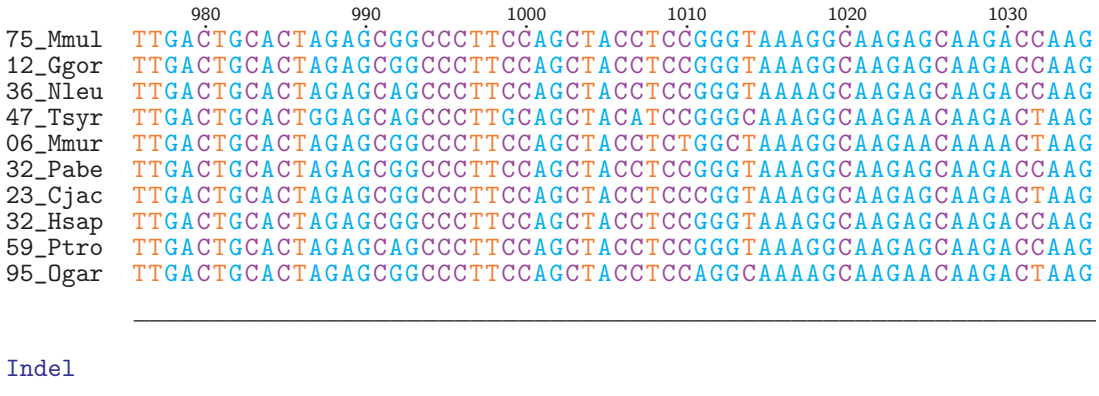
Indel



Indel



Indel


980
990
1000
1010
1020
1030
75_Mmul TTGACTGCACTAGAGCGGCCCTTCCAGCTACCTCCGGGTAAAGGCAAGAGCAAGACCAAG
12_Ggor TTGACTGCACTAGAGCGGCCCTTCCAGCTACCTCCGGGTAAAGGCAAGAGCAAGACCAAG
36_Nleu TTGACTGCACTAGAGCAGCCCTTCCAGCTACCTCCGGGTAAAAGCAAGAGCAAGACCAAG
47_Tsyr TTGACTGCACTGGAGCAGCCCTTGCAGCTACATCCGGGCAAAGGCAAGAACAAGACTAAG
06_Mmur TTGACTGCACTAGAGCGGCCCTTCCAGCTACCTCTGGCTAAAGGCAAGAACAAAACTAAG
32_Pabe TTGACTGCACTAGAGCGGCCCTTCCAGCTACCTCCGGGTAAAGGCAAGAGCAAGACCAAG
23_Cjac TTGACTGCACTAGAGCGGCCCTTCCAGCTACCTCCCGGTAAAGGCAAGAGCAAGACTAAG
32_Hsap TTGACTGCACTAGAGCGGCCCTTCCAGCTACCTCCGGGTAAAGGCAAGAGCAAGACCAAG
59_Ptro TTGACTGCACTAGAGCAGCCCTTCCAGCTACCTCCGGGTAAAGGCAAGAGCAAGACCAAG
95_Ogar TTGACTGCACTAGAGCGGCCCTTCCAGCTACCTCCAGGCAAAAGCAAGAACAAGACTAAG
Indel