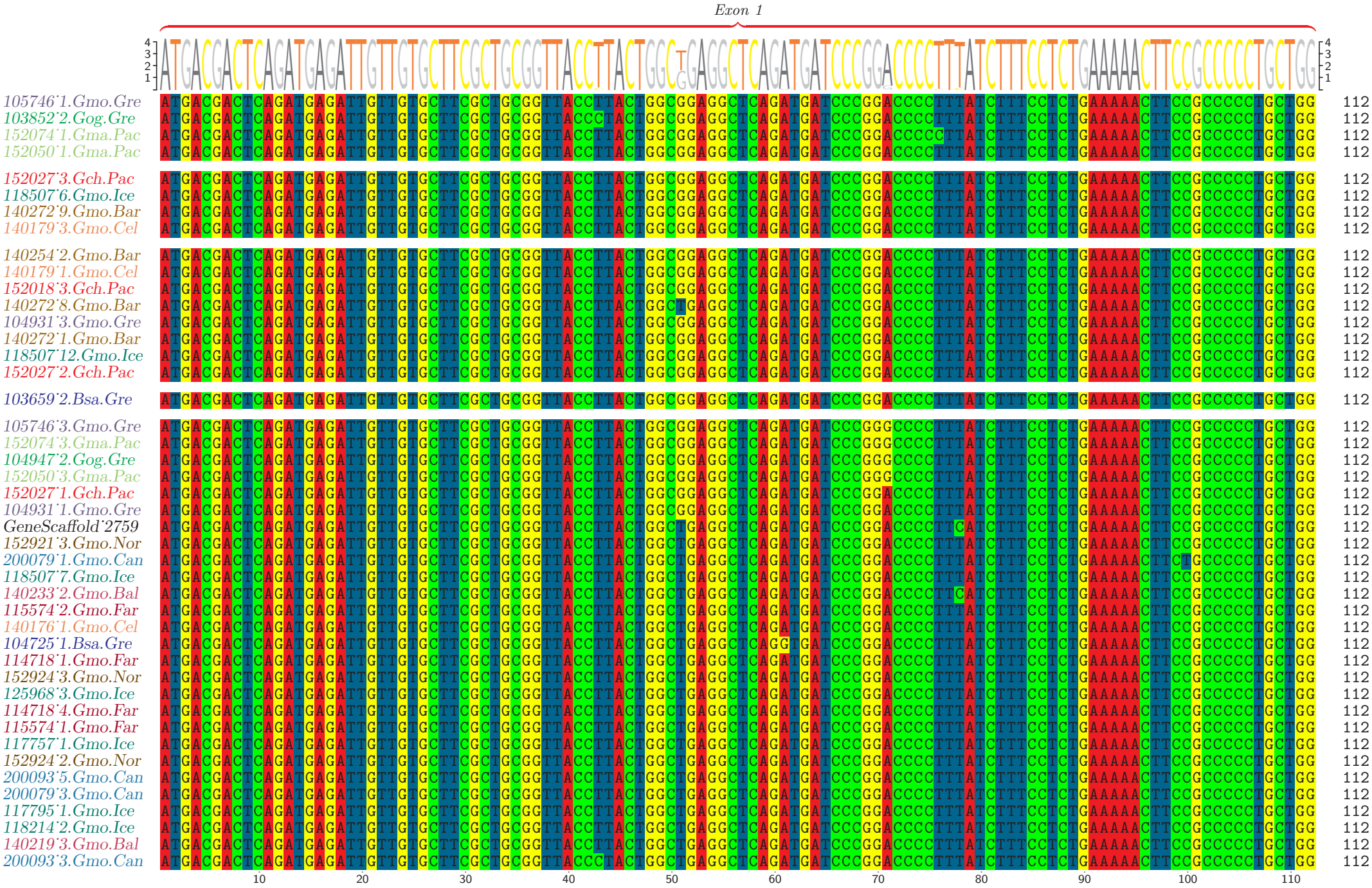
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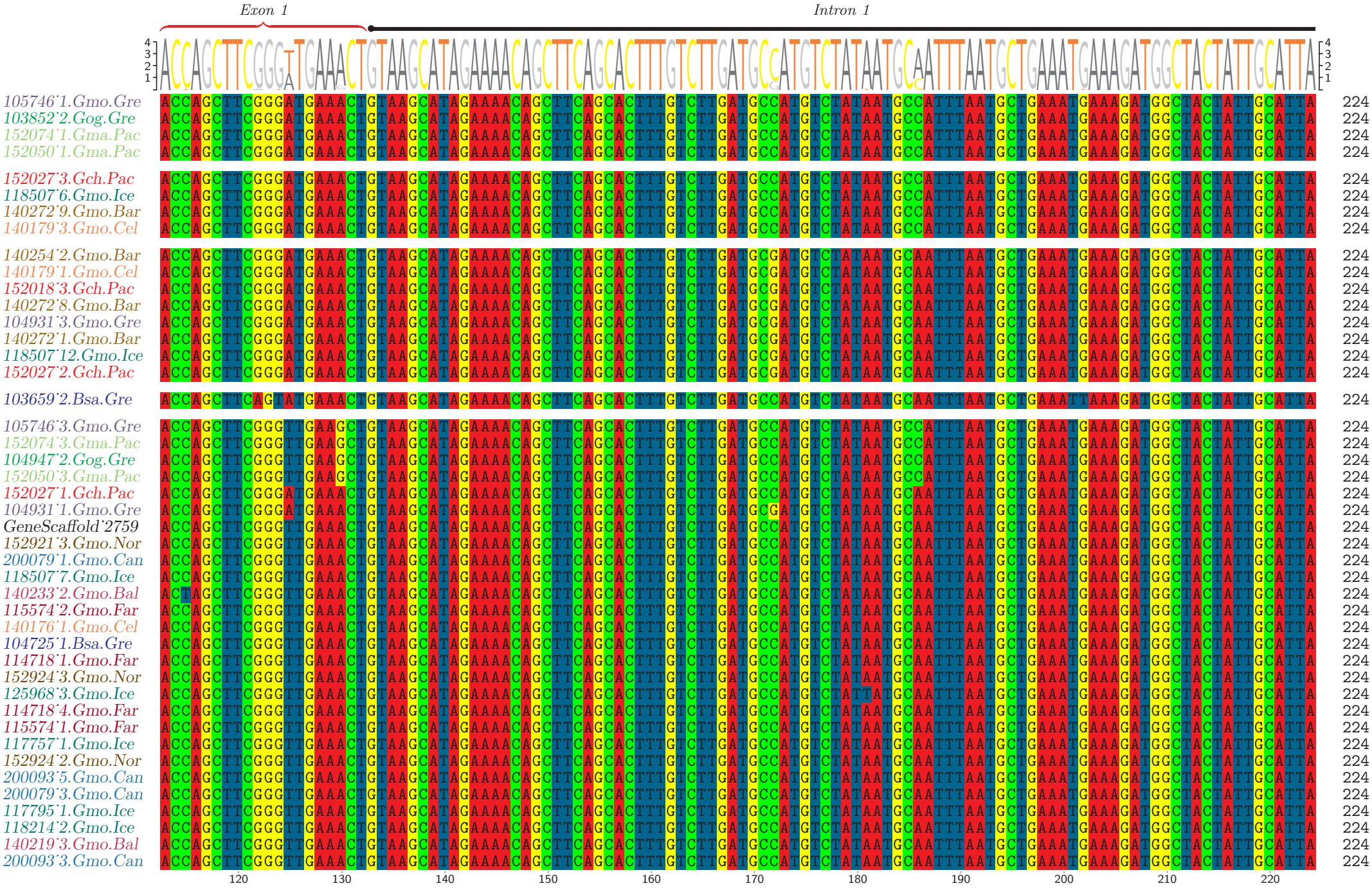
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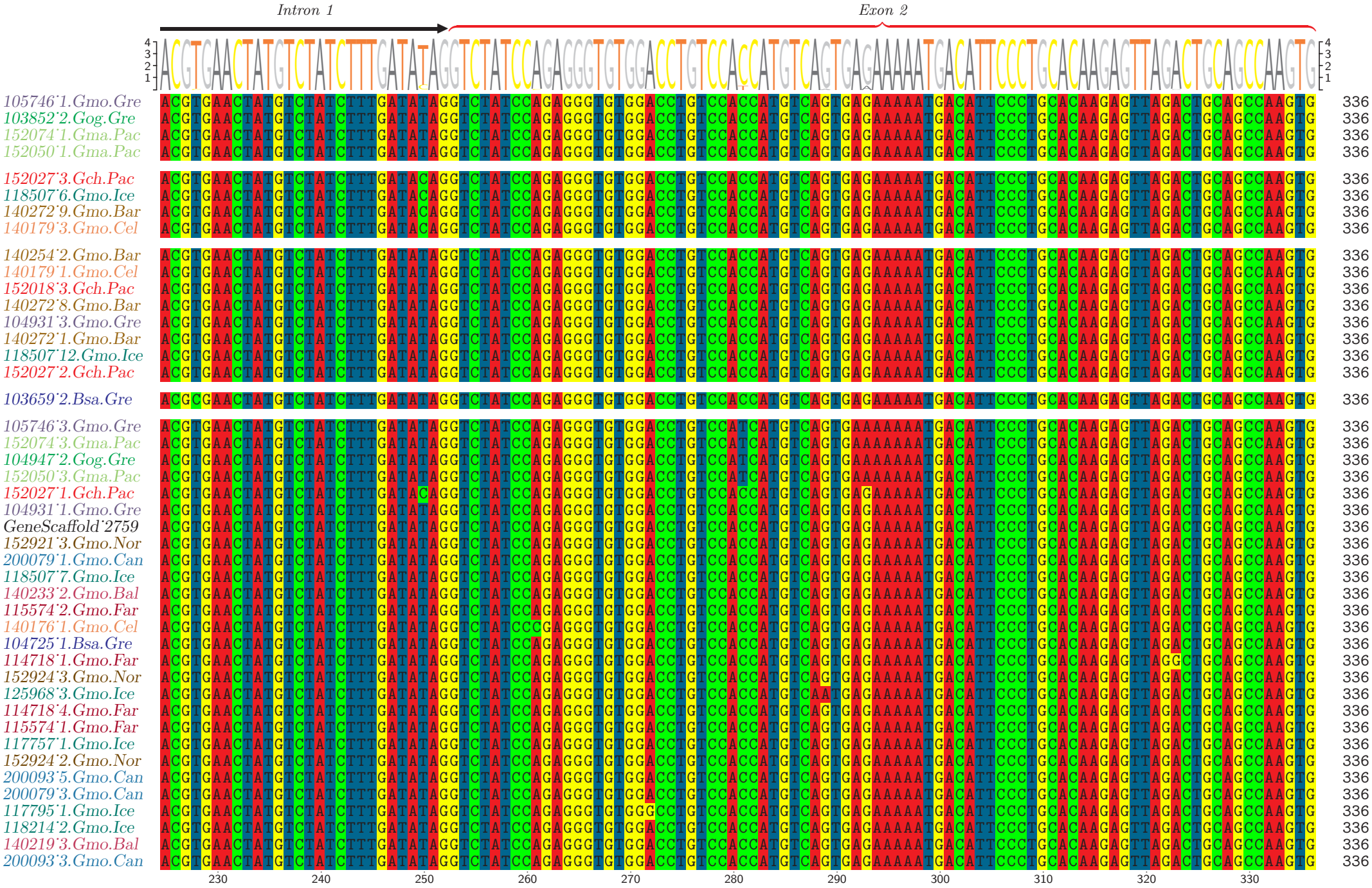
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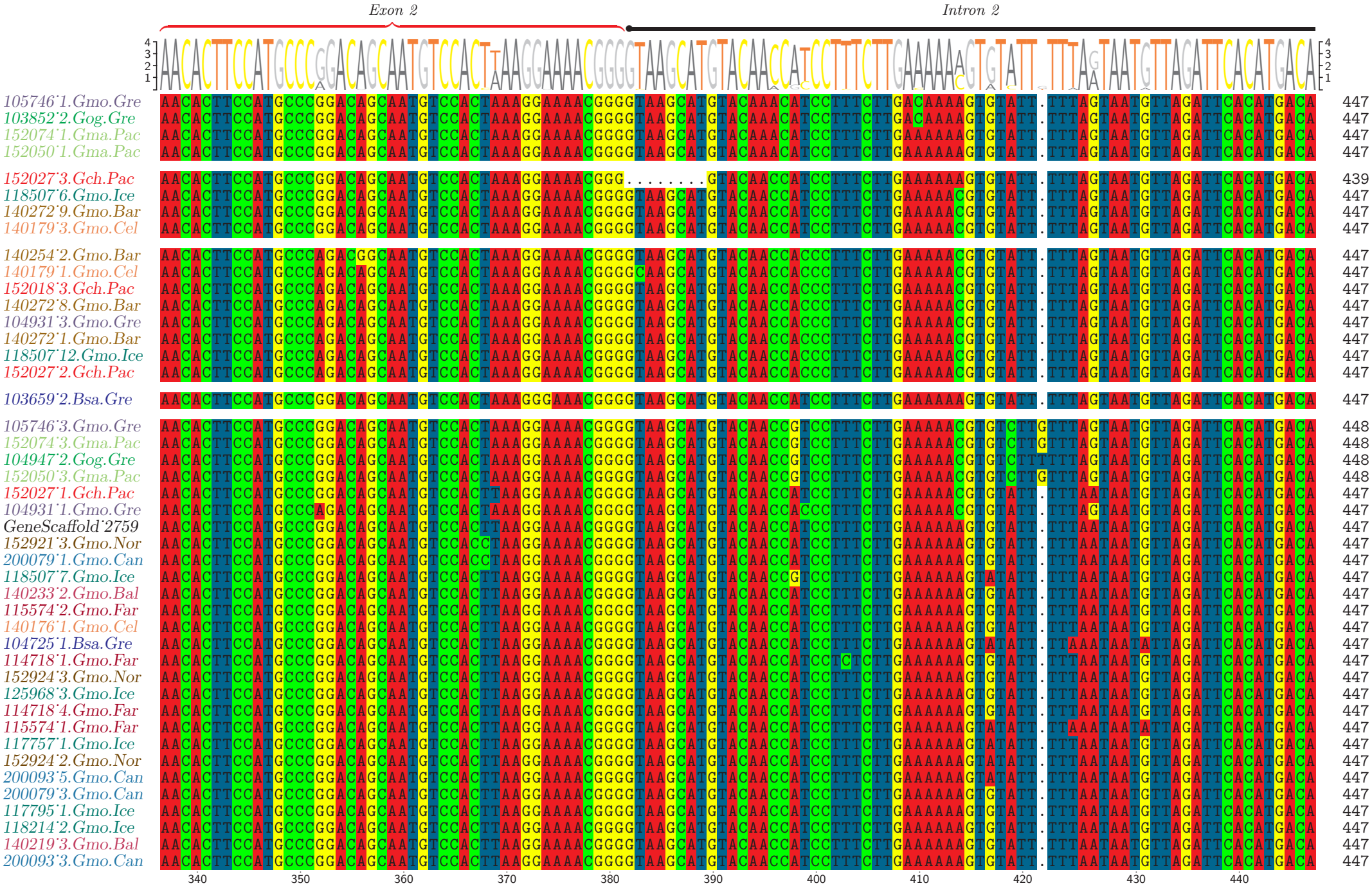
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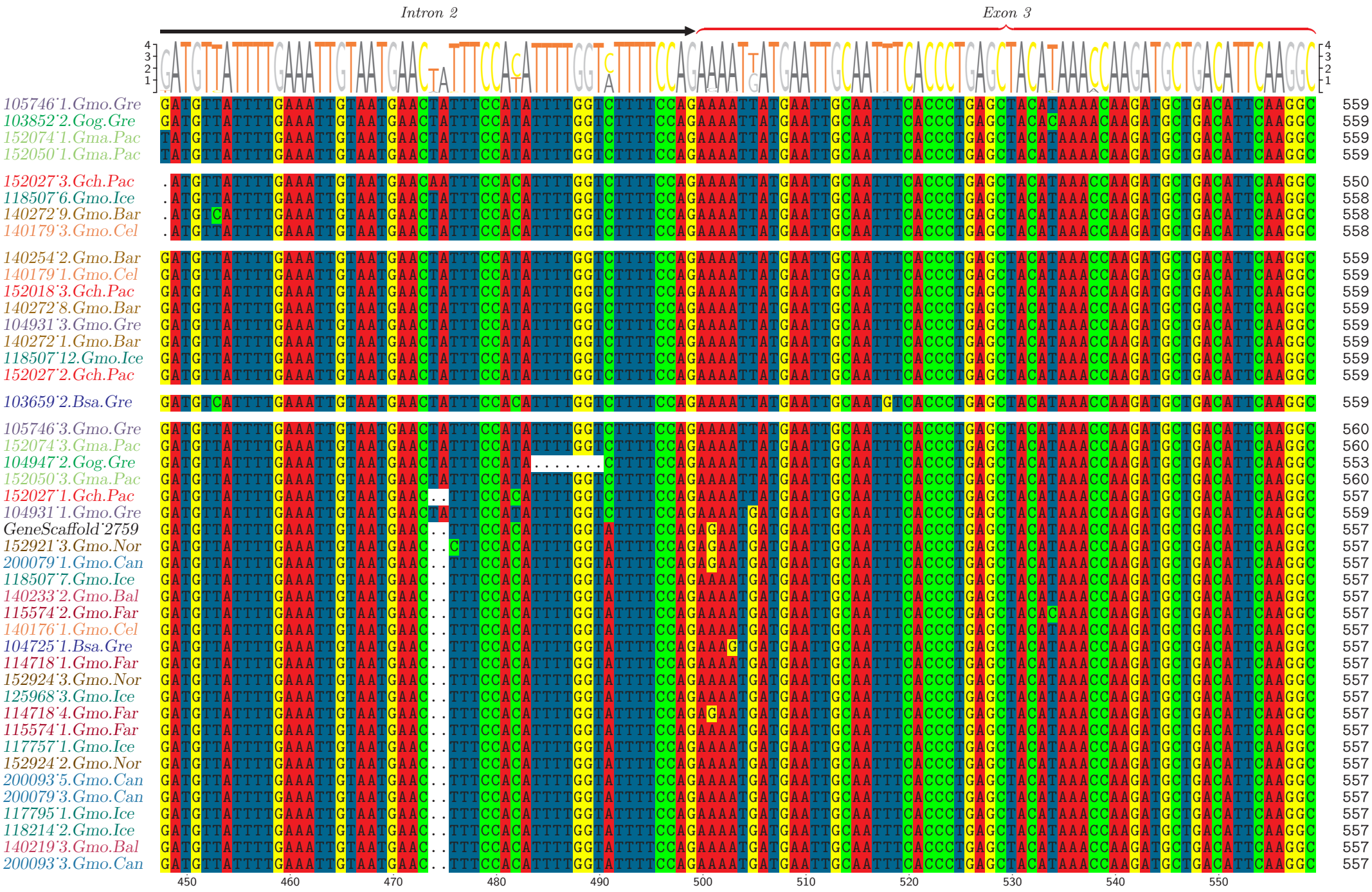
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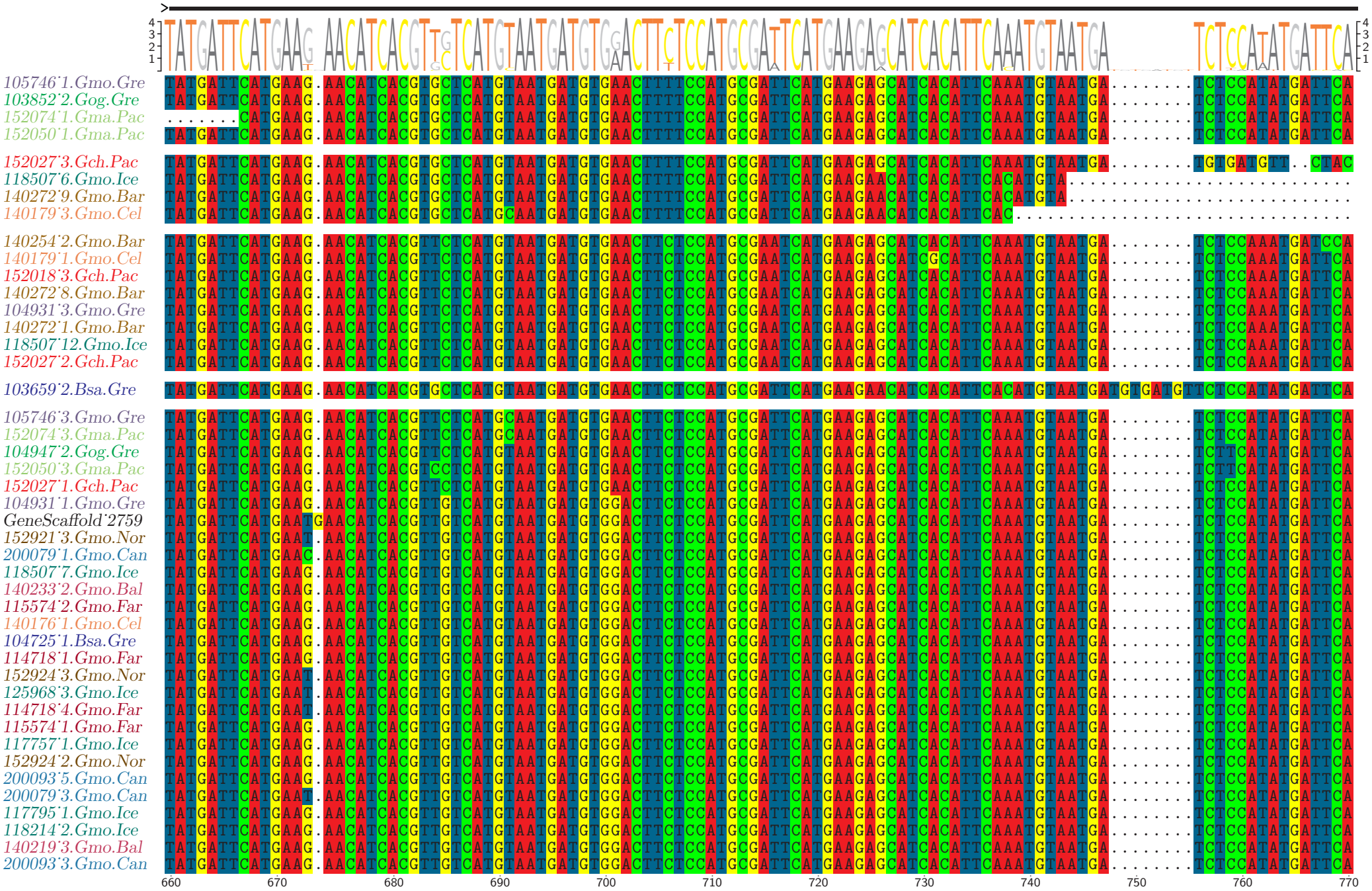
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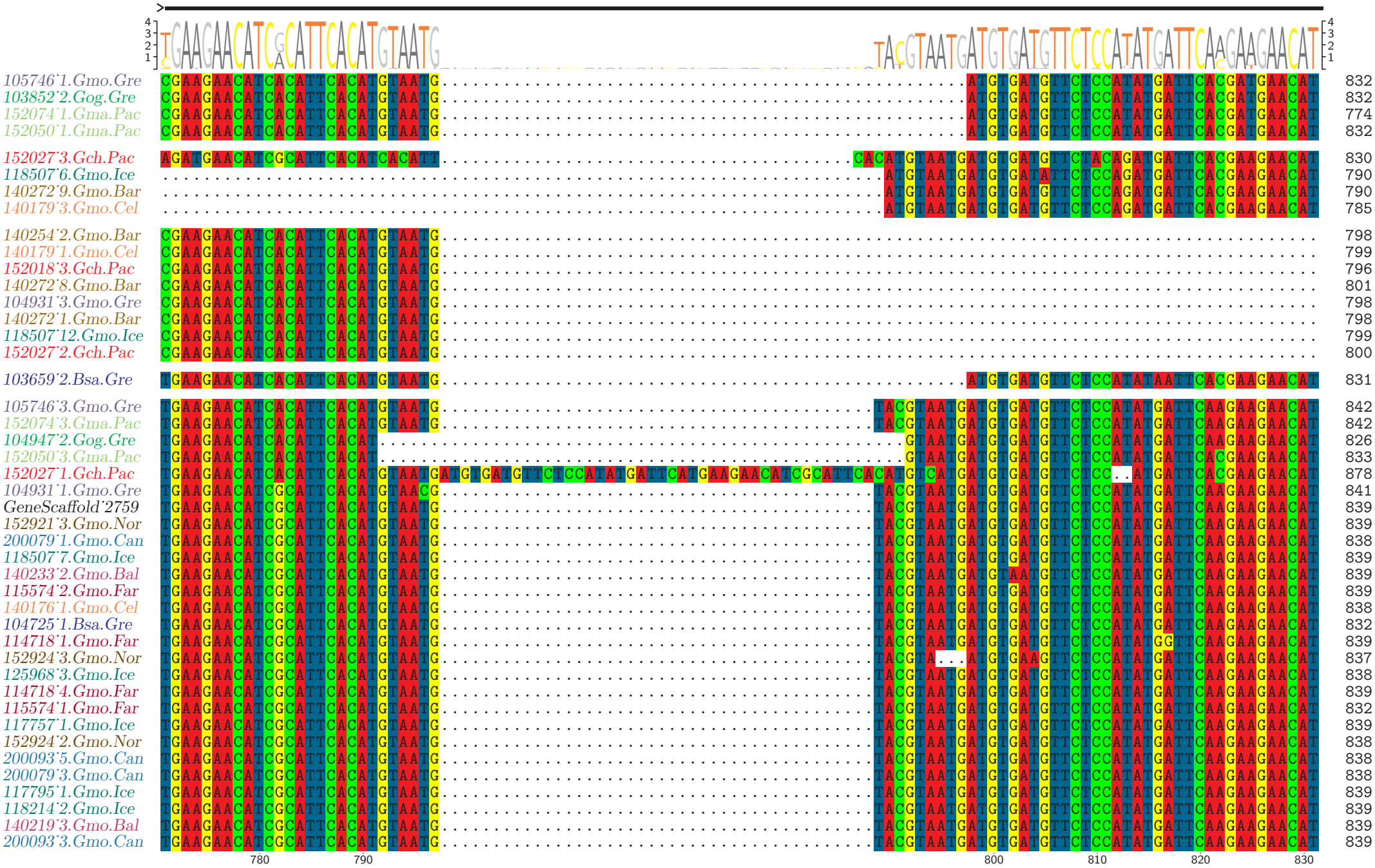
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Intron 3

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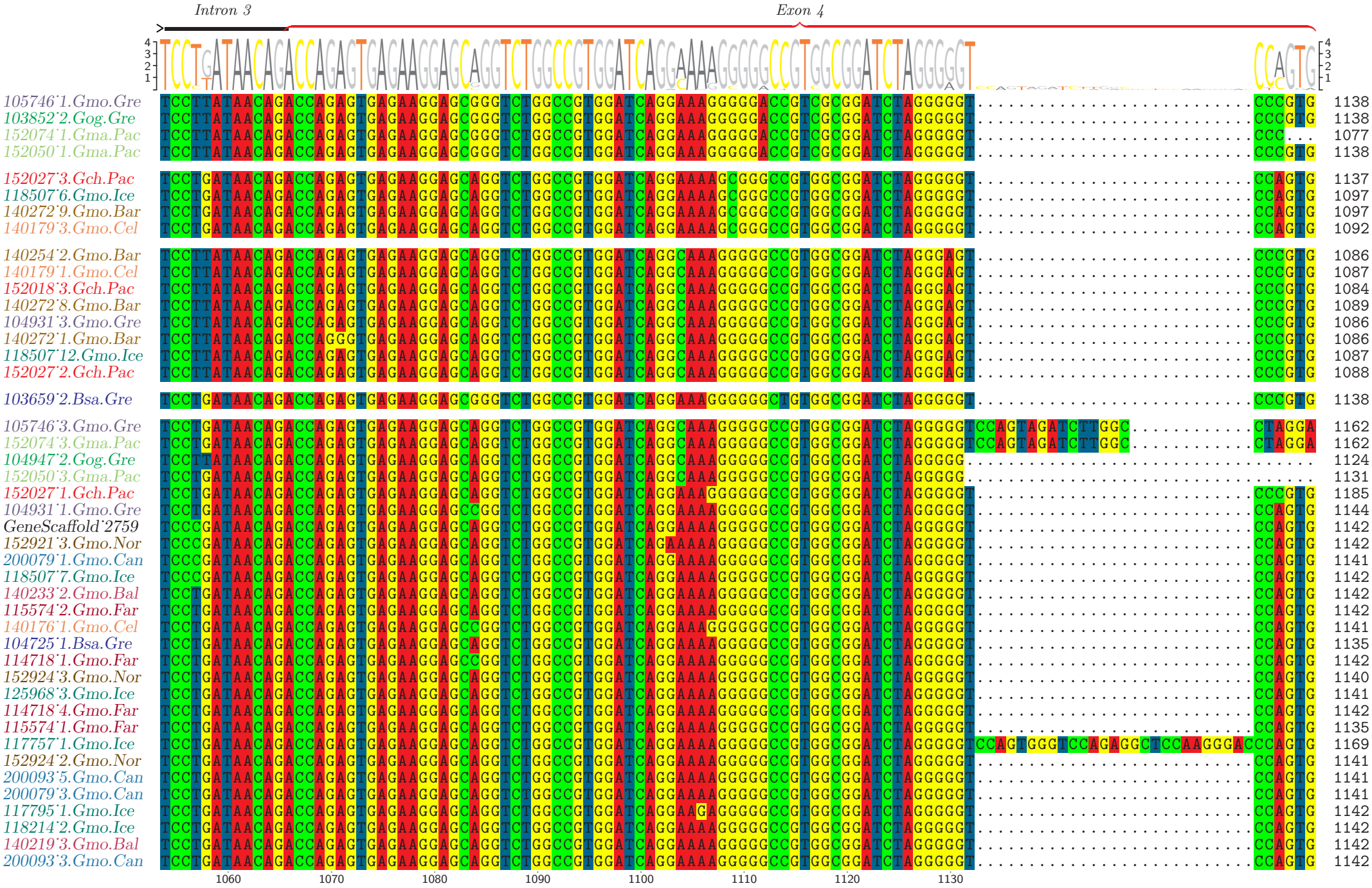
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118507˙6.Gmo.Ice
140272˙9.Gmo.Bar
140179˙3.Gmo.Cel
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Exon 4

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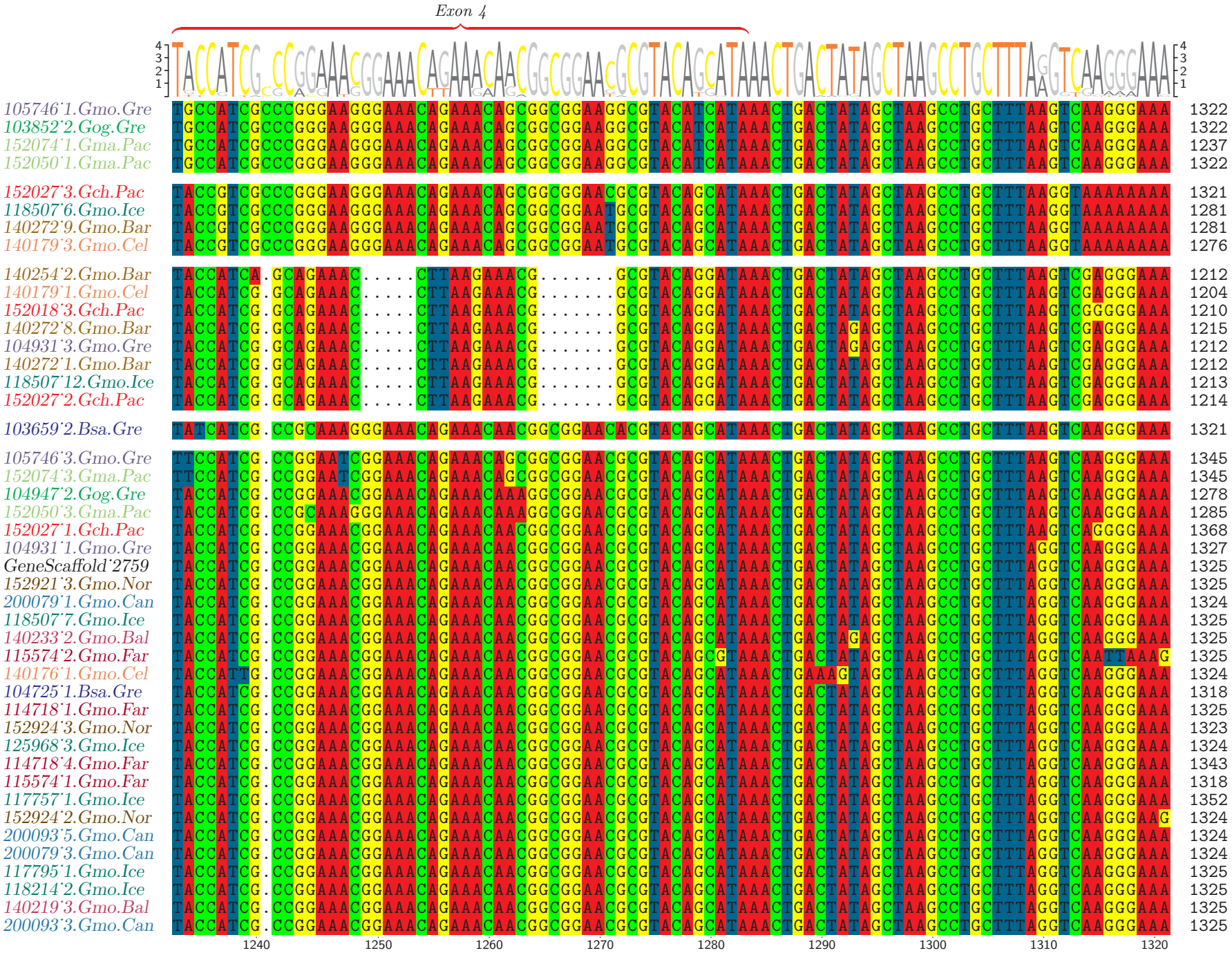
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140272˙1.Gmo.Bar TACCATCG.GCAGAAAC.....CTTAAGAAACG.....GCGTACAGGATAAACTGACTATAGCTAAGCCTGCTTTAAGTCGAGGGAAA 1212
118507˙12.Gmo.Ice TACCATCG.GCAGAAAC.....CTTAAGAAACG.....GCGTACAGGATAAACTGACTATAGCTAAGCCTGCTTTAAGTCGAGGGAAA 1213
152027˙2.Gch.Pac TACCATCG.GCAGAAAC.....CTTAAGAAACG.....GCGTACAGGATAAACTGACTATAGCTAAGCCTGCTTTAAGTCGAGGGAAA 1214
103659˙2.Bsa.Gre TATCATCG.CCGCAAAGGGAAACAGAAACAACGGCGGAACACGTACAGCATAAACTGACTATAGCTAAGCCTGCTTTAAGTCAAGGGAAA 1321
105746˙3.Gmo.Gre TTCCATCG.CCGGAATCGGAAACAGAAACAGCGGCGGAACGCGTACAGCATAAACTGACTATAGCTAAGCCTGCTTTAAGTCAAGGGAAA 1345
152074˙3.Gma.Pac TTCCATCG.CCGGAATCGGAAACAGAAACAGCGGCGGAACGCGTACAGCATAAACTGACTATAGCTAAGCCTGCTTTAAGTCAAGGGAAA 1345
104947˙2.Gog.Gre TACCATCG.CCGGAAACGGAAACAGAAACAAAGGCGGAACGCGTACAGCATAAACTGACTATAGCTAAGCCTGCTTTAAGTCAAGGGAAA 1278
152050˙3.Gma.Pac TACCATCG.CCGCAAAGGGAAACAGAAACAAAGGCGGAACGCGTACAGCATAAACTGACTATAGCTAAGCCTGCTTTAAGTCAAGGGAAA 1285
152027˙1.Gch.Pac TACCATCG.CCGGAAACGGAAACAGAAACAACGGCGGAACGCGTACAGCATAAACTGACTATAGCTAAGCCTGCTTTAAGTCAGGGGAAA 1368
104931˙1.Gmo.Gre TACCATCG.CCGGAAACGGAAACAGAAACAACGGCGGAACGCGTACAGCATAAACTGACTATAGCTAAGCCTGCTTTAGGTCAAGGGAAA 1327
GeneScaffold˙2759 TACCATCG.CCGGAAACGGAAACAGAAACAACGGCGGAACGCGTACAGCATAAACTGACTATAGCTAAGCCTGCTTTAGGTCAAGGGAAA 1325
152921˙3.Gmo.Nor TACCATCG.CCGGAAACGGAAACAGAAACAACGGCGGAACGCGTACAGCATAAACTGACTATAGCTAAGCCTGCTTTAGGTCAAGGGAAA 1325
200079˙1.Gmo.Can TACCATCG.CCGGAAACGGAAACAGAAACAACGGCGGAACGCGTACAGCATAAACTGACTATAGCTAAGCCTGCTTTAGGTCAAGGGAAA 1324
118507˙7.Gmo.Ice TACCATCG.CCGGAAACGGAAACAGAAACAACGGCGGAACGCGTACAGCATAAACTGACTATAGCTAAGCCTGCTTTAGGTCAAGGGAAA 1325
140233˙2.Gmo.Bal TACCATCG.CCGGAAACGGAAACAGAAACAACGGCGGAACGCGTACAGCATAAACTGACTAGAGCTAAGCCTGCTTTAGGTCAAGGGAAA 1325
115574˙2.Gmo.Far TACCATCG.CCGGAAACGGAAACAGAAACAACGGCGGAACGCGTACAGCGTAAACTGACTATAGCTAAGCCTGCTTTAGGTCAATTAAAG 1325
140176˙1.Gmo.Cel TACCATTG.CCGGAAACGGAAACAGAAACAACGGCGGAACGCGTACAGCATAAACTGAAAGTAGCTAAGCCTGCTTTAGGTCAAGGGAAA 1324
104725˙1.Bsa.Gre TACCATCG.CCGGAAACGGAAACAGAAACAACGGCGGAACGCGTACAGCATAAACTGACTATAGCTAAGCCTGCTTTAGGTCAAGGGAAA 1318
114718˙1.Gmo.Far TACCATCG.CCGGAAACGGAAACAGAAACAACGGCGGAACGCGTACAGCATAAACTGACTATAGCTAAGCCTGCTTTAGGTCAAGGGAAA 1325
152924˙3.Gmo.Nor TACCATCG.CCGGAAACGGAAACAGAAACAACGGCGGAACGCGTACAGCATAAACTGACTATAGCTAAGCCTGCTTTAGGTCAAGGGAAA 1323
125968˙3.Gmo.Ice TACCATCG.CCGGAAACGGAAACAGAAACAACGGCGGAACGCGTACAGCATAAACTGACTATAGCTAAGCCTGCTTTAGGTCAAGGGAAA 1324
114718˙4.Gmo.Far TACCATCG.CCGGAAACGGAAACAGAAACAACGGCGGAACGCGTACAGCATAAACTGACTATAGCTAAGCCTGCTTTAGGTCAAGGGAAA 1343
115574˙1.Gmo.Far TACCATCG.CCGGAAACGGAAACAGAAACAACGGCGGAACGCGTACAGCATAAACTGACTATAGCTAAGCCTGCTTTAGGTCAAGGGAAA 1318
117757˙1.Gmo.Ice TACCATCG.CCGGAAACGGAAACAGAAACAACGGCGGAACGCGTACAGCATAAACTGACTATAGCTAAGCCTGCTTTAGGTCAAGGGAAA 1352
152924˙2.Gmo.Nor TACCATCG.CCGGAAACGGAAACAGAAACAACGGCGGAACGCGTACAGCATAAACTGACTATAGCTAAGCCTGCTTTAGGTCAAGGGAAG 1324
200093˙5.Gmo.Can TACCATCG.CCGGAAACGGAAACAGAAACAACGGCGGAACGCGTACAGCATAAACTGACTATAGCTAAGCCTGCTTTAGGTCAAGGGAAA 1324
200079˙3.Gmo.Can TACCATCG.CCGGAAACGGAAACAGAAACAACGGCGGAACGCGTACAGCATAAACTGACTATAGCTAAGCCTGCTTTAGGTCAAGGGAAA 1324
117795˙1.Gmo.Ice TACCATCG.CCGGAAACGGAAACAGAAACAACGGCGGAACGCGTACAGCATAAACTGACTATAGCTAAGCCTGCTTTAGGTCAAGGGAAA 1325
118214˙2.Gmo.Ice TACCATCG.CCGGAAACGGAAACAGAAACAACGGCGGAACGCGTACAGCATAAACTGACTATAGCTAAGCCTGCTTTAGGTCAAGGGAAA 1325
140219˙3.Gmo.Bal TACCATCG.CCGGAAACGGAAACAGAAACAACGGCGGAACGCGTACAGCATAAACTGACTATAGCTAAGCCTGCTTTAGGTCAAGGGAAA 1325
200093˙3.Gmo.Can TACCATCG.CCGGAAACGGAAACAGAAACAACGGCGGAACGCGTACAGCATAAACTGACTATAGCTAAGCCTGCTTTAGGTCAAGGGAAA 1325
1240 1250 1260 1270 1280 1290 1300 1310 1320