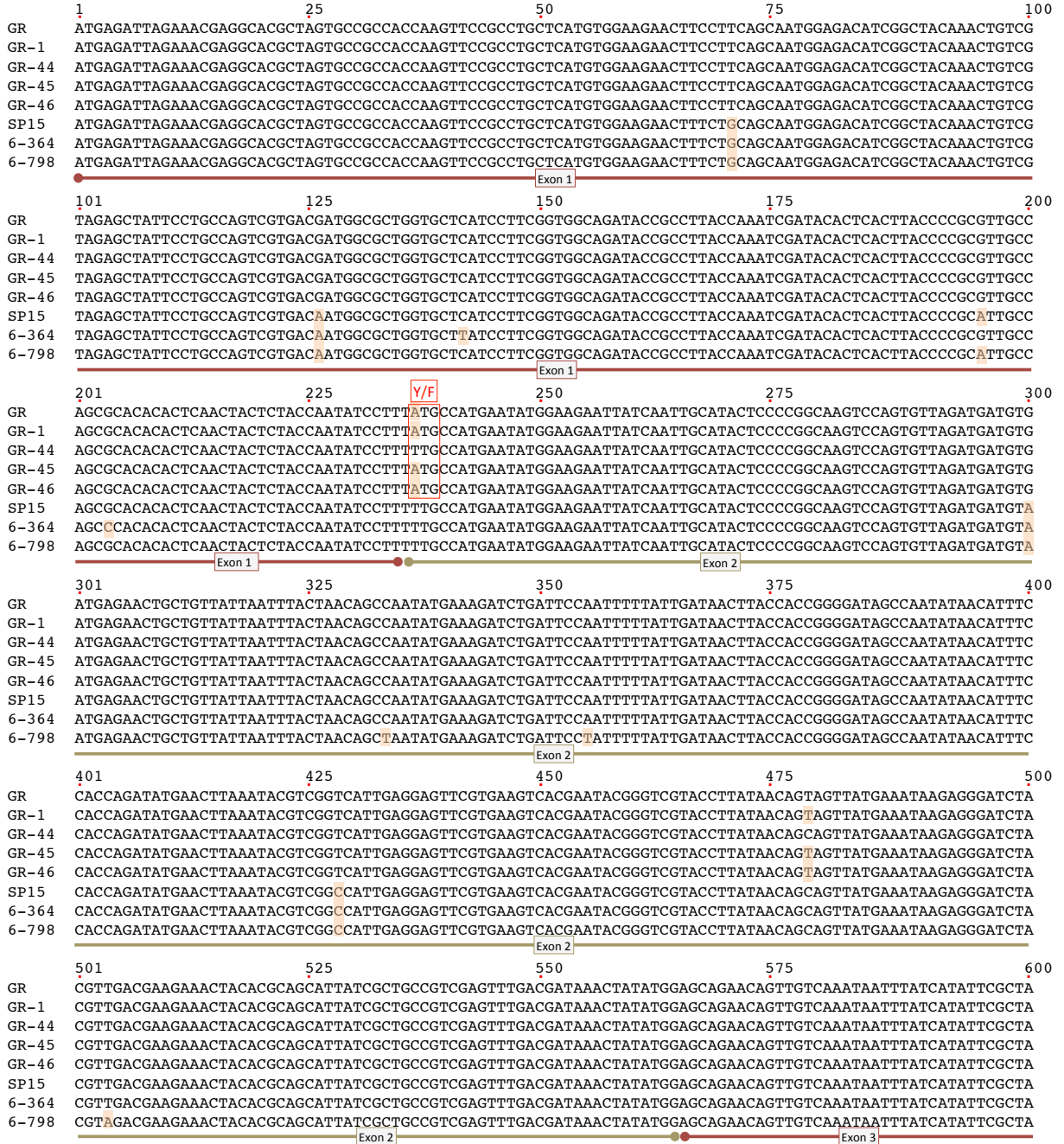


S2 Fig



601 625 650 675 700
GR CTTTTTCCTGAGAGACCCCGTCTCAATTCGTTTTTCCAAACTGGAGGGCGCACCTGGAGATCTGACGGAGTATCCCGGTTTTTCGAAGTGCCTGGGCCTA
GR-1 CTTTTTCCTGAGAGACCCCGTCTCAATTCGTTTTTCCAAACTGGAGGGCGCACCTGGAGATCTGACGGAGTATCCCGGTTTTTCGAAGTGCCTGGGCCTA
GR-44 CTTTTTCCTGAGAGACCCCGTCTCAATTCGTTTTTCCAAACTGGAGGGCGCACCTGGAGATCTGACGGAGTATCCCGGTTTTTCGAAGTGCCTGGGCCTA
GR-45 CTTTTTCCTGAGAGACCCCGTCTCAATTCGTTTTTCCAAACTGGAGGGCGCACCTGGAGATCTGACGGAGTATCCCGGTTTTTCGAAGTGCCTGGGCCTA
GR-46 CTTTTTCCTGAGAGACCCCGTCTCAATTCGTTTTTCCAAACTGGAGGGCGCACCTGGAGATCTGACGGAGTATCCCGGTTTTTCGAAGTGCCTGGGCCTA
SP15 CTTTTTCCTGAGAGACCCCGTCTCAATTCGTTTTTCCAAACTGGAGGGCGCACCTGGAGATCTGACGGAGTATCCCGGTTTTTCGAAGTGCCTGGGCCTA
6-364 CTTTTTCCTGAGAGACCCCGTCTCAATTCGTTTTTCCAAACTGGAGGGCGCACCTGGAGATCTGACGGAGTATCCCGGTTTTTCGAAGTGCCTGGGCCTA
6-798 CTTTTTCCTGAGAGACCCCGTCTCAATTCGTTTTTCCAAACTGGAGGGCGCACCTGGAGATCTGACGGAGTATCCCGGTTTTTCGAAGTGCCTGGGCCTA

701 725 750 775 800
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6-798 GATTTCCCTCACTCGTGGGAAGGTGGTAATGACCCAGGTTACGTTAACGAAATGTTCTGGCACTTCAACAAGTTATATCAACGGAGCTGGTATCAAGGGC

801 825 850 875 900
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6-798 AACCGGTTGGACTTGAAGTCATTTAGGGTGAACATTCAGAGGTACCCGCACCCCGCTACCTTACGACCAAGTTCAGTTGATCTGCTGCAGTTCATGTTT

901 925 950 975 1000
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6-798 CCCCTGTTTCATCATGTTGAGCTTCAGCTACACTCCGTCACATTCGACGGGCGGTACAGTTGAGAAGGAATTGCAATTAAGGAAACTATGAAAATTA

1001 1025 1050 1075 1100
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6-364 TGGGTCCTCCACATGTTGCACTGGACAGCATGGTTTGTAAACAGTTTATCTACCTATCAATCACAGCTGTTCTGCTAGTTGTGTTGCTAAAGGTAAA
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1101 1125 1150 1175 1200
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6-364 TTGGTTTACTAACGACGATGGCTTCAGCGAATATGCTGTATTTACTAATAACACCTTGGACGGTTTTGCTATTCTTCTTGATACTGTATTTATCTTGGCGG
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L/F

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GR 1301 1325 1350 1375 1400
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6-364 TCCTGGCCATGGATATCAATATGTCGACTGCGGTGCAGGTCATCAGTGCTTCAGTATTAACCTGCGATGTCCTATGGTTTCCAACCTAATGCTCGCTAA
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GR 1401 1425 1450 1475 1500
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6-364 GGAAAGCACTGGAGGCTGCAGTGGGGCGACTTCATGACGTCACCAGGACGGACACCACGCGCTTGGTGTTCGGCCACGTGGTCATCATGCTGGTAGTG
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GR 1601 1625 1650 1675 1700
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6-364 TCTGGTTCCCAAATATAAATCGAAAGATGCTGGATTAATTTTCGAAAATGATAACAGTGAATTTGATGATATTATAAAGAAAAGGATCCACAGACCA
6-798 TCTGGTTCCCAAATATAAATCGAAAGATGCTGGATTAATTTTCGAAAATGATAACAGTGAATTTGATGATATTATAAAGAAAAGGATCCACAGACCA

GR 1701 1725 1750 1775 1800
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6-798 CGAAGTTGGTGTAGAAATGCAAAATTTAACAAAAATCTTTGGGAATAACATAGCTGTAAACAATTTATCTTTGAATATCTATGACGATCAAATCACAGTT

2401 2425 2450 2475 2500
GR AAGGACACAGACTTCGACTTTGTGAAATGCTCCGTA...
GR-1 AAGGACACAGACTTCGACTTTGTGAAATGCTCCGTA...
GR-44 AAGGACACAGACTTCGACTTTGTGAAATGCTCCGTA...
GR-45 AAGGACACAGACTTCGACTTTGTGAAATGCTCCGTA...
GR-46 AAGGACACAGACTTCGACTTTGTGAAATGCTCCGTA...
SP15 AAGGACACAGACTTCGACTTTGTGAAATGCTCCGTA...
6-364 AAGGACACAGACTTCGACTTTGTGAAATGCTCCGTA...
6-798 AAGGACACAGACTTCGACTTTGTGAAATGCTCCGTA...

2501 2525 2550 2575 2600
GR ATTTGGTAAATGATTACTCACACGTTTTTGAAGAA...
GR-1 ATTTGGTAAATGATTACTCACACGTTTTTGAAGAA...
GR-44 ATTTGGTAAATGATTACTCACACGTTTTTGAAGAA...
GR-45 ATTTGGTAAATGATTACTCACACGTTTTTGAAGAA...
GR-46 ATTTGGTAAATGATTACTCACACGTTTTTGAAGAA...
SP15 ATTTGGTAAATGATTACTCACACGTTTTTGAAGAA...
6-364 ATTTGGTAAATGATTACTCACACGTTTTTGAAGAA...
6-798 ATTTGGTAAATGATTACTCACACGTTTTTGAAGAA...

2601 2625 2650 2675 2700
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GR-1 TACATTAGAAGATGCTTTATGTCGGTGGTGCCGACT...
GR-44 TACATTAGAAGATGCTTTATGTCGGTGGTGCCGACT...
GR-45 TACATTAGAAGATGCTTTATGTCGGTGGTGCCGACT...
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6-364 TACATTAGAAGATGCTTTATGTCGGTGGTGCCGACT...
6-798 TACATTAGAAGATGCTTTATGTCGGTGGTGCCGACT...

2701 2725 2750 2775 2800
GR GATATATTTAAAACAAGAAATCGATTCATCTTTGGA...
GR-1 GATATATTTAAAACAAGAAATCGATTCATCTTTGGA...
GR-44 GATATATTTAAAACAAGAAATCGATTCATCTTTGGA...
GR-45 GATATATTTAAAACAAGAAATCGATTCATCTTTGGA...
GR-46 GATATATTTAAAACAAGAAATCGATTCATCTTTGGA...
SP15 GATATATTTAAAACAAGAAATCGATTCATCTTTGGA...
6-364 GATATATTTAAAACAAGAAATCGATTCATCTTTGGA...
6-798 GATATATTTAAAACAAGAAATCGATTCATCTTTGGA...

2801 2825 2850 2875 2900
GR TATGGATGAAGCAGTGGCTGGTGTGATCCGCTCGCC...
GR-1 TATGGATGAAGCAGTGGCTGGTGTGATCCGCTCGCC...
GR-44 TATGGATGAAGCAGTGGCTGGTGTGATCCGCTCGCC...
GR-45 TATGGATGAAGCAGTGGCTGGTGTGATCCGCTCGCC...
GR-46 TATGGATGAAGCAGTGGCTGGTGTGATCCGCTCGCC...
SP15 TATGGATGAAGCAGTGGCTGGTGTGATCCGCTCGCC...
6-364 TATGGATGAAGCAGTGGCTGGTGTGATCCGCTCGCC...
6-798 TATGGATGAAGCAGTGGCTGGTGTGATCCGCTCGCC...

2901 2925 2950 2975 3000
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GR-44 GCGTTACGTCATGCTTTTATCACCACCATTAGA...
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GR-46 GCGTTACGTCATGCTTTTATCACCACCATTAGA...
SP15 GCGTTACGTCATGCTTTTATCACCACCATTAGA...
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6-798 GCGTTACGTCATGCTTTTATCACCACCATTAGA...

3001 3025 3050 3075 3100
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 GR-45 GTAGCGTCATCAGTGGGCGCCCTAGCGGCGCAAGCATACCAAAGCCTGTTTCGCCAATCTGGTGTATGGACATGGAAATCAACGCTATGGGAAGCCAGC
 GR-46 GTAGCGTCATCAGTGGGCGCCCTAGCGGCGCAAGCATACCAAAGCCTGTTTCGCCAATCTGGTGTATGGACATGGAAATCAACGCTATGGGAAGCCAGC
 SP15 GTATCGTCATCAGTGGGCGCCCTAGCGGCGCAAGCATACCAAAGCCTGTTTCGCCAATCTGGTGTATGGACATGGAAATCAACGCTATGGGAAGCCAGC
 6-364 GTAGCGTCATCAGTGGGCGCCCTAGCGGCGCAAGCATACCAAAGCCTGTTTCGCCAATCTGGTGTATGGACATGGAAATCAACGCTATGGGAAGCCAGC
 6-798 GTAGCGTCATCAGTGGGCGCCCTAGCGGCGCAAGCATACCAAAGCCTGTTTCGCCAATCTGGTGTATGGACATGGAAATCAACGCTATGGGAAGCCAGC
 Exon 17
 3101 3125 3150 3175 3200
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 GR-1 CAATAGAAGAATATTATCTAAATAGAACAAGTGATCCCGTTGTGATGGGTTTCGCTGCGGCACCCTGCTGATAGGCTCCACATTTGACGACGACTTCGC
 GR-44 CAATAGAAGAATATTATCTAAATAGAACAAGTGATCCCGTTGTGATGGGTTTCGCTGCGGCACCCTGCTGATAGGCTCCACATTTGACGACGACTTCGC
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 6-798 CAATAGAAAATATTATCTAAATAGAACAAGTGATCCCGTTGTGATGGGTTTCGCTGCGGCACCCTGCTGATAGGCTCCACATTTGACGACGACTTCGC
 P/A Exon 17 Exon 18
 3201 3225 3250 3275 3300
 GR TACCGCCTGGTTCAGTAATTTCCGGCTACCACGATGTTGCTACATCGCTTGCGGCAATCCACTCAGCTATTTCCAGATCTAAAAACTCGGATGCAGTACTC
 GR-1 TACCGCCTGGTTCAGTAATTTCCGGCTACCACGATGTTGCTACATCGCTTGCGGCAATCCACTCAGCTATTTCCAGATCTAAAAACTCGGATGCAGTACTC
 GR-44 TACCGCCTGGTTCAGTAATTTCCGGCTACCACGATGTTGCTACATCGCTTGCGGCAATCCACTCAGCTATTTCCAGATCTAAAAACTCGGATGCAGTACTC
 GR-45 TACCGCCTGGTTCAGTAATTTCCGGCTACCACGATGTTGCTACATCGCTTGCGGCAATCCACTCAGCTATTTCCAGATCTAAAAACTCGGATGCAGTACTC
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 6-798 TACCGCCTGGTTCAGTAATTTCCGGCTACCACGATGTTGCTACATCGCTTGCGGCAATCCACTCAGCTATTTCCAGATCTAAAAACTCGGATGCAGTACTC
 Exon 18
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 GR-1 AATGTATATAAATCATCCCTTGAAGCTTCTTACATAGATCAGAGTGACGTGCAGACTATGATAGCTATGTTGTCCATGCAGTTGTCCCTTGGCATCGGCA
 GR-44 AATGTATATAAATCATCCCTTGAAGCTTCTTACATAGATCAGAGTGACGTGCAGACTATGATAGCTATGTTGTCCATGCAGTTGTCCCTTGGCATCGGCA
 GR-45 AATGTATATAAATCATCCCTTGAAGCTTCTTACATAGATCAGAGTGACGTGCAGACTATGATAGCTATGTTGTCCATGCAGTTGTCCCTTGGCATCGGCA
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 6-798 AATGTATATAAATCATCCCTTGAAGCTTCTTACATAGATCAGAGTGACGTGCAGACTATGATAGCTATGTTGTCCATGCAGTTGTCCCTTGGCATCGGCA
 Exon 18 Exon 19
 3401 3425 3450 3475 3500
 GR GTAGTGTGAGCATTGTTAGTGCAGTTTTCATCATGTTTTATATCAAGGAACGATATGTCGGGGGCTAAACTTCTACAAAATGCAGCAGGCGTGGCCCTGC
 GR-1 GTAGTGTGAGCATTGTTAGTGCAGTTTTCATCATGTTTTATATCAAGGAACGATATGTCGGGGGCTAAACTTCTACAAAATGCAGCAGGCGTGGCCCTGC
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 GR-46 GTAGTGTGAGCATTGTTAGTGCAGTTTTCATCATGTTTTATATCAAGGAACGATATGTCGGGGGCTAAACTTCTACAAAATGCAGCAGGCGTGGCCCTGC
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 6-364 GTAGTGTGAGCATTGTTAGTGCAGTTTTCATCATGTTTTATATCAAGGAACGATATGTCGGGGGCTAAACTTCTACAAAATGCAGCAGGCGTGGCCCTGC
 6-798 GTAGTGTGAGCATTGTTAGTGCAGTTTTCATCATGTTTTATATCAAGGAACGATATGTCGGGGGCTAAACTTCTACAAAATGCAGCAGGCGTGGCCCTGC
 Exon 19 Exon 20
 3501 3525 3550 3575 3600
 GR TGTGCTGTGGGCGGCGCAGCGATCTCAATTGGTTTTGGTTCCTCATCACTTGTGTTTCCATCGTCATCTCGTGCCTCGCTTTTGATGTCATCGGGTTG
 GR-1 TGTGCTGTGGGCGGCGCAGCGATCTCAATTGGTTTTGGTTCCTCATCACTTGTGTTTCCATCGTCATCTCGTGCCTCGCTTTTGATGTCATCGGGTTG
 GR-44 TGTGCTGTGGGCGGCGCAGCGATCTCAATTGGTTTTGGTTCCTCATCACTTGTGTTTCCATCGTCATCTCGTGCCTCGCTTTTGATGTCATCGGGTTG
 GR-45 TGTGCTGTGGGCGGCGCAGCGATCTCAATTGGTTTTGGTTCCTCATCACTTGTGTTTCCATCGTCATCTCGTGCCTCGCTTTTGATGTCATCGGGTTG
 GR-46 TGTGCTGTGGGCGGCGCAGCGATCTCAATTGGTTTTGGTTCCTCATCACTTGTGTTTCCATCGTCATCTCGTGCCTCGCTTTTGATGTCATCGGGTTG
 SP15 TGTGCTGTGGGCGGCGCAGCGATCTCAATTGGTTTTGGTTCCTCATCACTTGTGTTTCCATCGTCATCTCGTGCCTCGCTTTTGATGTCATCGGGTTG
 6-364 TGTGCTGTGGGCGGCGCAGCGATCTCAATTGGTTTTGGTTCCTCATCACTTGTGTTTCCATCGTCATCTCGTGCCTCGCTTTTGATGTCATCGGGTTG
 6-798 TGTGCTGTGGGCGGCGCAGCGATCTCAATTGGTTTTGGTTCCTCATCACTTGTGTTTCCATCGTCATCTCGTGCCTCGCTTTTGATGTCATCGGGTTG
 Exon 20

3601 3625 3650 3675 3700
 GR TCGACCGTGCATGAATTAGGCCGAATGTTTCTGTGCGTCATGGTATACGGTGCAGCAATGTTGCCATTAGTGTACCTTCTGTCACTTAAGTTCAAGGGAC
 GR-1 TCGACCGTGCATGAATTAGGCCGAATGTTTCTGTGCGTCATGGTATACGGTGCAGCGATGTTGCCATTAGTGTACCTTCTGTGCGTTAAGTTCAAGGGAC
 GR-44 TCGACCGTGCATGAATTAGGCCGAATGTTTCTGTGCGTCATGGTATACGGTGCAGCGATGTTGCCATTAGTGTACCTTCTGTGCGTTAAGTTCAAGGGAC
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 6-364 TCGACCGTGCATGAATTAGGCCGAATGTTTCTGTGCGTCATGGTATACGGTGCAGCAATGTTGCCATTAGTGTACCTTCTGTCACTTAAGTTCAAGGGAC
 6-798 TCAACCGTGCATGAATTAGGCCAAATGTTTCTGTGCGTCATGGTATACGGTGCAGCAATGTTGCCATTAGTGTACCTTCTGTCACTTAAGTTCAAGGGAC

3701 3725 3750 3775 3800
 GR CAGCTGTCGGCTTCGTGGGTTTCTACTTCTCAACGTGCTTTTCGGTATGATGGGTGCGCAGGTGGTGGAGGCACATCTTCTCTATGCTGGACACGGA
 GR-1 CAGCTGTCGGCTTCGTGGGTTTCTACTTCTCAACGTGCTTTTCGGTATGATGGGTGCGCAGGTGGTGGAGGCACATCTCTCTCTGTCTGGACACGGA
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3801 3825 3850 3875 3900
 GR GCAAGCCGCCACATCTTCTGACTACTTACTGCAGTCTTACCCTGTTACAGCCTTGTGCACATCTATCAGGTTTTTGAATCAAGTCGGCTACGGGAGTAC
 GR-1 GCAAGCCGCCACATCTTCTGACTACTTACTGCAGTCTTACCCTGTTACAGCCTTGTGCACATCTATCAGGTTTTTGAATCAAGTCGGCTACGGGAGTAC
 GR-44 GCAAGCCGCCACATCTTCTGACTACTTACTGCAGTCTTACCCTGTTACAGCCTTGTGCACATCTATCAGGTTTTTGAATCAAGTCGGCTACGGGAGTAC
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 SP15 GCAAGCCGCCACATCTTCTGACTACTTACTGCAGTCTTACCCTGTTACAGCCTTGTGCACATCTATCAGGTTTTTGAATCAAGTCGGCTACGGGAGTAC
 6-364 GCAAGCCGCCACATCTTCTGACTACTTACTGCAGTCTTACCCTGTTACAGCCTTGTGCACATCTATCAGGTTTTTGAATCAAGTCGGCTACGGGAGTAC
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3901 3925 3950 3975 4000
 GR ACTTGCTTACAAGCCTGTGAGTACTTGAAGCAGTATACCCGAATCTAGAGTGCAACATGGCAAGCATGTGCGAATTCACAGTGACTGCTGCGTTCGTG
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 GR-44 ACTTGCTTACAAGCCTGTGAGTACTTGAAGCAGTATACCCGAATCTAGAGTGCAACATGGCAAGCATGTGCGAATTCACAGTGACTGCTGCGTTCGTG
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 6-364 ACTTGCTTACAAGCCTGTGAGTACTTGAAGCAGTATACCCGAATCTAGAGTGCAACATGGCAAGCATGTGCGAATTCACAGTGACTGCTGCGTTCGTG
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4001 4025 4050 4075 4100
 GR AAAATCCATACTTCGATTGGGAGGAACAGGCGCTTGTAGGTAAGTGTCTTCTCTGCCTAATGTTCTGGTTGCTGCTTATGACCAATTGA
 GR-1 AAAATCCATACTTCGATTGGGAGGAACAGGCGCTTGTAGGTAAGTGTCTTCTCTGCCTAATGTTCTGGTTGCTGCTTATGACCAATTGA
 GR-44 AAAATCCATACTTCGATTGGGAGGAACAGGCGCTTGTAGGTAAGTGTCTTCTCTGCCTAATGTTCTGGTTGCTGCTTATGACCAATTGA
 GR-45 AAAATCCATACTTCGATTGGGAGGAACAGGCGCTTGTAGGTAAGTGTCTTCTCTGCCTAATGTTCTGGTTGCTGCTTATGACCAATTGA
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 6-364 AAAATCCATACTTCGATTGGGAGGAACAGGCGCTTGTAGGTAAGTGTCTTCTCTGCCTAATGTTCTGGTTGCTGCTTATGACCAATTGA
 6-798 AAAATCCATACTTCGATTGGGAGGAACAGGCGCTTGTAGGTAAGTGTCTTCTCTGCCTAATGTTCTGGTTGCTGCTTATGACCAATTGA

4101 4125 4150 4175 4200
 GR ATATAGAGTGGTGCAAAAGGTGTTTCACATTCAGAAGACTCCTCCTCCAATAGACGAGAGCAGGTTGGACGAGGACGTTGATGACGAGGCGAGGCGCGG
 GR-1 ATATAGAGTGGTGCAAAAGGTGTTTCACATTCAGAAGACTCCTCCTCCAATAGACGAGAGCAGGTTGGACGAGGACGTTGATGACGAGGCGAGGCGCGG
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 6-798 ATACAGAGTGGTGCAAAAGGTGTTTCACATTCAGAAGACTCCTCCTCCAATAGACGAGAGCAGGTTGGACGAGGACGTTGATGACGAGGCGAGGCGCGG



K/T

4201 4225 4250 4275 4300

GR CCGTGGTCCACCGACACGACGCCGCGAACACGCCCTTCTAGCTCACGACCTCTCCAAGTACTACGGGAAACATCTCGCCGTAGACCAAGTCTCGTTTA
GR-1 CGCGTGGTCCACCGACACGACGCCGCGAACACGCCCTTCTAGCTCACGACCTCTCCAAGTACTACGGGAAACATCTCGCCGTAGACCAAGTCTCGTTTA
GR-44 CGCGTGGTCCACCGACACGACGCCGCGAACACGCCCTTCTAGCTCACGACCTCTCCAAGTACTACGGGAAACATCTCGCCGTAGACCAAGTCTCGTTTA
GR-45 CGCGTGGTCCACCGACACGACGCCGCGAACACGCCCTTCTAGCTCACGACCTCTCCAAGTACTACGGGAAACATCTCGCCGTAGACCAAGTCTCGTTTA
GR-46 CGCGTGGTCCACCGACACGACGCCGCGAACACGCCCTTCTAGCTCACGACCTCTCCAAGTACTACGGGAAACATCTCGCCGTAGACCAAGTCTCGTTTA
SP15 CGCGTGGTCCACCGACACGACGCCGCGAACACGCCCTTCTAGCTCACGACCTCTCCAAGTACTACGGGAAACATCTCGCCGTAGACCAAGTCTCGTTTA
6-364 CGCGTGGTCCACCGACACGACGCCGCGAACACGCCCTTCTAGCTCACGACCTCTCCAAGTACTACGGGAAACATCTCGCCGTAGACCAAGTCTCGTTTA
6-798 CGCGTGGTCCACCGACACGACGCCGCGAACACGCCCTTCTAGCTCACGACCTCTCCAAGTACTACGGGAAACATCTCGCCGTAGACCAAGTCTCGTTTA

Exon 25

4301 4325 4350 4375 4400

GR GTGTGAACGACGGCGAATGCTTCGGTCTATTGGGTGTGAATGGTGCCGGAAAAACAACCACCTTCAAGATGCTGATGGGTGATGAGTCCATTTCAAGCGG
GR-1 GTGTGAACGACGGCGAATGCTTCGGTCTATTGGGTGTGAATGGTGCCGGAAAAACAACCACCTTCAAGATGCTGATGGGTGATGAGTCCATTTCAAGCGG
GR-44 GTGTGAACGACGGCGAATGCTTCGGTCTATTGGGTGTGAATGGTGCCGGAAAAACAACCACCTTCAAGATGCTGATGGGTGATGAGTCCATTTCAAGCGG
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6-798 GTGTGAACGACGGCGAATGCTTCGGTCTATTGGGTGTGAATGGTGCCGGAAAAACAACCACCTTCAAGATGCTGATGGGTGATGAGTCCATTTCAAGCGG

Exon 26

4401 4425 4450 4475 4500

GR CGAGGCGTATGTCTCCGGGCACTCGGTGCAGAGGAATCTTGATAGAGTGCATGAGAATATTGGATATTGTCCGCAATTTGACGCATTATTTGGTGAGCTG
GR-1 CGAGGCGTATGTCTCCGGGCACTCGGTGCAGAGGAATCTTGATAGAGTGCATGAGAATATTGGATATTGTCCGCAATTTGATGCAATTTATTTGGTGAGCTG
GR-44 CGAGGCGTATGTCTCCGGGCACTCGGTGCAGAGGAATCTTGATAGAGTGCATGAGAATATTGGATATTGTCCGCAATTTGATGCAATTTATTTGGTGAGCTG
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SP15 CGAGGCGTATGTCTCCGGGCACTCGGTGCAGAGGAATCTTGATAGAGTGCATGAGAATATTGGATATTGTCCGCAATTTGACGCATTATTTGGTGAGCTG
6-364 CGAGGCGTATGTCTCCGGGCACTCGGTGCAGAGGAATCTTGATAGAGTGCATGAGAATATTGGATATTGTCCGCAATTTGATGCAATTTATTTGGTGAGCTG
6-798 CGAGGCGTATGTCTCCGGGCACTCGGTGCAGAGGAATCTTGATAGAGTGCATGAGAATATTGGATATTGTCCGCAATTTGACGCATTATTTGGTGAGCTG

Exon 27

4501 4525 4550 4575 4600

GR ACGGGTCGCCAGACACTCCACATGTTTGCCTTGATGCGCGGCTTGCCTTTGCGCACTGCGGCACCTCAGCTGAAACACTCGCACATGCGCTTGGCTTCT
GR-1 ACGGGTCGCCAGACACTCCACATGTTTGCCTTGATGCGCGGCTTGCCTTTGCGCACTGCGGCACCTCAGCTGAAACACTCGCACATGCGCTTGGCTTCT
GR-44 ACGGGTCGCCAGACACTCCACATGTTTGCCTTGATGCGCGGCTTGCCTTTGCGCACTGCGGCACCTCAGCTGAAACACTCGCACATGCGCTTGGCTTCT
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6-364 ACGGGTCGCCAGACACTCCACATGTTTGCCTTGATGCGCGGCTTGCCTTTGCGCACTGCGGCACCTCAGCTGAAACACTCGCACATGCGCTTGGCTTCT
6-798 ACGGGTCGCCAGACACTCCACATGTTTGCCTTGATGCGCGGCTTGCCTTTGCGCACTGCGGCACCTCAGCTGAAACACTCGCACATGCGCTTGGCTTCT

Exon 28

4601 4625 4650 4675 4700

GR TCAAAACATCTCGATAAAAAGGGTGCATCAGTATTTCAGGCGGCACGAAACGCAAGCTTAAACACGGCGATAGCATTTCATGGGACGAAACACGGCTTGTGTTTGT
GR-1 TCAAAACATCTCGATAAAAAGGGTGCATCAGTATTTCAGGCGGCACGAAACGCAAGCTTAAACACGGCGATAGCATTTCATGGGACGAAACACGGCTTGTGTTTGT
GR-44 TCAAAACATCTCGATAAAAAGGGTGCATCAGTATTTCAGGCGGCACGAAACGCAAGCTTAAACACGGCGATAGCATTTCATGGGACGAAACACGGCTTGTGTTTGT
GR-45 TCAAAACATCTCGATAAAAAGGGTGCATCAGTATTTCAGGCGGCACGAAACGCAAGCTTAAACACGGCGATAGCATTTCATGGGACGAAACACGGCTTGTGTTTGT
GR-46 TCAAAACATCTCGATAAAAAGGGTGCATCAGTATTTCAGGCGGCACGAAACGCAAGCTTAAACACGGCGATAGCATTTCATGGGACGAAACACGGCTTGTGTTTGT
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6-798 TCAAAACATCTAGATAAAAAGGGTGCATCAGTATTTCAGGCGGCACGAAACGCAAGCTTAAACACGGCGATAGCATTTCATGGGACGAAACACGGCTTGTGTTTGT

Exon 29

4701 4725 4750 4775 4800

GR TGATGAGCCTACAACGGAGTTCGATCCCGCCGCTAAACGCCACGATATGGCGCGCTACCCGCGCGCTGCAGCGAGCAGGTCGCGGCTGGTACTGACGTCA
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GR-44 TGATGAGCCTACAACGGAGTTCGATCCCGCCGCTAAACGCCACGATATGGCGCGCTACCCGCGCGCTGCAGCGAGCAGGTCGCGGCTGGTACTGACGTCA
GR-45 TGATGAGCCTACAACGGAGTTCGATCCCGCCGCTAAACGCCACGATATGGCGCGCTACCCGCGCGCTGCAGCGAGCAGGTCGCGGCTGGTACTGACGTCA
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4801 4825 4850 4875 4900
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6-798 CACAGTATGGAGGAGTGCAGGCGTTGTGTTTCGCGACTGACCATCATGGTCAATGGTCGCTTCCAGTGTCTGGGCACGCCACAACATCTCAAGAACAAT

4901 4925 4950 4975 5000
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5001 5025 5050 5075 5100
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5101 5125 5150 5175 5200
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5201 5225
GR ACCAAAGAGAAGAAGGAACGACGTTA
GR-1 ACCAAAGAGAAGAAGGAACGACGTTA
GR-44 ACCAAAGAGAAGAAGGAACGACGTTA
GR-45 ACCAAAGAGAAGAAGGAACGACGTTA
GR-46 ACCAAAGAGAAGAAGGAACGACGTTA
SP15 ACCAAAGAGAAGAAGGAACGACGTTA
6-364 ACCAAAGAGAAGAAGGAACGACGTTA
6-798 ACCAAAGAGAAGAAGGAACGACGTTA