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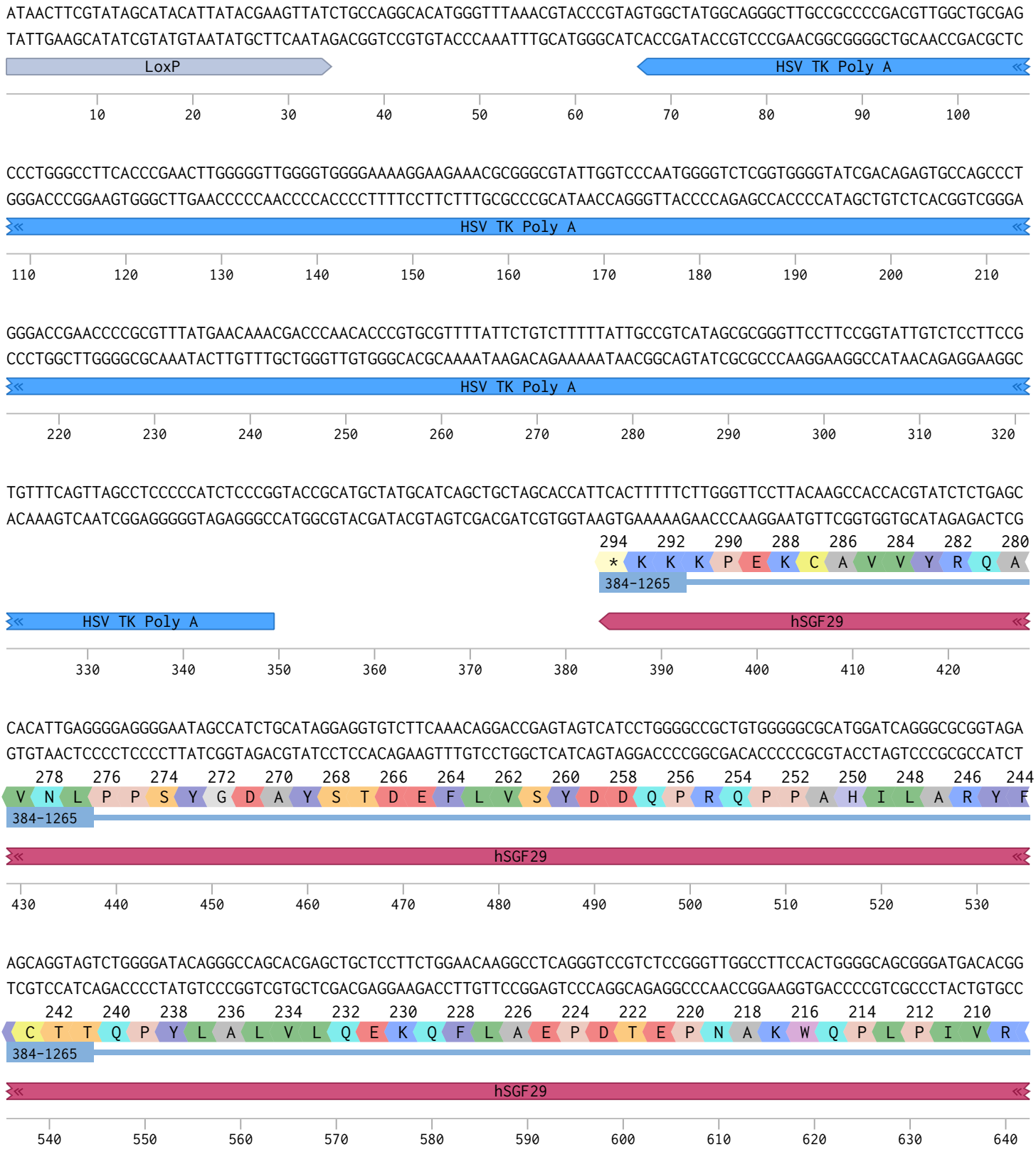

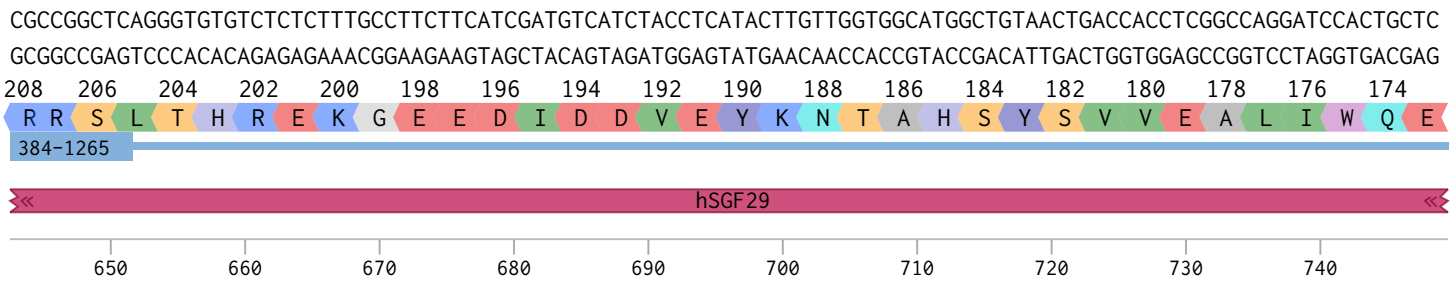
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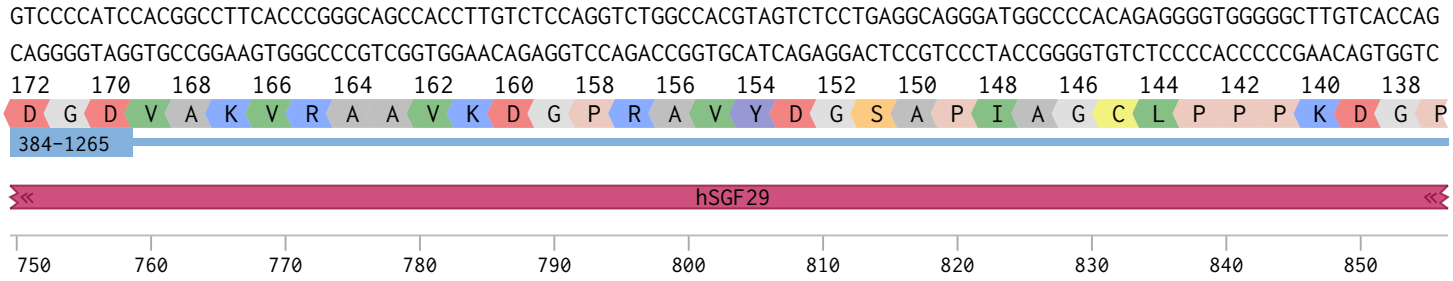
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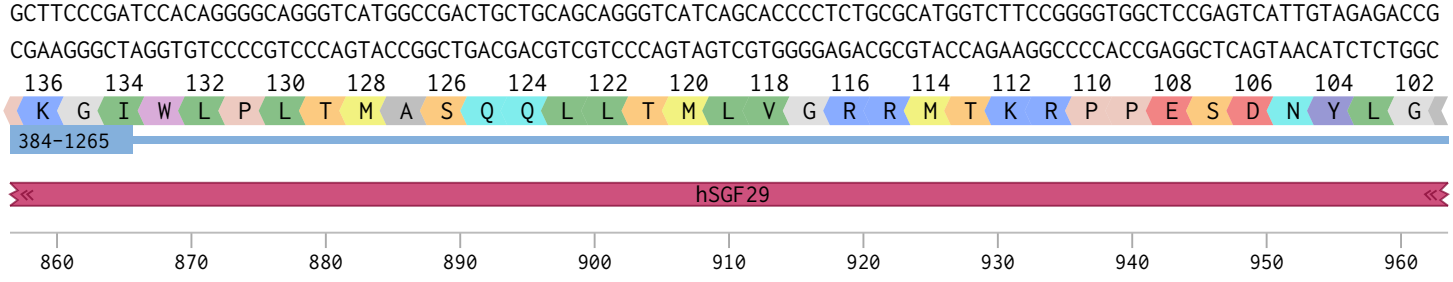
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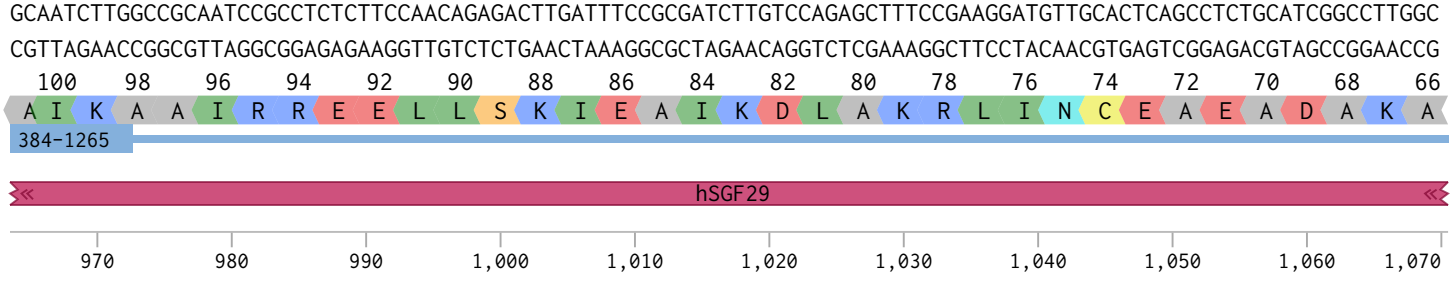
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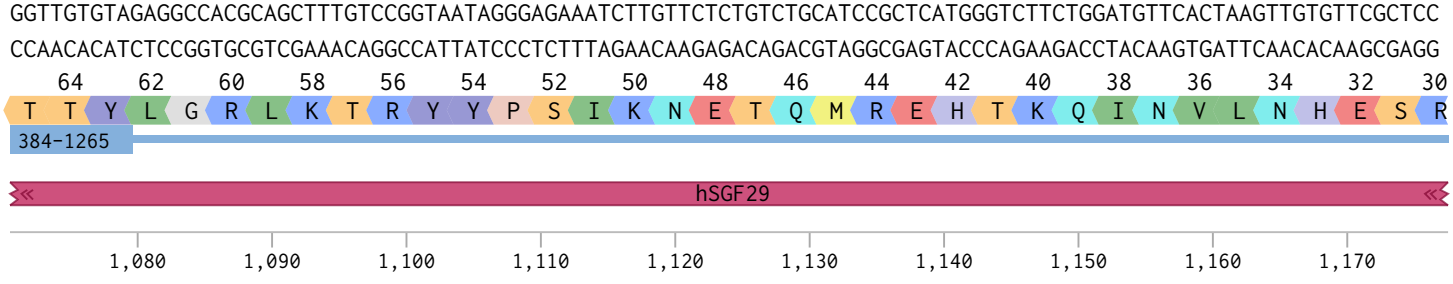
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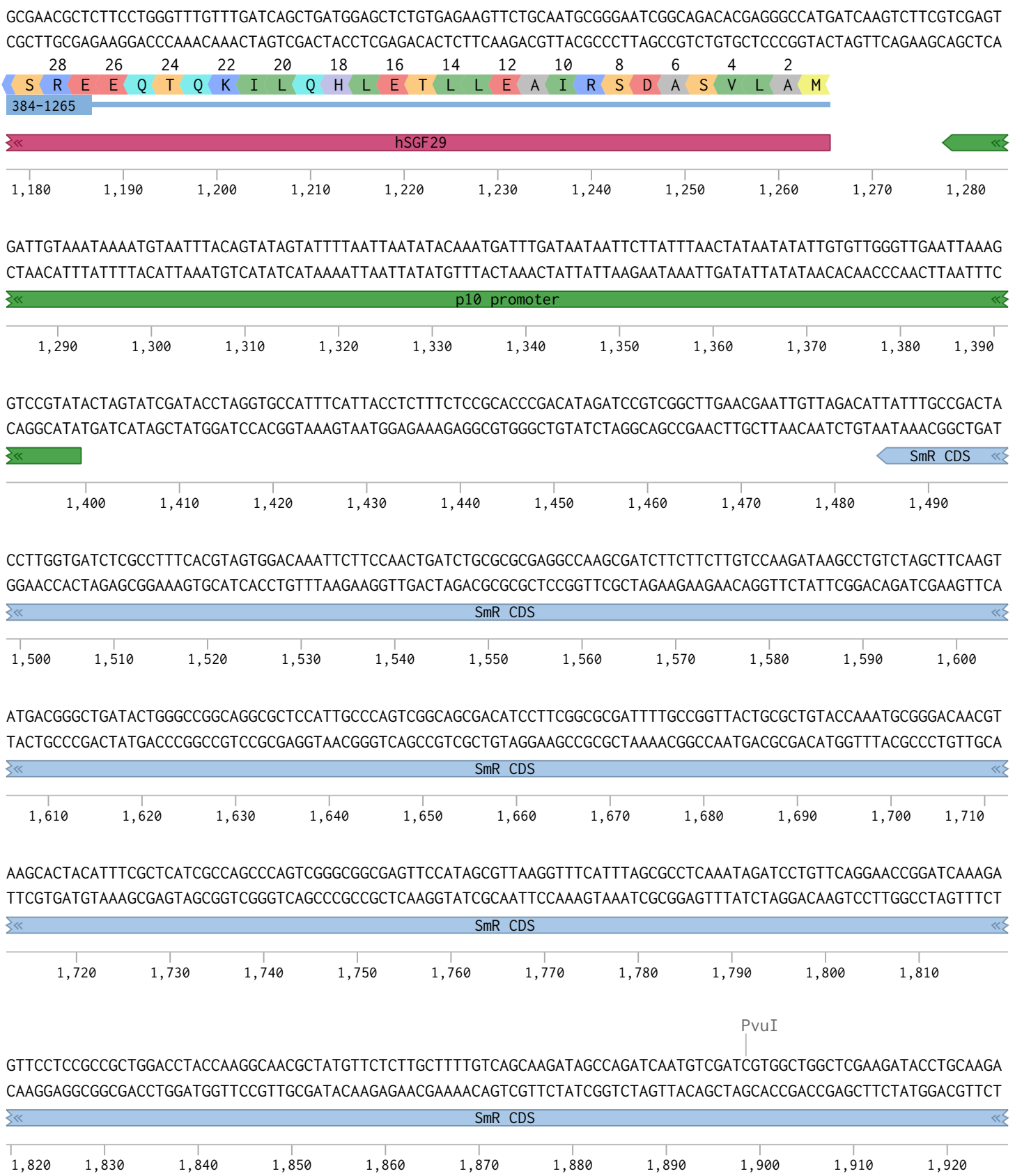
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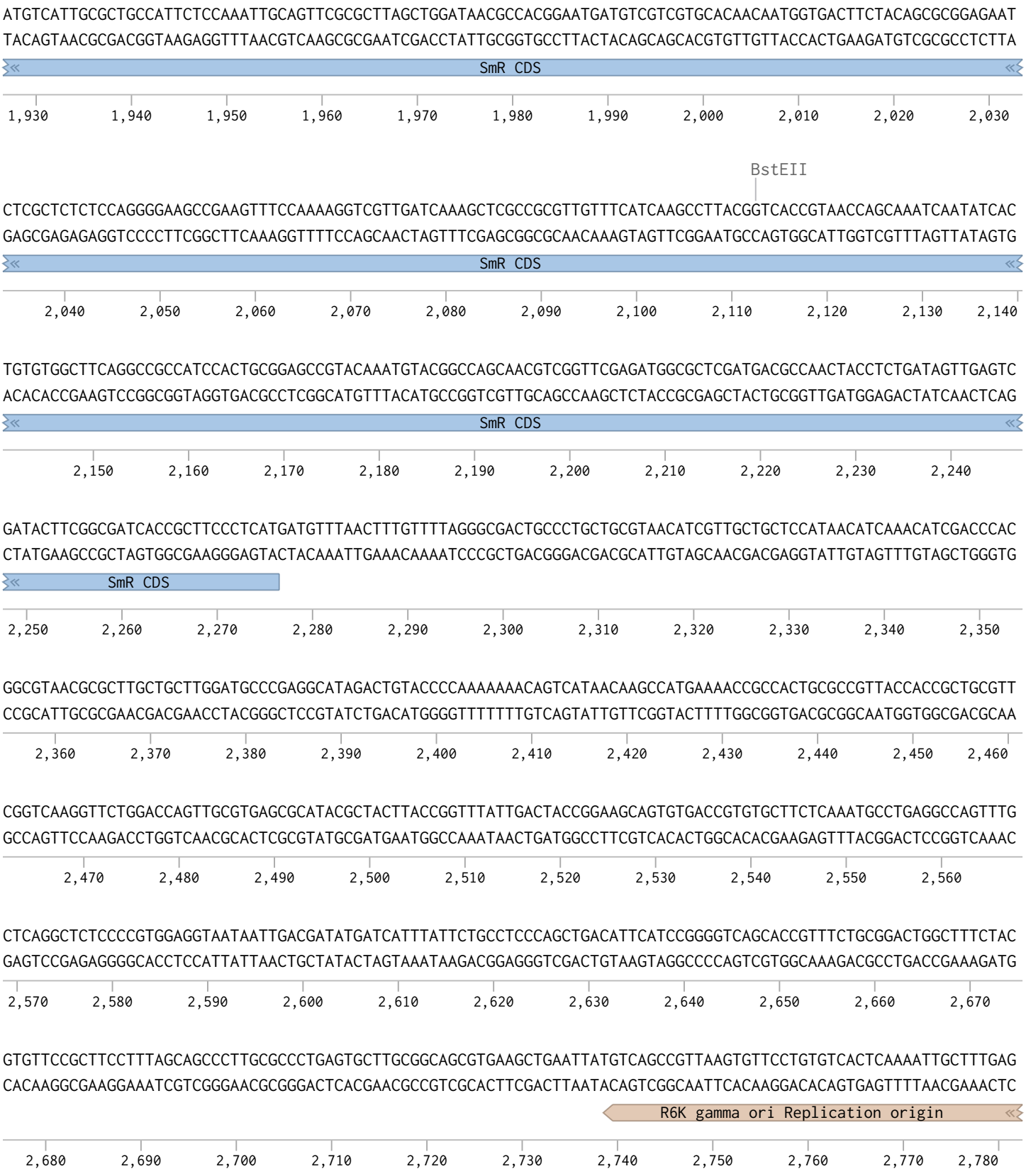
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R6K gamma ori Replication origin
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R6K gamma ori Replication origin

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R6K gamma ori Replication origin

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LoxP HSV TK Poly A

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* G D L L K L I S E R E K L T K W
3480-4778

HSV TK Poly A TADA3

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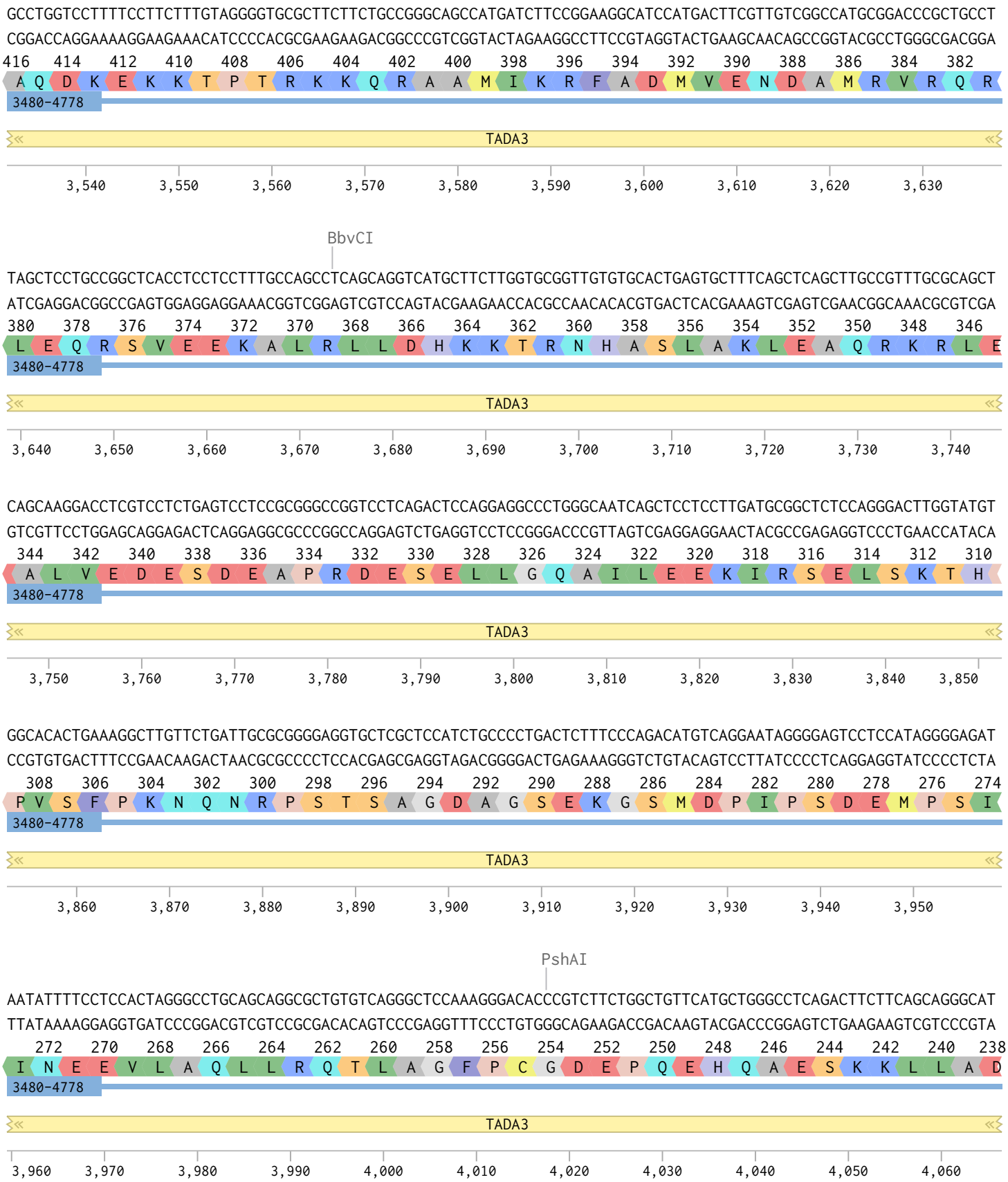
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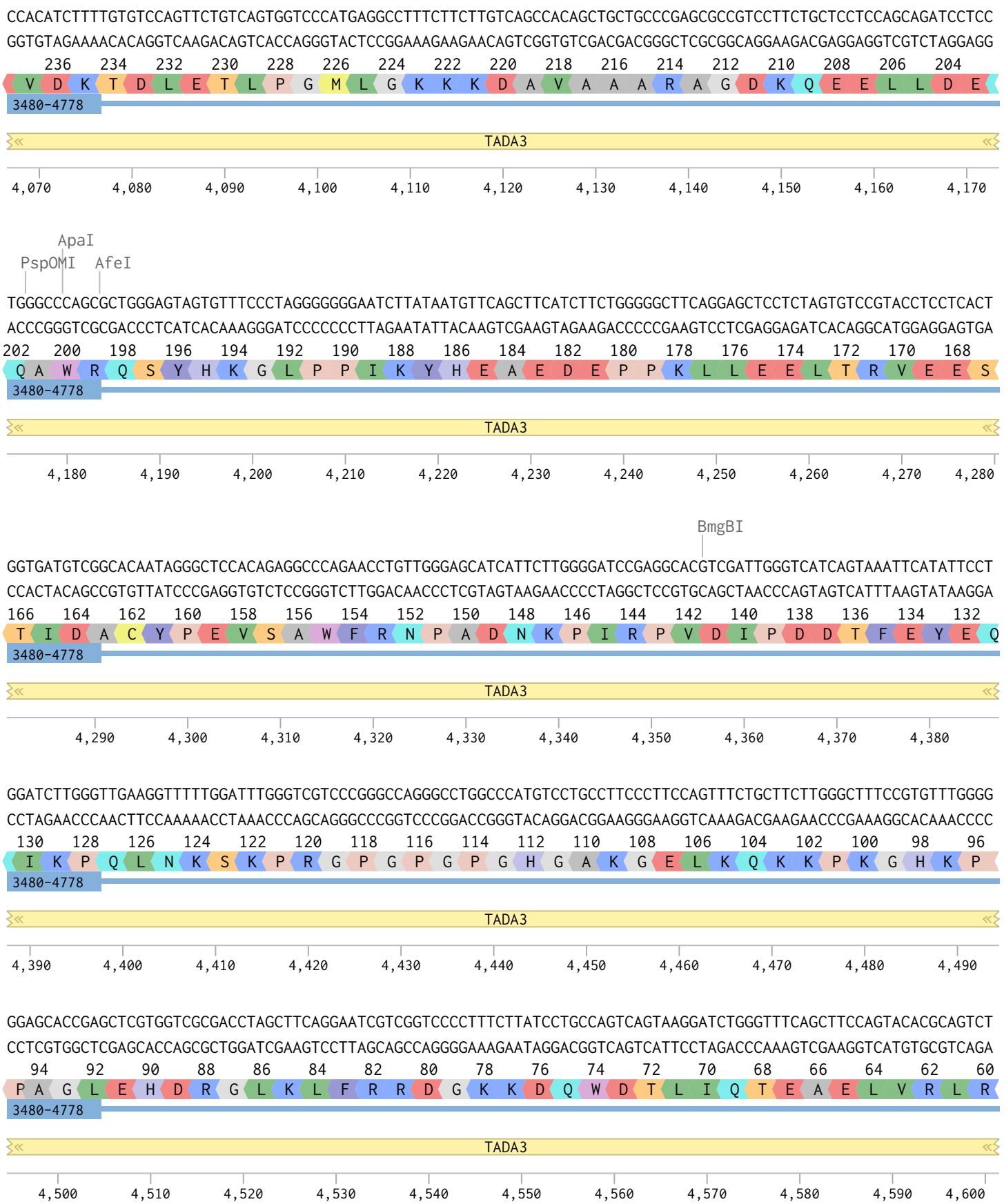
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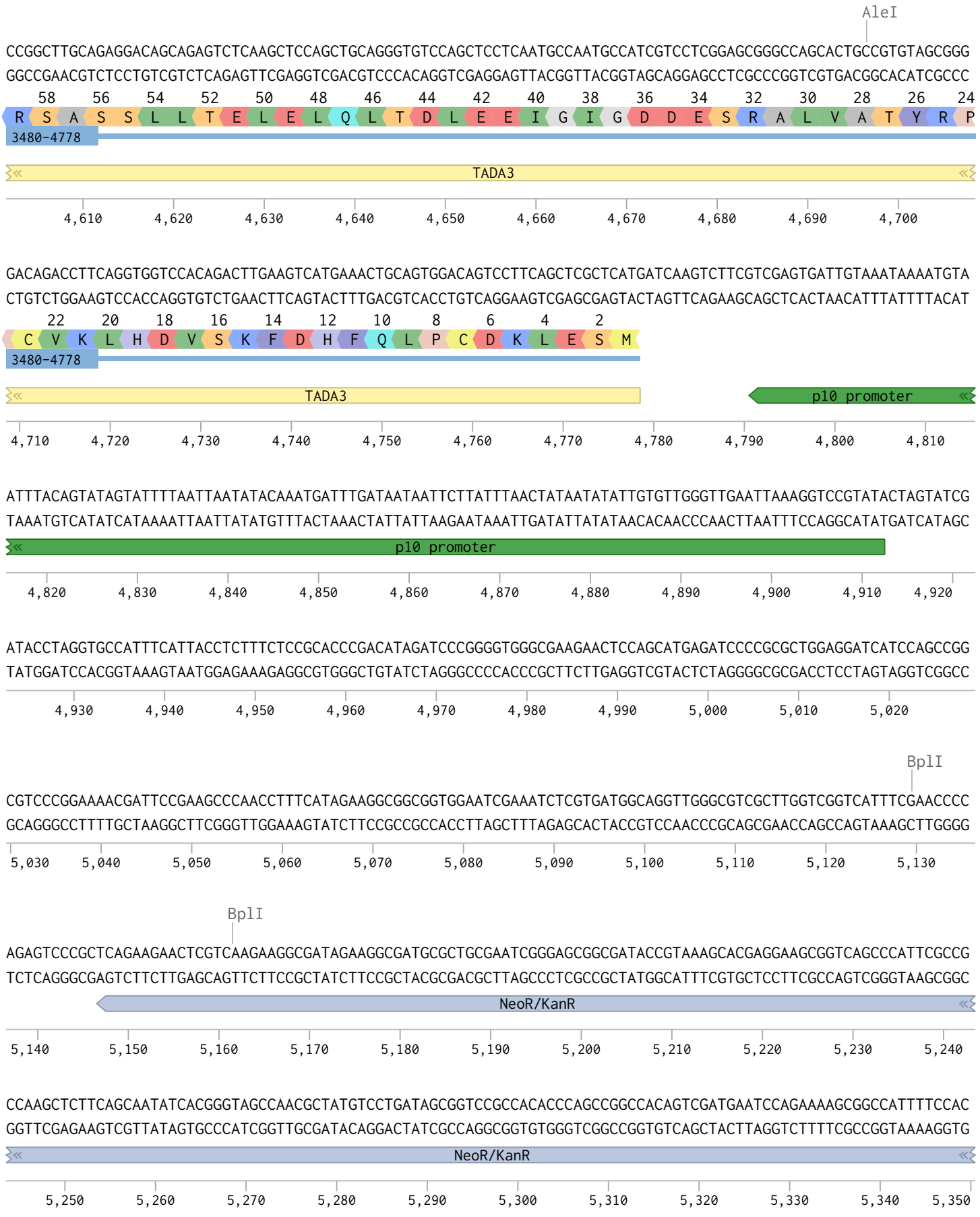
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p10 promoter
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4,930 4,940 4,950 4,960 4,970 4,980 4,990 5,000 5,010 5,020
BplI
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5,030 5,040 5,050 5,060 5,070 5,080 5,090 5,100 5,110 5,120 5,130
BplI
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NeoR/KanR
5,140 5,150 5,160 5,170 5,180 5,190 5,200 5,210 5,220 5,230 5,240
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NeoR/KanR
5,250 5,260 5,270 5,280 5,290 5,300 5,310 5,320 5,330 5,340 5,350

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NeoR/KanR

5,360 5,370 5,380 5,390 5,400 5,410 5,420 5,430 5,440 5,450

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NeoR/KanR

5,460 5,470 5,480 5,490 5,500 5,510 5,520 5,530 5,540 5,550 5,560

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NeoR/KanR

5,570 5,580 5,590 5,600 5,610 5,620 5,630 5,640 5,650 5,660 5,670

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NeoR/KanR

5,680 5,690 5,700 5,710 5,720 5,730 5,740 5,750 5,760 5,770

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NeoR/KanR

5,780 5,790 5,800 5,810 5,820 5,830 5,840 5,850 5,860 5,870 5,880

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XcmI

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6,000 6,010 6,020 6,030 6,040 6,050 6,060 6,070 6,080 6,090

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R6Kgamma

6,210 6,220 6,230 6,240 6,250 6,260 6,270 6,280 6,290 6,300 6,310

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6,320 6,330 6,340 6,350 6,360 6,370 6,380 6,390 6,400 6,410 6,420

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R6Kgamma

6,430 6,440 6,450 6,460 6,470 6,480 6,490 6,500 6,510 6,520

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R6Kgamma

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LoxP HSV TK Poly A

6,640 6,650 6,660 6,670 6,680 6,690 6,700 6,710 6,720 6,730 6,740

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HSV TK Poly A

6,750 6,760 6,770 6,780 6,790 6,800 6,810 6,820 6,830 6,840

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HSV TK Poly A

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HSV TK Poly A

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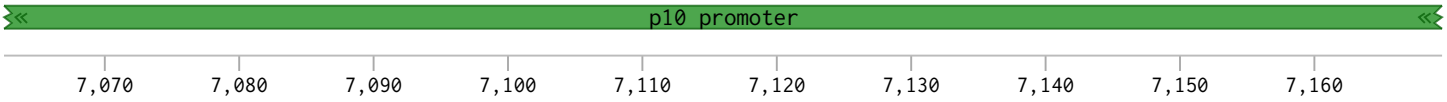

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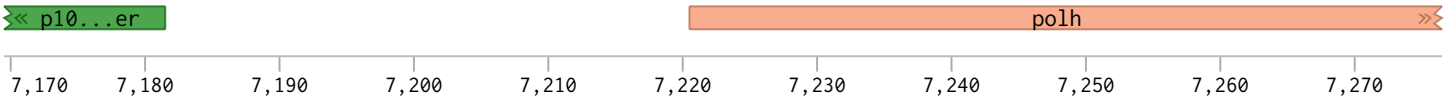

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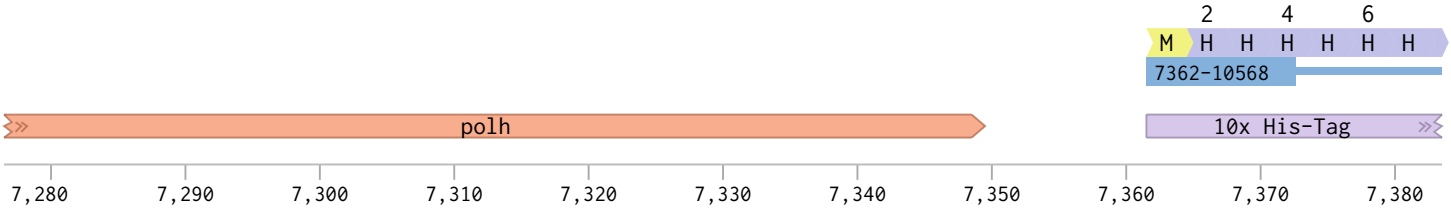

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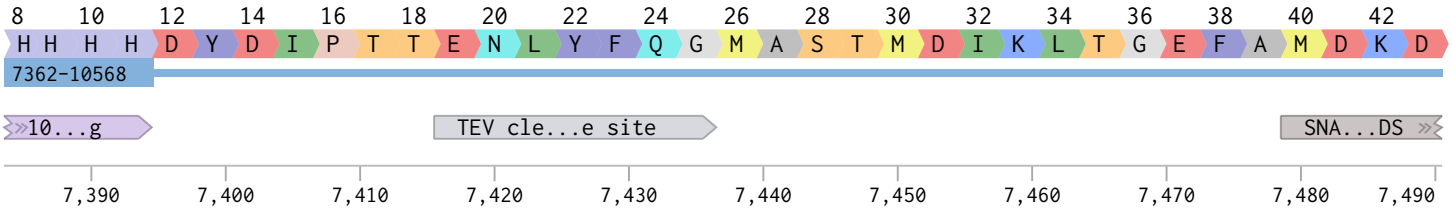

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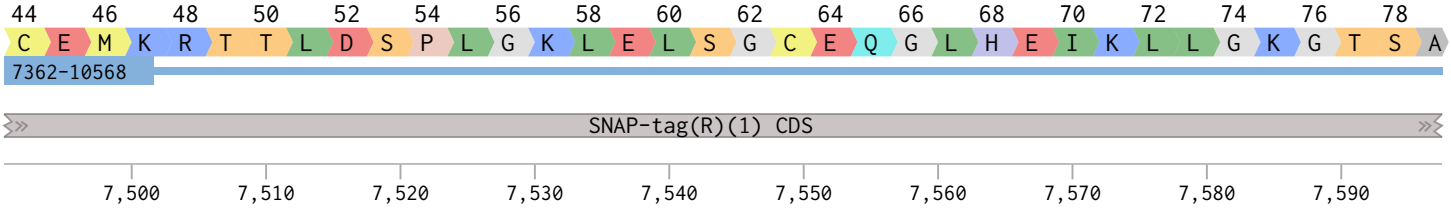

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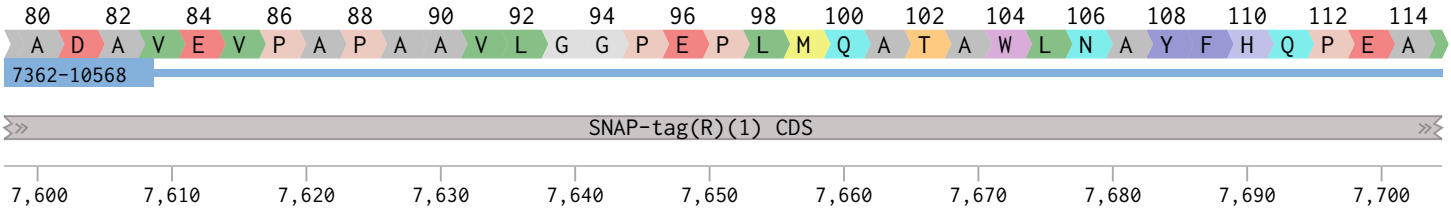

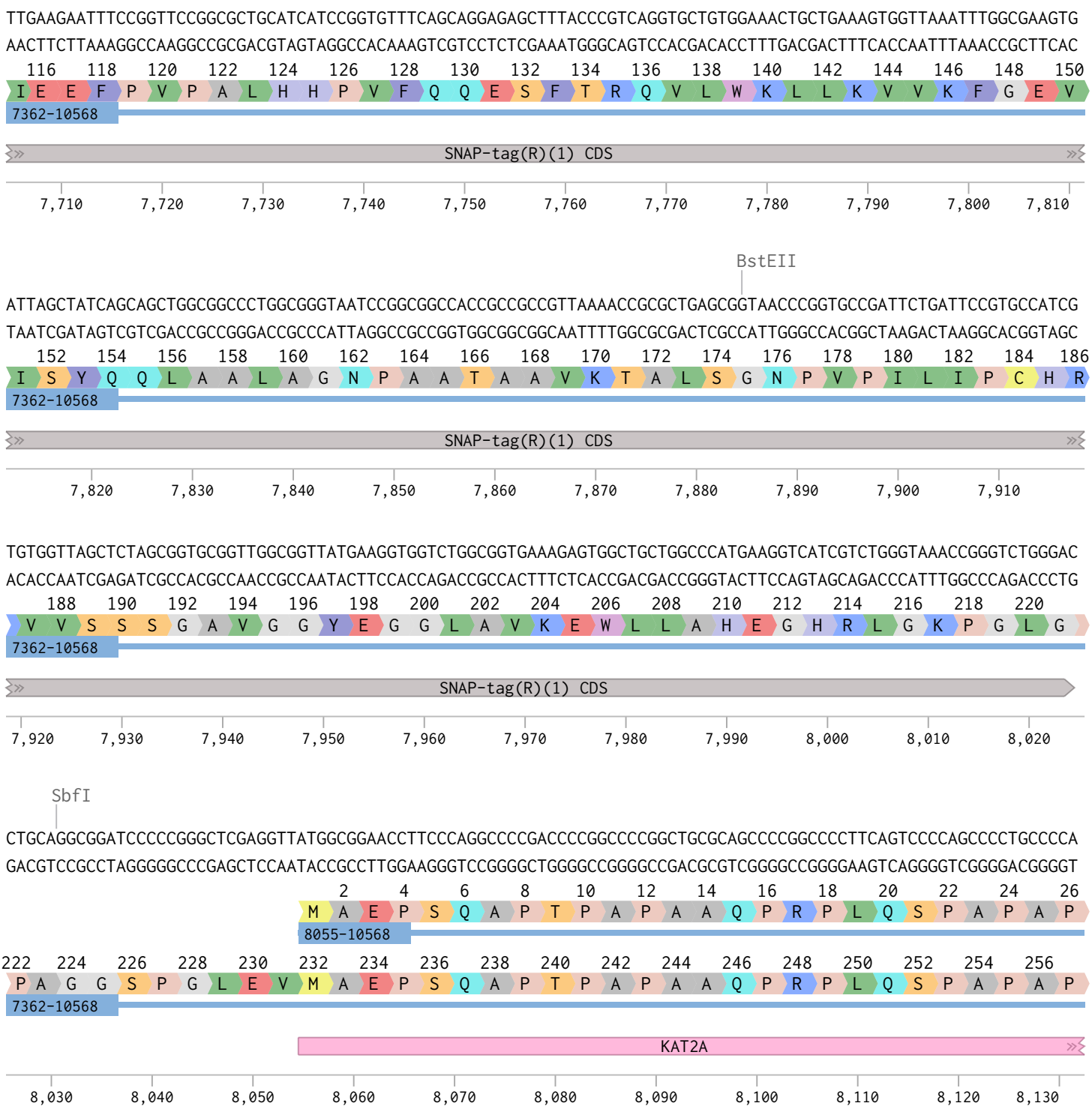
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8055-10568
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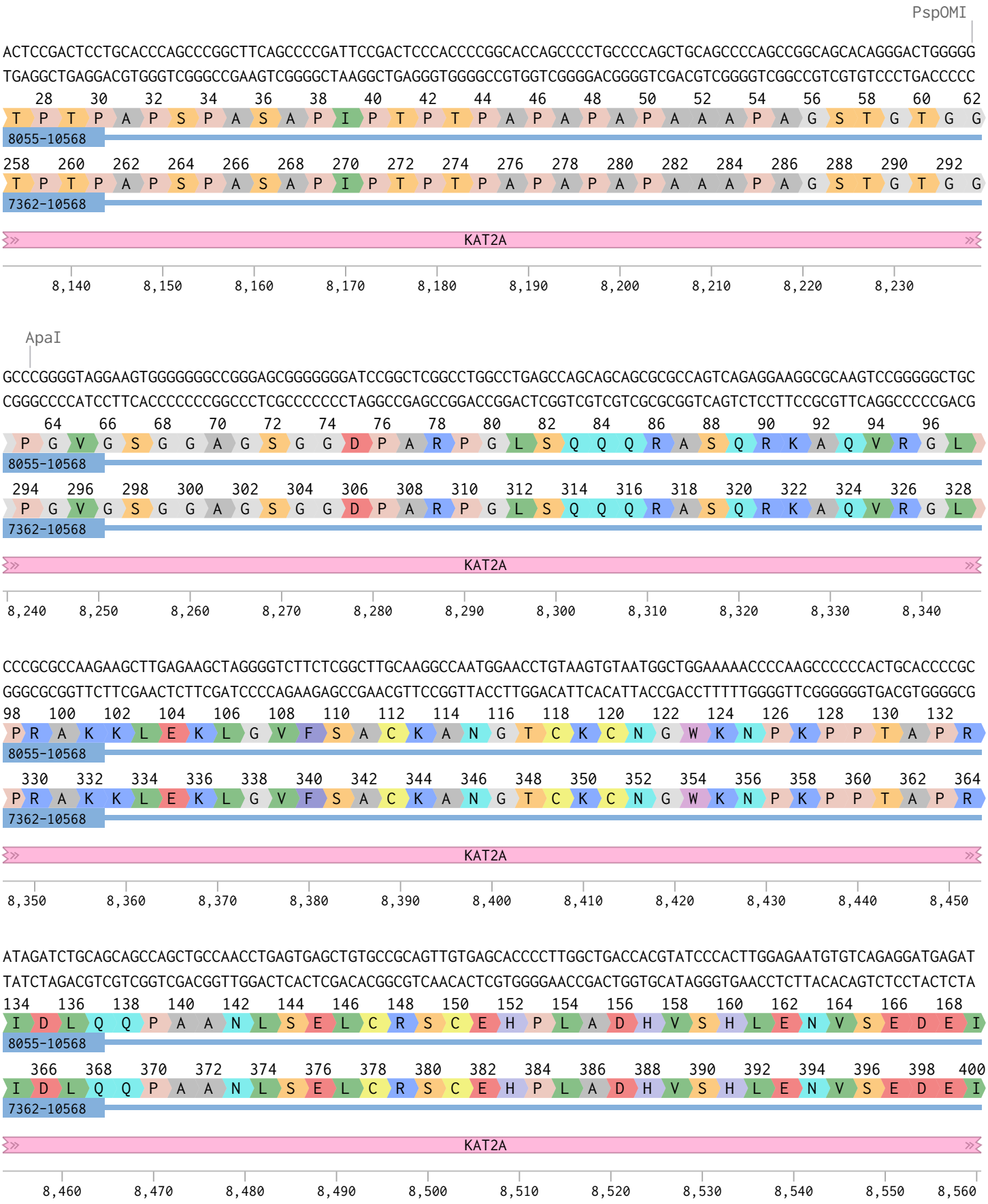
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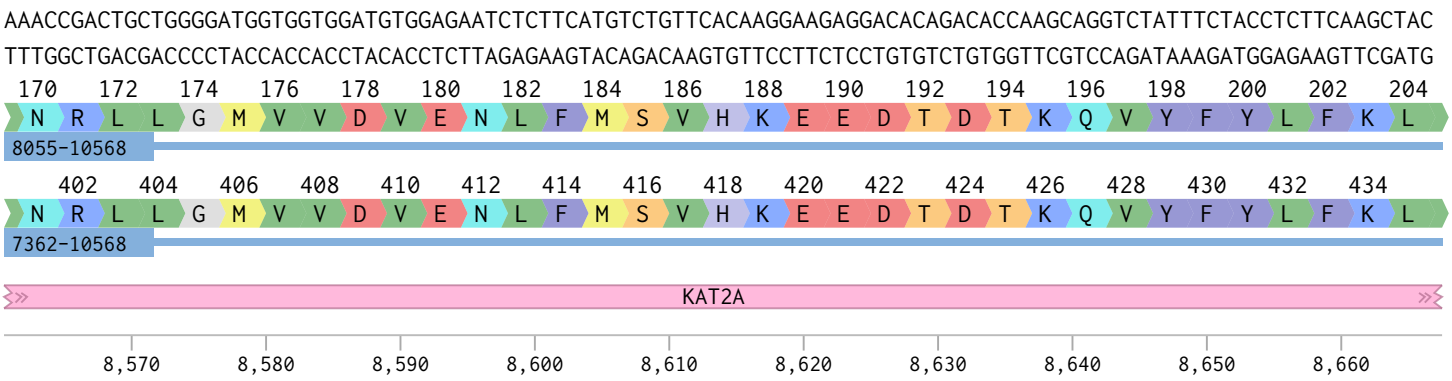
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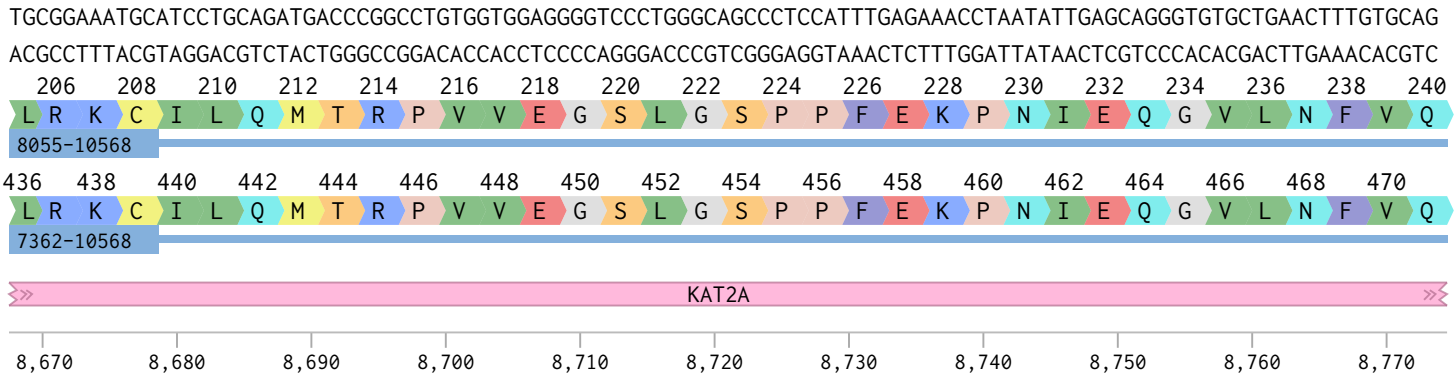
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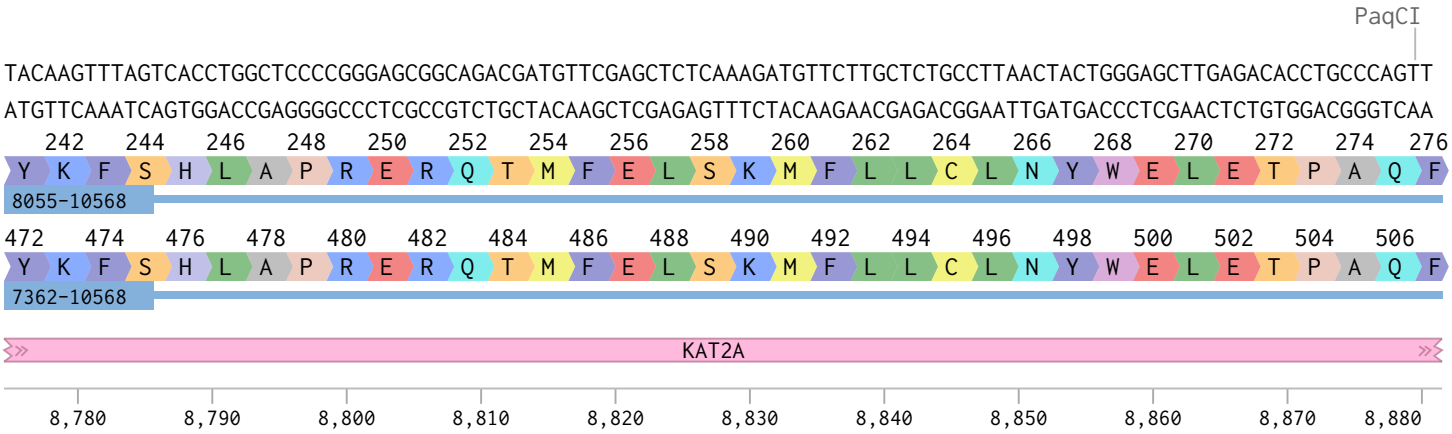
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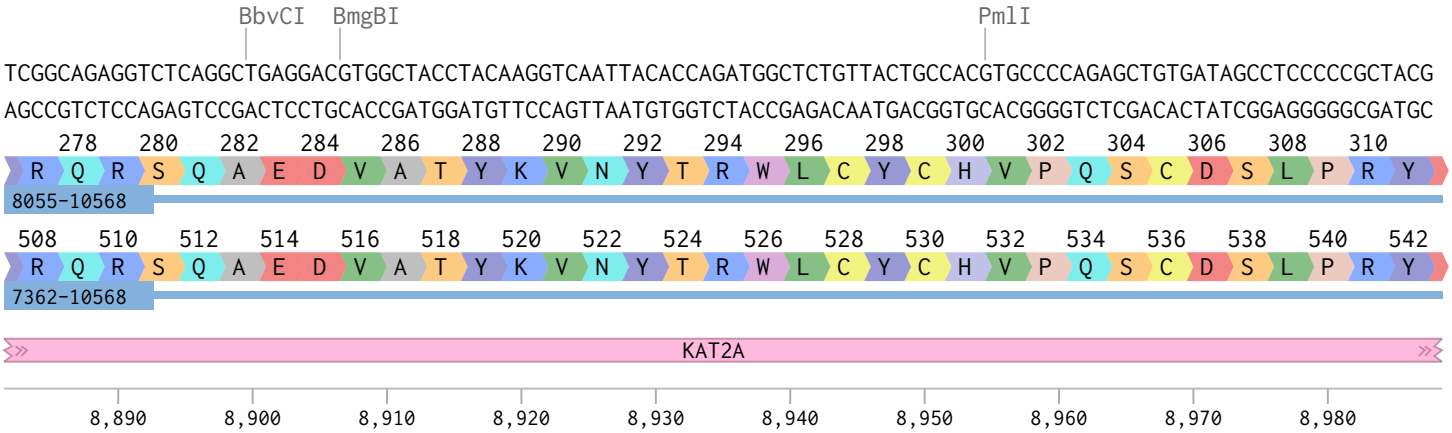
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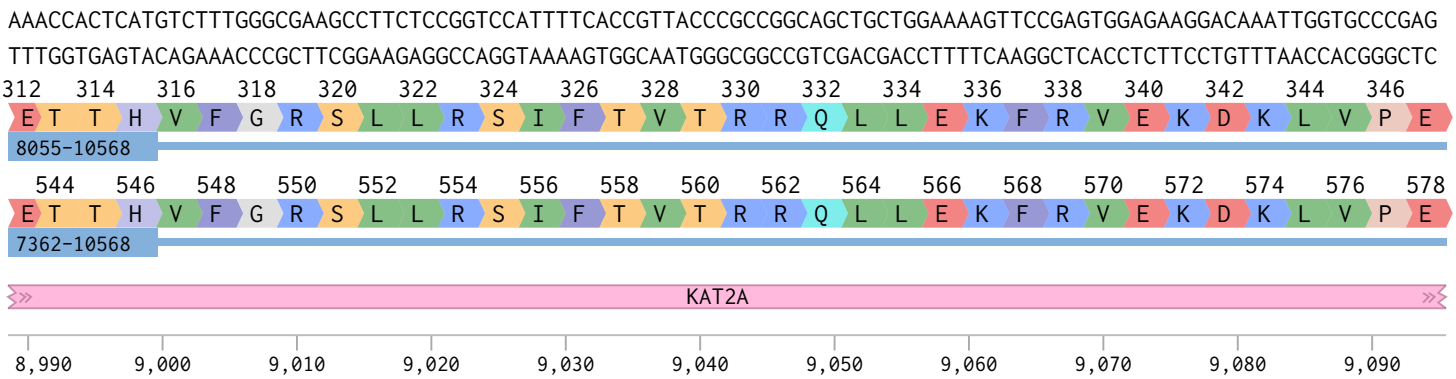
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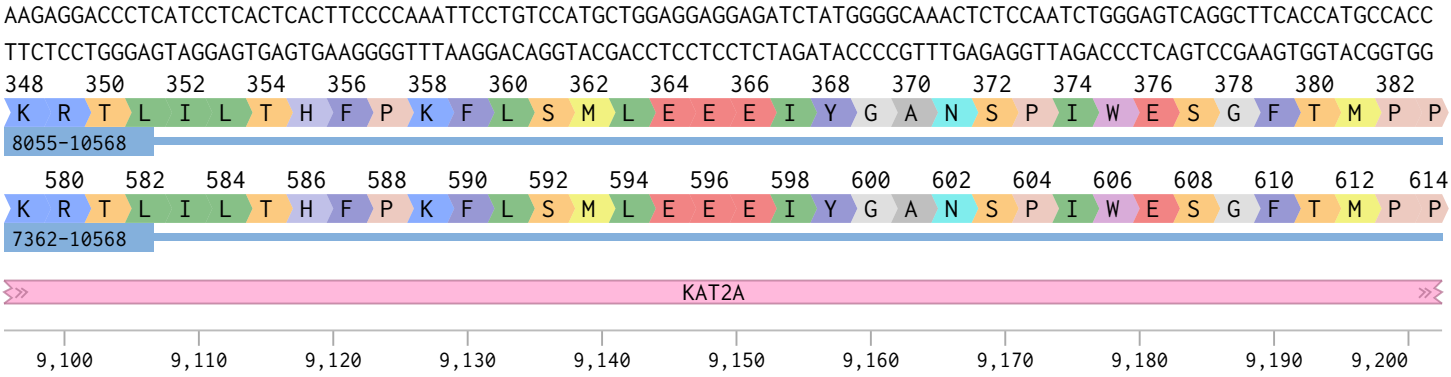
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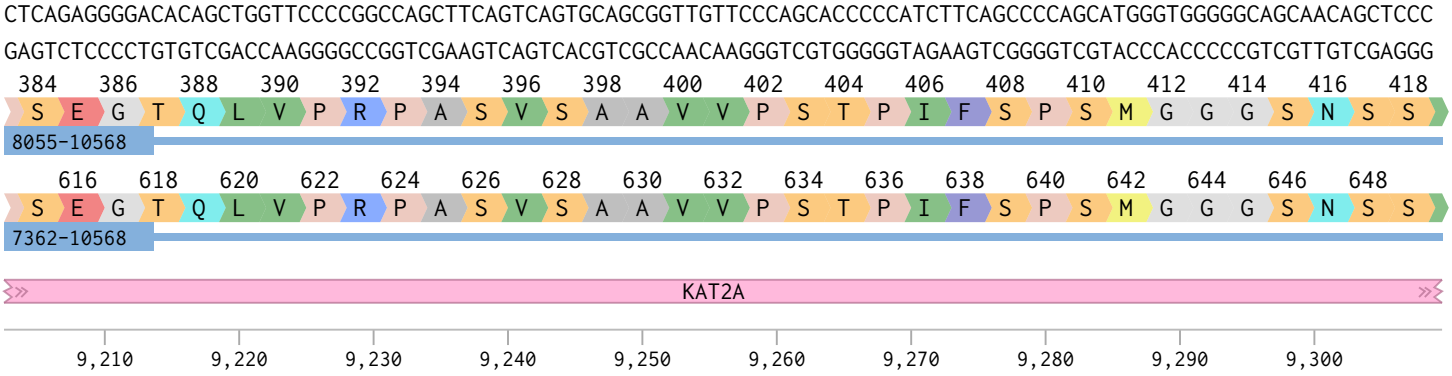
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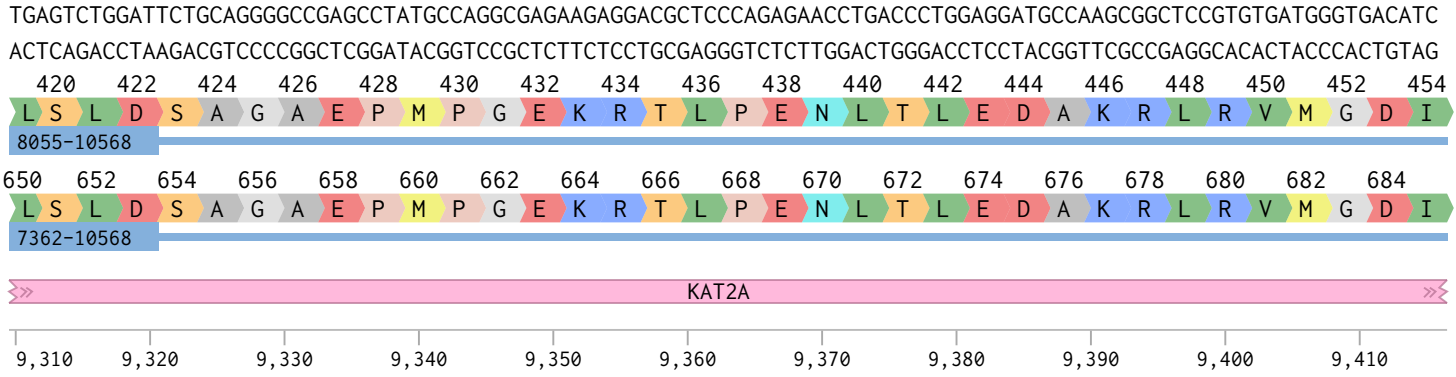
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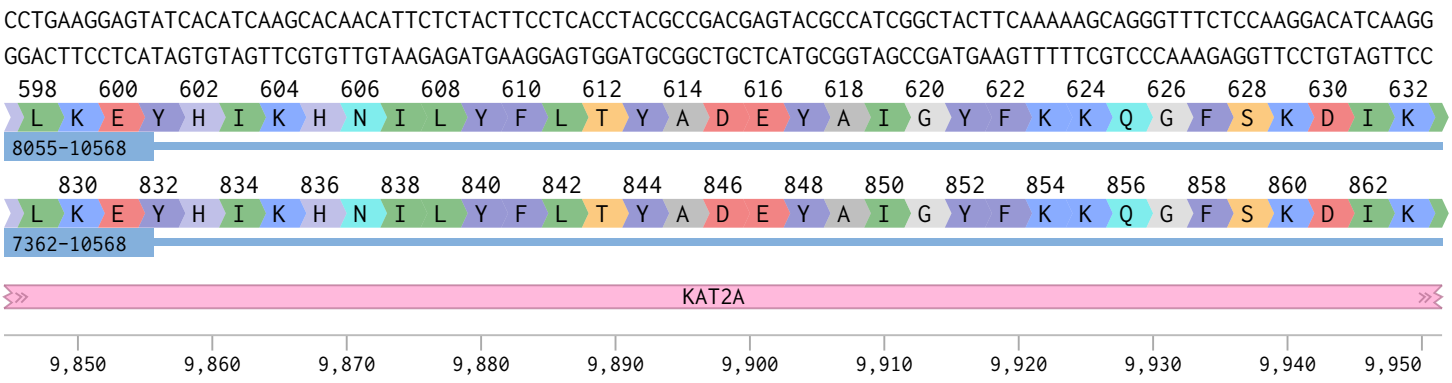
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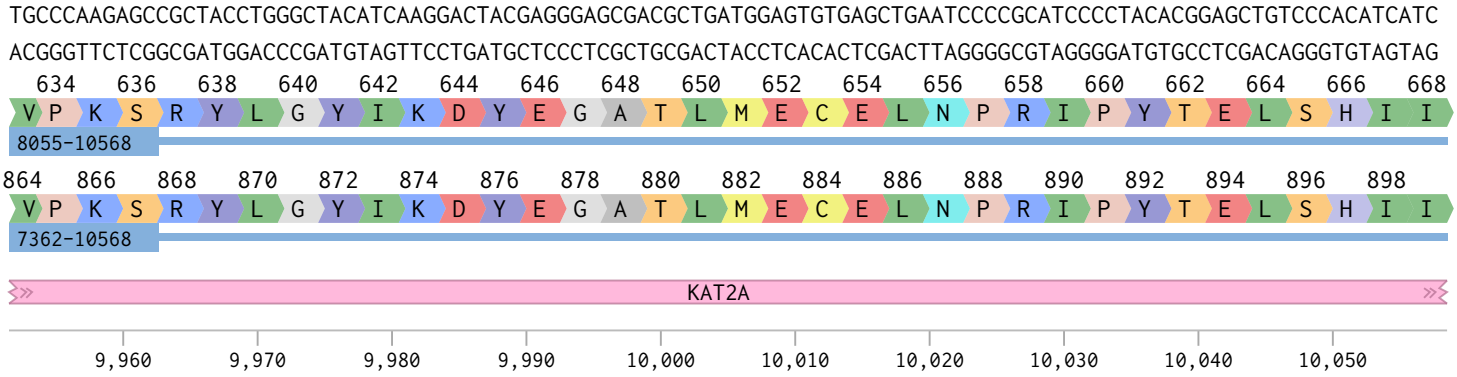
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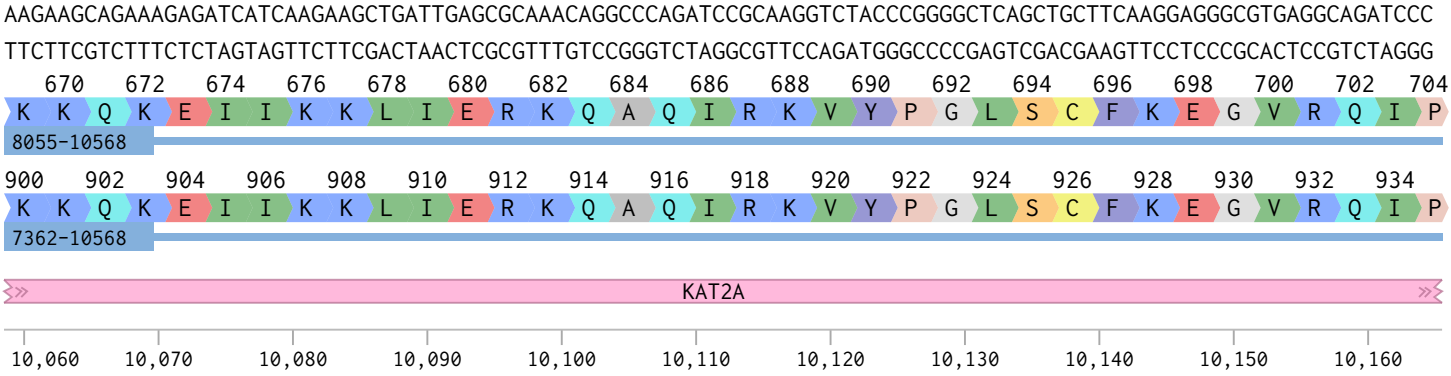
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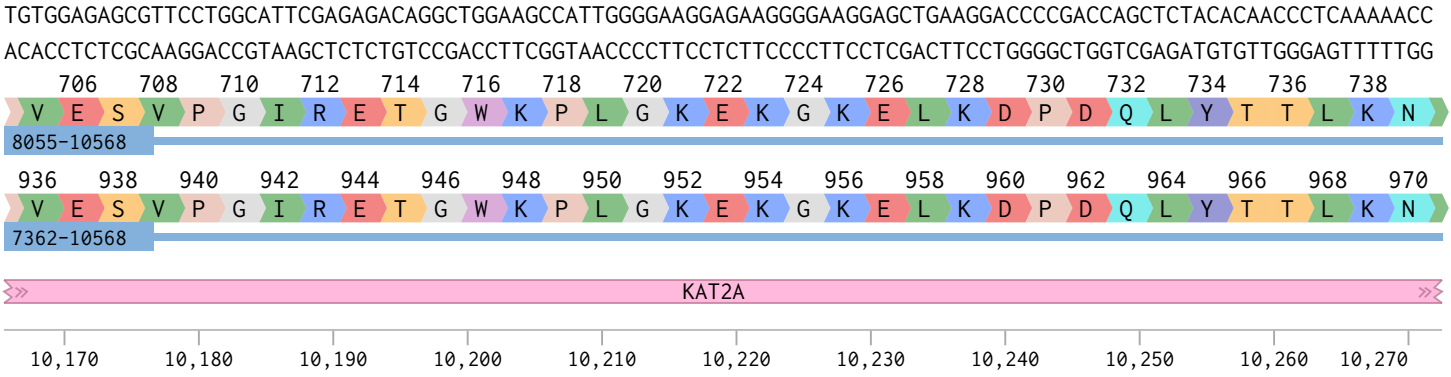
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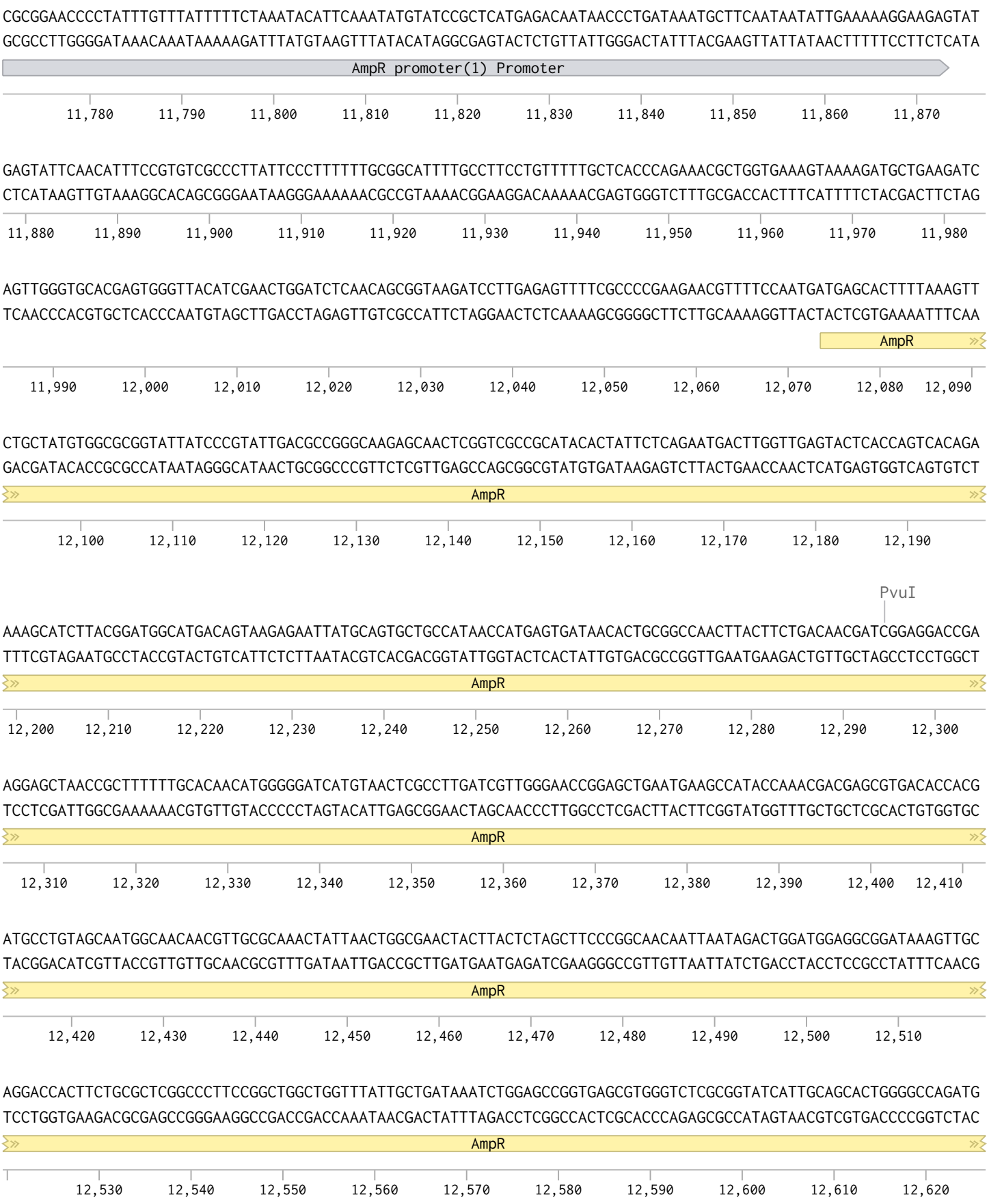
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AmpR
12,310 12,320 12,330 12,340 12,350 12,360 12,370 12,380 12,390 12,400 12,410
ATGCCTGTAGCAATGGCAACAACGTTGCGCAAACTATTAACTGGCGAACTACTTACTCTAGCTTCCCGGCAACAATTAATAGACTGGATGGAGGCGGATAAAGTTGC
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AmpR
12,420 12,430 12,440 12,450 12,460 12,470 12,480 12,490 12,500 12,510
AGGACCACTTCTGCGCTCGGCCCTTCCGGCTGGCTGGTTTATTGCTGATAAATCTGGAGCCGGTGAGCGTGGGTCTCGCGGTATCATTGCAGCACTGGGGCCAGATG
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AmpR
12,530 12,540 12,550 12,560 12,570 12,580 12,590 12,600 12,610 12,620

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CATTCGGGAGGGCATAGCATCAATAGATGTGCTGCCCCTCAGTCCGTTGATACCTACTTGCTTTATCTGTCTAGCGACTCTATCCACGGAGTGACTAATTCGTAACC

AmpR

12,630 12,640 12,650 12,660 12,670 12,680 12,690 12,700 12,710 12,720 12,730

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Co...n

12,740 12,750 12,760 12,770 12,780 12,790 12,800 12,810 12,820 12,830 12,840

AATCCCTTAACGTGAGTTTTCGTTCCACTGAGCGTCAGACCCCGTAGAAAAGATCAAAGGATCTTCTTGAGATCCTTTTTTTCTGCGCGTAATCTGCTGCTTGCAAA
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ColE1 origin

12,850 12,860 12,870 12,880 12,890 12,900 12,910 12,920 12,930 12,940

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ColE1 origin

12,950 12,960 12,970 12,980 12,990 13,000 13,010 13,020 13,030 13,040 13,050

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ColE1 origin

13,060 13,070 13,080 13,090 13,100 13,110 13,120 13,130 13,140 13,150 13,160

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ColE1 origin

13,170 13,180 13,190 13,200 13,210 13,220 13,230 13,240 13,250 13,260

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ColE1 origin

13,270 13,280 13,290 13,300 13,310 13,320 13,330 13,340 13,350 13,360 13,370

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ColE1 origin

13,380 13,390 13,400 13,410 13,420 13,430 13,440 13,450 13,460 13,470 13,480

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ColE1 origin

13,490 13,500 13,510 13,520 13,530 13,540 13,550 13,560 13,570 13,580

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13,600 13,610 13,620 13,630 13,640 13,650 13,660 13,670 13,680 13,690

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13,700 13,710 13,720 13,730 13,740 13,750 13,760 13,770 13,780 13,790 13,800

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Tn7R

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Tn7R

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Gent2

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Gent2

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Gent2

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Gent2
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Pc promoter Promoter
14,770 14,780 14,790 14,800 14,810 14,820 14,830 14,840 14,850 14,860 14,870
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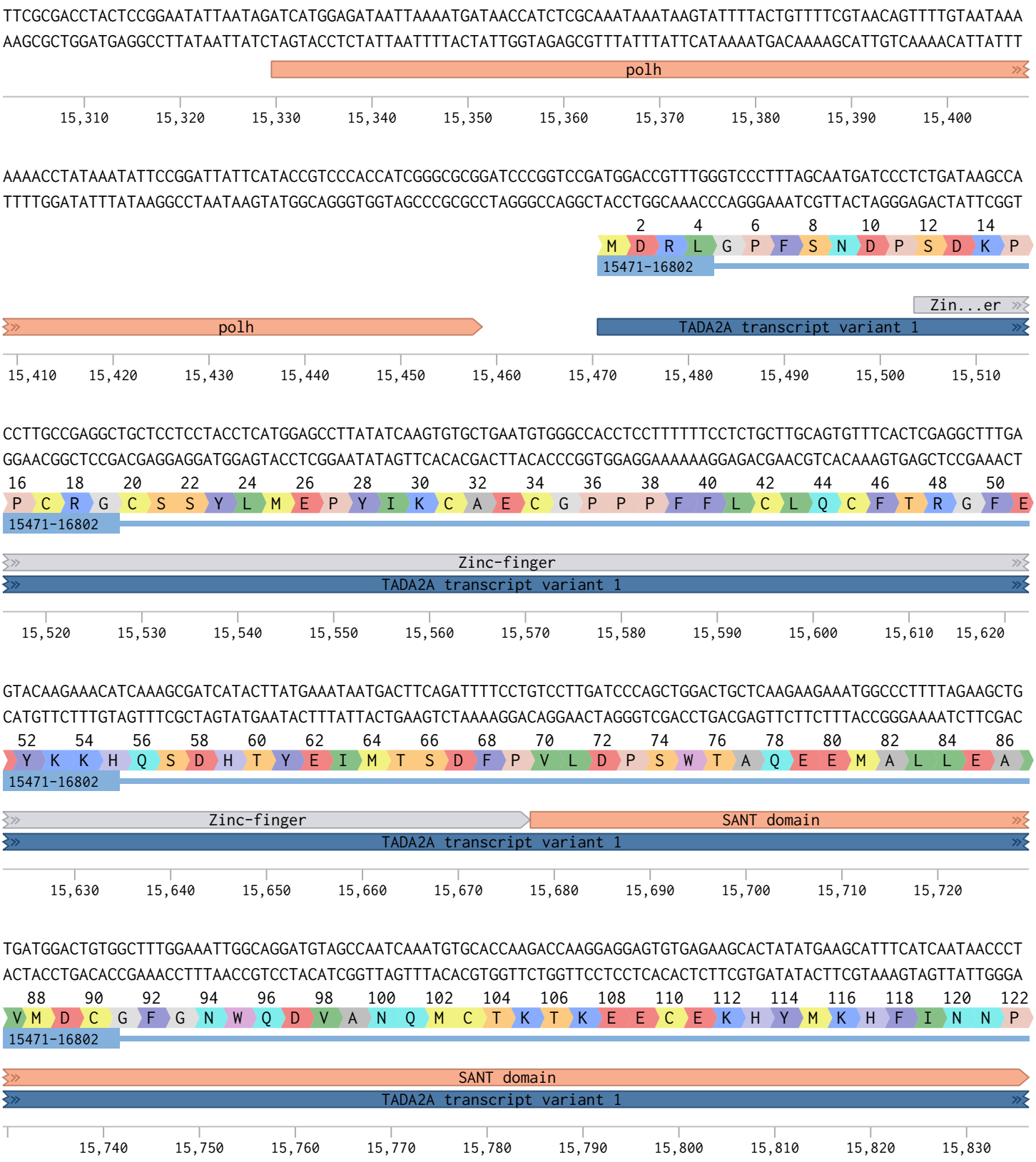
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15471-16802
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polh
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Zinc-finger
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Zinc-finger
SANT domain
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V M D C G F G N W Q D V A N Q M C T K T K E E C E K H Y M K H F I N N P
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SANT domain
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124 126 128 130 132 134 136 138 140 142 144 146 148 150 152 154 156 158
L F A S T L L N L K Q A E E A K T A D T A I P F H S T D D P P R P T F D
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194 196 198 200 202 204 206 208 210 212 214 216 218 220 222 224 226 228
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230 232 234 236 238 240 242 244 246 248 250 252 254 256 258 260 262 264
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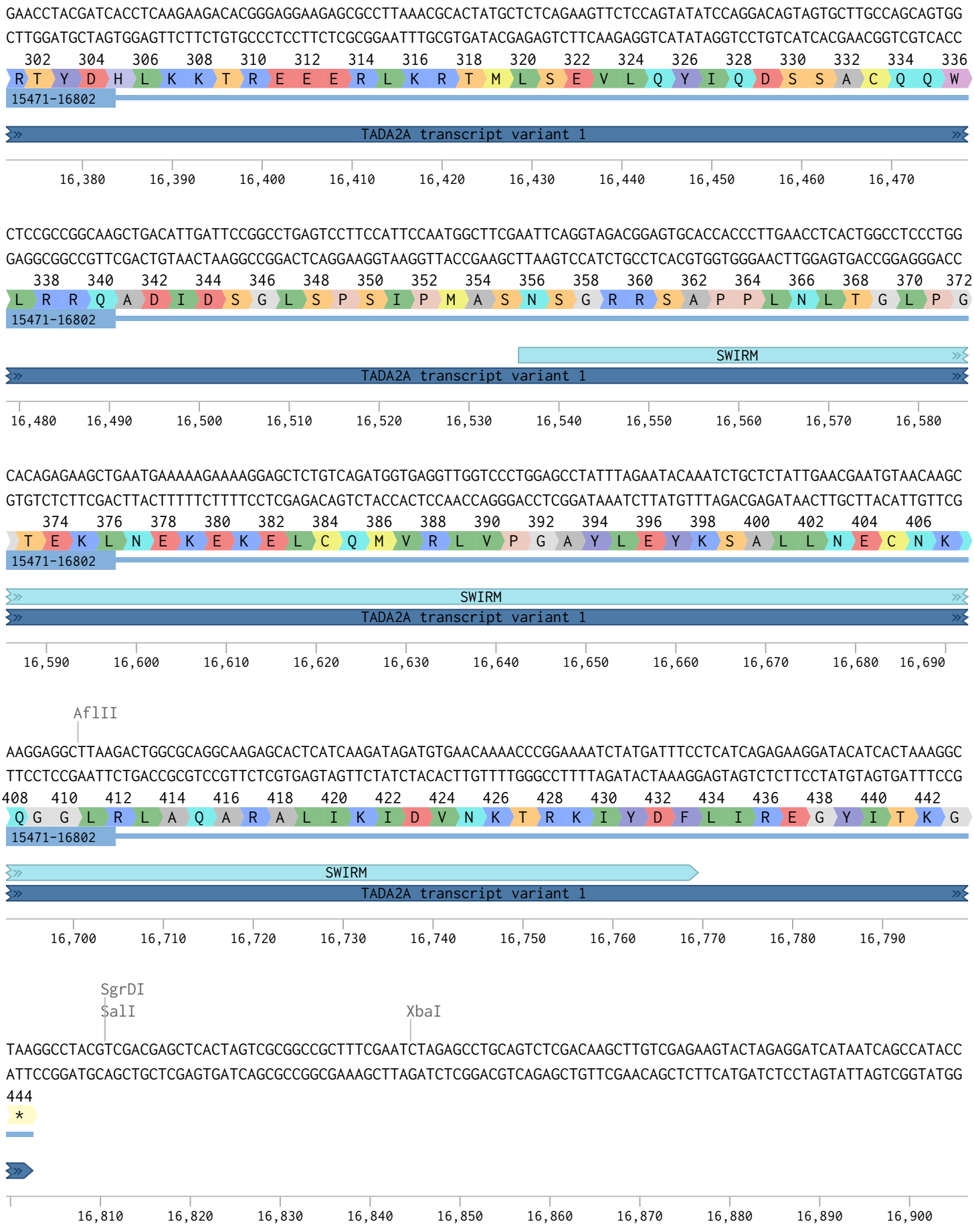
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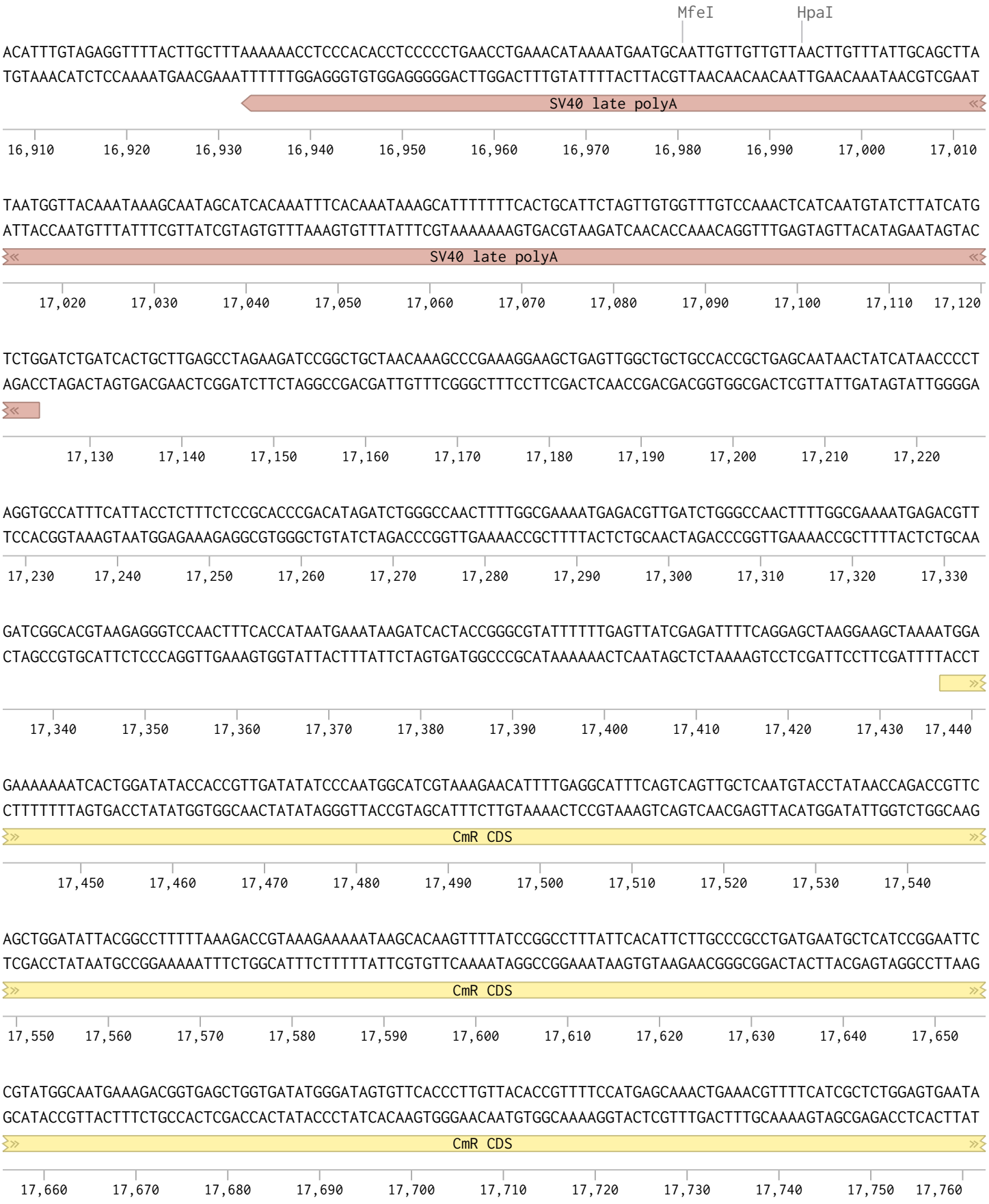
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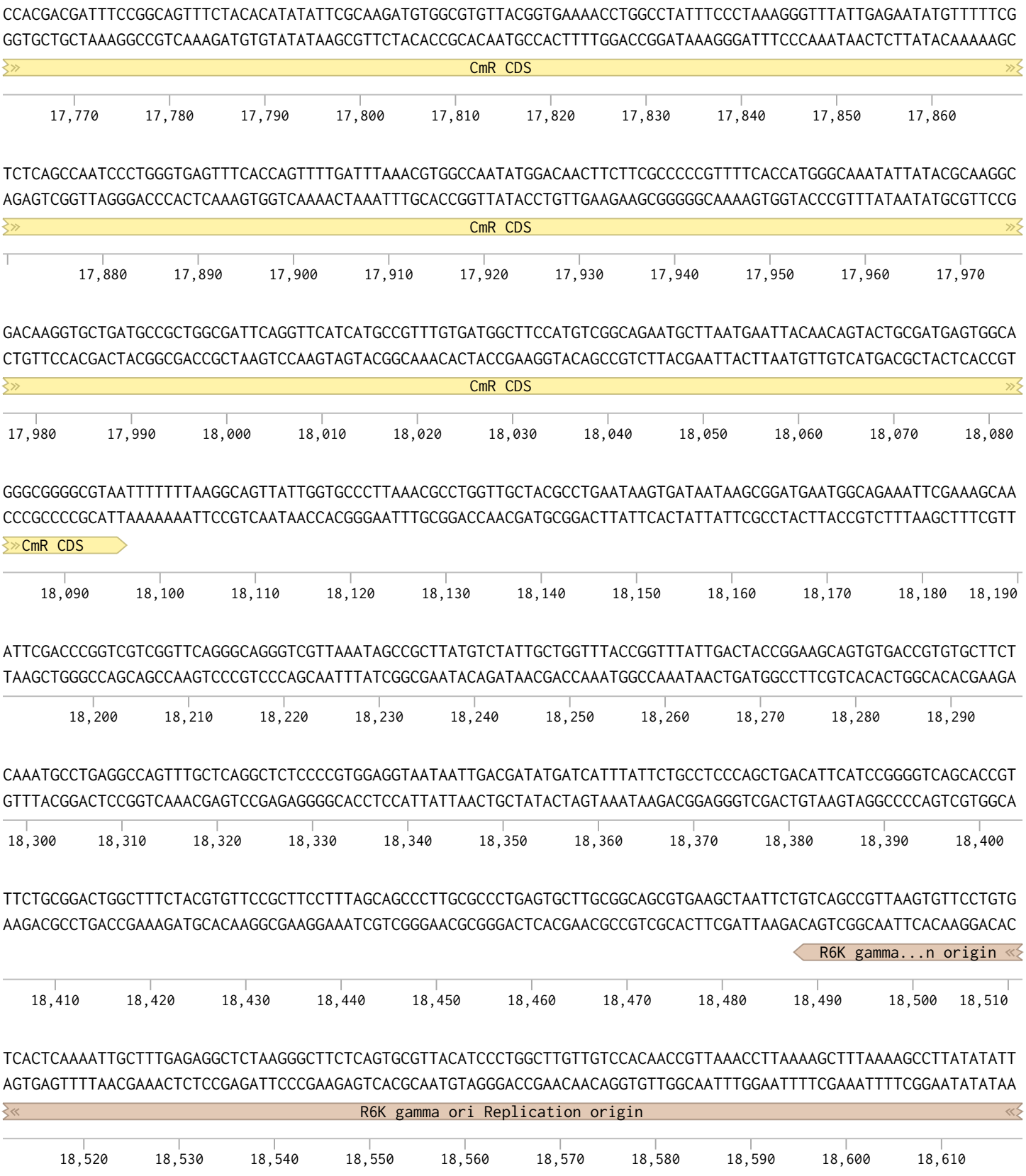
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17,770 17,780 17,790 17,800 17,810 17,820 17,830 17,840 17,850 17,860
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R6K gamma...n origin
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R6K gamma ori Replication origin
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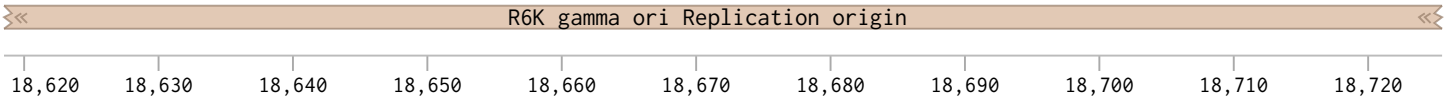

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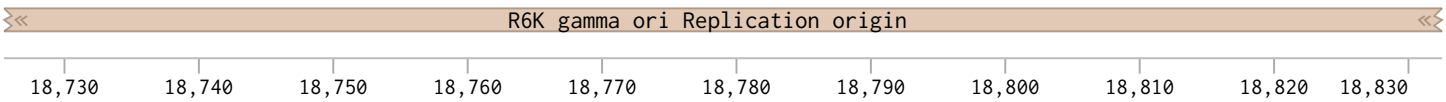

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