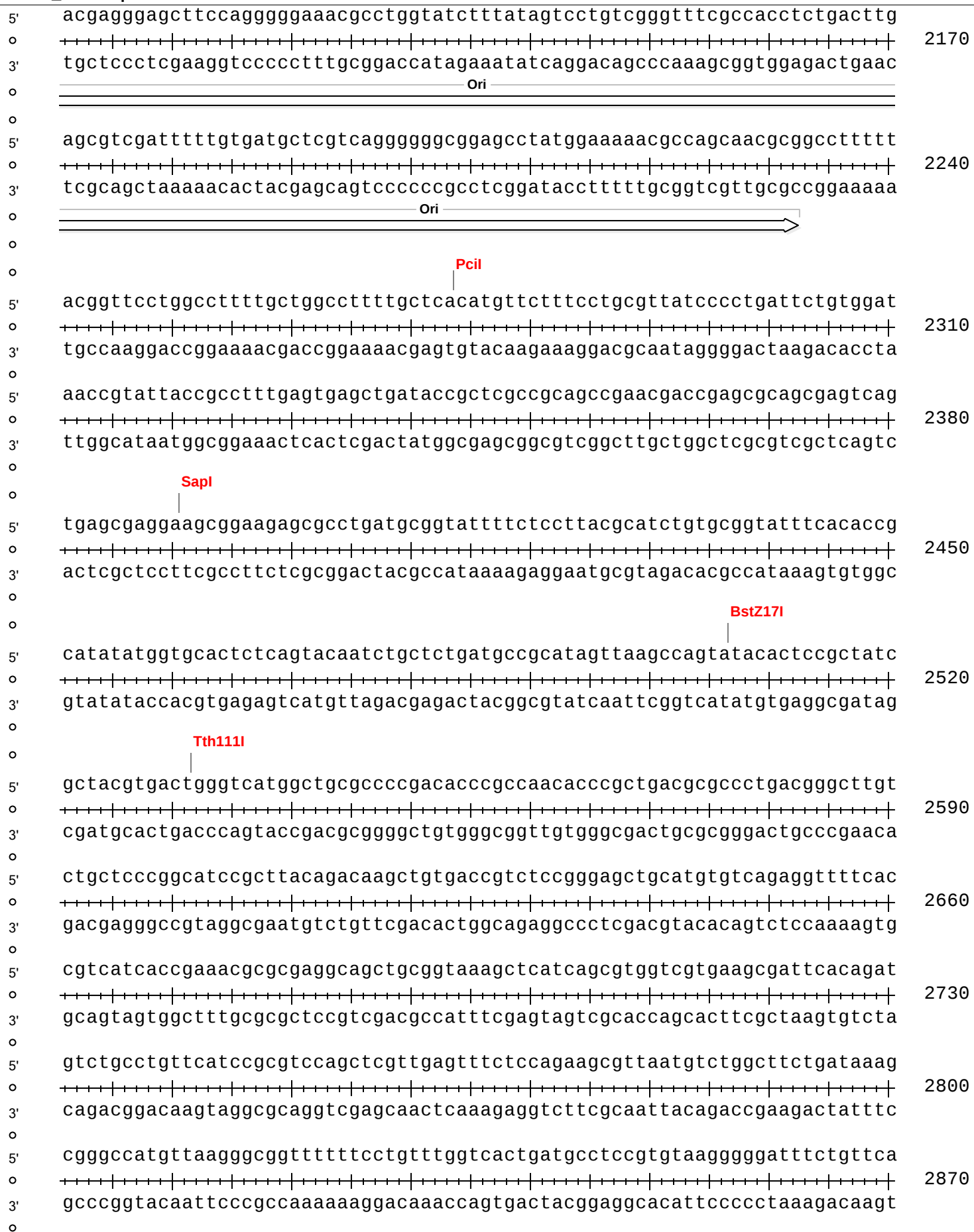


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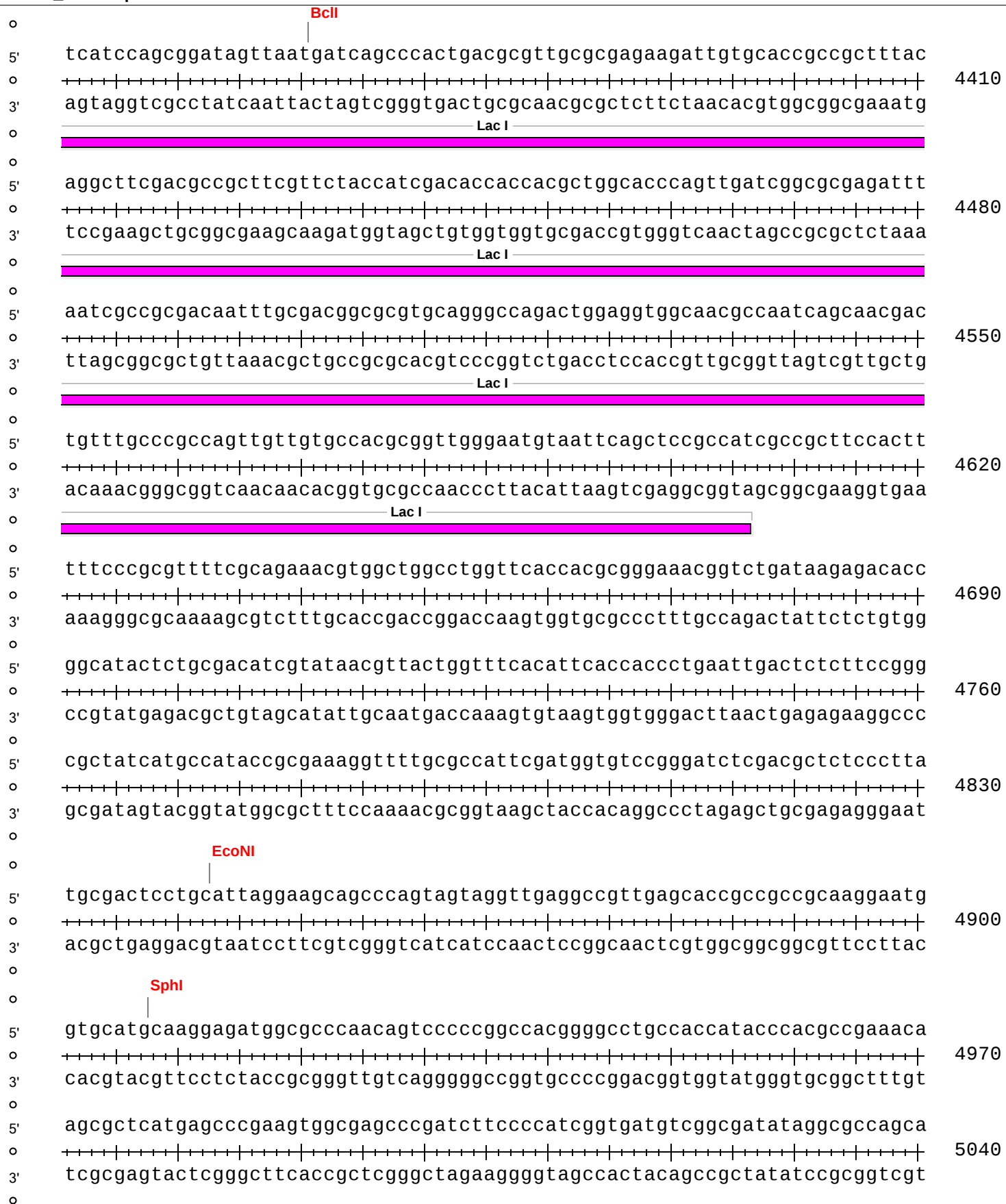
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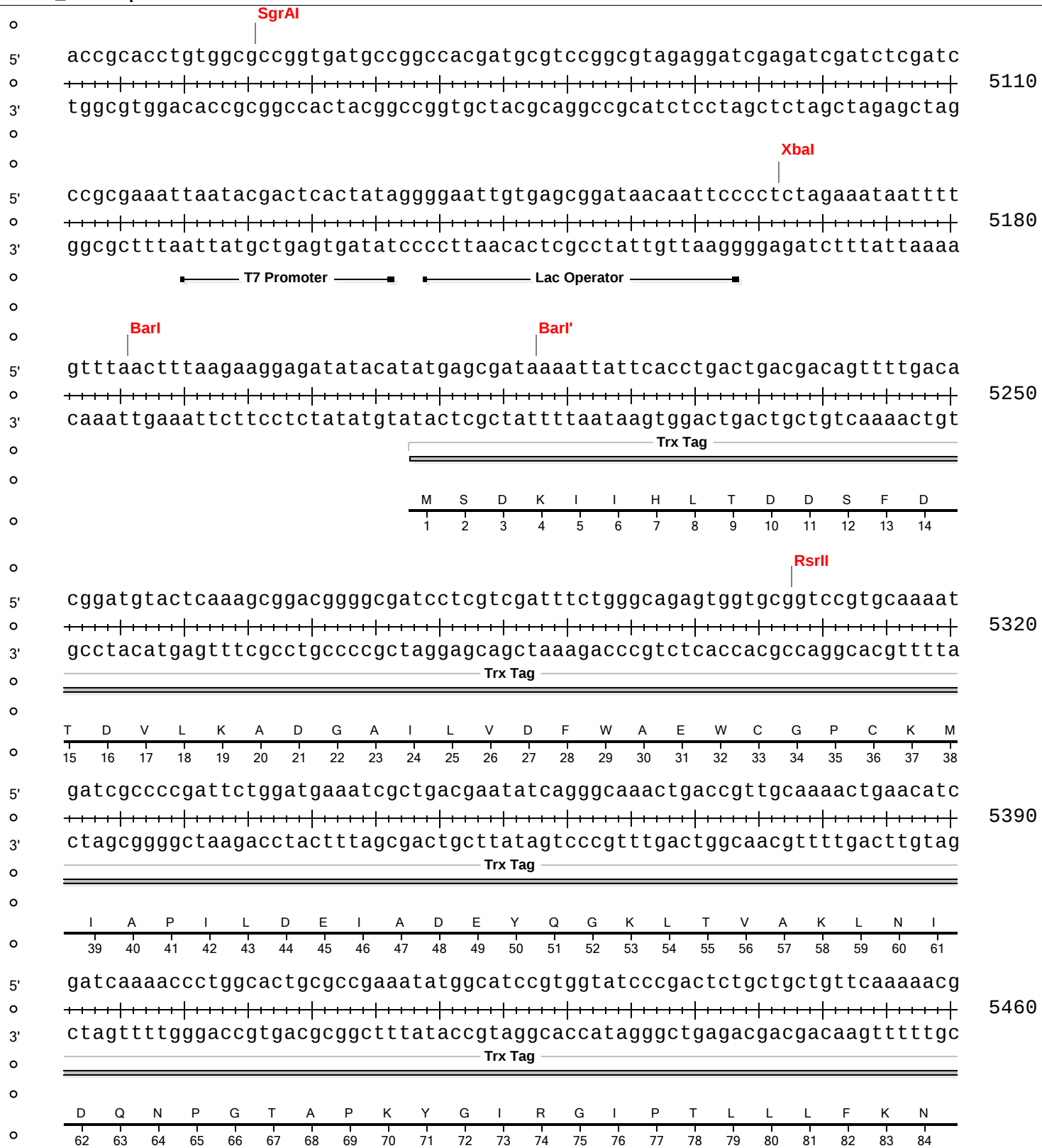
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←

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Lac I  
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Lac I  
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Lac I  
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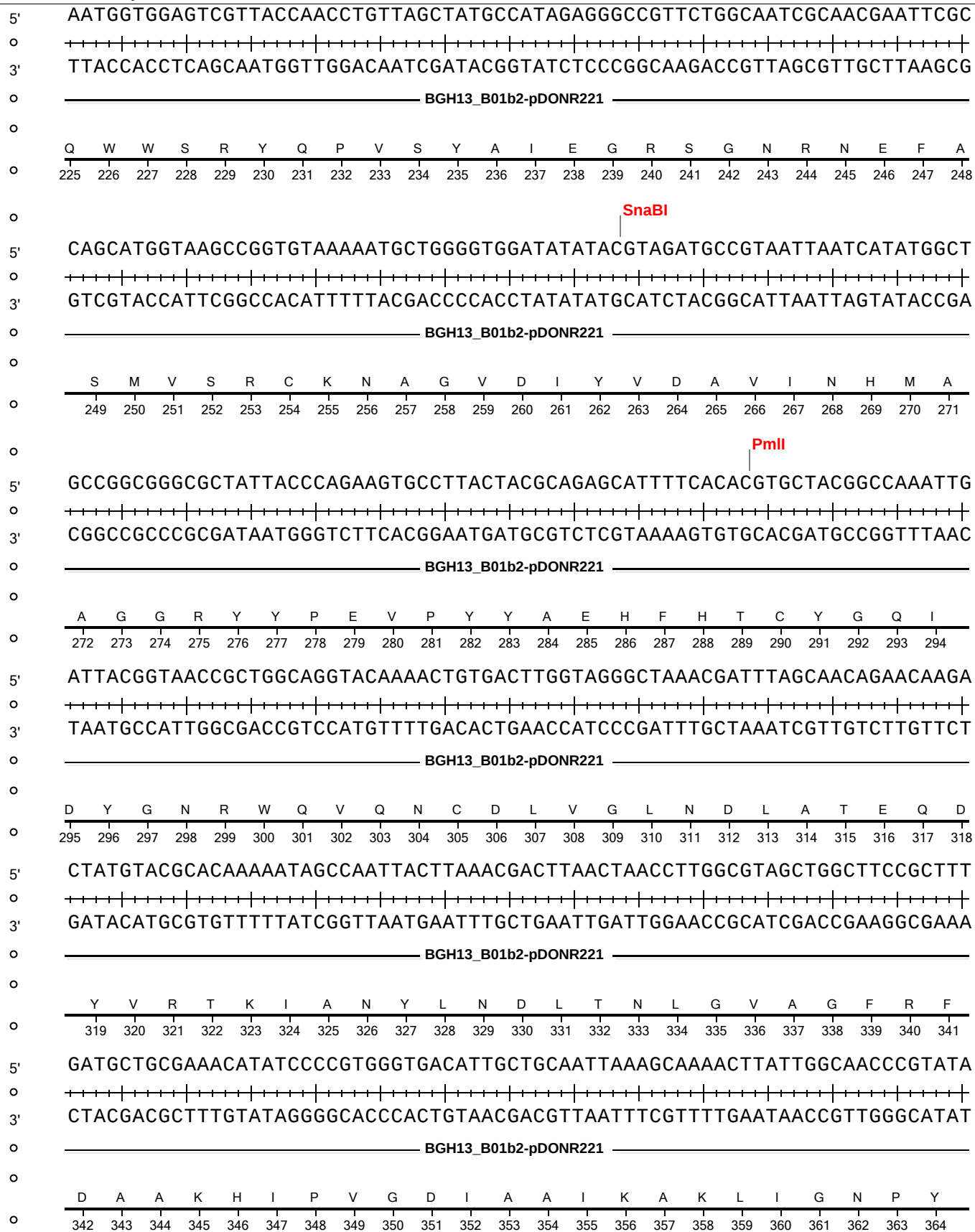
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Trx Tag  
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G E V A A T K V G A L S K G Q L K E F L D A N L  
85 86 87 88 89 90 91 92 93 94 95 96 97 98 99 100 101 102 103 104 105 106 107 108

The diagram illustrates a DNA sequence with a BglII restriction site and an enzyme binding site. The DNA sequence is shown in two orientations: 5' to 3' (top) and 3' to 5' (bottom). The 5' sequence is: ggtatgaaagaaaccgctgctgctaaattcgaacgccagcacatggacagcccagatctgggtaccgacg. The 3' sequence is: ccatactttctttggcgacgacgatttaagcttgcggtcgtgtacctgtcgggtctagacccatggctgc. A BglII restriction site is indicated by a vertical line at position 149, with the label "BglII" in red. An enzyme binding site is indicated by a pink box labeled "En...e" at the end of the sequence. Below the DNA sequence, a protein sequence is shown: G M K E T A A A K F E R Q H M D S P D L G T D, with positions 132 to 154 indicated below each amino acid.

5' aggcCAACCGCGAACCgcGTTTCGTTcATTtTGTTcGAATGGCAGTGGAAcGATATTGCTAGcGAGTGCgAA  
 3' tccgGTTGGCGCTTGGCGCAAGCAAGTAACAAGCTTACCGTCACCTTGCTATAACGATCGCTCACGCTT  
 BGH13\_B01b2-pDONR221  
 G Q P R T A F V H L F E W Q W N D I A S E C E  
 179 180 181 182 183 184 185 186 187 188 189 190 191 192 193 194 195 196 197 198 199 200 201

5' AATGTACTGGGGCCAAAAGGCTATGCGGCAGTGCAGGTATCCCCACCCCAAAAATCGGTTAGTAGCAGCC  
 3' TTACATGACCCCGGTTTTCCGATACGCCGTCACGTCCATAGGGGTGGGGTTTTTAGCCAATCATCGTCGG  
 BGH13\_B01b2-pDONR221  
 N V L G P K G Y A A V Q V S P P Q K S V S S S  
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## BGH13\_B01b2-pET32-DEST



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3' ATAAGGTTCTTCACTATCCACGCGGCCGCTCGGGTAGGTTGGTAGTCTTATATGGATGAATCGTCTGCA

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I F Q E V I G A P G E P I Q P S E Y T Y L A D V  
365 366 367 368 369 370 371 372 373 374 375 376 377 378 379 380 381 382 383 384 385 386 387 388

5' AACAGAGTTTAACTTTACCAAATCGGTTAGCCATTTTTTTAAAGGGCGCGGACCAATTAAAGAGTTAAAA  
+-----+  
3' TTGTCTCAAATTGAAATGGTTTAGCCAATCGGTAAAAAATTTCCTCGCGCCTGGTTAATTTCTCAATTTT

BGH13\_B01b2-pDONR221

T E F N F T K S V S H F F K G R G P I K E L K  
389 390 391 392 393 394 395 396 397 398 399 400 401 402 403 404 405 406 407 408 409 410 411

5' AATATTGGTACTTGGGGCGGTTGGGTTGCAAGTGCAGACGCCGTAACCTTTGTTGCCAACCCACGATAACC  
+-----+  
3' TTATAACCATGAACCCCGCCAACCCAACGTTACGTCTGCGGCATTGGAACAACGGTTGGTGCTATTGG

BGH13\_B01b2-pDONR221

N I G T W G G W V A S A D A V T F V A N H D N  
412 413 414 415 416 417 418 419 420 421 422 423 424 425 426 427 428 429 430 431 432 433 434

5' AGCGCCAGGATACAAACAACACCATTACCCATAAAGATGGTGATAACCGCAATACTATGGCGCACGTATT  
+-----+  
3' TCGCGGTCCTATGTTTGTGTGGTAATGGGTATTTCTACCACTATTGGCGTTATGATACCGCGTGCATAA

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Q R Q D T N N T I T H K D G D N R N T M A H V F  
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5' TATGCTGGGCTGGCCCTATGGGTACCCTAAAGTAATGTCTAGTTATGAGTGGTACGATCACGACCAAGGC  
+-----+  
3' ATACGACCCGACCGGGATACCCATGGGATTTTATTACAGATCAATACTCACCATGCTAGTGCTGGTTCCG

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M L G W P Y G Y P K V M S S Y E W Y D H D Q G  
459 460 461 462 463 464 465 466 467 468 469 470 471 472 473 474 475 476 477 478 479 480 481

5' CCTCCCGCCCACGGTGCTAGCAGCTGTAACAGCGGCTGGTTATGTGAACACCGCACGCCAGCTATCGCCA  
+-----+  
3' GGAGGGCGGGTGCCACGATCGTCGACATTGTCGCCGACCAATACACTTGTGGCGTGCGGTCGATAGCGGT

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P P A H G A S S C N S G W L C E H R T P A I A  
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## BGH13\_B01b2-pET32-DEST

