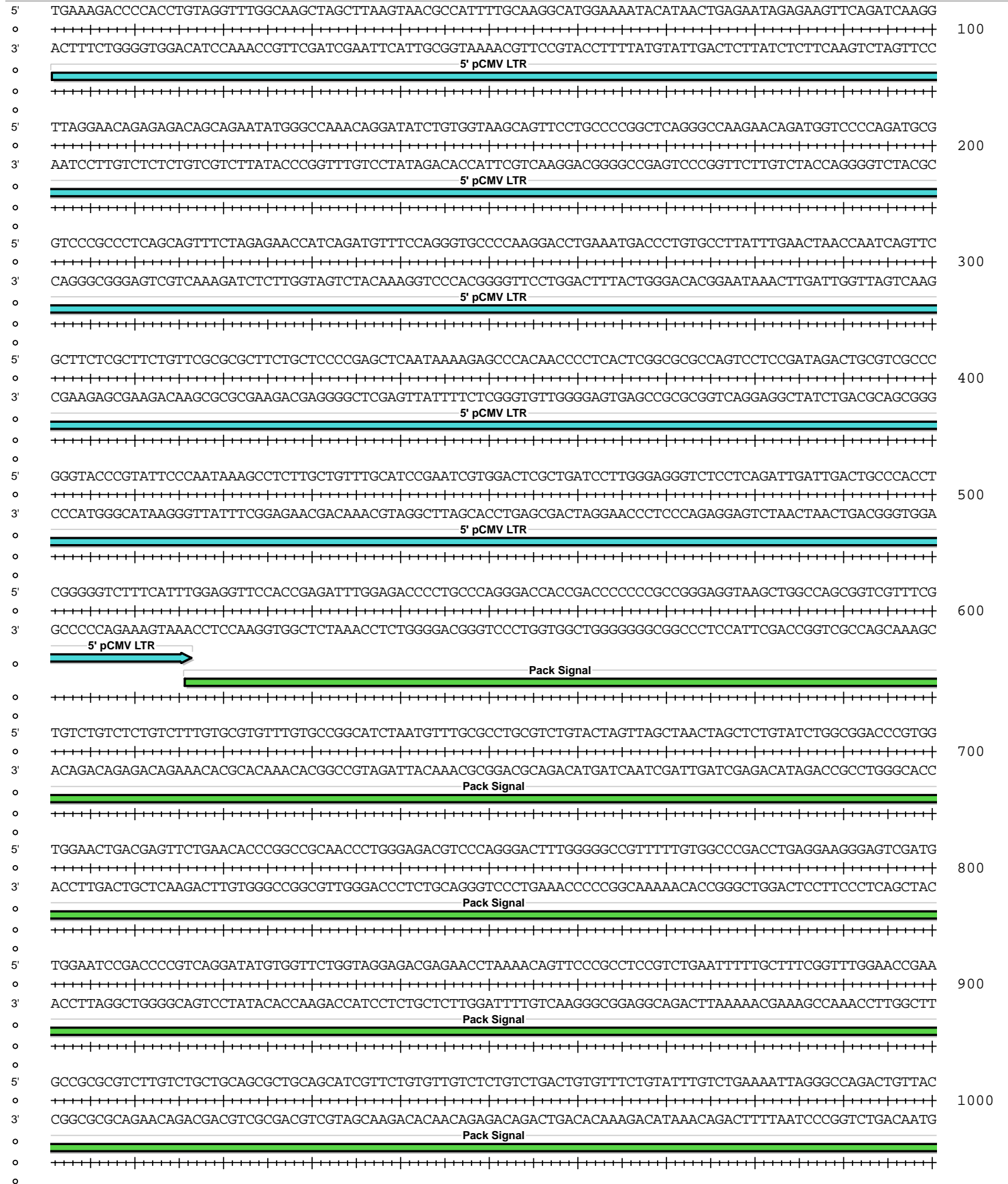
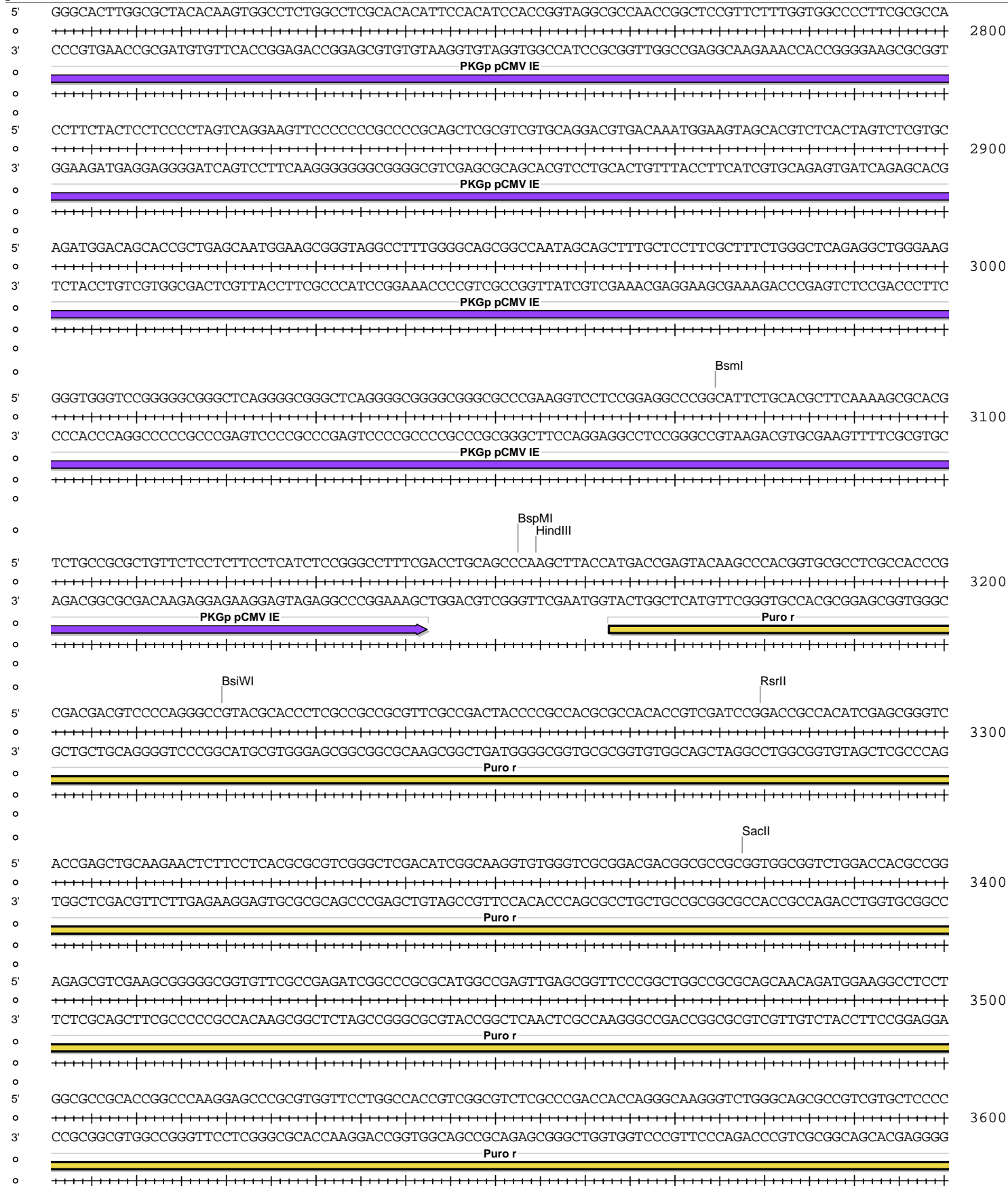


Absent Sites	0	AarI, AbsI, Accl, AjuI, AjuI', AlfI, AlfI', ApaI, AsiSI, AvrII, BamHI, BarI, BarI', BclI, BsaAI, BstBI, BstXI, BstZ17I, CspCI, CspCI', FseI, FspAI, HincII, HpaI, MauBI, MfeI, MluI, MreI, NruI, PacI, PfiMI, PmeI, PmlI, PshAI, PspOMI, PstI, PstI', Sall, SanDI, SbfI, SfiI, SgrDI, SnaBI, SrfI, SwaI, XcmI
AfIII	1	4887
Arsl	1	1732
Arsl'	1	1700
BbsI	1	2497
BglIII	1	1411
BsaBI	1	2336
BsiWI	1	3220
BsmI	1	3075
BspMI	1	3153
BtgZI	1	1559
Clal	1	3818
DrallI	1	3735
HindIII	1	3155
NdeI	1	6951
NotI	1	2158
NsiI	1	3817
PciI	1	4887
PsiI	1	2192
PspXI	1	2617
RsrII	1	3280
SacII	1	3378
Scal	1	6260
SgrAI	1	7323
XhoI	1	2617

















o  
5' GGCTGGCTTAACTATGCGGCATCAGAGCAGATTGTACTGAGAGTGCACCATATGCGGTGTGAAATACCGCACAGATGCGTAAGGAGAAAATACCGCATCA  
o ++++++  
3' CCGACCGAATTGATACGCCGTAGTCTCGTCTAACATGACTCTCACGTGGTATACGCCACACTTTATGGCGTGTCTACGCATTCTCTTTTATGGCGTAGT  
o ++++++ 7000  
o  
5' GGCGCCATTTCGCCATTTCAGGCTGCGCAACTGTTGGGAAGGGCGATCGGTGCGGGCCTCTTCGCTATTACGCCAGCTGGCGAAAGGGGGATGTGCTGCAAG  
o ++++++ 7100  
3' CCGCGGTAAGCGGTAAGTCCGACGCGTTGACAACCCCTTCCCGCTAGCCACGCCGAGAAGCGATAATGCGGTGACCCGCTTCCCCCTACACGACGTTT  
o ++++++  
o  
5' GCGATTAAGTTGGGTAACGCCAGGGTTTTCCAGTCACGACGTTGTAAAACGACGGCGCAAGGAATGGTGCATGCAAGGAGATGGCGCCCAACAGTCCCC  
o ++++++ 7200  
3' CGCTAATTCAACCCATTGCGGTCCCAAAGGGTCAGTGTGCAACATTTTGTCTGCCGCTTCTTACCACGTACGTTCTCTACCGCGGGTTGTCCAGGGG  
o ++++++  
o  
5' CGGCCACGGGGCCTGCCACCATACCCACGCCGAAACAAGCGCTCATGAGCCCGAAGTGGCGAGCCCGATCTTCCCATCGGTGATGTCCGGCATATAGGC  
o ++++++ 7300  
3' GCCGGTGCCCGGACGGTGGTATGGGTGCGGCTTTGTTTCGCGAGTACTCGGGCTTACCAGCTCGGGCTAGAAGGGGTAGCCACTACAGCCGCTATATCCG  
o ++++++  
o  
o  
o  
SgrAI  
5' GCCAGCAACCGCACCTGTGGCGCCGGTGTATGCCGGCCACGATGCGTCCGGCTAGAGGCGATTAGTCCAATTTGTTAAAGACAGGATATCAGTGGTCCAG  
o ++++++ 7400  
3' CCGTCGTTGGCGTGGACACCGCGCCACTACGGCCGGTGTCTACGAGCCGCATCTCCGCTAATCAGGTTAAACAATTTCTGTCTATAGTCACCAGGTC  
o ++++++  
o  
5' GCTCTAGTTTTGACTCAACAATATCACCAGCTGAAGCCTATAGAGTACGAGCCATAGATAAAAATAAAAGATTTTATTTAGTCTCCAGAAAAGGGGGAA  
o ++++++ 7500  
3' CGAGATCAAAACTGAGTTGTTATAGTGGTGCACCTTCGGATATCTCATGCTCGGTATCTATTTTATTTTCTAAAATAAATCAGAGGTCTTTTCCCCCTT  
o ++++++  
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