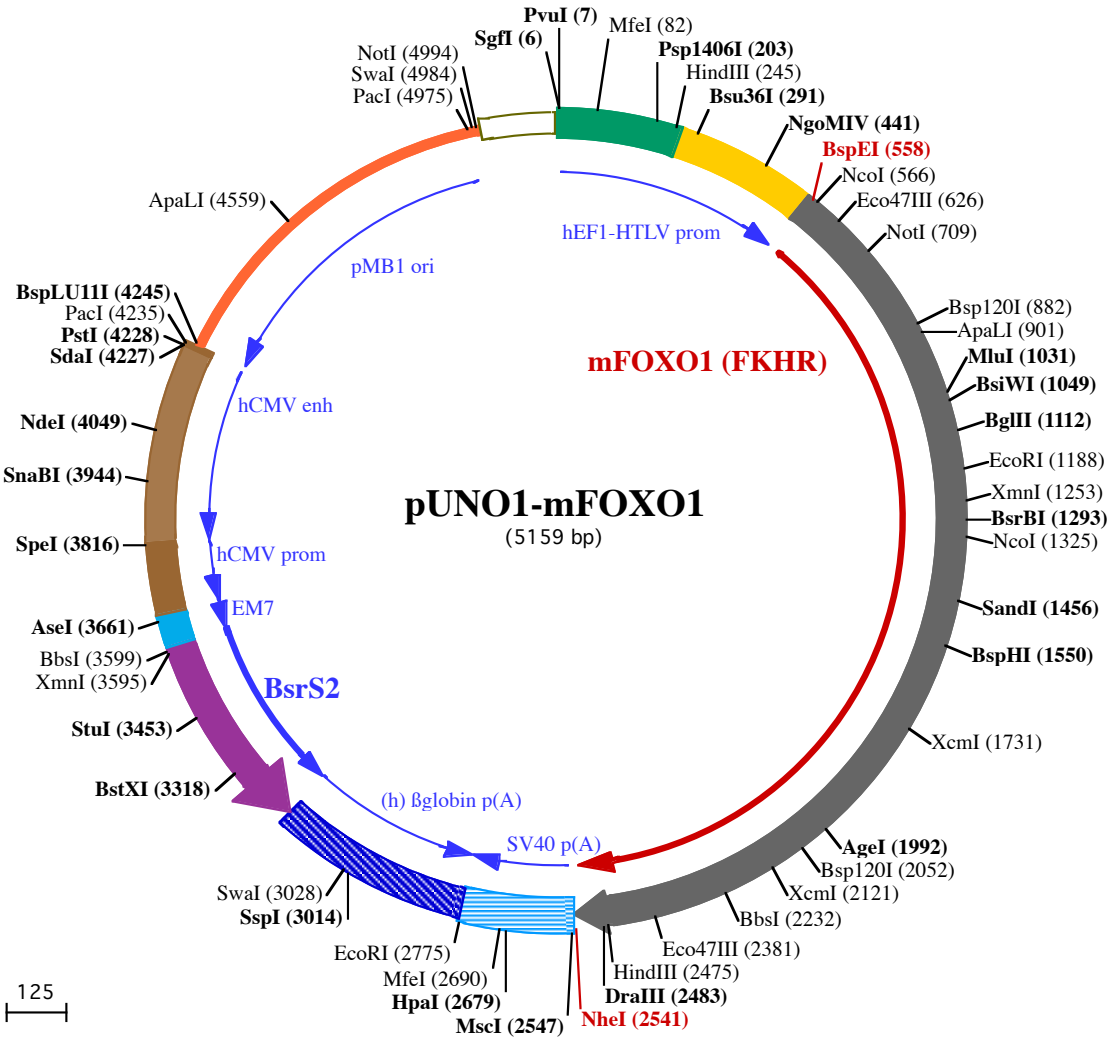
pUNO1-mFOXO1
(5159 bp)
mFOXO1 (FKHR)
hEF1-HTLV prom
pMB1 ori
hCMV enh
hCMV prom
EM7
BsrS2
(h) ßglobin p(A)
SV40 p(A)
PvuI (7)
SgfI (6)
MfeI (82)
Psp1406I (203)
HindIII (245)
Bsu36I (291)
NgoMIV (441)
BspEI (558)
NcoI (566)
Eco47III (626)
NotI (709)
Bsp120I (882)
ApaLI (901)
MluI (1031)
BsiWI (1049)
BglII (1112)
EcoRI (1188)
XmnI (1253)
BsrBI (1293)
NcoI (1325)
SandI (1456)
BspHI (1550)
XcmI (1731)
AgeI (1992)
Bsp120I (2052)
XcmI (2121)
BbsI (2232)
Eco47III (2381)
HindIII (2475)
DraIII (2483)
NheI (2541)
MscI (2547)
HpaI (2679)
MfeI (2690)
EcoRI (2775)
SspI (3014)
SwaI (3028)
StuI (3453)
BstXI (3318)
BbsI (3599)
XmnI (3595)
AseI (3661)
SpeI (3816)
SnaBI (3944)
NdeI (4049)
BspLU11I (4245)
PacI (4235)
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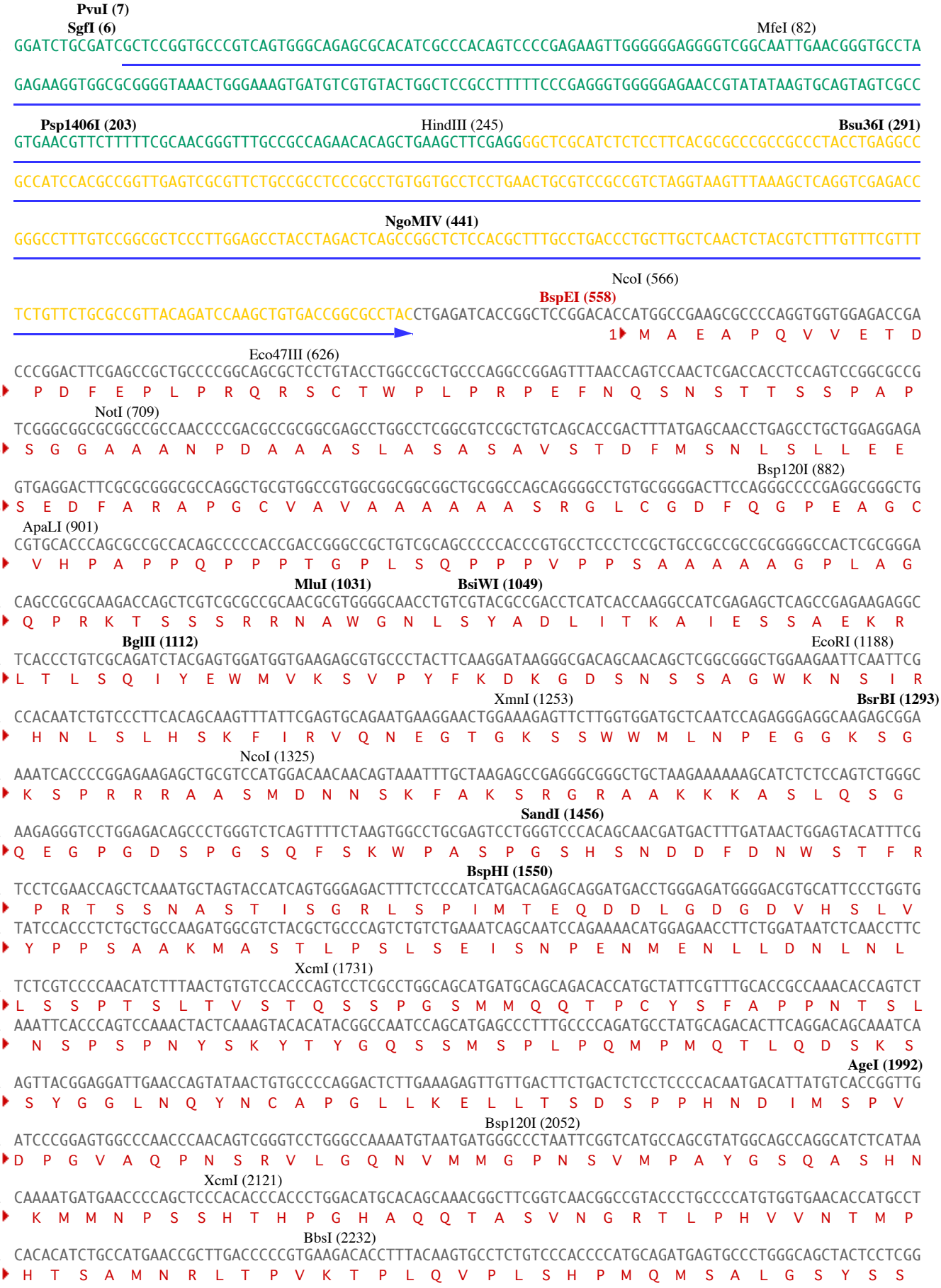
PacI (4975)
125


PvuI (7)
SgfI (6)
MfeI (82)
Psp1406I (203)
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BglII (1112)
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XmnI (1253)
BsrBI (1293)
NcoI (1325)
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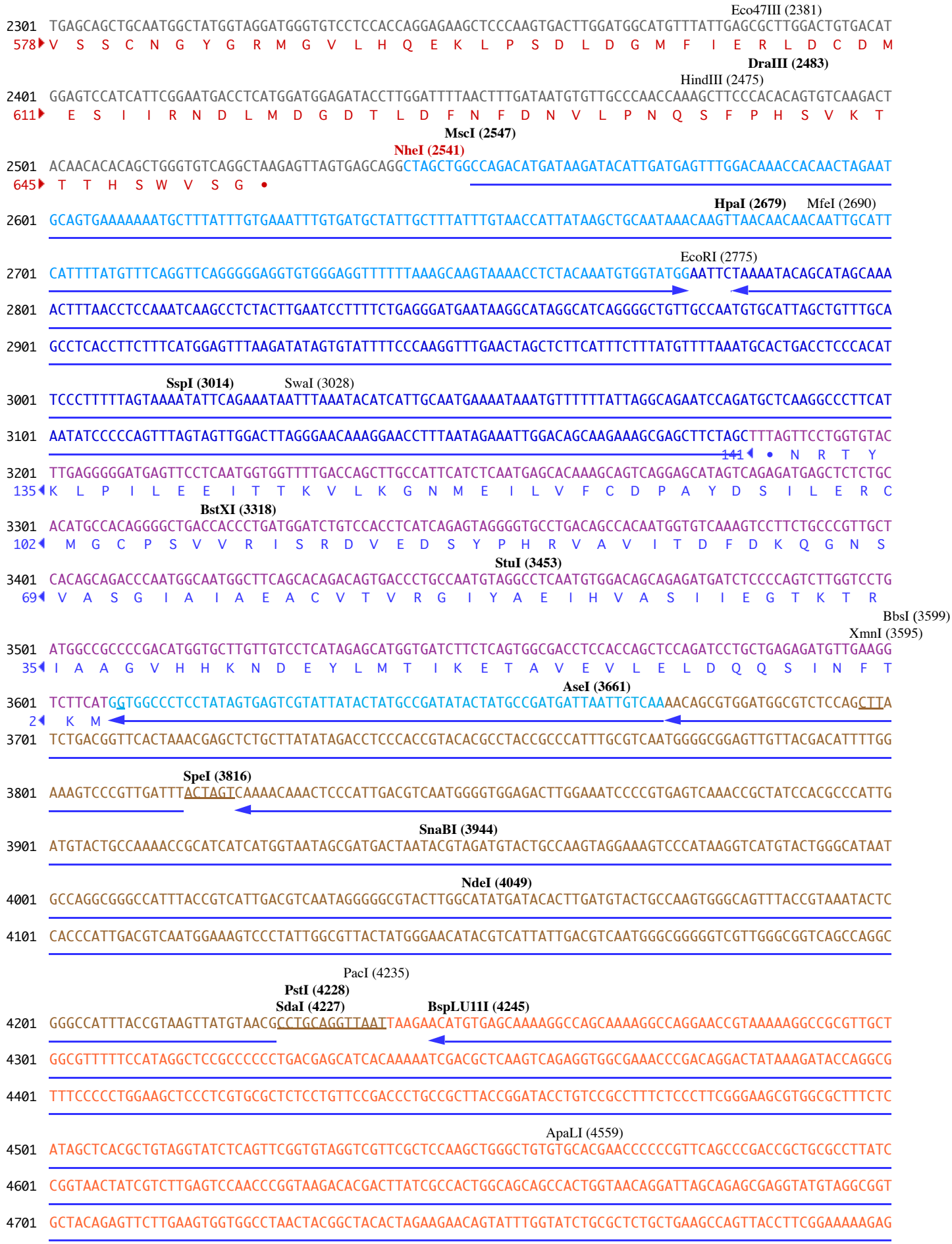
Eco47III (2381)
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578 V S S C N G Y G R M G V L H Q E K L P S D L D G M F I E R L D C D M
DraIII (2483)
HindIII (2475)
2401 GGAGTCCATCATTCGGAATGACCTCATGGATGGAGATACCTTGGATTTTAACTTTGATAATGTGTTGCCCAACCAAAGCTTCCCACACAGTGTCAAGACT
611 E S I I R N D L M D G D T L D F N F D N V L P N Q S F P H S V K T
MscI (2547)
NheI (2541)
2501 ACAACACACAGCTGGGTGTCAGGCTAAGAGTTAGTGAGCAGGCTAGCTGGCCAGACATGATAAGATACATTGATGAGTTTGGACAAACCACAACTAGAAT
645 T T H S W V S G •
HpaI (2679)
MfeI (2690)
2601 GCAGTGAAAAAAATGCTTTATTTGTGAAATTTGTGATGCTATTGCTTTATTTGTAACCATTATAAGCTGCAATAAACAAGTTAACAACAACAATTGCATT
EcoRI (2775)
2701 CATTTTATGTTTCAGGTTCAGGGGGAGGTGTGGGAGGTTTTTTAAAGCAAGTAAAACCTCTACAAATGTGGTATGGAATTCTAAAATACAGCATAGCAAA
2801 ACTTTAACCTCCAAATCAAGCCTCTACTTGAATCCTTTTCTGAGGGATGAATAAGGCATAGGCATCAGGGGCTGTTGCCAATGTGCATTAGCTGTTTGCA
2901 GCCTCACCTTCTTTCATGGAGTTTAAGATATAGTGTATTTTCCCAAGGTTTGAACTAGCTCTTCATTTCTTTATGTTTTAAATGCACTGACCTCCCACAT
SspI (3014)
SwaI (3028)
3001 TCCCTTTTTAGTAAAATATTCAGAAATAATTTAAATACATCATTGCAATGAAAATAAATGTTTTTTATTAGGCAGAATCCAGATGCTCAAGGCCCTTCAT
3101 AATATCCCCCAGTTTAGTAGTTGGACTTAGGGAACAAAGGAACCTTTAATAGAAATTGGACAGCAAGAAAGCGAGCTTCTAGCTTTAGTTCCTGGTGTAC
141 • N R T Y
3201 TTGAGGGGGATGAGTTCCTCAATGGTGGTTTTGACCAGCTTGCCATTCATCTCAATGAGCACAAAGCAGTCAGGAGCATAGTCAGAGATGAGCTCTCTGC
135 K L P I L E E I T T K V L K G N M E I L V F C D P A Y D S I L E R C
BstXI (3318)
3301 ACATGCCACAGGGGCTGACCACCCTGATGGATCTGTCCACCTCATCAGAGTAGGGGTGCCTGACAGCCACAATGGTGTCAAAGTCCTTCTGCCCGTTGCT
102 M G C P S V V R I S R D V E D S Y P H R V A V I T D F D K Q G N S
StuI (3453)
3401 CACAGCAGACCCAATGGCAATGGCTTCAGCACAGACAGTGACCCTGCCAATGTAGGCCTCAATGTGGACAGCAGAGATGATCTCCCCAGTCTTGGTCCTG
69 V A S G I A I A E A C V T V R G I Y A E I H V A S I I E G T K T R
BbsI (3599)
XmnI (3595)
3501 ATGGCCGCCCCGACATGGTGCTTGTTGTCCTCATAGAGCATGGTGATCTTCTCAGTGGCGACCTCCACCAGCTCCAGATCCTGCTGAGAGATGTTGAAGG
35 I A A G V H H K N D E Y L M T I K E T A V E V L E L D Q Q S I N F T
AseI (3661)
3601 TCTTCATGGTGGCCCTCCTATAGTGAGTCGTATTATACTATGCCGATATACTATGCCGATGATTAATTGTCAAAACAGCGTGGATGGCGTCTCCAGCTTA
2 K M
3701 TCTGACGGTTCACTAAACGAGCTCTGCTTATATAGACCTCCCACCGTACACGCCTACCGCCCATTTGCGTCAATGGGGCGGAGTTGTTACGACATTTTGG
SpeI (3816)
3801 AAAGTCCCGTTGATTTACTAGTCAAAACAAACTCCCATTGACGTCAATGGGGTGGAGACTTGGAAATCCCCGTGAGTCAAACCGCTATCCACGCCCATTG
SnaBI (3944)
3901 ATGTACTGCCAAAACCGCATCATCATGGTAATAGCGATGACTAATACGTAGATGTACTGCCAAGTAGGAAAGTCCCATAAGGTCATGTACTGGGCATAAT
NdeI (4049)
4001 GCCAGGCGGGCCATTTACCGTCATTGACGTCAATAGGGGGCGTACTTGGCATATGATACACTTGATGTACTGCCAAGTGGGCAGTTTACCGTAAATACTC
4101 CACCCATTGACGTCAATGGAAAGTCCCTATTGGCGTTACTATGGGAACATACGTCATTATTGACGTCAATGGGCGGGGGTCGTTGGGCGGTCAGCCAGGC
PacI (4235)
PstI (4228)
SdaI (4227)
BspLU11I (4245)
4201 GGGCCATTTACCGTAAGTTATGTAACGCCTGCAGGTTAATTAAGAACATGTGAGCAAAAGGCCAGCAAAAGGCCAGGAACCGTAAAAAGGCCGCGTTGCT
4301 GGCGTTTTTCCATAGGCTCCGCCCCCCTGACGAGCATCACAAAAATCGACGCTCAAGTCAGAGGTGGCGAAACCCGACAGGACTATAAAGATACCAGGCG
4401 TTTCCCCCTGGAAGCTCCCTCGTGCGCTCTCCTGTTCCGACCCTGCCGCTTACCGGATACCTGTCCGCCTTTCTCCCTTCGGGAAGCGTGGCGCTTTCTC
ApaLI (4559)
4501 ATAGCTCACGCTGTAGGTATCTCAGTTCGGTGTAGGTCGTTCGCTCCAAGCTGGGCTGTGTGCACGAACCCCCCGTTCAGCCCGACCGCTGCGCCTTATC
4601 CGGTAACTATCGTCTTGAGTCCAACCCGGTAAGACACGACTTATCGCCACTGGCAGCAGCCACTGGTAACAGGATTAGCAGAGCGAGGTATGTAGGCGGT
4701 GCTACAGAGTTCTTGAAGTGGTGGCCTAACTACGGCTACACTAGAAGAACAGTATTTGGTATCTGCGCTCTGCTGAAGCCAGTTACCTTCGGAAAAAGAG

4801 TTGGTAGCTCTTGATCCGGCAAACAAACCACCGCTGGTAGCGGTGGTTTTTTTGTTTGCAAGCAGCAGATTACGCGCAGAAAAAAAGGATCTCAAGAAGA

PacI (4975) SwaI (4984) NotI (4994)

4901 TCCTTTGATCTTTTCTACGGGGTCTGACGCTCAGTGGAACGAAAACTCACGTTAAGGGATTTTGGTCATGGCTAGTTAATTAACATTTAAATCAGCGGCC

5001 GCAATAAAATATCTTTATTTTCATTACATCTGTGTGTTGGTTTTTTGTGTGAATCGTAACTAACATACGCTCTCCATCAAAACAAAACGAAACAAAACAA

5101 ACTAGCAAAATAGGCTGTCCCCAGTGCAAGTGCAGGTGCCAGAACATTTCTCTATCGAA