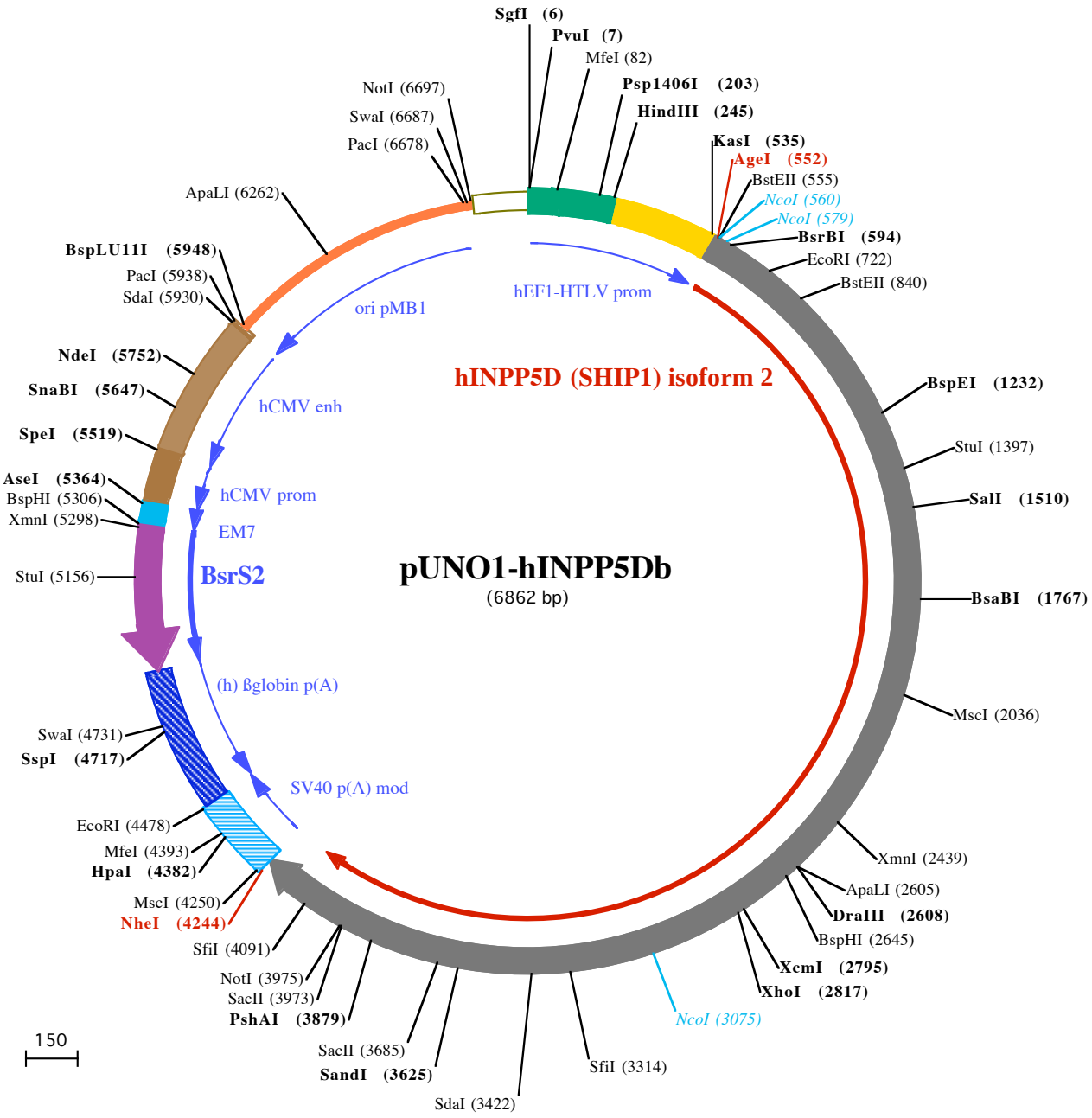
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(6862 bp)
hINPP5D (SHIP1) isoform 2
hEF1-HTLV prom
ori pMB1
hCMV enh
hCMV prom
EM7
BsrS2
(h) ßglobin p(A)
SV40 p(A) mod
SgfI (6)
PvuI (7)
MfeI (82)
Psp1406I (203)
HindIII (245)
KasI (535)
AgeI (552)
BstEII (555)
NcoI (560)
NcoI (579)
BsrBI (594)
EcoRI (722)
BstEII (840)
BspEI (1232)
StuI (1397)
SalI (1510)
BsaBI (1767)
MscI (2036)
XmnI (2439)
ApaLI (2605)
DraIII (2608)
BspHI (2645)
XcmI (2795)
XhoI (2817)
NcoI (3075)
SfiI (3314)
SdaI (3422)
SacII (3685)
SandI (3625)
NotI (3975)
SacII (3973)
PshAI (3879)
SfiI (4091)
NheI (4244)
MscI (4250)
HpaI (4382)
MfeI (4393)
EcoRI (4478)
SspI (4717)
SwaI (4731)
StuI (5156)
XmnI (5298)
BspHI (5306)
AseI (5364)
SpeI (5519)
SnaBI (5647)
NdeI (5752)
SdaI (5930)
PacI (5938)
BspLU11I (5948)
ApaLI (6262)
PacI (6678)
SwaI (6687)
NotI (6697)
150


PvuI (7)
SgfI (6)
MfeI (82)
1 GGATCTGCGATCGCTCCGGTGCCCGTCAGTGGGCAGAGCGCACATCGCCCACAGTCCCCGAGAAGTTGGGGGGAGGGGTCGGCAATTGAACGGGTGCCTA
101 GAGAAGGTGGCGCGGGGTAAACTGGGAAAGTGATGTCGTGTACTGGCTCCGCCTTTTTCCCGAGGGTGGGGGAGAACCGTATATAAGTGCAGTAGTCGCC
Psp1406I (203)
HindIII (245)
201 GTGAACGTTCTTTTTCGCAACGGGTTTGCCGCCAGAACACAGCTGAAGCTTCGAGGGGCTCGCATCTCTCCTTCACGCGCCCGCCGCCCTACCTGAGGCC
301 GCCATCCACGCCGGTTGAGTCGCGTTCTGCCGCCTCCCGCCTGTGGTGCCTCCTGAACTGCGTCCGCCGTCTAGGTAAGTTTAAAGCTCAGGTCGAGACC
401 GGGCCTTTGTCCGGCGCTCCCTTGGAGCCTACCTAGACTCAGCCGGCTCTCCACGCTTTGCCTGACCCTGCTTGCTCAACTCTACGTCTTTGTTTCGTTT
NcoI (560)
BstEII (555)
KasI (535)
AgeI (552)
NcoI (579)
BsrBI (594)
501 TCTGTTCTGCGCCGTTACAGATCCAAGCTGTGACCGGCGCCTACCTGAGATCACCGGTCACCATGGTCCCCTGCTGGAACCATGGCAACATCACCCGCTC
1 MetValProCysTrpAsnHisGlyAsnIleThrArgSe
601 CAAGGCGGAGGAGCTGCTTTCCAGGACAGGCAAGGACGGGAGCTTCCTCGTGCGTGCCAGCGAGTCCATCTCCCGGGCATACGCGCTCTGCGTGCTGTAT
13 rLysAlaGluGluLeuLeuSerArgThrGlyLysAspGlySerPheLeuValArgAlaSerGluSerIleSerArgAlaTyrAlaLeuCysValLeuTyr
EcoRI (722)
701 CGGAATTGCGTTTACACTTACAGAATTCTGCCCAATGAAGATGATAAATTCACTGTTCAGGCATCCGAAGGCGTCTCCATGAGGTTCTTCACCAAGCTGG
47 ArgAsnCysValTyrThrTyrArgIleLeuProAsnGluAspAspLysPheThrValGlnAlaSerGluGlyValSerMetArgPhePheThrLysLeuA
BstEII (840)
801 ACCAGCTCATCGAGTTTTACAAGAAGGAAAACATGGGGCTGGTGACCCATCTGCAATACCCTGTGCCGCTGGAGGAAGAGGACACAGGCGACGACCCTGA
80 spGlnLeuIleGluPheTyrLysLysGluAsnMetGlyLeuValThrHisLeuGlnTyrProValProLeuGluGluGluAspThrGlyAspAspProGl
901 GGAGGACACAGAAAGTGTCGTGTCTCCACCCGAGCTGCCCCCAAGAAACATCCCGCTGACTGCCAGCTCCTGTGAGGCCAAGGAGGTTCCTTTTTCAAAC
113 uGluAspThrGluSerValValSerProProGluLeuProProArgAsnIleProLeuThrAlaSerSerCysGluAlaLysGluValProPheSerAsn
1001 GAGAATCCCCGAGCGACCGAGACCAGCCGGCCGAGCCTCTCCGAGACATTGTTCCAGCGACTGCAAAGCATGGACACCAGTGGGCTTCCAGAAGAGCATC
147 GluAsnProArgAlaThrGluThrSerArgProSerLeuSerGluThrLeuPheGlnArgLeuGlnSerMetAspThrSerGlyLeuProGluGluHisL
1101 TTAAGGCCATCCAAGATTATTTAAGCACTCAGCTCGCCCAGGACTCTGAATTTGTGAAGACAGGGTCCAGCAGTCTTCCTCACCTGAAGAAACTGACCAC
180 euLysAlaIleGlnAspTyrLeuSerThrGlnLeuAlaGlnAspSerGluPheValLysThrGlySerSerSerLeuProHisLeuLysLysLeuThrTh
BspEI (1232)
1201 ACTGCTCTGCAAGGAGCTCTATGGAGAAGTCATCCGGACCCTCCCATCCCTGGAGTCTCTGCAGAGGTTATTTGACCAGCAGCTCTCCCCGGGCCTCCGT
213 rLeuLeuCysLysGluLeuTyrGlyGluValIleArgThrLeuProSerLeuGluSerLeuGlnArgLeuPheAspGlnGlnLeuSerProGlyLeuArg
StuI (1397)
1301 CCACGTCCTCAGGTTCCTGGTGAGGCCAATCCCATCAACATGGTGTCCAAGCTCAGCCAACTGACAAGCCTGTTGTCGTCCATTGAAGACAAGGTCAAGG
247 ProArgProGlnValProGlyGluAlaAsnProIleAsnMetValSerLysLeuSerGlnLeuThrSerLeuLeuSerSerIleGluAspLysValLysA
1401 CCTTGCTGCACGAGGGTCCTGAGTCTCCGCACCGGCCCTCCCTTATCCCTCCAGTCACCTTTGAGGTGAAGGCAGAGTCTCTGGGGATTCCTCAGAAAAT
280 laLeuLeuHisGluGlyProGluSerProHisArgProSerLeuIleProProValThrPheGluValLysAlaGluSerLeuGlyIleProGlnLysMe
SalI (1510)
1501 GCAGCTCAAAGTCGACGTTGAGTCTGGGAAACTGATCATTAAGAAGTCCAAGGATGGTTCTGAGGACAAGTTCTACAGCCACAAGAAAATCCTGCAGCTG
313 tGlnLeuLysValAspValGluSerGlyLysLeuIleIleLysLysSerLysAspGlySerGluAspLysPheTyrSerHisLysLysIleLeuGlnLeu
1601 ATTAAGTCACAGAAATTTCTGAATAAGTTGGTGATCTTGGTGGAAACAGAGAAGGAGAAGATCCTGCGGAAGGAATATGTTTTTGCTGACTCCAAAAAGA
347 IleLysSerGlnLysPheLeuAsnLysLeuValIleLeuValGluThrGluLysGluLysIleLeuArgLysGluTyrValPheAlaAspSerLysLysA
BsaBI (1767)
1701 GAGAAGGCTTCTGCCAGCTCCTGCAGCAGATGAAGAACAAGCACTCAGAGCAGCCGGAGCCCGACATGATCACCATCTTCATCGGCACCTGGAACATGGG
380 rgGluGlyPheCysGlnLeuLeuGlnGlnMetLysAsnLysHisSerGluGlnProGluProAspMetIleThrIlePheIleGlyThrTrpAsnMetGl
1801 TAACGCCCCCCCTCCCAAGAAGATCACGTCCTGGTTTCTCTCCAAGGGGCAGGGAAAGACGCGGGACGACTCTGCGGACTACATCCCCCATGACATTTAC
413 yAsnAlaProProProLysLysIleThrSerTrpPheLeuSerLysGlyGlnGlyLysThrArgAspAspSerAlaAspTyrIleProHisAspIleTyr
1901 GTGATCGGCACCCAAGAGGACCCCCTGAGTGAGAAGGAGTGGCTGGAGATCCTCAAACACTCCCTGCAAGAAATCACCAGTGTGACTTTTAAAACAGTCG
447 ValIleGlyThrGlnGluAspProLeuSerGluLysGluTrpLeuGluIleLeuLysHisSerLeuGlnGluIleThrSerValThrPheLysThrValA
MscI (2036)
2001 CCATCCACACGCTCTGGAACATCCGCATCGTGGTGCTGGCCAAGCCTGAGCACGAGAACCGGATCAGCCACATCTGTACTGACAACGTGAAGACAGGCAT
480 laIleHisThrLeuTrpAsnIleArgIleValValLeuAlaLysProGluHisGluAsnArgIleSerHisIleCysThrAspAsnValLysThrGlyIl
2101 TGCAAACACACTGGGGAACAAGGGAGCCGTGGGGGTGTCGTTCATGTTCAATGGAACCTCCTTAGGGTTCGTCAACAGCCACTTGACTTCAGGAAGTGAA
513 eAlaAsnThrLeuGlyAsnLysGlyAlaValGlyValSerPheMetPheAsnGlyThrSerLeuGlyPheValAsnSerHisLeuThrSerGlySerGlu
2201 AAGAAACTCAGGCGAAACCAAAACTATATGAACATTCTCCGGTTCCTGGCCCTGGGCGACAAGAAGCTGAGTCCCTTTAACATCACTCACCGCTTCACGC
547 LysLysLeuArgArgAsnGlnAsnTyrMetAsnIleLeuArgPheLeuAlaLeuGlyAspLysLysLeuSerProPheAsnIleThrHisArgPheThrH
2301 ACCTCTTCTGGTTTGGGGATCTTAACTACCGTGTGGATCTGCCTACCTGGGAGGCAGAAACCATCATCCAGAAAATCAAGCAGCAGCAGTACGCAGACCT
580 isLeuPheTrpPheGlyAspLeuAsnTyrArgValAspLeuProThrTrpGluAlaGluThrIleIleGlnLysIleLysGlnGlnGlnTyrAlaAspLe
XmnI (2439)
2401 CCTGTCCCACGACCAGCTGCTCACAGAGAGGAGGGAGCAGAAGGTCTTCCTACACTTCGAGGAGGAAGAAATCACGTTTGCCCCAACCTACCGTTTTGAG
613 uLeuSerHisAspGlnLeuLeuThrGluArgArgGluGlnLysValPheLeuHisPheGluGluGluGluIleThrPheAlaProThrTyrArgPheGlu
2501 AGACTGACTCGGGACAAATACGCCTACACCAAGCAGAAAGCGACAGGGATGAAGTACAACTTGCCTTCCTGGTGTGACCGAGTCCTCTGGAAGTCTTATC
647 ArgLeuThrArgAspLysTyrAlaTyrThrLysGlnLysAlaThrGlyMetLysTyrAsnLeuProSerTrpCysAspArgValLeuTrpLysSerTyrP
DraIII (2608)
ApaLI (2605)
BspHI (2645)
2601 CCCTGGTGCACGTGGTGTGTCAGTCTTATGGCAGTACCAGCGACATCATGACGAGTGACCACAGCCCTGTCTTTGCCACATTTGAGGCAGGAGTCACTTC
680 roLeuValHisValValCysGlnSerTyrGlySerThrSerAspIleMetThrSerAspHisSerProValPheAlaThrPheGluAlaGlyValThrSe
XcmI (2795)
2701 CCAGTTTGTCTCCAAGAACGGTCCCGGGACTGTTGACAGCCAAGGACAGATTGAGTTTCTCAGGTGCTATGCCACATTGAAGACCAAGTCCCAGACCAAA
713 rGlnPheValSerLysAsnGlyProGlyThrValAspSerGlnGlyGlnIleGluPheLeuArgCysTyrAlaThrLeuLysThrLysSerGlnThrLys


XhoI (2817)
NcoI (3075)
SfiI (3314)
SdaI (3422)
SandI (3625)
SacII (3685)
PshAI (3879)
NotI (3975)
SacII (3973)
SfiI (4091)
MscI (4250)
NheI (4244)
HpaI (4382)
MfeI (4393)
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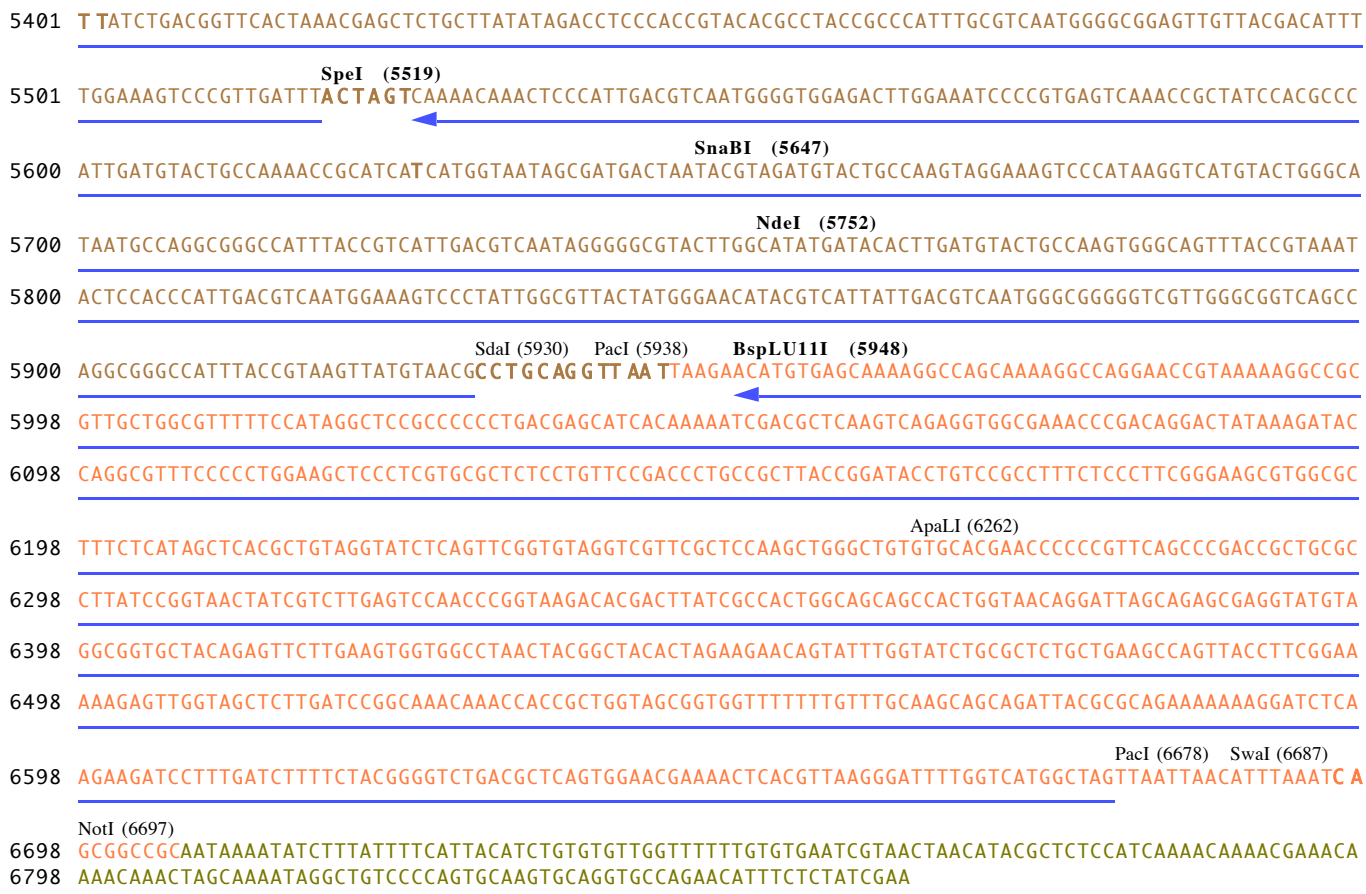
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SpeI (5519)
5501 TGGAAAGTCCCGTTGATTTACTAGTCAAAACAAACTCCCATTGACGTCAATGGGGTGGAGACTTGGAAATCCCCGTGAGTCAAACCGCTATCCACGCCC
SnaBI (5647)
5600 ATTGATGTACTGCCAAAACCGCATCATCATGGTAATAGCGATGACTAATACGTAGATGTACTGCCAAGTAGGAAAGTCCCATAAGGTCATGTACTGGGCA
NdeI (5752)
5700 TAATGCCAGGCGGGCCATTTACCGTCATTGACGTCAATAGGGGGCGTACTTGGCATATGATACACTTGATGTACTGCCAAGTGGGCAGTTTACCGTAAAT
5800 ACTCCACCCATTGACGTCAATGGAAAGTCCCTATTGGCGTTACTATGGGAACATACGTCATTATTGACGTCAATGGGCGGGGGTCGTTGGGCGGTCAGCC
SdaI (5930) PacI (5938) BspLU11I (5948)
5900 AGGCGGGCCATTTACCGTAAGTTATGTAACGCCTGCAGGTTAATTAAGAACATGTGAGCAAAAGGCCAGCAAAAGGCCAGGAACCGTAAAAAGGCCGC
5998 GTTGCTGGCGTTTTTCCATAGGCTCCGCCCCCCTGACGAGCATCACAAAAATCGACGCTCAAGTCAGAGGTGGCGAAACCCGACAGGACTATAAAGATAC
6098 CAGGCGTTTCCCCCTGGAAGCTCCCTCGTGCGCTCTCCTGTTCCGACCCTGCCGCTTACCGGATACCTGTCCGCCTTTCTCCCTTCGGGAAGCGTGGCGC
ApaLI (6262)
6198 TTTCTCATAGCTCACGCTGTAGGTATCTCAGTTCGGTGTAGGTCGTTCGCTCCAAGCTGGGCTGTGTGCACGAACCCCCCGTTCAGCCCGACCGCTGCGC
6298 CTTATCCGGTAACTATCGTCTTGAGTCCAACCCGGTAAGACACGACTTATCGCCACTGGCAGCAGCCACTGGTAACAGGATTAGCAGAGCGAGGTATGTA
6398 GGCGGTGCTACAGAGTTCTTGAAGTGGTGGCCTAACTACGGCTACACTAGAAGAACAGTATTTGGTATCTGCGCTCTGCTGAAGCCAGTTACCTTCGGAA
6498 AAAGAGTTGGTAGCTCTTGATCCGGCAAACAAACCACCGCTGGTAGCGGTGGTTTTTTTGTTTGCAAGCAGCAGATTACGCGCAGAAAAAAAGGATCTCA
PacI (6678) SwaI (6687)
6598 AGAAGATCCTTTGATCTTTTCTACGGGGTCTGACGCTCAGTGGAACGAAAACTCACGTTAAGGGATTTTGGTCATGGCTAGTTAATTAACATTTAAATCA
NotI (6697)
6698 GCGGCCGCAATAAAATATCTTTATTTTCATTACATCTGTGTGTTGGTTTTTTGTGTGAATCGTAACTAACATACGCTCTCCATCAAAACAAAACGAAACA
6798 AAACAAACTAGCAAAATAGGCTGTCCCCAGTGCAAGTGCAGGTGCCAGAACATTTCTCTATCGAA