





pDRAW32 - pAMSJ - 1500 bp. - [Sequence]

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1 GCACTTCACC AACAAAGGACC ATAGCATATG AAAATCGAAG AAGGTAAACT  
M K I E E G K L Frame 1

51 GGTAATCTGG ATTAACGGCG ATAAAGGCTA TAACGGTCTC GCTGAAGTCG  
V I W I N G D K G Y N G L A E V G Frame 1

101 GTAAGAAATT CGAGAAAGAT ACCGGAATTA AAGTCACCGT TGAGCATCCG  
K K F E K D T G I K V T V E H P Frame 1

151 GATAAACTGG AAGAGAAATT CCCACAGGTT GCGGCAACTG GCGATGGCCC  
D K L E E K F P Q V A A T G D G P Frame 1

201 TGACATTATC TTCGGGCGAC ACGACCGCTT TGGTGGCTAC GCTCAATCTG  
D I I F W A H D R F G G Y A Q S G Frame 1

251 GCCTGTTGGC TGAAATCACC CCGGACAAAG CGTTCCAGGA CAAGCTGTAT  
L L A E I T P D K A F Q D K L Y Frame 1

301 CCGTTTACCT GGGATGCCGT ACGTTACAAC GGCAAGCTGA TTGCTTACCC  
P F T W D A V R Y N G K L I A Y P Frame 1

DeiT BcIII

New DNA entry Analysis complete 12:00 من

pDRAW32 - pAMSJ - 1500 bp. - [Restriction analysis]

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pAMSJ, 1500 bp. (LINEAR)  
Restriction analysis 07/09/2014  
pDRAW32 revision 1.1.126  
UNLICENSED  
Rebase containing 277 enzymes.  
75 enzymes match enzyme selection criteria.

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**AccI** (GT'mk\_AC) [FblI,XmiI]  
Cuts 1 time.  
Cuts at position 1214.  
Fragment sizes 1214, 286.

**AcuI** (CTGAAGnnnnnnnnnnnn\_nn') [Eco57I]  
Cuts 2 times.  
Cuts at positions 113, 961.  
Fragment sizes 113, 848, 539.

**AfeI** (AGC'GCT) [Eco47III,Aor51HI]  
Cuts 1 time.  
Cuts at position 938.  
Fragment sizes 938, 562.

**AjuI** (GAAnnnnnnnTTGnnnnnn\_nnnnn')  
Cuts 1 time.  
Cuts at position 1411.  
Fragment sizes 1411, 89.

**AjuI** (CCAAAnnnnnnnTTCnnnnnn\_nnnnn')  
Cuts 1 time.  
Cuts at position 1379.  
Fragment sizes 1379, 121.

New DNA entry Analysis complete 12:00 من

