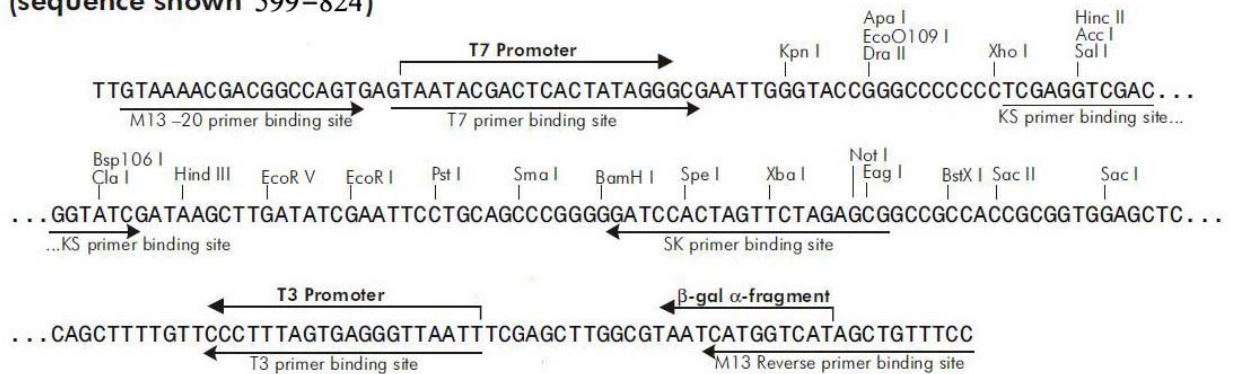


**pBluescript II SK (-) Multiple Cloning Site Region
(sequence shown 599–824)**



Seq:

LOCUS pBluescript II SK(-) 2959 bp DNA CIRCULAR SYN 19-MAR-2009

DEFINITION

ACCESSION pBluescript II SK(-)

KEYWORDS

SOURCE .
Unknown.

ORGANISM Unknown
Unclassified.

REFERENCE 1 (bases 1 to 2959)

AUTHORS Self

JOURNAL Unpublished.

COMMENT SECID/File created by SciEd Central, Scientific & Educational Software

FEATURES

Location/Qualifiers

CDS 21..327
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misc_feature 600..643
/gene="T7/M13F"
CDS complement (461..615)
/gene="lacZ"
misc_feature complement (770..826)
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CDS complement (1974..2831)
/gene="Amp"

BASE COUNT 707 a 755 c 731 g 766 t

ORIGIN

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121 CCACGTTTCGC CGGCTTTCCC CGTCAAGCTC TAAATCGGGG GCTCCCTTTA GGGTTCGGAT
181 TTAGTGCTTT ACGGCACCTC GACCCCAAAA AACTTGATTA GGGTGATGGT TCACGTAGTG
241 GGCCATCGCC CTGATAGACG GTTTTTCGCC CTTTGACGTT GGAGTCCACG TTCTTTAATA
301 GTGGACTCTT GTTCCAAACT GGAACAACAC TCAACCCTAT CTCGGTCTAT TCTTTTGATT
361 TATAAGGGAT TTTGCCGATT TCGGCCTATT GGTAAAAAA TGAGCTGATT TAACAAAAAT
421 TTAACGCGAA TTTTAACAAA ATATTAACGC TTACAATTTG CCATTGCGCA TTCAGGCTGC
481 GCAACTGTTG GGAAGGGCGA TCGGTGCGGG CCTCTTCGCT ATTACGCCAG CTGGCGAAAG
541 GGGGATGTGC TGCAAGGCGA TTAAGTTGGG TAACGCCAGG GTTTTCCCAG TCACGACGTT
601 GTAAAACGAC GGCCAGTGAA TTGTAATACG ACTCACTATA GGGCGAATTG GGTACCGGGC
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781 GGGTTAATTT CGAGCTTGGC GTAATCATGG TCATAGCTGT TTCCTGTGTG AAATTTGTTAT
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901 TAATGAGTGA GCTAACTCAC ATTAATTGCG TTGCGCTCAC TGCCCGCTTT CCAGTCGGGA
961 AACCTGTCGT GCCAGCTGCA TTAATGAATC GGCCAACGCG CGGGGAGAGG CGGTTTGCGT
1021 ATTGGGCGCT CTTCCGCTTC CTCGCTCACT GACTCGCTGC GCTCGGTCGT TCGGCTGCGG
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2041 CTCCCCGTCG TGTAGATAAC TACGATACGG GAGGGCTTAC CATCTGGCCC CAGTGCTGCA
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2161 GGAAGGGCCG AGCGCAGAAG TGGTCCTGCA ACTTTATCCG CCTCCATCCA GTCTATTAAT
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