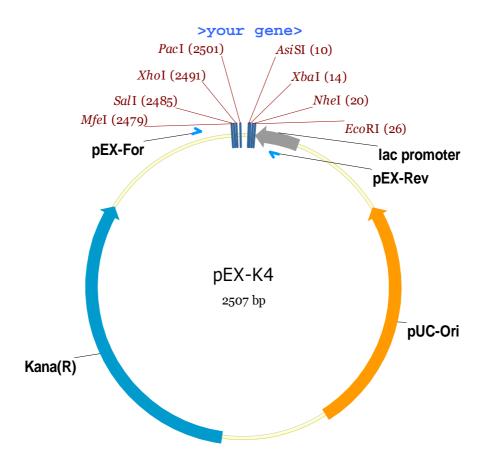
pEX-K4 vector map, multiple cloning site and full sequence



MCS of pEX-K4

GGAGCAGACAAGCCCGTCAGGGCGCGTCAGCGGGTGTTGGCGGGCC
TGGCTTAACTATGCGGCATCAGAGCAGATTGTACTGAGAGTGCACcaattgGT
CGACctcgagTTAATTAACGTA>your_gene>GTGGGCGATCGCtctagaGC
TAGCgaattcCTGTGTGAAATTGTTATCCGCTCACAATTCCACACAACATACG
AGCCGGAAGCATAAAGTGTAAAGCCTG

Start point: gene insertion site

(\underline{G} of $\underline{G}TGG$ 3' of "your gene" is bp no. 1)

Multiple Cloning Site: bases 2478-30

lac Promoter: bases 28-143
pUC origin: bases 404-1023

Kanamycin resistance ORF: bases 1309-2103 pEX-For priming site: bases 2380-2400 pEX-Rev priming site: bases 75-100

Full Sequence of pEX-K4 (2507 bp)

(1)

YOUR_GENE_gtgggcgatcgctctagagctagcgaattcctgtgtgaaattgttatccgctcacaa acattaattgcgttgcgctcactgcccgctttccagtcgggaaacctgtcgtgccagctgcattaatg $\verb| aatcggccaacgcgggggagaggggtttgcgtattgggcgctcttccgcttcctcgctcactgact| \\$ acagaatcaggggataacgcaggaaagaacatgtgagcaaaaggccagcaaaaggccaggaaccgtaa aaaggccgcgttgctggcgtttttccataggctccgccccctgacgagcatcacaaaaatcgacgct caagtcagaggtggcgaaacccgacaggactataaagataccaggcgtttccccctggaagctccctc gtgcgctctcctgttccgaccctgccgcttaccggatacctgtccgcctttctcccttcgggaagcgt ggcgctttctcatagctcacgctgtaggtatctcagttcggtgtaggtcgttcgctccaagctgggct gtgtgcacgaacccccgttcagcccgaccgctgcgccttatccggtaactatcgtcttgagtccaac ${\tt ccggtaagacacgacttatcgccactggcagcagccactggtaacaggattagcagagcgaggtatgt}$ aggcggtgctacagagttcttgaagtggtggcctaactacggctacactagaagaacagtatttggta agatcctttgatcttttctacggggtctgacgctcagtggaacgaaaactcacgttaagggattttgg taaagtatatatgagtaaaaatattccggaattgccagctggggcgcctctggtaaggttgggaagc $\verb|cctgcaaagtaaactggatggctttcttgccgccaaggatctgatggcgcaggggatcaagatctgat|\\$ caagagacaggatgaggatcgtttcgcatgattgaacaagatggattgcacgcaggttctccggccgc ttgggtggagaggctattcggctatgactgggcacaacagacaatcggctgctctgatgccgccgtgt ctgcaggacgaggcagcggctatcgtggctaggccacgacgggcgttccttgcgcagctgtgctcga cgttgtcactgaagcgggaagggactggctgctattgggcgaagtgccgggggcaggatctcctgtcat $\verb|cccaccttgctcctgccgagaaagtatccatcatggctgatgcaatgcggcggctgcatacgcttgat|$ cggtcttgtcgatcaggatgatctggacgaagagcatcaggggctcgcgccagccgaactgttcgcca ggctcaaggcgcgcatgcccgacggcgaggatctcgttgtcgtgacccatggcgatgcctgcttgccgaat atcatggtggaaaatggccgcttttctggattcatcgactgtggccggctgggtgtggcggaccgcta tcaggacatagcgttggctacccgtgatattgctgaagagcttggcggcgaatgggctgaccgcttcc tcgtgctttacggtatcgccgctcccgattcgcagcgcatcgccttctatcgccttcttgacgagttc ttctgaaccggtaatattattgaagcatttatcagggttattgtctcatgagcggatacatatttgaa tgtatttagaaaaataaacaaataggggttccgcgcacatttccccgaaaagtgccacctgacgtcta agaaaccattattatcatgacattaacctataaaaataggcgtatcacgaggccctttcgtctcgcgc gtttcggtgatgacggtgaaaacctctgacacatgcagctcccggagacggtcacagcttgtctgtaa ta (2507)