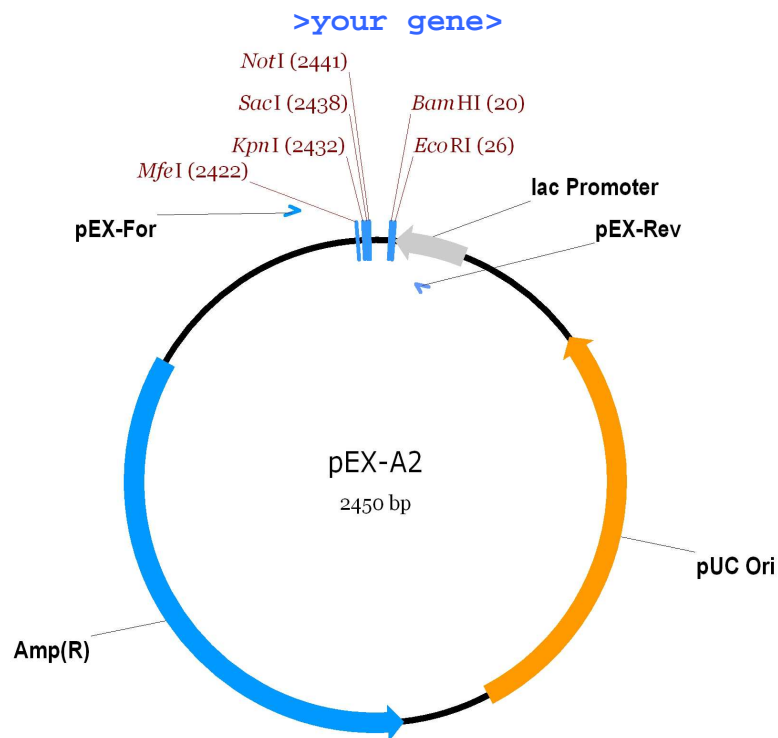


pEX-A2 vector map, multiple cloning site and full sequence



MCS of pEX-A2

```

GGAGCAGACAAGCCCGTCAGGGCGCGTCAGCGGGTGTGTCGGGGC
TGGCTTAACTATGCGGCATCAGAGCAGATTGTACTGAGAGTGCACcaattgGG
TACCgagctcGCGGCCGCAAGC>your_gene>ACCTGCTTTTGCTCGCTTgg
atccGAATTCCCTGTGTGAAATTGTTATCCGCTCACAATTCCACACAACATACG
AGCCGGAAGCATAAAGTGTAAGCCTG
  
```

Start point: gene insertion site

(A of ACCT 3' of "your gene" is bp no. 1)

Multiple Cloning Site: bases 2421-30

lac Promoter: bases 28 - 143

pUC origin: bases 360 - 1033

Ampicillin resistance ORF: bases 1178-2039

pEX-For priming site: bases 2323-2343

pEX-Rev priming site: bases 75 - 100

Full Sequence of pEX-A2 (2450 bp)

(1)

YOUR_GENE acctgcttttgctcgcttggatccgaattcctgtgtgaaattgttatccgc
tcacaattccacacaacatacagagccggaagcataaagtgtaaagcctgggggtgcctaata
gtgagctaactcacattaattgcttgcgctcactgcccgcctttccagtcgggaaacctgtc
gtgccagctgcattaatgaatcggccaacgcgcggggagaggcggtttgcgtattgggcgct
cttcgcgttcctcgctcactgactcgctgcgctcggtcgttcggctgcggcgagcggtatca
gctcactcaaaggcggttaatacggttatccacagaatcaggggataacgcaggaaagaacat
gtgagcaaaaggccagcaaaaggccaggaaccgtaaaaaggccgcgttgctggcggtttttcc
ataggctccgccccctgacgagcatcacaaaaatcgacgctcaagtcagagggtggcgaaac
ccgacaggactataaagataaccaggcggtttccccctggaagctccctcgctgcgctctcctgt
tccgaccctgccgcttaccggatacctgtccgcctttctcccttcgggaagcggtggcgcttt
ctcatagctcacgctgtaggtatctcagttcgggtgtaggtcgttcgctccaagctgggctgt
gtgcacgaaccccccggttcagcccagccgctgcgccttatccggtaactatcgctcttgagtc
caaccgcgtaagacacgacttatcgccactggcagcagccactggtaacaggattagcagag
cgaggtatgtaggcgggtgctacagagttcttgaagtgggtggcctaactacggctacactaga
agaacagtatTTTGGTATCTGCGCTCTGCTGAAGCCAGTTACCTTCGAAAAAGAGTTGGTAG
CTCTTGATCCGGCAAACAAACCACCGCTGGTAGCGGTGGTTTTTTTGTGTGCAAGCAGCAGA
TTACGCGCAGAAAAAAGGATCTCAAGAAGATCCTTTGATCTTTTCTACGGGGTCTGACGCT
CAGTGGAACGAAAACCTCACGTTAAGGGATTTTGGTCA TGAGATTATCAAAAAGGATCTTCAC
CTAGATCCTTTTAAATTA AAAATGAAGTTTTAAATCAATCTAAAGTATATATGAGTAAACTT
GGTCTGACAGTTACCAATGCTTAATCAGTGAGGCACCTATCTCAGCGATCTGTCTATTTCTGT
TCATCCATAGTTGCCTGACTCCCCGTCGTGTAGATAACTACGATACGGGAGGGCTTACCATC
TGGCCCCAGTGCTGCAATGATACC GCGACTCCACGCTCACCGGCTCCAGATTTATCAGCAA
TAAACCAGCCAGCCGGAAGGGCCGAGCGCAGAAGTGGTCTCTGCAACTTTATCCGCCTCCATC
CAGTCTATTAATTGTTGCCGGGAAGCTAGAGTAAGTAGTTCGCCAGTTAATAGTTTGCGCAA
CGTTGTTGCCATTGCTACAGGCATCGTGGTGTACGCTCGTCGTTTGGTATGGCTTCATTCA
GCTCCGGTTC CCAACGATCAAGGCGAGTTACATGATCCCCCATGTTGTGCAAAAAAGCGGT
TAGTCTCCTTCGGTCCTCCGATCGTTGT CAGAAGTAAGTTGGCCGCGAGTGTTATCACTCATGG
TTATGGCAGCACTGCATAATTCTCTTACTGT CATGCCATCCGTAAGATGCTTTTCTGTGACT
GGTGAGTACTCAACCAAGTCATTCTGAGAATAGTGTATGCGGCGACCGAGTTGCTCTTGCCC
GGCGTCAATACGGGATAATAACCGCGCCACATAGCAGAACCTTAAAAGTGCTCATCATTTGGAA
AACGTTCTTCGGGGCGAAAACCTCTCAAGGATCTTACC GCTGTTGAGATCCAGTTCGATGTAA
CCCCTCGTGCACCCAACCTGATCTTCAGCATCTTTTACTTTTACCAGCGTTTCTGGGTGAGC
AAAAACAGGAAGGCAAAATGCCGCAAAAAAGGGAATAAGGGCGACACGGAAATGTTGAATAC
TCATACTCTTCCTTTTTTCAATATTATTGAAGCATTTATCAGGGTTATTGTCTCATGAGCGGA
TACATATTTGAATGTATTTAGAAAAATAACAAATAGGGGTTCCGCGCACATTTCCCCGAAA
AGTGCCACCTGACGTCTAAGAAACCATTTATCATGACATTAACTATAAAAAATAGGCGTA
TCACGAGGCCCTTTCTGTCTCGCGCGTTTCGGTGATGACGGTGAAAACCTCTGACACATGCAG
CTCCCGGAGACGGTCACAGCTTGTCTGTAAGCGGATGCCGGGAGCAGACAAGCCCGTCAGGG
CGCGTCAGCGGGTGTGGCGGGTGTGGGGGCTGGCTTAAC TATGCGGCATCAGAGCAGATTG
TACTGAGAGTGCACCAATTGGGTACCAGAGCTCGCGGCCGCAAGC (2450)