Appendix

A. DNA sequences of the UZ3/4 TCR chains

TCR alpha 7 chain

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LOCUS       TCR alpha     822 bp    DNA
ACCESSION   TCR alpha 7
KEYWORDS    L + recombined V alpha 7.2 J alpha 18 + C alpha
SOURCE      CD8+ T cell clone UZ3/4.
  ORGANISM  mus musculus

REFERENCE   1  (bases 1 to 822)
  AUTHORS   Self
  JOURNAL   Unpublished.
FEATURES             Location/Qualifiers
BASE COUNT      226 a    211 c    181 g    204 t
ORIGIN
        1 ATGCCTCCTC ACAGCCTGCT CTGTGTGCTG GTGGCCTTGG CTTTCTCTGG ATCTAATGTG
       61 GCCCAGAAAG TGATTCAGGT CTGGTCAACA ACAAGCAGGC AGGAGGGCGA AAAACTCACA
      121 CTGGACTGTT CATATAAGAC AAGTCAGGTC TTATACCATC TTTTCTGGTA CAAGCACCTT
      181 CTTAGTGGAG AGATGGTTTT GCTTATTCGA CAAATGCCTT CTACTATTGC AATAGAGAGG
      241 AGCGGCCGCT ATTCTGTAGT CTTCCAGAAA TCACGCAAAT CCATCAGCCT TGTCATTTCA
      301 ACCTTACAAC CAGACGATTC GGGAAAGTAT TTCTGTGCTC TCTGGGAGCT GGACTATAAC
      361 CAGGGGAAGC TTATCTTTGG ACAGGGAACC AAGTTATCTA TCAAGCCCAA CATCCAGAAC
      421 CCAGAACCTG CTGTGTACCA GTTAAAAGAT CCTCGGTCTC AGGACAGCAC CCTCTGCCTG
      481 TTCACCGACT TTGACTCCCA AATCAATGTG CCGAAAACCA TGGAATCTGG AACGTTCATC
      541 ACTGACAAAA CTGTGCTGGA CATGAAAGCT ATGGATTCCA AGAGCAATGG GGCCATTGCC
      601 TGGAGCAACC AGACAAGCTT CACCTGCCAA GATATCTTCA AAGAGACCAA CGCCACCTAC
      661 CCCAGTTCAG ACGTTCCCTG TGATGCCACG TTGACTGAGA AAAGCTTTGA AACAGATATG
      721 AACCTAAACT TTCAAAACCT GTCAGTTATG GGACTCCGAA TCCTCCTGCT GAAAGTAGCC
      781 GGATTTAACC TGCTCATGAC GCTGAGGCTG TGGTCCAGTT GA

TRANSLATION
      MPPHSLLCVLVALAFSGSNVAQKVIQVWSTTSRQEGEKLTLDCSYKTSQVLYHL
      FWYKHLLSGEMVLLIRQMPSTIAIERSGRYSVVFQKSRKSISLVISTLQPDDSG
      KYFCALWELDYNQGKLIFGQGTKLSIKPNIQNPEPAVYQLK

      MPPHSLLCVLVALAFSGSNVAQKVIQVWSTTSRQEGEKLTLDCSYKTSQV
      LYHLFWYKHLLSGEMVLLIRQMPSTIAIERSGRYSVVFQKSRKSISLVIS
      TLQPDDSGKYFCALWELDYNQGKLIFGQGTKLSIKPNIQNPEPAVYQLKD
      PRSQDSTLCLFTDFDSQINVPKTMESGTFITDKTVLDMKAMDSKSNGAIA
      WSNQTSFTCQDIFKETNATYPSSDVPCDATLTEKSFETDMNLNFQNLSVM
      GLRILLLKVAGFNLLMTLRLWSS*
//
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TCR alpha 8 chain

```
LOCUS       TCR alpha     831 bp    DNA
ACCESSION   TCR alpha 8
KEYWORDS    L + recombined V alpha 8.2 J alpha 42 + C alpha
SOURCE      CD8+ T cell clone UZ3/4.
  ORGANISM  mus musculus

REFERENCE   1  (bases 1 to 831)
  AUTHORS   Self
  JOURNAL   Unpublished.
FEATURES             Location/Qualifiers
BASE COUNT      213 a    236 c    189 g    193 t
ORIGIN
        1 ATGCGTCCTG GCACCTGCTC AGTTCTTGTG CTCCTCCTAA TGCTCAGGAG GAGCAATGGA
       61 GATGGAGACT CAGTGACCCA GAAGGAAGGC CTGGTCACTC TCACCGAGGG GTTGCCTGTG
      121 ATGCTGAACT GCACCTATCA GACTATTTAC TCAAATGCTT TCCTTTTCTG GTATGTGCAC
      181 TATCTCAATG AATCCCCTCG GCTACTCCTG AAGAGCTCCA CAGACAACAA GAGGACCGAG
      241 CACCAAGGGT TCCACGCCAC TCTCCATAAG AGCAGCAGCT CCTTCCATCT GCAGAAGTCC
      301 TCAGCGCAGC TGTCAGACTC TGCCCTGTAC TACTGTGCTC TGAGTGATCG AGGGAGGGCA
      361 TCCTCCTCCT TCAGCAAGCT GGTGTTTGGG CAGGGGACAT CCTTATCAGT CGTTCCAAAC
      421 ATCCAGAACC CAGAACCTGC TGTGTACCAG TTAAAAGATC CTCGGTCTCA GGACAGCACC
      481 CTCTGCCTGT TCACCGACTT TGACTCCCAA ATCAATGTGC CGAAAACCAT GGAATCTGGA
      541 ACGTTCATCA CTGACAAAAC TGTGCTGGAC ATGAAAGCTA TGGATTCCAA GAGCAATGGG
      601 GCCATTGCCT GGAGCAACCA GACAAGCTTC ACCTGCCAAG ATATCTTCAA AGAGACCAAC
      661 GCCACCTACC CCAGTTCAGA CGTTCCCTGT GATGCCACGT TGACTGAGAA AAGCTTTGAA
      721 ACAGATATGA ACCTAAACTT TCAAAACCTG TCAGTTATGG GACTCCGAAT CCTCCTGCTG
      781 AAAGTAGCCG GATTTAACCT GCTCATGACG CTGAGGCTGT GGTCCAGTTG A

TRANSLATION
      MRPGTCSVLVLLLMLRRSNGDGDSVTQKEGLVTLTEGLPVMLNCTYQTIY
      SNAFLFWYVHYLNESPRLLLKSSTDNKRTEHQGFHATLHKSSSSFHLQKS
      SAQLSDSALYYCALSDRGRASSSFSKLVFGQGTSLSVVPNIQNPEPAVYQ
      LKDPRSQDSTLCLFTDFDSQINVPKTMESGTFITDKTVLDMKAMDSKSNG
      AIAWSNQTSFTCQDIFKETNATYPSSDVPCDATLTEKSFETDMNLNFQNL
      SVMGLRILLLKVAGFNLLMTLRLWSS*
//
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TCR beta 8 chain

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LOCUS       TCR beta    906 bp
ACCESSION   TCR beta 8
KEYWORDS    L + recombined V beta 8.1 D beta 1 J beta 1.1 + C beta
SOURCE      CD8+ T cell clone UZ3/4.
  ORGANISM  mus musculus

REFERENCE   1  (bases 1 to 906)
  AUTHORS   Self
  JOURNAL   Unpublished.
FEATURES             Location/Qualifiers
BASE COUNT      234 a    226 c    245 g    201 t
ORIGIN
        1 ATGGGCTCCA GACTCTTCTT TGTGGTTTTG ATTCTCCTGT GTGCAAAACA CATGGAGGCT
       61 GCAGTCACCC AAAGTCCAAG AAGCAAGGTG GCAGTAACAG GAGGAAAGGT GACATTGAGC
      121 TGTCACCAGA CTAATAACCA TGACTATATG TACTGGTATC GGCAGGACAC GGGGCATGGG
      181 CTGAGGCTGA TCCATTACTC ATATGTCGCT GACAGCACGG AGAAAGGAGA TATCCCTGAT
      241 GGGTACAAGG CCTCCAGACC AAGCCAAGAG AATTTCTCTC TCATTCTGGA GTTGGCTTCC
      301 CTTTCTCAGA CAGCTGTATA TTTCTGTGCC AGCAGTGTAG ACAGAACAGA AGTCTTCTTT
      361 GGTAAAGGAA CCAGACTCAC AGTTGTAGAG GATCTGAGAA ATGTGACTCC ACCCAAGGTC
      421 TCCTTGTTTG AGCCATCAAA AGCAGAGATT GCAAACAAAC AAAAGGCTAC CCTCGTGTGC
      481 TTGGCCAGGG GCTTCTTCCC TGACCACGTG GAGCTGAGCT GGTGGGTGAA TGGCAAGGAG
      541 GTCCACAGTG GGGTCAGCAC GGACCCTCAG GCCTACAAGG AGAGCAATTA TAGCTACTGC
      601 CTGAGCAGCC GCCTGAGGGT CTCTGCTACC TTCTGGCACA ATCCTCGAAA CCACTTCCGC
      661 TGCCAAGTGC AGTTCCATGG GCTTTCAGAG GAGGACAAGT GGCCAGAGGG CTCACCCAAA
      721 CCTGTCACAC AGAACATCAG TGCAGAGGCC TGGGGCCGAG CAGACTGTGG GATCACTTCA
      781 GCATCCTATC AGCAGGGGGT TCTGTCTGCA ACCATCCTCT ATGAGATCCT AGGGAAGGCC
      841 ACCCTATATG CTGTGCTGGT CAGTGGCCTG GTGCTGATGG CCATGGTCAA GAAAAAAAAT
      901 TCCTGA

TRANSLATION
      MGSRLFFVVLILLCAKHMEAAVTQSPRSKVAVTGGKVTLSCHQTNNHDYM
      YWYRQDTGHGLRLIHYSYVADSTEKGDIPDGYKASRPSQENFSLILELAS
      LSQTAVYFCASSVDRTEVFFGKGTRLTVVEDLRNVTPPKVSLFEPSKAEI
      ANKQKATLVCLARGFFPDHVELSWWVNGKEVHSGVSTDPQAYKESNYSYC
      LSSRLRVSATFWHNPRNHFRCQVQFHGLSEEDKWPEGSPKPVTQNISAEA
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      S*
//
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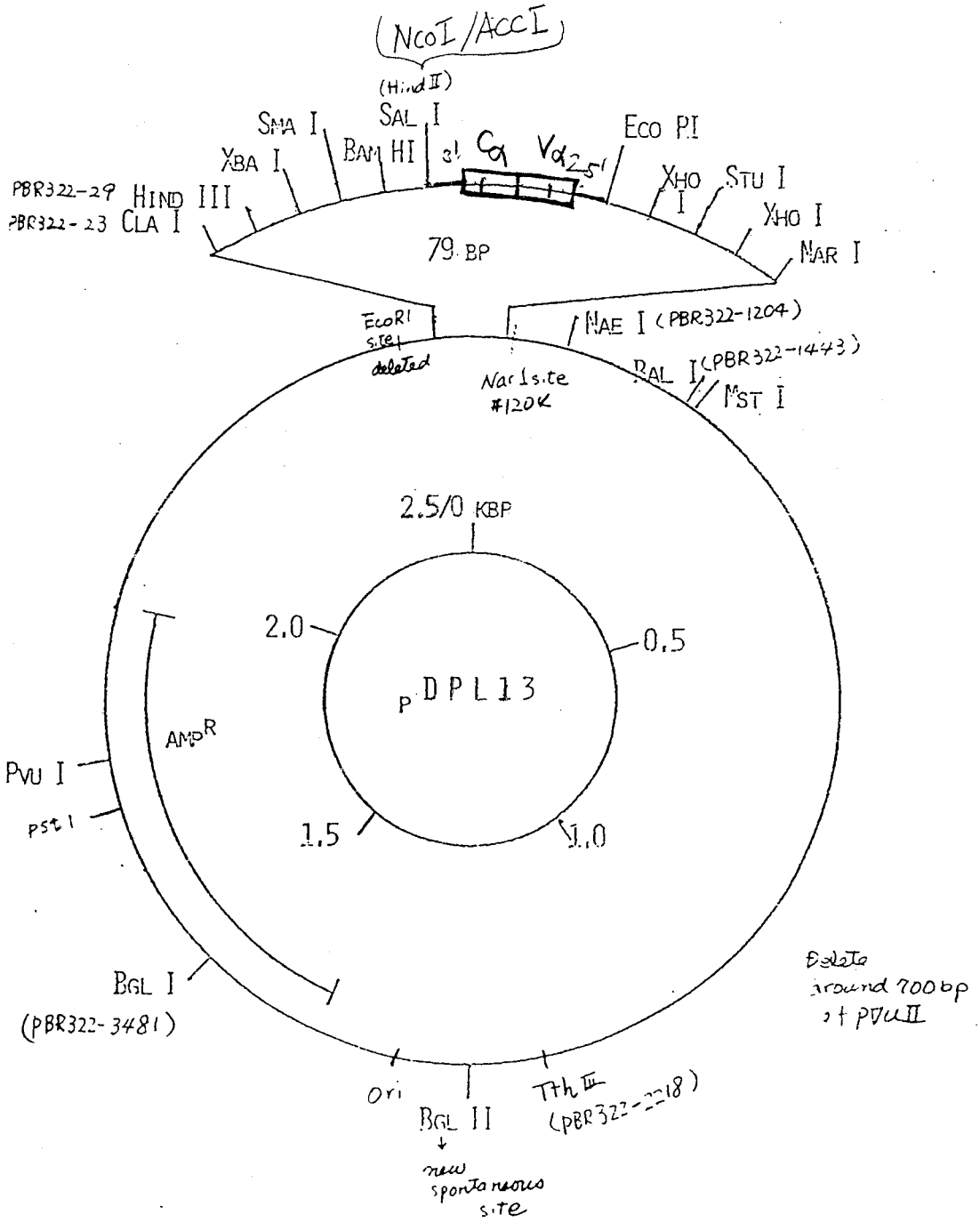
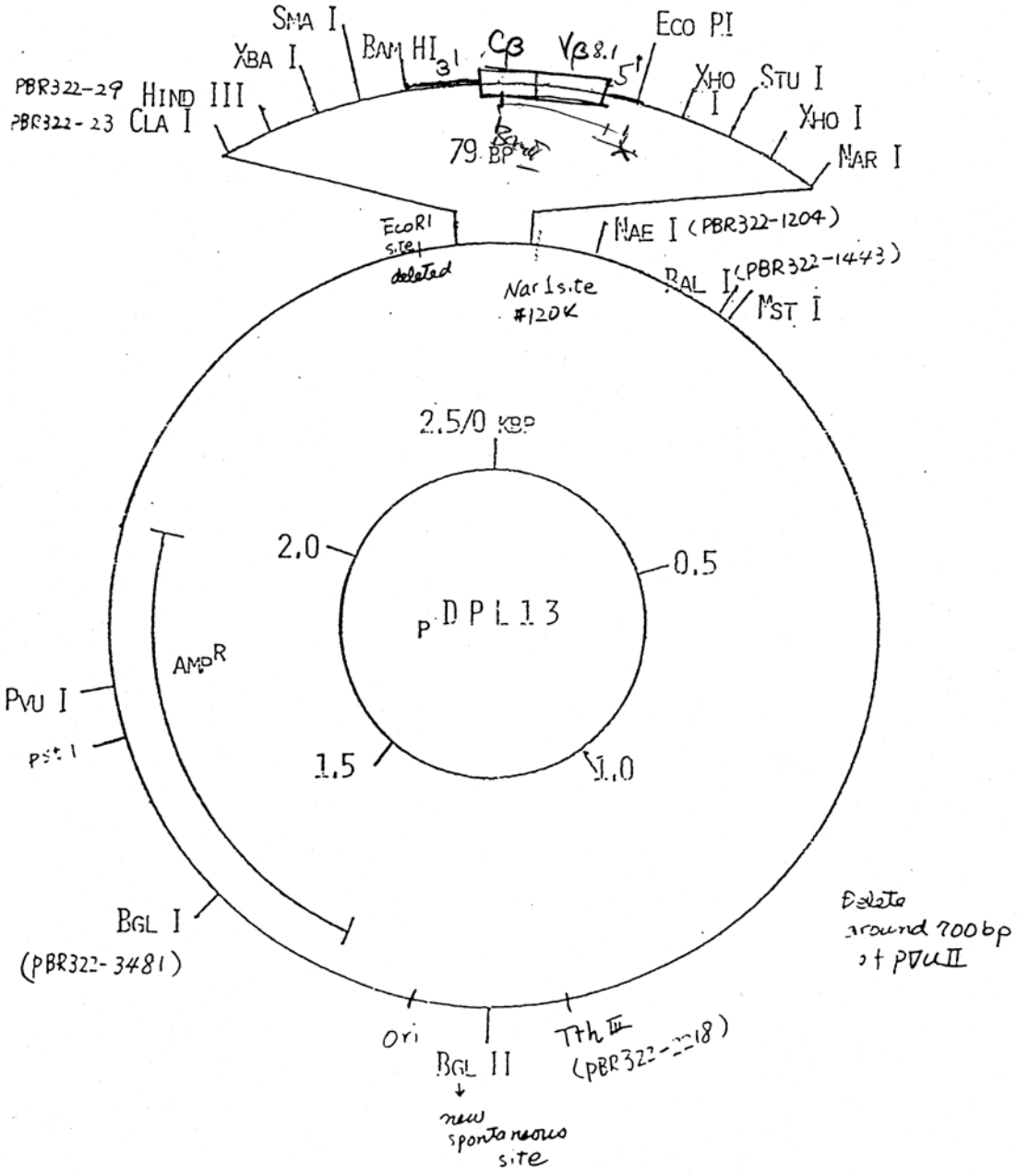
B. Plasmid charts

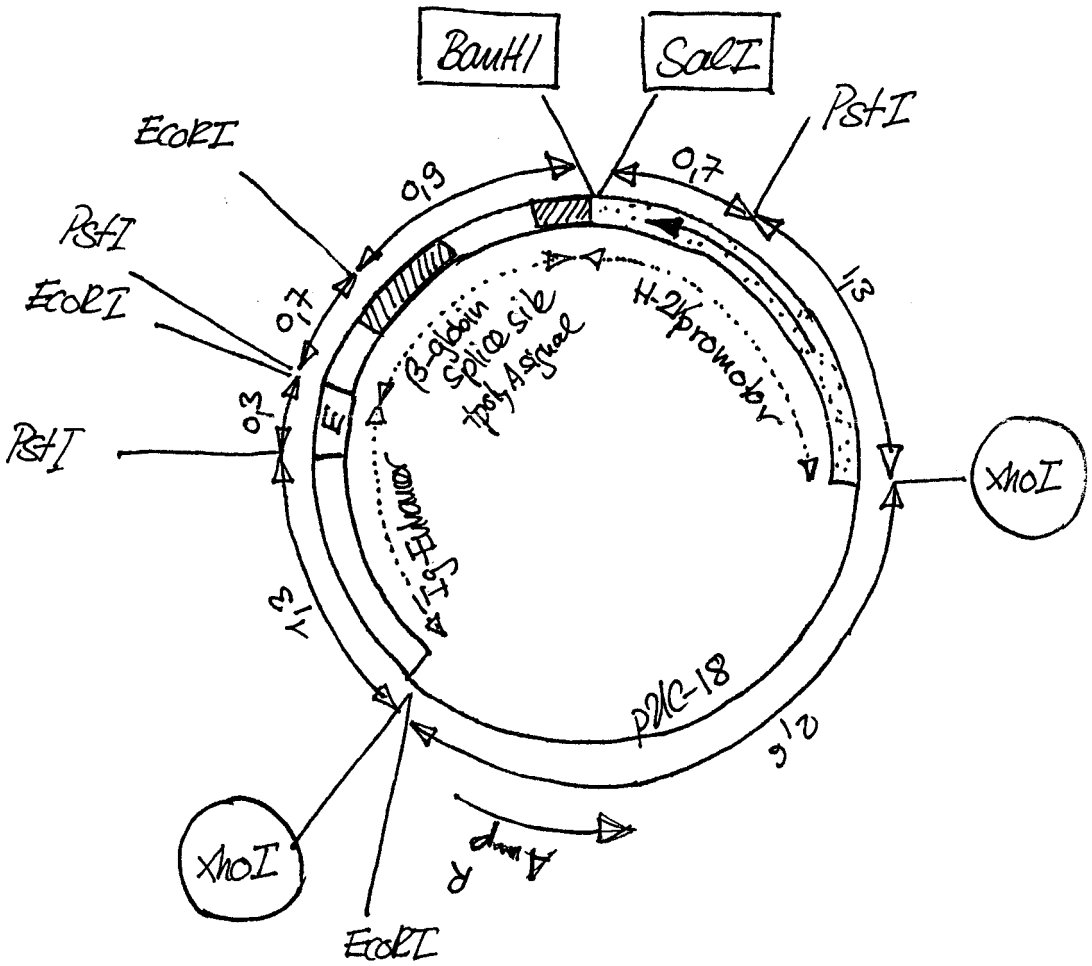
pP14αAR



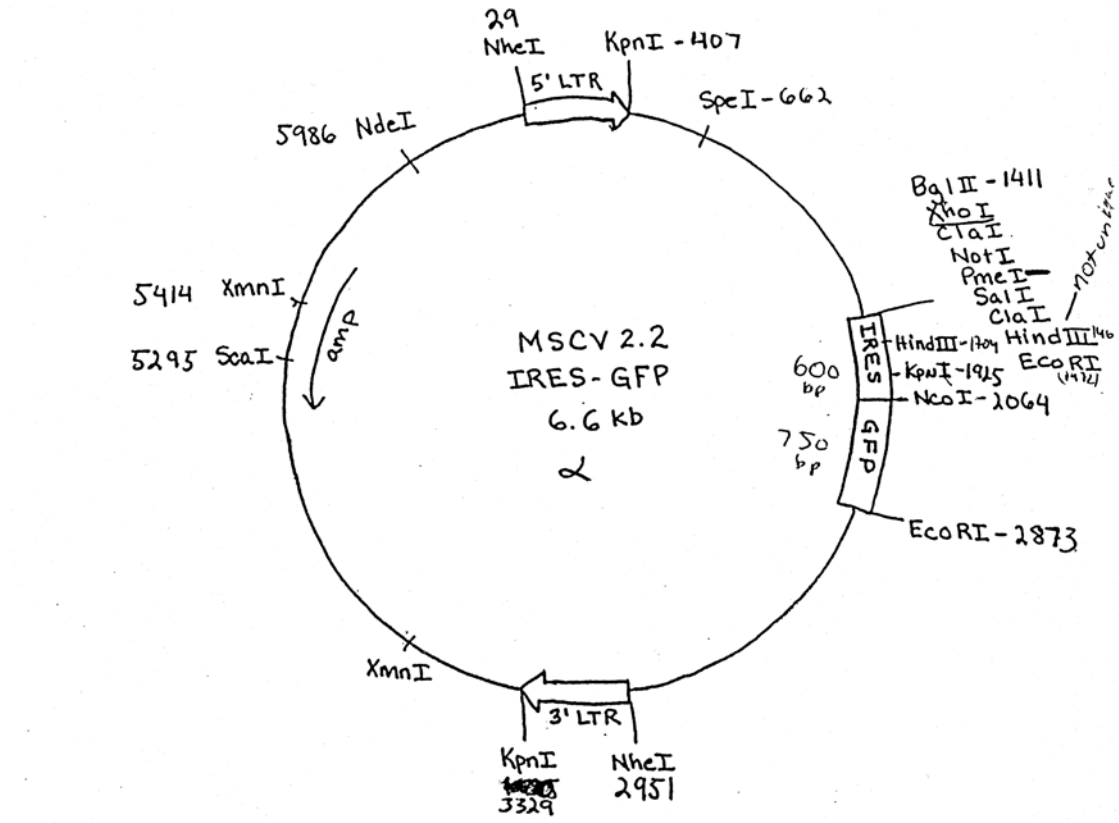
pP142β8AR

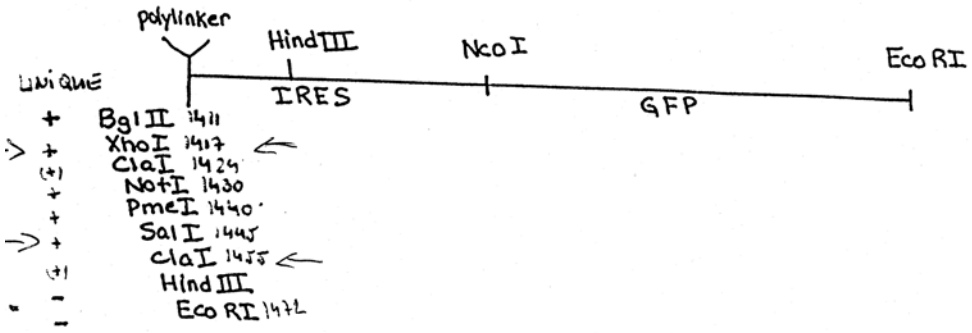

SMA I
BAM HI
Cβ
Vβ8.1
ECO PI
XBA I
PBR322-29 HIND III
PBR322-23 CLA I
XHO I
STU I
XHO I
NAR I
79 BP
EcoRI site deleted
NAE I (PBR322-1204)
Nar I site #120K
BAL I (PBR322-1443)
MST I
2.5/0 KBP
2.0
0.5
pDPL13
AMPR
1.5
1.0
PVU I
PST I
BGL I
(PBR322-3481)
Delete around 700bp of PVU II
ori
BGL II
new spontaneous site
Tth III (PBR322-2218)

pHSE3'

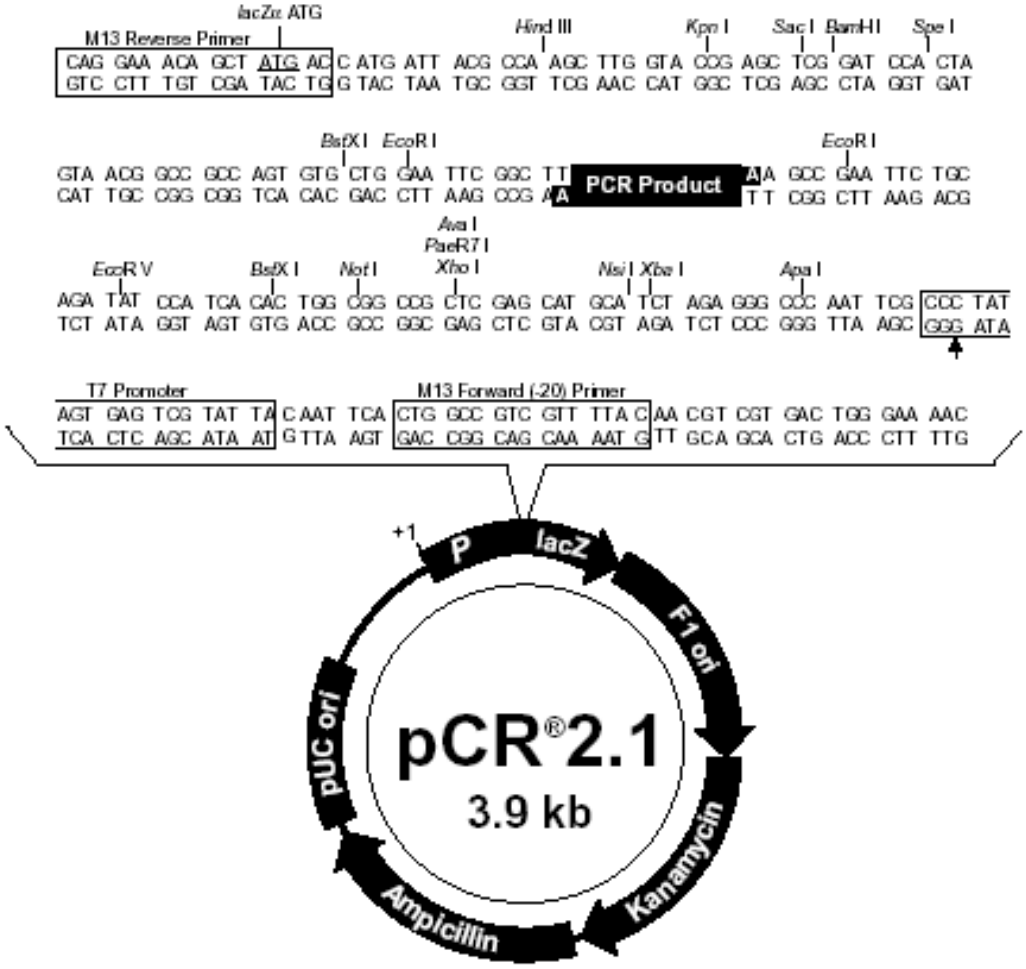

BamHI
SalI
PstI
EcoRI
PstI
EcoRI
PstI
XhoI
XhoI
EcoRI
0,9
0,7
0,7
0,3
1,3
1,3
0,7
1,3
0,6
E
β-globin splice site
poly A signal
H-2Kpromoter
pUC-18
Amp R

pMSCV2.2-IRES-GFP





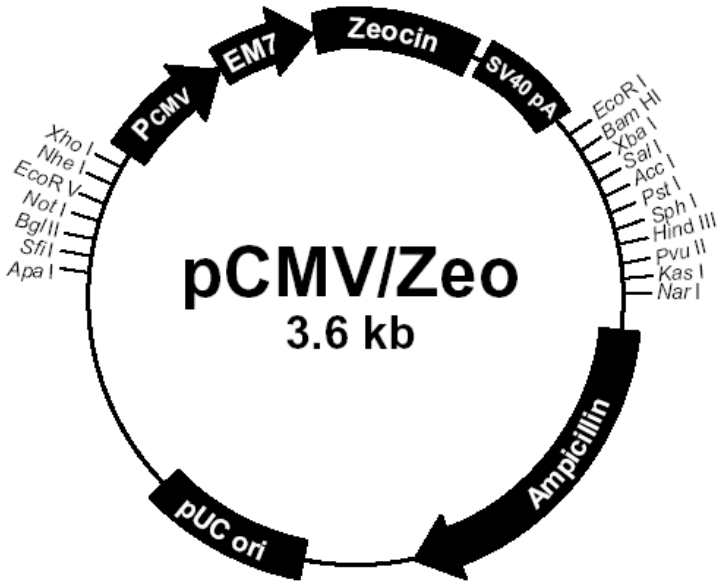
pCR2.1



Comments for pCR®2.1
3929 nucleotides

LacZα gene: bases 1-545
M13 Reverse priming site: bases 205-221
Multiple Cloning Site: bases 234-355
T7 promoter: bases 362-381
M13 (-20) Forward priming site: bases 389-404
f1 origin: bases 546-983
Kanamycin resistance ORF: bases 1317-2111
Ampicillin resistance ORF: bases 2129-2989
pUC origin: bases 3134-3807

pCMV/Zeo



Comments for pCMV/Zeo
3595 nucleotides

bla promoter (-10 region): bases 160-166
Ribosome binding site: bases 189-193
Beta-lactamase ORF: bases 201-1061
pUC origin: bases 1206-1879
5´ Polylinker: bases 1887-1937
CMV promoter: bases 1941-2449
EM7 promoter: bases 2565-2631
Sh ble ORF: bases 2632-3006
SV40 polyadenylation sequence: bases 3020-3101
3´ Polylinker: bases 3150-3359

C. Suppliers

Supplier	Location	URL
Amersham Biosciences Europe GmbH Amersham Biosciences Inc.	Munzinger Str. 9 D-79111 Freiburg; 800 Centennial Ave. Piscataway , NJ 08855-1327	www.amershambiosciences.com
Avidity LLC	1899 Gaylord St. Denver , CO 80206	www.avidity.com
BD Biosciences – PharMingen Clontech	Tullastr. 8-12 D-69126 Heidelberg; 10975 Torreyana Rd. San Diego , CA 92121	www.bdbiosciences.com
Bio-Rad Laboratories - Life Science Group Div.	Heidemannstr. 164 D-80937 München; 2000 Alfred Nobel Dr. Hercules , CA 94547	www.discover.bio-rad.com
GeneCraft ®	Tresckowstr. 10 D-48163 Münster	www.genecraft.de
ICN Biomedicals Inc.	Thüringer Str. 15 D-37269 Eschwege; 3300 Hyland Ave. Costa Mesa , CA 92626	www.icnbiomed.com
Invitrogen Corporation: Comprising products from Invitrogen, NOVEX, and Gibco BRL	Emmy-Noether-Str. 10 D-76131 Karlsruhe 1600 Faraday Avenue Carlsbad , CA 92008	www.invitrogen.com
Metabion GmbH	Am Klopferspitz 19 D-82152 Martinsried	www.metabion.com

Supplier	Location	URL
Mobitec	Lotzestr. 22a	www.probes.com
	D-37083 Göttingen	
Molecular Probes Inc.	4849 Pitchford Ave	
	Eugene , OR 97405-9165	
New England Biolabs	Bruningstr. 50	www.neb.com
GmbH	D-65926 Frankfurt a. M.	
New England Biolabs Inc.	32 Tozer Rd.	
	Beverly , MA 01915-5599	
QIAGEN GmbH	Max-Volmer-Str. 4	www.qiagen.com
	D-40724 Hiden	
QIAGEN	28159 Avenue Stanford	
	Valencia , CA 91355	
Roche Molecular	Sandhofer Str. 116	www.roche-applied-science.com
Biochemicals	D-68305 Mannheim	
Sigma-Aldrich	Eschenstr. 5	www.sigma-aldrich.com
	D-82024 Taufkirchen	
	3050 Spruce St.	
	St. Louis , MO 63103	
Stratagene Europe	Hogehilweg 15	www.stratagene.com
	NL-1101 CB Amsterdam	
Stratagene	11011 N. Torrey Pines Rd.	
	La Jolla , CA 92037	

D. Reprint

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