Sequence file available at www.neb.com See page 165 for ordering information.

There are no restriction sites for the following enzymes: AarI(x), Acc65I, AfeI, AflII, AgeI, AleI, AvrII, BaeI, BbsI, BbvCI, BclI, BglII, BlpI, BmgBI, Bpu10I, BseRI, BsiWI, BspDI, BstEII, BstXI, BstZ17I, Bsu36I, ClaI, CspCI, EcoNI, FspAI(x), I-CeuI, I-SceI, KpnI, MscI, NruI, NsiI, PI-PspI, PI-SceI, PmeI, PmII, SbfI, SexAI, SfiI, SgrAI, SmaI, SnaBI, SpeI, SrfI(x), SwaI, TspMI,

(x) = enzyme not available from NEB

pNEBR-X1Hygro is a plasmid cloning vector capable both of replication in E. coli and stable transfection of mammalian cells. It is designed for inducible expression of recombinant proteins in mammalian cells using the RheoSwitch Mammalian Inducible Expression System (NEB #E3000).

In E. coli, it replicates using the pMB1 origin of replication from pBR322 (although the rop gene is missing) and carries the bla (Ap^R) marker for selection with ampicillin. It also carries the hpt (HygR) marker under control of the thymidine kinase promoter; thus, following transfection into mammalian cells, it can be used to form stable cell lines by selection with hygromycin.

The multiple cloning site (MCS) is positioned downstream of a promoter apparatus consisting of 5 tandem copies of the yeast GAL4 response element (5XRE) followed by a minimal TATA box and a short leader sequence, and upstream of the SV40 polyadenylation (polyA) sequence. When co-transfected in mammalian cells with pNEBR-R1 (which encodes the RheoReceptor-1 protein), expression from the GAL4-derived promoter can be induced by the synthetic RSL1 ligand and controlled in a RSL1 concentration-dependent manner. A transcription terminafor the secreted Gaussia princeps luciferase) (1) cloned into the HindIII-NotI sites of pNEBR-X1.

Enzymes with unique restriction sites are shown in **bold** type. Location of sites of all NEB restriction enzymes can be found on the NEB web site (choose Technical Reference > DNA Sequences and Maps). Restriction site coordinates refer to the position of the 5'-most base on the top strand in each recognition sequence.

Open reading frame (ORF) coordinates are in the form "translational start – translational stop"; numbers refer to positions on the top (clockwise) strand, regardless of the direction of transcription and include the start and stop codons. Components of coordinated regions are indented below the region itself.

pMB1 origin of replication coordinates include the region from the -35 promoter sequence of the RNAII transcript to the RNA/DNA switch point. SV40 transcription terminator and polyA coordinates represent cloned regions and not necessarily the precise functional elements.

Coordinates

Source Tn3 S. hygroscopicus pMB1 SV40

S. cerevisiae

SV40

Feature

Stul 6044 Sali - Acci 6028 Sphi 6022 BsrGi 6015 BssHii 5995 Asci 5994	tor upstream of the 5XRE (not shown) prevents read-through transcription from other sources. pNEBR-X1 is identical to pNEBR-X1Hygro except that it does not contain the <i>hpt</i> (Hyg ⁿ) marker, so it is intended to be used for transient transfection. pNEBR-X1GLuc is a control plasmid with the reporter gene GLuc (the humanized coding sequence	bla (Ap ^R) 995-1855 aph-IV (hpt; Hyg ^R) 3672-2635 origin 4498-5086 SV40 TT region 5148-5606 expression region: 5X GAL4 RE 5743-5835 TATA box 5854-5860 transc. start 5889 (cw) MCS 5891-6049 SV40 polyA region 4-855	
Bmti - Nhel 5982 Noti 5976 PaeR7i - Tlii - Xhoi 5971 PspXi 5970 Fsel 5964 Pacl 5956 BamHi 5942 EcoRV 5935	PvuII 6153	(cw) = clockwise ori = origin of replication Ap = ampicillin Hyg = hygromycin RE = response element TT = transcriptional terminator	
HindIII 5921 Kasl - Nari - Sfoi 5915 Apal - PspOMI 5891		BsaBI 615	
SacI 5883 PciI 5142	pNEBR-X1 Hygro 6,268 bp	BcgI 1255	
BsaAI 4158		BspQI - SapI 2075	
BstBI 3808 — AatII - ZraI 364	'	PpuMI 2348 XcmI 2365	
References	SacII 2890 NAI - BspMI 3363		

References 435-443.