



MCS2														
NotI														
End of IRES2				EagI			BglII		BamHI		ClaI		SalI	
ATG	GCC	ACA	ACC	GCG	GCC	GCT	AGA	TCT	GGA	TCC	ATC	GAT	GTC	GAC
TAC	CGG	TGT	TGG	CGC	CGG	CGA	TCT	AGA	CCT	AGG	TAG	СТА	CAG	CTG
GAT	ATC	CAT	ATG	TCT	AGA									
	ATG TAC Ecc GAT	ATG GCC TAC CGG EcoRV GAT ATC	ATG GCC ACA   TAC CGG TGT   EcoRV NG   GAT ATC CAT	ATGGCCACAACCTACCGGTGTTGGEcoRVNdelGATATCCATATG	ATG GCC ACA ACC GCG     TAC CGG TGT TGG CGC     EcoRV   Ndel   Xb     GAT ATC CAT ATG TCT	End of IRES2 EagI   ATG GCC ACA ACC GCG GCC   TAC CGG TGT TGG CGC CGG   EcoRV NdeI XbaI   GAT ATC CAT ATG TCT AGA	End of IRES2 NotI   ATG GCC ACA ACC GCG GCC GCT   TAC CGG TGT TGG CGC CGC CGT   EcoRV Ndel Xbal   GAT ATC CAT ATG TCT AGA	NotI   End of IRES2 EagI Bg   ATG GCC ACA ACC GCG GCC GCT AGA   TAC CGG TGT TGG CGC CGG CGA TCT   EcoRV NdeI Xbal   GAT ATC CAT ATG TCT AGA	End of IRES2 NotI   ATG GCC ACA ACC GCG GCC GCT AGA TCT   TAC CGG TGT TGG CGC CGG CGA TCT AGA   EcoRV Ndel Xbal   GAT ATC CAT ATG TCT AGA	NotI   NotI   End of IRES2 NotI   ATG GCC ACA ACC GCG GCC AGA TCT GGA   TAC CGG TGT TGG CGC CGG CGA TCT AGA CCT   EcoRV NdeI XbaI TCT AGA GGA	NotI   End of IRES2 EagI BgIII BamHI   ATG GCC ACA ACC GCG GCC AGA TCT GGA TCC   TAC CGG TGT TGG CGC CGG CGA TCT AGA CCT AGG   EcoRV NdeI XbaI XbaI CAT ATG TCT AGA	End of IRES2 NotI   ATG GCC ACA   ACC GCG GCC   GCG TGT TGG   CGC CGC CGA   TAC CGG TGT   CGC CGC CGA   TAC CGG TGT   CGAT ATG Xbal   GAT ATC CAT	End of IRES2 Notl   ATG GCC ACA ACC GCG GCC GCT AGA TCT GGA TCC ATC GAT   TAC CGG TGT TGG CGC CGC GCA TCT AGA TCT AGG TCT AGG TCT AGG TCT AGG TCT AGG TAG CTA   EcoRV Ndel Xbal TCT AGA GGA TCT AGA CT AGG	End of IRES2   NotI   Ball   BamHI   ClaI   Same Same Same Same Same Same Same Same

pTRE-Dual1 Vector Map and Multiple Cloning Sites (MCS1 and MCS2). The internal start site (ATG) at the IRES2/ MCS2 junction is indicated in bold.



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## Description

pTRE-Dual1 is a tetracycline (Tet)-regulatable, mammalian expression vector designed to coexpress two genes of your choice under the control of  $P_{\text{Tight}}$ , a modified Tet-responsive promoter.  $P_{\text{Tight}}$  consists of a modified minimal CMV promoter, and seven direct repeats of a 36 bp regulatory sequence that contains the 19 bp tet operator sequence (*tetO*; 1). This vector is designed to be used with our Tet-On<sup>®</sup> Advanced and Tet-Off<sup>®</sup> Advanced Inducible Gene Expression Systems (Cat. Nos. 630930 and 630934). These systems provide the inducible gene expression strategy of Gossen & Bujard, with major improvements described by Urlinger, *et al.* (2–6).

pTRE-Dual1 allows inducible co-expression of two genes cloned into multiple cloning sites 1 and 2 (MCS1 and MCS2), respectively. An encephalomyocarditis virus (EMCV) internal ribosome entry site (IRES2), positioned between the two MCSs, facilitates cap-independent translation of the gene cloned into MCS2, from an internal start site at the IRES2/MCS2 junction (7). The vector also contains a ColE1 origin of replication and an ampicillin resistance gene (Amp<sup>r</sup>) to allow for propagation and selection in *E. coli*.

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### Use

pTRE-Dual1 is a mammalian expression vector that allows tightly regulated, doxycycline-controlled coexpression of two genes of your choice. Each gene must have both a start and a stop codon. For enhanced expression, the gene cloned into MCS2 should also be cloned in-frame with the start codon at the IRES2/ MCS2 junction (this codon is shown in bold in the MCS2 sequence on page 1).

In order to function, the system requires the presence of a tetracycline-controlled transcriptional activator (Tet-On Advanced or Tet-Off Advanced), supplied by a stable Tet-On Advanced or Tet-Off Advanced cell line that can be created with our Tet-On Advanced or Tet-Off Advanced Inducible Gene Expression Systems (Cat. Nos. 630930 and 630934).

## Location of features

- P<sub>Tight</sub> (modified Tet-responsive promoter): 8–321
- MČS1 (multiple cloning site 1): 323–366
- IRES2 (encephalomyocarditis virus internal ribosome entry site): 367-951
- MCS2 (multiple cloning site 2): 955–1005
- SV40 polyA signal: 1012-1194
- ColE1 origin of replication: 1370–1794
- Amp<sup>r</sup> (ampicillin resistance gene; β-lactamase): 1956–2879 (complementary)

# Propagation in *E. coli*

- Recommended host strain: DH5 $\alpha^{\text{TM}}$ , HB101, and other general purpose strains.
- Selectable marker: plasmid confers resistance to ampicillin (100 µg/ml) in *E. coli* hosts.
- E. coli replication origin: ColE1
- Plasmid incompatibility group: pMB1/CoIE1

#### References

- 1. pTRE-Tight Vectors (April 2003) *Clontechniques* XVIII(3):13–14.
- 2. Gossen, M. & Bujard, H. (1992) Proc. Natl. Acad. Sci USA 89(12):5547-5551.
- 3. Gossen, M., et al. (1995) Science 268(5218):1766–1769.
- 4. Urlinger, S. et al. (2000) Proc. Natl. Acad. Sci. USA 97(14):7963-7968.
- 5. Inducible Gene Expression Systems (January 2007) Clontechniques XXII(1):1-2.
- 6. Tet-On Advanced Inducible Gene Expression System (2006) Clontechniques XXI(2):1-3.
- 7. Jang, S. K. et al. (1988) J. Virol. 62(8):2636–2643.

**Note**: The vector sequence was compiled from information in the sequence databases, published literature, and other sources, together with partial sequences obtained by Clontech. This vector has not been completely sequenced.

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