

## Product datasheet for SC205715

## DcR1 (TNFRSF10C) (NM\_003841) Human 3' UTR Clone

## **Product data:**

## OriGene Technologies, Inc.

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Product Type:	3' UTR Clones
Product Name:	DcR1 (TNFRSF10C) (NM_003841) Human 3' UTR Clone
Symbol:	DcR1
Synonyms:	CD263; DCR1; DCR1-TNFR; LIT; TRAIL-R3; TRAILR3; TRID
Mammalian Cell Selection:	Neomycin
Vector:	pMirTarget (PS100062)
ACCN:	NM_003841
Insert Size:	444 bp
Insert Sequence:	<pre>&gt;SC205715 3'UTR clone of NM_003841 The sequence shown below is from the reference sequence of NM_003841. The complete sequence of this clone may contain minor differences, such as SNPs. Blue=Stop Codon Red=Cloning site GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAGGCCAAGAAGGGCGGAAAGATCGCCGTG TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC CTAATTGTGCTTCTGATTGTGTTTGTTTGAAAGACTTCACTGTGGAAGAAATTCCTTCC</pre>
<b>Restriction Sites:</b>	Sgfl-Mlul
OTI Disclaimer:	Our molecular clone sequence data has been matched to the sequence identifier above as a point of reference. Note that the complete sequence of this clone is largely the same as the reference sequence but may contain minor differences , e.g., single nucleotide polymorphisms (SNPs).
Components:	The cDNA clone is shipped in a 2-D bar-coded Matrix tube as 10 ug dried plasmid DNA. The package also includes 100 pmols of both the corresponding 5' and 3' vector primers in separate vials.



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	DcR1 (TNFRSF10C) (NM_003841) Human 3' UTR Clone – SC205715
RefSeq:	<u>NM 003841.5</u>
Summary:	The protein encoded by this gene is a member of the TNF-receptor superfamily. This receptor contains an extracellular TRAIL-binding domain and a transmembrane domain, but no cytoplasmic death domain. This receptor is not capable of inducing apoptosis, and is thought to function as an antagonistic receptor that protects cells from TRAIL-induced apoptosis. This gene was found to be a p53-regulated DNA damage-inducible gene. The expression of this gene was detected in many normal tissues but not in most cancer cell lines, which may explain the specific sensitivity of cancer cells to the apoptosis-inducing activity of TRAIL. [provided by RefSeq, Jul 2008]
Locus ID:	8794
MW:	16.4

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