

Product datasheet for SC200927

DcR3 (TNFRSF6B) (NM_032945) Human 3' UTR Clone

Product data:

Product Type: 3' UTR Clones **Product Name:** DcR3 (TNFRSF6B) (NM 032945) Human 3' UTR Clone Symbol: DcR3 DCR3; decoy receptor 3; DJ583P15.1.1; M68; OTTHUMP00000031583; TR6; tumor necrosis Synonyms: factor receptor superfamily, member 6b; tumor necrosis factor receptor superfamily, member 6b, decoy Mammalian Cell Neomycin Selection: Vector: pMirTarget (PS100062) ACCN: NM 032945 **Insert Size:** 151 bp >SC200927 3'UTR clone of NM_032945 **Insert Sequence:** The sequence shown below is from the reference sequence of NM_032945. The complete sequence of this clone may contain minor differences, such as SNPs. Blue=Stop Codon Red=Cloning site GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAGGCCAAGAAGGGCGGAAAGATCGCCGTG TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC CACTTGCACTGAAAGAGGCTTTTTTTTTTTAAATAGAAGAAATGAGGTTTCTTAAAGCTTATTTTTATAAAG CTTTTTCATAAAA CGAGATTTCGATTCCACCGCCGCCTTCTATGAAAGG **Restriction Sites:** 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. RefSeq: NM 032945.2



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Summary:	This gene belongs to the tumor necrosis factor receptor superfamily. The encoded protein is postulated to play a regulatory role in suppressing FasL- and LIGHT-mediated cell death. It acts as a decoy receptor that competes with death receptors for ligand binding. Over- expression of this gene has been noted in gastrointestinal tract tumors. Read-through transcription into this gene from the neighboring upstream gene, which encodes regulator of telomere elongation helicase 1 (RTEL1), generates a non-coding transcript. [provided by RefSeq, Feb 2011]
Locus ID:	8771
MW:	6.1

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