

## Product datasheet for **SC202389**

### CD163L1 (NM\_174941) Human 3' UTR Clone

#### Product data:

Product Type:	3' UTR Clones
Product Name:	CD163L1 (NM_174941) Human 3' UTR Clone
Vector:	pMirTarget (PS100062)
Symbol:	CD163L1
Synonyms:	CD163B; M160; SCARI2; WC1
ACCN:	NM_174941
Insert Size:	225 bp
Insert Sequence:	>SC202389 3'UTR clone of NM_174941 The sequence shown below is from the reference sequence of NM_174941. The complete sequence of this clone may contain minor differences, such as SNPs. Blue=Stop Codon Red=Cloning site  GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAAGCCAAGAAGGGCGGAAAGATCGCCGTG TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC GTTCTTCCTGCCTCTGAAGCCACAAAA <b>TG</b> ACTTTAGACTTCCAGGGCTCACCAGATCAACCTCTAAATA TCTTTGAAGGAGACAACAACCTTTTAAATGAATAAAGAGGAAGTCAAGTTGCCCTATGAAAACTTGTC AAATAACATTTCTTGAACAATAGGAGAACAGCTAAATTGATAAAGACTGGTGATAATAAAAAATTGAATT ATGTATATCACTGTAAA <b>ACGCGT</b> AAGCGGCCGCGCATCTAGATTGAAAGAAAATGACCGACCAAGCGACGCCCAACCTGCCATCA CGAGATTCGATTCCACCGCCCTTCTATGAAAGG
Restriction Sites:	Sgfl-MluI
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:	<u><a href="#">NM_174941.6</a></u>



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**Summary:** This gene encodes a member of the scavenger receptor cysteine-rich (SRCR) superfamily. Members of this family are secreted or membrane-anchored proteins mainly found in cells associated with the immune system. The SRCR family is defined by a 100-110 amino acid SRCR domain, which may mediate protein-protein interaction and ligand binding. The encoded protein contains twelve SRCR domains, a transmembrane region and a cytoplasmic domain. Alternative splicing results in multiple transcript variants encoding different isoforms. [provided by RefSeq, Jul 2014]

**Locus ID:** 283316

**MW:** 8.6