

Product datasheet for SC204609

LILRA5 (NM_021250) Human 3' UTR Clone

Product data:

OriGene Technologies, Inc.

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Product Type:	3' UTR Clones
Product Name:	LILRA5 (NM_021250) Human 3' UTR Clone
Symbol:	LILRA5
Synonyms:	CD85; CD85F; ILT-11; ILT11; LILRB7; LIR-9; LIR9
Mammalian Cell Selection:	Neomycin
Vector:	pMirTarget (PS100062)
ACCN:	NM_021250
Insert Size:	373 bp
Insert Sequence:	<pre>>SC204609 3'UTR clone of NM_021250 The sequence shown below is from the reference sequence of NM_021250. The complete sequence of this clone may contain minor differences, such as SNPs. Blue=Stop Codon Red=Cloning site GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAGGCCAAGAAGGGCGGAAAGATCGCCGTG TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC AGAAGCCCCCAAGCTGCAGAGTGGAAGGTGAACAGAAGAGAGAACAATGCACCATTGAATGCTGGAGCCT TGGAAGCCGCCAAGCTGCAGCTGGAAGGTGAACAGAAGAGAGAACAATGCACCATTGAATGCTGGAGCCT TGGAAGCGAATCTGATGGTCCTAGGAGGTTCGGGAAGACCATCTGAGGCCTATGCCATCTGGACTGTCT GCTGGCAATTCTTTTTTTTTT</pre>
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.



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	LILRA5 (NM_021250) Human 3' UTR Clone – SC204609
RefSeq:	<u>NM 021250.4</u>
Summary:	The protein encoded by this gene is a member of the leukocyte immunoglobulin-like receptor (LIR) family. LIR family members are known to have activating and inibitory functions in leukocytes. Crosslink of this receptor protein on the surface of monocytes has been shown to induce calcium flux and secretion of several proinflammatory cytokines, which suggests the roles of this protein in triggering innate immune responses. This gene is one of the leukocyte receptor genes that form a gene cluster on the chromosomal region 19q13.4. Four alternatively spliced transcript variants encoding distinct isoforms have been described. [provided by RefSeq, Jul 2008]
Locus ID:	353514
MW:	14.2

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