

Product datasheet for SC202754

CD130 (IL6ST) (NM_002184) Human 3' UTR Clone

Product data:

OriGene Technologies, Inc.

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Product Type:	3' UTR Clones
Product Name:	CD130 (IL6ST) (NM_002184) Human 3' UTR Clone
Symbol:	CD130
Synonyms:	CD130; CDW130; GP130; HIES4; IL-6RB; sGP130
Mammalian Cell Selection:	Neomycin
Vector:	pMirTarget (PS100062)
ACCN:	NM_002184
Insert Size:	232 bp
Insert Sequence:	>SC202754 3' UTR clone of NM_002184 The sequence shown below is from the reference sequence of NM_002184. The complete sequence of this clone may contain minor differences, such as SNPs. Red=Cloning site Blue=Stop Codon
	CAATTGGCAGAGCTCAGAATTCAAGCGATCGC
	TACCACAGACTGTACGGCAAGGCGGCTACATGCCTCAG TGA AGGACTAGTAGTTCCTGCTACAACTTCAG CAGTACCTATAAAGTAAAG
	ACGCGTAAGCGGCCGCGCATCTAGATTCGAAGAAAATGACCG
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:	<u>NM 002184.2</u>



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Summary:	The protein encoded by this gene is a signal transducer shared by many cytokines, including interleukin 6 (IL6), ciliary neurotrophic factor (CNTF), leukemia inhibitory factor (LIF), and oncostatin M (OSM). This protein functions as a part of the cytokine receptor complex. The activation of this protein is dependent upon the binding of cytokines to their receptors. vIL6, a protein related to IL6 and encoded by the Kaposi sarcoma-associated herpesvirus, can bypass the interleukin 6 receptor (IL6R) and directly activate this protein. Knockout studies in mice suggest that this gene plays a critical role in regulating myocyte apoptosis. Alternatively spliced transcript variants have been described. A related pseudogene has been identified on chromosome 17. [provided by RefSeq, May 2014]
Locus ID:	3572

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