

Product datasheet for SC208722

IL12RB1 (NM 153701) Human 3' UTR Clone

Product data:

Product Type: 3' UTR Clones

Product Name: IL12RB1 (NM_153701) Human 3' UTR Clone

Vector: pMirTarget (PS100062)

Symbol: IL12RB1

Synonyms: CD212; IL-12R-BETA1; IL12RB; IMD30

ACCN: NM_153701

Insert Size: 680 bp

Insert Sequence: >SC208722 3'UTR clone of NM_153701

The sequence shown below is from the reference sequence of NM_153701. The complete

sequence of this clone may contain minor differences, such as SNPs.

Blue=Stop Codon Red=Cloning site

GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAGGCCAAGAAGGGCGGAAAGATCGCCGTG

TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC

CAGCATTGGCTTGGCACATGCAGTTGGTACTCAATAAACGGCTGTTGCTATCCCCAGAA

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.



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RefSeq: <u>NM 153701.3</u>

Summary: The protein encoded by this gene is a type I transmembrane protein that belongs to the

hemopoietin receptor superfamily. This protein binds to interleukine 12 (IL12) with a low affinity, and is thought to be a part of IL12 receptor complex. This protein forms a disulfide-linked oligomer, which is required for its IL12 binding activity. The coexpression of this and IL12RB2 proteins was shown to lead to the formation of high-affinity IL12 binding sites and reconstitution of IL12 dependent signaling. Mutations in this gene impair the development of

interleukin-17-producing T lymphocytes and result in increased susceptibility to

mycobacterial and Salmonella infections. Alternative splicing results in multiple transcript

variants. [provided by RefSeq, Feb 2014]

Locus ID: 3594 MW: 25.6