

Product datasheet for SC210596

Ephrin A1 (EFNA1) (NM 182685) Human 3' UTR Clone

Product data:

Product Type: 3' UTR Clones

Product Name: Ephrin A1 (EFNA1) (NM_182685) Human 3' UTR Clone

Vector: pMirTarget (PS100062)

Symbol: EFNA1

Synonyms: B61; ECKLG; EFL1; EPLG1; GMAN; LERK-1; LERK1; TNFAIP4

ACCN: NM_182685

Insert Size: 862 bp

Insert Sequence: >SC210596 3'UTR clone of NM_182685

The sequence shown below is from the reference sequence of NM_182685. The complete

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

Blue=Stop Codon Red=Cloning site

GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAGGCCAAGAAGGGCGGAAAGATCGCCGTG

TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC

TCTGCCACTCCATATTAAAACATATGACCATTGA

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).



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Ephrin A1 (EFNA1) (NM_182685) Human 3' UTR Clone - SC210596

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 182685.2</u>

Summary: This gene encodes a member of the ephrin (EPH) family. The ephrins and EPH-related

receptors comprise the largest subfamily of receptor protein-tyrosine kinases and have been

implicated in mediating developmental events, especially in the nervous system and in

erythropoiesis. Based on their structures and sequence relationships, ephrins are divided into

the ephrin-A (EFNA) class, which are anchored to the membrane by a

glycosylphosphatidylinositol linkage, and the ephrin-B (EFNB) class, which are

transmembrane proteins. This gene encodes an EFNA class ephrin which binds to the EPHA2, EPHA4, EPHA5, EPHA6, and EPHA7 receptors. Two transcript variants that encode different

isoforms were identified through sequence analysis. [provided by RefSeq, Jul 2008]

Locus ID: 1942

MW: 31.9