

Product datasheet for **SC206839**

EVI2B (NM_006495) Human 3' UTR Clone

Product data:

Product Type:	3' UTR Clones
Symbol:	EVI2B
Synonyms:	CD361; D17S376; EVDB
Mammalian Cell Selection:	Neomycin
Vector:	pMirTarget (PSI00062)
ACCN:	NM_006495
Insert Size:	523 bp

Insert Sequence: >SC206839 3'UTR clone of NM_006495
The sequence shown below is from the reference sequence of NM_006495. The complete sequence of this clone may contain minor differences, such as SNPs.
Blue=Stop Codon **Red**=Cloning site

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GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAAGCCAAGAAGGGCGGAAAGATCGCCGTG
TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC
CTGCCACCTCCACCTGCAGAACTGTTAATAATTACAACTTGCTTTTAGCTGATCTTCCATCCTCAA
ATGACTCTTTTTCTTTATATGTTAACATATATAAAATGGCAACTGATAGTCAATTTTGATTTTATTC
AGGAACTATCTGAAATCTGCTCAGAGCCTATGTGCATAGATGAACTTTTTTAAAAAAGTTATTTA
ACAGTAATCTATTACTAATTATAGTACCTATCTTTAAAGTATAGTACATTTTACATATGTAATGGTA
TGTTTCAATAATTTAAGAACTCTGAAACAATCTACATATACTTATTACCCAGTACAGTTTTTTTCCCC
TGAAAAGCTGTGTATAAAATTATGGTGAATAAACTTTTATGTTTCCATTTCAAAGACCAGGGTGGAGAG
GAATAAGAGACTAAGTATATGCTTCAAGTTTAAATTAATACCTCAAGTATTAATAAATATTCCAAGT
TTGTGGGAATGGGAGATTAATGATGTTTGAGAAATAGA
ACGCGTAAGCGGCCGCGGCATCTAGATTGCAAGAAAATGACCGACCAAGCGACGCCAACCTGCCATCA
CGAGATTTTCGATTCCACCGCCGCTTCTATGAAAGG
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Restriction Sites: SgfI-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.
Note:	Plasmids are not sterile. For experiments where strict sterility is required, filtration with 0.22um filter is required.
RefSeq:	NM_006495.4
Summary:	Required for granulocyte differentiation and functionality of hematopoietic progenitor cells through the control of cell cycle progression and survival of hematopoietic progenitor cells. [UniProtKB/Swiss-Prot Function]
Locus ID:	2124
MW:	20.8