

Product datasheet for SC207465

STYXL2 (NM_001080426) Human 3' UTR Clone

Product data:

Product Type: 3' UTR Clones

Symbol: STYXL2

Synonyms: DUSP27

Mammalian Cell Selection: Neomycin

Vector: pMirTarget (PSI00062)

ACCN: NM_001080426

Insert Size: 587 bp

Insert Sequence: >SC207465 3'UTR clone of NM_001080426
The sequence shown below is from the reference sequence of NM_001080426. The complete sequence of this clone may contain minor differences, such as SNPs.
Blue=Stop Codon Red=Cloning site

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GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAAGCCAAGAAGGGCGGAAAGATCGCCGTG
TAACAATTGGCAGAGCTCAGAATTCAGCGATCGCC
ACCAAGCTGCAGAAAAGGAGGGAGGACTGAGCTGGGGAAAATCTGAGAACACTGAAAGAAACCACTCAC
GTTAGCATAGGGCTCAGGGCACACGTTGCCACCACTCATCGCAGGATGAGGATACAGAGAGGATCTTCC
AGAGGGGCAGAGCCAAATGAGAGGTACCAAGCATAAGGGCAGCAGAGGTGAGTAGGGAGGAGGCAAG
GAGGGGGAGAAACCATCAATACGAATACGAGGTCCGAATGCGGACCAACTGATACCATTTTCTGTTGCTC
AGCGCCCTCTAAGCTTTGGTGTTCACCTAATGTATTTGGCAGTGTTCATCACAGGCTAGAGAGGTGAG
CTTGAAAAGCACTGTAGTTTGTGAGAGCTCCAGTTTACATCCAGAAAGGCCATGAACATAGGACACG
CTTCTGTCTGTAGAGGCTTCATATGAGACCCAGAAAGTCTATCCTATGGCAAGTCTGACCTCTCCTGGC
AATGCTCAGTTCTGATTTTTTTTTTTTAAATGTTTGTGAGTCTCCATTAAAAATGGTATGTTGCAATA
GTTCTTCTTTATTCTGGTCATGTTTAACTTTTGG
ACGCGTAAGCGGCCCGGCATCTAGATTCTGAAGAAAATGACCGACCAAGCGACGCCAACCTGCCATCA
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_001080426.3
Summary:	May be required for myofiber maturation.[UniProtKB/Swiss-Prot Function]
Locus ID:	92235
MW:	22.3