

Product datasheet for **SC207395**

ESF1 (NM_016649) Human 3' UTR Clone

Product data:

Product Type:	3' UTR Clones
Symbol:	ESF1
Synonyms:	ABTAP; bA526K24.1; C20orf6; HDCMC28P
Mammalian Cell	Neomycin
Selection:	
Vector:	pMirTarget (PSI00062)
ACCN:	NM_016649
Insert Size:	579 bp

Insert Sequence: >SC207395 3'UTR clone of NM_016649
The sequence shown below is from the reference sequence of NM_016649. 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
CAAGCAAGAAAAAGCAAAAAGTCAAACTGGATGTTACTTATTTTTGAACTGAATACATCTTTTCC
TAAATGTACAAAAATAATAGGAGGGAATTTTATTGGGAACAAAGCTATCTTTCAAGAACATGAATAA
AATCTTTTCTGGACATAGTAAATTTTCTCCATAAATAATTGTACTTAATTGTGGATGACTGACAAA
TTTTTATTGTATATTCTACAGATCAGTCATAATTAAATTACCTGCATTATAGGGTTTATAAAATTTT
ATATTTTACAATGTTCACTTCTAACTAGTGAAAGTTACTCTAGCTTTTAAAAGGCTGTTTACAATTC
TGTGTAAGAAATAGAGCAGTATCTACTCAAGTTTGTGTAATGTTAGGGATAATTTGAAAAATATATATA
TTTAATACATTAATTTCTCTGGAAGCAGGAGGCATGTTTAAATACTATTTAAATAATTTATTTTCTA
GCCATAAAGGATGGAAGTCAAGAACTTTTGTGTTTAGTCATGTTAAGTATAGTTTATGAAATTAAT
TGTAATAAAAGTGTAATAATTTTCA
ACGCGTAAGCGGCCCGGCATCTAGATTGCAAGAAAATGACCGACCAAGCGACGCCAACCTGCCATCA
CGAGATTTGATTCCACCGCCGCTTCTATGAAAGG
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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:	<u>NM_016649.4</u>
Summary:	May constitute a novel regulatory system for basal transcription. Negatively regulates ABT1 (By similarity).[UniProtKB/Swiss-Prot Function]
Locus ID:	51575
MW:	22.9