

Product datasheet for **SC205317**

EAPP (NM_018453) Human 3' UTR Clone

Product data:

Product Type:	3' UTR Clones
Product Name:	EAPP (NM_018453) Human 3' UTR Clone
Symbol:	EAPP
Synonyms:	BM036; C14orf11
Mammalian Cell Selection:	Neomycin
Vector:	pMirTarget (PS100062)
ACCN:	NM_018453
Insert Size:	402 bp
Insert Sequence:	>SC205317 3'UTR clone of NM_018453 The sequence shown below is from the reference sequence of NM_018453. 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
TTTTTCAATGTTTTAGCAAGCCATTCCTAACAGCCCAACTGGCATTTAATTACCCAATACTGTATATA
AGGCAAATATGGACAGTTACTTTCTCTTGCCTGTTTCATATCCTTCAGTGACATTGAGGAAGCAGTGT
TCTCTTTTTAAAGGAGAATAGTTGTCAACCTTCATTCATCTCTTACATCTCTCACCTCTCCTTTTTTT
TTTCTTTGATTTTCCCCCTTATTGATGGGACTGATATTCTGTTTTTGGTGAACATTTGGAAACTG
TCGGGCTTTTTATTAAAGCTCTGTAGAATAAAATGTTCTGGAATTATAAGCAATCTTTGTTTCTGAGT
GTTATTTTTATTTGGATATAGCTTTGATGTAATTAACCTGGAAACTCTTTCCTGAA
ACGCGTAAGCGGCCGCGGCATCTAGATTGAAAGAAAATGACCGACCAAGCGACGCCCAACCTGCCATCA
CGAGATTCGATTCCACCGCCGCTTCTATGAAAGG
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Restriction Sites:	Sgfl-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.



[View online »](#)

RefSeq: [NM_018453.4](#)

Summary: This gene encodes a phosphoprotein that interacts with several members of the E2F family of proteins. The protein localizes to the nucleus, and is present throughout the cell cycle except during mitosis. It functions to modulate E2F-regulated transcription and stimulate proliferation. Alternative splicing results in multiple transcript variants. [provided by RefSeq, Jan 2016]

Locus ID: 55837

MW: 15.8