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Product datasheet for SC206746

TACC2 (NM_006997) Human 3' UTR Clone

Product data:

Product Type:	3' UTR Clones
Product Name:	TACC2 (NM_006997) Human 3' UTR Clone
Symbol:	TACC2
Synonyms:	AZU-1; ECTACC
Mammalian Cell Selection:	Neomycin
Vector:	pMirTarget (PS100062)
ACCN:	NM_006997
Insert Size:	516 bp
Insert Sequence:	>SC206746 3'UTR clone of NM_006997 The sequence shown below is from the reference sequence of NM_006997. The complete sequence of this clone may contain minor differences, such as SNPs. Blue=Stop Codon Red=Cloning site GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAGGCCAAGAAGGGCGGAAAGATCGCCGTG TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC GAACTGATTGCCAAAATGGGGAAAAGCTAACTCTGAACCGAATGTTTTGGACTTAACTGTTGCGTGCAA TATGACCGTCGGCACACTGCTGTTCCTCCAGTTCCATGGACAGGTTCTGTTTTCACTTTTTCGTATGCA CTACTGTATTTCCTTTCTAAATAAAATTGATTTGATTGTATGCAGTACTAAGGAGACTATCAGAATTTC TTGCTATTGGTTTGCATTTTCCTAGTATAATTCATAGCAAGTTGACCTCAGAGTTCCTGTATCAGGGAG ATTGTCTGATTCTCTAATAAAAGACACATTGCTGACCTTGGCCTTGCCCTTTGTACACAAGTTCCCAGG GTGAGCAGCTTTTGGATTTAATATGAACATGTACAGCGTGCATAGGGACTCTTGCCTTAAGGAGTGTAA ACTTGATCTGCATTTGCTGATTTGTTTTTAAAAAAACAAGAAATGCATGTTTCAAATAAAATTCTCTAT TGTAAATAAAATTTTTTCTTTGGATCTTGGCAA ACGCGTAAGCGGCCGCGGCATCTAGATTCGAAGAAAATGACCGACCAAGCGACGCCCAACCTGCCATCA CGAGATTTCGATTCCACCGCCGCCTTCTATGAAAGG
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).

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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:	NM_006997.4
Summary:	Transforming acidic coiled-coil proteins are a conserved family of centrosome- and microtubule-interacting proteins that are implicated in cancer. This gene encodes a protein that concentrates at centrosomes throughout the cell cycle. This gene lies within a chromosomal region associated with tumorigenesis. Expression of this gene is induced by erythropoietin and is thought to affect the progression of breast tumors. Several transcript variants encoding different isoforms have been found for this gene. [provided by RefSeq, Jul 2008]
Locus ID:	10579
MW:	19.8