

Product datasheet for **SC213753**

JKAMP (NM_016475) Human 3' UTR Clone

Product data:

Product Type:	3' UTR Clones
Product Name:	JKAMP (NM_016475) Human 3' UTR Clone
Vector:	pMirTarget (PS100062)
Symbol:	JKAMP
Synonyms:	C14orf100; C24orf100; CDA06; HSPC213; HSPC327; JAMP
ACCN:	NM_016475
Insert Size:	1368 bp
Insert Sequence:	>SC213753 3'UTR clone of NM_016475 The sequence shown below is from the reference sequence of NM_016475. The complete sequence of this clone may contain minor differences, such as SNPs. Blue=Stop Codon Red=Cloning site

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GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAAGCCAAGAAGGGCGGAAAGATCGCCGTG
TAACAATTGGCAGAGCTCAGAATTCAAACGATCGCC
ATACTCTCAGAAGGACCAATGGACACTGTAGTGTAGACATGTGAAATGCCAAAAACCTGAGAAGTGCTC
CTAATAAAAAAGTAAATCAATCTTAACAGTGTATGAGAATCTATTCTATCATATATGGGAACAAGATTGT
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AAACTCTGTAATACTCTGTTACACAGGGTAATATTATCTGCTACACTGGAAGGCCCTAGGAAGCCCTT
GCTTCTCTCAACAGTTCAGCTGTTCTTTAGGGCAAAATCATGTTTCTGTGTACCTAGCAATGTGTTCCC
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AACATAGAAATGATGTATATTGTTTTTGTATCTATTTATTTTCATCAATACAGTATTTTGATGTATT
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TTGTTCCGTGTAAGAAAACAATAATAAAAAGCTTACCTACATAAAATTTCAATGTTTGGACTTAATT
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TTCACAGCAGTTGTATCCTTGAGTTACTTTGTTAATGTATTTTCTCAGTACATTTAACCACTGGGAAA
TGAACCCCTGTACGAATGTGTTTCTTCTCTGTAGGAATAAAAAATAAATATAAAAAATTTTATTGT
ATTGCACACATTTGGGGGTTATTAGTGTTCAATAAAATTTCTGAGTTGCCCAAAGAA
ACGCGTAAGCGGCCGCGCATCTAGATTCGAAGAAAATGACCGACCAAGCGACGCCCAACCTGCCATCA
CGAGATTCGATTCCACCGCCGCTTCTATGAAAGG
  
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Restriction Sites:	Sgfl-Mlul
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.
RefSeq:	NM_016475.5
Summary:	May be a regulator of the duration of MAPK8 activity in response to various stress stimuli. Facilitates degradation of misfolded endoplasmic reticulum (ER) luminal proteins through the recruitment of components of the proteasome and endoplasmic reticulum-associated degradation (ERAD) system (By similarity).[UniProtKB/Swiss-Prot Function]
Locus ID:	51528
MW:	53.4