

Product datasheet for SC206399

MINDY1 (NM_001163260) Human 3' UTR Clone

Product data:

Product Type:	3' UTR Clones
Product Name:	MINDY1 (NM_001163260) Human 3' UTR Clone
Symbol:	MINDY1
Synonyms:	FAM63A; MINDY-1
Mammalian Cell Selection:	Neomycin
Vector:	pMirTarget (PS100062)
ACCN:	NM_001163260
Insert Size:	482 bp
Insert Sequence:	>SC206399 3'UTR clone of NM_001163260 The sequence shown below is from the reference sequence of NM_001163260. The complete sequence of this clone may contain minor differences, such as SNPs. Blue =Stop Codon Red =Cloning site
	GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAAGCCAAGAAGGGCGGAAAGATCGCCGTG TAACAATTGGCAGAGCTCAGAATTCAA GCGATCGCC AAGCACGAGTCAGACTGCATTCTGCTGT AG CTCTGCCCCAGTGCCAGGCTGGCCTGCCCTTCTTCCAG AGGCTATGGCTAGTTGGCTTGTCCCCGCCTCCACCCCTGAGATGTGCTGGATAACTTATTTATGGAC TGTTGGGGATGAGAGCAGGCAACAAATGCCAAGGTCAGACTTGGTAATGCCTTGACCTCACGTGCTGC TGCCTTCTGCCTCCCATCCAGGGCAACACTAGGATTGGTGGGTTTCTGGTTCTCAACTCCCGGTCCC TGAATAGTCACACGTATGTACAGACTGAGGCTCTGGGGTGAGGTCCTATCCAGAATGCATCTTCTG CTTCCCATCCCTGCTGCCTGGATGCTCCTGATCACCTAGGCAGGCTGTCTCCAGTTGTTTCAGAGCTT AATTTGGGTTTCTATCTTATTGTAATGCCTTCTGGGGTTTGGAAATAAACTTCTGGCCGGGCA ACGCGT AAGCGGCCGCGCATCTAGATTGGAAGAAAATGACCGACCAAGCGACGCCCAACCTGCCATCA CGAGATTCGATTCCACCGCCGCTTCTATGAAAGG
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.



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RefSeq: [NM_001163260.2](#)

Summary: Hydrolase that can specifically remove 'Lys-48'-linked conjugated ubiquitin from proteins. Has exodeubiquitinase activity and has a preference for long polyubiquitin chains. May play a regulatory role at the level of protein turnover.[UniProtKB/Swiss-Prot Function]

Locus ID: 55793

MW: 17.8