OriGene Technologies, Inc.
9620 Medical Center Drive, Ste 200
Rockville, MD 20850, US
Phone: +1-888-267-4436
https://www.origene.com
techsupport@origene.com
EU: info-de@origene.com
CN: techsupport@origene.cn

Product datasheet for SC203973

FAIM1 (FAIM) (NM_001033032) Human 3' UTR Clone

Product data:

Product Type:	3' UTR Clones
Product Name:	FAIM1 (FAIM) (NM_001033032) Human 3' UTR Clone
Symbol:	FAIM1
Synonyms:	FAIM1
Mammalian Cell Selection:	Neomycin
Vector:	pMirTarget (PS100062)
ACCN:	NM_001033032
Insert Size:	327 bp
Insert Sequence:	>SC203973 3'UTR clone of NM_001033032 The sequence shown below is from the reference sequence of NM_001033032. The complete sequence of this clone may contain minor differences, such as SNPs. Blue=Stop Codon Red=Cloning site GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAGGCCAAGAAGGGCGGAAAGATCGCCGTG TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC AATAGAGAAATCCCAGAGATTGCAAGTTAATGAATTTTCATCTTAAGAAGTAAAGATCAGGACTTTTTA ATTACTGTGGTAATTAAATGTGTTCAGTATGTACTTATCAGTACATTTAGTCTGCAATGTTTTAATTTT TTAAAAAGTTACATGAAACTAACATTCCAAGGGTCAGGAAAAAAACCAATTATGTATAGTCATAAAAAT TACAATTTATGATGCAAATAATGTAAAATGTTTTAAAGACAAATGGCAAATAAGATATGGACCAAAGTC ACTAATGTTTTACAACAGTAACCTTTACTATAATAAATACTTTTAAAAAAA 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).
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_001033032.2

View online »

Summary: The protein encoded by this gene protects against death receptor-triggered apoptosis and regulates B-cell signaling and differentiation. Several transcript variants encoding different isoforms have been found for this gene. [provided by RefSeq, Jul 2011]

Locus ID: 55179

MW: 12.9