

Product datasheet for SC310396

MAEL (NM_032858) Human Untagged Clone


Product data:

Product Type:	Expression Plasmids
Product Name:	MAEL (NM_032858) Human Untagged Clone
Tag:	Tag Free
Symbol:	MAEL
Synonyms:	CT128; SPATA35
Mammalian Cell Selection:	Neomycin
Vector:	pCMV6-Entry (PS100001)
E. coli Selection:	Kanamycin (25 ug/mL)
Fully Sequenced ORF:	>SC310396 representing NM_032858. Blue=Insert sequence Red=Cloning site Green=Tag(s)

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GCTCGTTTGTAGTGAACCGTCAGAATTTTGTAAACGACTCACTATAGGGCGCCGGGAATTCGTCGACTG
GATCCGGTACCGAGGAGATCTGCCGCCGCGATCGCC
ATGCCGAACCGTAAGGCCAGCCGGAATGCTTACTATTTCTTCGTGCAGGAGAAGATCCCCGAACACGG
CGACGAGGCCCTGCCTGTGGCTCGCGTTGCTGATGCCATCCCTTACTGCTCCTCAGACTGGGCGCTTCTG
AGGGAGGAAGAAAAGGAGAAATACGCAGAAATGGCTCGAGAATGGAGGGCCGCTCAGGAAAGGACCCT
GGGCCCTCAGAGAAGCAGAAACCTGTTTTACACCACTGAGGAGGCCAGGCATGCTTGTACCAAAGCAG
AATGTTTTACCTCCAGATATGTCAGCTTTGTCTTTAAAAGGTGATCAAGCTCTCCTTGGAGGCATTTTT
TATTTTTGAACATTTTTAGCCATGGCGAGCTACCTCCTCATTGTGAACAGCGCTTCCTCCCTTGTGAA
ATTGGCTGTGTTAAGTATTCTCTCCAAGAAGGTATTATGGCAGATTTCCACAGTTTTATAAATCCTGGT
GAAATTCACGAGGATTCGATTTTCATTGTGAGGCTGCAAGTATTCTAGTCACAAGATTCCTATTTCA
AATTTTGAACGTGGGCATAACCAAGCAACTGTGTTACAAAACCTTTATAGATTTATTCATCCCAACCCA
GGGAACCTGGCCACCTATCTACTGCAAGTCTGATGATAGAACCAGAGTCAACTGGTGTGTTGAAGCATATG
GCAAAGGCATCAGAAATCAGGCAAGATCTACAACCTTCTACTGTAGAGGACCTTGTAGTGGGGATCTAC
CAACAAAATTTCTCAAGGAGCCCTCTAAGACTTGGATTGCAAGCCTCCTAGATGTGGCCATGTGGGAT
TATTCTAGCAACACAAGGTGCAAGTGGCATGAAGAAAATGATATTCTTCTGTGCTTTAGCTGTTTGC
AAGAAGATTGCGTACTGCATCAGTAATTCTTGCCACTCTCTTGGAAATCCAGCTCACAGAGGCTCAT
GTACCACTACAAGATTATGAGGCCAGCAATAGTGTGACACCCAAAATGGTTGATTGGATGCAGGGCGT
TACCAGAAGCTAAGGGTTGGGAGTTCAGGATTCTCTCATTTCACCTTCTAATGAGGAACAAAGATCA
AACACACCATTGGTACTACCCATCTAGGGCAAAAATTTCTGGCCAAAACAGCAGCGTTTCGGGGAAGA
GGAATTACCCGCTTACTAGAGAGCATTTCCAATTCTCCAGCAATATCCACAAAATCTCCAACCTGTGAC
ACTTCACTCTCACCTTACATGTCCCAAAAAGATGGATACAAATCTTTCTTCTTCTTATCTTAA
ACGCGTACGCGGCCGCTCGAGCAGAAACTCATCTCAGAAGAGGATCTGGCAGCAATGATATCCTGGAT
TACAAGGATGACGACGATAAGGTTTAAACGGCCGGC
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Restriction Sites:	Sgfl-Mlul
Plasmid Map:	
ACCN:	NM_032858
Insert Size:	1305 bp
OTI Disclaimer:	Our molecular clone sequence data has been matched to the reference identifier above as a point of reference. Note that the complete sequence of our molecular clones may differ from the sequence published for this corresponding reference, e.g., by representing an alternative RNA splicing form or single nucleotide polymorphism (SNP).
OTI Annotation:	This TrueClone is provided through our Custom Cloning Process that includes sub-cloning into OriGene's pCMV6 vector and full sequencing to provide a non-variant match to the expected reference without frameshifts, and is delivered as lyophilized plasmid DNA.
Components:	The ORF clone is ion-exchange column purified and shipped in a 2D barcoded Matrix tube containing 10ug of transfection-ready, dried plasmid DNA (reconstitute with 100 ul of water).
Reconstitution Method:	<ol style="list-style-type: none"> 1. Centrifuge at 5,000xg for 5min. 2. Carefully open the tube and add 100ul of sterile water to dissolve the DNA. 3. Close the tube and incubate for 10 minutes at room temperature. 4. Briefly vortex the tube and then do a quick spin (less than 5000xg) to concentrate the liquid at the bottom. 5. Store the suspended plasmid at -20°C. The DNA is stable for at least one year from date of shipping when stored at -20°C.
RefSeq:	<u>NM_032858.2</u>
RefSeq Size:	1942 bp
RefSeq ORF:	1305 bp
Locus ID:	84944
UniProt ID:	<u>Q96JY0</u>
Cytogenetics:	1q24.1
MW:	49.2 kDa

Gene Summary:

Plays a central role during spermatogenesis by repressing transposable elements and preventing their mobilization, which is essential for the germline integrity. Acts via the piRNA metabolic process, which mediates the repression of transposable elements during meiosis by forming complexes composed of piRNAs and Piwi proteins and governs the methylation and subsequent repression of transposons. Its association with piP-bodies suggests a participation in the secondary piRNAs metabolic process. Required for the localization of germ-cell factors to the meiotic nuage (By similarity).[UniProtKB/Swiss-Prot Function]

Transcript Variant: This variant (1) represents the longest transcript and encodes the longest isoform (1). Sequence Note: This RefSeq record was created from transcript and genomic sequence data to make the sequence consistent with the reference genome assembly. The genomic coordinates used for the transcript record were based on transcript alignments.