

Product datasheet for MC227116

Tmem173 (NM_001289592) Mouse Untagged Clone

Product data:

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| Product Type: | Expression Plasmids |
| Product Name: | Tmem173 (NM_001289592) Mouse Untagged Clone |
| Tag: | Tag Free |
| Symbol: | Tmem173 |
| Synonyms: | 2610307O08Rik; ERIS; Mita; MPYS; STING |
| Mammalian Cell Selection: | Neomycin |
| Vector: | pCMV6-Entry (PS100001) |
| E. coli Selection: | Kanamycin (25 ug/mL) |
| Fully Sequenced ORF: | >MC227116 representing NM_001289592 Red=Cloning site Blue=ORF Orange=Stop codon |

TTTTGTAATACGACTCACTATAGGGCGGCCGGAATTCGTCGACTGGATCCGGTACCGAGGAGATCTGCC
GCC**CGATCGC**C

ATGCCATACTCCAACCTGCATCCAGCCATCCCACGGCCAGAGGTCACCGCTCCAAATATGTAGCCCTCA
TCTTTCTGGTGGCCAGCCTGATGATCCTTTGGGTGGCAAAGGATCCACCAATCACACTCTGAAGTACCT
AGCACTTCACCTAGCCTCGCACGAACCTGGACTACTGTTGAAAAACCTCTGCTGTCTGGCTGAAGAGCTG
TGCCATGTCCAGTCCAGTTGGATGTTTGGCCTTCTGGTCTCTATAAGTCCCTAAGCATGCTCCTGGGCC
TTCAGAGCTTGACTCCAGCGGAAGTCTCTGCAGTCTGTGAAGAAAAGAAGTTAAATGTTGCCACGGGCT
GGCCTGGTCATACTACATTGGGTACTTGCAGTTGATCTTACCAGGGCTCCAGGCCCGGATCCGAATGTTT
AATCAGCTACATAACAACATGCTCAGTGGTGCAGGGAGCCGAAGACTGTACATCCTCTTTCCATTGGACT
GTGGGGTGCCTGACAACCTGAGTGTAGTTGACCCCAACATTTCGATTCCGAGATATGCTGCCCCAGCAAAA
CATCGACCGTGTCTGGCATCAAGAATCGGGTTTATTCCAACAGCGTCTACGAGATTCTGGAGAACGGACAG
CCAGCAGGCGTCTGTATCCTGGAGTACGCCACCCCTTGACAGCCCTGTTTGCCATGTACAGGATGCCA
AAGCTGGCTTCAGTCGGGAGGATCGGCTTGAGCAGGCTAACTCTTCTGCCGGACACTTGAGGAAATCCT
GGAAGATGTCCCCGAGTCTCGAAATAACTGCCGCTCATTGTCTACCAAGAACCACAGACGGAACAGT
TTCTCACTGTCTCAGGAGGTGCTCCGGCACATTCTGCAGGAAGAAAAGGAGGAGTTACCATGAATGCC
CCATGACCTCAGTGGCACCTCTCCCTCCGTAAGTGTCCCAAGACCAAGACTCCTCATCAGTGGTATGGA
TCAGCCTCTCCCACTCCGCACTGACCTCATC**TGA**

ACGCGTACGCGGCCGCTCGAGCAGAAACTCATCTCAGAAGAGGATCTGGCAGCAAATGATATCCTGGATT
ACAAGGATGACGACGATAAGGTTTAA

Chromatograms: https://cdn.origene.com/chromatograms/ja3236_h04.zip


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| Restriction Sites: | Sgfl-Mlul |
| ACCN: | NM_001289592 |
| Insert Size: | 1014 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: | Clone contains native stop codon, and expresses the complete ORF without any c-terminal tag. |
| 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. |
| Note: | Plasmids are not sterile. For experiments where strict sterility is required, filtration with 0.22um filter is required. |
| RefSeq: | <u>NM_001289592.1, NP_001276521.1</u> |
| RefSeq Size: | 2179 bp |
| RefSeq ORF: | 1014 bp |
| Locus ID: | 72512 |
| UniProt ID: | <u>Q3TBT3</u> |
| Cytogenetics: | 18 |

Gene Summary:

Facilitator of innate immune signaling that acts as a sensor of cytosolic DNA from bacteria and viruses and promotes the production of type I interferon (IFN-alpha and IFN-beta) (PubMed:18818105, PubMed:19433799, PubMed:19776740, PubMed:26229117, PubMed:26669264). Innate immune response is triggered in response to non-CpG double-stranded DNA from viruses and bacteria delivered to the cytoplasm (PubMed:18818105, PubMed:19433799, PubMed:19776740, PubMed:26229117, PubMed:26669264). Acts by binding cyclic dinucleotides: recognizes and binds cyclic di-GMP (c-di-GMP), a second messenger produced by bacteria, and cyclic GMP-AMP (cGAMP), a messenger produced by CGAS in response to DNA virus in the cytosol (PubMed:21947006, PubMed:23722158, PubMed:23258412, PubMed:23519410, PubMed:23910378). Upon binding of c-di-GMP or cGAMP, TMEM173/STING oligomerizes, translocates from the endoplasmic reticulum and is phosphorylated by TBK1 on the pLxIS motif, leading to recruitment and subsequent activation of the transcription factor IRF3 to induce expression of type I interferon and exert a potent anti-viral state (PubMed:25636800). In addition to promote the production of type I interferons, plays a direct role in autophagy (PubMed:30568238). Following cGAMP-binding, TMEM173/STING buds from the endoplasmic reticulum into COPII vesicles, which then form the endoplasmic reticulum-Golgi intermediate compartment (ERGIC) (By similarity). The ERGIC serves as the membrane source for WIPI2 recruitment and LC3 lipidation, leading to formation of autophagosomes that target cytosolic DNA or DNA viruses for degradation by the lysosome (By similarity). The autophagy- and interferon-inducing activities can be uncoupled and autophagy induction is independent of TBK1 phosphorylation (By similarity). Autophagy is also triggered upon infection by bacteria: following c-di-GMP-binding, which is produced by live Gram-positive bacteria, promotes reticulophagy (PubMed:29056340). Exhibits 2',3' phosphodiester linkage-specific ligand recognition: can bind both 2'-3' linked cGAMP (2'-3'-cGAMP) and 3'-3' linked cGAMP but is preferentially activated by 2'-3' linked cGAMP (PubMed:26300263). The preference for 2'-3'-cGAMP, compared to other linkage isomers is probably due to the ligand itself, whichs adopts an organized free-ligand conformation that resembles the TMEM173/STING-bound conformation and pays low energy costs in changing into the active conformation (By similarity). May be involved in translocon function, the translocon possibly being able to influence the induction of type I interferons (By similarity). May be involved in transduction of apoptotic signals via its association with the major histocompatibility complex class II (MHC-II) (PubMed:18559423).[UniProtKB/Swiss-Prot Function]

Transcript Variant: This variant (3) uses an alternate in-frame splice site in the 5' coding region, compared to variant 1. The encoded isoform (3) is shorter than isoform 1.