

## Product datasheet for SC111322

### TMLHE (NM\_018196) Human Untagged Clone

#### Product data:

**Product Type:** Expression Plasmids  
**Product Name:** TMLHE (NM\_018196) Human Untagged Clone  
**Tag:** Tag Free  
**Symbol:** TMLHE  
**Synonyms:** AUTSX6; BBOX2; TMLD; TMLH; TMLHED; XAP130  
**Mammalian Cell Selection:** None  
**Vector:** pCMV6-XL5  
**E. coli Selection:** Ampicillin (100 ug/mL)

**Fully Sequenced ORF:** >OriGene ORF sequence for NM\_018196 edited  
ATGTGGTACCACAGATTGTCCCACCTACACAGCAGGCTTCAGGACTTGCTGAAGGGAGGA  
GTCATATATCCGGCCCTCCACAGCCCAACTTCAAAGCTTACTTCCTTTAGCTGTCCAT  
TGGCACCATACAGCCTCCAAGTCTCTGACTTGTGCTTGGCAGCAACATGAAGATCATTTT  
GAGCTGAAATATGCTAATACCGTGATGCGCTTTGATTACGTCTGGCTTCGAGACCACTGC  
CGCTCAGCATCGTGCTACAACCTAAGACTCACCAGCGCAGCCTGGATACTGCCAGTGTG  
GATTTATGTATCAAGCCAAAGACCATTCTGCTGGATGAGACCACACTCTTTTCACTTGG  
CCAGATGGTCATGTGACTAAATATGATTTGAATTGGCTGGTAAAAACAGCTATGAAGGG  
CAGAAAACAAAAGTCATCCAGCCTAGAATACTATGGAATGCTGAAATCTACCAGCAAGCC  
CAAGTTCATCGGTAGATTGCCAGAGCTTCTTAGAAAACCAACGAGGGACTGAAGAAGTTT  
CTGCAAACTTTCTGCTCTATGGAATTGCATTCTGATAGAAAATGTCCCTCCCACTCAAGAG  
CACACAGAGAAGTTGGCAGAAAGGATCAGCTTAATCAGAGAAACATTTATGGGAGGATG  
TGGTATTTCACTTCAGACTTCTCCAGAGGTGACACTGCGTACACCAAGCTAGCTCTGGAT  
CGGCACACTGACACTACCTATTTTCAAGAGCCCTGTGGCATTCAAGTGTTTCATTGTCTT  
AAACATGAAGGAACTGGTGGCAGGACACTGCTAGTAGATGGATTCTATGCAGCAGAACAG  
GTACTTCAAAGGCACCTGAGGAATTTGAACTCCTCAGTAAAGTGCCATTGAAGCATGAA  
TATATTGAAGATGTTGGAGAATGCACAACCACATGATTGGGATTGGGCCAGTCTTAAAT  
ATCTACCATGGAATAAAGAGCTGTATTTGATCAGGTACAACAACATATGACCGGGCTGTC  
ATCAATACCGTTCCTTATGATGTCGTCATCGCTGGTATACAGCACACCGGACTTAACG  
ATAGAGTTGAGGAGACCTGAGAATGAGTTTTGGGTCAAACATAAAGCCTGGCAGGGTCCTA  
TTTATAGACAACCTGGCGTGCCTACATGGCAGGGAATGCTTCACTGGCTACCGCCAACCTG  
TGTGGCTGCTATTTAACAAGAGATGATGTATTAACACTGCTCGCCTCTTGGGGCTTCAG  
GCTTAA



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**5' Read Nucleotide Sequence:**

>OriGene 5' read for NM\_018196 unedited  
 AATACGACTCACTATAGGGCGGCCGGAATTCGGCACGAGCGGAGCGGGCTGGGGAGGG  
 GAGCGTGGGGCCGACAGTTTTGGGGGTGAAAAGGCAAAAGCGGGTAAAAGGCTGCCTCC  
 CGAGACTCTCCTTGCTTGGAAATCTGCCACTCTGCGGAGTTAGCAGTCACGACCTCCAG  
 CACAGGATGTGGTACCACAGATTGTCCCACCTACACAGCAGGCTTCAGGACTTGTGAAG  
 GGAGGAGTCATATATCCGGCCCTCCACAGCCAACTTCAAAGCTTACTTCCTTAGCT  
 GTCCATTGGCACCATACAGCCTCCAAGTCTCTGACTTGTGCTTGGCAGCAACATGAAGAT  
 CATTTTGGAGCTGAAATATGCTAATACCGTGATGCGCTTTGATTACGTCTGGCTTCGAGAC  
 CACTGCCGCTCAGCATCGTGCTACAACCTAAGACTCACAGCGCAGCCTGGATACTGCC  
 AGTGTGGATTTATGTATCAAGCCAAAGACCATTCTGCTGGATGAGACCACACTCTTTTTTC  
 ACTTGGCCAGATGGTCATGTGACTAAATATGATTTGAATTGGCTGGTAAAAACAGCTAT  
 GAAGGGCAGAAAACAAAAGTCATCCAGCCTAGAATACTATGGAATGCTGAAATCTACCAG  
 CAAGCCCAAGTTCATCGGTAGATTGCCAGAGCTTCTTAAAAACAACGAGGGACTGAAGA  
 AATTTCTGCAAACTTTCTGCTCTATGGAANTGCATTCTGAGAANATGTCCCTCCCCTC  
 AGACACACAGAGAAGNTGGGCAGAAGGATCAGCTTATCAGAGAACCATTTATGGGGAGA  
 TGTGGTATTTCACTTCAGACTCTNCAGAGTGACCTGCGTACACCAGCTAGCTCTGATCGG  
 CCACTGACCTACTATTTAGAGCCTGGGCATCAGTGTTTCATGGCTAAAA

**3' Read Nucleotide Sequence:**

>OriGene 3' read for NM\_018196 unedited  
 GCTATGGACCGCGGCCCGCTGCTAAATACGGTTAATCCCTTTTTTTTTTTTGGACTCATT  
 AAATTTTGGCAATTAGCATGTATAGTATGCCTATGAGAGAGAGAGAGAAAATTAGCAGA  
 AATGGAGGAAAACATATTACATCCTAAATACCACCAAATTGCCAAAATACACATTTGGGT  
 AGTAATCTCCAGAGAACTCTCTGGAGGCACAGCATCTATCATGCTGATTTCCCTGAACAA  
 TATTACAGAATAGCCCTGTGTACTTTCTTTGGGGGGGGAGGTGTAAAGAATTATCATA  
 TGGAACTCACAAGTCTCTAAAACCTTTCTGCAATTATCCCTCATAGGATTACCACACTG  
 ATGAGTTTGTTCACAGGCGCTGTTGTGACTTCCTAATTTAACTCCTATAGAGATGATTT  
 GTGTGGAAGATAAAAACACTATATTTGAACTCAATTTCCACCTTATCCCTGAGCTTATGCT  
 ACCAAGCAATTTTACAATGTTTGGAGTTCTATTAGACTACAGAGAAGAACTAGGGTTT  
 CAGCTAAACCTTTAATCCACAGGCCTAGGTAAGGACTGTCTCAAGGTAGTCGCTAGCC  
 CATGGGTTCTCAAAGTATGGTCTGAAGACCCCTGGAGGAGTCCCTGAGACCCTTTTCGTG  
 GGTCTGCAGATTAATAATTATTTTGTATAATAACACCAAGATGAGTTACATTCAATTAGG  
 CATTTTTCTTTTGAATTAAGCAGACTTTTGGAACTAAGATGTGCATAGCCCTGTAGTA  
 AGCTAAGTGGAAAGTTACCAAGGCAGTAGTAGCCTTTTTTCCCCCTCTCGNTATCTCATG  
 AAGATACAGAGGACTTTTCCAGAGCTACATGACATATGGTTATGGCATCTCTCATAGC  
 TGACGGATGTGTGCTTCCGCGCTCCCTTCTGAGCTCTTAAAATTTCTCAGTCCTTATACC  
 CTAATAATTAGTACTCCAAGATAGGACCCACN

**Restriction Sites:**

NotI-NotI

**ACCN:**

NM\_018196

**Insert Size:**

2880 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).

**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:**

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:** [NM\\_018196.1](#), [NP\\_060666.1](#)

**RefSeq Size:** 2836 bp

**RefSeq ORF:** 1266 bp

**Locus ID:** 55217

**UniProt ID:** [Q9NVH6](#)

**Cytogenetics:** Xq28

**Domains:** Gamma-BBH

**Protein Pathways:** Lysine degradation

**Gene Summary:** This gene encodes the protein trimethyllysine dioxygenase which is the first enzyme in the carnitine biosynthesis pathway. Carnitine play an essential role in the transport of activated fatty acids across the inner mitochondrial membrane. The encoded protein converts trimethyllysine into hydroxytrimethyllysine. A pseudogene of this gene is found on chromosome X. Alternate splicing results in multiple transcript variants.[provided by RefSeq, May 2010]

Transcript Variant: This variant (1) represents the longer transcript and encodes the longer 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.