

Product datasheet for **SC117349**

EML1 (NM_004434) Human Untagged Clone

Product data:

Product Type:	Expression Plasmids
Product Name:	EML1 (NM_004434) Human Untagged Clone
Tag:	Tag Free
Symbol:	EML1
Synonyms:	BH; ELP79; EMAP; EMAP-1; EMAPL
Mammalian Cell Selection:	None
Vector:	<u>pCMV6-XL4</u>
E. coli Selection:	Ampicillin (100 ug/mL)



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Fully Sequenced ORF: >NCBI ORF sequence for NM_004434, the custom clone sequence may differ by one or more nucleotides

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ATGGAGGACGGCTTCTCCAGCTACAGCAGCCTGTACGACACGTCCTCGTCTCCAGTTCTGCAACGATG
ACAGCGCTTCTGCTGCAAGTAGCATGGAGGTGACAGACCGCATTGCTTCACTGGAGCAGAGAGTCCAGAT
GCAAGAAGACGACATCCAGCTGCTCAAATCAGCTCTAGCTGATGTGGTTCCGGCGGCTGAACATTACTGAG
GAACAGCAGGCCGTGCTTAACAGGAAAGGACCTACCAAAGCAAGACCACTGATGCAGACCCCTGCCTTTAA
GAACCACGGTCAACAATGGCACTGTGTTACCAAAGAAACCTACTGGCTCTCTACCATCCCCCTCCGGGGT
CAGGAAAGAAACTGCTGTGCCAGCAACCAAAAGTAAATCAAGAGGACCAGCTCTTCTGAACGAGTGTCT
CCTGGGGTGCAGAGGAAAGCAATGGGGATTCCAGAGGAAACCGAATCGCACAGGCTCCACCAGCAGCT
CTTCCAGTGGCAAAAAGAACAGTAAAGCAAACCAAGGAGCCTGTATTCACTGCAGAAGAAGGCTATGT
AAAAATGTTTCTCGTGGACGCCCTGTTACCATGTACATGCCCAAAGATCAAGTGGATTCTTACAGCTTG
GAAGCAAAAGTAGAACTCCAACCAAGAGACTCAAGCTGGAATGGGTCTATGGGTACAGGGTTCGAGACT
GCCGTAACAACCTGTACTTGGCTCCGACGGGAGAGACCGTCTACTTCATCGCATCCGTGGTGGTGTATA
CAACGTGGAGGAGCAACTGCAGAGGCATTACGCTGGCCACAACGATGACGTGAAGTGCCTAGCAGTTCAT
CCTGATCGGATCAGATAGCAACAGGACAAGTTGCGGGCACATCGAAGGATGAAAAACAATTGCCCCAC
ATGTGCGCATCTGGGATTCTGTGACATTGAATACTCTCCACGTCATTGGAATAGGTTTTTTGACCGAGC
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CATTACTCAGGGTCACACTGATGAGCTCTGGGACTGGCCATCCATGCCTCAAAATCTCAGTTCTTGACC
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AGGATCCAGCTCAGTCTTCTGGTTTTTCATCCTCAGGGTCTGTGGTTGCAGTCGGAACACTCACTGGGAG
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GCGTTAGTGACAACGGGAGGAAGTACACGCGAGTGGGCAAGTGTCTCGGGTCACTCCAGCTTCATTACTCA
CCTGGACTGGTCTGTAAACTCACAGTTCCTCGTGTCAAATTCGGGAGACTACGAAATCCTCTACTGGGT
CCCTCTGCCTGTAAAGCAAGTCTAAGTGTGAAACTACAAGAGACATTGAATGGGCTACCTATACCTGCA
CTTTGGGATTCATGTTTTGGAGTGTGGCCAGAAAGGCTCGGACGGAACCGACATCAATGCCGTCTGTGCG
GGCCATGAGAAGAACTCCTGTCAACAGGCGACGACTTTGGCAAAGTGCACCTCTTCTCATACCCCTGC
TCGCAGTTCAGGGCTCCAAGCCACATCTACGGCGGGCACAGCAGCCATGTCACCAATGTCGATTTCTCT
GTGAAGACAGCCACCTCATCTCCACGGGGGAAAGACACAAGCATCATGCAGTGGCGGTCATTTAG
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5' Read Nucleotide Sequence: >OriGene 5' read for NM_004434 unedited
 CCGCCGCGAATTCGGCAGCAGCTGGGTGGGTGACAGCCGCTGAGGCTCTGAGTGACCCTG
 CAGGCTGGTGACCAGCCGCGCCCATCTTCCACACCATGTGGACGGCGAGGGCCCTC
 CGCCGATGACAGCGCTTCTGCTGCAAGTAGCATGGAGGTGACAGACCGCATTGCTTCACT
 GGAGCAGAGAGTCCAGATGCAAGAAGACGACATCCAGCTGCTCAAATCAGCTCTAGCTGA
 TGTGTTCCGGCGGCTGAACATTACTGAGAACAGCAGGCCGTGCTTAACAGGAAAGGACC
 TACCAAAGCAAGACCACCTGATGCAGACCCTGCCTTTAAGAACCAGGTCAACAATGGCAC
 TGTGTTACCAAAGAAACCTACTGGCTCTCTACCATCCCCCTCCGGGTGTCAGGAAAGAAAC
 TGCTGTGCCAGCAACCAAAGTAAACATCAAGAGGACCAGCTCTTCTGAACGAGTGTCTCC
 TGGGGGTGCAAGGAAAGCAATGGGGATTCCAGAGGAAACCGGAATCGCACAGGCTCCAC
 CAGCAGCTCTTCCAGTGGCANAAAGAACAGTGAAAGCAAACCCAAGGAGCCTGTATTTCAG
 TGCAGAAGAAGGCTATGTAATAATGTTTCTTCTGAGCAGCCCTGTACCATGTACATGCC
 CAAAGATCAAGTGGATTCTTACAGCTTGGAAAGCAAAGTAGAACTCCACCAAGAGACTCN
 AGCTGGAATG

3' Read Nucleotide Sequence: >OriGene 3' read for NM_004434 unedited
 TCCGGCTCCCCTCCCCCTTCCACCCAAACCTTGGTGCCTTTAACTCTTTTTTAAAA
 ATTGAGATCTTCCCATGTAACATAAAATCAATCTGTCACACATCGAAATGGATAAGGCC
 TTCTTGACAAATTTCTGCCACCTCCGTTTAAACGCATCAGAATCAATCTTATCTCATACT
 GTAATGGGTGCGCCATGCACAAAACACAGGAAAAGCAGCTGCCCTTGGTACCTGGTGGAG
 CCTTTCGAGACTTCTCTGTCAGGTTTATGGTGCCTCGAGAGAGATCTTCTGACTTGG
 TCTACTCGGTACATACATAAAATCTCCCCTGTTTCATCAGCCACCACTTCTAAGCCCTAC
 CCTTACATACCTATTTGGAATCCCTTCAATCCTCTCCTATTTCCACCTCCTCCTGCC
 CTTCTTCTTTTACAATTATGCGTTTACCCTCCTCCCCACATCACTCCCCCTTCCC
 AACTAGTTTCCCTATCTTATCCCCCAATCTCACCATCTCATCCTTACTATCCCCCTT
 TTTCTTACACCTTCTTCCCTTCCATCTCCCTCCCGATCGTTCCCTCCCTCTTCTCC
 CATAACCTCCCCCTCCCATACACACTTACATCCCACCTCCATTCCCACCTTTCCCTA
 CCATTTCTATTCTACCCTCTAACATAAATATATATATTTCTCCACCCCTCTCCACTTGG
 TCCACCCCATCATTTCCTTATCTCTACATATCTTACCATTCTATCCACCCCTCCCCC
 CCCCCACACCTCTCCCTTATCTCTATCCACCTCCCACTTCCCATTCTCAAATCACC
 TCCCCTACTACTACATGCACCCACCCCTCACATTTACATCATCTCCACCCCGCAAA
 CCTATCTTTTTATCCCTTCTGCTACCTACATCTCCCTTCTCCTTCTCCCTTCTACTACCT
 CTACNCCTCTCCTCCCCCACCCCTTCTCCCAACCTTCTCTCTCACCACATCTCTCAC
 TCTTCTTTTCTCCTCGCACTACTATTCTCTCTCTCCTCTCTCCTCTCTCCTCTCTAT
 CCACCTTCACTCACATCTCCACCAACACACCACCG

Restriction Sites: NotI-NotI

ACCN: NM_004434

Insert Size: 3890 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:	<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:	NM_004434.1 , NP_004425.1
RefSeq Size:	4479 bp
RefSeq ORF:	4479 bp
Locus ID:	2009
UniProt ID:	O00423
Cytogenetics:	14q32.2
Domains:	WD40, HELP
Gene Summary:	<p>Human echinoderm microtubule-associated protein-like is a strong candidate for the Usher syndrome type 1A gene. Usher syndromes (USHs) are a group of genetic disorders consisting of congenital deafness, retinitis pigmentosa, and vestibular dysfunction of variable onset and severity depending on the genetic type. The disease process in USHs involves the entire brain and is not limited to the posterior fossa or auditory and visual systems. The USHs are categorized as type I (USH1A, USH1B, USH1C, USH1D, USH1E and USH1F), type II (USH2A and USH2B) and type III (USH3). The type I is the most severe form. Gene loci responsible for these three types are all mapped. Two transcript variants encoding different isoforms have been found for this gene. [provided by RefSeq, Jul 2008]</p> <p>Transcript Variant: This variant (2) lacks an alternate in-frame exon compared to variant 1. The resulting isoform (b) has the same N- and C-termini but is shorter compared to isoform a.</p>