

## Product datasheet for **SC337915**

### CHL1 (NM\_001253388) Human Untagged Clone

#### Product data:

Product Type:	Expression Plasmids
Product Name:	CHL1 (NM_001253388) Human Untagged Clone
Tag:	Tag Free
Symbol:	CHL1
Synonyms:	CALL; L1CAM2
Mammalian Cell Selection:	Neomycin
Vector:	pCMV6-Entry (PS100001)
E. coli Selection:	Kanamycin (25 ug/mL)
Fully Sequenced ORF:	>NCBI ORF sequence for NM_001253388, the custom clone sequence may differ by one or more nucleotides

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ATGGAGCCGCTTTTACTTGAAGAGGACTAATCGTATATCTAATGTTCCCTCCTGTTAAAATTCTCAAAG
CAATTGAAATACCATCTTCAGTTC AACAGGTTCCAACAATCATAAAACAGTCAAAAGTCCAAGTTCCTT
TCCCTTCGATGAGTATTTTCAAATTGAATGTGAAGCTAAAGGAAATCCAGAACCAACATTTTCGTGGACT
AAGGATGGCAACCCTTTTATTTCACTGACCATCGGATAATTCCATCGAACAAATTCAGGAACATTCAGGA
TCCCAAACGAGGGGCACATATCTCACTTTCAAGGGAAATACCGCTGCTTTGCTTCAAATAAACTGGGAAT
CGCTATGTCAGAAGAAATAGAATTTATAGTTC AAGTGTCCAAAATCCCAAAGAAAAAATTGACCCCT
CTTGAAGTGGAGGAGGAGATCCAATTGCTCCCATGCAATCCTCCCAAAGGCCTCCCACCTTTACACA
TTTATTGGATGAATATTGAATTAGAACACATCGAACAAAGATGAAAGAGTATACATGAGCCAAAAGGGAGA
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GAAGGCTTGCCAACTCCACAGGTTGATTGGAACAAAATGGTGGTGACTTACCAAAGGGGAGAGAAAACAA
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CTTCCCCAGGAAATCAGTTTACCAACCTTCAACCAAATCATACTGCTGTGTACCAGTGTGAAGCCTCA
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TAAGAATCCTCGTATCCCCAAATTGCATATGCTTGAATTACATTGTGAAAGCAAATGTGACTCACATTTG  
 AACACAGTTTGAAGTTGTCTGGAGTAAAGATGGAGAAGCCTTTGAAATTAATGGCACAGAAGATGGCA  
 GGATAATTATTGATGGAGCTAATTTGACCATATCTAATGTAACCTTAGAGGACCAAGGATTTACTGCTG  
 TTCAGCTCATACTGCTCTAGACAGTGTCCGATATAACTCAAGTAACTGTTCTTGATGTTCCGGATCCA  
 CCAGAAAACCTTCACTTGTCTGAAAGACAGAACAGGAGTGTTCGGCTGACCTGGGAAGCTGGAGCTGACC  
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 TTAATGATATTAACATTACAACCCATCAAAGCCAGCTGGCACCTCTCAAACCTGAATGCAACTACCA  
 AGTACAAATCTACTTGAGGGCTGCACCTCACAGGGCTGTGGAAAACCGATCACGGAGGAAAGCTCCAC  
 CTTAGGAGAAGGAAATATGCTGGTTTATATGATGACATCTCCACTCAAGGCTGGTTATTGGACTGATG  
 GTGCGATTGCTCTTCTCACACTACTATTATTAACCTGTTTGTGTTGTAAGAGGAATAGAGGTGGAAGT  
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 TGAATACAGTGACAGTATGAAAAGCCTCTCAAAGGAAGCCTTCGGTCCCTTAATAGGGATATGCAGCCT  
 ACTGAAAGTGCTGACAGCTTAGTCGAATACGGAGAGGGAGACCATGGTCTCTTCAAGTGAAGATGGATCAT  
 TTATTGGTGCCTACGCTGGATCTAAGGAGAAGGGATCTGTTGAAAGCAATGGAAGTTCTACAGCAACTTT  
 TCCCTTCGGGCA<sup>TAA</sup>

- Restriction Sites:** SgfI-MluI
- ACCN:** NM\_001253388
- 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\\_001253388.1](#), [NP\\_001240317.1](#)
- RefSeq Size:** 7324 bp

RefSeq ORF: 3516 bp

Locus ID: 10752

Cytogenetics: 3p26.3

Protein Families: Transmembrane

**Gene Summary:** The protein encoded by this gene is a member of the L1 gene family of neural cell adhesion molecules. It is a neural recognition molecule that may be involved in signal transduction pathways. The deletion of one copy of this gene may be responsible for mental defects in patients with 3p- syndrome. This protein may also play a role in the growth of certain cancers. Alternate splicing results in both coding and non-coding variants. [provided by RefSeq, Nov 2011]

Transcript Variant: This variant (3) lacks an in-frame exon in the coding region, compared to variant 1. This variant encodes isoform 3, which is shorter than 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.