

Product datasheet for **SC325107**

CHRD1 (NM_001143982) Human Untagged Clone

Product data:

Product Type:	Expression Plasmids
Product Name:	CHRD1 (NM_001143982) Human Untagged Clone
Tag:	Tag Free
Symbol:	CHRD1
Synonyms:	CHL; dA141H5.1; MGC1; MGCN; NRLN1; VOPT
Vector:	<u>pCMV6 series</u>
Fully Sequenced ORF:	>NCBI ORF sequence for NM_001143982, the custom clone sequence may differ by one or more nucleotides ATGAGAAAAAGTGGAAATGGGAGGCATGAAATACATCTTTTCGTTGTTGTTCTTTCTT TTGCTAGAAAGGAGGCAAAACAGAGCAAGTAAAACATTCAGAGACATATTGCATGTTTCAA GACAAGAAGTACAGAGTGGGTGAGAGATGGCATCCTTACCTGGAACCTTATGGGTGGTT TACTGCGTGAACGCATCTGCTCAGAGAATGGGAATGTGCTTTCAGCCGAGTCAGATGT CCAAATGTTTCATTGCCTTCTCCTGTGCATATTCTCATCTGTGCTGCCCTCGCTGCCCA GACTCCTTACCCCAAGTGAACAATAAGGTGACCAGCAAGTCTTGCAGTACAATGGGACA ACTTACCAACATGGAGAGCTGTTTCGTAGCTGAAGGGCTCTTTCAGAAATCGGCAACCCAAT CAATGCACCCAGTGCAGCTGTTTCGGAGGAAACGTGTATTGTGGTCTCAAGACTTGCCCC AAATTAACCTGTGCCCTCCAGTCTCTGTTCCAGATTCCTGTGCTGCCGGGTATGCAGAGGA GATGGAGAAGTGTATGGGAACATTCTGATGGTGATATCTTCCGGCAACCTGCCAACAGA GAAGCAAGACATTCTTACCACCGCTCTCACTATGATCCTCCACCAAGCCGACAGGCTGGA GGTCTGTCCCCTTTCTGGGGCCAGAAGTACCAGGGGAGCTCTTATGGATTCCCAGCAA GCATCAGGAACCATTTGTCAAATTTGTCATCAATAACAAACACAAGCATGGACAAGTGTGT GTTTCCAATGGAAAGACCTATTCTCATGGCGAGTCTTGGCACCAAGCAAGAGTGAAGAAAATC CACTGCCCAATCGATACCCCTGCAAGTATCCTCAAAAAATAGACGGAAAATGCTGCAAG GTGTGTCCAGGTAAAAAGCAAAAGAAGAACTTCCAGGCCAAAGCTTTGACAATAAAGGC TACTTCTGCGGGGAAGAAACGATGCCTGTGTATGAGTCTGTATTTCATGGAGGATGGGGAG ACAACCAGAAAAATAGCACTGGAGACTGAGAGACCACCTCAGGTAGAGGTCCACGTTTGG ACTATTCGAAAGGGCATTCTCCAGCACTCCATATTGAGAAGATCTCCAAGAGGATGTTT GAGGAGCTTCTCACTTCAAGCTGGTGACCAGAACAACCCTGAGCCAGTGAAGATCTTC ACCGAAGGAGAAGCTCAGATCAGCCAGATGTGTTCAAGTCGTGTATGCAGAACAGAGCTT GAAGATTTAGTCAAGTTTTGTACCTGGAGAGATCTGAAAAGGGCCACTGT
Restriction Sites:	Please inquire
ACCN:	NM_001143982



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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_001143982.1</u> , <u>NP_001137454.1</u>
RefSeq Size:	4098 bp
RefSeq ORF:	1374 bp
Locus ID:	91851
UniProt ID:	<u>Q9BU40</u>
Cytogenetics:	Xq23
Protein Families:	ES Cell Differentiation/IPS, Secreted Protein
Gene Summary:	<p>This gene encodes an antagonist of bone morphogenetic protein 4. The encoded protein may play a role in topographic retinotectal projection and in the regulation of retinal angiogenesis in response to hypoxia. Alternatively spliced transcript variants encoding different isoforms have been described. [provided by RefSeq, Jan 2009]</p> <p>Transcript Variant: This variant (2) uses an alternate in-frame splice site in the 5' coding region compared to variant 1. The resulting protein (isoform 2) is shorter but has the identical N- and C-termini compared to 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.</p>