

Product datasheet for **SC332818**

ZNF211 (NM_001265599) Human Untagged Clone

Product data:

Product Type: Expression Plasmids
Product Name: ZNF211 (NM_001265599) Human Untagged Clone
Tag: Tag Free
Symbol: ZNF211
Synonyms: C2H2-25; CH2H2-25; ZNF-25
Vector: pCMV6-Entry (PS100001)
Fully Sequenced ORF: >SC332818 representing NM_001265599.
Blue=Insert sequence Red=Cloning site Green=Tag(s)

```
ATGCTGGAGAACCTTGCACCTACGTCCTCCCTGGGACTTATATCATCCTGGTCTCATGTAGTTGCTCAA
CTAGGGCTAGGGGAAGTGCCTCTGTTCTTACAGGATGTTTCACTCCAGCCTCAGCAAGATGGGAT
CAGAGAGGGCCTGGCCTACATGAATGGCACTTGGGAAAAGGCATGTCATCAGGTTGTTGGTGTGGAGTG
GAACATGAGGAAACACCTTCTGAACAGAGAATTTCTGGAGAAAGAGTGCCACAGTTCAGGACTTCCAAA
GAAGGTTTCATCTCCAGAAATGCCGACTCCTGTGAAATATGTTGCCTGGTCTTGAGAGATATTTTGCAC
TTGGCTGAACACCAAGGAACAACTGCGGGCAGAACTACACACATGTGAAAACAATTCTACATCAGT
GCAAACTTCAACAGCACCAGAGGCAGCACATTACAGAGGCACCTTTCAGAAATGTTGAGACACTGCC
TCGTTTACACAGAGTTGCATAGTCCATGTGTCCGAGAAACCCCTTACCTGCAGGGAGATCAGGAAAGAC
TTCTGGCCAACATGAGGTTTCTCCATCAAGACGCCACTCAAACAGGGGAGAAGCCAAATAACAGTAAC
AAGTGTGCGGTGGCCTTTTACAGTGGAAAAAGTCATCACAACCTGGGAAAATGCAGTAAAGCCTTTAGC
CACATAGACACACTTGTTCAGGACCAGAGAATCCTCACTAGAGAAGGACTTTTTGAGTGCAGTAAATGT
GGGAAAGCATGTACGCGAAGATGTAACCTCATTACAGCACCAGAAAGTCCACAGTGAAGAAAGGCCTTAT
GAATGCAATGAATGTGGAAAATCTTTACCTACTACTCCAGTTTTCATTATACATCAGAGAGTTCATACT
GGAGAAAGCCTTATGCGTGCCCTGAATGTGGGAAATCGTTTATGTCAGATATACAGCCTCAATAGCCAT
AGGAAAGTTCACACTGGAGAAAGGCCTTATGAATGTGGGAAATGTGGGAAATCTTTAGCCAAAGGTCC
AACCTCATGCAGCATCGCAGAGTTCACACTGGAGAAAGGCCTTATGAATGCAGCGAATGTGGGAAATCT
TTAGCCAAAACCTTTAGCCTGATCTACCACCAGAGAGTTCACACTGGAGAAAGACCTCATGAGTGAAT
GAATGTGGAAAATCCTTTAGCCGAAAGCTCCAGCCTCATTACACCACCGAGACTTCACACTGGAGAAAAG
CCCTATGAGTGCAGTAAATGTGGAAAGTCATTTAAGCAAAGCTCCAGTTCAGTTCACATCGGAAAGTC
CACACAGGGGAAAGGCCTTATGTGTGTGGGAAATGTGGGAAATCCTTTAGCCATAGCTCCAACCTTAAG
AACCACCAGAGAGTTCACACTGGAGAAAGACCTGTTGAGTGCAGTGAATGTAGCAAATCCTTTAGCTGT
AAATCTAACCTCATTAAACACCTGAGAGTTCACACTGGAGAAAGGCCTTATGAGTGCAGTGAATGTGGG
AAATCCTTTAGCCAAAGTCTAGCCTCATTCAACACCGCAGAGTTCACACGGGAAAAGGCCTTATCAG
TGAGTCAATGTGGGAAATCCTTTGGCTGCAAATCTGTCTCATTCAACACCAGAGAGTTCACATTGGA
GAAAAGCCTTAG
```

Restriction Sites: Sgfl-Mlul
ACCN: NM_001265599



[View online »](#)

Insert Size:	1668 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_001265599.1
RefSeq Size:	2478 bp
RefSeq ORF:	1668 bp
Locus ID:	10520
UniProt ID:	Q13398
Cytogenetics:	19q13.43
Protein Families:	Transcription Factors
MW:	63.2 kDa
Gene Summary:	<p>This gene encodes a protein containing a Kruppel-associated box domain and multiple zinc finger domains. This protein may play a role in developmental processes. Alternative splicing results in multiple transcript variants. [provided by RefSeq, Mar 2014]</p> <p>Transcript Variant: This variant (5) differs in the 5' UTR, initiates translation at a downstream, in-frame start codon and lacks an exon in the coding region, but maintains the reading frame, compared to variant 1. The encoded isoform (5) 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.</p>