

Product datasheet for **SC332816**

ZNF211 (NM_001265597) Human Untagged Clone

Product data:

Product Type: Expression Plasmids
Product Name: ZNF211 (NM_001265597) Human Untagged Clone
Tag: Tag Free
Symbol: ZNF211
Synonyms: C2H2-25; CH2H2-25; ZNF-25
Vector: pCMV6-Entry (PS100001)
Fully Sequenced ORF: >SC332816 representing NM_001265597.
 Blue=Insert sequence Red=Cloning site Green=Tag(s)

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ATGCTCGGGTCCCCCGGGTCGCCCGAGCTCCCGGTCCAGCTCCGCCACAGACTCGGATGGCGACC
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Restriction Sites:	Sgfl-Mlul
ACCN:	NM_001265597
Insert Size:	1890 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_001265597.1
RefSeq Size:	2668 bp
RefSeq ORF:	1890 bp
Locus ID:	10520
UniProt ID:	Q13398
Cytogenetics:	19q13.43
Protein Families:	Transcription Factors
MW:	71.7 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 (3) contains an alternate exon in the coding region, but maintains the reading frame, compared to variant 1. The encoded isoform (3) is longer 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>