

Product datasheet for SC315768

ZNF331 (NM 001079906) Human Untagged Clone

Product data:

Product Type: Expression Plasmids

Product Name: ZNF331 (NM_001079906) Human Untagged Clone

Tag: Tag Free **ZNF331** Symbol:

Synonyms: RITA; ZNF361; ZNF463

Vector: pCMV6 series

>NCBI ORF sequence for NM_001079906, the custom clone sequence may differ by one or **Fully Sequenced ORF:**

more nucleotides

ATGGCCCAGGGTTTGGTGACGTTCGCCGACGTAGCCATAGACTTTTCTCAGGAGGAGTGG GCCTGTCTGAACTCTGCTCAGAGGGACCTGTACTGGGACGTGATGCTGGAGAACTACAGT AACTTGGTCTCACTGGATTTGGAGTCAGCATATGAAAATAAGAGTTTACCTACAGAAAAA AACATTCATGAAATAAGGGCTTCCAAAAGGAATTCAGATAGAAGAAGTAAATCCCTTGGC AATCAGATGATCATCAATTATGTCAAAAGACCTGCTACTAGAGAAGGCACCCCTCCTAGA ACACATCAGAGACATCATAAGGAGAATTCCTTTGAATGTAAGGACTGTGGGAAGGCCTTT AGTCGTGGCTATCAACTTAGTCAACATCAGAAAATCCATACTGGTGAGAAACCTTATGAA TGTAAAGAATGTAAGAAGGCCTTCCGTTGGGGCAATCAGCTTACTCAACATCAAAAAATT CATACTGGGGAGAGCCCTACGAATGTAAAGACTGTGGGAAGGCTTTTCGATGGGGCTCA AGCCTCGTTATTCATAAGAGGATTCATACTGGTGAAAAACCCTATGAATGTAAAGACTGT GGAAAGGCCTTTCGGCGTGGTGATGAGCTCACTCAGCACCAGAGATTCCACACTGGGGAG AAAGACTACGAATGCAAAGACTGTGGGAAGACCTTTAGCCGTGTGTATAAACTTATTCAG CACAAGAGAATTCATAGTGGGGAGAAGCCTTACGAGTGTAAAGACTGTGGGAAGGCTTTT ATTTGTGGTTCAAGCCTCATTCAGCATAAAAGAATTCACACAGGTGAGAAACCCTATGAA TGTCAAGAATGTGGGAAGGCCTTTACTCGAGTCAATTACCTTACTCAGCATCAGAAGATC CACACCGGTGAGAAGCCTCACGAATGTAAGGAGTGTGGGAAGGCCTTTCGCTGGGGTTCG AGCCTCGTTAAGCACGAGAGGATACATACGGGCGAGAAGCCGTACAAGTGCACAGAATGT GGGAAGGCCTTCAATTGTGGCTATCACCTCACTCAGCACGAGAGAATCCACACAGGCGAA ACCCCGTATAAATGTAAGGAGTGTGGGAAGGCTTTCATTTATGGATCGAGCCTCGTGAAA CATGAGAGAATTCATACCGGGGTGAAACCCTATGGGTGTACAGAATGTGGGAAGAGCTTT AGTCACGGCCATCAGCTTACACAACATCAGAAAACGCACAGTGGGGCGAAATCCTACGAA TGTAAGGAGTGCGGGAAGGCATGTAACCACCTAAACCATCTCCGAGAACATCAGAGGATC CACAACAGT

Restriction Sites: Please inquire ACCN: NM 001079906



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ZNF331 (NM_001079906) Human Untagged Clone - SC315768

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: 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 001079906.1</u>, <u>NP 001073375.1</u>

19q13.42

 RefSeq Size:
 4151 bp

 RefSeq ORF:
 1392 bp

 Locus ID:
 55422

 UniProt ID:
 Q9NQX6

Cytogenetics:

Protein Families: Transcription Factors

Gene Summary: This gene encodes a zinc finger protein containing a KRAB (Kruppel-associated box) domain

found in transcriptional repressors. This gene may be methylated and silenced in cancer cells. This gene is located within a differentially methylated region (DMR) and shows allelespecific expression in placenta. Alternative splicing and the use of alternative promoters results in multiple transcript variants encoding the same protein. [provided by RefSeq, Nov

2015]

Transcript Variant: This variant (4, also known as e) differs in the 5' UTR compared to variant 1. All variants encode the same protein. 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.