

Product datasheet for **SC328929**

ZNF430 (NM_001172671) Human Untagged Clone

Product data:

Product Type:	Expression Plasmids
Product Name:	ZNF430 (NM_001172671) Human Untagged Clone
Tag:	Tag Free
Symbol:	ZNF430
Mammalian Cell Selection:	None
Vector:	<u>pCMV6-XL5</u>
E. coli Selection:	Ampicillin (100 ug/mL)



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Fully Sequenced ORF:	>NCBI ORF sequence for NM_001172671, the custom clone sequence may differ by one or more nucleotides <pre> ATGGAGAACCTGAAGTCTGGAGTGTATCCTCTCAAGGAAGCAAGTGGATGCCCTGGGGCT GACAGGAATCTTCTGGTGTACTCTTTTATGAAAAGGGGCCATTGACATTTAGGGATGTG GCCATAGAATTTTCTCTGGAGGAGTGGCAATGCCTGGACACTGCTCAGCAGGATTTGTAT AGAAAAGTGATGTTAGAGAACTACAGAAACCTGGTCTTCTTGGGTATTGCTGTTTCTAAG CCAGACCTGATCACCTGTCTAGAGCAAGGAAAAGAGCCCTGGAATATGAAGAGACATGCG ATGGTAGATCAACCCCGAGTTACATATTCTCATTGTCCTGCAAGACCTTTGGCCAGAGCAG GGCATAAAAGATTCTTTCCAAGAAGTCATATTGAGAAGATATGGAAAATGTGGACATGAA GATTTACAGTTAAGAACAGGCTGTAAAAGTGTGGATGAGTGAATCTGCACAAAGAATGT TATGATGAACTAAACAGTGTGTTGACAACTACCCAGAGTGAAATATTTCAATATGATAAA TATGTGAATGTCTTTATAAATTTCAAATCCAAATATACAAAAGATAAGACATACTGGA AAGAAGCCTTTCAAATGTAAAAATGTGACAAATCGTTTTGCATGCTTTTACACCTAACT CAACATAAAAGAATTCATATTAGGGAATTTCTTACCAATGTGAAGAATGTGGTAAAGTC TTTAACTGGTTCTCAACCCTTAGACACAGAAGAATTCATACTGGAGAGAAACCTTAC AAATGTGAACAATGTGGCAAAGCTTTTAAGCAGTCCTCAACCCTTACTACACATAAGATA ATTCATACTGGAGAGAAACCTACAGATGTGAAGAATGTGGCAAACCTTTAACCAGTCC TCACACCTTACTACATAAAAGAATTCATACTGGAGAGAAACCTACAGATGTGAAGAA TGTGGCAGAGCTTTTAACCGGTCTCACACCTTACTACACATAAGATTATTCATACTGGA GAGAAACCTACAAATGTGAAGAATGTGGCAAAGCTTTTAACCAATCCTCAACCCTTACT ACACATAAGATAATTCATGCTGGAGAGAAACCTTACAAATGTGAAGAATGTGGCAAAGCT TTTTACCGATTCTCATACCTTACTAAACATAAGATAATTCATACTGGAGAGAAATTTCTAC AAATGTGAAGAATGTGGCAAAGGCTTTAATTGGTCCTCGACCCTTACTAAACATAAAAGA ATTCATACTGGAGAGAAACCTACAAATGTGAACAATGTGGCAAAGCTTTTAATGAGTCC TCAAACCTTACTGCACATAAGATAATTCATACTGGAGAGAAACCTACAAATGTGAAGAA TGTGGCAAAGCCTTTAACCGGTCCCAAACTTACTGCACATAAGGTAATTCATTCTGGA GAGAAACCTACAAATGTGAAGAATGTGGCAAAGCTTTTAACCAATTCTCAAACCTTACT AAACATAAGATAACTCATATTGGAGATACATCTTACAAATACCTAGAATGTGATAAAGCC TTTAGCCAGTCTTCAACTCTTACTAAACATAAGGTAATTCATACTGGAGAGAAACCTTAC AACTGTGAAGAATACGGCAAAGCTTTCAACCAGTCCTCAACCTTATTGAACAAAGTAAT TCATACTGGAGAGAAACCTACAAATGTGA </pre>
Restriction Sites:	Please inquire
ACCN:	NM_001172671
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_001172671.1, NP_001166142.1</u>
RefSeq Size:	3916 bp
RefSeq ORF:	1710 bp
Locus ID:	80264
Cytogenetics:	19p12
Protein Families:	Transcription Factors
Gene Summary:	May be involved in transcriptional regulation.[UniProtKB/Swiss-Prot Function] Transcript Variant: This variant (2) uses an alternate in-frame splice site in the coding region, compared to variant 1. The resulting isoform (2) lacks one internal amino acid, 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.