

Product datasheet for **SC331877**

ANKRD6 (NM_001242813) Human Untagged Clone

Product data:

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|---------------|--|
| Product Type: | Expression Plasmids |
| Product Name: | ANKRD6 (NM_001242813) Human Untagged Clone |
| Tag: | Tag Free |
| Symbol: | ANKRD6 |
| Vector: | pCMV6-Entry (PS100001) |



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Fully Sequenced ORF: >SC331877 representing NM_001242813.
 Blue=Insert sequence Red=Cloning site Green=Tag(s)

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ATGAGCCAGCAAGATGCGGTGCTGCACCTTCAGAGCGCCTTCTCGTAGCTGCGTACAAAGGCCAAACA
GAGAATGTGGTTCAGCTCATCAACAAGGGCGCCAGGGTAGCGGTTACCAAGCATGGCCGGACTCCCCTG
CATCTTGCTGCCAATAAGGGCCATCTTCTGTGGTCCAGATCTTGCTGAAGGCTGGCTGCGACCTTGAT
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TCCTGGCATGGTTTCAGCCAGTCAGCCAAGCTGCTCATTAAAGCAGGAGCCAACGTGCTTGCCAAGAAC
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GAAAAGTGGTATGAAAGGAAGATTGAAGAAGCAGGAAGCCAAGCCAATCAGAAAGCCAGCAAGATAAG
GCTACATTGAAGGAACACATTA AAAAGTTTGAAGAGGAACCTGCCAAACTAAGGACTAGGGTGCAGAAAG
GAAAATTAG
  
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Restriction Sites: SgfI-MluI

ACCN: NM_001242813

Insert Size: 2079 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:

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_001242813.1](#)

RefSeq Size: 5250 bp

RefSeq ORF: 2079 bp

Locus ID: 22881

UniProt ID: [Q9Y2G4](#)

Cytogenetics: 6q15

MW: 75.7 kDa

Gene Summary: Recruits CKI-epsilon to the beta-catenin degradation complex that consists of AXN1 or AXN2 and GSK3-beta and allows efficient phosphorylation of beta-catenin, thereby inhibiting beta-catenin/Tcf signals.[UniProtKB/Swiss-Prot Function]
Transcript Variant: This variant (4) lacks an in-frame exon in the coding region, compared to variant 1. The resulting isoform (c) lacks an internal segment, compared to isoform a.
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.