

Product datasheet for **SC331469**

WDR27 (NM_001202550) Human Untagged Clone

Product data:

Product Type:	Expression Plasmids
Product Name:	WDR27 (NM_001202550) Human Untagged Clone
Tag:	Tag Free
Symbol:	WDR27
Vector:	pCMV6-Entry (PS100001)



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Fully Sequenced ORF: >SC331469 representing NM_001202550.
 Blue=Insert sequence Red=Cloning site Green=Tag(s)

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ATGAAAAATCCCCAAGACATTTTCTCAAGTAATGGTGGCTGTCTAAGTGATATAGTTATAGAAAAATAC
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CACCATCAGCCAATTACTGCTATGGCTTTTGGAAATAAAGTGAACCCACTTCTAATCTGCTCAGCATCA
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CAGCTTGCCACAGCTACCTTGGATGGTAAACTCCAGCTCTTTCTAGCTGAGTAA
  
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Restriction Sites: SgfI-MluI

ACCN: NM_001202550

Insert Size: 2193 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_001202550.1](#)

RefSeq Size: 3096 bp

RefSeq ORF: 2193 bp

Locus ID: 253769

UniProt ID: [A2RRH5](#)

Cytogenetics: 6q27

MW: 79.7 kDa

Gene Summary: This gene encodes a protein with multiple WD repeats. Proteins with these repeats may form scaffolds for protein-protein interaction and play key roles in cell signalling. Alternative splicing results in multiple transcript variants, but the full-length structure of some of these variants cannot be determined. [provided by RefSeq, Nov 2015]
Transcript Variant: This variant (2) lacks three alternate exons that result in the loss of an in-frame segment in the 5' coding region, and lacks another alternate exon that results in a frameshift in the 3' coding region, compared to variant 1. The encoded isoform (2) has a distinct C-terminus and 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.