

Product datasheet for SC330855

CA6 (NM_001270502) Human Untagged Clone

Product data:

Product Type: Expression Plasmids

Product Name: CA6 (NM_001270502) Human Untagged Clone

Tag: Tag Free Symbol: CA6

Synonyms: CA-VI; GUSTIN

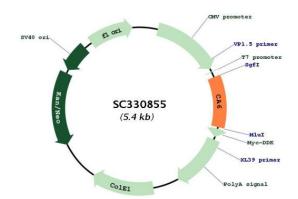
Vector: pCMV6-Entry (PS100001)

Fully Sequenced ORF: >SC330855 representing NM_001270502.

Blue=Insert sequence Red=Cloning site Green=Tag(s)

Restriction Sites: Sgfl-Mlul

Plasmid Map:





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CA6 (NM_001270502) Human Untagged Clone - SC330855

ACCN: NM 001270502

Insert Size: 543 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 001270502.1

RefSeq Size:1049 bpRefSeq ORF:543 bpLocus ID:765

Cytogenetics: 1p36.23

Protein Families: Druggable Genome, Secreted Protein

Protein Pathways: Nitrogen metabolism

MW: 21.2 kDa

Gene Summary: The protein encoded by this gene is one of several isozymes of carbonic anhydrase. This

protein is found only in salivary glands and saliva and protein may play a role in the

reversible hydratation of carbon dioxide though its function in saliva is unknown. [provided

by RefSeq, Jul 2008]

Transcript Variant: This variant (4) lacks two consecutive exons in the 5' coding region, compared to variant 1. The resulting isoform (4) has a shorter and distinct N-terminus,

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.