

## **Product datasheet for SC200856**

## CRYBB2 (NM\_000496) Human 3' UTR Clone

**Product data:** 

Product Type: 3' UTR Clones

Symbol: CRYBB2

Synonyms: CCA2; CRYB2; CRYB2A; CTRCT3; D22S665

Mammalian Cell Neomycin

Selection:

**Vector:** pMirTarget (PS100062)

ACCN: NM\_000496

Insert Size: 127 bp

Insert Sequence: >SC200856 3'UTR clone of NM\_000496

The sequence shown below is from the reference sequence of NM\_000496. The complete sequence of

this clone may contain minor differences, such as SNPs.

Blue=Stop Codon Red=Cloning site

GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAGGCCAAGAAGGGCGGAAAGATCGCCGTG

TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC

 ${\tt CAACGTGGTGCCTTCCACCCCTCCAACTAGTGCCCTCCCCACCATGCCTCCTTCCCAGGACCCAGGTCT}$ 

GCTGCCCAGGAACCCTCCAGACCTCCCAGAGAGTGAATAAAGTGTGACTTGCAACTTG

CGAGATTTCGATTCCACCGCCGCCTTCTATGAAAGG

Restriction Sites: Sgfl-Mlul

OTI Disclaimer: Our molecular clone sequence data has been matched to the sequence identifier above as a

point of reference. Note that the complete sequence of this clone is largely the same as the reference sequence but may contain minor differences, e.g., single nucleotide polymorphisms

(SNPs).

Components: The cDNA clone is shipped in a 2-D bar-coded Matrix tube as 10 ug dried plasmid DNA. The

package also includes 100 pmols of both the corresponding 5' and 3' vector primers in

separate vials.

**Note:** Plasmids are not sterile. For experiments where strict sterility is required, filtration with 0.22um

filter is required.



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**RefSeq:** <u>NM\_000496.3</u>

Summary: Crystallins are separated into two classes: taxon-specific, or enzyme, and ubiquitous. The

cause type 2 cerulean cataracts. [provided by RefSeq, Jul 2008]

latter class constitutes the major proteins of vertebrate eye lens and maintains the transparency and refractive index of the lens. Since lens central fiber cells lose their nuclei during development, these crystallins are made and then retained throughout life, making them extremely stable proteins. Mammalian lens crystallins are divided into alpha, beta, and gamma families; beta and gamma crystallins are also considered as a superfamily. Alpha and beta families are further divided into acidic and basic groups. Seven protein regions exist in crystallins: four homologous motifs, a connecting peptide, and N- and C-terminal extensions. Beta-crystallins, the most heterogeneous, differ by the presence of the C-terminal extension (present in the basic group, none in the acidic group). Beta-crystallins form aggregates of different sizes and are able to self-associate to form dimers or to form heterodimers with other beta-crystallins. This gene, a beta basic group member, is part of a gene cluster with beta-A4, beta-B1, and beta-B3. A chain-terminating mutation was found to

**Locus ID:** 1415

**MW:** 4.7