

Product datasheet for SC205194

OriGene Technologies, Inc.

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Sodium bicarbonate transporter like protein 11 (SLC4A11) (NM_032034) Human 3' UTR Clone

Product data:

Product Type: 3' UTR Clones

Product Name: Sodium bicarbonate transporter like protein 11 (SLC4A11) (NM_032034) Human 3' UTR Clone

Symbol: Sodium bicarbonate transporter like protein 11
Synonyms: BTR1; CDPD1; CHED; CHED2; dJ794l6.2; NABC1

Mammalian Cell

Selection:

Neomycin

Vector: pMirTarget (PS100062)

ACCN: NM_032034

Insert Size: 395 bp

Insert Sequence: >SC205194 3'UTR clone of NM_032034

The sequence shown below is from the reference sequence of NM_032034. The complete

sequence of this clone may contain minor differences, such as SNPs.

Blue=Stop Codon Red=Cloning site

GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAGGCCAAGAAGGGCGGAAAGATCGCCGTG

TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC

TAAGGAACTTGCTGGACGCACATTAGAGAATAAAGCTCCTGTTTCTAGGC

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.





Sodium bicarbonate transporter like protein 11 (SLC4A11) (NM_032034) Human 3' UTR Clone – SC205194

RefSeq: <u>NM 032034.4</u>

Summary: This gene encodes a voltage-regulated, electrogenic sodium-coupled borate cotransporter

that is essential for borate homeostasis, cell growth and cell proliferation. Mutations in this gene have been associated with a number of endothelial corneal dystrophies including recessive corneal endothelial dystrophy 2, corneal dystrophy and perceptive deafness, and Fuchs endothelial corneal dystrophy. Multiple transcript variants encoding different isoforms

have been described. [provided by RefSeq, Mar 2010]

Locus ID: 83959 **MW:** 13.9