

Product datasheet for SC212095

CHRFAM7A (NM 148911) Human 3' UTR Clone

Product data:

Product Type: 3' UTR Clones

Product Name: CHRFAM7A (NM_148911) Human 3' UTR Clone

Symbol: CHRFAM7A

Synonyms: CHRNA7; CHRNA7-DR1; D-10; NACHRA7

Mammalian Cell

Selection:

Neomycin

Vector: pMirTarget (PS100062)

ACCN: NM_148911

Insert Size: 1083 bp

Insert Sequence: >SC212095 3'UTR clone of NM_148911

The sequence shown below is from the reference sequence of NM_148911. The complete

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

Blue=Stop Codon Red=Cloning site

GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAGGCCAAGAAGGGCGGAAAGATCGCCGTG

TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC

GGGCCAGGAAGTAGAGGCTGGGAACTCTTCTGGTCCCCAGTATGGCAG

CGAGATTTCGATTCCACCGCCGCCTTCTATGAAAGG

Restriction Sites: Sgfl-Mlul



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CHRFAM7A (NM_148911) Human 3' UTR Clone - SC212095

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.

RefSeq: <u>NM 148911.1</u>

Summary: The nicotinic acetylcholine receptors (nAChRs) are members of a superfamily of ligand-gated

ion channels that mediate fast signal transmission at synapses. The family member CHRNA7, which is located on chromosome 15 in a region associated with several neuropsychiatric disorders, is partially duplicated and forms a hybrid with a novel gene from the family with sequence similarity 7 (FAM7A). Alternative splicing has been observed, and two variants exist, for this hybrid gene. The N-terminally truncated products predicted by the largest open reading frames for each variant would lack the majority of the neurotransmitter-gated ion-channel ligand binding domain but retain the transmembrane region that forms the ion channel. Although current evidence supports transcription of this hybrid gene, translation of the nicotinic acetylcholine receptor-like protein-encoding open reading frames has not been

confirmed. [provided by RefSeq, Jul 2008]

Locus ID: 89832 MW: 41.3