

# Product datasheet for SC313352

## CHRFAM7A (NM\_139320) Human Untagged Clone

### **Product data:**

#### OriGene Technologies, Inc.

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Product Type:	Expression Plasmids
Product Name:	CHRFAM7A (NM_139320) Human Untagged Clone
Tag:	Tag Free
Symbol:	CHRFAM7A
Synonyms:	CHRNA7; CHRNA7-DR1; D-10; NACHRA7
Mammalian Cell Selection:	Neomycin
Vector:	pCMV6-Entry (PS100001)
E. coli Selection:	Kanamycin (25 ug/mL)
Fully Sequenced ORF:	<pre>&gt;SC313352 representing NM_139320. Blue=Insert sequence Red=Cloning site Green=Tag(s)</pre>
	GCTCGTTTAGTGAACCGTCAGAATTTTGTAATACGACTCACTATAGGGCGGCCGGGAATTCGTCGACTG GATCCGGTACCGAGGAGATCTGCCGCCGCGATCGCC ATGCAAAAATATTGCATCTACCAGCATTTCAGTTCCAATTGCTAATCCAGCATTTGTGGATAGCTGCA AACTGCGATATTGCTGATGAGCGCTTTGACGCCACATTCCACACTAACGTGTTGGTGAATTCTTCTGGG CATTGCCAGTACCTGCCTCCAGGCATATTCAAGAGTTCCTGCTACATCGATGTACGCTGGTTTCCCTTT GATGTGCAGCACTGCAAACTGAAGTTTGGGTCCTGGTCTTACGGAGGCTGGTCCTTGGATCTGCAGATG CAGGAGGCAGATATCAGTGGCTATATCCCCAATGGAGAATGGGACCTAGTGGGAATCCCCGGCAAGAGG AGTGAAAGGTTCTATGAGTGCTGCAAAGAGCCCTACCCTGATGTCACCTTCACAGTGACCATGGCCGC AGGACGCTCTACTATGGCCTCAACCTGCTGATCCCCTGTGTGCTCATCTCCGGCCCTGGCGC AGGACGCTCTACTATGGCCTCAACCTGCTGATCCCCTGGTGGTCCATCTCCGCCCTGCTGGTG TTCCTGCTTCCTGCAGATTCCGGGGGAGAAGATTTCCCTGGGGATAACAGTCTTACTCTCTCT
<b>Restriction Sites:</b>	Sgfl-Mlul



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CHRFAM7A (NM_139320) Human Untagged Clone – SC313352	
ACCN:	NM_139320
Insert Size:	1239 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).
OTI Annotation:	This TrueClone is provided through our Custom Cloning Process that includes sub-cloning into OriGene's pCMV6 vector and full sequencing to provide a non-variant match to the expected reference without frameshifts, and is delivered as lyophilized plasmid DNA.
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:	<ol> <li>Centrifuge at 5,000xg for 5min.</li> <li>Carefully open the tube and add 100ul of sterile water to dissolve the DNA.</li> <li>Close the tube and incubate for 10 minutes at room temperature.</li> <li>Briefly vortex the tube and then do a quick spin (less than 5000xg) to concentrate the liquid at the bottom.</li> <li>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.</li> </ol>
RefSeq:	<u>NM 139320.1</u>
RefSeq Size:	2858 bp
RefSeq ORF:	1239 bp
Locus ID:	89832
UniProt ID:	<u>Q494W8</u>
Cytogenetics:	15q13.2
Domains:	Neur_chan_memb, Neur_chan_LBD
Protein Families:	Druggable Genome, Ion Channels: Other, Transmembrane
MW:	46.2 kDa

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#### **GRIGENE** CHRFAM7A (NM\_139320) Human Untagged Clone – SC313352

Gene Summary:The nicotinic acetylcholine receptors (nAChRs) are members of a superfamily of ligand-gated<br/>ion channels that mediate fast signal transmission at synapses. The family member CHRNA7,<br/>which is located on chromosome 15 in a region associated with several neuropsychiatric<br/>disorders, is partially duplicated and forms a hybrid with a novel gene from the family with<br/>sequence similarity 7 (FAM7A). Alternative splicing has been observed, and two variants exist,<br/>for this hybrid gene. The N-terminally truncated products predicted by the largest open<br/>reading frames for each variant would lack the majority of the neurotransmitter-gated ion-<br/>channel ligand binding domain but retain the transmembrane region that forms the ion<br/>channel. Although current evidence supports transcription of this hybrid gene, translation of<br/>the nicotinic acetylcholine receptor-like protein-encoding open reading frames has not been<br/>confirmed. [provided by RefSeq, Jul 2008]<br/>Transcript Variant: This variant (1) is the full-length variant and encodes the longer isoform (1).

Sequence Note: The RefSeq transcript and protein were derived from genomic sequence to make the sequence consistent with the reference genome assembly. The genomic coordinates used for the transcript record were based on alignments.

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