

OriGene Technologies, Inc.

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Product datasheet for SC212593

KCNQ1 (NM_181798) Human 3' UTR Clone

Product data:

Product Type:	3' UTR Clones
Product Name:	KCNQ1 (NM_181798) Human 3' UTR Clone
Vector:	pMirTarget (PS100062)
Symbol:	KCNQ1
Synonyms:	ATFB1; ATFB3; JLNS1; KCNA8; KCNA9; Kv1.9; Kv7.1; KVLQT1; LQT; LQT1; RWS; SQT2; WRS
ACCN:	NM_181798
Insert Size:	1153 bp
Insert Sequence:	>SC212593 3'UTR clone of NM_181798 The sequence shown below is from the reference sequence of NM_181798. The complete sequence of this clone may contain minor differences, such as SNPs. Blue=Stop Codon Red=Cloning site
	GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAGGCCAAGAAGGGCGGAAAGATCGCCGTG TAACAATTGGCAGAGCTCAGAATTCCAAGCGATCGCC CCCAGGAGGGCCCCGATGAGGGGTCCTGAGGAGAGGGGATGGGGCTGGGGGATGGGCCTGAGTGAG
Restriction Sites:	Sgfl-Mlul



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	KCNQ1 (NM_181798) Human 3' UTR Clone – SC212593
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 181798.1</u>
Summary:	This gene encodes a voltage-gated potassium channel required for repolarization phase of the cardiac action potential. This protein can form heteromultimers with two other potassium channel proteins, KCNE1 and KCNE3. Mutations in this gene are associated with hereditary long QT syndrome 1 (also known as Romano-Ward syndrome), Jervell and Lange-Nielsen syndrome, and familial atrial fibrillation. This gene exhibits tissue-specific imprinting, with preferential expression from the maternal allele in some tissues, and biallelic expression in others. This gene is located in a region of chromosome 11 amongst other imprinted genes that are associated with Beckwith-Wiedemann syndrome (BWS), and itself has been shown to be disrupted by chromosomal rearrangements in patients with BWS. Alternatively spliced transcript variants have been found for this gene. [provided by RefSeq, Aug 2011]
Locus ID:	3784
MW:	41.8

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