

Product datasheet for **RG217175**

Nav1.5 (SCN5A) (NM_001099404) Human Tagged ORF Clone

Product data:

Product Type:	Expression Plasmids
Product Name:	Nav1.5 (SCN5A) (NM_001099404) Human Tagged ORF Clone
Tag:	TurboGFP
Symbol:	SCN5A
Synonyms:	CDCD2; CMD1E; CMPD2; HB1; HB2; HBBB; HH1; ICCD; IVF; LQT3; Nav1.5; PFHB1; SSS1; VF1
Mammalian Cell Selection:	Neomycin
Vector:	pCMV6-AC-GFP (PS100010)
E. coli Selection:	Ampicillin (100 ug/mL)
ORF Nucleotide Sequence:	>RG217175 representing NM_001099404 Red=Cloning site Blue=ORF Green=Tags(s)

TTTTGTAATACGACTCACTATAGGGCGGCCGGAATTCGTCGACTGGATCCGGTACCGAGGAGATCTGCC
GCC**CGATCGCC**

ATGGCAAACCTTCTATTACCTCGGGCACCAGCAGCTTCCGCAGGTTACACGGGAGTCCCTGGCAGCCA
TCGAGAAGCGCATGGCAGAGAAGCAAGCCCGGGCTCAACCACCTTCAGGAGAGCCGAGAGGGGCTGCC
CGAGGAGGAGGCTCCCCGGCCCCAGCTGGACCTGCAGGCCCTCAAAAAGCTGCCAGATCTCTATGGCAAT
CCACCCCAAGAGCTCATCGGAGAGCCCCGGAGGACCTGGACCCCTTCTATAGCACCCAAAAGACTTTCA
TCGTACTGAATAAAGGCAAGACCATCTCCGGTTCAGTGCCACCAACGCCTTGTATGTCTCAGTCCCTT
CCACCCCATCCGGAGAGCGGCTGTGAAGATTCTGGTTCAGTCCGCTTCAACATGCTCATATGTGACC
ATCCTCACCAACTGCGTGTTCATGGCCAGCACACCCTCCACCTGGACCAAGTATGTCGAGTACACCT
TCACCGCCATTTACACCTTTGAGTCTCTGGTCAAGATTCTGGCTCGAGGCTTCTGCCTGCACGCTTAC
TTTCTTCGGGACCCATGGAACCTGGCTGGACTTTAGTGTGATTATCATGGCGTATGTATCAGAAAATATA
AACTAGGCAATTTGTCGGCTCTTCGAACCTTCAGAGTCTGAGAGCTCTAAAACTATTTAGTTATCC
CAGGGTGAAGACCATCGTGGGGCCCTGATCCAGTCTGTGAAGAAGCTGGCTGATGTGATGGTCCCTCAC
AGTCTTCTGCCTCAGCGTCTTTGCCCTCATCGGCTGCAGCTCTCATGGCAACCTAAGGCACAAGTGC
GTGCGCAACTTCACAGCGCTCAACGGCACCAACGGCTCCGTGGAGGCCGACGGCTTGGTCTGGGAATCCC
TGGACCTTACCTCAGTGATCCAGAAAATTACCTGCTCAAGAACGGCACCTCTGATGTGTTACTGTGTGG
GAACAGCTCTGACGCTGGACATGTCCGGAGGGTACCGTGCCTAAAGGCAGGCGAGAACCCCGACCAC
GGCTACACCAGCTTCGATTCCTTTGCCTGGCCTTTCTTGCACTCTCCGCTGATGACGCAAGGACTGCT
GGGAGCGCTCTATCAGCAGACCCTCAGGTCCGCAGGGAAGATCTACATGATCTTCTTATGCTTGTGAT
CTTCTGGGGTCTTCTACCTGGTGAACCTGATCTGGCCGTGGTGCATGGCCATGAGGAGCAAAAAC
CAAGCCACCATCGCTGAGACCGAGGAGAAGGAAAAGCGCTTCCAGGAGGCCATGGAAATGCTCAAGAAA
AACACGAGGCCCTCACCATCAGGGGTGTGGATACCGTGTCCGTAAGCTCCTTGGAGATGTCCCTTTGGC
CCCAGTAAACAGCCATGAGAGAAGAAGCAAGAGGAGAAAACGGATGTCTCAGGAACTGAGGAGTGTGG



[View online »](#)

GAGGACAGGCTCCCCAAGTCTGACTCAGAAGATGGTCCCAGAGCAATGAATCATCTCAGCCTCACCCGTG
 GCCTCAGCAGGACTTCTATGAAGCCACGTTCCAGCCGCGGGAGCATTTCACCTTCGCAGGCGAGACCT
 GGGTTCTGAAGCAGATTTTGAGATGATGAAAACAGCACAGCGGGGAGAGCGAGAGCCACCACATCA
 CTGCTGGTGCCTGGCCCTGCGCCGACCAGTGCCAGGGACAGCCAGTCCCGAACTCGGCTCCTG
 GCCAGCCCTCCATGGCAAAAAGAACAGCACTGTGGACTGCAATGGGGTGGTCTCATTACTGGGGCAGG
 CGACCCAGAGGCCACATCCCCAGGAAGCCACTCTCCGCCCTGTGATGCTAGAGCACCCGCCAGACAG
 ACCACGCCATCGAGAGGCCAGGCGGGCCAGATGCTGACCTCCAGGCTCCGTGTGATGGCTTCG
 AGGAGCCAGGAGCACGGCAGCGGCCCTCAGCGAGTCAGCGTCCACCAGCGCACTGGAAGAGTTAGA
 GGAGTCTCGCCACAAGTGTCCACCATGTGGAACCGTCTCGCCAGCGCTACCTGATCTGGGAGTGTGC
 CCGCTGTGGATGTCCATCAAGCAGGGAGTGAAGTTGGTGGTCATGGACCCGTTTACTGACCTCACCATCA
 CTATGTGCATCGTACTCAACACTCTTTCATGGCGCTGGAGCACTACAACATGACAAGTGAATTCGAGGA
 GATGCTGCAGGTCGAAACCTGGTCTTACAGGGATTTTACAGCAGAGATGACCTTCAAGATCATTGCC
 CTCGACCCCTACTACTTCCAACAGGGCTGGAACATCTTCGACAGCATCATCGTCATCCTTAGCCTCA
 TGGAGCTGGGCCGTCCCGCATGAGCAACTTGTGGTGTGCGCTCCTCCGCCCTGCTGCGGGTCTTCAA
 GCTGGCCAAATCATGGCCACCCTGAACACTCATCAAGATCATCGGAACTCAGTGGGGCAGTGGGG
 AACCTGACACTGGTGTAGCCATCATCGTGTTCATCTTTGCTGTGGTGGGATGCAGCTCTTTGGCAAGA
 ACTACTCGGAGCTGAGGGACAGCGACTCAGGCCTGCTGCCTCGCTGGCACATGATGGACTCTTTTATGC
 CTTCCTCATCATCTTCCGCATCCTCTGTGGAGAGTGGATCGAGACCATGTGGACTGCATGGAGGTGTG
 GGGCAGTCATTATGCCTGCTGGTCTTCTTGCTTGTATGGTCATTGGCAACCTTGTGGTCTGAATCTCT
 TCCTGGCCTTGTGCTCAGTCCCTCAGTGCAGACAACCTCACAGCCCTGATGAGGACAGAGAGATGAA
 CAACCTCCAGCTGGCCCTGGCCCGCATCCAGAGGGGCTGCGCTTGTCAAGCGGACCACCTGGGATTTCT
 TGCTGTGGTCTCCTGCGCAGCGGCCCTCAGAAGCCCGCAGCCCTTGGCCCGGAGGCCAGCTGCCAGCT
 GCATGCCACCCCTACTCCCCGCCACCCAGAGAGAGAGAGAGAGTGCCTCCCACCCGAAAGAACAGC
 GTTTGAGGAAGGCGAGCAACCAGGCCAGGGCACCCCGGGATCCAGAGCCCGTGTGTGCCCATCGCT
 GTGGCCGAGTCAGACACAGATGACCAAGAAGAAGATGAGGAGAACAGCCTGGGCACGAGGAGGAGTCCA
 GCAAGCAGCAGGAATCCCAGCCTGTGTCCGGTGGCCAGAGGCCCTCCGGATCCAGGACCTGGAGCCA
 GGTGTGAGGACTGCCTCCTCTGAGGCCAGGCCAGTGCATCTCAGGCCACTGGCGGCAGCAGTGGAAA
 GCGGAACCCAGGCCAGGGTGCAGTGCAGACCCAGAGGACAGTTGCTCCGAGGGCAGCACAGCAGACA
 TGACCAACACCGCTGAGCTCCTGGAGCAGATCCCTGACCTCGGCCAGGATGTCAAGGACCCAGAGGACTG
 CTTCACTGAAGGCTGTGTCGGCGCTGTCCCTGCTGTGCGGTGGACACCACAGGCCCCAGGGAAGGTC
 TGGTGGCGGTTGGCAAGACCTGTACACATCGTGGAGCACAGCTGGTTCGAGACATTCATCATCTTCA
 TGATCCTACTCAGCAGTGGAGCGCTGGCCCTTCAGGACATCTACCTAGAGGAGCGGAAGACCATCAAGGT
 TCTGCTTGAGTATGCCGACAAGATGTTACATATGTCTTCTGCTGGAGATGCTGCTCAAGTGGGTGGCC
 TACGGCTTCAAGAAGTACTTACCAATGCCTGGTGTGGCTCGACTTCTCATCGTAGACGTCTCTCTGG
 TCAGCCTGGTGGCCAACACCCCTGGGCTTTCGCCAGATGGGCCCATCAAGTCACTGCGGACGCTGCGTGC
 ACTCCGCTCCTCTGAGAGCTCTGTACGATTTGAGGGCATGAGGGTGGTGGTCAATGCCCTGGTGGCGCC
 ATCCCCTCCATCATGAACGTCTCCTCGTCTGCCTCATCTTCTGGCTCATCTTCAAGCATCATGGCGTGA
 ACCTCTTTGCGGGGAAGTTTGGGAGTGCATCAACCAGACAGAGGGAGACTTGCCTTTGAACTACCCAT
 CGTGAACAACAAGAGCCAGTGTGAGTCTTGAACCTTGACCGGAGAATTGACTGGACCAAGTGAAGTC
 AACCTTTGACAACGTGGGGCCGGTACCTGGCCCTTCTGCAAGTGGCAACATTTAAAGGCTGGATGGACA
 TTATGTATGCAGCTGTGGACTCCAGGGGTATGAAGAGCAGCCTCAGTGGGAATACAACCTCATATGTA
 CATCTATTTTGTCAATTTTTCATCATCTTTGGGTCTTTCTTACCCTGAACCTCTTTATTGGTGTATCATT
 GACAACCTCAACCAACAGAAAGAAAGTTAGGGGGCCAGGACATCTTCATGACAGAGGAGCAGAAGAAGT
 ACTACAATGCCATGAAGAAGCTGGGCTCCAAGAAGCCCCAGAAGCCATCCCACGGCCCTGAACAAGTA
 CCAGGGCTTCATATTGACATTGTGACCAAGCAGGCCCTTGGAGTCAACATCATGTTTCTGATCTGCTTG
 AATATGGTGAACATGATGGTGGAGACAGATGACCAAAAGTCTGAGAAAATCAACATCTTGGCCAAGATCA
 ACCTGCTCTTTGTGGCCATCTTACAGGGGAGTATTGTCAAGCTGGCTGCCCTGCGCCACTACTACTT
 CACCAACAGCTGGAATATCTTCCACTTCGTGGTGTGCATCCTCTCCATCGTGGGCACTGTCTCTCGGAC
 ATCATCCAGAAGTACTTCTTCTCCCGACGCTCTTCCGAGTCATCCGCCTGGCCCAATAGGCCGATCC
 TCAGACTGATCCGAGGGGCAAGGGGATCCGCACGCTGCTCTTTGCCCTCATGATGTCCCTGCCTGCCCT
 CTTCAACATCGGGCTGCTGCTCTTCTCGTGTGTTCACTACTCCATCTTTGGCATGGCCAACCTCGCT
 TATGTCAAGTGGGAGGCTGGCATCGACGACATGTTCAACTCCAGACCTTCGCCAACAGCATGCTGTGCC

TCTTCCAGATCACCACGTGCGCCGGCTGGGATGGCCTCCTCAGCCCCATCCTCAACACTGGGCCGCCCTA
 CTGCGACCCCACTCTGCCAACAGCAATGGCTCTCGGGGGGACTGCGGGAGCCAGCCGTGGGCATCCTC
 TTCTTACCACCTACATCATCTCTTCTCATCGTGGTCAACATGTACATTGCCATCATCTGGAGA
 ACTTCAGCGTGGCCACGGAGGAGCACCAGCCCTGAGTGAGGACGACTTCGATATGTTCTATGAGAT
 CTGGGAGAAATTTGACCCAGAGGCCACTCAGTTTATTGAGTATTCGGTCTGTCTGACTTTGCCGATGCC
 CTGTCTGAGCCACTCCGTATCGCCAAGCCCAACCAGATAAGCCTCATCAACATGGACCTGCCATGGTGA
 GTGGGGACCGCATCCATTGCATGGACATCTCTTTGCCTTACCCAAAAGGGTCTCGGGGAGCTGGGGA
 GATGGAGCCCTGAAGATCCAGATGGAGGAGAAGTTCATGGCAGCCAACCCATCCAAGATCTCTACGAG
 CCCATCACCACCACTCCGCGCAAGCACGAAGAGGTGTCGGCCATGGTTATCCAGAGAGCCTTCCGCA
 GGCACCTGCTGCAACGCTCTTTGAAGCATGCCTCCTTCTTCCGTGAGCAGGCGGGCAGCGCCCTCTC
 CGAAGAGGATGCCCTGAGCGAGAGGGCCTCATCGCTACGTGATGAGTGAGAATTCTCCGACCCCTT
 GGCCACCCTCCAGCTCTCCATCTCTCCACTTCTTCCACCCTCTATGACAGTGTACTAGAGCCA
 CCAGCGATAACCTCCAGTGGGGGTCTGACTACAGCCACAGTGAAGATCTCGCCGACTCCCCCTTC
 TCCGGACAGGACCGTGAGTCCATCGTG

AGCGGACCGACGCGTACGCGGCCGCTCGAG - GFP Tag - GTTTAA

Protein Sequence:

>RG217175 representing NM_001099404
 Red=Cloning site Green=Tags(s)

MANFLLPRGTSSFRFTRESLAAIEKRMAEKQARGSTTLQESREGLPEEEAPRPQLDLQASKKLPDLYGN
 PPQELIGEPLDLPFYSTQKTFIVLNKGTIFRFSATNALYVLSPPFHPIRRAAVKILVHSLFNMLIMCT
 ILTNCVMAQHDPWPWKYVEYFTAIYTFESLVKILARGFCLHAFTFLRDPWNWLDVSVIIMAYVSENI
 KLGNSALRTRFRVLRALKTISVIPGLKTIYVIGALIQSVKKLADVMVLTVFCLSVFALIGLQLFMGNLRHKC
 VRNFTALNGTNGSVEADGLVWESLDLSDPENYLLKNGTSDVLLCGNSSDAGTCEGYRCLKAGENPDH
 GYTSFDSFAWAFALFRLMTQDCWERLYQQTLSRAGKIYMIFFMLVIFLGSFYLVNLIILAVVAMAYEEQN
 QATIAETEKEKRFQEMEMLKKEHEALTIRGVDTVSRSSLEMSPLAPVNSHERRSKRRKRMSSGTEECG
 EDRLPKSDSEDPGRAMNHLSTRGLSRTSMKPRSSRGSIFTFRRRDLGSEADFADDENSTAGESESHHTS
 LLVWPWLRRTSAQGQPSPGTSAPGHALHGKKNSTVDCNGVVSLGAGDPEATSPGSHLLRPVMLEHPPDT
 TTPSEEPGGPQMLTSQAPCVDGFEEPGARQRALSAVSVLTSALIEEESRHKCPPCWNRLAQRYL IWECC
 PLWMSIKQGVKLVVMDPFTDLTITMCIVLNTLFMALEHYNMTSEFEEMLVQGNLVFTGIFTAEMTFKIIA
 LDPYYFYQGWNI FDSIIIVILSMLGLSRMSNLVLSRFRLLRVFKLAKSWPTLNTL IKIIGNSVGALG
 NLTVLVAIIIVFIFAVVGMQLFGKNYSEL RSDSGLLPRWHMDFHAFLLIIFRILCGEWIETMWDCEVS
 GQSLCCLLVFLVMVIGNLVVNLFLALLSSFSADNLTAPDEDREMNNLQALARIQRGLRFVKRTTWDF
 CCGLLRQRPQKPAALAAQGQLPSCIATPYSPPPTEKVPTRKETRFEEGEQPGQGTGPDPEPVCVPIA
 VAESD TDDQEEDEENSLGTEEESKQESQPVSGGPEAPPDSRTWSQVSATASSEAEASASQADWRQWK
 AEPQAPGCGETPEDSCSEGSTADMTNTAELLEQIPDLGQDVKDPEDCFTEGCVRRCPCCAVDTTQAPGKV
 WWRLRKTCYHIVEHSWFETFIIFMILLSSGALAFEDIYLEERKTIKVLLLEYADKMFYVFLVLEMLLKWVA
 YGFKKYFTNAWCWLDLFLIVDVSLVSLVANTLGF AEMGPIKSLRTRLRALRPLRALSRFEGMRVVVNLVGA
 IPSIMVLLVCLIFWLIFSIMGVNLFAGKFGRCINQTEGDLPLNYTIVNNKSQCESLNLGELYWTKVKV
 NFDNVGAGYLALLQVATFKGWMDIMYAAVDSRGYEEQPQWEYNLYMYIYFVIFIIIFGSFFTLNLFIGVII
 DNFNQKQKLLGGQDIFMTEEQKYYNAMKKGSKKPQKPIPRPLNKYQGFIFDIVTKQAFDVTIMFLICL
 NMVTMMVETDDQSPEKINILAKINLLFVAIFTGECIVKLAALRHHYFTNSWNI FDFVVVILSIVGTVLS
 IIQKYFFSPTLFRVIRLARIGRILRLIRGAKGIRTLFFALMMLPALFNIGLLLFLVMFIYSIFGMANFA
 YVWEAGIDDMFNFTFANSMLCLFQITTSAGWDGLLSPILNTGPPYCDPTLPNSNGSRGDCGSPAVGIL
 FFTTYIIISFLIVNMYIAIILENF SVATEESTEPL SEDDFDMFYEIWEKFDPEATQFIEYSVLSDFADA
 LSEPLRIAKPNQISLINMDLPMVSGDRICMDILFAFTKRVLGEGEMDALKIQMEEKFMAANPSKISYE
 PITTTTLRRKHEEVSAMVIQRAFRRHLLQRSLKHASFLFRQQAGSGLSEEDAPEREGLIAYMSENF SRPL
 GPPSSSISSTSFPPSYDSVTRATSDNLQVRGSDYSHSEDLADFPSPDRDRESIV

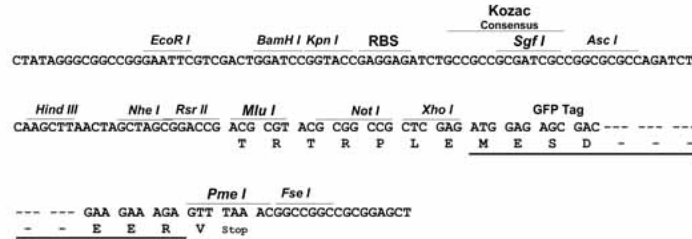
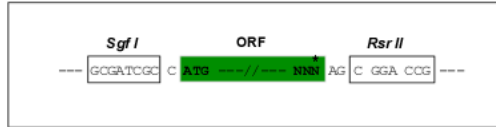
SGPTRRRLE - GFP Tag - V

Restriction Sites:

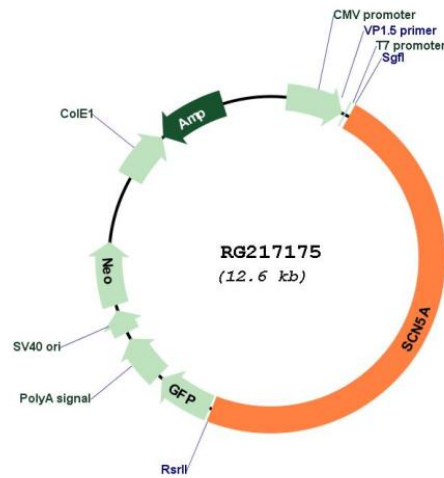
Sgfl-RsrII

Cloning Scheme:

Cloning sites used for ORF Shuttling:



Plasmid Map:



ACCN: NM_001099404

ORF Size: 6048 bp

OTI Disclaimer: The molecular sequence of this clone aligns with the gene accession number as a point of reference only. However, individual transcript sequences of the same gene can differ through naturally occurring variations (e.g. polymorphisms), each with its own valid existence. This clone is substantially in agreement with the reference, but a complete review of all prevailing variants is recommended prior to use. [More info](#)

OTI Annotation: This clone was engineered to express the complete ORF with an expression tag. Expression varies depending on the nature of the gene.

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 style="list-style-type: none">1. Centrifuge at 5,000xg for 5min.2. Carefully open the tube and add 100ul of sterile water to dissolve the DNA.3. Close the tube and incubate for 10 minutes at room temperature.4. Briefly vortex the tube and then do a quick spin (less than 5000xg) to concentrate the liquid at the bottom.5. 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.
RefSeq:	<u>NM_001099404.2</u>
RefSeq Size:	8504 bp
RefSeq ORF:	6051 bp
Locus ID:	6331
UniProt ID:	<u>Q14524</u>
Cytogenetics:	3p22.2
Protein Families:	Druggable Genome, Ion Channels: Sodium, Transmembrane
Gene Summary:	The protein encoded by this gene is an integral membrane protein and tetrodotoxin-resistant voltage-gated sodium channel subunit. This protein is found primarily in cardiac muscle and is responsible for the initial upstroke of the action potential in an electrocardiogram. Defects in this gene are a cause of long QT syndrome type 3 (LQT3), an autosomal dominant cardiac disease. Alternative splicing results in several transcript variants encoding different isoforms. [provided by RefSeq, Jul 2008]