

Product datasheet for MR215244

Scn2a (NM_001099298) Mouse Tagged ORF Clone

Product data:

Product Type:	Expression Plasmids
Product Name:	Scn2a (NM_001099298) Mouse Tagged ORF Clone
Tag:	Myc-DDK
Symbol:	Scn2a
Synonyms:	6430408L10; A230052E19Rik; Nav; Nav1.2; Scn; Scn2a1
Mammalian Cell Selection:	Neomycin
Vector:	pCMV6-Entry (PS100001)
E. coli Selection:	Kanamycin (25 ug/mL)
ORF Nucleotide Sequence:	>MR215244 representing NM_001099298 Red=Cloning site Blue=ORF Green=Tags(s)

TTTTGTAATACGACTCACTATAGGGCGGCCGGAATTCGTCGACTGGATCCGGTACCGAGGAGATCTGCC
GCCCGATCGCC

ATGGCACAATCAGTGCTGGTACCGCCAGGACCTGACAGCTTCGGCTTCTTTACCAGGGAATCCCTTGCTG
CTATTGAACAACGCATTGCAGAAGAGAAAGCTAAGAGACCCAAACAGGAACGCAAGGACGAAGACGACGA
AAATGGCCCAAAGCCAAATAGTGACTTGGAAAGCAGGAAAATCCCTTCCTTTTATTTATGGAGACATTCCT
CCAGAGATGGTGTGAGAGCCCTGGAGGACCTGGACCCCTACTATATCAATAAGAAAACCTTTATAGTAT
TGAAATAAGGAAAGCAATCTCTCGGTTGAGTCCACCTCGCCCTGTACATTTAACTCCCTTCAACCC
CATTAGAAAATTAGCTATTAAGATTTTGGTACATTCCTTTATTCAATGTGCTTATCATGTGACAATTCTT
ACCAACTGTGATTTATGACCATGAGCAACCTCCAGACTGGACAAGAATGTGGAGTATACTTTTACAG
GAATTTATACTTTTGAATCACTTATTAATACTTGAAGGGGCTTTTGTCTAGAAGATTTACATTTCT
ACGGGATCCCTGGAACCTGGCTGGATTTACAGTCACTACCTTTGCGTATGTAACAGAATTTGAAACCTA
GGCAATGTTTCAGCTCTTCGAACCTTCAGAGTCTTGAGAGCTTTGAAAATTTCTGTAATTCAGGCC
TGAAGACCATCGTGGGGCCCTGATCCAGTCCGTGAAGAAGCTGTCTGACGTGATGACTCACTGTGTT
CTGTCTGAGTGTCTTTGCTCTCATCGGGCTGCAGCTCTTCATGGCAACCTGAGGAATAAATGCTTGCAG
TGGCCCCAGACAATTCTACCTTTGAAATAAACATCACTTCCTTTTAACTCATTGGATTGGAATG
GTACTGCCTTCAATAGGACAATGAACATGTTTAACTGGGATGAATATATTGAAGATAAAAGTCACTTTTA
TTTTTTGGAAGGACAAAACGATGCTCTGCTTTGTGGTAACAGCTCGGATGCTGGCCAGTGTCCGGAAGGA
TACATCTGTGTGAAGGCTGGGAGAAACCCCAACTATGGCTACACGAGTTTGGACCTTCAGCTGGGCCCT
TCTTGTCTTATTTGCTTATGACTCAAGACTTCTGGGAAAACCTTTATCAGCTGACCTTGCCTGCTGC
TGGGAAAACATATATGATTTTTCTGCTTGTGATTTTCTGGGCTCATTCTACCTAATAAATTTGATC
CTGGCTGTGGTGGCCATGGCCTATGAAGAACAGAACCAGGCCACGCTGGAGGAAGCTGAACAGAAGGAGG
CAGAGTTTCAGCAGATGCTGGAGCACTTAAGAAGCAGCAAGAAGAAGCTCAGGCAGCAGCTGCAGCAGC
GTCTGCAGAAATCCAGAGACTTCAGTGGAGCAGGTGGGATAGGTGTTTTCTCGGAGAGTTCTCCGTAGCT



[View online »](#)

TCAAAGTTAAGTTCCAAGAGTGAAAAGGAGCTGAAAAACAGAAGGAAGAAAAAGAAACAGAAAGAGCAGG
 CTGGAGAGGAAGAGAAGGAAGATGCTGTGCGGAAATCTGCCTCTGAAGACAGCATACGAAAGAAAGGCTT
 CCGGTTTTCCCTAGAAGGGAGTAGACTGACCTACGAGAAGAGGTTTTCTCTCCACACCAGTCTCTTTG
 AGCATCCGAGGCTCCCTTTTTCTCCGAGACGCAACAGTAGAGCAAGCCTTTTCAGTTCAAAGGTCCGG
 TGAAGGATATTGGTTCTGAAAATGACTTTGCGGACGATGAACACAGCACGTTTGAGGACAACGACAGCAG
 GAGAGACTCTCTATTTGTACCACACAGACATGGAGAAAGGCGTCCAGCAACGTTAGCCAGGCCAGCCGT
 GCCTCCCGGGGATACCCACTCTACCCATGAATGGGAAGATGCACAGTGCAGTGGACTGCAACCGTGTGG
 TCTCCCTAGTTGGAGCCCTTCTGCTCTCACATCTCCTGTGGGGCAGCTACTACCAGAGGCCACAACACTAC
 TGAACAGAAAT AAGGAAGAGGAGTCCAGTCTTACCATGTTTCTATGGACTTGTGGAAGACCTTACA
 TCAAGGCAAAGAGCAATGAGTATGGCCAGCATTTTGACCAACACTATGGAAGAATTGAAGAATCCAGAC
 AGAAATGCCCTCCATGCTGGTATAAATTTGCTAATATGTGCCTGATTTGGGACTGTTGTAAGCCATGGCT
 AAAGTAAACATGTTGTCAATCTGGTAGTGATGGATCCATTGTTGACCTGGCCATCACCATCTGCATT
 GTGTTAAATACACTCTTCATGGCCATGGAGCACTACCCGATGACGGAGCAGTTCAGCAGTGTGCTGTCTG
 TTGGGAATTTGGTCTTCACTGGGATCTTACCAGCAAAATGTTTCTCAAGATAATAGCCATGGATCCATA
 TTATTACTTTCAAGAAGGCTGGAATATCTTTGATGGCTTTATAGTGAGCCTTAGTTTAAATGGAACCTCGG
 TTGGCGAATGTGGAAGGACTGTCAAGTCTCCGATCTTTCCGACTGCTTAGAGTCTCAAGTTGGCAAAAT
 CCTGGCCACACTGAATATGCTCATTAAAGATCATCGGCAACTCGTGGGCGCGCTGGGCAACCTGACCC
 GGTGCTGGCCATCATCGTCTTATTTTTGCCGTGGTCCGATGCAGCTGTTTGGAAAGAGCTACAAGGAG
 TGTGTTTGCAAGATTTCCAATGATTGTGAGCTGCCGCGCTGGCACATGCATGACTTCTTCCACTCGTTCC
 TGATCGTGTCCGCGTGTGTGGGAGTGGATAGAGACCATGTGGGACTGCATGGAGGTCCGAGGCCA
 GACCATGTGTCTACTGTCTTATGATGGTTCATGGTATTGGGAACCTTGTGTTCTGAACCTCTTCTTG
 GCCTTGTCTCAGTCTTTTCCAGCTCGGACAACCTGGCTGCCACAGATGACGATAACGAAATGAACAACC
 TCCAGATAGCCGTGGGAAGGATGCAAGAAGGGATTGATTTTGTAAAAGGAAGATACGTGAATCATTCA
 GAAAGCCTTTGTGAGAAAGCAGAAAGCTTAGATGAAATCAAACCCCTTGAAGATCTGAATAACAAGAAA
 GACAGTTGTATCTCAACCACACAACCATAGAAAATAGGCAAGGACCTGAATTACCTCAAAGATGGAACG
 GGACGACCAGTGGCATAGGCAGCAGTGTGGAGAAGTACGTGGTAGATGAGAGTGATTACATGTCATTTCAT
 CAACAACCCAGCCTCACCGTACTGTGCCATTGCTGTGGGAGAGTCTGACTTTGAAAACCTTAAATACA
 GAAGAATTCAGCAGTGAATCAGATATGGAAGAAAGCAAGGAGAAATGAAATGCAACTAGTTCATCTGAAG
 GCAGTACAGTTGATATAGGTGCTCCTGCAGAGGGAGCAACCAGAGGCCGAACCAAGAGTCCCTTGA
 GCCAGAAGCCTGTTTACAGAAGACTGTGTAAGAAAGTCAAGTGTGTCAGATAAGCATAGAAGAAGGC
 AAAGGCAAACCTGGTGGAACTTGAGGAAAACCTGCTACAAGATAGTAGAGCACAACCTGGTTTGAACCT
 TCATCGTCTTATGATTCTGCTCAGCAGCGGTGCTCTGGCCTTTGAAGACATTTACATCGAGCAGCGAAA
 AACCATCAAGACCATGCTGGAGTATGCTGACAAGTCTTCACTTACATCTTCATCCTGGAGATGTTGCTA
 AAGTGGTTGCATATGGTTTCAAATGACTTCAACATGCCTGGTGGTGGCTGGACTTCTGATCGTTG
 ATGCTCACTGGTTAGCTTAACTGCAAAATGCCTTGGGCTATTGAGAACTTGGTGCCATCAAATCCCTCCG
 AACATTAAGAGCACTGAGGCCTTACGAGCCTTATCCCGATTTGAAGGAATGAGGGTTGTTGTAACCGT
 CTCCTAGGTGCAATCCCATCCATAATGAACGACTTCTGGTCTGCCTGATCTTTTGGCTAATATTCAGTA
 TCATGGGGTGAATCTCTTTGCCGGCAAGTCTATCATTGCATCAACTACACCACCGGGAAATGTTTGA
 TGTGAGCGTGGTCAACAACACTACAGTGAAGTCCAGGCTCATAGAGAGCAATCAGACAGCCAGTGGGAAG
 AATGTGAAAGTCAACTTTGACAACGTGGGACTTGGATATCTTTCTGCTTCAAGTAGCCACATTTAAAG
 GATGGATGGATATCATGTATGCAGCTGTTGACTCAAGAAATGTAGAAGTCAACCCAAATATGAAGACAA
 TCTATACATGTACCTTTACTTTGTCTCTTTATTTTTCGGCTCATTCTTCACTGAACCTGTTTCATC
 GGCGTCATCATAGATAACTTCAACCAACAGAAGAAGAAGTTGGAGGTCAAGACATTTTTATGACAGAAG
 AACAGAAGAAATACTACAATGCAATGAAGAAGCTAGGCTCAAAGAAACCACAAAACCGATACCTCGCCC
 TGCTAACAAAGTTCAAGGGATGGTCTTTGATTTTGAACCAACAAGTGTGACATCAGCATCATGATC
 CTCATCTGCCTCAACATGGTGACCATGATGGTGGAAACGGACGACCAGAGCAAGAGATGACCAACATCC
 TGTATTGGATCAACCTGGTGTTCATCGTCTGTTACCGGCGAGTGTGTGCTCAAGCTTATCTCACTCCG
 TCATTATTTTACCATTGGATGGAACATTTTGTATTTGTGGTGGTCACTCTCCATTGTAGGGATG
 TTTCTCGCGGAGCTGATAGAGAAGTATTTCTGTGCTCCCTACCCTGTTCCGAGTCACTCCGCTGGCCAGGA
 TTGGACGAATCCTACGCCTGATCAAAGGCGCAAGGGGATCCGCACGCTGCTCTTTGCTCTGATGATGTC
 CCTTCTGCGCTGTTAACATCGGCCTCTGCTTTTCTGGTCACTGTTTCTACGCCATCTTTGGGATG
 TCCAACCTTGCCTACGTTAAGAGGGAAGTTGGGATTGATGACATGTTCAACTTTGAGACCTTTGGCAACA

GCATGATCTGCCTGTTCCAAATCACCACTTCTGCGGGCTGGGATGGACTGCTGGCCCTATACTAAATAG
 TGGACCTCCCATTGTGACCCTGAAAAAGATCATCCTGGAAGCTCGGTGAAGGGAGACTGTGGAAACCCA
 TCTGTGGGGATTTTCTTTTTGTGAGCTACATCATATCCTTCTGTTGTGGTGAACATGTACATTG
 CTGTATCCTGGAGAAGTTCAGCGTTGCCACAGAAAGTGCAGAGCCTCTGAGTGAGGACGACTTTGA
 GATGTTCTACGAAGTCTGGGAGAAGTTCGACCTGACGCCACCCAGTTCATAGAGTCTGCAAGCTCTCT
 GACTTTGCAGCTGCCCTGGATCCTCCCTCCTCATCGAAAGCCAAACAAAGTGCAGCTCATTGCCATGG
 ACCTGCCATGGTGGAGTGGAGACCCGATCCACTGCCTGGACATCTTATTTGCTTTTACAAAGCGGTCTC
 GGTGAGAGTGGAGAGATGGATGCTCTTGAATCCAGATGGAAGAGCGGTTTCATGGCTTCCAATCCCTCC
 AAGGTCTCTTATGAGCCATTACCACCACTCTGAAGCGCAAACAAGAGGAGGTGTCTGCTATTGTATCC
 AGCGAGCTTACAGACGCTATCTTCTGAAACAGAAAGTTAAGAAGGTTTCGTCTATATATAAAAAAGACAA
 GGGTAAAGAAGATGAGGGAACGCCCATCAAAGAAGACATCATCACGGATAAACTGAATGAGAATCAACT
 CCAGAGAAGACTGACGTGACACCCTCCACCACTTCGCCACCTTCTACGATAGCGTGACCAAACCGGAGA
 AGGAAAAATTTGAAAAGGACAAATCAGAAAAAGAGGACAAGGGGAAAGATATCAGGGAAAGTAAAAAG

ACGCGTACGCGGCCGCTCGAGCAGAAACTCATCTCAGAAGAGGATCTGGCAGCAAATGATATCCTGGATT
 ACAAGGATGACGACGATAAGGTTTAA

Protein Sequence:

>MR215244 representing NM_001099298
 Red=Cloning site Green=Tags(s)

MAQSVLVPPGPDSEFRFFTTRESLAAIEQRIAEKAKRPKQERKDEDDENGPKPNSDLEAGKSLPFIYGDIP
 PEMVSEPLEDLDPYYINKKTFIVLNKGAISRFSATSALYILTPFNPIRKLAIKILVHSLFNVLIMCTIL
 TNCVFMNTSNPPDWTKNVEYTFGTIYTFESLIKILARGFCLEDFTLRDPWNWLDFTVITFAYVTEFVNL
 GNVSALRTFRVLRALKTISVIPGLKTIIVGALIQSVKLSVDMILTVFCLSVFALIGLQLFMGNLRNKCLQ
 WPPDNSTFEINITSFFNNSLDWNGTAFNRMTMFMNWEYIEDKSHFYFLEGQNDALLCGNSSDAGQCPEG
 YICVKAGRNPNGYTSFDTFSWAFSLFRLMTQDFWENLYQLTLRAAGKTYMIFVFLVIFLGSFYLINLI
 LAVVAMAYEEQNQATLEAEQKEAEFQQMLEQLKKQEEAQAASRDFSGAGGIGVFSSESSVA
 SKLSSKSEKELNRRKKKKQKEQAGEEKEKEDAVRKSASEDSIRKKGFRFSLEGSRLTYEKRFSSPHQSL
 SIRGSLFSPRRNSRSLFSFKGRVKDIGSENDFADEHSTFEDNDSRRDSLFPVPHRHGERRPSNVSQASR
 ASRGIPTLPMNGKMHSAVDCNGVVSIVGGPSALTSVPVQLPEGTTTETEIRKRRSSSYHVSMDLLEDPT
 SRQRAMSMASILTNTMEELEESRQKCPPCWYKFNAMCLIWDCCKPWLKVKHVNLVVMDFVDLAITICI
 VLNTLFMAMEHYPMTEQFSSVLSVGNLVFTGIFTAEMFLKIIAMDPIYFQEGWNIFDGFIVLSLMELG
 LANVEGLSVLRSFRLLRVFKLAKSWPTLNMLIKIIGNSVGLGNLTLVLAIIIVFIFAVVGMQLFGKSYKE
 CVCKISNDCELPRWHMDFHFSFLIVFRVLGCEWIEIEMWDCMEVAGQTMCLTVFMMVMVIGNLVVNLFL
 ALLLSSFSSDNLAAATDDDNEMNQLIAGVGMQKIDFVKRKIREFIQKAFVRKQKALDEIKPLEDLNKK
 DSCISNHTTIEIGKDLNLYKDGNGTTSIGSSVEKYVVDSDYMSFINNPSLTVTVPIAVGESDFENLNT
 EEFSSSESDMEESKEKLNATSSSEGSTVDIGAPAEQEPEAEPEESLEPEACFTEDCVRKFKCCQISIEEG
 KGKLVWNLRKTCYKIVEHNWFETIVFMILLSSGALAFEDIYIEQRKTIKTMLEYADKVFYIFILEMLL
 KWWAYGFQMYFTNAWCWLDLIVDVSLVSLTANALGYSELGAIKSLRTRLRALRPLRALSREFEGMRVVNA
 LLGAIPSIMNVLLVCLIFWLIFSIMGVNLFAGKFYHCINYYTGMFVSVVNNYSECQALIESNQATARWK
 NVKVNFDNVGLGYLSLLQVATFKGWMDIMYAAVDSRNVELQPKYEDNLYMYLYFVIFIFIGSFTLNLFI
 GVIIDNFNQKKKFGQDIFMTEEQKYYNAMKKGSKPKQPIPRPANKFQGMVDFVTKQVDFDISIMI
 LICLNMTMMVETDDQSQEMTNILYWINLVFIVLFTGECVLKLSLRHYFTIGWNIIDFVIVVILSIVGM
 FLAELIEKYFVSPFLFRVIRLARIGRILRLIKGAKGIRTLFALMMSLPALFNIGLLFLVMFIYAIIFGM
 SNFAYVKREVGIDDMFNFTFGNSMICLFIQITTSAGWDGLLAPILNSGPPDCPEKDHPGSSVKGDGCPN
 SVGIFFFVSYIIISFLVVNMYIAVILENFVATEESAELPSEDDFEMFYEVEKFDPDATQFIEFCKLS
 DFAAALDPPLLIAKPNKQVLIAMDLPVSGDRIHCLDILFAFTKRVLGESGEMDALRIQMEERFMASNPS
 KVSYPITTTLKRKQEEVSAIIVIQRAYRRYLLKQKVKVSSIIYKKDKGKEDEGTPIKEDIITDKLNENST
 PEKTDVTPSTTSPPSYDSVTKPEKEKFEKDKSEKEDKGDIRESKK

TRTRPLEQKLISEEDLAANDILDYKDDDDKV

Restriction Sites:

Sgfl-MluI

Cloning Scheme:



ACCN: NM_001099298

ORF Size: 6018 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:

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: [NM_001099298.3](#)

RefSeq Size: 8690 bp

RefSeq ORF: 6021 bp

Locus ID: 110876

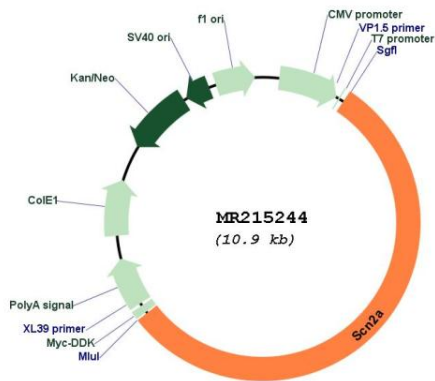
UniProt ID: [B1AWN6](#)

Cytogenetics: 2 38.61 cM

MW: 227.9 kDa

Gene Summary: Voltage-gated sodium channels are transmembrane glycoprotein complexes composed of a large alpha subunit with four repeat domains, each of which is composed of six membrane-spanning segments, and one or more regulatory beta subunits. Voltage-gated sodium channels are responsible for the generation and propagation of action potentials in neurons and muscle. This gene encodes one member of the sodium channel alpha subunit gene family. In humans, variants of this gene are associated with seizure disorders and autism spectrum disorder. Mice homozygous for a knockout mutation die with severe hypoxia and extensive neuronal cell death, while gain of function mutations result in progressive seizure disorder. Alternative splicing results in multiple transcript variants. [provided by RefSeq, Nov 2016]

Product images:



Circular map for MR215244