

Product datasheet for RN217560

Slc4a7 (NM_001270860) Rat Untagged Clone

Product data:

Product Type: Expression Plasmids
Product Name: Slc4a7 (NM_001270860) Rat Untagged Clone
Tag: Tag Free
Symbol: Slc4a7
Synonyms: NBC3; NBCn1
Vector: pCMV6-Entry (PS100001)
E. coli Selection: Kanamycin (25 ug/mL)
Cell Selection: Neomycin
Fully Sequenced ORF: >RN217560 representing NM_001270860
 Red=Cloning site Blue=ORF Orange=Stop codon

TTTTGTAATACGACTCACTATAGGGCGGCCGGAATTCGTCGACTGGATCCGGTACCGAGGAGATCTGCC
 GCCGCGATCGCC

ATGGAGGCAGACGGGGCCGGCGAGCAGATGAGACCGCTACTCACGCGGGTCCCGATGAAGAAGCTGTTG
 TGGATCTTGGCAAACACTAGCTCAACCGTGAACACCAAGTTTAAAAAGAAGAATTAGAGAGTCATCGAGC
 TGTATATGTTGGTGTTCATGTACCGTTTAGTAAAGAGAGTCGTCGGCGTACAGGCATCGAGGGCACAAA
 CATCACACCAGGAGAAGAAAAGACAAAGACTCAGATAAGGAAGATGGACGGGAGTCTCCTCTTATGACA
 CGCCATCGCAGAGGGTGCAGTTCATCCTTGGGACTGAAGACGATGATGAGGAGCACATCCCCACGACCT
 CTTACGGAGATGGACGAGCTCTGCTACCGAGACGGGAAGAGTACGAGTGGAAAGAGACAGCCAGGTGG
 CTGAAGTTCGAAGAGGATGTTGAGGATGGTGGTGACCGATGGAGTAAACCGTATGTGGCCACTCTGTCTT
 TGCACAGTCTCTTTGAGCTGAGGAGTGTATCCTAAATGGAACAGTCATGCTGGATATGAGAGCAAGCAC
 TCTGGATGAAATAGCAGATATGGTGTAGACAACATGATAGCGTCTGGCCAGCTAGATGACTCCATAAGG
 GAGAATGTTGAGAAGCTCTTCTGAAGAGACATCATCAAAATGAGAAAAGGTTTACAAGTCGGATTC
 CCCTCGTTCGATCCTTTGCAGATATAGGGGAAGGCCTGTCAGCCTCCCGCATTCTTTCGCAACAGGTCT
 GTCTGCCTCAAACCTTTCTTGAGAGGAGAATCGCCTTTATCCCTTCTTCTCAGTCATCTTCTTCTTCT
 TCAAGAGCTGGCACCCCTGCAGGCTCAAGGTGTACCACCCAGTACCACCCCAAGAACAGTCCCTCCTT
 CCAGCCCCAGCTTAAGTCGCCTGACCTCCAGAAGTTCCCAACAGACTCAGCCTCAGGCCCAAGTACT
 GGTGTACCTGACAGGGATGATATCCAGAGTAGTAATTCATCCGCTGAGGAAGACATAGAAGCACTG
 AAAGGCCAAGAGCAGAAGAATGAGGAAAATACTGACTTCACTCCAGGGATTTTGGCTTCTCCACAGTCTG
 CTCCTGAAAACCTGGACAGTAGTAAAAGTGGTGAAGTAAAGGTAACGGAAGTGGAGGAAGCAGAGAAAA
 TAGTACTGTTGACTTTAGCAAGGTTGACATGAATTTTCATGAGGAAAATTCCTACAGGAGCTGAGGCATCC
 AATGTTTTGGTAGGAGAGGTGGATTTCTTGAGAGACCTATCATTGCATTTGTGAGACTGGCTCCTGCAG
 TTCTCCTCTCAGGTTGACTGAGGTCCCTGTGCCCACTAGGTTTTTGTCTGTTACTGGGCCAGCAGG
 AAAGGCTCCACAGTACCATGAAATTGGCAGATCCATAGCAACTCTAATGACAGATGAGATTTTTTCATGAT
 GTAGCTTATAAGCAAAGATCGAAATGACCTCTATCTGGAATTGATGAATTTTATGATCAAGTAACTG



TTCTTCCTCCAGGAGAGTGGGATCCTTCCATACGCATAGAGCCACCAAAAAGTGTTCCTTCTCAGGAGAA
 AAGGAAGATTCCTGCGTTTCCCAATGGATCTGCTCCAGTGTCTGCTGACCCTCCTAAGGAGGCTGATCAC
 CACGCTGGGCCTGAGCTGCAGAGGACTGGACGGCTTTTGGTGGTTTGATACTTGACATCAAAGGAAAAG
 CACTTTTTTCTTGAGTGACTTCAAGGATGCATTAAGTCTGCAGTGCCTGGCCTCGATTCTTTTCTATA
 CTGTGCCTGTATGTCTCCTGTAATCACTTTTGGAGGGCTGCTTGGAGAAGCTACAGAAGGCAGAATAAGT
 GCAATAGAGTCTCTTTTGGAGCATCATTAACTGGGATTGCCTATTCATTGTTTGGTGGCAACCTCTAA
 CAATACTGGGAGCACGGTCCAGTCTAGTGTGGAAAAATTTTAAATTCGTAGAGATTATCA
 CCTATCCTATCTATCATTAAAGAACCAGTATTGGTCTGTGGACTTCTTTCTGTGCATTGTGTGGTCGCA
 ACAGATGCCAGCAGCCTTGTGTGTACATTACTCGTTTACAGAAGAAGCTTTCGCCGACTCATTTGTA
 TCATCTTCATCTATGAAGCCTTGGAGAACTCTTCACTTAGGAGAAAATATATGCATTTAACATGCACAA
 CAACTTGGATGCATTGACCAGTTACACATGTGTATGTGCTGAGCCATCTAATCCTAGCAACGAACTGTA
 GAGCTGTGGGAGAGGAAGAACGTGACAGCAGCCAGTATTTCTGGGCAAACCTTACCCTGTCTGAGTGA
 AGACCTCCACGGTATGTTTGTGGGATCAGCTTGTGGCCTCACGGCCTTATGTTCCCGATGTGCTCTT
 CTGGTGTGTCGTCTGTTTTTACAACGTTCTTCTGTCTTCATTCTCAAGCAGTTAAGACCAAGAGA
 TATTTTCTACTAAGGTGCGATCAACAATCAGTGACTTTGCTGTATTTCTACAATAGTAATAATGGTTG
 CAATTGACTACCTGTAGGAATTCATCTCCTAAACTTCATGTTCTGAAAAGTTTGGAGCCTACTGATCC
 AAGCAGGGGCTGGATCATAAGCCCTTGGGAGATAACCCTTGGTGGACCTTACTAATTGCAGCTGTTCCA
 GCTCTCCTTTGTACCATCTCATCTTATGACCAGCAGATCACAGCTGTGATCATCAACAGGAAAGAAC
 ACAAACTGAAGAAAGGAGCTGGCTATCACCTGGACTTGTCTATGGTTGCTGTGATGTTGGGAGTCTGCTC
 CATCATGGGCCTGCCGTGGTTTGTGGTGAACAGTGTGTTCTATAAGTCATGTCAACAGCTTAAAAGTA
 GAGTCTGAATGTTCTGCTCCAGGGGAACAGCCCAAGTTCCTCGGAATTCGTGAACAGCGGTTACAGGGC
 TGATGATTTTTATCCTGATGGCCTCTCTGTGTTCACTGACTTCACTATTAAGTTTATTCGGATGCCAGT
 TCTATATGGTGTTCCTTTATATGGGAGTTTCTTCTGAAAGGAATTCAGTTTTTTGACCGTATCAAA
 TTATTTGGAATGCCTGCCAAGCACCAGCCGACCTGATCTACCTCGCTATGTCCCTCTCTGGAAGGTGC
 ACGTGTTCACGGTCGTCCAGCTGACCTGCCTGGTCTGCTCTGGGTGATCAAAGCCTCTGCTGCTGCAGT
 AGTTTTTCCCATGATGGTCTTGCATTAGTCTTTGTGCGCAAGCTCATGGATCTGTGCTTCAAAAGAGA
 GAACTCAGTTGGCTTGTGACCTCATGCCAGAAAGTAAGAAAAAGAAAGATGACAAGAAGAAGG
 AGAAGGAGGAAGCTGAGCGGATGCTTCAAGGTGACGGGGATACTGTGCACCTCCCATTGAAAGGGGAG
 TCTCTACAAATCCAGTTAAGACCCTAAAATATAGTATTGACCCTTCACTGTTTAAACATATCAGATGAA
 ATGGCCAAAACCTGCCAGTGAAGGCACCTTCCATGAATACTGAGAATGCCAAAGTAACCAGACCTAACA
 TGAGCCCTGAAAAGCCTGTGAGTGTGACAATAAATTCGAAGATGAACCATCAAAAAATACATGGATGC
 TGAACCTCATTGTAA

ACGCGTACGCGGCCGCTCGAGCAGAAACTCATCTCAGAAGAGGATCTGGCAGCAAATGATATCCTGGATT
 ACAAGGATGACGACGATAAGGTTTAA

Restriction Sites:

Sgfl-Mlul

ACCN:

NM_001270860

Insert Size:

3726 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:

Clone contains native stop codon, and expresses the complete ORF without any c-terminal tag.

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_001270860.1](#), [NP_001257789.1](#)

RefSeq Size: 4002 bp

RefSeq ORF: 3726 bp

Locus ID: 117955

Cytogenetics: 15p16

Gene Summary: regulates intracellular pH in different cells along with Na⁺-driven HCO₃⁻ transporters, the Na⁺-driven Cl⁻/HCO₃⁻ exchanger [RGD, Feb 2006]
Transcript Variant: This variant (2) uses an alternate in-frame splice junction at the 3' end of an exon compared to variant 1. The resulting isoform (2) has the same N- and C-termini but is shorter compared to 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.