

Product datasheet for **MC229404**

Slc4a10 (NM_001242378) Mouse Untagged Clone

Product data:

Product Type: Expression Plasmids
Product Name: Slc4a10 (NM_001242378) Mouse Untagged Clone
Tag: Tag Free
Symbol: Slc4a10
Synonyms: mKIAA4136; NCBE
Vector: pCMV6-Entry (PS100001)
E. coli Selection: Kanamycin (25 ug/mL)
Cell Selection: Neomycin
Fully Sequenced ORF: >MC229404 representing NM_001242378
 Red=Cloning site Blue=ORF Orange=Stop codon

TTTTGTAATACGACTCACTATAGGGCGCCGGGAATTCGTCGACTGGATCCGGTACCGAGGAGATCTGCC
 GCC**CGATCGCC**

ATGGAGATTAAGACCAGGGAGCCCAAATGGAGCCGCTGCTGCCTACGAGAAATGATGAAGAAGCCGTTG
 TGGATAGAGGTGGAACACGCTCTATTCTCAAACACATTTTGAGAAAGAAGATTTAGAAGGTCATCGGAC
 ATTATTTATTGGAGTTCATGTGCCCTGGGTGGAAGAAAAGCCATCGTCGTCACAGGCATCGTGGTCAT
 AAGCACAGAAAGAGGGACAGAGAGAGAGATTCCGGACTGGAGGATGGAAGAGAGTCCCCTCTTTTGACA
 CCCCATCGCAGAGGGTGCAGTTTATTCTTGGAAGTGGAGACGATGATGAGGAGCACCTCCCTCATGACCT
 TTTCACAGAGCTGGATGAGATTTGCTGGCGTGAAGGGGAAGATGCTGAGTGGCGAGAGACAGCCAGGTGG
 TTGAAATTTGAAGAGGATGTGGAAGATGGAGGAGAAAGATGGAGTAAGCCCTATGTGGCCACGCTTTCAT
 TACACAGCTTGTTTGAGTTGAGAAGCTGCATCCTGAATGGAAGTGTGCTACTGGACATGCATGCCAACAC
 GATAGAAGAAATGCAGATATGGTCCTTGACCAGCAGGTCAGCTCAGGCCAGCTGAATGAAGATGTTCCG
 CACAGGGTCCACGAAGCATTGATGAAGCAGCATCATCACCAGAATCAGAAAAAAGTGGCTAACAGGATTC
 CTATTGTCCGATCTTTGCTGATATTGGCAAGAAACAATCAGAACAAATTCATGGATAAAAAAGTGCAGC
 TCAGGTTGTTCTCCTCAGTCTGCTCCAGCCTGTGCTGAGAATAAAAAATGATGTCAGCAGGGAAAAACAGC
 ACTGTAGACTTCAGCAAGGGACTGGGAGCCAAACAAAAGGGGCATACTAGTCCATGTGGGATGAAACAAA
 GGCTTGACAAAGGACCTCCACACCAGCAGGAGAGAGAGTTGATCTGCATTTTATGAAAAAGATTCTCC
 GGGTGCTGAAGCTTCAAACATCTTGGTAGGAGAACTGGAGTTCCTAGACAGAACTGTGGTTGCCCTTGTG
 AGGTTGTCTCCAGCTGTCTTCTCAAGGACTTGCTGAAGTTCCAATCCCAAGCAGATTTCTGTTTCATCC
 TTCTGGGACCCCTGGGAAAGGTC AACAGTACCACGAGATTGGCAGATCGATTGCGACCTTAATGACTGA
 TGAGGTGTTTCATGATGTTGCTTACAAAGCTAAAGACCGCAATGACTTGGTATCAGGAATTGATGAGTTT
 CTGGATCAGGTTACCGTTCTTCTCCTGGAGAATGGGATCCAAGCATAACGAATAGAACCTCCAAAAATG
 TCCTTCCAGGAGAAGAGGAAGATTCTGCTGTACCAAATGGAACAGCAGCTCATGGCGAAGCTGAGCC
 ACATGGAGGACACAGCGGACCTGAACTCCAGCGAACTGGGAGGATTTTGGGGGACTTATATTAGATATC
 AAAAGAAAGGCTCCATTCTTCTGGAGTGACTTCAGGGATGCTTTCAGCCTGCAGTCTTAGCATCGTTCC



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TGTTTCTCTACTGTGCATGCATGTCTCCTGTCATCACATTTGGAGGACTGTTGGGAGAAGCAACTGAAGG
 TCGTATAAGTGAATCGAATCACTCTTTGGAGCATCTATGACCGGGATAGCCTATTCTCTTTTTGGTGGA
 CAGCCCCGACCATATTAGGCAGCACAGGACCTGTTTTGGTGTTGAAAAGATCTTGTTAAAGTTTTGCA
 AGGAATACGGCCTGTCGTA CTACTTGTCTTACGGGCCAGCATTGGGCTCTGGACTGCAACACTGTGCATCAT
 CCTTGTGGCCACGGACGCGAGCTCACTCGTCTGTACATCACCCGGTTTACCGAAGAGGCTTTTGCTTCT
 CTCATTTGCATCATTTTTATCTATGAAGCCCTGGAGAAGTTGTTTGAGCTCAGTAAAACCTATCCAATCA
 ATATGCACAATGATTTGGAAGTGTGACACAATACTCATGTAACCTGTATGGAGCCACATAGTCCCAGCAA
 TGACACACTGAAGGAATGGCGGGAGTCCAACCTTTCTGCCTCTGACATAA TCTGGGGGAACCTA ACTGTG
 TCAGAGTGCAGATCACTGCACGGGGAGTATGTCGGGCGAGCCTGTGGCCATGGCCACCCCTACGTGCCAG
 ATGTTCTCTTCTGGTGGTGTATCCTGTTCTTCTCCACAGTTACCATGTCAGCCACCCTGAAGCAGTTCAA
 GACCAGCCGCTATTTCCCAACCAAGTTTCGATCCATAGTGAGTGATTTTGGCGTTTTTCTTACAATTCTG
 TGTATGGTTTTAATTGACTATGCCATTGGGATCCCATCACAAAACCTACAAGTACCAAGCGTTTTCAAGC
 CGACCAGAGACGACCGTGGCTGGTTTGTACACCTTTGGGTCAAACCCATGGTGGACAATCATAGCTGC
 CATCATCCAGCTTTACTCTGTACTATTCTGATTTTCATGGACCAGCAGATTACAGCTGTCATCATCAAC
 AGAAAAGAGCACAAAGCTAAAGAAAGTTGTGGCTATCACCTGGATCTGTTAATGGTGGCAGTCATGCTCG
 GGGTCTGCTCCATTATGGGCCTGCCATGGTTTGTGGCTGCCACAGTTCTCTCCATCACTCATGTCAACAG
 CCTCAAGCTCGAATCAGAGTGCTCTGCTCCAGGAGAACAACCCAAGTTTCTCGGCATTGGGAGCAGAGG
 GTGACCGGGCTCATGATTTTTATTCTTATGGGTTTCATCCGTTTTTCATGACCAGCATTCTGAAGTTTATCC
 CCATGCCAGTGTATACGGAGTGTCTTTTATATGGGTGCTTCGCTCTCAAAGGAATTCAGTTATTTGA
 TAGAATAAAGCTCTTCTGGATGCCAGCCAAACATCAACCAGATTTTCATCTATCTAAGGCACGTGCCCTC
 CGGAAAGTCCATCTTTCACAGTCATTCAGATGAGTTGTCTCGGCCTTCTGTGGATAATCAAAGTTTGA
 GAGCTGCTATTGTCTTCTATGATGGTGTGGCACTAGTGTGTTGAGAAAAGTTGATGGACTTCTTGT
 TACCAAACGGGAACTCAGCTGGCTTGTGATTTAATGCCTGAGAGTAAAAGAAGAACTTGAAGATGCT
 GAGAAAGAAGAAGAACAAAGTATGCTAGCCATGGAGGACGAGGGCACAGTACAACCTCCCACTGGAGGGAC
 ACTACAGAGACGACCCGCTGTGATCAATATTTCTGATGAAATGTCAAAGACTGCCATGTGGGGAACT
 TCTAGTTACTGCTGACAACTCAAAGAAAAGGAGTCACGCTTTCCTTCTAAAAGCAATGAAAGCCGAAAG
 GAGAAGAAAGCTGACTCAGGAAAGCGTTGACAGGGAGACTTGTCTATGA

ACGCGTACGCGGCCGCTCGAGCAGAACTCATCTCAGAAGAGGATCTGGCAGCAAATGATATCCTGGATT
 ACAAGGATGACGACGATAAGGTTTAA

- Restriction Sites:** SgfI-MluI
- ACCN:** NM_001242378
- Insert Size:** 3411 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_001242378.1](#), [NP_001229307.1](#)

RefSeq Size: 5514 bp

RefSeq ORF: 3411 bp

Locus ID: 94229

UniProt ID: [Q5DTL9](#)

Cytogenetics: 2 C1.3

Gene Summary: Sodium/bicarbonate cotransporter which plays an important role in regulating intracellular pH (PubMed:10993873, PubMed:20566632). Has been shown to act as a sodium/bicarbonate cotransporter in exchange for intracellular chloride (PubMed:10993873, PubMed:20566632). Has also been shown to act as a sodium/bicarbonate cotransporter which is not responsible for net efflux of chloride, with the observed chloride efflux being due to chloride self-exchange (By similarity). Controls neuronal pH and may contribute to the secretion of cerebrospinal fluid (PubMed:18165320). Reduces the excitability of CA1 pyramidal neurons and modulates short-term synaptic plasticity (PubMed:26136660). Required in retinal cells to maintain normal pH which is necessary for normal vision (PubMed:23056253). In the kidney, likely to mediate bicarbonate reclamation in the apical membrane of the proximal tubules (By similarity).[UniProtKB/Swiss-Prot Function]

Transcript Variant: This variant (1) encodes the longest isoform (1). Sequence Note: This RefSeq record was created from transcript and genomic sequence data to make the sequence consistent with the reference genome assembly. The genomic coordinates used for the transcript record were based on transcript alignments.