

Product datasheet for **MC219928**

Bcas1 (NM_029815) Mouse Untagged Clone

Product data:

Product Type:	Expression Plasmids
Product Name:	Bcas1 (NM_029815) Mouse Untagged Clone
Tag:	Tag Free
Symbol:	Bcas1
Synonyms:	2210416M21Rik; 9030223A09Rik; AI841227; NABC1
Vector:	pCMV6-Entry (PS100001)
E. coli Selection:	Kanamycin (25 ug/mL)
Cell Selection:	Neomycin



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Fully Sequenced ORF: >MC219928 representing NM_029815
 Red=Cloning site Blue=ORF Orange=Stop codon

TTTTGTAATACGACTCACTATAGGGCGGCCGGAATTCGTCGACTGGATCCGGTACCGAGGAGATCTGCC
 GCC**CGGATCGCC**

ATGGGAAACCAGATGAGCGTCCCACTAAGCGCTGGAGACCAGGAGCAGCACCCAGGAGCAGACACTTGCA
 AGGTGACGTGAGACAATGAGTGTGTTTCAAGTGGGAACCCAGTTGTGCTGTCCACTCGTGTAAATCAACA
 CTATGAGGAAGTCGACTTGGGAATCAGCAGCTCCAAGGATAATGTGGCTACTTCTCCCCAAGACAATG
 GAGGCCAGGCTGTGGCGATGCCAGCGAAAGAATCTTGGAAAGAAGCCAAGACCAAGGCACCGAGCTG
 CTAGATCCATTTTTTCTTGACTCTCTCGGCCTGTACCAGGACGTCCCGGGACCAAGGCACGGATTC
 ATCGGCTGCATCAGGGAGGTTTGTATGTCAGCCCAAGCGCAGCGCCTGAGAACAAGACCCAAGTGAGCAC
 GGGGCACTTCCGGTGGCAGCTGCACCAGGCAGGCTCCAGATAAAACCCAGGGTGCCCGAGGCCAAGC
 AGCAGACCCTCCCTGCCACCGGCCATTGGCACCTCACCACCTGAGTCCAGGCAGAAGCCCCGGCCCA
 GGACAAGGATTTGGCTTTTTGAACAGATTCTTTAAACTGGACAAGGAAGAGAAAGCGCCGGTGAAC
 AGCCAGCCCAAAGAAGCGAAAGGCTCGGAAGACCCAGAACAGGCCACAGAGGCTCCTGCCGTGCCAGGGA
 ATCCCCATGGTGTCTCTGCAGGGGAGGACATAGTCGACAGCGAGCAGAGAGGACAAGACGTTGACTCT
 GAGTTATTCTGTCCCTGGGGATCCAGAGGTGCCGGGACCAAGGAGGACCCGACGTTGGTGCATACC
 ACAGAGAACAGCAGCTCCATCATGAGCTTCTTCAAGACTGGTTTACCTAACAAAGACTGAAACAAAGA
 AGGACCCAGAAGACAGCAAGGCAACCAAGGCAGACAGTGTCTGTGATGGACACGCTGTGGCCAGAAGAT
 GTCGGAGACGCAGGCTAAAAGCAAGAAGAAGCGCCTGGACAGCCCCAGGCTAGGACTCTCCTTTAGAAAG
 CTCTTTAGGCACAAGGATACTGAGAATCACCACGACTTCAGCCAATCTCAAGTCGGACAAGCCAAT
 TCACACCCAGGAGACCCGAGGGAAGACAAGGCTACCAAAAGCTGCAGCCCCCACCACCTCCTGA
 ACCCACAGCGAAGGGAGAGACAGCGGCAAGGAGAAGGCAGGGCCACCTCACTACCGCTGGAAAGTTG
 TTTTGGAAAGAAGTCAGTTAAGGAGGATACACTTTCCACAGGTGCAGAGGAGAACGCGGTGTGTGAGTCA
 CAGTAGAGACCGTAAGGCTTGGGAAGTAGAATCCAGCTTACAACTGTGGATCTCAGTGAAGAGACCCA
 GCCTGAACCCACAGACGTAAAAGTCAAAGAAGAAAGCAAAACCCGGAAGACCCCTCTGATGGCGTTTCTC
 AGACAAATGTCAGTGAGATCGAGCGAAGGGATCCCCCGCTCGGAAGAAAGTAACGTGAAAGACTCCAGT
 GCCAAACGTCAAACCTCCGTGGAAAAGACGCCCTACCCCCAGAGCCAGAGCCCGGGAACAGCTCAGAA
 AAACAAGGAGACCTCCTCTCGAAGGACAAGAAATCAGTGGACAAGAAGTCAGCGACTGAGAACAGCAAG
 CAGAAGAACGGCAACAGGAAGTCAGAGAGCCGGCGCGTGTGTGAGCCGCCACAGTGGAGGCAACG
 CAATGCAGACTGGGGACAAGACCCCAAAGAAGTCTGAGAAGCGGCAGAGTCCCTCGGGGCTTCTCGAA
 GGGCTGGGACCAAGCGGATGTCGGATGCTCAGGTGCAAAACGGACCCGGTCTCCATCGGACCAGTTGGA
 AAATCCAAG**TAA**

ACGCGTACGCGGCCGCTCGAGCAGAAACTCATCTCAGAAGAGGATCTGGCAGCAAATGATATCCTGGATT
 ACAAGGATGACGACGATAAGGTTTAA

- Restriction Sites:** SgfI-MluI
- ACCN:** NM_029815
- Insert Size:** 1902 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).
- 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_029815.2](#), [NP_084091.2](#)

RefSeq Size: 3090 bp

RefSeq ORF: 1902 bp

Locus ID: 76960

UniProt ID: [Q80YN3](#)

Cytogenetics: 2 H3

Gene Summary: Required for myelination.[UniProtKB/Swiss-Prot Function]

Transcript Variant: This variant (1) represents the longer transcript and encodes the longer 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.