

Product datasheet for **MC205705**

Bcap29 (NM_007530) Mouse Untagged Clone

Product data:

| | |
|---------------------------|---|
| Product Type: | Expression Plasmids |
| Product Name: | Bcap29 (NM_007530) Mouse Untagged Clone |
| Tag: | Tag Free |
| Symbol: | Bcap29 |
| Synonyms: | AW208404; Bap29 |
| Mammalian Cell Selection: | Neomycin |
| Vector: | PCMV6-Kan/Neo (PCMV6KN) |
| E. coli Selection: | Kanamycin (25 ug/mL) |



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Fully Sequenced ORF: >BC021661
 CCACGCGTCCGTGACGGGCGGGCGGGCCCGCTGCTGACCTCGCGTCCGGCGGCTCGGGCCGGGCGT
 TCTGCGGCCCAGCCACCTCGGGACCGGGGAAGAACCTGCGGGGCGTCCGGCGCACTGCCTCGTACCC
 GGAGTTAGGCGAACTCCCGGCAGGCCACGCCGCGTGTCTGGCCGTTACCCCATGTTTCTCTAAAATTG
 AAATCTTCTCTGTATAGGTGTGAAGAAAAACAAAACAAACGATGACAATCCAATGGGCGGCGTTGC
 CAGCTTTCTTTATGCTGAAATAGGACTCATTTACTCTTCTGTCTGCCCTTATTCTCCACAGAGATGG
 CAGAAGATTTTTTCATTTAGTGTCTGGGTAAGATTGCAAGTTTTTGAACAAAGCTTTTCTTACCATTA
 TAATACTATTGATCATTCTGTTTCTAGATGCCGTGAGAGAAGTAAGAAAGTATTCTGCCAATGTTGT
 AGAGAAGAACTCAGCCATCAGACCGAGTGCCTTTGAGCACACACAGATGAAGCTCTTCAGGTCTCAAAGA
 AATCTGTATATTTCTGGATTCTCATTATTTTTTGGCTAGTGTGAGACGTCTGGTTACGCTTACTC
 AGCTGGCAAAAGAAATAGCAAACAAAGGAGTGCTTAAATTCAGCAGAAAATACCAATAAGGCTGCCAA
 GAAATTCATGGAAGAGAATGAAAACTAAAATTGGGGCTAAGAAATGACAATGCTGAAGAACACCTTTG
 GAAGCTGAAAATAAAAACTAATAGAAAGCAAGGAGAACCTGAAAACCGAGCTGAAGAAGGCTCCGATG
 CCCTTTTGAAGGCCAAAATGATGTCATGACTATGAAGATTCAGTCAGAGAGACTTTCTAAAGATATGA
 CCGACTCTGAAAGAACACTCAGAACTCAGAACCGTTTAGAGAAGAGAAGAAGAAAGGCTGTGAACT
 AACTTTATAAAAGAAGACTGTAATACACCTTGCAAAAGTTCAGTTGAAAACAAAGAAAGCTGTGCTAGCG
 CCTGCAGACCCGCACAGGCTATACAGTAGTGACATTTCCAAAATATTTGATAATGTTTCAGATTTAGAT
 ATAAAGTATGGATTTAGCTCTTTTGGAAAATATAAGCATGTTAAATACCATATTTACATATTGCTAATA
 GCATTGATATATGGTGGCTCTTTACCCATAAAAGGAAATTTTCATTTGCTAGCATTTACTTCCCAAATTA
 GATCTGAAGTTGCATGAAACCATTATTAGCTTTGAAAGCTTTGATGTGTTTTAGTAATAATCAGATGAT
 GGTAGTGGTAAAAATGATGTGTTGAAAAGAGAAGTCAAGTATCAGAAAATGTTTTACAGGGAACAAGG
 ATGAGTAACCACAATTAGCCACTGCGCTTATCGAGAAGAGTCAGAGAACAGCGAAAACACTACGGCCTTAGG
 GCTCTTCAAGCCTCATGATTGTACAACCTACTCGAAATATGACAGCACTGTTGATGTGTGTTGCTAC
 TGTGCTTCTGTGCCTATAAGACTATTGATAATTTGTTTGTAGTCATATATGGTCAGTAGTCTGCTTTTCAA
 GGTGATCTGCTAATAGAATGTATGCCATTTGCAAACTGTCATACCACTGAACTGAGCTTATGTTGCAC
 TGTGGATTTCTCCACTGACAAAGCTTATTTACGACATCCTAAAAGCAGCGTTTTTTCTATTTGTGAAAGT
 TACTGATTGACTGGTATTACCTCAGTATTTCATGCACTGGTATCATAACAGATCTCTACTATGTTTCTA
 TATTCTCAAAGTTAAATACAGGTTTTTCATGCAAAATTCATGTAGAAAGTAGTGAGTGAATAATATGATT
 TAGGTTATTTAGTGTTTAACTGTGAAAAGCCACTGTCTTAAATGCTTAAATATAAAAGAAAATGCAA
 AATAAAAAAAAAAAAAAAAAAAG

Restriction Sites: RsrII-NotI

ACCN: NM_007530

Insert Size: 723 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: [BC021661](#), [AAH21661](#)

RefSeq Size: 1983 bp

RefSeq ORF: 723 bp

Locus ID: 12033

UniProt ID: [Q61334](#)

Cytogenetics: 12 13.63 cM

Gene Summary: May play a role in anterograde transport of membrane proteins from the endoplasmic reticulum to the Golgi. May be involved in CASP8-mediated apoptosis (By similarity). [UniProtKB/Swiss-Prot Function]
Transcript Variant: This variant (1) represents the longer transcript. Variants 1 and 2 both encode the same protein. 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.