

## Product datasheet for **MC201469**

### **Bfar (NM\_025976) Mouse Untagged Clone**

#### **Product data:**

Product Type:	Expression Plasmids
Product Name:	Bfar (NM_025976) Mouse Untagged Clone
Tag:	Tag Free
Symbol:	Bfar
Synonyms:	3010001A07Rik; 3110001I22Rik; AI666707; AW107665; Bar; Rnf47
Mammalian Cell Selection:	Neomycin
Vector:	PCMV6-Kan/Neo (PCMV6KN)
E. coli Selection:	Kanamycin (25 ug/mL)



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**Fully Sequenced ORF:** >BC027221 sequence for NM\_025976  
TCTACTCCGCGGGTCTTGGCAGGTGGCGGTGACAGAGTGGGTACCTGGACTAGTGGCGAGACTTTATG  
ATGCATTGCAGCAGTGTCTGAGTCTGAAAGTGTCTATGTGTATGGATCCCAGAAATTTGCTGAGAGATGG  
AGGAGCCTCAGAAAAATGACCTGAGCATGAGAGAGCAGGAGGAAGAGCATCCTGTGAGAAGCAGCGGCC  
TCAGATTTCTGTGAGCGAGTTCTCTGCCACTGCTGCTACGACACCCTGGTTAACCCACACCTTGAAC  
TGTGGCCACAGCTTCTGCCGCACTGCCTGGCTTTATGGTGGATGTCTTCAAAGAAGACAGAGTGTCCAG  
AATGCAGAGAAAAATGGGAAGGTTTTCTAAAGTCAACATTCTCCTCAGGGATGCCATTGAAAAGTTATT  
TCCTGATGCCATTAGAATGCGAGTTGAAGACATTGAGCAGAATAATGATGTAGTCCAAAGTCTTGACAGCC  
TTTCAGAAATATGGGAATGATCAGAATCCCTTAGCTCCCAGCACAGGGGAGTAAATCCTCAGAGAGGAG  
GGGGATTCTTCTCGGTGTTCTCACAGCTTAACTGGTGTGGCAGTCATCCTGCTTGTGTATCATTGGCG  
CAGCAGAGAATCTGAACATGGCCTCCTGGTGCACAAGGCTGTAGATAAGTGGACGATGGAAGAAGTTGTC  
CTCTGGTTAGAACAGCTAGGACCTTGGGCTCCCTGTACAGAGACAGGTTCTTATCTGAAAGAGTAAATG  
GAAGGTTGCTTTAACTTTGACAGAAGAAGAGTTTTCCAGGGCACCTTATACCATAGAGAACAGTAGCCA  
CAGAAGGGTCATCCTCACGGAGCTGGAGCGTGTGAGAGCCCTGGGAGTGAAGCCACCGCAGAATCTCTGG  
GAGTACAAGGCTGTCAACCCGGGTAGATCCCTATTCTTGTCTACGCTCTCAAGAGCTCACCCAGACTTG  
GCTTGCTATACCTGTACCTGTTTACTACACAGACTGCTTCTACCTTATCCACACCATCTGCCCTCT  
GCAGGAAAACAGCTCTGGCGAGGACATTTTACCAGCTTCTGGACCTGAGGGAGCCACGTTGGAAGCAG  
TGGAGAGAATTCCTCGTCAAGTACTCCTTTCTTCCGTACCAGCTGATCGCGGAGTTTGCCTGGGACTGGC  
TGGAGGTCACACTACTGGACATCTCGCTTCTGATCGTCAATGCCGTGCTGCTCTCTGTTCTGGAGTTGTT  
CTCTTTTTGGAGGATCTGGTTCGAGAAGTGAAGTGAAGACAGTGCACAGAGGATGTGGAGCCACTTCTGG  
AAAGTATCGACACAGGGGCTTTTTCATGGCCATGTTCTGGCCGCTTATTCTCAGTTTGTCTGCAACTGTT  
TATTTTACTGGGCTCTGACTTCAACCAATTATTAACATTGATCTGGTGGTCAAGGAAGTCCGACGGCT  
GGAAACCAAGTGTATGATGGGCAGCAAGGCCGAGCGAGCCGTTGAATCTACCTGACAACCAAGCTGT  
GACATTCAGAGACAAGGGACTCGTTTTCTACCCCTGGTAAATGCGAGGTGCTGCTTCTGTGTCAAGGC  
TTCAACAGGTCCTTCTCTGACTCTGGCACCATGCCAGCTGACTGCTACCCAGGTCAGATTGGACT  
CATGGTCAATACCCAGCCAGTCTTTTTACAGGGACTTGAGAGGCTCAGGCTACTATGACCAATAGAAGTG  
AAAGTCTTGGTCATAATTTGGGGTACATGCTGTCAAAGGCTAATATAGACTGGGTGTTGAGTGCACA  
CCTTGAATCTGAGCACTAGGAAAACAGGCAGGTGGATTTTTGAGTTCAAGACCAGAGTGGTCTACAGAGT  
GAGTTCTACAACAGCCAGGGCTACATAGAGGAACCATGTCTCTAAAAAAAAAAAAAAAAAGAAAGAAAGAAA  
AAGAAAAGAAAAAAAAAAAAAAAA

**Restriction Sites:** RsrII-NotI

**ACCN:** NM\_025976

**Insert Size:** 1353 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: [BC027221](#), [AAH27221](#)

RefSeq Size: 1983 bp

RefSeq ORF: 1353 bp

Locus ID: 67118

UniProt ID: [Q8R079](#)

Cytogenetics: 16 A1

**Gene Summary:** Apoptosis regulator. Has anti-apoptotic activity, both for apoptosis triggered via death-receptors and via mitochondrial factors (By similarity).[UniProtKB/Swiss-Prot Function]  
Transcript Variant: This variant (1) represents the longest transcript and 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.