

## Product datasheet for **SC321746**

### **RIMKLB (NM\_020734) Human Untagged Clone**

#### **Product data:**

Product Type:	Expression Plasmids
Product Name:	RIMKLB (NM_020734) Human Untagged Clone
Tag:	Tag Free
Symbol:	RIMKLB
Synonyms:	FAM80B; NAAGS; NAAGS-I
Mammalian Cell Selection:	Neomycin
Vector:	pCMV6-AC (PS100020)
E. coli Selection:	Ampicillin (100 ug/mL)



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**Fully Sequenced ORF:** >OriGene sequence for NM\_020734.1  
 CCACGCGTCCGATGGATTTCGTGGGCTGCTTCCACCTGCTAGGAGGGTGGTGTACTCTAAC  
 TCAGGGACAGAAGCCCCTGTCTGTGCTCAGGACTCTTGCAGACCTCTTTACCTGGCTGTT  
 CATCTTCATAATCAACTGGTAGACGTTACATCCAAGAGGAAATAATCCAGGCAAGGAAG  
 CACAAGCTGATCAAGATGTGTAGTTCTGTGGCTGCCAAGTTGTGGTTTTTGACAGATCGT  
 CGCATCAGGGAAGACTATCCTCAAAAAGAGATTTTACGAGCATTGAAGGCCAAATGTTGT  
 GAGGAGAACTGGACTTTAGGGCTGTGGTATGGATGAGGTGGTCTGACAATCGAGCAA  
 GGAAACCTGGGTCTGCGGATCAATGGAGAGCTAATCACTGCCATCCACACAAGTGGTGTA  
 GTCAGAGTACCAACCCCTTGGGTGCAAAGTGATAGTGACATCACTGTTTTGCGCCATCTA  
 GAGAAGATGGGATGTCGGTTAATGAACCGACCTCAAGCCATCCTGAACTGCGTTAATAAG  
 TTCTGGACATTTCAAGAGTTGGCTGGCCATGGTGTTCCTCTGCCGGATACTTTCTCTTAT  
 GGTGGCCACGAAAATTTGCTAAAATGATTGATGAGGCTGAAGTTCTGGAGTTCCCAATG  
 GTAGTAAAGAATACGCGGGTACAGAGGTAAGCTGTTTTCTGGCTCGAGATAAGCAC  
 CATTTGGCTGATCTAAGCCATCTTATTCGCCATGAAGCGCCATACCTGTTCCAGAAGTAT  
 GTTAAAGAGTCTCATGGACGGGATGTACGTGTCATTGTCGTGGGAGGCCGTGGTTGGC  
 ACCATGTTACGTTGTTCAACAGATGGGAGAATGCAAAGCAACTGCTCATTAGGTGGTGTG  
 GGGATGATGTGCTCATTGAGTGAACAAGGGAAGCAGCTAGCTATCCAGGTGTCTAATATC  
 CTGGGGATGGATGTGTGGCATTGACCTGCTGATGAAAGATGACGGCTCCTTCTGCGTC  
 TGTGAGGCCAATGCAAATCCTTTTCAGGAGCCAAAAACAAATACAAACAAACAAAAA  
 ATACCCAGAGAAAAAATTTCTTCTTCCCCCTTCTGATGATGAGTGAGAGTATTGAG  
 AACTTTCCGGGTCAGTGCCTTCTAAATCCCCTTCCCCCAATAATGCAGCTGTATAATG  
 AATGGTAAATGCACCGGTTTGGATTTCAGGCACAGCCCCAGTCTGCCTACAGCAGACAAT  
 AAATGGTAGGTGGCAGCAGAGGAAGGAATTTTCAGATTGATGGCTACTGACTTAAAGT  
 CTGCATAGGAGAAGAGATAGAGCATGACATACCAAGTGAAGCAAGGGGGACAAAAATGGA  
 AACAGGACTAAGATTCCTGCATTTTACTCTAAATGGTTCTTTTGAAGTAAACAGTGATC  
 TTTTTGCTTACCCTCATCAACAGAATGGATGAAGATAGATTTGACTGTGTGCTTTTTCA  
 AGTGGAGAATATGAATATACAGAACAAGAAAGCAATAACTCCAACCTGTTTGATTCCGTC  
 TGTTTTCTAAATAAAGACACTATGCGCTGGAAATAAAAAAAAAAAAAAAAA

**Restriction Sites:** Please inquire

**ACCN:** NM\_020734

**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:** This TrueClone is provided through our Custom Cloning Process that includes sub-cloning into OriGene's pCMV6 vector and full sequencing to provide a non-variant match to the expected reference without frameshifts, and is delivered as lyophilized plasmid DNA.

**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\\_020734.1](#), [NP\\_065785.1](#)

**RefSeq Size:** 1610 bp

**RefSeq ORF:** 924 bp

**Locus ID:** 57494

**UniProt ID:** [Q9ULI2](#)

**Cytogenetics:** 12p13.31

**Gene Summary:** Catalyzes the synthesis of beta-citryl-L-glutamate and N-acetyl-L-aspartyl-L-glutamate. Beta-citryl-L-glutamate is synthesized more efficiently than N-acetyl-L-aspartyl-L-glutamate. [UniProtKB/Swiss-Prot Function]

Transcript Variant: This variant (1) represents the longest transcript and encodes the functional protein. All of the protein-coding variants encode the same protein. 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.