

Product datasheet for **SC332687**

CLPB (NM_001258392) Human Untagged Clone

Product data:

Product Type:	Expression Plasmids
Product Name:	CLPB (NM_001258392) Human Untagged Clone
Tag:	Tag Free
Symbol:	CLPB
Synonyms:	ANKCLB; HSP78; MEGCANN; MGCA7; SKD3
Vector:	pCMV6-Entry (PS100001)



[View online »](#)

Fully Sequenced ORF: >SC332687 representing NM_001258392.
 Blue=Insert sequence Red=Cloning site Green=Tag(s)

```

ATGCTGGGGTCCCTGGTGTGAGGAGAAAAGCACTGGCGCCACGGCTACTCCTCCGGCTGCTCAGGTCC
CCAACGCTCCGGGGCCATGGAGGTGCTTCCGGCCGGAATGTGACTACTGGGAGTCTCGGGAGCCGCAG
TGGTGTAGGGTAGCCACCGGGGGGCGCCCTGGAACATCGCCGGCCTTGTCTCCGGACGTGGGGCAGCC
ACCGGGGGGCGCCAGGGAGGACGCTTCGATACCAAAATGCCTCGCGGCTGCCACTTGGGGACGCCTTCT
GGTCCCGAAGAAAACACTCCAGGACAGGACAGCTGGAACGGGGTCCCGAGCAGGGCCGGACTGGGCATG
TGCGCCCTGGCCGACGCGTGGTGGTTCATTGCTACAGCAAGAGTCCGTCACAAAGGATGCAGCCCTG
TTGGAAGCTGCCCGTCCCAACAATATGCAAGAAGTACAGAGGCTGTTGTGAGAAGGTGCAGATGTCAAT
GCAAAGCACAGACTTGGCTGGACAGCACTATGGTGGCAGCCATCAACCGAAAACAAGTGTGGTACAG
GTCCTGCTTGTGCTGGGGCTGATCCAAACCTTGGAGATGATTTACAGAGTGTTTACAAGACTGCCAAG
GAACAGGGAATCCATTCTTTGGAAGTCTGATCACCCGAGAGGATGACTCAACAACAGGCTGAACAAC
CGCGCCAGTTTCAAGGGCTGCACGGCTTGCATATGCTGTTCTTGTGATGACTACCGCACTGTCAAG
GAGCTGCTTGTGAGGAGCCAAACCCCTGCAGAGGAATGAAATGGGACACACCCCTTGGATTATGCC
CGAGAAGGGGAAGTGATGAAGCTTCTGAGGACTTCTGAAGCCAAGTACCAAGAGAAGCAGCGGAAGCGT
GAGGCTGAGGAGCGGGCGCGCTTCCCCCTGGAGCAGCGACTAAAGGAGCAGATCATTGGCCAGGAGAGC
GCCATCGCCACAGTGGGTGCTGCGATCCGGAGGAAGGAGAATGGCTGGTACGATGAAGAACCCTCTG
GTCTTCTCTTCTGGGATCATCTGGAATAGGAAAAACAGAGCTGGCCAAGCAGACAGCCAAATATATG
CACAAAGATGCTAAAAGGGCTTCATCAGGCTGGACATGTCCGAGTCCAGGAGCGACACGAGGTGGCC
AAGTTTATTGGGTCTCCACCAGGCTACGTTGGCCATGAGGAGGGTGGCCAGCTGACCAAGAAGTTGAAG
CAGTGGCCCAATGCTGTGGTGTCTTTGATGAAGTAGACAAGGCCATCCAGATGTGCTCACCATCATG
CTGCAGCTGTTTGTGAGGGCCGGCTGACAGATGGAAGGGAAGACCATTGATTGCAAGGACGCCATC
TTCATCATGACCTCCAATGTGGCCAGCGACGAGATCGCACAGCAGCGCTGCAGCTGAGGCAGGAAGCT
TTGGAGATGAGCCGTAACCGTATTGCCGAAAACCTGGGGATGTCCAGATAAGTGACAAGATCACCATC
TCAAAGAACTTCAAGGAGAATGTGATTCGCCCTATCCTGAAAGCTCACTCCGGAGGGATGAGTTTCTG
GGACGGATCAATGAGATCGTCTACTTCTCCCTTCTGCCACTCGGAGCTCATCAAACCTCGTCAACAAG
GAACTAACTTCTGGCCAAGAGAGCCAAAGCAAGGCACAACATCACGCTGCTCTGGACCGCGAGGTG
GCAGATGTGCTGGTGCAGCGCTACAATGTGCACTATGGCGCCGCTCCATCAAACATGAGGTAGAACGC
CGTGTGGTGAACCAGCTGGCAGCAGCTATGAGCAGGACCTGCTGCCAGGGGGCTGACTTTGCGCATC
ACGGTGGAGGACTCAGACAAGCAGCTACTCAAAGCCAGAACTGCCCTCACCCAGGCTGAGAAGCGC
CTCCCAAGCTGCGTCTGGAGATCATCGACAAGGACAGCAAGACTCGCAGACTGGACATCCGGGCACCA
CTGCACCTGAGAAGGTGTGCAACCATCTAG
  
```

Restriction Sites: Sgfl-Mlul

ACCN: NM_001258392

Insert Size: 2034 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_001258392.1](#)

RefSeq Size: 3182 bp

RefSeq ORF: 2034 bp

Locus ID: 81570

UniProt ID: [Q9H078](#)

Cytogenetics: 11q13.4

MW: 75.4 kDa

Gene Summary: This gene belongs to the ATP-ases associated with diverse cellular activities (AAA+) superfamily. Members of this superfamily form ring-shaped homo-hexamers and have highly conserved ATPase domains that are involved in various processes including DNA replication, protein degradation and reactivation of misfolded proteins. All members of this family hydrolyze ATP through their AAA+ domains and use the energy generated through ATP hydrolysis to exert mechanical force on their substrates. In addition to an AAA+ domain, the protein encoded by this gene contains a C-terminal D2 domain, which is characteristic of the AAA+ subfamily of Caseinolytic peptidases to which this protein belongs. It cooperates with Hsp70 in the disaggregation of protein aggregates. Allelic variants of this gene are associated with 3-methylglutaconic aciduria, which causes cataracts and neutropenia. Alternative splicing results in multiple transcript variants. [provided by RefSeq, Apr 2015]

Transcript Variant: This variant (2) lacks an in-frame exon in the 5' coding region compared to variant 1. The encoded isoform (2) is shorter than 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.