

## Product datasheet for **TR307338**

### CLPB Human shRNA Plasmid Kit (Locus ID 81570)

#### Product data:

Product Type:	shRNA Plasmids
Product Name:	CLPB Human shRNA Plasmid Kit (Locus ID 81570)
Locus ID:	81570
Synonyms:	ANKCLB; HSP78; MEGCANN; MGCA7; SKD3
Vector:	pRS (TR20003)
E. coli Selection:	Ampicillin
Mammalian Cell Selection:	Puromycin
Format:	Retroviral plasmids
Components:	CLPB - Human, 4 unique 29mer shRNA constructs in retroviral untagged vector(Gene ID = 81570). 5µg purified plasmid DNA per construct 29-mer scrambled shRNA cassette in pRS Vector, TR30012, included for free.
RefSeq:	<a href="#">NM_001258392</a> , <a href="#">NM_001258393</a> , <a href="#">NM_001258394</a> , <a href="#">NM_030813</a> , <a href="#">NM_030813.2</a> , <a href="#">NM_030813.3</a> , <a href="#">NM_030813.4</a> , <a href="#">NM_030813.5</a> , <a href="#">NM_001258394.1</a> , <a href="#">NM_001258394.2</a> , <a href="#">NM_001258393.1</a> , <a href="#">NM_001258393.2</a> , <a href="#">NM_001258392.1</a> , <a href="#">NM_001258392.2</a> , <a href="#">BC006404</a> , <a href="#">BC006404.2</a> , <a href="#">BC006257</a> , <a href="#">NM_001258392.3</a> , <a href="#">NM_030813.6</a> , <a href="#">NM_001258393.3</a>
UniProt ID:	<a href="#">Q9H078</a>
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]



[View online »](#)

- shRNA Design:** These shRNA constructs were designed against multiple splice variants at this gene locus. To be certain that your variant of interest is targeted, please contact [techsupport@origene.com](mailto:techsupport@origene.com). If you need a special design or shRNA sequence, please utilize our [custom shRNA service](#).
- Performance Guaranteed:** OriGene guarantees that the sequences in the shRNA expression cassettes are verified to correspond to the target gene with 100% identity. One of the four constructs at minimum are guaranteed to produce 70% or more gene expression knock-down provided a minimum transfection efficiency of 80% is achieved. Western Blot data is recommended over qPCR to evaluate the silencing effect of the shRNA constructs 72 hrs post transfection. To properly assess knockdown, the gene expression level from the included scramble control vector must be used in comparison with the target-specific shRNA transfected samples.
- For non-conforming shRNA, requests for replacement product must be made within ninety (90) days from the date of delivery of the shRNA kit. To arrange for a free replacement with newly designed constructs, please contact Technical Services at [techsupport@origene.com](mailto:techsupport@origene.com). Please provide your data indicating the transfection efficiency and measurement of gene expression knockdown compared to the scrambled shRNA control (Western Blot data preferred).