

# Product datasheet for KN209057BN

## PICALM Human Gene Knockout Kit (CRISPR)

### **Product data:**

#### OriGene Technologies, Inc.

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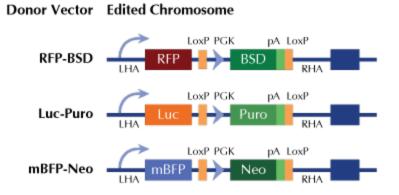
Product Type:	Knockout Kits (CRISPR)
Format:	2 gRNA vectors, 1 mBFP-Neo donor, 1 scramble control
Donor DNA:	mBFP-Neo
Symbol:	PICALM
Locus ID:	8301
Components:	<b>KN209057G1</b> , PICALM gRNA vector 1 in pCas-Guide CRISPR vector (GE100002) <b>KN209057G2</b> , PICALM gRNA vector 2 in pCas-Guide CRISPR vector (GE100002) <b>KN209057BND</b> , donor DNA containing left and right homologous arms and mBFP-Neo functional cassette. <b>GE100003</b> , scramble sequence in pCas-Guide vector
Disclaimer:	These products are manufactured and supplied by OriGene under license from ERS. The kit is designed based on the best knowledge of CRISPR technology. The system has been functionally validated for knocking-in the cassette downstream the native promoter. The efficiency of the knock-out varies due to the nature of the biology and the complexity of the experimental process.
RefSeq:	<u>NM 001008660, NM 001206946, NM 001206947, NM 007166</u>
UniProt ID:	<u>Q13492</u>
Synonyms:	CALM; CLTH; LAP
Summary:	This gene encodes a clathrin assembly protein, which recruits clathrin and adaptor protein complex 2 (AP2) to cell membranes at sites of coated-pit formation and clathrin-vesicle assembly. The protein may be required to determine the amount of membrane to be recycled, possibly by regulating the size of the clathrin cage. The protein is involved in AP2-dependent clathrin-mediated endocytosis at the neuromuscular junction. A chromosomal translocation t(10;11)(p13;q14) leading to the fusion of this gene and the MLLT10 gene is found in acute lymphoblastic leukemia, acute myeloid leukemia and malignant lymphomas. The polymorphisms of this gene are associated with the risk of Alzheimer disease. Multiple alternatively spliced transcript variants encoding different isoforms have been found for this



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gene. [provided by RefSeq, May 2011]

#### **Product images:**



RFP, Luc, and mBFP will be under native gene promoter

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