

## **Product datasheet for TL308485**

#### OriGene Technologies, Inc.

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### **UNC45B Human shRNA Plasmid Kit (Locus ID 146862)**

#### **Product data:**

**Product Type:** shRNA Plasmids

**Product Name:** UNC45B Human shRNA Plasmid Kit (Locus ID 146862)

**Locus ID:** 146862

Synonyms: CMYA4; CTRCT43; MFM11; SMUNC45; UNC-45B; UNC45

Vector: pGFP-C-shLenti (TR30023)

E. coli Selection: Chloramphenicol (34 ug/ml)

**Mammalian Cell** 

Puromycin

Selection: Format:

Lentiviral plasmids

Components: UNC45B - Human, 4 unique 29mer shRNA constructs in lentiviral GFP vector(Gene ID =

146862). 5µg purified plasmid DNA per construct

29-mer scrambled shRNA cassette in pGFP-C-shLenti Vector, TR30021, included for free.

**RefSeq:** NM 001033576, NM 001267052, NM 001308281, NM 173167, NM 001033576.1,

NM 173167.1, NM 173167.2, NM 173167.3, NM 001267052.1, BC101062, BC101063,

NM 001267052.2, NM 001033576.2

UniProt ID: Q8IWX7

**Summary:** This gene encodes a co-chaperone required for folding and accumulation of type II myosins.

The protein consists of three tetratricopeptide repeat motifs at the N-terminus that form a complex with heat shock protein 90, a central region of unknown function that is conserved in all Unc-45 proteins, and a C-terminal Unc-45/Cro1/She4 domain. The protein is expressed at high levels in striated muscle, where its muscle myosin chaperone activity is dependent on heat shock protein 90 acting as a co-chaperone. A missense mutation in this gene has been associated with cataract development. Alternative splicing results in multiple transcript

variants. [provided by RefSeq, Apr 2015]

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 <u>techsupport@origene.com</u>. 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. Please provide your data indicating the transfection efficiency and measurement of gene expression knockdown compared to the scrambled shRNA control (Western Blot data preferred).