

### **Product datasheet for TR313944**

## OriGene Technologies, Inc.

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#### **CHD3 Human shRNA Plasmid Kit (Locus ID 1107)**

#### **Product data:**

**Product Type:** shRNA Plasmids

**Product Name:** CHD3 Human shRNA Plasmid Kit (Locus ID 1107)

Locus ID: 1107

Synonyms: Mi-2a; Mi2-ALPHA; SNIBCPS; ZFH

**Vector:** pRS (TR20003)

E. coli Selection: Ampicillin

Mammalian Cell Puromycin

Selection:

Format: Retroviral plasmids

Components: CHD3 - Human, 4 unique 29mer shRNA constructs in retroviral untagged vector(Gene ID =

1107). 5µg purified plasmid DNA per construct

29-mer scrambled shRNA cassette in pRS Vector, TR30012, included for free.

RefSeq: NM 001005271, NM 001005273, NM 001272, NM 005852, NM 001005271.1,

NM 001005271.2, NM 001005273.1, NM 001005273.2, NM 005852.2, NM 005852.3, BC110648, BC156472, BM554923, NM 005852.4, NM 001005271.3, NM 001005273.3

UniProt ID: Q12873

**Summary:** This gene encodes a member of the CHD family of proteins which are characterized by the

presence of chromo (chromatin organization modifier) domains and SNF2-related

helicase/ATPase domains. This protein is one of the components of a histone deacetylase complex referred to as the Mi-2/NuRD complex which participates in the remodeling of chromatin by deacetylating histones. Chromatin remodeling is essential for many processes including transcription. Autoantibodies against this protein are found in a subset of patients with dermatomyositis. Three alternatively spliced transcripts encoding different isoforms

have been described. [provided by RefSeq, Jul 2008]

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).