

## **Product datasheet for TR312455**

## OriGene Technologies, Inc.

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## HIP2 (UBE2K) Human shRNA Plasmid Kit (Locus ID 3093)

**Product data:** 

**Product Type:** shRNA Plasmids

Product Name: HIP2 (UBE2K) Human shRNA Plasmid Kit (Locus ID 3093)

**Locus ID:** 3093

**Synonyms:** E2-25K; HIP2; HYPG; LIG; UBC1

**Vector:** pRS (TR20003)

E. coli Selection: Ampicillin

Mammalian Cell Puromycin

Selection:

i di omycin

Format: Retroviral plasmids

Components: UBE2K - Human, 4 unique 29mer shRNA constructs in retroviral untagged vector(Gene ID =

3093). 5µg purified plasmid DNA per construct

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

**RefSeq:** <u>BC006419</u>, <u>NM 0011111112</u>, <u>NM 0011111113</u>, <u>NM 001312646</u>, <u>NM 001312647</u>, <u>NM 001312648</u>,

NM 005339, NM 005339.1, NM 005339.2, NM 005339.3, NM 005339.4, NM 001111113.1, NM 001111112.1, BC050600, BC050600.1, BC022804, NM 001111112.2, NM 001111113.2,

NM 005339.5

UniProt ID: P61086

**Summary:** The protein encoded by this gene belongs to the ubiquitin-conjugating enzyme family. This

protein interacts with RING finger proteins, and it can ubiquitinate huntingtin, the gene product for Huntington's disease. Known functions for this protein include a role in aggregate

formation of expanded polyglutamine proteins and the suppression of apoptosis in polyglutamine diseases, a role in the dislocation of newly synthesized MHC class I heavy chains from the endoplasmic reticulum, and involvement in foam cell formation. Multiple transcript variants encoding different isoforms have been identified for this gene. [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 <u>custom shRNA service</u>.







## 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).