

## **Product datasheet for TR311602**

## Product datasneet for TRS 11602

MAGED2 Human shRNA Plasmid Kit (Locus ID 10916)

## Product data:

**Product Type:** shRNA Plasmids

**Product Name:** MAGED2 Human shRNA Plasmid Kit (Locus ID 10916)

**Locus ID:** 10916

Synonyms: 11B6; BARTS5; BCG-1; BCG1; HCA10; MAGE-D2

Vector: pRS (TR20003)

E. coli Selection: Ampicillin

Mammalian Cell Puromycin

Selection:

Format: Retroviral plasmids

Components: MAGED2 - Human, 4 unique 29mer shRNA constructs in retroviral untagged vector(Gene ID =

10916). 5µg purified plasmid DNA per construct

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

**RefSeq:** NM 006787, NM 014599, NM 177433, NM 201222, NM 014599.3, NM 014599.4,

NM 014599.5, NM 201222.1, NM 201222.2, NM 177433.1, NM 177433.2, BC000304,

BC000304.2, BC091503, BM043994, BM803170, NM 177433.3, NM 014599.6, NM 201222.3

UniProt ID: Q9UNF1

**Summary:** This gene is a member of the MAGED gene family. The MAGED genes are clustered on

chromosome Xp11. This gene is located in Xp11.2, a hot spot for X-linked intellectual disability (XLID). Mutations in this gene cause a form of transient antenatal Bartter's

syndrome. This gene may also be involved in several types of cancer, including breast cancer

and melanoma. The protein encoded by this gene is progressively recruited from the cytoplasm to the nucleoplasm during the interphase and after nucleolar stress and is thus thought to play a role in cell cycle regulation. Alternative splicing results in multiple transcript

variants. [provided by RefSeq, Jul 2017]

**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.



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## 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).