

Product datasheet for TL502615

Psma7 Mouse shRNA Plasmid (Locus ID 26444)

Product data:

Product Type: shRNA Plasmids

Product Name: Psma7 Mouse shRNA Plasmid (Locus ID 26444)

Locus ID: 26444 Synonyms: C6-I

Vector:pGFP-C-shLenti (TR30023)E. coli Selection:Chloramphenicol (34 ug/ml)

Mammalian Cell

Selection:

Puromycin

Format: Lentiviral plasmids

Components: Psma7 - Mouse, 4 unique 29mer shRNA constructs in lentiviral GFP vector(Gene ID = 26444).

5µg purified plasmid DNA per construct

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

RefSeq: <u>BC008222</u>, <u>NM 001289476</u>, <u>NM 011969</u>, <u>NM 011969.1</u>, <u>NM 011969.2</u>, <u>NM 001289476.1</u>,

BC014752

UniProt ID: 09Z2U0

Summary: Component of the 20S core proteasome complex involved in the proteolytic degradation of

most intracellular proteins. This complex plays numerous essential roles within the cell by associating with different regulatory particles. Associated with two 19S regulatory particles, forms the 26S proteasome and thus participates in the ATP-dependent degradation of ubiquitinated proteins. The 26S proteasome plays a key role in the maintenance of protein

homeostasis by removing misfolded or damaged proteins that could impair cellular

functions, and by removing proteins whose functions are no longer required. Associated with the PA200 or PA28, the 20S proteasome mediates ubiquitin-independent protein

degradation. This type of proteolysis is required in several pathways including

spermatogenesis (20S-PA200 complex) or generation of a subset of MHC class I-presented

antigenic peptides (20S-PA28 complex).[UniProtKB/Swiss-Prot Function]

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>.



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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).