

## Product datasheet for **TL502622**

### Homer1 Mouse shRNA Plasmid (Locus ID 26556)

#### Product data:

Product Type:	shRNA Plasmids
Product Name:	Homer1 Mouse shRNA Plasmid (Locus ID 26556)
Locus ID:	26556
Synonyms:	homer-1; PSD-Zip45; SYN47; Ves-1; vesl-1
Vector:	pGFP-C-shLenti (TR30023)
E. coli Selection:	Chloramphenicol (34 ug/ml)
Mammalian Cell Selection:	Puromycin
Format:	Lentiviral plasmids
Components:	Homer1 - Mouse, 4 unique 29mer shRNA constructs in lentiviral GFP vector(Gene ID = 26556). 5µg purified plasmid DNA per construct 29-mer scrambled shRNA cassette in pGFP-C-shLenti Vector, TR30021, included for free.
RefSeq:	<a href="#">BC064041</a> , <a href="#">NM_001284189</a> , <a href="#">NM_001347598</a> , <a href="#">NM_011982</a> , <a href="#">NM_147176</a> , <a href="#">NM_152134</a> , <a href="#">NM_011982.1</a> , <a href="#">NM_011982.2</a> , <a href="#">NM_011982.3</a> , <a href="#">NM_011982.4</a> , <a href="#">NM_152134.1</a> , <a href="#">NM_152134.2</a> , <a href="#">NM_152134.3</a> , <a href="#">NM_147176.1</a> , <a href="#">NM_147176.2</a> , <a href="#">NM_147176.3</a> , <a href="#">NM_147176.4</a> , <a href="#">NM_001284189.1</a> , <a href="#">NM_001284189.2</a>
UniProt ID:	<a href="#">Q9Z2Y3</a>



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<b>Summary:</b>	<p>Postsynaptic density scaffolding protein. Binds and cross-links cytoplasmic regions of GRM1, GRM5, ITPR1, DNM3, RYR1, RYR2, SHANK1 and SHANK3. By physically linking GRM1 and GRM5 with ER-associated ITPR1 receptors, it aids the coupling of surface receptors to intracellular calcium release. May also couple GRM1 to PI3 kinase through its interaction with AGAP2. Isoform 1 regulates the trafficking and surface expression of GRM5. Differentially regulates the functions of the calcium activated channel ryanodine receptors RYR1 and RYR2. Isoform 1 decreases the activity of RYR2, and increases the activity of RYR1, whereas isoform 5 counteracts the effects by competing for binding sites. Isoform 3 regulates the trafficking and surface expression of GRM5. Isoform 5 acts as a natural dominant negative, in dynamic competition with constitutively expressed isoform 1, isoform 2 and isoform 3 to regulate synaptic metabotropic glutamate function. Isoform 5, may be involved in the structural changes that occur at synapses during long-lasting neuronal plasticity and development (By similarity). Forms a high-order complex with SHANK1, which in turn is necessary for the structural and functional integrity of dendritic spines (By similarity). Negatively regulates T cell activation by inhibiting the calcineurin-NFAT pathway. Acts by competing with calcineurin/PPP3CA for NFAT protein binding, hence preventing NFAT activation by PPP3CA (By similarity).[UniProtKB/Swiss-Prot Function]</p>
<b>shRNA Design:</b>	<p>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 <a href="mailto:techsupport@origene.com">techsupport@origene.com</a>. If you need a special design or shRNA sequence, please utilize our <a href="#">custom shRNA service</a>.</p>
<b>Performance Guaranteed:</b>	<p>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.</p> <p>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 <a href="mailto:techsupport@origene.com">techsupport@origene.com</a>. Please provide your data indicating the transfection efficiency and measurement of gene expression knockdown compared to the scrambled shRNA control (Western Blot data preferred).</p>