

## Product datasheet for **TL502806**

### Mapk8ip3 Mouse shRNA Plasmid (Locus ID 30957)

#### Product data:

Product Type:	shRNA Plasmids
Product Name:	Mapk8ip3 Mouse shRNA Plasmid (Locus ID 30957)
Locus ID:	30957
Synonyms:	BB120594; D17Wsu15e; JIP-3; Jip3; JSAP1; JSAP1a; JSAP1b; JSAP1c; JSAP1d; mKIAA1066; Syd2
Vector:	pGFP-C-shLenti (TR30023)
E. coli Selection:	Chloramphenicol (34 ug/ml)
Mammalian Cell Selection:	Puromycin
Format:	Lentiviral plasmids
Components:	Mapk8ip3 - Mouse, 4 unique 29mer shRNA constructs in lentiviral GFP vector(Gene ID = 30957). 5µg purified plasmid DNA per construct 29-mer scrambled shRNA cassette in pGFP-C-shLenti Vector, TR30021, included for free.
RefSeq:	<a href="#">BC060603</a> , <a href="#">BC065105</a> , <a href="#">BC068312</a> , <a href="#">NM_001163447</a> , <a href="#">NM_001163448</a> , <a href="#">NM_001163449</a> , <a href="#">NM_001163450</a> , <a href="#">NM_001163451</a> , <a href="#">NM_001163453</a> , <a href="#">NM_013931</a> , <a href="#">NM_001163449.1</a> , <a href="#">NM_001163447.1</a> , <a href="#">NM_013931.1</a> , <a href="#">NM_013931.2</a> , <a href="#">NM_013931.3</a> , <a href="#">NM_013931.4</a> , <a href="#">NM_001163450.1</a> , <a href="#">NM_001163448.1</a> , <a href="#">NM_001163451.1</a> , <a href="#">NM_001163453.1</a> , <a href="#">BC004003</a> , <a href="#">BM899006</a>
UniProt ID:	<a href="#">Q9ESN9</a>



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**Summary:**

The JNK-interacting protein (JIP) group of scaffold proteins selectively mediates JNK signaling by aggregating specific components of the MAPK cascade to form a functional JNK signaling module. May function as a regulator of vesicle transport, through interactions with the JNK-signaling components and motor proteins (PubMed:10523642, PubMed:10629060). Promotes neuronal axon elongation in a kinesin- and JNK-dependent manner (PubMed:23576431, PubMed:25944905, PubMed:28259553). Activates cofilin at axon tips via local activation of JNK, thereby regulating filopodial dynamics and enhancing axon elongation (PubMed:23576431, PubMed:25944905, PubMed:28259553). Its binding to kinesin heavy chains (KHC), promotes kinesin-1 motility along microtubules and is essential for axon elongation and regeneration (PubMed:23576431, PubMed:25944905, PubMed:28259553). Regulates cortical neuronal migration by mediating NTRK2/TRKB anterograde axonal transport during brain development (PubMed:23576431, PubMed:25944905, PubMed:28259553). Acts as an adapter that bridges the interaction between NTRK2/TRKB and KLC1 and drives NTRK2/TRKB axonal but not dendritic anterograde transport, which is essential for subsequent BDNF-triggered signaling and filopodia formation (PubMed:23576431, PubMed:25944905, PubMed:28259553).[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 [techsupport@origene.com](mailto:techsupport@origene.com). 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](mailto: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).