

OriGene Technologies, Inc.

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Product datasheet for TP502531

Rab27a (NM_023635) Mouse Recombinant Protein

Product data:

Product Type:	Recombinant Proteins
Description:	Purified recombinant protein of Mouse RAB27A, member RAS oncogene family (Rab27a), with C-terminal MYC/DDK tag, expressed in HEK293T cells, 20ug
Species:	Mouse
Expression Host:	HEK293T
Expression cDNA Clone or AA Sequence:	>MR202531 protein sequence <mark>Red</mark> =Cloning site Green=Tags(s)
	MSDGDYDYLIKFLALGDSGVGKTSVLYQYTDGKFNSKFITTVGIDFREKRVVYRANGPDGAVGRGQRIHL QLWDTAGQERFRSLTTAFFRDAMGFLLLFDLTNEQSFLNVRNWISQLQMHAYCENPDIVLCGNKSDLEDQ RAVKEEEARELAEKYGIPYFETSAANGTNISHAIEMLLDLIMKRMERCVDKSWIPEGVVRSNGHTSADQL SEEKEKGLCGC
	SGPTRTRPLEQKLISEEDLAANDILDYKDDDDKV
Tag:	C-MYC/DDK
Predicted MW:	25 kDa
Concentration:	>0.05 µg/µL as determined by microplate BCA method
Purity:	> 80% as determined by SDS-PAGE and Coomassie blue staining
Buffer:	25 mM Tris-HCl, 100 mM glycine, pH 7.3, 10% glycerol
Note:	For testing in cell culture applications, please filter before use. Note that you may experience some loss of protein during the filtration process.
Storage:	Store at -80°C after receiving vials.
Stability:	Stable for 12 months from the date of receipt of the product under proper storage and handling conditions. Avoid repeated freeze-thaw cycles.
RefSeq:	<u>NP 076124</u>
Locus ID:	11891
UniProt ID:	<u>Q9ERI2, Q544U7</u>
RefSeq Size:	3066



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	Rab27a (NM_023635) Mouse Recombinant Protein – TP502531
Cytogenetics:	9 40.08 cM
RefSeq ORF:	666
Synonyms:	2210402C08Rik; 2410003M20Rik; 4933437C11Rik; ash
Summary:	The protein encoded by this gene is a member of the Rab family of proteins, which is the largest family within the Ras superfamily of GTPases. This gene product is thought to regulate vesicular transport, together with its specific effectors. Mutations in this gene cause several defects, including actin-based melanosome transport defects and immunodeficiency. Mutations in the human ortholog of this gene are associated with Griscelli syndrome type 2. Alternative splicing results in multiple transcript variants. [provided by RefSeq, Jul 2014]

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