

## Product datasheet for TP503764

## Fhl2 (NM\_010212) Mouse Recombinant Protein

## **Product data:**

## OriGene Technologies, Inc.

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Product Type:	Recombinant Proteins
Description:	Purified recombinant protein of Mouse four and a half LIM domains 2 (Fhl2), with C-terminal MYC/DDK tag, expressed in HEK293T cells, 20ug
Species:	Mouse
Expression Host:	HEK293T
Expression cDNA Clone or AA Sequence:	>MR203764 protein sequence <mark>Red</mark> =Cloning site Green=Tags(s)
	MTERFDCHHCNESLYGKKYILKEENPHCVACFEEPYANTCEECGTPIGCDCKDLSYKDRHWHEGCFHCSR CGSSLVDKPFAAKEEQLLCTDCYSNEYSSKCQECKKTIMPGTRKMEYKGSSWHETCFTCQRCQQPIGTKS FIPKENQNFCVPCYEKQYALQCVQCKKPITTGGVTYREQPWHKECFVCTACKKQLSGQRFTARDEFPYCL TCFCDLYAKKCAGCTNPISGLGGTKYISFEERQWHNDCFNCKKCSLSLVGRGFLTERDDILCPDCGKDI
	TRTRPLEQKLISEEDLAANDILDYKDDDDKV
Tag:	C-MYC/DDK
Predicted MW:	32.1 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 034342</u>
Locus ID:	14200
UniProt ID:	<u>O70433, Q543D7</u>
RefSeq Size:	1566



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	Fhl2 (NM_010212) Mouse Recombinant Protein – TP503764
Cytogenetics:	1 C1.1
RefSeq ORF:	840
Synonyms:	FHL-2; SL; SLIM-3; SLIM3
Summary:	This gene encodes a member of the four-and-a-half-LIM-only protein family. The encoded protein functions as a regulator in numerous signaling pathways and cellular processes in development and cellular differentiation, including development and maintenance of the cardiovascular system and striated muscles. This gene also plays a role in bone formation and regulates and bone mineral content and bone mineral density. Alternative splicing results in multiple transcript variants. [provided by RefSeq, Jan 2014]

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