

## **Product datasheet for TP505854**

## OriGene Technologies, Inc.

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## Rnf34 (NM\_030564) Mouse Recombinant Protein

**Product data:** 

**Product Type:** Recombinant Proteins

**Description:** Purified recombinant protein of Mouse ring finger protein 34 (Rnf34), with C-terminal

MYC/DDK tag, expressed in HEK293T cells, 20ug

Species: Mouse

**Expression Host:** HEK293T

**Expression cDNA Clone** >MR205854 protein sequence or AA Sequence: Red=Cloning site Green=Tags(s)

**TKCGKRMSECPICRQYVVRAVHVFKS** 

**TRTRPLEQKLISEEDLAANDILDYKDDDDKV** 

Tag: C-MYC/DDK

Predicted MW: 42 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:** NP 085041

Locus ID: 80751 UniProt ID: Q99KR6





## Rnf34 (NM\_030564) Mouse Recombinant Protein - TP505854

RefSeq Size: 1980

Cytogenetics: 5 F
RefSeq ORF: 1131

**Synonyms:** AW061037; AW536122; BC004042; C88279; RIFF

Summary: E3 ubiquitin-protein ligase that regulates several biological processes through the ubiquitin-

mediated proteasomal degradation of various target proteins. Ubiquitinates the caspases CASP8 and CASP10, promoting their proteasomal degradation, to negatively regulate cell death downstream of death domain receptors in the extrinsic pathway of apoptosis. May mediate 'Lys-48'-linked polyubiquitination of RIPK1 and its subsequent proteasomal degradation thereby indirectly regulating the tumor necrosis factor-mediated signaling pathway. Negatively regulates p53/TP53 through its direct ubiquitination and targeting to proteasomal

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degradation. Indirectly, may also negatively regulate p53/TP53 through ubiquitination and degradation of SFN. Mediates PPARGC1A proteasomal degradation probably through

ubiquitination thereby indirectly regulating the metabolism of brown fat cells (PubMed:22064484). Possibly involved in innate immunity, through 'Lys-48'-linked

polyubiquitination of NOD1 and its subsequent proteasomal degradation.[UniProtKB/Swiss-

Prot Function]