

**TRIM37 Antibody (C-term)**  
**Affinity Purified Rabbit Polyclonal Antibody (Pab)**  
**Catalog # AP13288b**

### Specification

#### TRIM37 Antibody (C-term) - Product Information

Application	WB,E
Primary Accession	<a href="#">O94972</a>
Other Accession	<a href="#">NP_056109.1</a> , <a href="#">NP_001005207.1</a>
Reactivity	Human
Host	Rabbit
Clonality	Polyclonal
Isotype	Rabbit IgG
Calculated MW	107906
Antigen Region	936-964

#### TRIM37 Antibody (C-term) - Additional Information

##### Gene ID 4591

##### Other Names

E3 ubiquitin-protein ligase TRIM37, 632-, Mulibrey nanism protein, Tripartite motif-containing protein 37, TRIM37, KIAA0898, MUL, POB1

##### Target/Specificity

This TRIM37 antibody is generated from rabbits immunized with a KLH conjugated synthetic peptide between 936-964 amino acids from the C-terminal region of human TRIM37.

##### Dilution

WB~~1:1000

E~~Use at an assay dependent concentration.

##### Format

Purified polyclonal antibody supplied in PBS with 0.09% (W/V) sodium azide. This antibody is purified through a protein A column, followed by peptide affinity purification.

##### Storage

Maintain refrigerated at 2-8°C for up to 2 weeks. For long term storage store at -20°C in small aliquots to prevent freeze-thaw cycles.

##### Precautions

TRIM37 Antibody (C-term) is for research use only and not for use in diagnostic or therapeutic procedures.

#### TRIM37 Antibody (C-term) - Protein Information

Name TRIM37 {ECO:0000303|PubMed:28724525, ECO:0000312|HGNC:HGNC:7523}

**Function** E3 ubiquitin-protein ligase required to prevent centriole reduplication (PubMed:[15885686](#), PubMed:[23769972](#)). Probably acts by ubiquitinating positive regulators of centriole reduplication (PubMed:[23769972](#)). Mediates monoubiquitination of 'Lys-119' of histone H2A (H2AK119Ub), a specific tag for epigenetic transcriptional repression: associates with some Polycomb group (PcG) multiprotein PRC2-like complex and mediates repression of target genes (PubMed:[25470042](#)). Also acts as a positive regulator of peroxisome import by mediating monoubiquitination of PEX5 at 'Lys-472': monoubiquitination promotes PEX5 stabilization by preventing its polyubiquitination and degradation by the proteasome (PubMed:[28724525](#)). Has anti-HIV activity (PubMed:[24317724](#)).

#### Cellular Location

Chromosome. Cytoplasm, perinuclear region. Peroxisome membrane; Peripheral membrane protein. Note=Found in vesicles of the peroxisome. Aggregates as aggresomes, a perinuclear region where certain misfolded or aggregated proteins are sequestered for proteasomal degradation.

#### Tissue Location

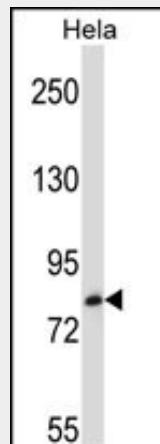
Ubiquitous (PubMed:10888877). Highly expressed in testis, while it is weakly expressed in other tissues (PubMed:16310976).

### TRIM37 Antibody (C-term) - Protocols

Provided below are standard protocols that you may find useful for product applications.

- [Western Blot](#)
- [Blocking Peptides](#)
- [Dot Blot](#)
- [Immunohistochemistry](#)
- [Immunofluorescence](#)
- [Immunoprecipitation](#)
- [Flow Cytometry](#)
- [Cell Culture](#)

### TRIM37 Antibody (C-term) - Images



TRIM37 Antibody (C-term) (Cat. #AP13288b) western blot analysis in HeLa cell line lysates (35ug/lane). This demonstrates the TRIM37 antibody detected the TRIM37 protein (arrow).

### TRIM37 Antibody (C-term) - Background

This gene encodes a member of the tripartite motif (TRIM) family, whose members are involved in diverse cellular functions such as developmental patterning and oncogenesis. The TRIM motif includes zinc-binding domains, a RING finger region, a B-box motif and a coiled-coil domain. The RING finger and B-box domains chelate zinc and might be involved in protein-protein and/or protein-nucleic acid interactions. The gene mutations are associated with mulibrey (muscle-liver-brain-eye) nanism, an autosomal recessive disorder that involves several tissues of mesodermal origin. Alternatively spliced transcript variants encoding the same protein have been identified. [provided by RefSeq].

**TRIM37 Antibody (C-term) - References**

Xin, X., et al. Genome Res. 19(7):1262-1269(2009)  
Karlberg, S., et al. Mod. Pathol. 22(4):570-578(2009)  
Doganc, T., et al. Clin. Dysmorphol. 16(3):173-176(2007)  
Hamalainen, R.H., et al. Clin. Genet. 70(6):473-479(2006)  
Olsen, J.V., et al. Cell 127(3):635-648(2006)