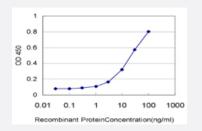


# TRIM37 monoclonal antibody (M01), clone 2D11

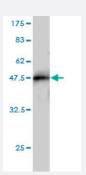
Catalog # H00004591-M01 Size 100 ug

### Applications



#### Sandwich ELISA (Recombinant protein)

Detection limit for recombinant GST tagged TRIM37 is approximately 0.3ng/ml as a capture antibody.



Western Blot detection against Immunogen (36.74 KDa).

Specification	
Product Description	Mouse monoclonal antibody raised against a partial recombinant TRIM37.
Immunogen	TRIM37 (NP_056109, 865 a.a. ~ 964 a.a) partial recombinant protein with GST tag. MW of the GST t ag alone is 26 KDa.
Sequence	GHLEGLQMTDLENNSETGELQPVLPEGASAAPEEGMSSDSDIECDTENEEQEEHTSVGGFHDS FMVMTQPPDEDTHSSFPDGEQIGPEDLSFNTDENSGR
Host	Mouse
Reactivity	Human



### **Product Information**

Interspecies Antigen Sequence	Mouse (77); Rat (77)
Isotype	lgG2a Kappa
Quality Control Testing	Antibody Reactive Against Recombinant Protein. Western Blot detection against Immunogen (36.74 KDa) .
Storage Buffer	In 1x PBS, pH 7.4
Storage Instruction	Store at -20°C or lower. Aliquot to avoid repeated freezing and thawing.

## Applications

- Western Blot (Recombinant protein)
  <u>Protocol Download</u>
- Sandwich ELISA (Recombinant protein)
  Detection limit for recombinant GST tagged TRIM37 is approximately 0.3ng/ml as a capture antibody.
  <u>Protocol Download</u>
- ELISA

### Gene Info — TRIM37

Entrez GenelD	<u>4591</u>
GeneBank Accession#	<u>NM_015294</u>
Protein Accession#	<u>NP_056109</u>
Gene Name	TRIM37
Gene Alias	KIAA0898, MUL, POB1, TEF3
Gene Description	tripartite motif-containing 37
Omim ID	<u>253250 605073</u>
Gene Ontology	Hyperlink



**Gene Summary** 

#### **Product Information**

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 incl udes zinc-binding domains, a RING finger region, a B-box motif and a coiled-coil domain. The RI NG 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-ey e) nanism, an autosomal recessive disorder that involves several tissues of mesodermal origin. Al ternatively spliced transcript variants encoding the same protein have been identified. [provided b y RefSeq

**Other Designations** 

RING-B-box-coiled-coil protein|tripartite motif-containing 37 protein

#### Pathway

<u>Ubiquitin mediated proteolysis</u>