

## Anti-PD-L1 Polyclonal Antibody 3

### Product Details

Ig Type:	IgG
Reactivity:	Human,Mouse,Rat (predicted:Pig,Cow,Horse,Sheep)
Molecular Weight:	Theoretical: 32 kDa.
Purification:	Protein A purified

### Applications

#### Sample:

LOVO (Human) Cell Lysate at 30  $\mu$ g

Heart (Mouse) Lysate at 40  $\mu$ g

Kidney (Mouse) Lysate at 40  $\mu$ g

Muscle (Mouse) Lysate at 40  $\mu$ g

#### Verified Activity:

Heart (Rat) Lysate at 40  $\mu$ g

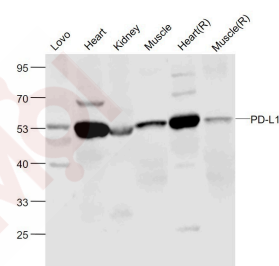
Muscle (Rat) Lysate at 40  $\mu$ g

Primary: Anti-PD-L1 (TMAB-01361) at 1/500 dilution

Secondary: IRDye800CW Goat Anti-Rabbit IgG at 1/20000 dilution

Predicted band size: 50 kDa

Observed band size: 53 kDa



Application: WB

Recommended WB: 1:500-2000

### Properties

Stability & Storage: Store at -20°C or -80°C for 12 months. Avoid repeated freeze-thaw cycles.

Shipping: Shipping with blue ice.

### Antigen Details

Immunogen:	KLH conjugated synthetic peptide: human B7-H1/PDL1/CD274
Antigen Species:	Human
Gene ID:	29126
Uniprot ID:	Q9NZQ7
Synonyms:	Pdcd1lg1;Pdcd1l1;CD274 molecule;A530045L16Rik;Pdl1;B7h1
Biology Area:	Tumor antigens,Tumor-associated antigens,Non-lineage,Autoimmune,pd-l1

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### Research Background

This gene encodes an immune inhibitory receptor ligand that is expressed by hematopoietic and non-hematopoietic cells, such as T cells and B cells and various types of tumor cells. The encoded protein is a type I transmembrane protein that has immunoglobulin V-like and C-like domains. Interaction of this ligand with its receptor inhibits T-cell activation and cytokine production. During infection or inflammation of normal tissue, this interaction is important for preventing autoimmunity by maintaining homeostasis of the immune response. In tumor microenvironments, this interaction provides an immune escape for tumor cells through cytotoxic T-cell inactivation. Expression of this gene in tumor cells is considered to be prognostic in many types of human malignancies, including colon cancer and renal cell carcinoma. Alternative splicing results in multiple transcript variants. [provided by RefSeq, Sep 2015]

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