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# RUNX1 (PT0706R) PT® Rabbit mAb

CatalogNo: YM8515 Recombinant R

## Key Features

Host Species

Rabbit

MW • 49kD (Calculated) 43-55kD (Observed) Reactivity

Human,Mouse,Rat

Isotype

IgG,Kappa

Applications
• WB,IHC,IF,IP,ELISA

### **Recommended Dilution Ratios**

IHC 1:200-1:1000 WB 1:2000-1:10000 IF 1:200-1:1000 ELISA 1:5000-1:20000 IP 1:50-1:200

## **Storage**

Storage*	-15°C to -25°C/1 year(Do not lower than -25°C)
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Formulation PBS, 50% glycerol, 0.05% Proclin 300, 0.05% BSA

### **Basic Information**

Clonality	Monoclonal
Clone Number	PT0706R

### Immunogen Information

Specificity Endogenous

## Target Information

	Gene na	ame	RUNX1
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#### **Protein Name**

#### Name Runt-related transcription factor 1

Organism	Gene ID	UniProt ID
Human	<u>861;</u>	<u>Q01196;</u>
Mouse	<u>12394;</u>	<u>Q03347;</u>
Rat	<u>50662;</u>	<u>Q63046;</u>

#### Cellular Nucleus

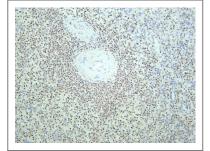
#### Localization

- **Tissue specificity** Expressed in all tissues examined except brain and heart. Highest levels in thymus, bone marrow and peripheral blood.
- Function Alternative products: Additional isoforms seem to exist, Caution: The fusion of AML1 with EAP in T-MDS induces a change of reading frame in the latter resulting in 17 AA unrelated to those of EAP., Disease: A chromosomal aberration involving RUNX1/AML1 is a cause of chronic myelogenous leukemia (CML). Translocation t(3;21)(g26;g22) with EAP, MSD1 or EVI1., Disease: A chromosomal aberration involving RUNX1/AML1 is a cause of chronic myelomonocytic leukemia. Inversion inv(21)(g21:g22) with USP16. Disease: A chromosomal aberration involving RUNX1/AML1 is a cause of M2 type acute myeloid leukemia (AML-M2). Translocation t(8;21)(q22;q22) with RUNX1T1/MTG8/ETO., Disease: A chromosomal aberration involving RUNX1/AML1 is a cause of therapy-related myelodysplastic syndrome (T-MDS). Translocation t(3;21)(g26;g22) with EAP, MSD1 or EVI1., Disease: A chromosomal aberration involving RUNX1/AML1 is found in childhood acute lymphoblastic leukemia (ALL). Translocation t(12;21)(p13;q22) with TEL. The translocation fuses the 3'-end of TEL to the alternate 5'-exon of AML-1H., Disease: A chromosomal aberration involving RUNX1/AML1 is found in therapy-related myeloid malignancies. Translocation t(16;21)(g24;g22) that forms a RUNX1-CBFA2T3 fusion protein., Disease: Defects in RUNX1 are the cause of familial platelet disorder with associated myeloid malignancy (FPDMM) [MIM:601399]. FPDMM is an autosomal dominant disease characterized by qualitative and quantitative platelet defects, and propensity to develop acute myelogenous leukemia., Domain: A proline/serine/threonine rich region at the C-terminus is necessary for transcriptional activation of target genes., Function: CBF binds to the core site, 5'-PYGPYGGT-3', of a number of enhancers and promoters, including murine leukemia virus, polyomavirus enhancer, T-cell receptor enhancers, LCK, IL-3 and GM-CSF promoters. The alpha subunit binds DNA and appears to have a role in the development of normal hematopoiesis. Isoform AML-1L interferes with the transactivation activity of RUNX1. Acts synergistically with ELF4 to transactivate the IL-3 promoter and with ELF2 to transactivate the mouse BLK promoter. Inhibits MYST4dependent transcriptional activation., PTM: Methylated., PTM: Phosphorylated in its C-terminus upon IL-6 treatment. Phosphorylation enhances interaction with MYST3., similarity: Contains 1 Runt domain., subunit: Heterodimer with CBFB. RUNX1 binds DNA as a monomer and through the Runt domain. DNA-binding is increased by heterodimerization. Isoform AML-1L can neither bind DNA nor heterodimerize. Interacts with TLE1 and THOC4. Interacts with ELF1, ELF2 and SPI1. Interacts via its Runt domain with the ELF4 N-terminal region. Interaction with ELF2 isoform 2 (NERF-1a) may act to repress RUNX1-mediated transactivation. Interacts with MYST3 and MYST4. Interacts with SUV39H1. leading to abrogate the transactivating and DNA-binding properties of RUNX1., tissue specificity: Expressed in all tissues examined except brain and heart. Highest levels in thymus, bone marrow and peripheral blood.,

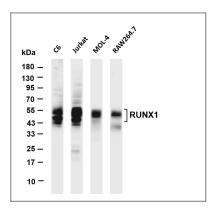
## Validation Data



Human tonsil was stained with anti-RUNX1 rabbit antibody



Human spleen was stained with anti-RUNX1 rabbit antibody



Various whole cell lysates were separated by 4-20% SDS-PAGE, and the membrane was blotted with anti-RUNX1 antibody. The HRP-conjugated Goat anti-Rabbit IgG(H + L) antibody was used to detect the antibody. Lane 1: C6 Lane 2: Jurkat Lane 3: MOL-4 Lane 4: RAW264.7 Predicted band size: 49kDa Observed band size: 43-55kDa

## **Contact information**

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Please scan the QR code to access additional product information: **RUNX1 (PT0706R) PT® Rabbit mAb** 

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