

TWIST1 (PT0464R) PT® Rabbit mAb

CatalogNo: YM8299 **Recombinant** 

Key Features

Host Species

- Rabbit

Reactivity

- WB,IF,IP,ELISA

Applications

- Human,Mouse,Rat,

MW

- 21kD (Calculated)
- 26kD (Observed)

Isotype

- IgG,Kappa

Recommended Dilution Ratios

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)

Basic Information

Clonality Monoclonal**Clone Number** PT0464R

Immunogen Information

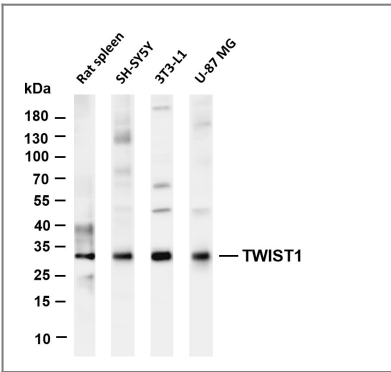
Specificity Endogenous

Target Information

Gene name TWIST1 BHLHA38 TWIST

Protein Name	Twist-related protein 1 (Class A basic helix-loop-helix protein 38) (bHLHa38) (H-twist)		
	Organism	Gene ID	UniProt ID
	Human	7291 ;	Q15672 ;
	Mouse		P26687 ;
Cellular Localization	Nucleus		
Tissue specificity	Subset of mesodermal cells.		
Function	<p>Disease:Defects in TWIST1 are a cause of Saethre-Chotzen syndrome (SCS) [MIM:101400]; also known as acrocephalosyndactyly type 3 (ACS3). SCS is a craniosynostosis syndrome characterized by coronal synostosis, brachycephaly, low frontal hairline, facial asymmetry, hypertelorism, broad halluces, and clinodactyly.,Disease:Defects in TWIST1 are the cause of craniosynostosis type 1 (CRS1) [MIM:123100]. Craniosynostosis consists of premature fusion of one or more cranial sutures, resulting in an abnormal head shape.,Disease:Defects in TWIST1 are the cause of Robinow-Sorauf syndrome (RSS) [MIM:180750]; also known as craniosynostosis-bifid hallux syndrome. RSS is an autosomal dominant defect characterized by minor skull and limb anomalies which is very similar to Saethre-Chotzen syndrome.,Function:Probable transcription factor, which seems to be involved in the negative regulation of cellular determination and in the differentiation of several lineages including myogenesis, osteogenesis, and neurogenesis. Inhibits myogenesis by sequestering E proteins, inhibiting trans-activation by MEF2, and inhibiting DNA-binding by MYOD1 through physical interaction. This interaction probably involves the basic domains of both proteins (By similarity). Also represses expression of proinflammatory cytokines such as TNFA and IL1B.,similarity:Contains 1 basic helix-loop-helix (bHLH) domain.,subunit:Efficient DNA binding requires dimerization with another bHLH protein. Homodimer.,tissue specificity:Subset of mesodermal cells.,</p>		

| Validation Data



Various whole cell lysates were separated by 4-20% SDS-PAGE, and the membrane was blotted with anti-TWIST1 (PT0464R) antibody. The HRP-conjugated Goat anti-Rabbit IgG(H + L) antibody was used to detect the antibody. Lane 1: Rat spleen Lane 2: SH-SY5Y Lane 3: 3T3-L1 Lane 4: U-87 MG Predicted band size: 21kDa Observed band size: 26kDa

| Contact information

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