

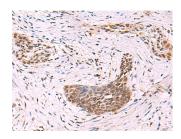
Tel:240-252-7368(USA) Fax: 240-252-7376(USA) techsupport@elabscience.com Website: www.elabscience.com

# **CREBBP Polyclonal Antibody**

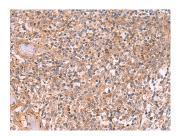
Catalog No.E-AB-52070ReactivityH,M,RStorageStore at -20°C. Avoid freeze / thaw cycles.HostRabbitApplicationsIHC,ELISAIsotypeIgG

**Note:** Centrifuge before opening to ensure complete recovery of vial contents.

# **Images**



Immunohistochemistry of paraffinembedded Human esophagus cancer tissue using CREBBP Polyclonal Antibody at dilution of 1:35(×200)



Immunohistochemistry of paraffinembedded Human tonsil tissue using CREBBP Polyclonal Antibody at dilution of 1:35(×200)

# **Immunogen Information**

Immunogen Synthetic peptide of human CREBBP

**Gene Accession** NP004371 **Swissprot** Q92793

**Synonyms** CBP,CBP,CREB binding protein,CREB-binding

protein, Crebbp, Cyclic AMP responsive enhancer binding protein, KAT3A, RSTS, RTS, Rubinstein Taybi

syndrome

#### **Product Information**

**Buffer** PBS with 0.05% NaN3 and 40% Glycerol,pH7.4

**Purify** Antigen affinity purification

**Dilution** IHC 1:30-1:150, ELISA 1:5000-1:10000

# **Background**

This gene is ubiquitously expressed and is involved in the transcriptional coactivation of many different transcription factors. First isolated as a nuclear protein that binds to cAMP-response element binding protein (CREB), this gene is now known to play critical roles in embryonic development, growth control, and homeostasis by coupling chromatin remodeling to transcription factor recognition. The protein encoded by this gene has intrinsic histone acetyltransferase activity and also acts as a scaffold to stabilize additional protein interactions with the transcription complex. This protein acetylates both histone and non-histone proteins. This protein shares regions of very high sequence similarity with protein p300 in its bromodomain, cysteine-histidine-rich regions, and histone acetyltransferase domain. Mutations in this gene cause Rubinstein-Taybi syndrome (RTS). Chromosomal translocations involving this gene have been associated with acute myeloid leukemia. Alternative splicing results in multiple transcript variants encoding different isoforms.