

# Anti-RUNX3 Antibody

Rabbit polyclonal antibody to RUNX3 Catalog # AP59501

#### **Product Information**

Application WB Primary Accession Q13761

**Reactivity** Human, Chicken

HostRabbitClonalityPolyclonalCalculated MW44356

#### **Additional Information**

Gene ID 864

Other Names AML2; CBFA3; PEBP2A3; Runt-related transcription factor 3; Acute myeloid

leukemia 2 protein; Core-binding factor subunit alpha-3; CBF-alpha-3; Oncogene AML-2; Polyomavirus enhancer-binding protein 2 alpha C subunit; PEA2-alpha C; PEBP2-alpha C; SL3-3 enhancer factor 1 alpha C subunit;

SL3/AKV core-binding factor alpha C subunit

**Target/Specificity** KLH-conjugated synthetic peptide encompassing a sequence within the center

region of human RUNX3. The exact sequence is proprietary.

**Dilution** WB~~WB (1/500 - 1/1000)

**Format** Liquid in 0.42% Potassium phosphate, 0.87% Sodium chloride, pH 7.3, 30%

glycerol, and 0.09% (W/V) sodium azide.

**Storage** Store at -20 °C.Stable for 12 months from date of receipt

### **Protein Information**

Name RUNX3

**Synonyms** AML2, CBFA3, PEBP2A3

**Function** Forms the heterodimeric complex core-binding factor (CBF) with CBFB.

RUNX members modulate the transcription of their target genes through recognizing the core consensus binding sequence 5'- TGTGGT-3', or very rarely, 5'-TGCGGT-3', within their regulatory regions via their runt domain, while CBFB is a non-DNA-binding regulatory subunit that allosterically enhances the sequence-specific DNA-binding capacity of RUNX. The

heterodimers bind to the core site of a number of enhancers and promoters, including murine leukemia virus, polyomavirus enhancer, T-cell receptor enhancers, LCK, IL3 and GM-CSF promoters (By similarity). May be involved in

the control of cellular proliferation and/or differentiation. In association with ZFHX3, up- regulates CDKN1A promoter activity following TGF-beta stimulation (PubMed:20599712). CBF complexes repress ZBTB7B transcription factor during cytotoxic (CD8+) T cell development. They bind to RUNX-binding sequence within the ZBTB7B locus acting as transcriptional silencer and allowing for cytotoxic T cell differentiation. CBF complexes binding to the transcriptional silencer is essential for recruitment of nuclear protein complexes that catalyze epigenetic modifications to establish epigenetic ZBTB7B silencing (By similarity). Necessary for the development and survival of sensory neurons expressing parvalbumin (By similarity).

**Cellular Location** 

Nucleus {ECO:0000255 | PROSITE-ProRule:PRU00399, ECO:0000269 | PubMed:20100835, ECO:0000269 | PubMed:20599712}. Cytoplasm. Note=The tyrosine phosphorylated form localizes to the cytoplasm. Translocates from the cytoplasm to the nucleus following TGF-beta stimulation

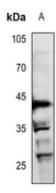
**Tissue Location** 

Expressed in gastric cancer tissues (at protein level).

## **Background**

KLH-conjugated synthetic peptide encompassing a sequence within the center region of human RUNX3. The exact sequence is proprietary.

## **Images**



Western blot analysis of RUNX3 expression in EC9706 (A) whole cell lysates.

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