



Catalog #: APRab06188



# Summary

FoxO4 (Acetyl Lys189) Rabbit Polyclonal Antibody **Production Name** 

Description Rabbit Polyclonal Antibody

Rabbit Host **Application** WB.ELISA

Reactivity Human, Rat, Mouse

### **Performance**

Conjugation Unconjugated Modification Acetylated

Isotype IgG

**Clonality** Polyclonal **Form** Liquid

Store at 4°C short term. Aliquot and store at -20°C long term. Avoid freeze/thaw cycles. Storage

Liquid in PBS containing 50% glycerol, 0.5% protective protein and 0.02% New type

preservative N.

**Purification** Affinity purification

## **Immunogen**

**Buffer** 

Gene Name FOXO4 AFX AFX1 MLLT7

**Alternative Names** Forkhead box protein O4 (Fork head domain transcription factor AFX1)

Gene ID 4303.0

SwissProt ID P98177. Synthetic Acetyl peptide from human protein at AA range: 189

# **Application**

**Dilution Ratio** WB 1:500-2000, ELISA 1:10000-20000

**Molecular Weight** 55kDa

# **Background**

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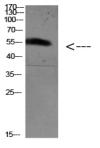
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This gene encodes a member of the O class of winged helix/forkhead transcription factor family. Proteins encoded by this class are regulated by factors involved in growth and differentiation indicating they play a role in these processes. A translocation involving this gene on chromosome X and the homolog of the Drosophila trithorax gene, encoding a DNA binding protein, located on chromosome 11 is associated with leukemia. Multiple transcript variants encoding different isoforms have been found for this gene. [provided by RefSeq, Jan 2010], disease: A chromosomal aberration involving FOXO4 is found in acute leukemias. Translocation t(X;11)(q13;q23) with MLL/HRX. The result is a rogue activator protein, function: Transcription factor involved in the regulation of the insulin signaling pathway. Binds to insulin-response elements (IREs) and can activate transcription of IGFBP1. Down-regulates expression of HIF1A and suppresses hypoxiainduced transcriptional activation of HIF1A-modulated genes. Also involved in negative regulation of the cell cycle, pharmaceutical: A constitutively active FOXO4 mutant where phosphorylation sites Thr-32, Ser-187 and Ser-262 have been mutated to alanine may have therapeutic potential in ERBB2/HER2-overexpressing cancers as it inhibits ERBB2mediated cell survival, transformation and tumorigenicity, PTM: Acetylation by CBP, which is induced by peroxidase stress, inhibits transcriptional activity. Deacetylation by SIRT1 is NAD-dependent and stimulates transcriptional activity, PTM: Phosphorylation by PKB/AKT1 inhibits transcriptional activity and is responsible for cytoplasmic localization., similarity: Contains 1 fork-head DNA-binding domain., subcellular location: When phosphorylated, translocated from nucleus to cytoplasm. Dephosphorylation triggers nuclear translocation., subunit: Interacts with CBP, MYOCD, SIRT1, SRF and YWHAZ. Acetylated by CBP and deacetylated by SIRT1. Binding of YWHAZ inhibits DNA-binding, tissue specificity: Heart, brain, placenta, lung, liver, skeletal muscle, kidney and pancreas. Isoform zeta is most abundant in the liver, kidney, and pancreas.,

#### Research Area

## **Image Data**



Western blot analysis of 3T3 mouse-kidney KB K562 Hela lysate, antibody was diluted at 500. Secondary antibody was diluted at 1:20000

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## Note

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