

CA IX Polyclonal Antibody

YT0575 Catalog No:

Reactivity: Human; Mouse

Applications: WB;ELISA

CAIX Target:

Fields: >>Nitrogen metabolism;>>Metabolic pathways

Gene Name: CA9

Protein Name: Carbonic anhydrase 9

Q16790

Q8VHB5

Human Gene Id: 768

Human Swiss Prot

No:

Mouse Swiss Prot

No:

Immunogen: The antiserum was produced against synthesized peptide derived from human

CA IX. AA range:33-82

CA IX Polyclonal Antibody detects endogenous levels of CA IX protein. **Specificity:**

Formulation: Liquid in PBS containing 50% glycerol, 0.5% BSA and 0.02% sodium azide.

Source: Polyclonal, Rabbit, IgG

Dilution: WB 1:500 - 1:2000. ELISA: 1:10000. Not yet tested in other applications.

The antibody was affinity-purified from rabbit antiserum by affinity-**Purification:**

chromatography using epitope-specific immunogen.

Concentration: 1 mg/ml

-15°C to -25°C/1 year(Do not lower than -25°C) **Storage Stability:**

1/3



Observed Band: 58kD

Cell Pathway : Nitrogen metabolism;

Background:

Carbonic anhydrases (CAs) are a large family of zinc metalloenzymes that catalyze the reversible hydration of carbon dioxide. They participate in a variety of biological processes, including respiration, calcification, acid-base balance, bone resorption, and the formation of aqueous humor, cerebrospinal fluid, saliva, and gastric acid. They show extensive diversity in tissue distribution and in their subcellular localization. CA IX is a transmembrane protein and is one of only two tumor-associated carbonic anhydrase isoenzymes known. It is expressed in all clear-cell renal cell carcinoma, but is not detected in normal kidney or most other normal tissues. It may be involved in cell proliferation and transformation. This gene was mapped to 17q21.2 by fluorescence in situ hybridization, however, radiation hybrid mapping localized it to 9p13-p12. [provided by RefSeq, Jun 2014],

Function:

catalytic activity: H(2)CO(3) = CO(2) + H(2)O., cofactor: Zinc., function: Reversible hydration of carbon dioxide. Participates in pH regulation. May be involved in the control of cell proliferation and transformation. Appears to be a novel specific biomarker for a cervical neoplasia., induction: By hypoxia., PTM: Asn-346 bears high-mannose type glycan structures., similarity: Belongs to the alpha-carbonic anhydrase family., subcellular location: Found on the surface microvilli and in the nucleus, particularly in nucleolus., subunit: Forms oligomers linked by disulfide bonds., tissue specificity: Expressed primarily in carcinoma cells lines. Expression is restricted to very few normal tissues and the most abundant expression is found in the epithelial cells of gastric mucosa.,

Subcellular Location:

Nucleus . Nucleus, nucleolus . Cell membrane ; Single-pass type I membrane protein . Cell projection, microvillus membrane ; Single-pass type I membrane protein . Found on the surface microvilli and in the nucleus, particularly in nucleolus.

Expression:

Expressed primarily in carcinoma cells lines. Expression is restricted to very few normal tissues and the most abundant expression is found in the epithelial cells of gastric mucosa.

Sort: 2965

No4:

Host: Rabbit

Modifications: Unmodified



Products Images