

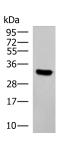
Tel:240-252-7368(USA) Fax: 240-252-7376(USA) techsupport@elabscience.com Website: www.elabscience.com

# **KCTD7 Polyclonal Antibody**

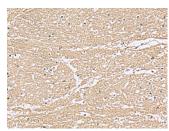
Catalog No.E-AB-18176ReactivityH,M,RStorageStore at -20°C. Avoid freeze / thaw cycles.HostRabbitApplicationsWB,IHC,ELISAIsotypeIgG

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

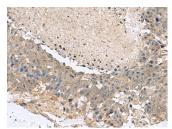
# **Images**



Western blot analysis of Hela cell lysate using KCTD7 Polyclonal Antibody at dilution of 1:1200



Immunohistochemistry of paraffinembedded Human brain tissue using KCTD7 Polyclonal Antibody at dilution of 1:55(×200)



Immunohistochemistry of paraffinembedded Human liver cancer tissue using KCTD7 Polyclonal Antibody at dilution of 1:55(×200)

# **Immunogen Information**

**Immunogen** Synthetic peptide of human KCTD7

**Gene Accession** NP694578 **Swissprot** Q96MP8

**Synonyms** BTB/POZ domain containing protein

KCTD7,EPM3,FLJ32069,Potassium channel

tetramerisation domain containing 7

#### **Product Information**

Calculated MW 33 kDa

**Observed MW** Refer to figures

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

**Purify** Antigen affinity purification

**Dilution** WB 1:1000-1:5000, IHC 1:50-1:300, ELISA

1:5000-1:10000

# **Background**

KCTD7 (Potassium Channel Tetramerization Domain Containing 7) is a Protein Coding gene. Diseases associated with KCTD7 include Epilepsy, Progressive Myoclonic 3, With Or Without Intracellular Inclusions and Cln14 Disease. Among its related pathways are Neuropathic Pain-Signaling in Dorsal Horn Neurons and Innate Immune System. An important paralog of this gene is KCTD14. This gene encodes a member of the potassium channel tetramerization domain-containing protein family. Family members are identified on a structural basis and contain an amino-terminal domain similar to the T1 domain present in the voltage-gated potassium channel. Mutations in this gene have been associated with progressive myoclonic epilepsy-3. Alternative splicing results in multiple transcript variants.

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