

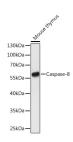
Tel:240-252-7368(USA) Fax: 240-252-7376(USA) techsupport@elabscience.com Website: www.elabscience.com

# **Caspase-8 Polyclonal Antibody**

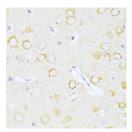
Catalog No.E-AB-63511ReactivityH,M,RStorageStore at -20°C. Avoid freeze / thaw cycles.HostRabbitApplicationsWB,IHC,IFIsotypeIgG

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

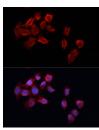
# **Images**



Western blot analysis of extracts of Mouse thymus using Caspase-8 Polyclonal Antibody at dilution of 1:1000.



Immunohistochemistry of paraffinembedded Rat brain using Caspase-8 Polyclonal Antibody at dilution of 1:100 (40x lens).



Immunofluorescence analysis of HeLa cells using Caspase-8 Polyclonal Antibody at dilution of 1:100 (40x lens). Blue: DAPI for nuclear staining.

# **Immunogen Information**

Immunogen Recombinant protein of mouse Caspase-8

**GeneID** 12370 **Swissprot** Q14790

Synonyms CASP8, ALPS2B, CAP4, Casp-8, FLICE, MACH, MCH

5,caspase-8,Caspase 8,Casp8

#### **Product Information**

Calculated MW 55kDa
Observed MW 58kDa

**Buffer** PBS with 0.02% sodium azide, 50% glycerol, pH7.3.

**Purify** Affinity purification

**Dilution** WB 1:500-1:2000 IHC 1:50-1:200 IF 1:50-1:200

# **Background**

This gene encodes a member of the cysteine-aspartic acid protease (caspase) family. Sequential activation of caspases plays a central role in the execution-phase of cell apoptosis. Caspases exist as inactive proenzymes composed of a prodomain, a large protease subunit, and a small protease subunit. Activation of caspases requires proteolytic processing at conserved internal aspartic residues to generate a heterodimeric enzyme consisting of the large and small subunits. This protein is involved in the programmed cell death induced by Fas and various apoptotic stimuli. The N-terminal FADD-like death effector domain of this protein suggests that it may interact with Fas-interacting protein FADD. This protein was detected in the insoluble fraction of the affected brain region from Huntington disease patients but not in those from normal controls, which implicated the role in neurodegenerative diseases. Many alternatively spliced transcript variants encoding different isoforms have been described, although not all variants have had their fulllength sequences determined.

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