

Cleaved-Caspase-5 p10 (S331) Polyclonal Antibody

Catalog No :	YC0030
Reactivity :	Human;
Applications :	WB;ELISA
Target :	Caspase-5
Fields :	>>NOD-like receptor signaling pathway;>>Salmonella infection
Gene Name :	CASP5
Protein Name :	Caspase5
Human Gene Id :	838
Human Swiss Prot	P51878
No : Immunogen :	The antiserum was produced against synthesized peptide derived from human Caspase 5. AA range:312-361
Specificity :	Cleaved-Caspase-5 p10 (S331) Polyclonal Antibody detects endogenous levels of fragment of activated Caspase-5 p10 protein resulting from cleavage adjacent to S331.
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: 1:20000. Not yet tested in other applications.
Purification :	The antibody was affinity-purified from rabbit antiserum by affinity- chromatography using epitope-specific immunogen.
Concentration :	1 mg/ml
Storage Stability :	-15°C to -25°C/1 year(Do not lower than -25°C)



Best Tools for immunology Research **Observed Band :** 10kD **Cell Pathway :** NOD-like receptor; This gene encodes a member of the cysteine-aspartic acid protease (caspase) **Background**: family. Sequential activation of caspases plays a central role in the executionphase of cell apoptosis. Caspases exist as inactive proenzymes which undergo proteolytic processing at conserved aspartic residues to produce two subunits, large and small, that dimerize to form the active enzyme. Overexpression of the active form of this enzyme induces apoptosis in fibroblasts. Max, a central component of the Myc/Max/Mad transcription regulation network important for cell growth, differentiation, and apoptosis, is cleaved by this protein; this process requires Fas-mediated dephosphorylation of Max. The expression of this gene is regulated by interferon-gamma and lipopolysaccharide. Alternatively spliced transcript variants have been identified for this gene. [provided by RefSeq, Aug 2010], catalytic activity:Strict requirement for Asp at the P1 position. It has a preferred **Function:** cleavage sequence of Tyr-Val-Ala-Asp-I- but also cleaves at Asp-Glu-Val-Asp-[-.,function:Mediator of programmed cell death (apoptosis).,PTM:The two subunits are derived from the precursor sequence by an autocatalytic mechanism., similarity: Belongs to the peptidase C14A family., similarity: Contains 1 CARD domain.,subunit:Heterotetramer that consists of two anti-parallel arranged heterodimers, each one formed by a 20 kDa (p20) and a 10 kDa (p10) subunits.,tissue specificity:Expressed in barely detectable amounts in most tissues except brain, highest levels being found in lung, liver and skeletal muscle. Subcellular neuron projection, neuronal cell body, IPAF inflammasome complex, NLRP1 inflammasome complex,NLRP3 inflammasome complex,AIM2 inflammasome Location : complex, **Expression**: Expressed in barely detectable amounts in most tissues except brain, highest levels being found in lung, liver and skeletal muscle.

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