

bs-3436R**[Primary Antibody]****phospho-TAK1 (Thr184) Rabbit pAb****BioSS**
ANTIBODIES

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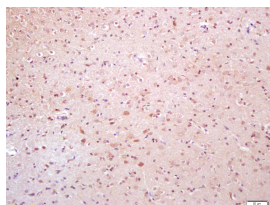
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— DATASHEET —

Host: Rabbit	Isotype: IgG	Applications: IHC-P (1:100-500) IHC-F (1:100-500)
Clonality: Polyclonal		
GeneID: 6885	SWISS: O43318	Reactivity: Rat (predicted: Human, Mouse, Rabbit, Pig, Cow, Chicken, Horse)
Target: TAK1 (Thr184)		
Immunogen: KLH conjugated Synthesised phosphopeptide derived from human TAK1 around the phosphorylation site of Thr184: IQ(p-T)HM.		
Purification: affinity purified by Protein A		Predicted MW.: 67 kDa
Concentration: 1mg/ml		Subcellular Location: Cell membrane ,Cytoplasm
Storage: 0.01M TBS (pH7.4) with 1% BSA, 0.02% Proclin300 and 50% Glycerol. Shipped at 4°C. Store at -20°C for one year. Avoid repeated freeze/thaw cycles.		
Background: The protein encoded by this gene is a member of the serine/threonine protein kinase family. This kinase mediates the signaling transduction induced by TGF beta and morphogenetic protein (BMP), and controls a variety of cell functions including transcription regulation and apoptosis. In response to IL-1, this protein forms a kinase complex including TRAF6, MAP3K7P1/TAB1 and MAP3K7P2/TAB2; this complex is required for the activation of nuclear factor kappa B. This kinase can also activate MAPK8/JNK, MAP2K4/MKK4, and thus plays a role in the cell response to environmental stresses. Four alternatively spliced transcript variants encoding distinct isoforms have been reported. [provided by RefSeq, Jul 2008]		

— VALIDATION IMAGES —

Tissue/cell: rat brain tissue; 4% Paraformaldehyde-fixed and paraffin-embedded; Antigen retrieval: citrate buffer (0.01M, pH 6.0), Boiling bathing for 15min; Block endogenous peroxidase by 3% Hydrogen peroxide for 30min; Blocking buffer (normal goat serum, C-0005) at 37°C for 20 min; Incubation: Anti-Phospho-TAK1(Thr184) Polyclonal Antibody, Unconjugated(bs-3436R) 1:200, overnight at 4°C, followed by conjugation to the secondary antibody(SP-0023) and DAB(C-0010) staining

— SELECTED CITATIONS —

- **[IF=3.36]** Johnson Chacko L et al. Early appearance of key transcription factors influence the spatiotemporal development of the human inner ear. Cell Tissue Res. 2019 Dec 2. IF ;Human fetuses. 31788757