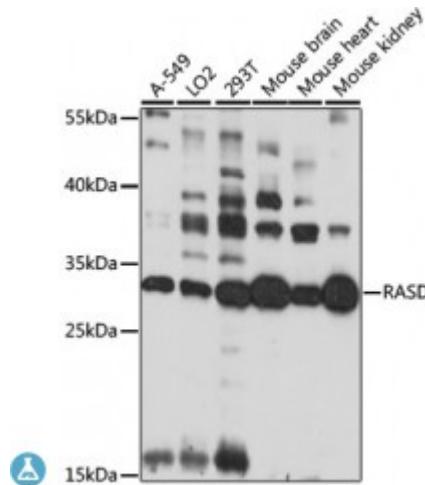


## Anti-RASD1 Antibody



### Description

This gene encodes a member of the Ras superfamily of small GTPases and is induced by dexamethasone. The encoded protein is an activator of G-protein signaling and acts as a direct nucleotide exchange factor for Gi-Go proteins. This protein interacts with the neuronal nitric oxide adaptor protein CAPON, and a nuclear adaptor protein FE65, which interacts with the Alzheimer's disease amyloid precursor protein. This gene may play a role in dexamethasone-induced alterations in cell morphology, growth and cell-extracellular matrix interactions. Epigenetic inactivation of this gene is closely correlated with resistance to dexamethasone in multiple myeloma cells. Alternatively spliced transcript variants encoding different isoforms have been found for this gene.

<b>Model</b>	STJ117646
<b>Host</b>	Rabbit
<b>Reactivity</b>	Human, Mouse
<b>Applications</b>	WB
<b>Immunogen</b>	Recombinant fusion protein containing a sequence corresponding to amino acids 132-281 of human RASD1 (NP_057168.1).
<b>Gene ID</b>	<a href="#">51655</a>
<b>Gene Symbol</b>	<a href="#">RASD1</a>
<b>Dilution range</b>	WB 1:200 - 1:2000
<b>Tissue Specificity</b>	Expressed in a variety of tissues including heart, cardiovascular tissues, brain, placenta, lung, liver, skeletal muscle, kidney, pancreas, gastrointestinal and reproductive tissues

<b>Purification</b>	Affinity purification
<b>Note</b>	For Research Use Only (RUO).
<b>Protein Name</b>	Dexamethasone-induced Ras-related protein 1 Activator of G-protein signaling 1
<b>Molecular Weight</b>	31.642 kDa
<b>Clonality</b>	Polyclonal
<b>Conjugation</b>	Unconjugated
<b>Isotype</b>	IgG
<b>Formulation</b>	PBS with 0.02% sodium azide, 50% glycerol, pH7.3.
<b>Storage Instruction</b>	Store at -20C. Avoid freeze / thaw cycles.
<b>Database Links</b>	<a href="#">HGNC:15828</a> <a href="#">OMIM:605550</a>
<b>Alternative Names</b>	Dexamethasone-induced Ras-related protein 1 Activator of G-protein signaling 1
<b>Function</b>	Small GTPase, Negatively regulates the transcription regulation activity of the APBB1/FE65-APP complex via its interaction with APBB1/FE65 ,
<b>Cellular Localization</b>	Cell membrane
<b>Post-translational Modifications</b>	S-nitrosylation stimulates guanine-nucleotide exchange activity,

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