

## Anti-TRIM11 Antibody

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**Description**

The protein encoded by this gene is a member of the tripartite motif (TRIM) family. The TRIM motif includes three zinc-binding domains, a RING, a B-box type 1 and a B-box type 2, and a coiled-coil region. This protein localizes to the nucleus and the cytoplasm. Its function has not been identified.

<b>Model</b>	STJ115826
<b>Host</b>	Rabbit
<b>Reactivity</b>	Human
<b>Applications</b>	WB
<b>Immunogen</b>	Recombinant fusion protein containing a sequence corresponding to amino acids 267-468 of human TRIM11 (NP_660215.1).
<b>Gene ID</b>	<a href="#">81559</a>
<b>Gene Symbol</b>	<a href="#">TRIM11</a>
<b>Dilution range</b>	WB 1:500 - 1:2000
<b>Tissue Specificity</b>	Ubiquitous
<b>Purification</b>	Affinity purification
<b>Note</b>	For Research Use Only (RUO).
<b>Protein Name</b>	E3 ubiquitin-protein ligase TRIM11
<b>Molecular Weight</b>	52.774 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:16281</a> <a href="#">OMIM:607868</a> <a href="#">Reactome:R-HSA-983168</a>
<b>Alternative Names</b>	E3 ubiquitin-protein ligase TRIM11
<b>Function</b>	E3 ubiquitin-protein ligase that promotes the degradation of insoluble ubiquitinated proteins, including insoluble PAX6, poly-Gln repeat expanded HTT and poly-Ala repeat expanded ARX, Mediates PAX6 ubiquitination leading to proteasomal degradation, thereby modulating cortical neurogenesis, May also inhibit PAX6 transcriptional activity, possibly in part by preventing the binding of PAX6 to its consensus sequences, May contribute to the regulation of the intracellular level of HN (humanin) or HN-containing proteins through the proteasomal degradation pathway, Mediates MED15 ubiquitination leading to proteasomal degradation, May contribute to the innate restriction of retroviruses, Upon overexpression, reduces HIV-1 and murine leukemia virus infectivity, by suppressing viral gene expression, Antiviral activity depends on a functional E3 ubiquitin-protein ligase domain, May regulate TRIM5 turnover via the proteasome pathway, thus counteracting the TRIM5-mediated cross-species restriction of retroviral infection at early stages of the retroviral life cycle,
<b>Cellular Localization</b>	Cytoplasm, Nucleus