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TSG101 (ANT0040R) Rabbit mAb

CatalogNo: ANT8150 Recombinant R

Formulation: PBS,50%glycerol,0.05%Proclin 300,0.05%BSA

Quantity: 100 ug/vial

Host Species Reactivity Applications

Rabbit
 Human, Mouse, Rat,
 WB, IF, IP, ELISA

MW Isotype

• 44kD (Calculated) • IgG,Kappa

44kD (Observed)

Recommended Dilution Ratios

WB 1:1000-1:5000 IF 1:200-1:1000 ELISA 1:5000-1:20000 IP 1:50-1:200,

Storage

Storage* -15°C to -25°C/1 year(Do not lower than -25°C)

Basic Information

Clonality Monoclonal

Clone Number ANT0040R

Immunogen Information

Specificity Endogenous

Target Information

Gene name Protein Name

TSG101

Tumor susceptibility gene 101 protein

Organis	sm	Gene ID	UniProt ID	
	Human	<u>7251</u> ;	<u>Q99816</u> ;	
	Mouse	<u>22088</u> ;	<u>Q61187</u> ;	
	Rat	<u>292925</u> ;	<u>Q6IRE4</u> ;	

Cellular Localization Cytoplasm, Nuclear

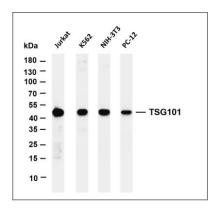
Tissue specificity Heart, brain, placenta, lung, liver, skeletal, kidney and pancreas.

Function

Alternative products: Additional isoforms seem to exist. Several shorter isoforms are detected in primary breast cancers and other tumors, Domain: The coiled coil domain may interact with stathmin., Domain: The UEV domain binds ubiquitin and P-[ST]-A-P peptide motif independently., Domain: The UEV domain is required for the interaction of the complex with ubiquitin. It also mediates the interaction with ANTAP/PSAP motifs of HIV-1 P6 protein and human spumaretrovirus Gag protein., Function: Component of the ESCRT-I complex, a regulator of vesicular trafficking process. Binds to ubiquitinated cargo proteins and is required for the sorting of endocytic ubiquitinated cargos into multivesicular bodies (MVBs). Mediates the association between the ESCRT-0 and ESCRT-I complex. Required for completion of cytokinesis; the function requires CEP55. May be involved in cell growth and differentiation. Acts as a negative growth regulator. Involved in the budding of many viruses through an interaction with viral proteins that contain a late-budding motif P-[ST]-A-P. This interaction is essential for viral particle budding of numerous retroviruses., PTM: Monoubiquitinated at multiple sites by LRSAM1. Ubiquitination inactivates it, possibly by regulating its shuttling between an active membrane-bound protein and an inactive soluble form., similarity: Belongs to the ubiquitin-conjugating enzyme family. UEV subfamily., similarity: Contains 1 SB (steadiness box) domain., similarity: Contains 1 UEV (ubiquitin E2 variant) domain.,subcellular location:Mainly cytoplasmic. Membraneassociated when active and soluble when inactive. Depending on the stage of the cell cycle, detected in the nucleus. Colocalized with CEP55 in the midbody during cytokinesis., subunit: Component of the ESCRT-I complex (endosomal sorting complex required for transport I) which consists of TSG101, VPS28, a VPS37 protein (VPS37A to -D) and a FAM125/MVB12 protein (FAM125A or -B) in a 1:1:1:1 stoechiometry. Interacts with VPS37A, VPS37B and VPS37C. Interacts with ubiquitin, stathmin, GMCL, DMAP1 and AATF

(By similarity). Interacts with HGS; the interaction mediates the association with the ESCRT-0 complex. Interacts with GGA1 and GGA3. Interacts (via UEV domain) with PDCD6IP/AIP1. Interacts with VPS28, SNF8 and VPS36. Self-associates. Interacts with FAM125A/MVB12A; the association appears to be mediated by the TSG101-VPS37 binary subcomplex. Interacts with VPS37D. Interacts with LRSAM1. Interacts with CEP55; the interaction is required for cytokinesis but not for viral budding. Interacts with HIV-1 p6. Interacts with human spumavirus Gag. Interacts with HTLV-1 Gag. Interacts with Ebola virus

VP40. Interacts with EIAV p9; the interaction has been shown in vitro., tissue specificity: Heart, brain, placenta, lung, liver, skeletal, kidney and pancreas.,



Validation Data

Various whole cell lysates were separated by 4-20% SDS-PAGE, and the membrane was blotted with anti-TSG101 (ANT0040R) antibody. The HRPconjugated Goat anti-Rabbit IgG(H+L) antibody was used to detect the antibody. Lane 1: Jurkat Lane 2: K562 Lane 3: NIH-3T3 Lane 4: PC-12

Predicted band size: 44kDa Observed band size: 44kDa

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