



Applications

WB,IF,IP,ELISA

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53BP1 (ANT0056R) Rabbit mAb

CatalogNo: ANT8293 Recombinant R

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

Quantity: 100 ug/vial

Host SpeciesRabbitReactivityHuman,

MW Isotype
• 214kD (Calculated) • IgG,Kappa

450kD (Observed)

Recommended Dilution Ratios

WB 1:2000-1:10000 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 ANT0056R

Immunogen Information

Specificity Endogenous

Target Information

Gene name TP53BP1

| Organism | Gene ID | | UniProt ID | | |
|----------|---------|---------------|------------|-----------------|---------|
| | Human | <u>7158</u> ; | | <u>Q12888</u> ; | |
| | Mouse | | 27223; | | P70399; |

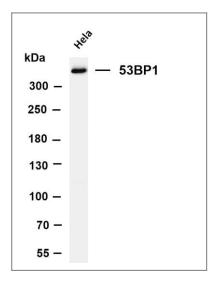
Cellular Localization Nucleus

Tissue specificity Cerebellum, Cervix, Epithelium, Myeloid leukemia cell, Skeletal muscle,

Function

Function: May have a role in checkpoint signaling during mitosis (By similarity). Enhances TP53-mediated transcriptional activation. Plays a role in the response to DNA damage., ANTM: Asymmetrically dimethylated on Arg residues by PRMT1. Methylation is required for DNA binding., PTM: Phosphorylated at basal level in the absence of DNA damage. Hyper-phosphorylated in an ATM-dependent manner in response to DNA damage induced by ionizing radiation. Hyper-phosphorylated in an ATR-dependent manner in response to DNA damage induced by UV irradiation., similarity: Contains 2 BRCT domains., subcellular location: Associated with kinetochores. Both nuclear and cytoplasmic in some cells. Recruited to sites of DNA damage, such as double stand breaks. Methylation of histone H4 at 'Lys-20' is required for efficient localization to double strand breaks., subunit: Interacts with IFI202A (By similarity). Binds to the central domain of TP53/p53. May form homo-oligomers. Interacts with DCLRE1C. Interacts with histone H2AFX and this requires phosphorylation of H2AFX on 'Ser-139'. Interacts with histone H4 that has been dimethylated at 'Lys-20'. Has low affinity for histone H4 containing monomethylated 'Lys-20'. Does not bind histone H4 containing unmethylated or trimethylated 'Lys-20'. Has low affinity for histone H3 that has been dimethylated on 'Lys-79'. Has very low affinity for histone H3 that has been monomethylated on 'Lys-79' (in vitro). Does not bind unmethylated histone H3.,

Validation Data



Various whole cell lysates were separated by 4-8% SDS-PAGE, and the membrane was blotted with anti-53BP1 (ANT0056R) antibody. The HRPconjugated Goat anti-Rabbit IgG(H + L) antibody was used to detect the antibody. Lane 1: Hela Predicted band size: 214kDa Observed band size: 450kDa