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Mouse Monoclonal Antibody 53BP1 conjugated to Sepharose Beads

CatalogNo: ANT8293-S

Size 200ul

Storage Store at 4 °C for frequent use

Description

This Antagene antibody is immobilized via covalent binding of primary amino groups to N-hydroxysuccinimide (NHS)-activated sepharose beads. It is useful for immunoprecipitation assays.

53BP1 (ANTO056R) Rabbit mAb

Formulation: 50% slurry in PBS pH 7.2 with 0.01mg NaN3a3 preservative.

Host Species Reactivity Applications
• Rabbit • Human, • WB,IF,IP,ELISA

MW Isotype
• 214kD (Calculated) • IgG,Kappa
450kD (Observed)

#### Recommended Dilution Ratios

ΙP

#### **Basic Information**

**Clonality** Monoclonal

Clone Number ANT0056R

## Immunogen Information

**Specificity** Endogenous

# **Target Information**

Gene name
Protein Name

TP53BP1

Tumor suppressor p53-binding protein 1

| Organism | Gene ID |               | UniProt ID     |            |
|----------|---------|---------------|----------------|------------|
|          | Human   | <u>7158</u> ; | Q12888         | <u>3</u> ; |
|          | Mouse   |               | <u>27223</u> ; | P70399;    |

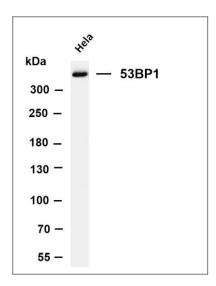
Cellular Nucleus
Localization

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

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