

## 53BP1 (Phospho Ser1618) Rabbit pAb

CatalogNo: YP1253

### Key Features

#### Host Species

- Rabbit

#### Reactivity

- Human, Rat

#### Applications

- WB

#### MW

- 213kD (Observed)

#### Isotype

- IgG

### Storage

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

**Formulation** Liquid in PBS containing 50% glycerol, 0.5% BSA and 0.02% sodium azide.

### Recommended Dilution Ratios

**WB 1:1000-2000**

### Basic Information

**Clonality** Polyclonal

### Immunogen Information

**Immunogen** Synthesized phospho peptide around human 53BP1 (Ser1618)

**Specificity** This antibody detects endogenous levels of Human Rat 53BP1 (phospho-Ser1618)

### Target Information

**Gene name** TP53BP1

**Protein Name** 53BP1 (Ser1618)

Organism	Gene ID	UniProt ID
Human	<a href="#">7158;</a>	<a href="#">Q12888;</a>
Mouse	<a href="#">27223;</a>	<a href="#">P70399;</a>

**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.,PTM: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 strand 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

## Contact information

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