

**53BP1 (Phospho Ser1618) rabbit pAb**

<b>Catalog No :</b>	YP1253
<b>Reactivity :</b>	Human;Rat
<b>Applications :</b>	WB
<b>Target :</b>	53BP1
<b>Fields :</b>	>>NOD-like receptor signaling pathway
<b>Gene Name :</b>	TP53BP1
<b>Protein Name :</b>	53BP1 (Ser1618)
<b>Human Gene Id :</b>	7158
<b>Human Swiss Prot No :</b>	Q12888
<b>Mouse Gene Id :</b>	27223
<b>Mouse Swiss Prot No :</b>	P70399
<b>Immunogen :</b>	Synthesized phospho peptide around human 53BP1 (Ser1618)
<b>Specificity :</b>	This antibody detects endogenous levels of Human Rat 53BP1 (phospho-Ser1618)
<b>Formulation :</b>	Liquid in PBS containing 50% glycerol, 0.5% BSA and 0.02% sodium azide.
<b>Source :</b>	Polyclonal, Rabbit,IgG
<b>Dilution :</b>	WB 1:1000-2000
<b>Purification :</b>	The antibody was affinity-purified from rabbit serum by affinity-chromatography using specific immunogen.
<b>Concentration :</b>	1 mg/ml

<b>Storage Stability :</b>	-15 °C to -25 °C/1 year(Do not lower than -25 °C)
<b>Observed Band :</b>	213kD
<b>Function :</b>	<p>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 th</p>
<b>Subcellular Location :</b>	<p>Nucleus . Chromosome . Chromosome, centromere, kinetochore . Localizes to the nucleus in absence of DNA damage (PubMed:28241136). Following DNA damage, recruited to sites of DNA damage, such as double strand breaks (DSBs): recognizes and binds histone H2A monoubiquitinated at 'Lys-15' (H2AK15Ub) and histone H4 dimethylated at 'Lys-20' (H4K20me2), two histone marks that are present at DSBs sites (PubMed:23333306, PubMed:23760478, PubMed:24703952, PubMed:28241136, PubMed:17190600). Associated with kinetochores during mitosis (By similarity). .</p>
<b>Expression :</b>	Cerebellum,Cervix,Epithelium,Myeloid leukemia cell,Skeletal muscle,
<b>Sort :</b>	1528
<b>No4 :</b>	1

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