

**Product Datasheet** 

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## Phospho-RPA32/RPA2 (Ser33) Rabbit Polyclonal Antibody

Catalog #: EAB10888

Host/Isotype	Clonality	Applications	MW (kDa)	Reactivity	
Rabbit IgG	Polyclonal	WB, IHC-P	29	Human, Mouse, Rat	

## **Applications Dilutions**

The application notes include recommended starting dilutions; optimal dilutions/concentrations should be determined by the end user.

**WB**(Western Blotting) 1:500-2000 **IHC-P**(Immunohistochemistry-Paraffin) 1:50-300

## **Product Information**

Conjugate Unconjugate

Specificity

Phospho-RPA32/RPA2 (Ser33) Rabbit Polyclonal Antibody detects endogenous levels of

RPA32/RPA2 protein only when phosphorylated at Ser33.

**Purification** Affinity purification

Concentration1mg/mlFormatLiquid

Formulation In PBS, pH 7.4, Containing 0.02% sodium azide, 0.5% BSA and 50% Glycerol

**Shipping** Gel Pack

Storage Storag

Aliquots may be stored at +4°C for 1-2 weeks

 UniProt ID
 P15927

 Entrez-Gene Id
 6118

## **Product Description**

This gene encodes a subunit of the heterotrimeric Replication Protein A (RPA) complex, which binds to single-stranded DNA (ssDNA), forming a nucleoprotein complex that plays an important role in DNA metabolism, being involved in DNA replication, repair, recombination, telomere maintenance, and co-ordinating the cellular response to DNA damage through activation of the ataxia telangiectasia and Rad3-related protein (ATR) kinase. The RPA complex protects single-stranded DNA from nucleases, prevents formation of secondary structures that would interfere with repair, and co-ordinates the recruitment and departure of different genome maintenance factors. The heterotrimeric complex has two different modes of ssDNA binding, a low-affinity and high-affinity mode, determined by which oligonucleotide/oligosaccharide-binding (OB) domains of the complex are utilized, and differing in the length of DNA bound. This subunit contains a single OB domain that participates in high-affinity DNA binding and also contains a winged helix domain at its carboxy terminus, which interacts with many genome maintenance protein. Post-translational modifications of the RPA complex also plays a role in coordinating different damage response pathways.