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<b>Product name:</b>	Histone H2A.X (phospho Ser139) Rabbit Polyclonal Antibody
<b>Cat number:</b>	ABN04775
<b>Conjugate:</b>	Unconjugated
<b>Size:</b>	100µL
<b>Clone:</b>	POLY
<b>Concentration:</b>	1mg/ml
<b>Host:</b>	Rabbit
<b>Isotype:</b>	IgG
<b>Immunogen:</b>	H2A.X around the phosphorylation site of Ser139. AA range:94-143
<b>Reactivity:</b>	Human,Mouse,Rat,Hamster
<b>Applications:</b>	WB 1:500-1:2000,IHC 1:100-1:300,ICC/IF 1:50-1:200,ELISA 1:5000-1:20000
<b>Molecular Weight:</b>	19kDa
<b>Purification:</b>	Affinity purification
<b>Form:</b>	liquid
<b>Buffer:</b>	Liquid in PBS containing 50% glycerol, 0.5% protective protein and 0.02% New type preservative N.
<b>Storage:</b>	Store at 4°C short term. Aliquot and store at -20°C long term. Avoid freeze/thaw cycles.
<b>Synonyms:</b>	H2AFX; H2AX; Histone H2A.x; H2a/x
<b>Source:</b>	Rabbit
<b>Background:</b>	<p>Histones are basic nuclear proteins that are responsible for the nucleosome structure of the chromosomal fiber in eukaryotes. Two molecules of each of the four core histones (H2A, H2B, H3, and H4) form an octamer, around which approximately 146 bp of DNA is wrapped in repeating units, called nucleosomes. The linker histone, H1, interacts with linker DNA between nucleosomes and functions in the compaction of chromatin into higher order structures. This gene encodes a replication-independent histone that is a member of the histone H2A family, and generates two transcripts through the use of the conserved stem-loop termination motif, and the polyA addition motif. [provided by RefSeq, Oct 2015],developmental stage:Synthesized in G1 as well as in S-phase.,domain:The [ST]-Q motif constitutes a recognition sequence for kinases from the PI3/PI4-kinase family.,function:Variant histone H2A which replaces conventional H2A in a subset of nucleosomes. Nucleosomes wrap and compact DNA into chromatin, limiting DNA accessibility to the cellular machineries which require DNA as a template. Histones thereby play a central role in transcription regulation, DNA repair,</p>

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DNA replication and chromosomal stability. DNA accessibility is regulated via a complex set of post-translational modifications of histones, also called histone code, and nucleosome remodeling. Required for checkpoint-mediated arrest of cell cycle progression in response to low doses of ionizing radiation and for efficient repair of DNA double strand breaks (DSBs) specifically when modified by C-terminal phosphorylation. PTM: Monoubiquitination of Lys-120 (H2AXK119ub) by RING1 and RNF2/RING2 complex gives a specific tag for epigenetic transcriptional repression. Following DNA double-strand breaks (DSBs), it is ubiquitinated through 'Lys-63' linkage of ubiquitin moieties by the E2 ligase UBE2N and the E3 ligases RNF8 and RNF168, leading to the recruitment of repair proteins to sites of DNA damage. Monoubiquitination and ionizing radiation-induced 'Lys-63'-linked ubiquitination are distinct events. PTM: Phosphorylated on Ser-140 (to form gamma-H2AFX or H2AX139ph) in response to DNA double strand breaks (DSBs) generated by exogenous genotoxic agents and by stalled replication forks, and may also occur during meiotic recombination events and immunoglobulin class switching in lymphocytes. Phosphorylation can extend up to several thousand nucleosomes from the actual site of the DSB and may mark the surrounding chromatin for recruitment of proteins required for DNA damage signaling and repair. Widespread phosphorylation may also serve to amplify the damage signal or aid repair of persistent lesions. Phosphorylation of Ser-140 (H2AX139ph) in response to ionizing radiation is mediated by both ATM and PRKDC while defects in DNA replication induce Ser-140 phosphorylation (H2AX139ph) subsequent to activation of ATR and PRKDC. Dephosphorylation of Ser-140 by PP2A is required for DNA DSB repair. In meiosis, Ser-140 phosphorylation (H2AX139ph) may occur at synaptonemal complexes during leptotene as an ATM-dependent response to the formation of programmed DSBs by SPO11. Ser-140 phosphorylation (H2AX139ph) may subsequently occur at unsynapsed regions of both autosomes and the XY bivalent during zygotene, downstream of ATR and BRCA1 activation. Ser-140 phosphorylation (H2AX139ph) may also be required for transcriptional repression of unsynapsed chromatin and meiotic sex chromosome inactivation (MSCI), whereby the X and Y chromosomes condense in pachytene to form the heterochromatic XY-body. During immunoglobulin class switch recombination in lymphocytes, Ser-140 phosphorylation (H2AX139ph) may occur at sites of DNA-recombination subsequent to activation of the activation-induced cytidine deaminase AICDA. Phosphorylation at Tyr-143 (H2AXY142ph) by BAZ1B/WSTF determines the relative recruitment of either DNA repair or pro-apoptotic factors. Phosphorylation at Tyr-143 (H2AXY142ph) favors the recruitment of APBB1/FE65 and pro-apoptosis factors such as MAPK8/JNK1, triggering apoptosis. In contrast, dephosphorylation of Tyr-143 by EYA proteins (EYA1, EYA2, EYA3 or EYA4) favors the recruitment of MDC1-containing DNA repair complexes to the tail of phosphorylated Ser-140 (H2AX139ph). similarity: Belongs to the histone H2A family. subunit: The nucleosome is a histone octamer containing two molecules each of H2A, H2B, H3 and H4 assembled in one

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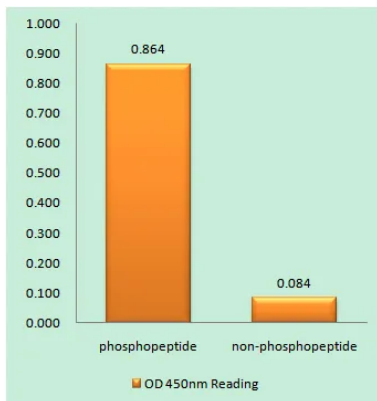
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H3-H4 heterotetramer and two H2A-H2B heterodimers. The octamer wraps approximately 147 bp of DNA. Interacts with numerous proteins required for DNA damage signaling and repair when phosphorylated on Ser-140. These include MDC1, TP53BP1, BRCA1 and the MRN complex, composed of MRE11A, RAD50, and NBN. Interaction with the MRN complex is mediated at least in part by NBN. Also interacts with DHX9/NDHII when phosphorylated on Ser-140.,

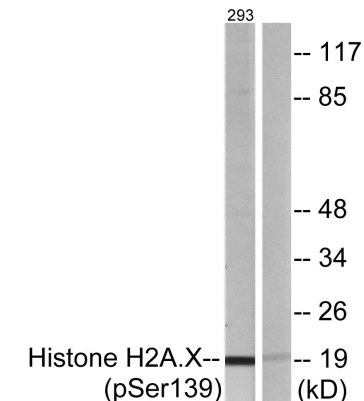
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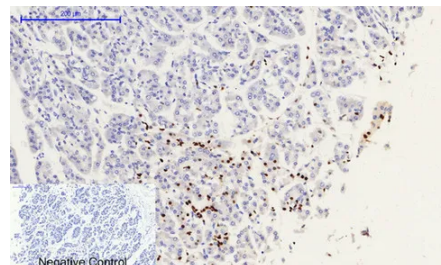
Web-site: <https://immunologicalsciences.com> - E-mail: [info@immunologicalsciences.com](mailto:info@immunologicalsciences.com)



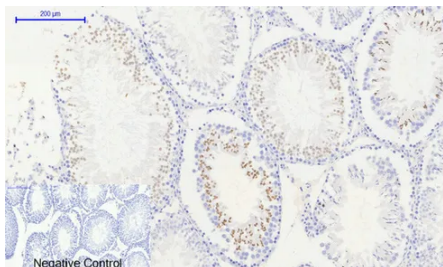
Enzyme-Linked Immunosorbent Assay Phospho-ELISA for Immunogen Phosphopeptide Phospho-left and Non-Phosphopeptide Phospho-right, using Histone H2A.X Phospho-Ser139 Antibody



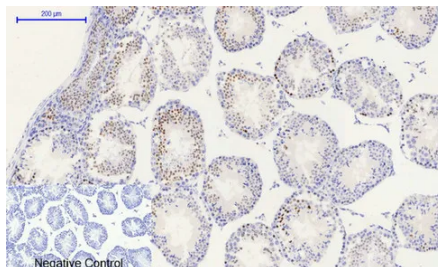
Western blot analysis of lysates from 293 cells treated with heat shock, using Histone H2A.X Phospho-Ser139 Antibody. The lane on the right is blocked with the phospho peptide.



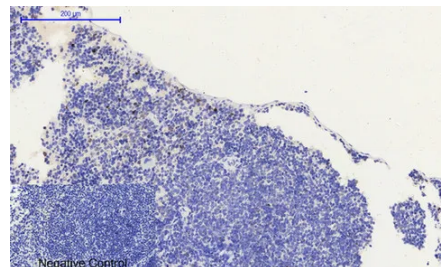
Immunohistochemical analysis of paraffin-embedded Human-stomach-cancer tissue. 1, Histone H2A.X phospho Ser139 Polyclonal Antibody was diluted at 1:200 4°C, overnight. 2, Sodium citrate pH 6.0 was used for antibody retrieval >98°C, 20min. 3, Secondary antibody was diluted at 1:200 room temperature, 30min. Negative control was used by secondary antibody only.



Immunohistochemical analysis of paraffin-embedded Rat-testis tissue. 1, Histone H2A.X phospho Ser139 Polyclonal Antibody was diluted at 1:200 4°C, overnight. 2, Sodium citrate pH 6.0 was used for antibody retrieval >98°C, 20min. 3, Secondary antibody was diluted at 1:200 room temperature, 30min. Negative control was used by secondary antibody only.



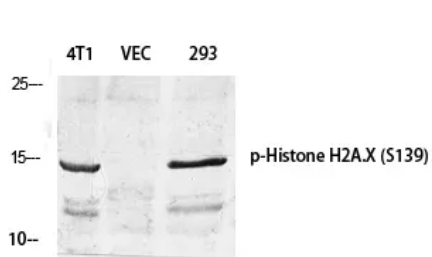
Immunohistochemical analysis of paraffin-embedded Mouse-testis tissue. 1, Histone H2A.X phospho Ser139 Polyclonal Antibody was diluted at 1:200 4°C, overnight. 2, Sodium citrate pH 6.0 was used for antibody retrieval >98°C, 20min. 3, Secondary antibody was diluted at 1:200 room temperature, 30min. Negative control was used by secondary antibody only.



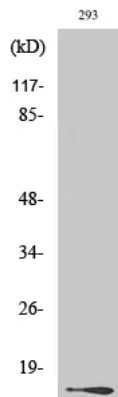
Immunohistochemical analysis of paraffin-embedded Mouse-spleen tissue. 1, Histone H2A.X phospho Ser139 Polyclonal Antibody was diluted at 1:200 4°C, overnight. 2, Sodium citrate pH 6.0 was used for antibody retrieval >98°C, 20min. 3, Secondary antibody was diluted at 1:200 room temperature, 30min. Negative control was used by secondary antibody only.

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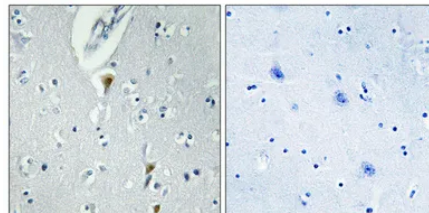
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Western Blot analysis of various cells using Phospho-Histone H2A.X S139 Polyclonal Antibody diluted at 1500.



Western Blot analysis of 293 cells using Phospho-Histone H2A.X S139 Polyclonal Antibody diluted at 1500.



Immunohistochemical analysis of paraffin-embedded Human brain. Antibody was diluted at 1:1004°, overnight. High-pressure and temperature Tris-EDTA, pH8.0 was used for antigen retrieval. Negative contrl right obtained from antibody was pre-absorbed by immunogen peptide.

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