

### Human gamma H2A.X (phospho S139) peptide ab15645

**3 References**   [1 图像](#)

#### 描述

**产品名称** 人gamma H2A.X (phospho S139)多肽  
**纯度** > 90 % HPLC.

**Accession** **P16104**  
**无动物成分** No  
**性质** Synthetic  
**种属** Human  
**修饰** phospho S139

#### 技术指标

Our **Abpromise guarantee** covers the use of **ab15645** in the following tested applications.

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

**应用** Blocking - Blocking peptide for Anti-gamma H2A.X (phospho S139) antibody (**ab2893**)

**形式** Lyophilized

**补充说明**

- First try to dissolve a small amount of peptide in either water or buffer. The more charged residues on a peptide, the more soluble it is in aqueous solutions.
- If the peptide doesn't dissolve try an organic solvent e.g. DMSO, then dilute using water or buffer.
- Consider that any solvent used must be compatible with your assay. If a peptide does not dissolve and you need to recover it, lyophilise to remove the solvent.
- Gentle warming and sonication can effectively aid peptide solubilisation. If the solution is cloudy or has gelled the peptide may be in suspension rather than solubilised.
- Peptides containing cysteine are easily oxidised, so should be prepared in solution just prior to use.

#### 制备和贮存

**稳定性和存储** Shipped at 4°C. Store at -20°C.  
Information available upon request.

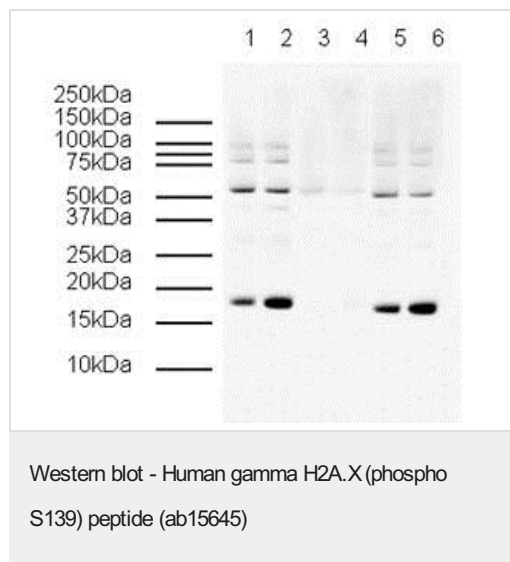
## 常规信息

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功能	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, 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.
序列相似性	Belongs to the histone H2A family.
发展阶段	Synthesized in G1 as well as in S-phase.
结构域	The [ST]-Q motif constitutes a recognition sequence for kinases from the PI3/PI4-kinase family.
翻译后修饰	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 occurs 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). 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.
细胞定位	Nucleus. Chromosome.

## 图片

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**All lanes :** Anti-gamma H2A.X (phospho S139) antibody (**ab2893**) at 1/500 dilution

**Lane 1 :** Control HeLa (Human epithelial cell line from cervix adenocarcinoma) whole cell lysate Histone preparation

**Lane 2 :** Colcemid treated HeLa whole cell lysate Histone preparation

**Lane 3 :** Control HeLa (Human epithelial cell line from cervix adenocarcinoma) whole cell lysate Histone preparation with Human gamma H2A.X (phospho S139) peptide (ab15645) at 1 µg

**Lane 4 :** Colcemid treated HeLa whole cell lysate Histone preparation with Human gamma H2A.X (phospho S139) peptide (ab15645) at 1 µg

**Lane 5 :** Control HeLa (Human epithelial cell line from cervix adenocarcinoma) whole cell lysate Histone preparation with Human Histone H2A.X (unmodified ) peptide (**ab15646**) at 1 µg

**Lane 6 :** Colcemid treated HeLa whole cell lysate Histone preparation with Human Histone H2A.X (unmodified ) peptide (**ab15646**) at 1 µg

### Secondary

**All lanes :** Goat Anti-Rabbit IgG H&L (HRP) (**ab6721**) at 1/5000 dilution

**Please note:** All products are "FOR RESEARCH USE ONLY. NOT FOR USE IN DIAGNOSTIC PROCEDURES"

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