

H2AZ Polyclonal Antibody

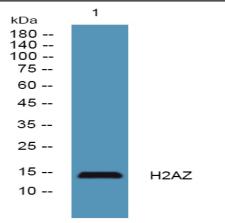
Catalog No :	YN2795
Reactivity :	Human;Rat;Mouse;
Applications :	WB;ELISA
Target :	Histone H2A
Fields :	>>Necroptosis;>>Neutrophil extracellular trap formation;>>Alcoholism;>>Systemic lupus erythematosus
Gene Name :	H2AFZ H2AZ
Protein Name :	Histone H2A.Z (H2A/z)
Human Gene Id :	3015
Human Swiss Prot	P0C0S5
No : Mouse Swiss Prot	P0C0S6
No : Rat Swiss Prot No :	P0C0S7
Immunogen :	Synthesized peptide derived from part region of human protein AA range: 1-80
Specificity :	H2AZ Polyclonal Antibody detects endogenous levels of protein.
Formulation :	Liquid in PBS containing 50% glycerol, and 0.02% sodium azide.
Source :	Polyclonal, Rabbit,IgG
Dilution :	WB 1:500-2000 ELISA 1:5000-20000
Purification :	The antibody was affinity-purified from rabbit antiserum by affinity- chromatography using epitope-specific immunogen.
Concentration :	1 mg/ml



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Storage Stability :	-15°C to -25°C/1 year(Do not lower than -25°C)
Observed Band :	14kD
Cell Pathway :	Systemic lupus erythematosus;
Background :	Histones are basic nuclear proteins that are responsible for the nucleosome structure of the chromosomal fiber in eukaryotes. Nucleosomes consist of approximately 146 bp of DNA wrapped around a histone octamer composed of pairs of each of the four core histones (H2A, H2B, H3, and H4). The chromatin fiber is further compacted through the interaction of a linker histone, H1, with the DNA between the nucleosomes to form higher order chromatin structures. This gene encodes a replication-independent member of the histone H2A family that is distinct from other members of the family. Studies in mice have shown that this particular histone H2A leads to embryonic lethality. [provided by RefSeq, Jul 2008],
Function :	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, 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. May be involved in the formation of constitutive heterochromatin. May be required for chromosome segregation during cell division.,mass spectrometry:Monoisotopic, not modified PubMed:16457589,PTM:Acetylated on Lys-5, Lys-8 and Lys-12 during interphase. Acetylation disappears at mitosis.,PTM:Monoubiquitination of Lys-122 gives a specific tag for epigenetic transcriptional repression.,PTM:Not phosphor
Subcellular Location :	Nucleus. Chromosome.
Expression :	Brain,Epithelium,Skeletal muscle,Uterus,

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Western blot analysis of lysates from A431 cells, primary antibody was diluted at 1:1000, 4° over night