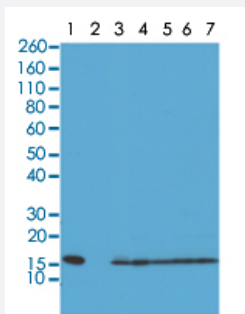


RecomAb™

Histone H2A monoclonal antibody, clone RM225

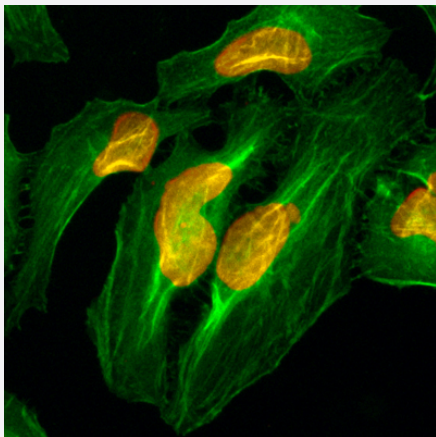
Catalog # MAB15118 Size 100 ug

Applications



Western Blot

Western blot analysis of Lane 1: recombinant Histone H2A, Lane 2: recombinant Histone H2B, Lane 3: HeLa, Lane 4: A375, Lane 5: SK-MEL-2, Lane 6: A431, Lane 7: K562 whole cell lysates with Histone H2A monoclonal antibody, clone RM225 (Cat # MAB15118) at 0.5 ug/mL working concentration.



Immunocytochemistry

Immunocytochemistry staining of HeLa cells with Histone H2A monoclonal antibody, clone RM225 (Cat # MAB15118) (Red). Actin filaments were labeled with fluorescein phalloidin (Green).

Specification

Product Description	Rabbit recombinant monoclonal antibody raised against of human histone H2A.
Antibody Species	Rabbit
Immunogen	Original antibody is raised against a synthetic peptide corresponding to C-terminus of human Histone H2A.
Sequence	N/A

Reactivity	Human
Form	Liquid
Purification	Protein A affinity purification
Isotype	IgG
Recommend Usage	ELISA (0.2-1 ug/mL) Immunocytochemistry (1-2 ug/mL) Multiplex (0.2-1 ug/mL) Western Blot (0.5-2 ug/mL) The optimal working dilution should be determined by the end user.
Storage Buffer	In PBS (50% glycerol, 1% BSA, 0.09% sodium azide)
Storage Instruction	Store at -20°C. Aliquot to avoid repeated freezing and thawing.
Note	This product contains sodium azide: a POISONOUS AND HAZARDOUS SUBSTANCE which should be handled by trained staff only.

Applications

- Western Blot

Western blot analysis of Lane 1: recombinant Histone H2A, Lane 2: recombinant Histone H2B, Lane 3: HeLa, Lane 4: A375, Lane 5: SK-MEL-2, Lane 6: A431, Lane 7: K562 whole cell lysates with Histone H2A monoclonal antibody, clone RM225 (Cat # MAB15118) at 0.5 ug/mL working concentration.

- Immunocytochemistry

Immunocytochemistry staining of HeLa cells with Histone H2A monoclonal antibody, clone RM225 (Cat # MAB15118) (Red). Actin filaments was labeled with fluorescein phalloidin (Green).

- Enzyme-linked Immunoabsorbent Assay

Gene Info — HIST1H2AE

Entrez GeneID	3012
Protein Accession#	P04908 ; Q93077 ; Q7L7L0 ; Q6FI13 ; P20671 ; P0C0S8
Gene Name	HIST1H2AE
Gene Alias	H2A.1, H2A.2, H2A/a, H2AFA

Gene Description	histone cluster 1, H2ae
Omim ID	602786
Gene Ontology	Hyperlink
Gene Summary	<p>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 is intronless and encodes a member of the histone H2A family. Transcripts from this gene lack polyA tails; instead, they contain a palindromic termination element. This gene is found in the large histone gene cluster on chromosome 6p22-p21.3. [provided by RefSeq]</p>
Other Designations	H2A histone family, member A histone 1, H2ae histone H2AE

Gene Info — HIST1H2AD

Entrez GeneID	3013
Protein Accession#	P04908 ; Q93077 ; Q7L7L0 ; Q6FI13 ; P20671 ; P0C0S8
Gene Name	HIST1H2AD
Gene Alias	H2A.3, H2A/g, H2AFG, HIST1H3D
Gene Description	histone cluster 1, H2ad
Omim ID	602792
Gene Ontology	Hyperlink
Gene Summary	<p>Histones are basic nuclear proteins that are responsible for the nucleosome structure of the chromosomal fiber in eukaryotes. This structure consists of approximately 146 bp of DNA wrapped around a nucleosome, an 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 is intronless and encodes a member of the histone H2A family. Transcripts from this gene lack poly A tails; instead, they contain a palindromic termination element. This gene is found in the large histone gene cluster on chromosome 6p22-p21.3. [provided by RefSeq]</p>
Other Designations	H2A histone family, member G OTTHUMP00000016153 histone 1, H2ad histone H2AD

Gene Info — HIST1H2AI

Entrez GeneID	8329
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Protein Accession#	P04908; Q93077; Q7L7L0; Q6FI13; P20671; P0C0S8
Gene Name	HIST1H2AI
Gene Alias	FLJ92027, H2A/c, H2AFC
Gene Description	histone cluster 1, H2ai
Omim ID	602787
Gene Ontology	Hyperlink
Gene Summary	<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 is intronless and encodes a member of the histone H2A family. Transcripts from this gene lack polyA tails but instead contain a palindromic termination element. This gene is found in the small histone gene cluster on chromosome 6p22-p21.3. [provided by RefSeq]</p>
Other Designations	H2A histone family, member C OTTHUMP00000016183 histone 1, H2ai

Gene Info — HIST1H2AC

Entrez GeneID	8334
Protein Accession#	P04908; Q93077; Q7L7L0; Q6FI13; P20671; P0C0S8
Gene Name	HIST1H2AC
Gene Alias	H2A/I, H2AFL, MGC99519, dJ221C16.4
Gene Description	histone cluster 1, H2ac
Omim ID	602794
Gene Ontology	Hyperlink
Gene Summary	<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 is intronless and encodes a member of the histone H2A family. Transcripts from this gene lack polyA tails but instead contain a palindromic termination element. This gene is found in the large histone gene cluster on chromosome 6. [provided by RefSeq]</p>
Other Designations	H2A histone family, member L OTTHUMP00000016145 histone 1, H2ac histone H2AC

Gene Info — HIST2H2AA4

Entrez GeneID [723790](#)

Protein Accession# [P04908; Q93077; Q7L7L0; Q6FI13; P20671; P0C0S8](#)

Gene Name HIST2H2AA4

Gene Alias H2A/R

Gene Description histone cluster 2, H2aa4

Gene Ontology [Hyperlink](#)

Gene Summary

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 is intronless and encodes a member of the histone H2A family. Transcripts from this gene lack polyA tails but instead contain a palindromic termination element. This gene is found in a histone cluster on chromosome 1. This gene is one of four histone genes in the cluster that are duplicated; this record represents the telomeric copy. [provided by RefSeq]

Other Designations OTTHUMP00000013922|histone 2, H2aa4|histone H2A/r

Pathway

- [Systemic lupus erythematosus](#)
- [Systemic lupus erythematosus](#)
- [Systemic lupus erythematosus](#)
- [Systemic lupus erythematosus](#)
- [Systemic lupus erythematosus](#)