## abcam

### Product datasheet

# Histone H4 (tri-methyl K20) Quantification Kit (Colorimetric) abl 15097

#### Overview

Product name Histone H4 (tri-methyl K20) Quantification Kit (Colorimetric)

**Detection method**Colorimetric

Sample type Tissue, Adherent cells, Suspension cells

Sensitivity 5 ng/well

Range 20 ng/well - 2000 ng/well

Assay time 2h 30m

Species reactivity Reacts with: Mouse, Human

Predicted to work with: Mammals

Product overview Methylation of histone H4 at lysine 20 is catalyzed by SET9 and SUV4.20h among other methyl

transferases in mammalian cells. H4 (tri-methyl K20) acts as a passive feature or structure determinant for chromatin degradation and release, as well as being an epigenetic marker of early apoptosis. H4 (tri-methyl K20) is also considered as a common hallmark of human cancer.

Histone H4 (tri-methyl K20) Quantification Kit (Colorimetric) (ab115097) enables the user to measure global tri-methylation of histone H4K20. The global H4K20 tri-methylation can also be changed by inhibiton or activation of HMTs, making quantitative detection of global tri-methyl

histone H4K20 a useful tool for better understanding epigenetic regulation of gene

activation/repression. ab115097 is suitable for use with a variety of mammalian cells including

fresh and frozen tissues, and cultured adherent and suspension cells.

**Platform** Microplate reader

#### **Properties**

**Storage instructions** Please refer to protocols.

10X Wash Buffer 1 x 10ml 1 x 20ml	Components	Identifier	48 tests	96 tests
9 Wall Cample String (with Frame)	10X Wash Buffer		1 x 10ml	1 x 20ml
o-vveii Sampie Surps (with Frame) 4 units 9 units	8-Well Sample Strips (with Frame)		4 units	9 units
8-Well Standard Control Strips Green Ringed 2 units 3 units	8-Well Standard Control Strips	Green Ringed	2 units	3 units

Components	Identifier	48 tests	96 tests
Antibody Buffer		1 x 6ml	1 x 12ml
Color Developer		1 x 5ml	1 x 10ml
Detecting Antibody, 1 mg/mL		1 x 5µl	1 x 10µl
Signal Enhancer		1 x 120µl	1 x 240µl
Signal Report Solution		1 x 5µl	1 x 10µl
Standard Control, 100 μg/mL		1 x 10µl	1 x 20µl
Stop Solution		1 x 3ml	1 x 6ml

#### **Function**

Core component of nucleosome. 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.

#### Sequence similarities

## Post-translational modifications

Belongs to the histone H4 family.

Acetylation at Lys-6 (H4K5ac), Lys-9 (H4K8ac), Lys-13 (H4K12ac) and Lys-17 (H4K16ac) occurs in coding regions of the genome but not in heterochromatin.

Citrullination at Arg-4 (H4R3ci) by PADI4 impairs methylation.

Monomethylation and asymmetric dimethylation at Arg-4 (H4R3me1 and H4R3me2a, respectively) by PRMT1 favors acetylation at Lys-9 (H4K8ac) and Lys-13 (H4K12ac).

Demethylation is performed by JMJD6. Symmetric dimethylation on Arg-4 (H4R3me2s) by the

PRDM1/PRMT5 complex may play a crucial role in the germ-cell lineage.

Monomethylated, dimethylated or trimethylated at Lys-21 (H4K20me1, H4K20me2, H4K20me3).

Monomethylation is performed by SET8. Trimethylation is performed by SUV420H1 and

SUV420H2 and induces gene silencing.

Ubiquitinated by the CUL4-DDB-RBX1 complex in response to ultraviolet irradiation. This may weaken the interaction between histones and DNA and facilitate DNA accessibility to repair proteins. Monoubiquitinated at Lys-92 of histone H4 (H4K91ub1) in response to DNA damage. The exact role of H4K91ub1 in DNA damage response is still unclear but it may function as a licensing signal for additional histone H4 post-translational modifications such as H4 Lys-21

methylation (H4K20me).

Sumoylated, which is associated with transcriptional repression.

#### **Cellular localization**

Nucleus. Chromosome.

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