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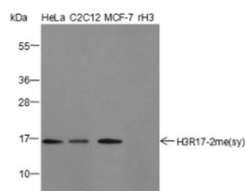
sales@bioss.com.cn

techsupport@bioss.com.cn

400-901-9800

**Symmetric Di-Methyl-Histone H3 (Arg17) Rabbit pAb****— DATASHEET —**

<b>Host:</b> Rabbit	<b>Isotype:</b> IgG	<b>Applications:</b> WB (1:500-1:2000)
<b>Clonality:</b> Polyclonal		<b>Reactivity:</b> Human (predicted: Mouse, Rat)
<b>Target:</b> Symmetric Di-Methyl-Histone H3 (Arg17)		<b>Subcellular Location:</b> Nucleus
<b>Purification:</b> Antigen affinity purification		
<b>Concentration:</b> 1mg/ml		
<b>Storage:</b> 0.01M TBS (pH7.4) with 1% BSA, 0.02% Proclin300 and 50% Glycerol. Shipped at 4°C. Store at -20°C for one year. Avoid repeated freeze/thaw cycles.		
<b>Background:</b> Modulation of the chromatin structure plays an important role in the regulation of transcription in eukaryotes. The nucleosome, made up of four core histone proteins (H2A, H2B, H3 and H4), is the primary building block of chromatin. The N-terminal tail of core histones undergoes different posttranslational modifications including acetylation, phosphorylation and methylation. These modifications occur in response to cell signal stimuli and have a direct effect on gene expression. In most species, the histone H2B is primarily acetylated at lysines 5, 12, 15 and 20. Histone H3 is primarily acetylated at lysines 9, 14, 18 and 23. Acetylation at lysine 9 appears to have a dominant role in histone deposition and chromatin assembly in some organisms. Phosphorylation at Ser10 of histone H3 is tightly correlated with chromosome condensation during both mitosis and meiosis.		

**— VALIDATION IMAGES —**

Blocking buffer: 5% NFDM/TBST Primary ab  
dilution: 1:2000 Primary ab incubation  
condition: 2 hours at room temperature  
Secondary ab: Goat Anti-Rabbit IgG H&L (HRP)  
Lysate: HeLa, C2C12, MCF-7 Protein loading  
quantity: 20 µg Exposure time: 60 s Predicted  
MW: 17 kDa Observed MW: 17 kDa