### Histone H2A/H2B Dimer Human, Recombinant

**M2508S**

- **Quantity:** 2 nmol
- **Concentration:** 20 µM
- **Lot:** 0011212
- **Storage:** Store at –20°C
- **Expiration:** Exp: 12/13

**Description:** Histone H2A combines with Histone H2B to form the H2A/H2B heterodimer. Two H2A/H2B heterodimers interact with an H3/H4 tetramer to form the histone octamer (1,2). Histones are also modified by various enzymes and can act as substrates for them. These modifications have been shown to be important in gene regulation. Because the histones are folded with their subunit partners, the dimer may be a better substrate for specific enzymes and modifications (3).

**Source:** Purified H2A and H2B (NEB #M2502 and NWB #M2505) were denatured, refolded and purified by gel filtration.

### SDS-PAGE analysis of Histone H2A/H2B Dimer Human, Recombinant

<table>
<thead>
<tr>
<th>Lane</th>
<th>Description</th>
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<tbody>
<tr>
<td>1 &amp; 7</td>
<td>NEB Protein Ladder (NEB #P7703)</td>
</tr>
<tr>
<td>2</td>
<td>Histone H2A (NEB #M2502S)</td>
</tr>
<tr>
<td>3</td>
<td>Histone H2B (NEB #M2505S)</td>
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<tr>
<td>4–6</td>
<td>2, 4, 8 µg Histone H2A/H2B Dimer</td>
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**Note:** To use as a substrate in an enzyme modification assay or other salt sensitive protocol, use 1 to 2 µl of the dimer in a minimum 20 µl reaction so that the salt concentration in the reaction ≤ 200 mM.

### Quality Control Assays:

**Protein Concentration:** 20 µM (0.55 mg/ml) calculated using the molar extinction coefficient of Histone Dimer (11,920) and its absorbance at 280 nm (4.5).

**Endonuclease Assay:** Incubation of a 50 µl reaction containing 10 µg (~360 pmol) of Histone H2A/H2B Dimer Human, Recombinant with 1 µg of X174 RF I (supercoiled) plasmid DNA for 4 hours at 37°C resulted in < 5.0% conversion to RF II form (nicked circle) as determined by agarose gel electrophoresis.

### References: