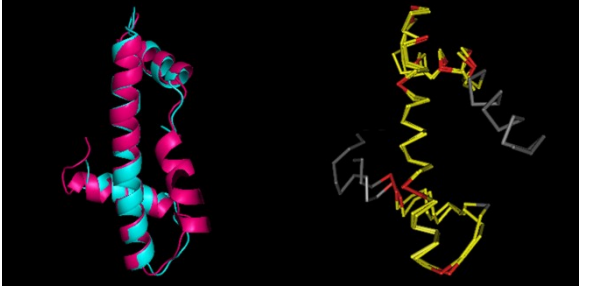
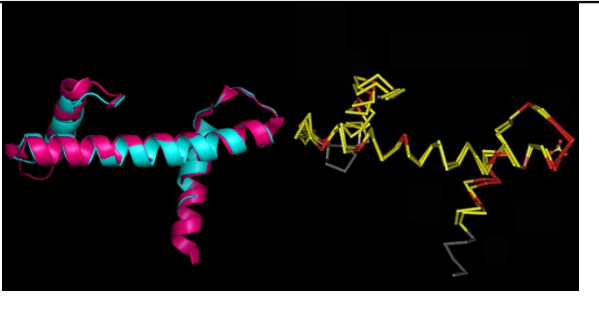
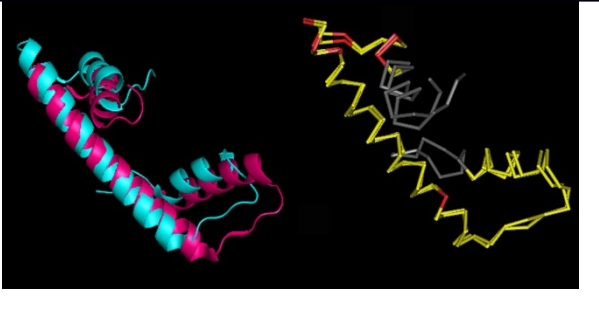
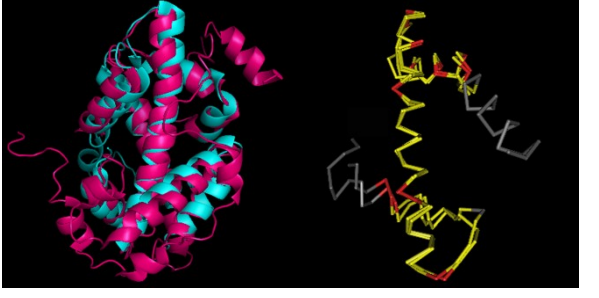


PDB # - Function	RMSD - % identity	Images from PyMOL and Cn3D	References
<p><b>Example 1</b></p> <ul style="list-style-type: none"> <li>- 1BH8_Chain B: TATA binding protein associated factor (TAF)II28 in <i>Homo sapiens</i></li> <li>- 1A7W: Histone HmfB from archaea <i>Methanothermus fervidus</i></li> </ul>	<p>RMSD: 1.0 Identity: 22.6%</p>		<p>37, 69</p>
<p><b>Example 2</b></p> <ul style="list-style-type: none"> <li>- 3R45_Chain A: Histone H3-like centromeric protein A in <i>Homo sapiens</i></li> <li>- 1B6W: Histone HmfA from archaea <i>Methanothermus fervidus</i></li> </ul>	<p>RMSD: 1.1 Identity: 24.6%</p>		<p>37, 70</p>
<p><b>Example 3</b></p> <ul style="list-style-type: none"> <li>- 3AFA_Chain E: Histone 3 in <i>Homo sapiens</i></li> <li>- 1H3O_Chain D: Transcription Initiation Factor (TFIID) in <i>Homo sapiens</i></li> </ul>	<p>RMSD: 1.0 Identity: 12.3%</p>		<p>71, 72</p>
<p><b>Example 4</b></p> <ul style="list-style-type: none"> <li>- 2JSS: Yeast (<i>Saccharomyces cerevisiae</i>) Histone H2A/H2B</li> <li>- 1Q9C: Double histone fold of <i>Homo sapiens</i> Son of sevenless (hSos)</li> </ul>	<p>RMSD: 1.9 Identity: 22.7%</p>		<p>73, 74</p>