Supplementary Table 2: Interaction change in histone monomers with respect to key findings

Simulation type	1	Observed structural change in histone	Monomer 1	Monomer 2
	number			
Intact nucleosome	1	Formation of α-helix in H3 tail	V	X
H3 tail truncated	1	Change of interaction in Arg81 of H2A	X	
	1	Change of interaction in Aigor of 112A	X	V
nucleosome	2		X	V
	3		X	X
	1	Change of interaction of Arg88 of H2A	X	√
	2		X	$\sqrt{}$
	3		X	V
	1	Destabilization of H2A docking domain contacts with closely lying residues	X	V
	2		x	$\sqrt{}$
	3		Х	√
H2A tail truncated	1	Change of interaction of Arg88 of H2A	√	√
nucleosome	2		V	V
	1	Destabilization of H2A docking domain contacts with closely lying residues	V	V
	2		٧	X