

Table S2: Compare the LeNup predictions with the MNase-seg results

Threshold	number of nucleosome	number of nucleosome	ratio
	MNase-seg	LeNup prediction	
2	143196	109930	0.7677
5	119857	99107	0.8269
10	67569	61730	0.9136
15	28706	27467	0.9568
20	11209	10794	0.9630
25	3807	3744	0.9835
30	1375	1362	0.9905