

Additional File 2: Performance of various machine-learning algorithms to classify enriched and depleted CTCF sites (ENCODE peak list for K562 cell line) using multiple histone marks.

	SVM - RBF kernel	SVM – Polynomial kernel	Random forest (RF)
auROc	82.39	79.01	81.52
Time (s)	1024.89	6127.795	566.916