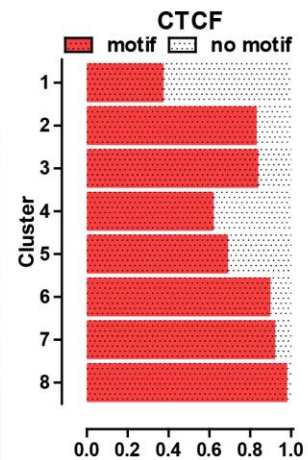
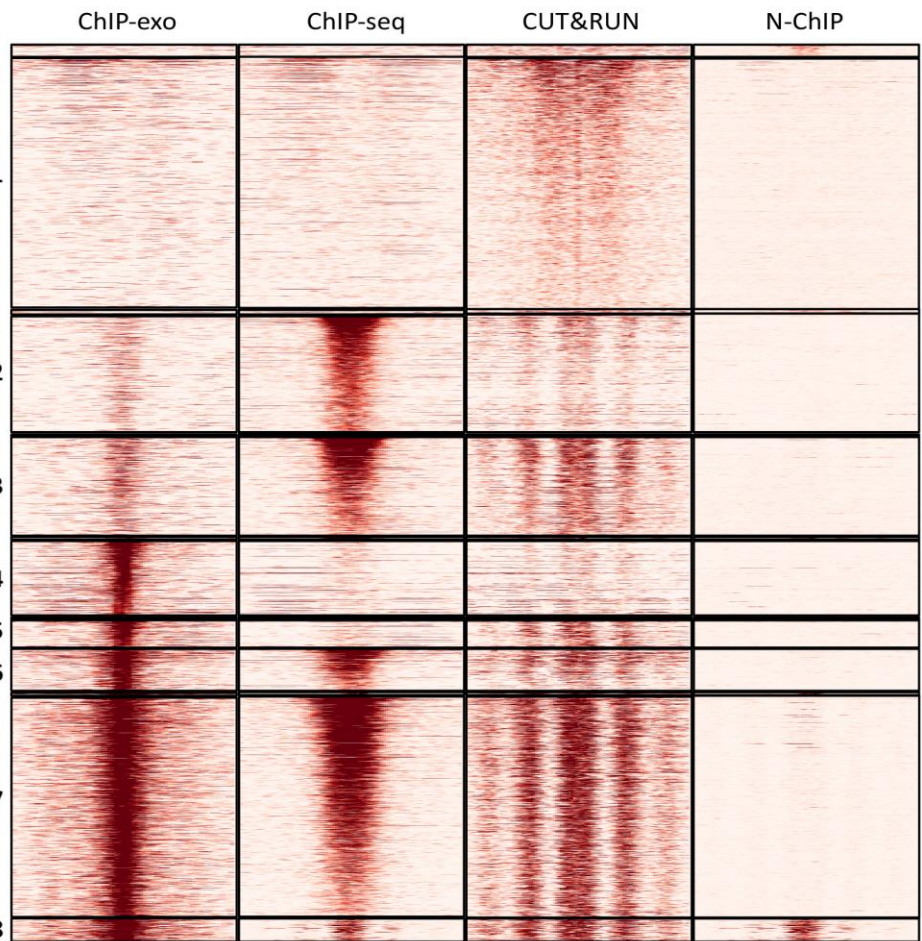
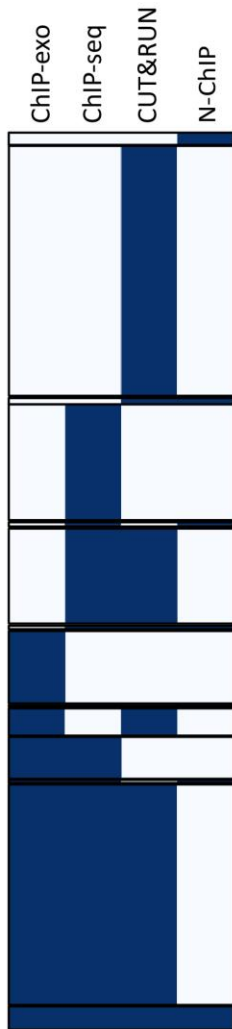


CTCF

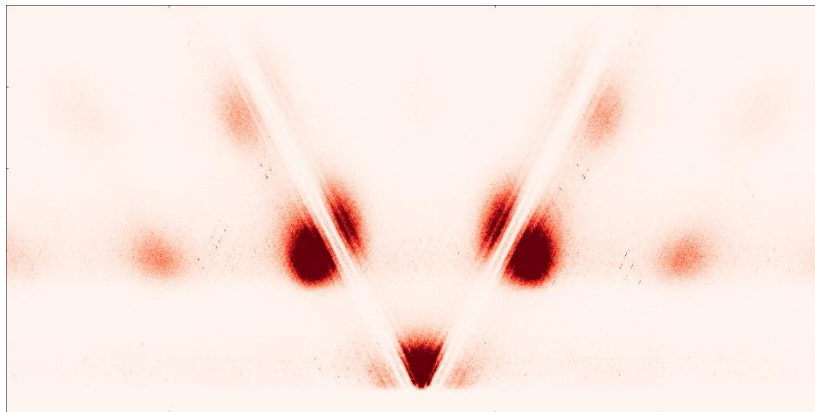


CTCF

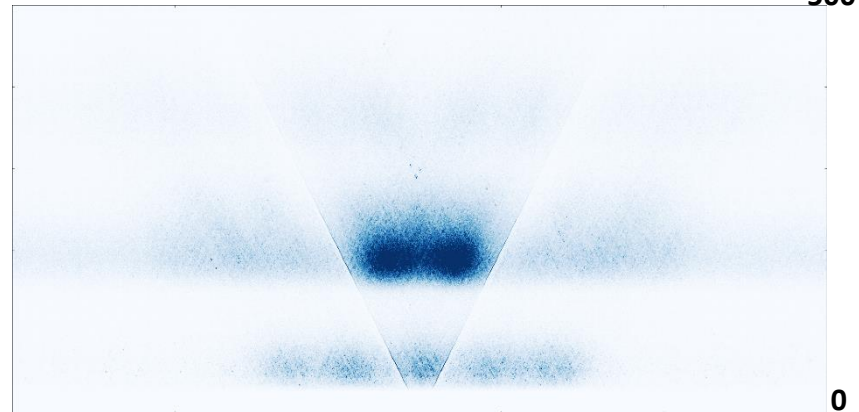
motif-centered

MACS2 peak call-centered

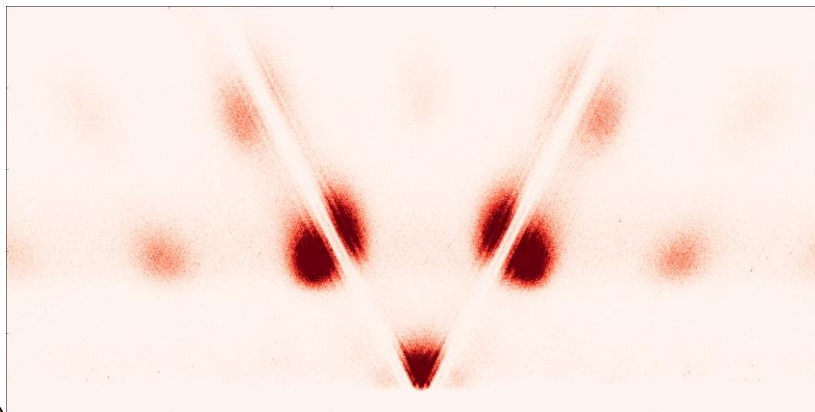
ChIP-common sites
with motif



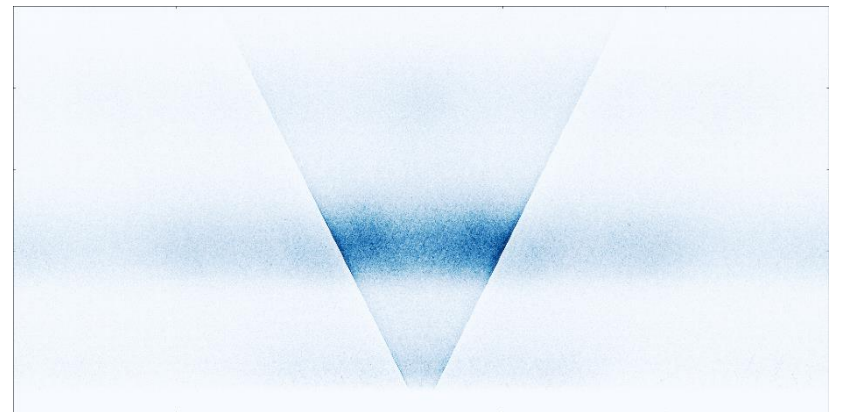
ChIP-common sites
with motif



CUTnRUN-only sites
with motif



CUTnRUN-only sites
no motif



GATA1

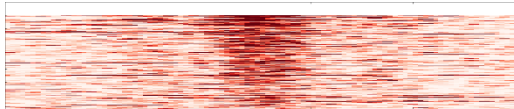
number peaks

	number peaks			
	ChIP peak		no ChIP peak	
	motif	no motif	motif	no motif
ENCODE K562 GATA1	6,328	8,277		
GATA1_1_100_250k_neb	511	558	117	525
GATA1_1_2500_250k_neb	213	261	22	167
GATA1_1_2500_50k_neb	595	568	101	476
GATA1_1_500_250k_neb	1,678	1,613	788	2,064
GATA1_1_500_50k_neb	65	111	9	82

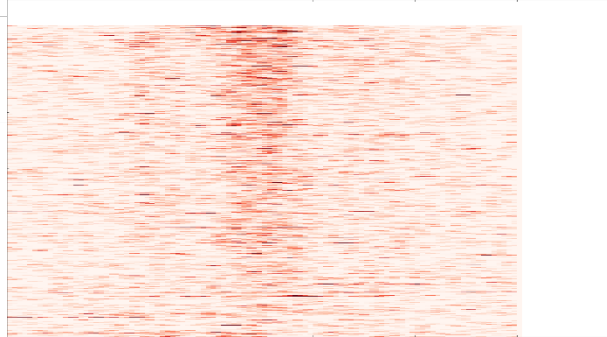
fraction

	ChIP peak		no ChIP peak		fraction	
	ChIP peak		no ChIP peak		fraction	
	motif	no motif	motif	no motif	motif	no motif
ENCODE K562 GATA1	1.00	0.00	0.43	0.57	n/a	n/a
GATA1_1_100_250k_neb	0.62	0.38	0.48	0.52	0.18	0.82
GATA1_1_2500_250k_neb	0.71	0.29	0.45	0.55	0.12	0.88
GATA1_1_2500_50k_neb	0.67	0.33	0.51	0.49	0.18	0.82
GATA1_1_500_250k_neb	0.54	0.46	0.51	0.49	0.28	0.72
GATA1_1_500_50k_neb	0.66	0.34	0.37	0.63	0.10	0.90

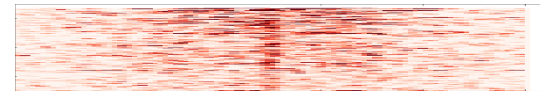
common sites



ChIP-only sites



CUT&RUN only sites

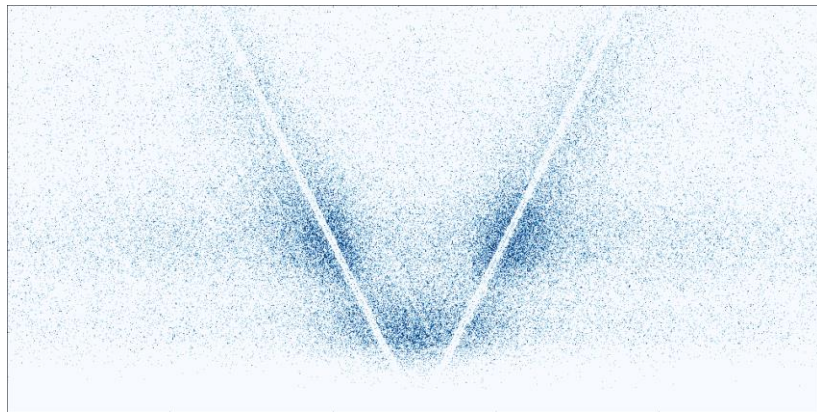


GATA1

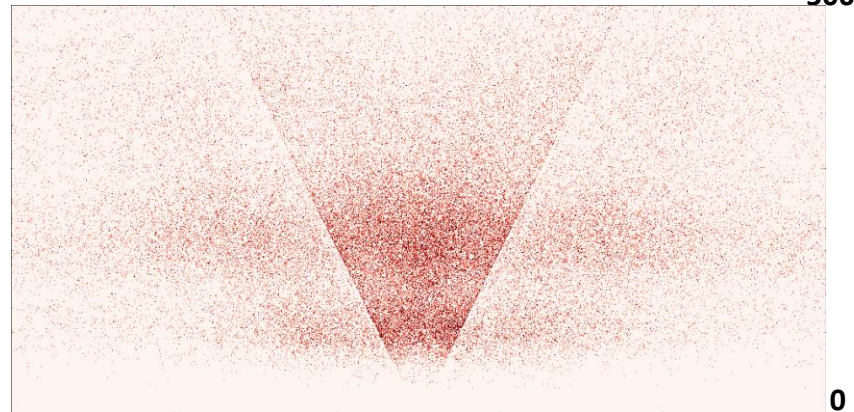
motif-centered

MACS2 peak call-centered

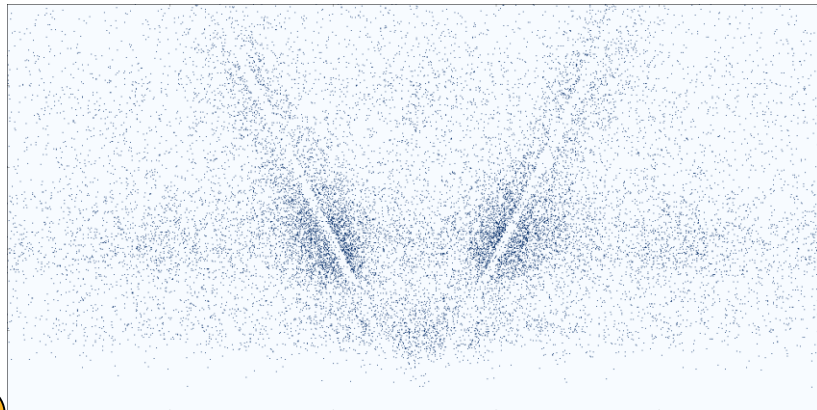
ChIP-common sites
with motif



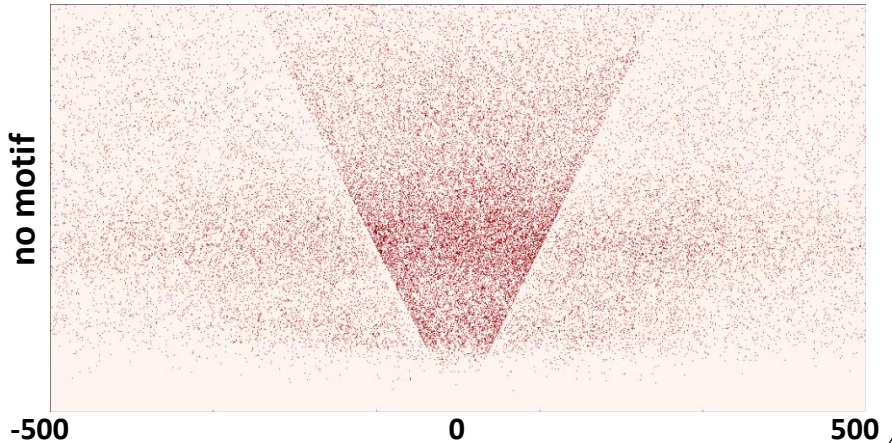
ChIP-common sites
with motif

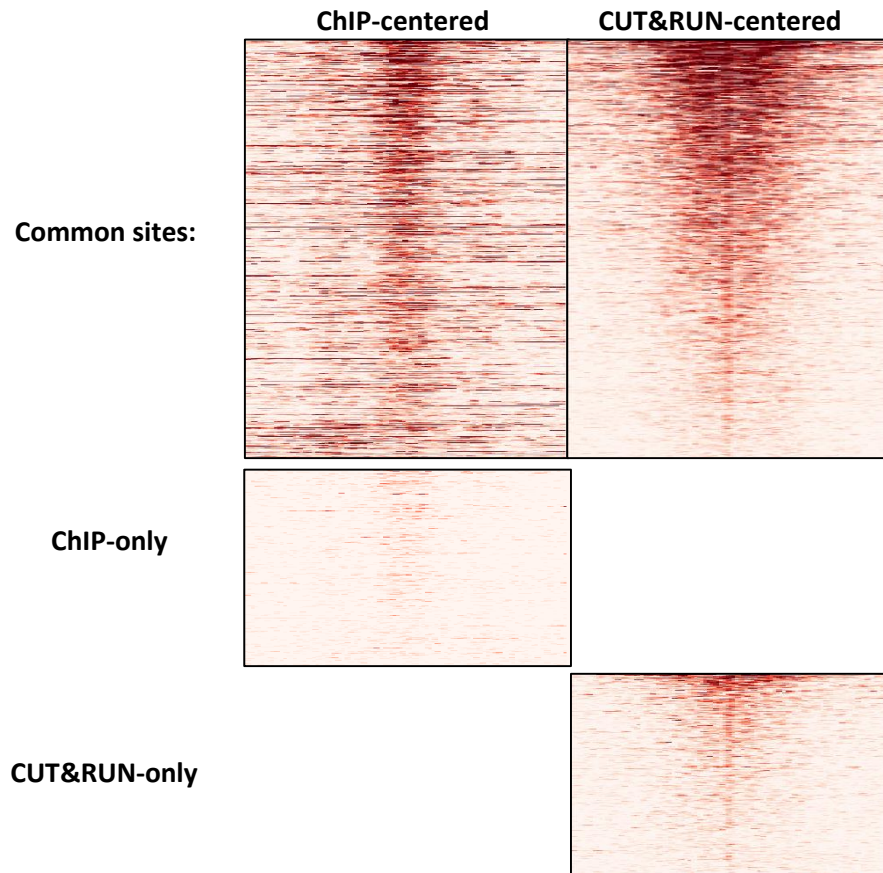


CUTnRUN-only sites
with motif



CUTnRUN-only sites
no motif





MYC

