ctcf_binding_site_rank

	tf_binding_site_ra	0.75 0.50	dnasei_rank			
1	exon	exon_rank	intron	intron_rank	open_chromatin_region	open_chromatin_
	81770.0	0.75	66090.0	0.5	49230.0	
	99711.0	1.0	37435.0	0.75	13967.0	
	33009.0	0.5	123379.0	1.0	101846.0	
•	2640.0	0.25	16380.0	0.25	100174.0	
•	120810.0	1.0	7056.0	0.25	102356.0	
•	93109.0	0.75	39055.0	0.25	19035.0	
	promoter_rank \	X	/ CAOII_IUIIK		•	•

promoter_flanking_region_rank

intron_rank

open_chromatin_region_rank