

# Dataset stats

Species	#	Raw fragments	Read Length	Unique	Unique dedup	Multi	Fraction aligned	Complexity	NSC	RSC	QC
hg38	ChIP-seq_JUNB_Act.5h-GSM3258570	54,581,168	1x36	41,882,020			0.77	0.51	1.330	1.010	1

ChIP-seq\_JUNB\_Act.5h-GSM3258570.1x36mers.unique.ba

