Table S5. MACS program parameters

MACS		
Name	Value	Description
mfold	5 (5)	threshold for high quality peaks when determining d
gsize	3.66E8	effective genome size
tsize	36	tag length
bw	120	bandwidth (length of sheared fragments)
pvalue	0.05	p-value to determine significant peaks

This table shows the parameters that were used when running MACS. Default parameters were used for all other values.