

Genome Browser

hg38

Genome Version

tracks

H1_hg38

K562_TSA_SON

H1_TSA_SON

K562_TSA_LaminB

H1_TSA_LaminB

K562_DamID_LaminB

H1_DamID_LaminB

K562_DamID_Nucleolus

H1_DamID_Nucleolus

SPIN_H1

SPIN_K562

omero

2-stage_Repli-seq.hg38

0/54

4DNFI3KHL5UL

4DNFI4BSJRMF

4DNFI5BZJXDE

4DNFI5LSTPKO

4DNFI6TILVWX

4DNFI7EOU166

4DNFI8HG51KP

4DNFI9FTS684

4DNFI9R1MQ7U

4DNFI9UM7MDC

4DNFIBH7GURD

4DNFIBIZK6EY

4DNFIBKO9LI7

4DNFIC4VLF86

4DNFI3KHL5UL

4DNFI4BSJRMF

4DNFI5BZJXDE

4DNFI5LSTPKO

4DNFI6TILVWX

4DNFI7EOU166

4DNFI8HG51KP

4DNFI9FTS684

4DNFI9R1MQ7U

4DNFI9UM7MDC

4DNFIBH7GURD

4DNFIBIZK6EY

4DNFIBKO9LI7

4DNFIC4VLF86

2-stage_Repli-seq.hg38

0/54

3D - SPIN

Structure

Structure URL

/static/data/1

H1_TSA_LaminB

CLEAR

Genome: hg38

Structure : 1

Binsize: 200000

Signal : H1_TSA_LaminB

Dropping

Current

chr18:38,175,001-38,350,000;chr18:38,725,001-39,150,000;chr18:39,575,001-41,650,000;chr18:42,150,001-44,525,000;chr18:51,775,001-54,075,000

Highlight

chr18:38,175,001-38,350,000;chr18:38,725,001-39,150,000;chr18:39,575,001-41,650,000;chr18:42,150,001-44,525,000;chr18:51,775,001-54,075,000

