
Repeating Elements by RepeatMasker
MT-ND5
MT-TL2
MT-ND4
MT-ND4L
MT-TR
MT-CO3
MT-ATP6
MT-ATP8
MT-TK
MT-CO2
MT-TD
MT-TS1
MT-CO1
MT-TY
MT-TC
MT-TN
MT-TA
MT-TW
MT-ND2
MT-TM
MT-TQ
MT-TI
MT-CYB
MT-TE
MT-ND1
MT-TL1
MT-ND6
MT-TS2
MT-TH
MT-RNR2
MT-TV
MT-ND3
MT-TG
MT-RNR1
MT-TF
ChIP_BRCA1
KD_DDX5
Gene Expression in 54 tissues from GTEx RNA-seq of 17382 samples, 948 donors (V8, Aug 2019)
K562_expt_1811457
K562_expt_1811458
HepG2_expt_1011229
K562_expt_1811449
K562_expt_1811450
K562_expt_1811451
HepG2_expt_1011130
HepG2_expt_1011131
K562_expt_1811453
K562_expt_1811454
K562_expt_1811455
K562_expt_1811456
K562_expt_1811547
K562_expt_1811548
HepG2_expt_1010921
HepG2_expt_1010922
K562_expt_1811222
HepG2_expt_1010923
HepG2_expt_1010924
HepG2_expt_1010925
HepG2_expt_1010926
HepG2_expt_1010927
K562_expt_1811223
K562_expt_1811224
HepG2_expt_1010928
K562_expt_1811225
HepG2_expt_1010929
K562_expt_1811226
HepG2_expt_1011000
HepG2_expt_1011001
HepG2_expt_1011002
HepG2_expt_1011003
HepG2_expt_1011004
HepG2_expt_1011005
K562_expt_1811227
K562_expt_1811228
K562_expt_1811229
HepG2_expt_1011006
HepG2_expt_1011007
HepG2_expt_1011008
HepG2_expt_1011009
HepG2_expt_1011010
HepG2_expt_1011011
HepG2_expt_1011012
HepG2_expt_1011013
K562_expt_1811230
HepG2_expt_1011014
K562_expt_1811231
HepG2_expt_1011015
HepG2_expt_1011016
HepG2_expt_1011017
HepG2_expt_1011018
HepG2_expt_1011019
HepG2_expt_1011020
HepG2_expt_1011021
HepG2_expt_1011022
HepG2_expt_1011023
K562_expt_1811232
K562_expt_1811233
HepG2_expt_1011024
HepG2_expt_1011025
K562_expt_1811234
HepG2_expt_1011026
K562_expt_1811235
K562_expt_1811236
HepG2_expt_1011027
HepG2_expt_1011028
K562_expt_1811237
HepG2_expt_1011029
K562_expt_1811238
K562_expt_1811239
K562_expt_1811240
K562_expt_1811241
K562_expt_1811242
K562_expt_1811243
K562_expt_1811244
K562_expt_1811245
Scale
chrM:
1,000
2,000
3,000
4,000
5,000
6,000
7,000
8,000
9,000
10,000
11,000
12,000
13,000
14,000
15,000
16,000
5 kb
hg38
GENCODE V49
RefSeq Curated
RefSeq genes from NCBI
Stacked Quantitative R-loop Score
Quantitative R-loop Score
Quantitative R-loop Zone
HEK293 Quantitative R-loop Region
HEK293 Quantitative R-loop Score
HeLa Quantitative R-loop Region
HeLa Quantitative R-loop Score
HeLa Score
V2-Level-1
V2-Level-4
V2-Level-8
bigwig_99_HEK293_Control_DRIP-seq_ChIP_NS_1_SE
DRIP-seq_HEK293
bigbed_99_HEK293_Control_DRIP-seq_ChIP_NS_1_SE
bigwig_229_HEK293T_2HBD-Anti-GST_CUT&Tag_ChIP_NS_2_PE
CUT&Tag_HEK293T
bigbed_229_HEK293T_2HBD-Anti-GST_CUT&Tag_ChIP_NS_2_PE
bigwig_127_HEK293T_D210N_R-ChIP-SS_ChIP_SS_1_SE_fwd
R-ChIP-SS_HEK293T.fwd
bigwig_127_HEK293T_D210N_R-ChIP-SS_ChIP_SS_1_SE_rev
R-ChIP-SS_HEK293T.rev
bigbed_127_HEK293T_D210N_R-ChIP-SS_input_SS_1_SE
bigwig_DeepER_Human_fwd
DeepER_Human.fwd
bigbed_DeepER_Human
bigwig_DeepER_Human_rev
DeepER_Human.rev
GC-skew
AT-skew
HEK293T_Control_GRO-seq.fwd
GRO-seq_HEK293T.fwd
HEK293T_Control_GRO-seq.rev
GRO-seq_HEK293T.rev
RLFS_RloopDB
CLIP_AQR
K562_expt_1811546
K562_expt_1811547
K562_expt_1811548
K562_expt_1811549
K562_expt_1811550
K562_expt_1811551
K562_expt_1811552
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K562_expt_1811554
K562_expt_1811555
K562_expt_1811556
K562_expt_1811557
K562_expt_1811558
HepG2_expt_1011120
HepG2_expt_1011121
HepG2_expt_1011122
HepG2_expt_1011123
K562_expt_1811231
K562_expt_1811232
K562_expt_1811233
HepG2_expt_1011124
HepG2_expt_1011125
K562_expt_1811234
HepG2_expt_1011126
K562_expt_1811235
K562_expt_1811236
HepG2_expt_1011127
K562_expt_1811237
HepG2_expt_1011128
K562_expt_1811238
K562_expt_1811239
K562_expt_1811240
K562_expt_1811241
K562_expt_1811242
K562_expt_1811243
K562_expt_1811244
K562_expt_1811245