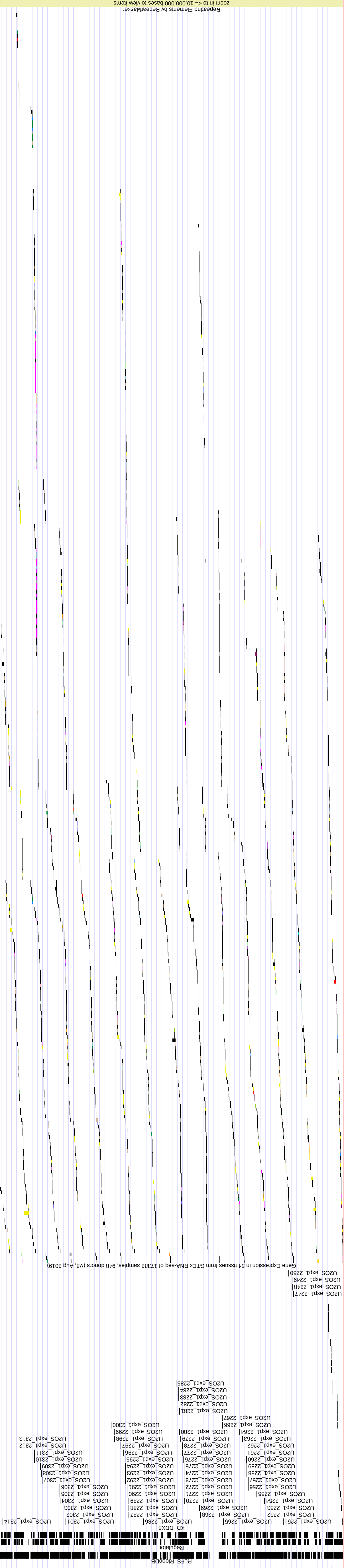

RepeatMasker
Repeating Elements by RepeatMasker
zoom in to <= 10,000,000 bases to view items
RLFS_RloopDB
CLIP_AQR
ChIP_BRCA1
KD_DDX5
U2OS_exp1_2214
U2OS_exp1_2251
U2OS_exp1_2265
U2OS_exp1_2286
U2OS_exp1_2301
U2OS_exp1_2314
U2OS_exp1_2215
U2OS_exp1_2252
U2OS_exp1_2268
U2OS_exp1_2287
U2OS_exp1_2302
U2OS_exp1_2216
U2OS_exp1_2253
U2OS_exp1_2269
U2OS_exp1_2288
U2OS_exp1_2303
U2OS_exp1_2217
U2OS_exp1_2254
U2OS_exp1_2270
U2OS_exp1_2289
U2OS_exp1_2304
U2OS_exp1_2218
U2OS_exp1_2255
U2OS_exp1_2271
U2OS_exp1_2290
U2OS_exp1_2305
U2OS_exp1_2219
U2OS_exp1_2256
U2OS_exp1_2272
U2OS_exp1_2291
U2OS_exp1_2306
U2OS_exp1_2220
U2OS_exp1_2257
U2OS_exp1_2273
U2OS_exp1_2292
U2OS_exp1_2307
U2OS_exp1_2221
U2OS_exp1_2258
U2OS_exp1_2274
U2OS_exp1_2293
U2OS_exp1_2308
U2OS_exp1_2222
U2OS_exp1_2259
U2OS_exp1_2275
U2OS_exp1_2294
U2OS_exp1_2309
U2OS_exp1_2223
U2OS_exp1_2260
U2OS_exp1_2276
U2OS_exp1_2295
U2OS_exp1_2310
U2OS_exp1_2224
U2OS_exp1_2261
U2OS_exp1_2277
U2OS_exp1_2296
U2OS_exp1_2311
U2OS_exp1_2225
U2OS_exp1_2262
U2OS_exp1_2278
U2OS_exp1_2297
U2OS_exp1_2312
U2OS_exp1_2226
U2OS_exp1_2263
U2OS_exp1_2279
U2OS_exp1_2298
U2OS_exp1_2313
U2OS_exp1_2227
U2OS_exp1_2264
U2OS_exp1_2280
U2OS_exp1_2299
U2OS_exp1_2228
U2OS_exp1_2266
U2OS_exp1_2300
U2OS_exp1_2229
U2OS_exp1_2267
U2OS_exp1_2230
U2OS_exp1_2281
U2OS_exp1_2231
U2OS_exp1_2282
U2OS_exp1_2232
U2OS_exp1_2283
U2OS_exp1_2233
U2OS_exp1_2284
U2OS_exp1_2234
U2OS_exp1_2285
U2OS_exp1_2235
U2OS_exp1_2236
U2OS_exp1_2237
U2OS_exp1_2238
U2OS_exp1_2239
U2OS_exp1_2240
U2OS_exp1_2241
U2OS_exp1_2242
U2OS_exp1_2243
U2OS_exp1_2244
U2OS_exp1_2245
U2OS_exp1_2246
U2OS_exp1_2247
U2OS_exp1_2248
U2OS_exp1_2249
U2OS_exp1_2250
Gene Expression in 54 tissues from GTEx RNA-seq of 17382 samples, 948 donors (V8, Aug 2019)
RefSeq Curated
3.57423
Stacked Quantitative R-loop Score
Quantitative R-loop Score
0
Quantitative R-loop Zone
Quantitative R-loop Zone
HEK293 Region
16.4556
HEK293 Quantitative R-loop Region
HEK293 Score
HEK293 Quantitative R-loop Score
0
HeLa Region
10.6566
HeLa Quantitative R-loop Region
HeLa Score
HeLa Quantitative R-loop Score
0
R-loop Zone Level
V2-Level-1
V2-Level-4
V2-Level-8
DRIP-seq_HEK293
6.642
bigwig_99_HEK293_Control_DRIP-seq_ChIP_NS_1_SE
0
R-loop Signal
DRIP-seq_HEK293
31.5237
bigwig_229_HEK293T_2HBD-Anti-GST_CUT&Tag_ChIP_NS_2_PE
0
R-loop Signal
CUT&Tag_HEK293T
16.764
bigwig_127_HEK293T_D210N_R-ChIP-SS_ChIP_SS_1_SE_fwd
R-ChIP-SS_HEK293T.fwd
0
bigwig 127 HEK293T D210N R-ChIP-SS ChIP SS 1 SE rev
-0
R-ChIP-SS_HEK293T.rev
-12.954
bigbed 127 HEK293T D210N R-ChIP-SS Input SS 1 SE
bigwig DeepER Human fwd
0.9999
DeepER_Human.fwd
1e-05
DeepER
bigwig DeepER Human rev
DeepER_Human
-1e-05
DeepER_Human.rev
-0.99989
GC-skew
1
GC-skew
-1
AT-skew
1
AT-skew
-1
HEK293T_Control_GRO-seq.fwd
10.614
GRO-seq_HEK293T.fwd
0.1769
HEK293T Control GRO-seq.rev
-0.1769
GRO-seq_HEK293T.rev
-11.6754
Scale
chr7:
50 Mb
hg38
50,000,000
100,000,000
150,000,000