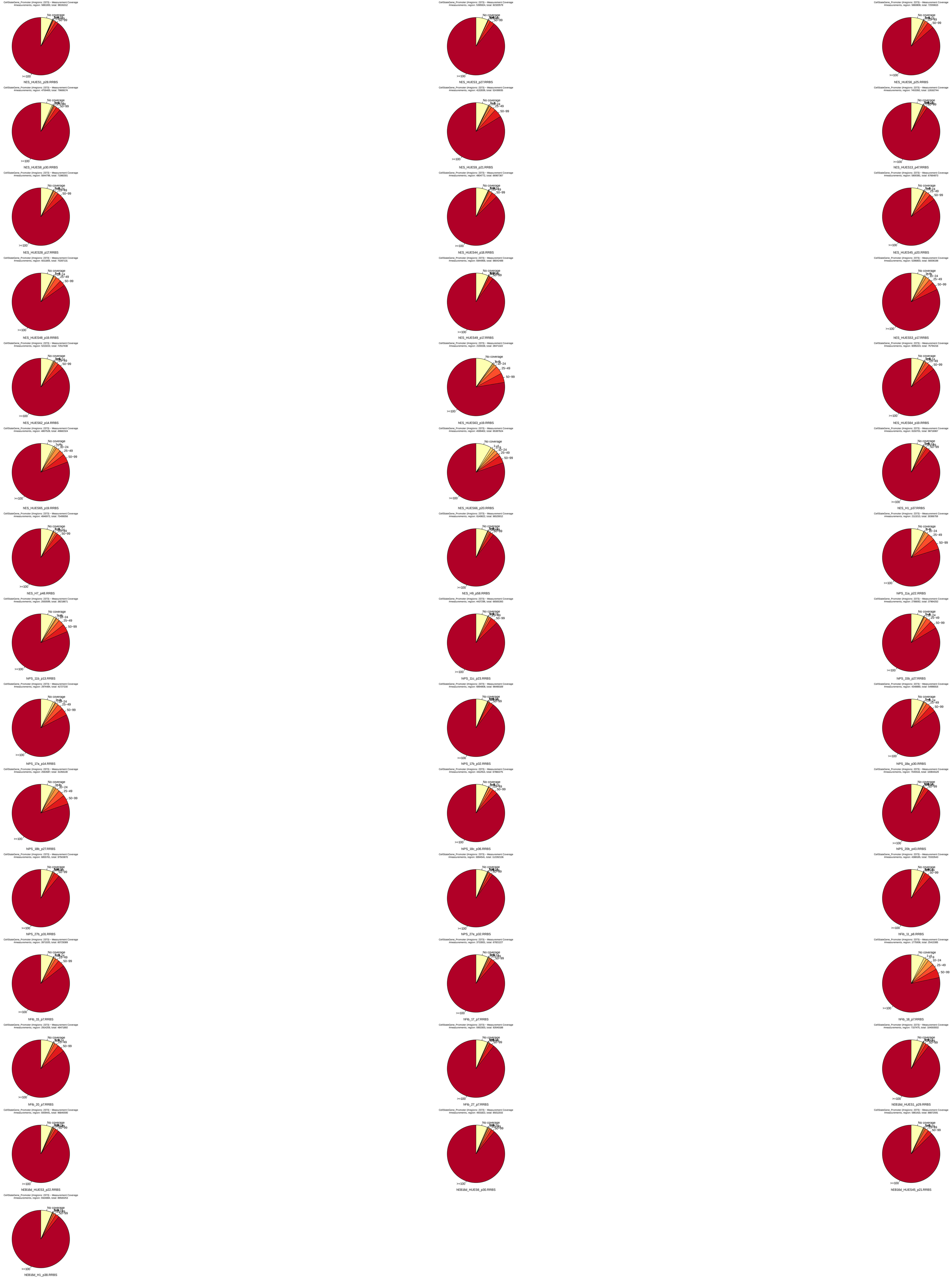
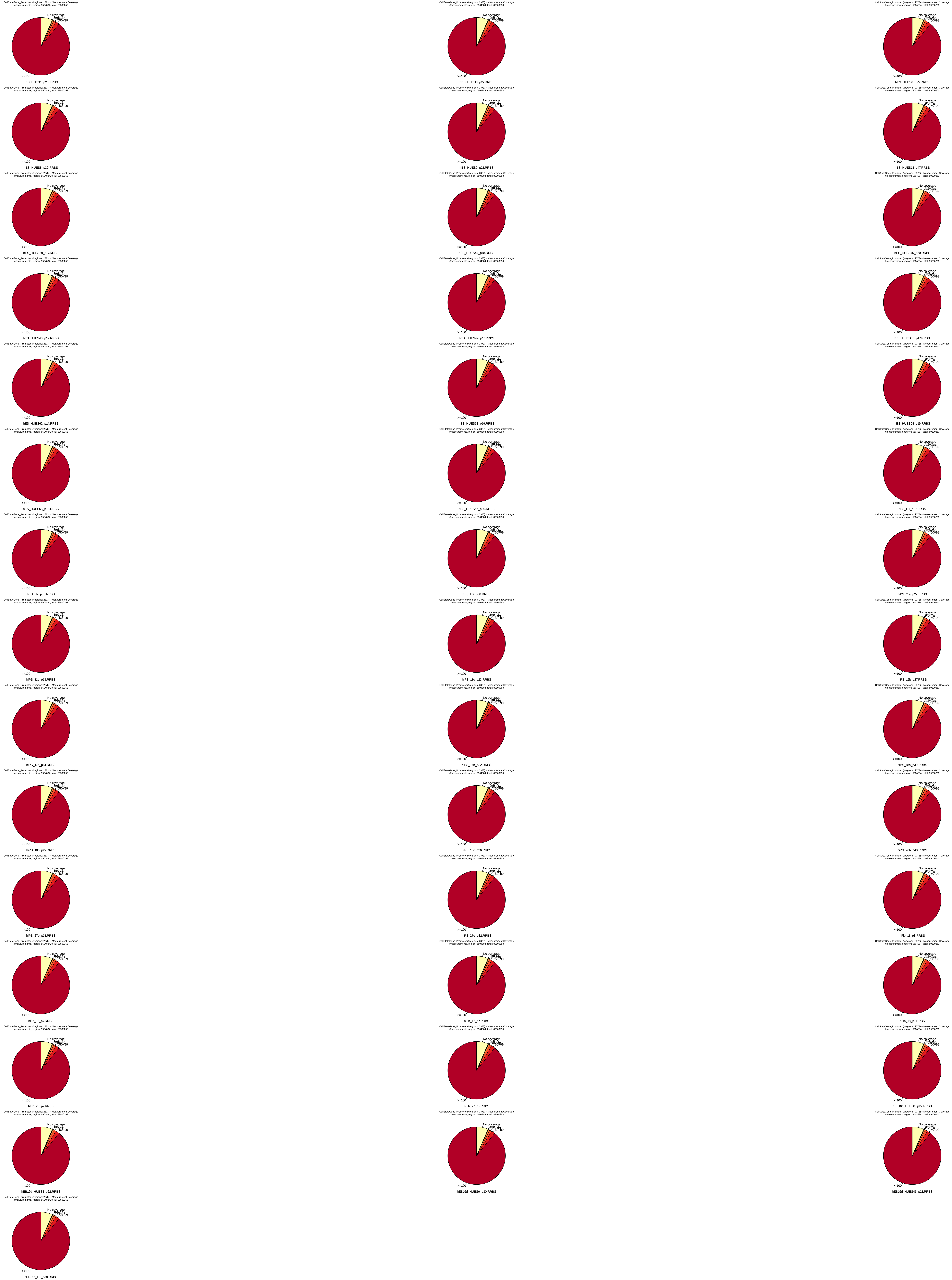


CellStateGene_Promoter (#regions: 2373) − Measurement Coverage
#measurements, region: 5881000, total: 98191012
No coverage
50−99
>=100
hES_HUES1_p28.RRBS
CellStateGene_Promoter (#regions: 2373) − Measurement Coverage
#measurements, region: 5395924, total: 82320579
No coverage
50−99
>=100
hES_HUES3_p27.RRBS
CellStateGene_Promoter (#regions: 2373) − Measurement Coverage
#measurements, region: 5860808, total: 72559615
No coverage
25−49
50−99
>=100
hES_HUES6_p25.RRBS
CellStateGene_Promoter (#regions: 2373) − Measurement Coverage
#measurements, region: 4759400, total: 79669174
No coverage
50−99
>=100
hES_HUES8_p30.RRBS
CellStateGene_Promoter (#regions: 2373) − Measurement Coverage
#measurements, region: 4132839, total: 52436935
No coverage
10−24
25−49
50−99
>=100
hES_HUES9_p21.RRBS
CellStateGene_Promoter (#regions: 2373) − Measurement Coverage
#measurements, region: 7453392, total: 119162744
No coverage
50−99
>=100
hES_HUES13_p47.RRBS
CellStateGene_Promoter (#regions: 2373) − Measurement Coverage
#measurements, region: 5844796, total: 71980301
No coverage
25−49
50−99
>=100
hES_HUES28_p17.RRBS
CellStateGene_Promoter (#regions: 2373) − Measurement Coverage
#measurements, region: 4804772, total: 66967367
No coverage
25−49
50−99
>=100
hES_HUES44_p18.RRBS
CellStateGene_Promoter (#regions: 2373) − Measurement Coverage
#measurements, region: 5800381, total: 67664873
No coverage
10−24
25−49
50−99
>=100
hES_HUES45_p20.RRBS
CellStateGene_Promoter (#regions: 2373) − Measurement Coverage
#measurements, region: 6011805, total: 70287131
No coverage
10−24
25−49
50−99
>=100
hES_HUES48_p19.RRBS
CellStateGene_Promoter (#regions: 2373) − Measurement Coverage
#measurements, region: 5944958, total: 98042489
No coverage
50−99
>=100
hES_HUES49_p17.RRBS
CellStateGene_Promoter (#regions: 2373) − Measurement Coverage
#measurements, region: 5286803, total: 56006198
No coverage
10−24
25−49
50−99
>=100
hES_HUES53_p17.RRBS
CellStateGene_Promoter (#regions: 2373) − Measurement Coverage
#measurements, region: 5210222, total: 72517438
No coverage
25−49
50−99
>=100
hES_HUES62_p14.RRBS
CellStateGene_Promoter (#regions: 2373) − Measurement Coverage
#measurements, region: 2159158, total: 28471322
No coverage
10−24
25−49
50−99
>=100
hES_HUES63_p19.RRBS
CellStateGene_Promoter (#regions: 2373) − Measurement Coverage
#measurements, region: 6095223, total: 75794216
No coverage
25−49
50−99
>=100
hES_HUES64_p19.RRBS
CellStateGene_Promoter (#regions: 2373) − Measurement Coverage
#measurements, region: 4807528, total: 49682324
No coverage
10−24
25−49
50−99
>=100
hES_HUES65_p19.RRBS
CellStateGene_Promoter (#regions: 2373) − Measurement Coverage
#measurements, region: 4599402, total: 65387624
No coverage
1−4
5−9
10−24
25−49
50−99
>=100
hES_HUES66_p20.RRBS
CellStateGene_Promoter (#regions: 2373) − Measurement Coverage
#measurements, region: 6150701, total: 98716967
No coverage
50−99
>=100
hES_H1_p37.RRBS
CellStateGene_Promoter (#regions: 2373) − Measurement Coverage
#measurements, region: 4846972, total: 75499056
No coverage
50−99
>=100
hES_H7_p48.RRBS
CellStateGene_Promoter (#regions: 2373) − Measurement Coverage
#measurements, region: 6149822, total: 96529012
No coverage
50−99
>=100
hES_H9_p58.RRBS
CellStateGene_Promoter (#regions: 2373) − Measurement Coverage
#measurements, region: 2113213, total: 30366759
No coverage
10−24
25−49
50−99
>=100
hiPS_11a_p22.RRBS
CellStateGene_Promoter (#regions: 2373) − Measurement Coverage
#measurements, region: 2930599, total: 38218671
No coverage
10−24
25−49
50−99
>=100
hiPS_11b_p13.RRBS
CellStateGene_Promoter (#regions: 2373) − Measurement Coverage
#measurements, region: 4437296, total: 66565393
No coverage
25−49
50−99
>=100
hiPS_11c_p23.RRBS
CellStateGene_Promoter (#regions: 2373) − Measurement Coverage
#measurements, region: 2788062, total: 37964252
No coverage
10−24
25−49
50−99
>=100
hiPS_15b_p27.RRBS
CellStateGene_Promoter (#regions: 2373) − Measurement Coverage
#measurements, region: 2974484, total: 41727216
No coverage
10−24
25−49
50−99
>=100
hiPS_17a_p14.RRBS
CellStateGene_Promoter (#regions: 2373) − Measurement Coverage
#measurements, region: 6894808, total: 98490169
No coverage
50−99
>=100
hiPS_17b_p32.RRBS
CellStateGene_Promoter (#regions: 2373) − Measurement Coverage
#measurements, region: 4348880, total: 54986816
No coverage
10−24
25−49
50−99
>=100
hiPS_18a_p30.RRBS
CellStateGene_Promoter (#regions: 2373) − Measurement Coverage
#measurements, region: 2563587, total: 31356130
No coverage
10−24
25−49
50−99
>=100
hiPS_18b_p27.RRBS
CellStateGene_Promoter (#regions: 2373) − Measurement Coverage
#measurements, region: 4412913, total: 67882275
No coverage
25−49
50−99
>=100
hiPS_18c_p36.RRBS
CellStateGene_Promoter (#regions: 2373) − Measurement Coverage
#measurements, region: 7045516, total: 100844129
No coverage
50−99
>=100
hiPS_20b_p43.RRBS
CellStateGene_Promoter (#regions: 2373) − Measurement Coverage
#measurements, region: 6855761, total: 97503870
No coverage
50−99
>=100
hiPS_27b_p31.RRBS
CellStateGene_Promoter (#regions: 2373) − Measurement Coverage
#measurements, region: 6994541, total: 112282136
No coverage
50−99
>=100
hiPS_27e_p32.RRBS
CellStateGene_Promoter (#regions: 2373) − Measurement Coverage
#measurements, region: 4388185, total: 70333543
No coverage
50−99
>=100
hFib_11_p8.RRBS
CellStateGene_Promoter (#regions: 2373) − Measurement Coverage
#measurements, region: 3971020, total: 60729389
No coverage
25−49
50−99
>=100
hFib_15_p7.RRBS
CellStateGene_Promoter (#regions: 2373) − Measurement Coverage
#measurements, region: 3733931, total: 67821227
No coverage
50−99
>=100
hFib_17_p7.RRBS
CellStateGene_Promoter (#regions: 2373) − Measurement Coverage
#measurements, region: 1775936, total: 25413385
No coverage
1−4
5−9
10−24
25−49
50−99
>=100
hFib_18_p7.RRBS
CellStateGene_Promoter (#regions: 2373) − Measurement Coverage
#measurements, region: 2914259, total: 48471892
No coverage
25−49
50−99
>=100
hFib_20_p7.RRBS
CellStateGene_Promoter (#regions: 2373) − Measurement Coverage
#measurements, region: 5953303, total: 92640188
No coverage
50−99
>=100
hFib_27_p7.RRBS
CellStateGene_Promoter (#regions: 2373) − Measurement Coverage
#measurements, region: 7157475, total: 104593502
No coverage
50−99
>=100
hEB16d_HUES1_p29.RRBS
CellStateGene_Promoter (#regions: 2373) − Measurement Coverage
#measurements, region: 5939441, total: 96845590
No coverage
50−99
>=100
hEB16d_HUES3_p22.RRBS
CellStateGene_Promoter (#regions: 2373) − Measurement Coverage
#measurements, region: 4931822, total: 85012015
No coverage
50−99
>=100
hEB16d_HUES6_p30.RRBS
CellStateGene_Promoter (#regions: 2373) − Measurement Coverage
#measurements, region: 5861410, total: 88872591
No coverage
25−49
50−99
>=100
hEB16d_HUES45_p21.RRBS
CellStateGene_Promoter (#regions: 2373) − Measurement Coverage
#measurements, region: 5504884, total: 89500253
No coverage
50−99
>=100
hEB16d_H1_p38.RRBS


CellStateGene_Promoter (#regions: 2373) ~ Measurement Coverage
#measurements, region: 5504884, total: 89500253
No coverage
50−99
>=100
hES_HUES1_p28.RRBS
hES_HUES3_p27.RRBS
hES_HUES6_p25.RRBS
hES_HUES8_p30.RRBS
hES_HUES9_p21.RRBS
hES_HUES13_p47.RRBS
hES_HUES28_p17.RRBS
hES_HUES44_p18.RRBS
hES_HUES45_p20.RRBS
hES_HUES48_p19.RRBS
hES_HUES49_p17.RRBS
hES_HUES53_p17.RRBS
hES_HUES62_p14.RRBS
hES_HUES63_p19.RRBS
hES_HUES64_p19.RRBS
hES_HUES65_p19.RRBS
hES_HUES66_p20.RRBS
hES_H1_p37.RRBS
hES_H7_p48.RRBS
hES_H9_p58.RRBS
hiPS_11a_p22.RRBS
hiPS_11b_p13.RRBS
hiPS_11c_p23.RRBS
hiPS_15b_p27.RRBS
hiPS_17a_p14.RRBS
hiPS_17b_p32.RRBS
hiPS_18a_p30.RRBS
hiPS_18b_p27.RRBS
hiPS_18c_p36.RRBS
hiPS_20b_p43.RRBS
hiPS_27b_p31.RRBS
hiPS_27e_p32.RRBS
hFib_11_p8.RRBS
hFib_15_p7.RRBS
hFib_17_p7.RRBS
hFib_18_p7.RRBS
hFib_20_p7.RRBS
hFib_27_p7.RRBS
hEB16d_HUES1_p29.RRBS
hEB16d_HUES3_p22.RRBS
hEB16d_HUES6_p30.RRBS
hEB16d_HUES45_p21.RRBS
hEB16d_H1_p38.RRBS