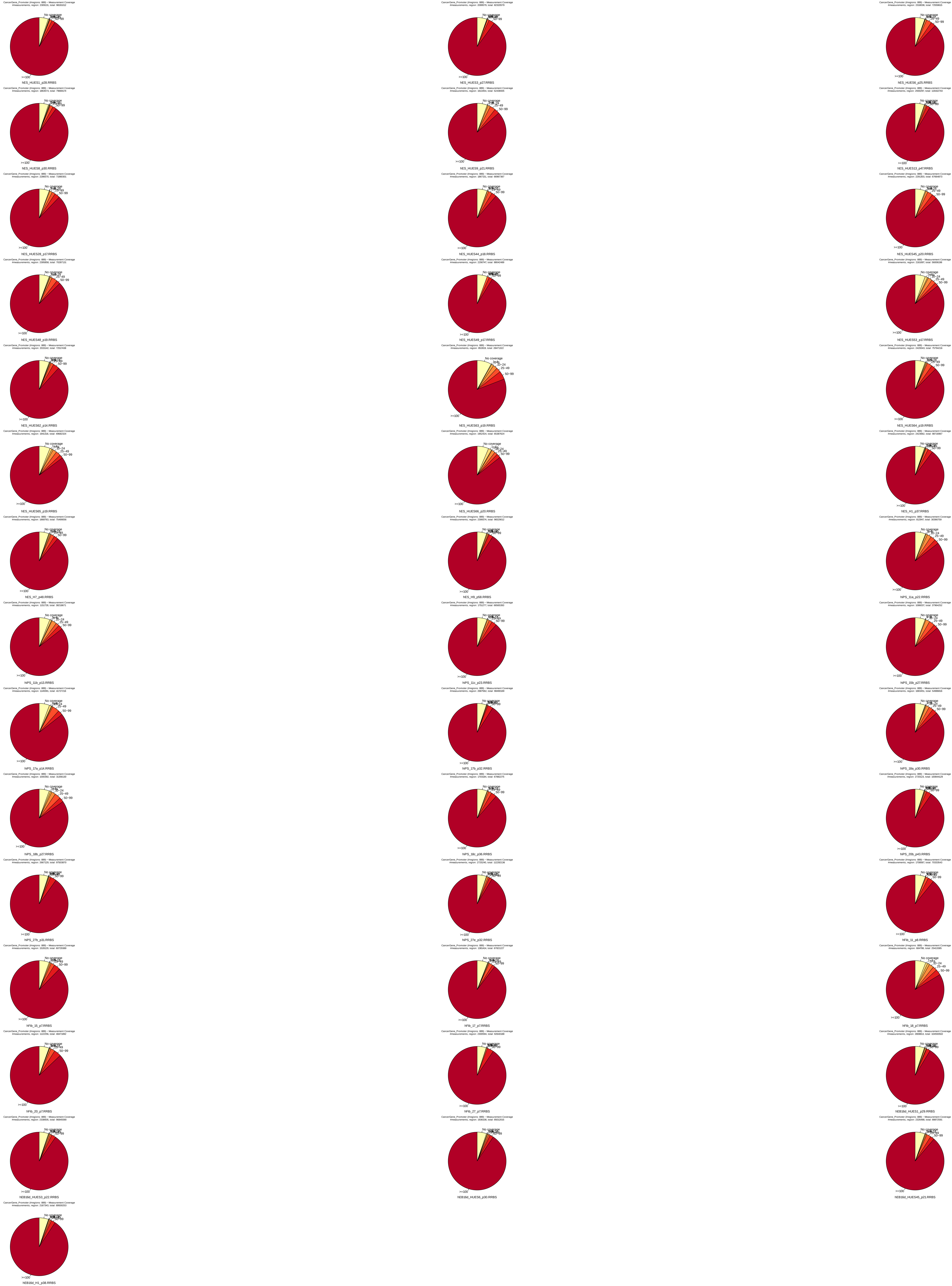
CancerGene_Promoter (#regions: 889) − Measurement Coverage
#measurements, region: 2309131, total: 98191012
hES_HUES1_p28.RRBS
CancerGene_Promoter (#regions: 889) − Measurement Coverage
#measurements, region: 2099579, total: 82320579
hES_HUES3_p27.RRBS
CancerGene_Promoter (#regions: 889) − Measurement Coverage
#measurements, region: 2318038, total: 72559615
hES_HUES6_p25.RRBS
CancerGene_Promoter (#regions: 889) − Measurement Coverage
#measurements, region: 1863573, total: 79669174
hES_HUES8_p30.RRBS
CancerGene_Promoter (#regions: 889) − Measurement Coverage
#measurements, region: 1613454, total: 52436935
hES_HUES9_p21.RRBS
CancerGene_Promoter (#regions: 889) − Measurement Coverage
#measurements, region: 2930297, total: 119162744
hES_HUES13_p47.RRBS
CancerGene_Promoter (#regions: 889) − Measurement Coverage
#measurements, region: 2286570, total: 71980301
hES_HUES28_p17.RRBS
CancerGene_Promoter (#regions: 889) − Measurement Coverage
#measurements, region: 1867151, total: 66967367
hES_HUES44_p18.RRBS
CancerGene_Promoter (#regions: 889) − Measurement Coverage
#measurements, region: 2291303, total: 67664873
hES_HUES45_p20.RRBS
CancerGene_Promoter (#regions: 889) − Measurement Coverage
#measurements, region: 2395858, total: 70287131
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CancerGene_Promoter (#regions: 889) − Measurement Coverage
#measurements, region: 2290747, total: 98042489
hES_HUES49_p17.RRBS
CancerGene_Promoter (#regions: 889) − Measurement Coverage
#measurements, region: 2161697, total: 56006198
hES_HUES53_p17.RRBS
CancerGene_Promoter (#regions: 889) − Measurement Coverage
#measurements, region: 2015142, total: 72517438
hES_HUES62_p14.RRBS
CancerGene_Promoter (#regions: 889) − Measurement Coverage
#measurements, region: 852819, total: 28471322
hES_HUES63_p19.RRBS
CancerGene_Promoter (#regions: 889) − Measurement Coverage
#measurements, region: 2420043, total: 75794216
hES_HUES64_p19.RRBS
CancerGene_Promoter (#regions: 889) − Measurement Coverage
#measurements, region: 1941318, total: 49682324
hES_HUES65_p19.RRBS
CancerGene_Promoter (#regions: 889) − Measurement Coverage
#measurements, region: 1952424, total: 65387624
hES_HUES66_p20.RRBS
CancerGene_Promoter (#regions: 889) − Measurement Coverage
#measurements, region: 2423662, total: 98716967
hES_H1_p37.RRBS
CancerGene_Promoter (#regions: 889) − Measurement Coverage
#measurements, region: 1868763, total: 75499056
hES_H7_p48.RRBS
CancerGene_Promoter (#regions: 889) − Measurement Coverage
#measurements, region: 2399374, total: 96529012
hES_H9_p58.RRBS
CancerGene_Promoter (#regions: 889) − Measurement Coverage
#measurements, region: 812947, total: 30366759
hiPS_11a_p22.RRBS
CancerGene_Promoter (#regions: 889) − Measurement Coverage
#measurements, region: 1151728, total: 38218671
hiPS_11b_p13.RRBS
CancerGene_Promoter (#regions: 889) − Measurement Coverage
#measurements, region: 1751277, total: 66565393
hiPS_11c_p23.RRBS
CancerGene_Promoter (#regions: 889) − Measurement Coverage
#measurements, region: 1086037, total: 37964252
hiPS_15b_p27.RRBS
CancerGene_Promoter (#regions: 889) − Measurement Coverage
#measurements, region: 1149391, total: 41727216
hiPS_17a_p14.RRBS
CancerGene_Promoter (#regions: 889) − Measurement Coverage
#measurements, region: 2687562, total: 98490169
hiPS_17b_p32.RRBS
CancerGene_Promoter (#regions: 889) − Measurement Coverage
#measurements, region: 1683451, total: 54986816
hiPS_18a_p30.RRBS
CancerGene_Promoter (#regions: 889) − Measurement Coverage
#measurements, region: 1000392, total: 31356130
hiPS_18b_p27.RRBS
CancerGene_Promoter (#regions: 889) − Measurement Coverage
#measurements, region: 1703184, total: 67882275
hiPS_18c_p36.RRBS
CancerGene_Promoter (#regions: 889) − Measurement Coverage
#measurements, region: 2720123, total: 100844129
hiPS_20b_p43.RRBS
CancerGene_Promoter (#regions: 889) − Measurement Coverage
#measurements, region: 2667129, total: 97503870
hiPS_27b_p31.RRBS
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hiPS_27e_p32.RRBS
CancerGene_Promoter (#regions: 889) − Measurement Coverage
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hFib_11_p8.RRBS
CancerGene_Promoter (#regions: 889) − Measurement Coverage
#measurements, region: 1528129, total: 60729389
hFib_15_p7.RRBS
CancerGene_Promoter (#regions: 889) − Measurement Coverage
#measurements, region: 1381414, total: 67821227
hFib_17_p7.RRBS
CancerGene_Promoter (#regions: 889) − Measurement Coverage
#measurements, region: 684786, total: 25413385
hFib_18_p7.RRBS
CancerGene_Promoter (#regions: 889) − Measurement Coverage
#measurements, region: 1122256, total: 48471892
hFib_20_p7.RRBS
CancerGene_Promoter (#regions: 889) − Measurement Coverage
#measurements, region: 2300594, total: 92640188
hFib_27_p7.RRBS
CancerGene_Promoter (#regions: 889) − Measurement Coverage
#measurements, region: 2808813, total: 104593502
hEB16d_HUES1_p29.RRBS
CancerGene_Promoter (#regions: 889) − Measurement Coverage
#measurements, region: 2338935, total: 96845590
hEB16d_HUES3_p22.RRBS
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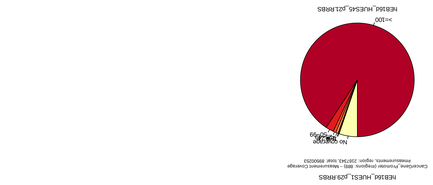
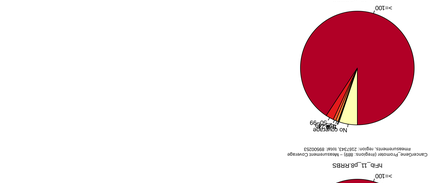
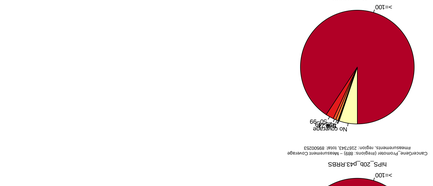
#measurements, region: 1936108, total: 85012015
hEB16d_HUES6_p30.RRBS
CancerGene_Promoter (#regions: 889) − Measurement Coverage
#measurements, region: 2326488, total: 88872591
hEB16d_HUES45_p21.RRBS
CancerGene_Promoter (#regions: 889) − Measurement Coverage
#measurements, region: 2167343, total: 89500253
hEB16d_H1_p38.RRBS
No coverage
1−4
5−9
10−24
25−49
50−99
>=100

NPS_13r_J13.RRBS
>100
No coverage
Characterizing Promoter Divergence Between RRS - Melanomonas Coverage
Pseudomonoecia, region: 210734..210734, total: 89900253

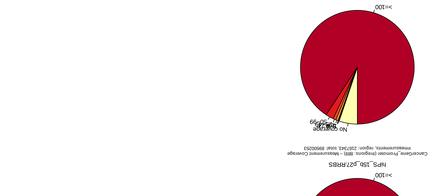

NPS_18r_J23.RRBS
>100
No coverage
Characterizing Promoter Divergence Between RRS - Melanomonas Coverage
Pseudomonoecia, region: 210734..210734, total: 89900253

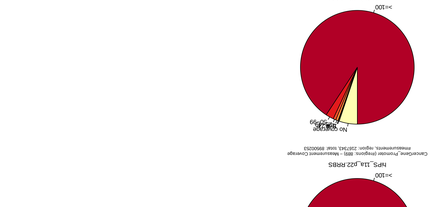
NPS_20r_J33.RRBS
>100
No coverage
Characterizing Promoter Divergence Between RRS - Melanomonas Coverage
Pseudomonoecia, region: 210734..210734, total: 89900253


NPS_22r_J43.RRBS
>100
No coverage
Characterizing Promoter Divergence Between RRS - Melanomonas Coverage
Pseudomonoecia, region: 210734..210734, total: 89900253

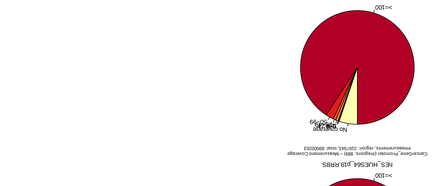

NPS_24r_J53.RRBS
>100
No coverage
Characterizing Promoter Divergence Between RRS - Melanomonas Coverage
Pseudomonoecia, region: 210734..210734, total: 89900253

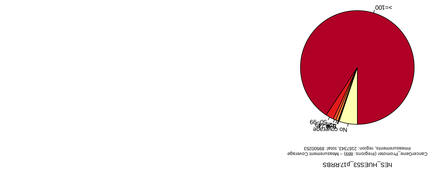
NPS_26r_J63.RRBS
>100
No coverage
Characterizing Promoter Divergence Between RRS - Melanomonas Coverage
Pseudomonoecia, region: 210734..210734, total: 89900253


NPS_28r_J73.RRBS
>100
No coverage
Characterizing Promoter Divergence Between RRS - Melanomonas Coverage
Pseudomonoecia, region: 210734..210734, total: 89900253


NPS_30r_J83.RRBS
>100
No coverage
Characterizing Promoter Divergence Between RRS - Melanomonas Coverage
Pseudomonoecia, region: 210734..210734, total: 89900253

NES_1r_J93.RRBS
>100
No coverage
Characterizing Promoter Divergence Between RRS - Melanomonas Coverage
Pseudomonoecia, region: 210734..210734, total: 89900253


NES_3r_J103.RRBS
>100
No coverage
Characterizing Promoter Divergence Between RRS - Melanomonas Coverage
Pseudomonoecia, region: 210734..210734, total: 89900253


NES_5r_J113.RRBS
>100
No coverage
Characterizing Promoter Divergence Between RRS - Melanomonas Coverage
Pseudomonoecia, region: 210734..210734, total: 89900253

NES_7r_J123.RRBS
>100
No coverage
Characterizing Promoter Divergence Between RRS - Melanomonas Coverage
Pseudomonoecia, region: 210734..210734, total: 89900253

NES_9r_J133.RRBS
>100
No coverage
Characterizing Promoter Divergence Between RRS - Melanomonas Coverage
Pseudomonoecia, region: 210734..210734, total: 89900253

NES_11r_J143.RRBS
>100
No coverage
Characterizing Promoter Divergence Between RRS - Melanomonas Coverage
Pseudomonoecia, region: 210734..210734, total: 89900253

NES_13r_J153.RRBS
>100
No coverage
Characterizing Promoter Divergence Between RRS - Melanomonas Coverage
Pseudomonoecia, region: 210734..210734, total: 89900253

NES_15r_J163.RRBS
>100
No coverage
Characterizing Promoter Divergence Between RRS - Melanomonas Coverage
Pseudomonoecia, region: 210734..210734, total: 89900253

NES_17r_J173.RRBS
>100
No coverage
Characterizing Promoter Divergence Between RRS - Melanomonas Coverage
Pseudomonoecia, region: 210734..210734, total: 89900253

NES_19r_J183.RRBS
>100
No coverage
Characterizing Promoter Divergence Between RRS - Melanomonas Coverage
Pseudomonoecia, region: 210734..210734, total: 89900253

NES_21r_J193.RRBS
>100
No coverage
Characterizing Promoter Divergence Between RRS - Melanomonas Coverage
Pseudomonoecia, region: 210734..210734, total: 89900253

NES_23r_J203.RRBS
>100
No coverage
Characterizing Promoter Divergence Between RRS - Melanomonas Coverage
Pseudomonoecia, region: 210734..210734, total: 89900253

NES_25r_J213.RRBS
>100
No coverage
Characterizing Promoter Divergence Between RRS - Melanomonas Coverage
Pseudomonoecia, region: 210734..210734, total: 89900253

NES_27r_J223.RRBS
>100
No coverage
Characterizing Promoter Divergence Between RRS - Melanomonas Coverage
Pseudomonoecia, region: 210734..210734, total: 89900253

NES_29r_J233.RRBS
>100
No coverage
Characterizing Promoter Divergence Between RRS - Melanomonas Coverage
Pseudomonoecia, region: 210734..210734, total: 89900253

NES_31r_J243.RRBS
>100
No coverage
Characterizing Promoter Divergence Between RRS - Melanomonas Coverage
Pseudomonoecia, region: 210734..210734, total: 89900253

NES_33r_J253.RRBS
>100
No coverage
Characterizing Promoter Divergence Between RRS - Melanomonas Coverage
Pseudomonoecia, region: 210734..210734, total: 89900253

NES_35r_J263.RRBS
>100
No coverage
Characterizing Promoter Divergence Between RRS - Melanomonas Coverage
Pseudomonoecia, region: 210734..210734, total: 89900253

NES_37r_J273.RRBS
>100
No coverage
Characterizing Promoter Divergence Between RRS - Melanomonas Coverage
Pseudomonoecia, region: 210734..210734, total: 89900253

NES_39r_J283.RRBS
>100
No coverage
Characterizing Promoter Divergence Between RRS - Melanomonas Coverage
Pseudomonoecia, region: 210734..210734, total: 89900253

NES_41r_J293.RRBS
>100
No coverage
Characterizing Promoter Divergence Between RRS - Melanomonas Coverage
Pseudomonoecia, region: 210734..210734, total: 89900253

NES_43r_J303.RRBS
>100
No coverage
Characterizing Promoter Divergence Between RRS - Melanomonas Coverage
Pseudomonoecia, region: 210734..210734, total: 89900253

NPS_11r_J13.RRBS
>100
No coverage
Characterizing Promoter Divergence Between RRS - Melanomonas Coverage
Pseudomonoecia, region: 210734..210734, total: 89900253

NPS_16r_J23.RRBS
>100
No coverage
Characterizing Promoter Divergence Between RRS - Melanomonas Coverage
Pseudomonoecia, region: 210734..210734, total: 89900253

NPS_18r_J33.RRBS
>100
No coverage
Characterizing Promoter Divergence Between RRS - Melanomonas Coverage
Pseudomonoecia, region: 210734..210734, total: 89900253

NPS_20r_J43.RRBS
>100
No coverage
Characterizing Promoter Divergence Between RRS - Melanomonas Coverage
Pseudomonoecia, region: 210734..210734, total: 89900253

NPS_22r_J53.RRBS
>100
No coverage
Characterizing Promoter Divergence Between RRS - Melanomonas Coverage
Pseudomonoecia, region: 210734..210734, total: 89900253

NPS_24r_J63.RRBS
>100
No coverage
Characterizing Promoter Divergence Between RRS - Melanomonas Coverage
Pseudomonoecia, region: 210734..210734, total: 89900253

NPS_26r_J73.RRBS
>100
No coverage
Characterizing Promoter Divergence Between RRS - Melanomonas Coverage
Pseudomonoecia, region: 210734..210734, total: 89900253

NPS_28r_J83.RRBS
>100
No coverage
Characterizing Promoter Divergence Between RRS - Melanomonas Coverage
Pseudomonoecia, region: 210734..210734, total: 89900253

NPS_30r_J93.RRBS
>100
No coverage
Characterizing Promoter Divergence Between RRS - Melanomonas Coverage
Pseudomonoecia, region: 210734..210734, total: 89900253

NPS_32r_J103.RRBS
>100
No coverage
Characterizing Promoter Divergence Between RRS - Melanomonas Coverage
Pseudomonoecia, region: 210734..210734, total: 89900253

NPS_34r_J113.RRBS
>100
No coverage
Characterizing Promoter Divergence Between RRS - Melanomonas Coverage
Pseudomonoecia, region: 210734..210734, total: 89900253

NPS_36r_J123.RRBS
>100
No coverage
Characterizing Promoter Divergence Between RRS - Melanomonas Coverage
Pseudomonoecia, region: 210734..210734, total: 89900253

NPS_38r_J133.RRBS
>100
No coverage
Characterizing Promoter Divergence Between RRS - Melanomonas Coverage
Pseudomonoecia, region: 210734..210734, total: 89900253

NPS_40r_J143.RRBS
>100
No coverage
Characterizing Promoter Divergence Between RRS - Melanomonas Coverage
Pseudomonoecia, region: 210734..210734, total: 89900253

NPS_42r_J153.RRBS
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No coverage
Characterizing Promoter Divergence Between RRS - Melanomonas Coverage
Pseudomonoecia, region: 210734..210734, total: 89900253

NPS_44r_J163.RRBS
>100
No coverage
Characterizing Promoter Divergence Between RRS - Melanomonas Coverage
Pseudomonoecia, region: 210734..210734, total: 89900253

NES_2r_J98.RRBS
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No coverage
Characterizing Promoter Divergence Between RRS - Melanomonas Coverage
Pseudomonoecia, region: 210734..210734, total: 89900253

NES_4r_J108.RRBS
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No coverage
Characterizing Promoter Divergence Between RRS - Melanomonas Coverage
Pseudomonoecia, region: 210734..210734, total: 89900253

NES_6r_J118.RRBS
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No coverage
Characterizing Promoter Divergence Between RRS - Melanomonas Coverage
Pseudomonoecia, region: 210734..210734, total: 89900253

NES_8r_J128.RRBS
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No coverage
Characterizing Promoter Divergence Between RRS - Melanomonas Coverage
Pseudomonoecia, region: 210734..210734, total: 89900253

NES_10r_J138.RRBS
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No coverage
Characterizing Promoter Divergence Between RRS - Melanomonas Coverage
Pseudomonoecia, region: 210734..210734, total: 89900253

NES_12r_J148.RRBS
>100
No coverage
Characterizing Promoter Divergence Between RRS - Melanomonas Coverage
Pseudomonoecia, region: 210734..210734, total: 89900253

NES_14r_J158.RRBS
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No coverage
Characterizing Promoter Divergence Between RRS - Melanomonas Coverage
Pseudomonoecia, region: 210734..210734, total: 89900253

NES_16r_J168.RRBS
>100
No coverage
Characterizing Promoter Divergence Between RRS - Melanomonas Coverage
Pseudomonoecia, region: 210734..210734, total: 89900253

NES_18r_J178.RRBS
>100
No coverage
Characterizing Promoter Divergence Between RRS - Melanomonas Coverage
Pseudomonoecia, region: 210734..210734, total: 89900253

NES_20r_J188.RRBS
>100
No coverage
Characterizing Promoter Divergence Between RRS - Melanomonas Coverage
Pseudomonoecia, region: 210734..210734, total: 89900253

NES_22r_J198.RRBS
>100
No coverage
Characterizing Promoter Divergence Between RRS - Melanomonas Coverage
Pseudomonoecia, region: 210734..210734, total: 89900253

NES_24r_J208.RRBS
>100
No coverage
Characterizing Promoter Divergence Between RRS - Melanomonas Coverage
Pseudomonoecia, region: 210734..210734, total: 89900253

NES_26r_J218.RRBS
>100
No coverage
Characterizing Promoter Divergence Between RRS - Melanomonas Coverage
Pseudomonoecia, region: 210734..210734, total: 89900253

NES_28r_J228.RRBS
>100
No coverage
Characterizing Promoter Divergence Between RRS - Melanomonas Coverage
Pseudomonoecia, region: 210734..210734, total: 89900253

NEB1b6_H1_J28.RRBS
>100
No coverage
Characterizing Promoter Divergence Between RRS - Melanomonas Coverage
Pseudomonoecia, region: 210734..210734, total: 89900253

NEB1b6_HUES3_J22.RRBS
>100
No coverage
Characterizing Promoter Divergence Between RRS - Melanomonas Coverage
Pseudomonoecia, region: 210734..210734, total: 89900253

NFib_20_J17.RRBS
>100
No coverage
Characterizing Promoter Divergence Between RRS - Melanomonas Coverage
Pseudomonoecia, region: 210734..210734, total: 89900253

NFib_15_J17.RRBS
>100
No coverage
Characterizing Promoter Divergence Between RRS - Melanomonas Coverage
Pseudomonoecia, region: 210734..210734, total: 89900253

NPS_27r_J22.RRBS
>100
No coverage
Characterizing Promoter Divergence Between RRS - Melanomonas Coverage
Pseudomonoecia, region: 210734..210734, total: 89900253

NPS_18r_J36.RRBS
>100
No coverage
Characterizing Promoter Divergence Between RRS - Melanomonas Coverage
Pseudomonoecia, region: 210734..210734, total: 89900253

NPS_17r_J29.RRBS
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No coverage
Characterizing Promoter Divergence Between RRS - Melanomonas Coverage
Pseudomonoecia, region: 210734..210734, total: 89900253

NES_9r_J49.RRBS
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No coverage
Characterizing Promoter Divergence Between RRS - Melanomonas Coverage
Pseudomonoecia, region: 210734..210734, total: 89900253

NES_7r_J49.RRBS
>100
No coverage
Characterizing Promoter Divergence Between RRS - Melanomonas Coverage
Pseudomonoecia, region: 210734..210734, total: 89900253

NES_5r_J39.RRBS
>100
No coverage
Characterizing Promoter Divergence Between RRS - Melanomonas Coverage
Pseudomonoecia, region: 210734..210734, total: 89900253

NES_3r_J29.RRBS
>100
No coverage
Characterizing Promoter Divergence Between RRS - Melanomonas Coverage
Pseudomonoecia, region: 210734..210734, total: 89900253

NES_1r_J19.RRBS
>100
No coverage
Characterizing Promoter Divergence Between RRS - Melanomonas Coverage
Pseudomonoecia, region: 210734..210734, total: 89900253

NEB1b6_HUES45_J21.RRBS
>100
No coverage
Characterizing Promoter Divergence Between RRS - Melanomonas Coverage
Pseudomonoecia, region: 210734..210734, total: 89900253

NEB1b6_HUES56_J30.RRBS
>100
No coverage
Characterizing Promoter Divergence Between RRS - Melanomonas Coverage
Pseudomonoecia, region: 210734..210734, total: 89900253

NEB1b6_HUES53_J29.RRBS
>100
No coverage
Characterizing Promoter Divergence Between RRS - Melanomonas Coverage
Pseudomonoecia, region: 210734..210734, total: 89900253

NPS_15r_J27.RRBS
>100
No coverage
Characterizing Promoter Divergence Between RRS - Melanomonas Coverage
Pseudomonoecia, region: 210734..210734, total: 89900253

NFib_13_J27.RRBS
>100
No coverage
Characterizing Promoter Divergence Between RRS - Melanomonas Coverage
Pseudomonoecia, region: 210734..210734, total: 89900253

NFib_11_J27.RRBS
>100
No coverage
Characterizing Promoter Divergence Between RRS - Melanomonas Coverage
Pseudomonoecia, region: 210734..210734, total: 89900253

NFib_9_J27.RRBS
>100
No coverage
Characterizing Promoter Divergence Between RRS - Melanomonas Coverage
Pseudomonoecia, region: 210734..210734, total: 89900253

NFib_7_J27.RRBS
>100
No coverage
Characterizing Promoter Divergence Between RRS - Melanomonas Coverage
Pseudomonoecia, region: 210734..210734, total: 89900253

NFib_5_J27.RRBS
>100
No coverage
Characterizing Promoter Divergence Between RRS - Melanomonas Coverage
Pseudomonoecia, region: 210734..210734, total: 89900253

NFib_3_J27.RRBS
>100
No coverage
Characterizing Promoter Divergence Between RRS - Melanomonas Coverage
Pseudomonoecia, region: 210734..210734, total: 89900253

NFib_1_J27.RRBS
>100
No coverage
Characterizing Promoter Divergence Between RRS - Melanomonas Coverage
Pseudomonoecia, region: 210734..210734, total: 89900253

NPS_13r_J27.RRBS
>100
No coverage
Characterizing Promoter Divergence Between RRS - Melanomonas Coverage
Pseudomonoecia, region: 210734..210734, total: 89900253

NPS_11r_J27.RRBS
>100
No coverage
Characterizing Promoter Divergence Between RRS - Melanomonas Coverage
Pseudomonoecia, region: 210734..210734, total: 89900253

NES_2r_J27.RRBS
>100
No coverage
Characterizing Promoter Divergence Between RRS - Melanomonas Coverage
Pseudomonoecia, region: 210734..210734, total: 89900253

NES_4r_J27.RRBS
>100
No coverage
Characterizing Promoter Divergence Between RRS - Melanomonas Coverage
Pseudomonoecia, region: 210734..210734, total: 89900253

NES_6r_J27.RRBS
>100
No coverage
Characterizing Promoter Divergence Between RRS - Melanomonas Coverage
Pseudomonoecia, region: 210734..210734, total: 89900253

NES_8r_J27.RRBS
>100
No coverage
Characterizing Promoter Divergence Between RRS - Melanomonas Coverage
Pseudomonoecia, region: 210734..210734, total: 89900253

NES_10r_J27.RRBS
>100
No coverage
Characterizing Promoter Divergence Between RRS - Melanomonas Coverage
Pseudomonoecia, region: 210734..210734, total: 89900253