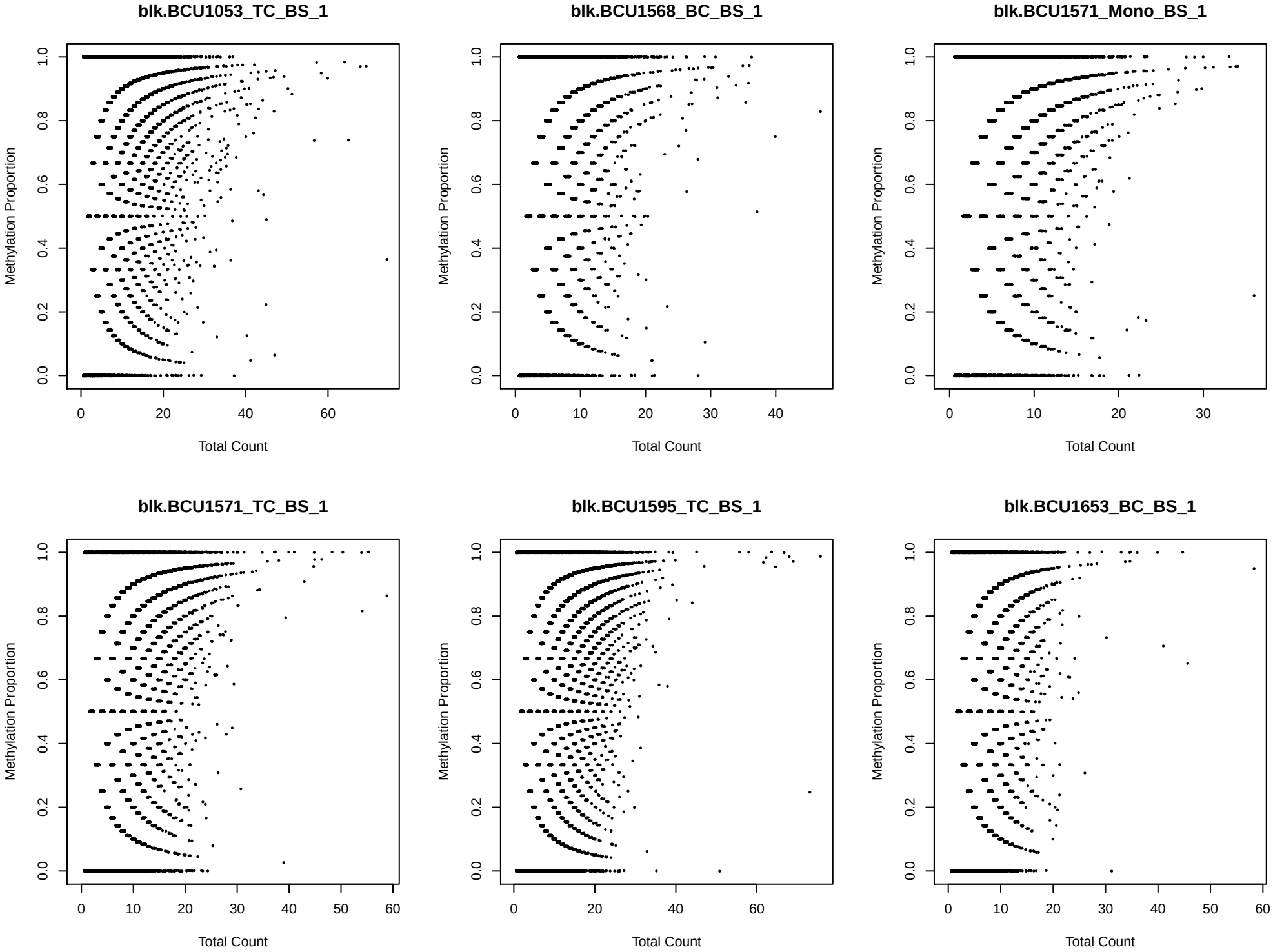
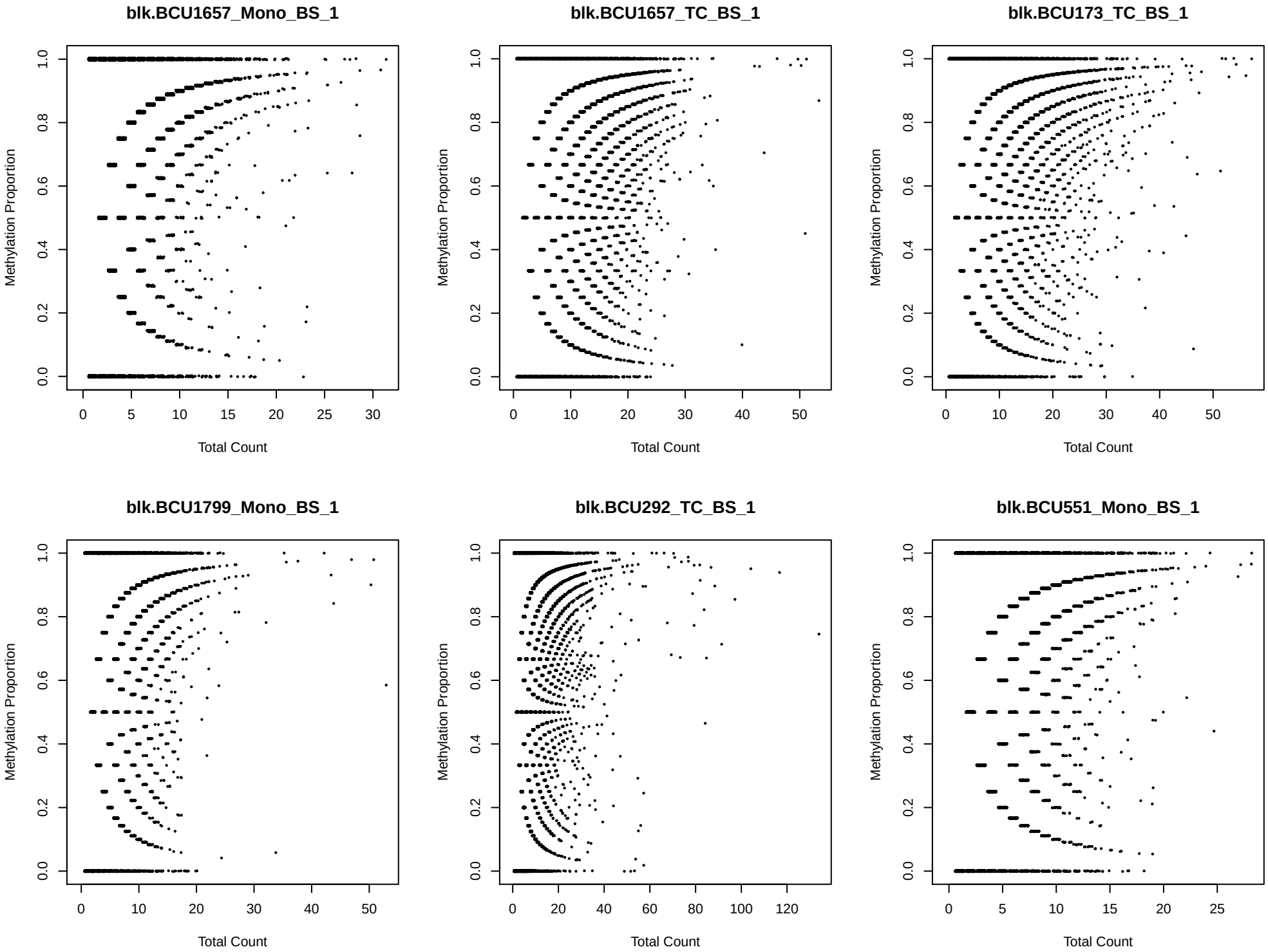
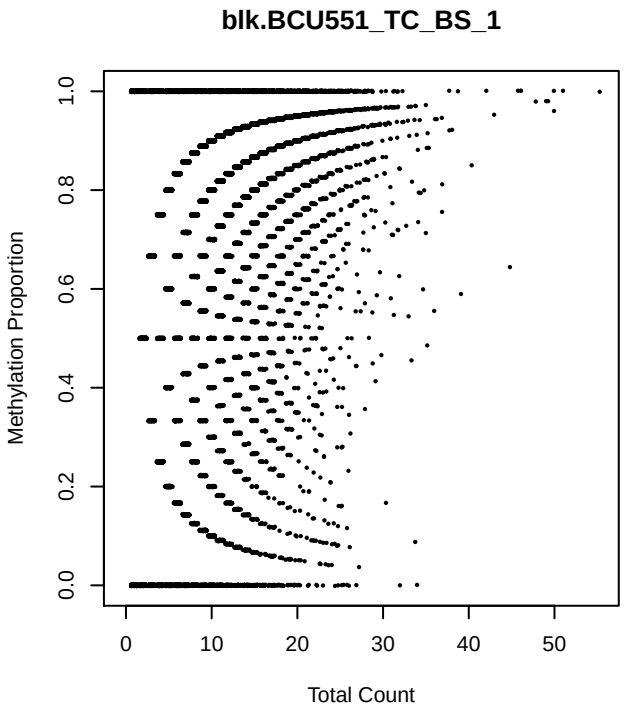
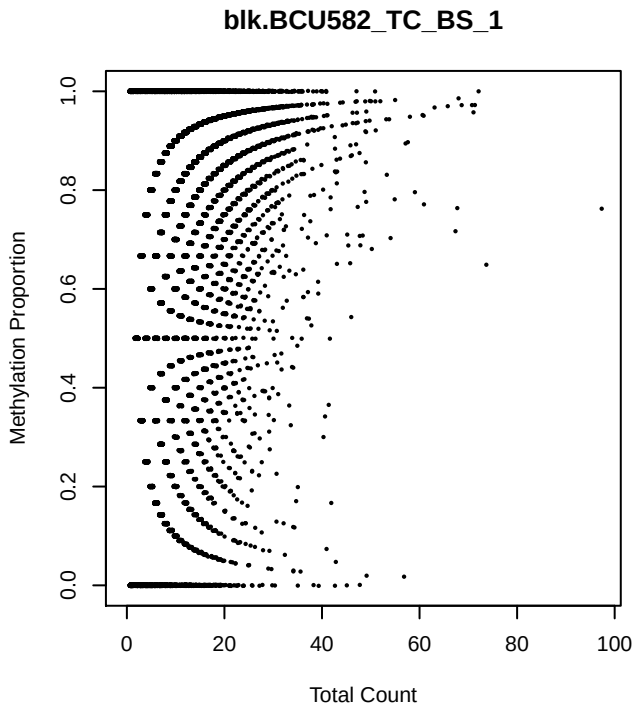
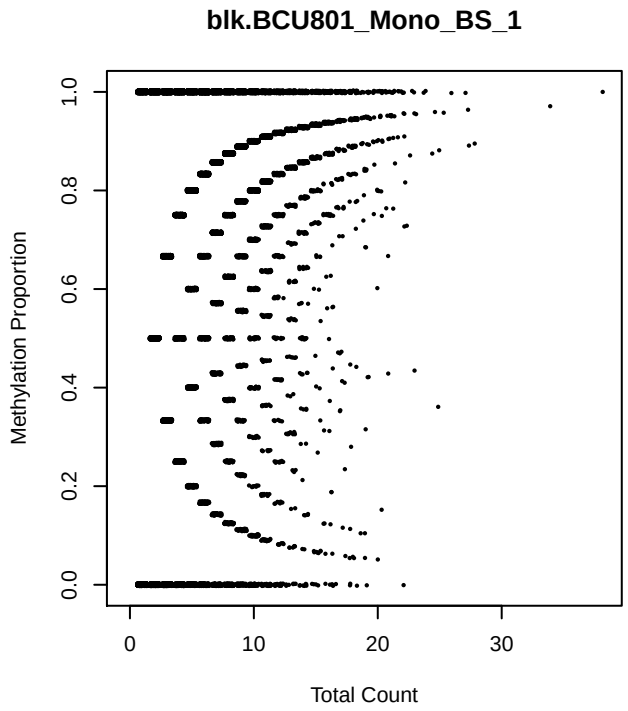
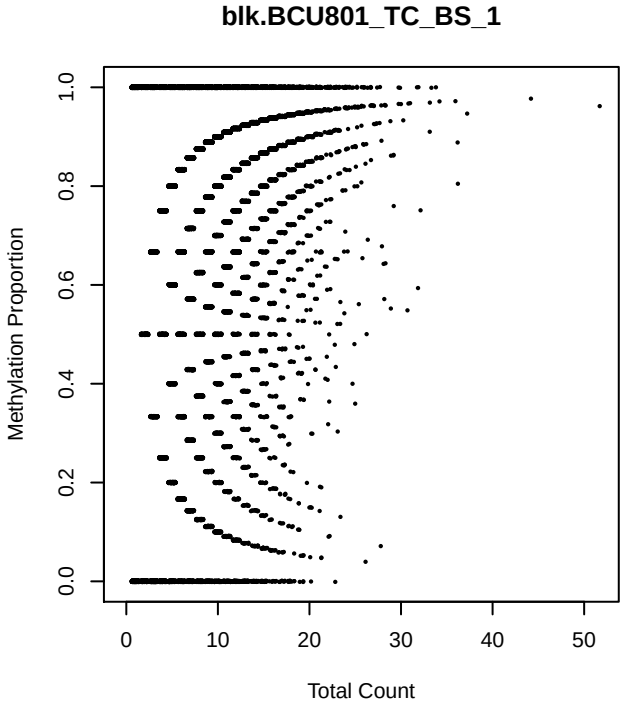

blk.BCU1053_TC_BS_1
blk.BCU1568_BC_BS_1
blk.BCU1571_Mono_BS_1
blk.BCU1571_TC_BS_1
blk.BCU1595_TC_BS_1
blk.BCU1653_BC_BS_1
Methylation Proportion
Total Count

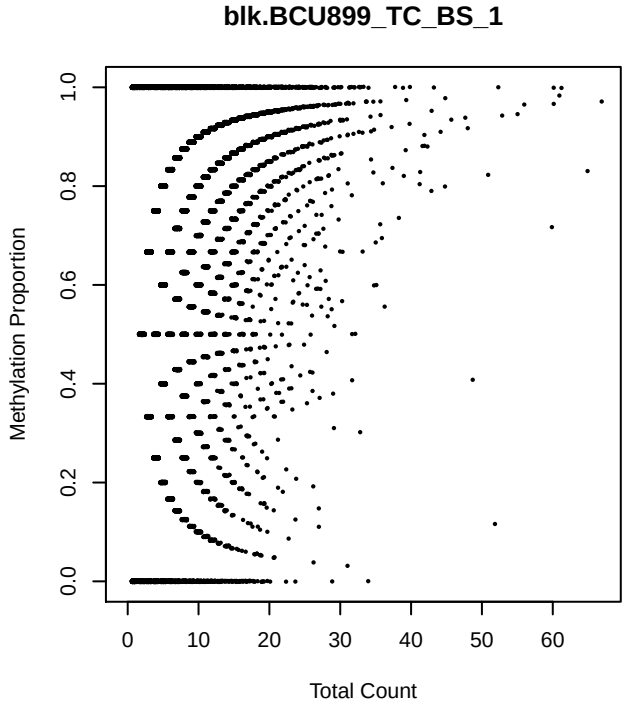

blk.BCU1657_Mono_BS_1
blk.BCU1657_TC_BS_1
blk.BCU173_TC_BS_1
blk.BCU1799_Mono_BS_1
blk.BCU292_TC_BS_1
blk.BCU551_Mono_BS_1
Methylation Proportion
Total Count

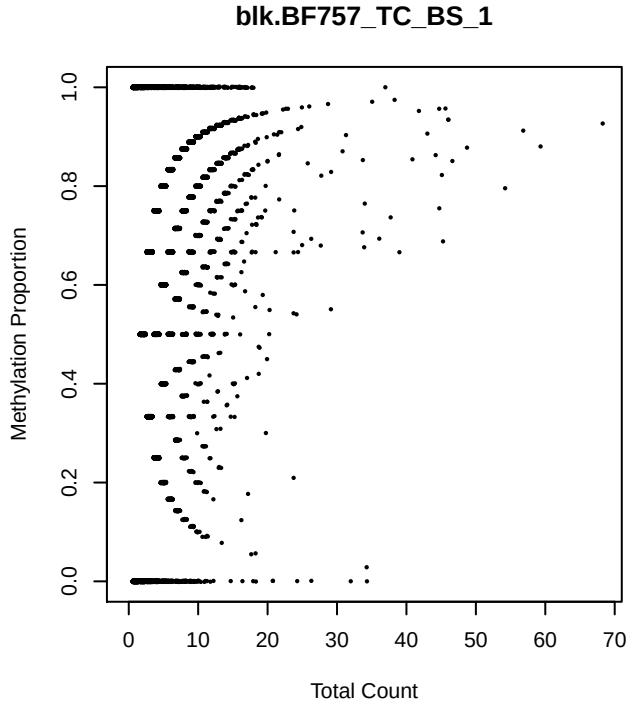

blk.BCU551_TC_BS_1
Methylation Proportion
Total Count

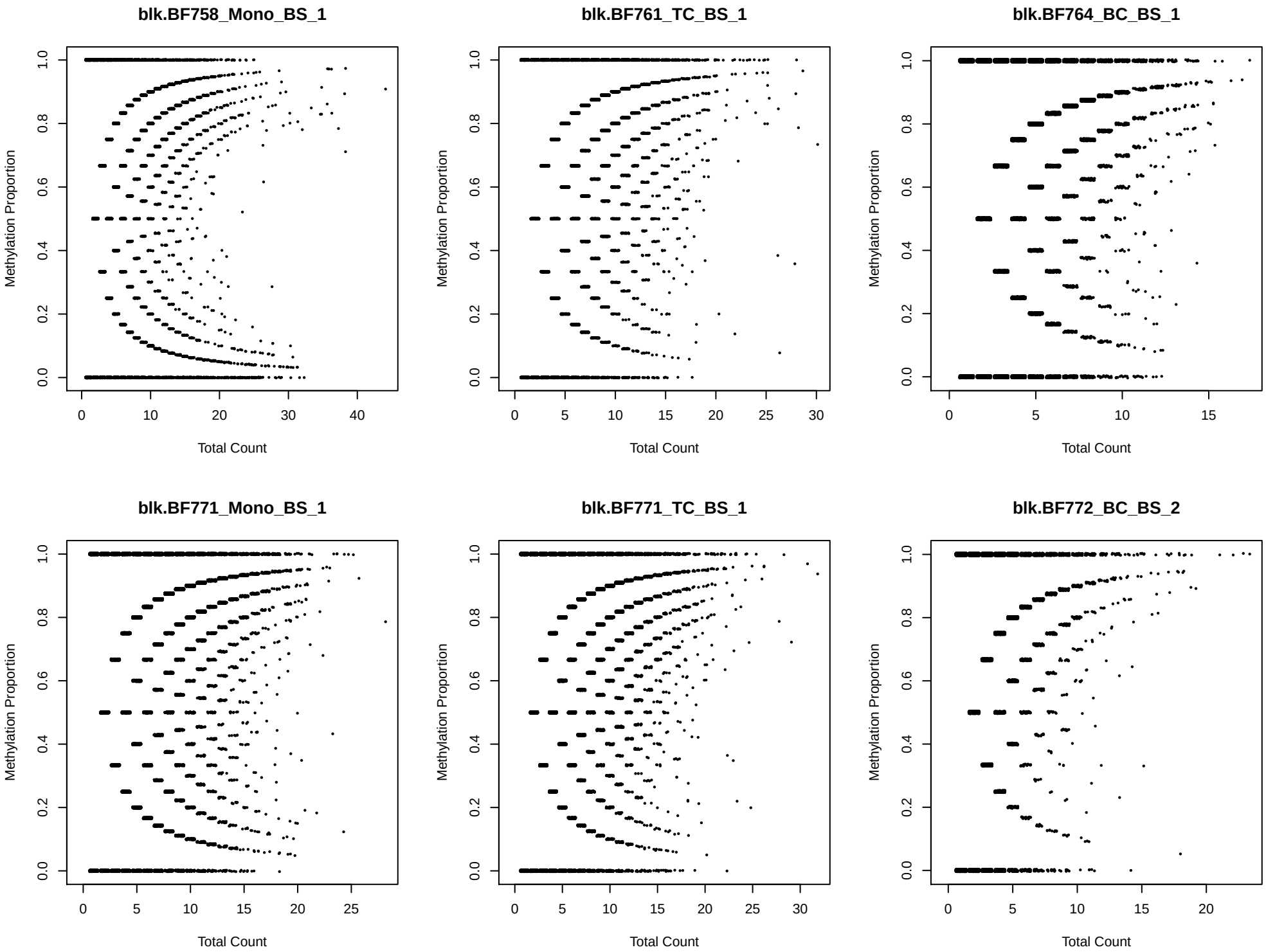

blk.BCU582_TC_BS_1
Methylation Proportion
Total Count

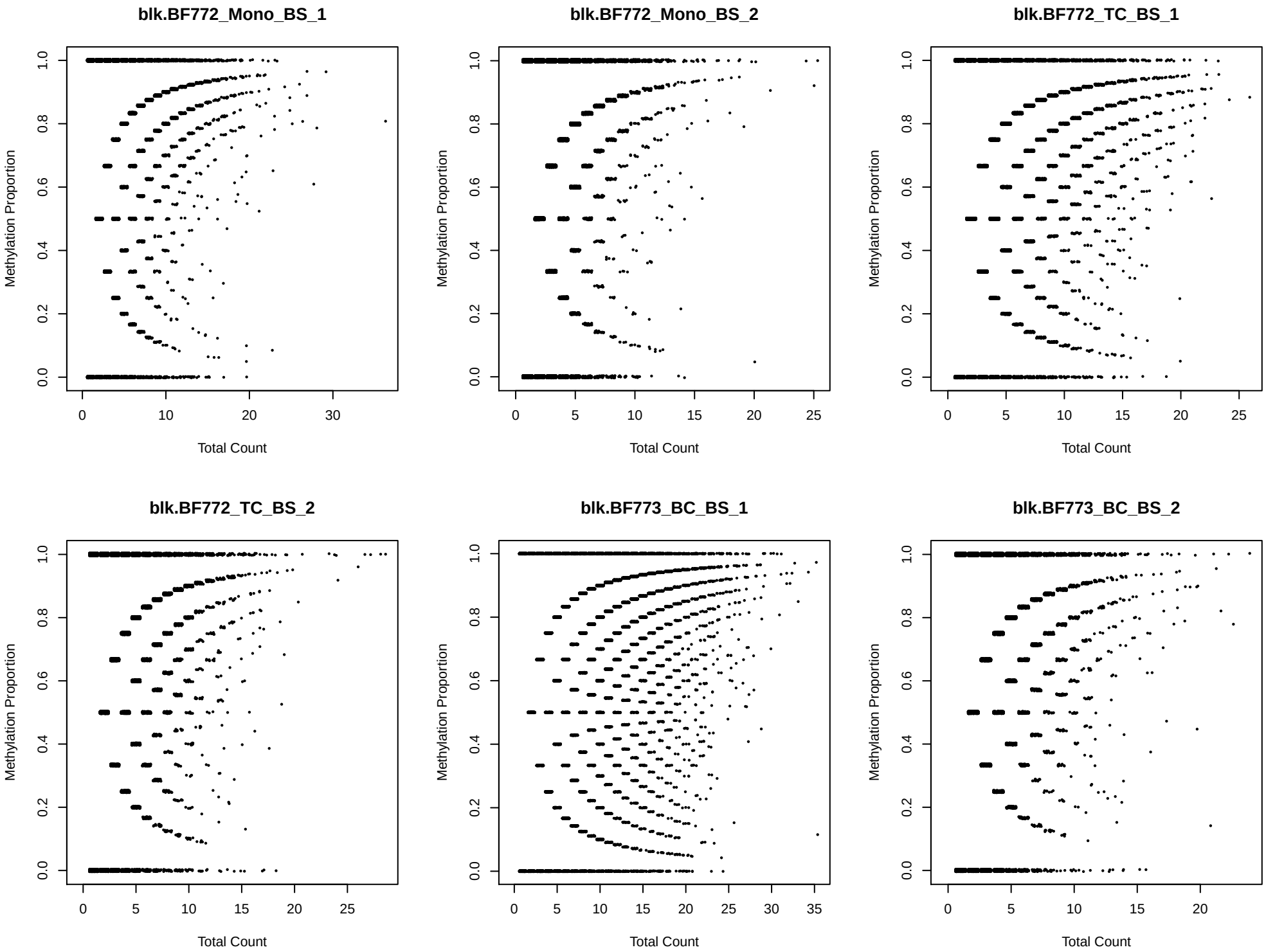

blk.BCU801_Mono_BS_1
Methylation Proportion
Total Count

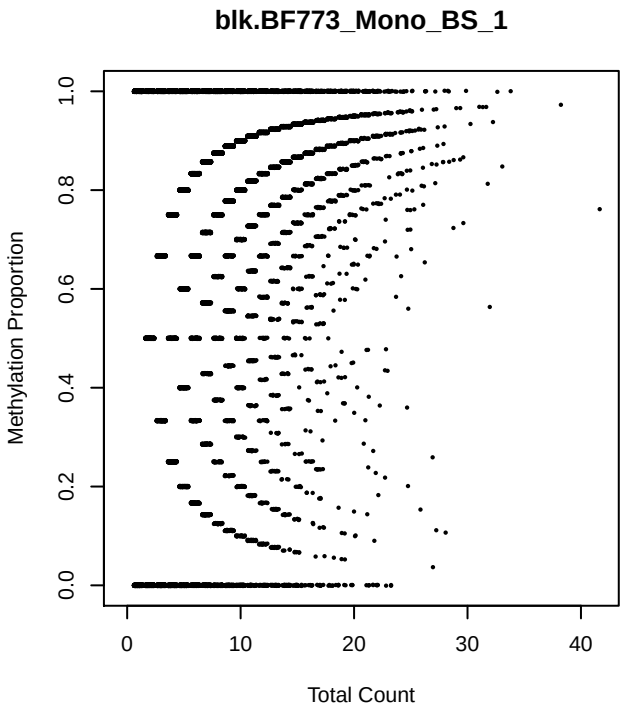

blk.BCU801_TC_BS_1
Methylation Proportion
Total Count

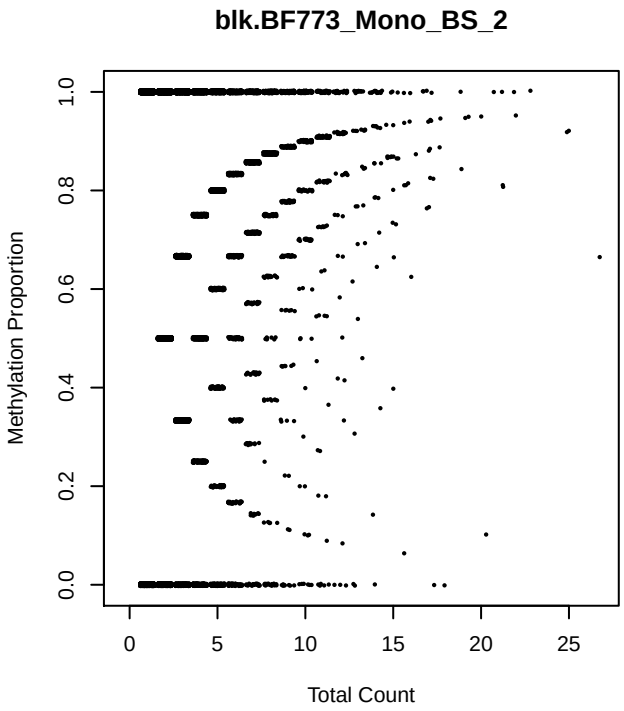

blk.BCU899_TC_BS_1
Methylation Proportion
Total Count

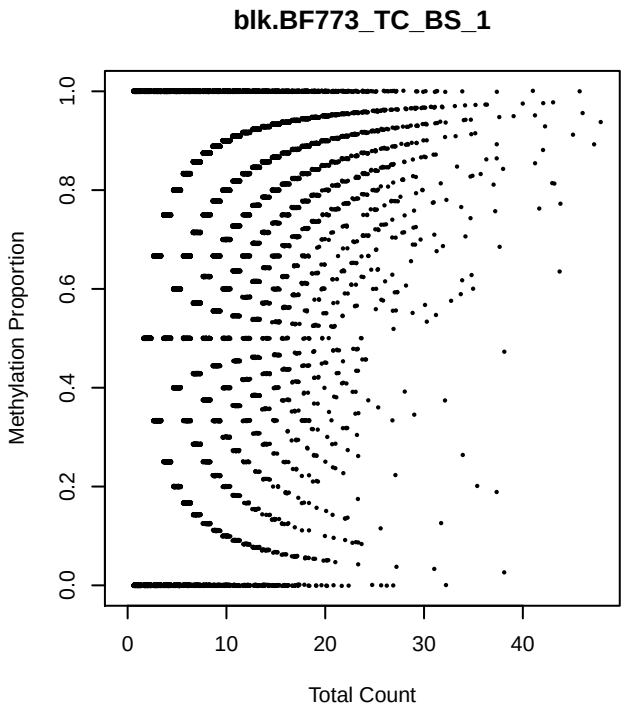

blk.BF757_TC_BS_1
Methylation Proportion
Total Count

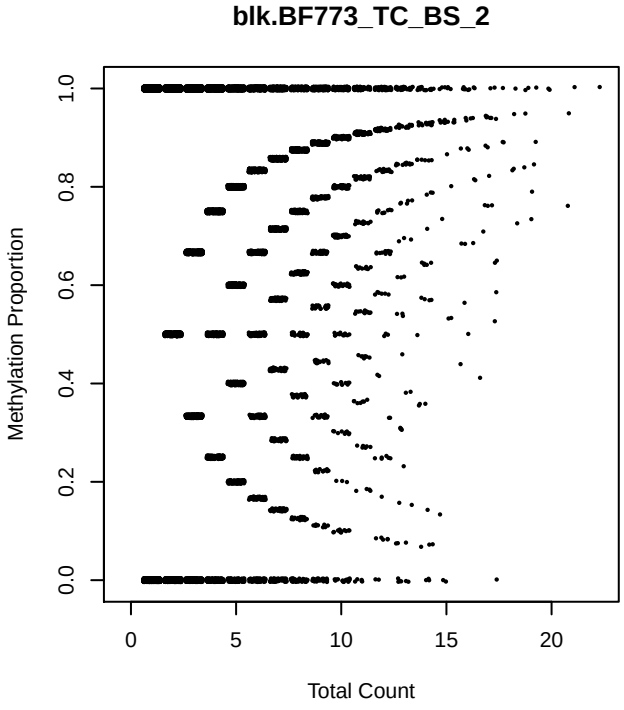

blk.BF758_Mono_BS_1
blk.BF761_TC_BS_1
blk.BF764_BC_BS_1
blk.BF771_Mono_BS_1
blk.BF771_TC_BS_1
blk.BF772_BC_BS_2
Methylation Proportion
Total Count

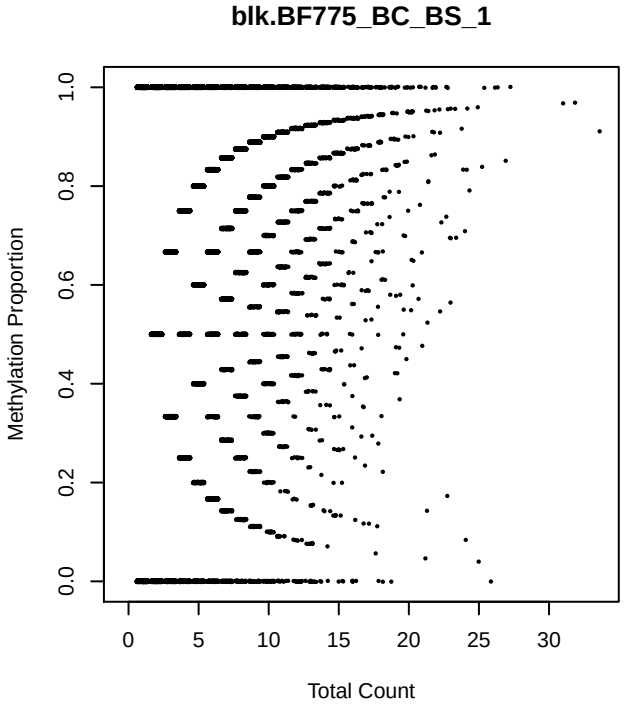

blk.BF772_Mono_BS_1
blk.BF772_Mono_BS_2
blk.BF772_TC_BS_1
blk.BF772_TC_BS_2
blk.BF773_BC_BS_1
blk.BF773_BC_BS_2
Methylation Proportion
Total Count

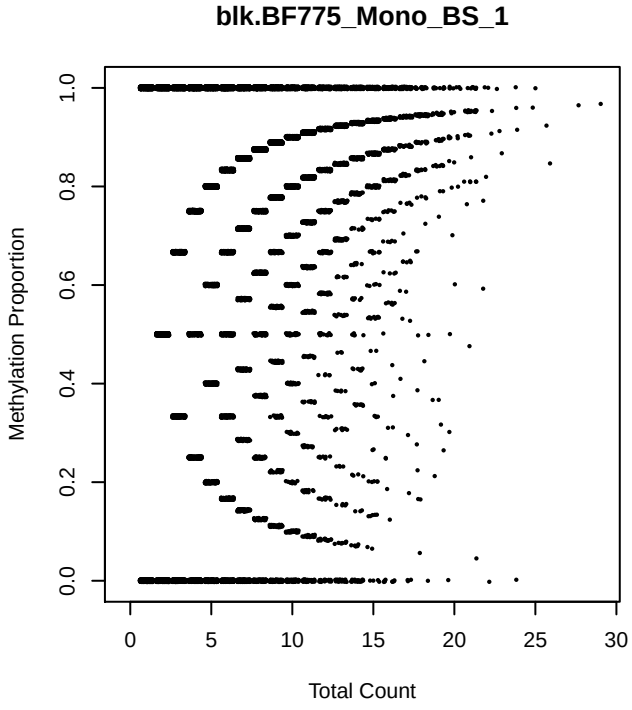

blk.BF773_Mono_BS_1
Methylation Proportion
Total Count

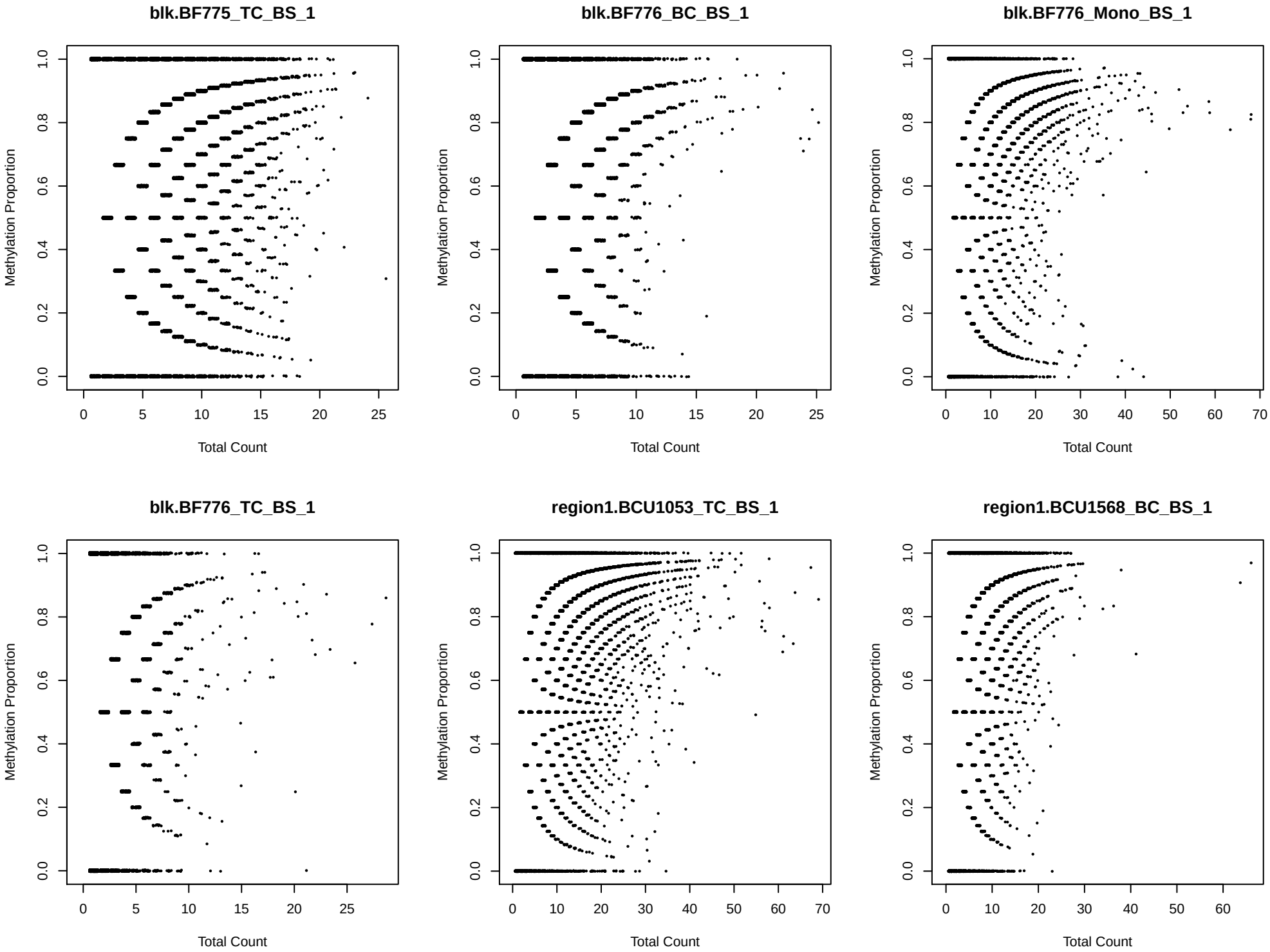

blk.BF773_Mono_BS_2
Methylation Proportion
Total Count

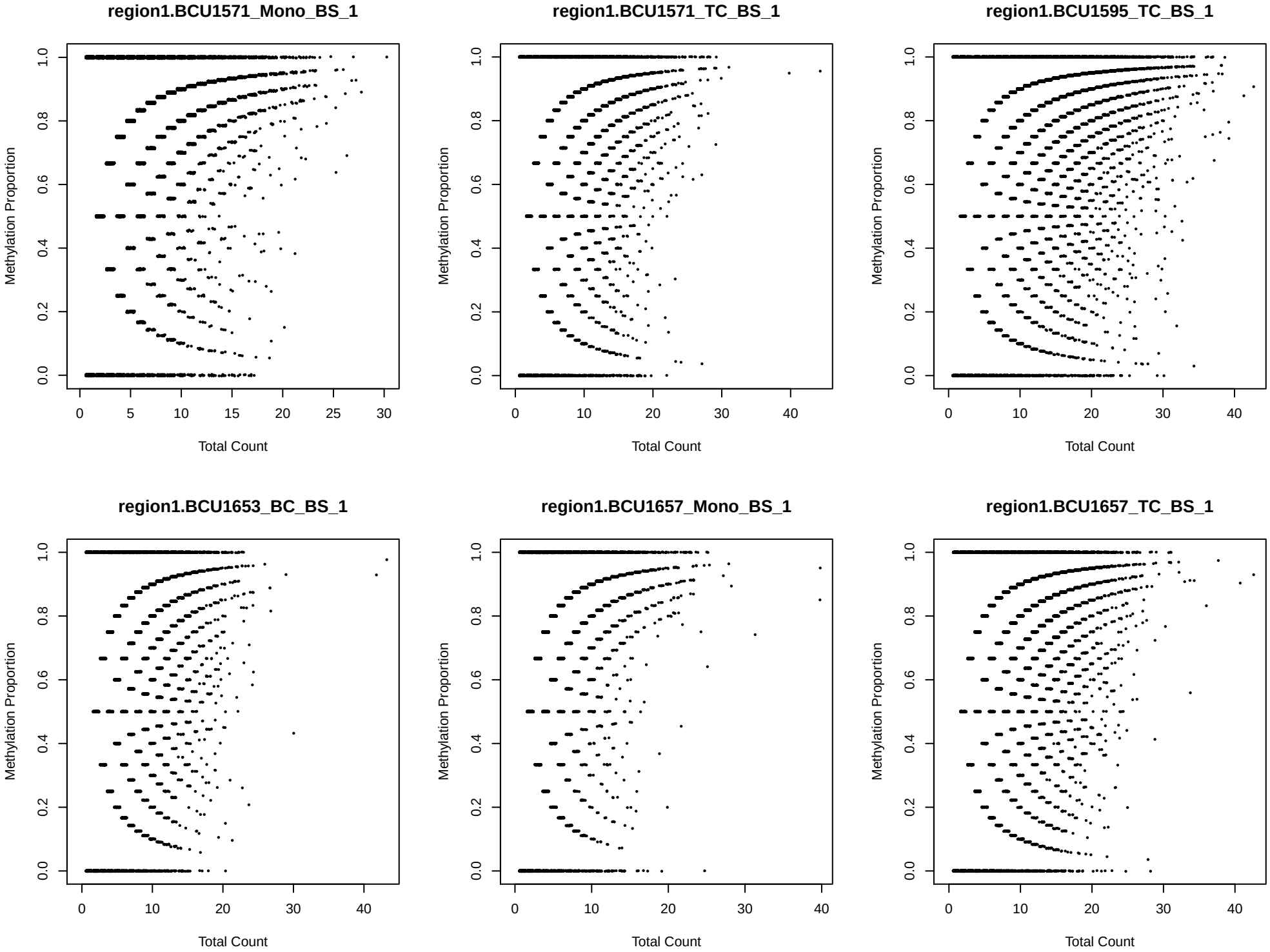

blk.BF773_TC_BS_1
Methylation Proportion
Total Count

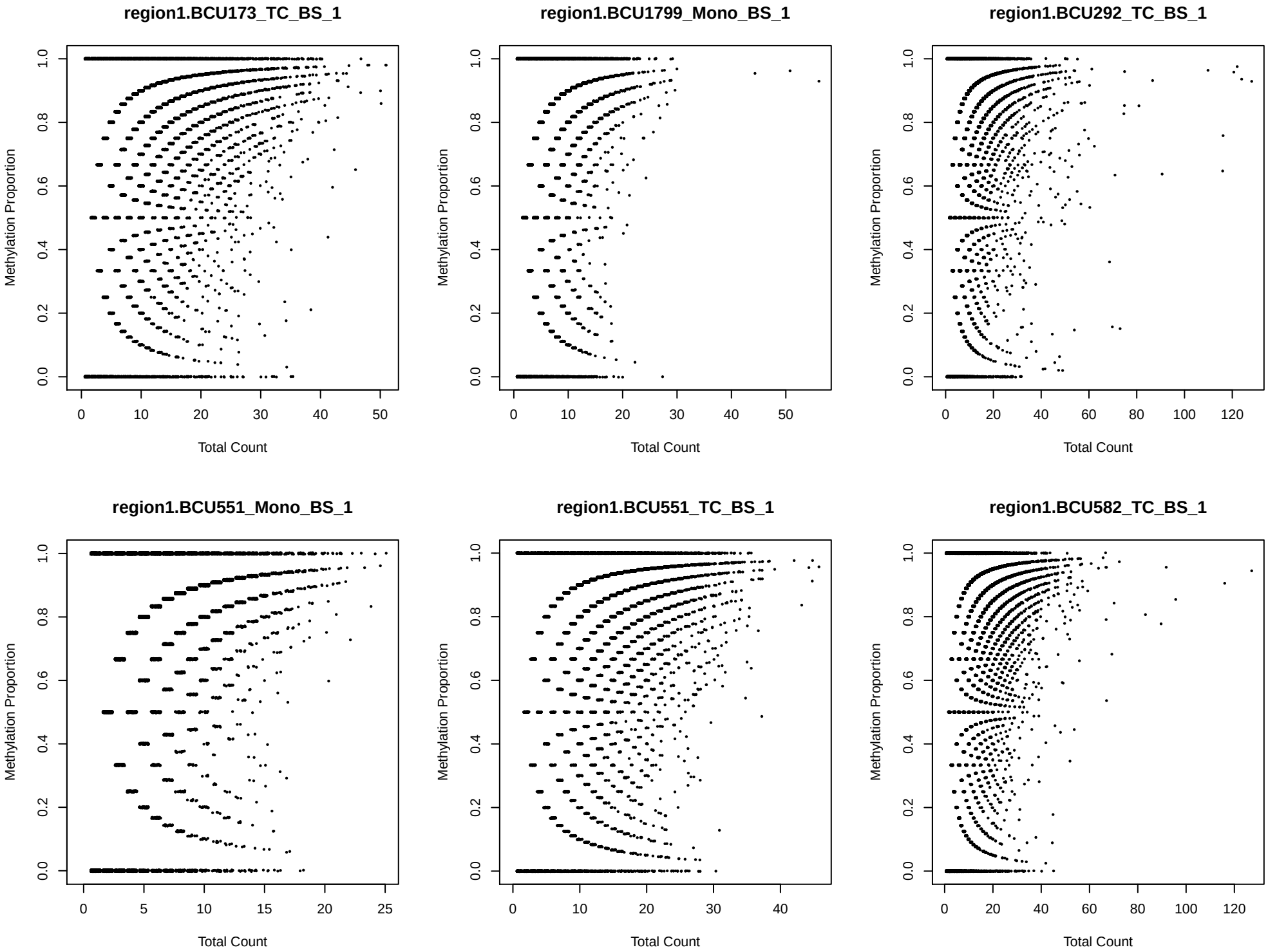

blk.BF773_TC_BS_2
Methylation Proportion
Total Count

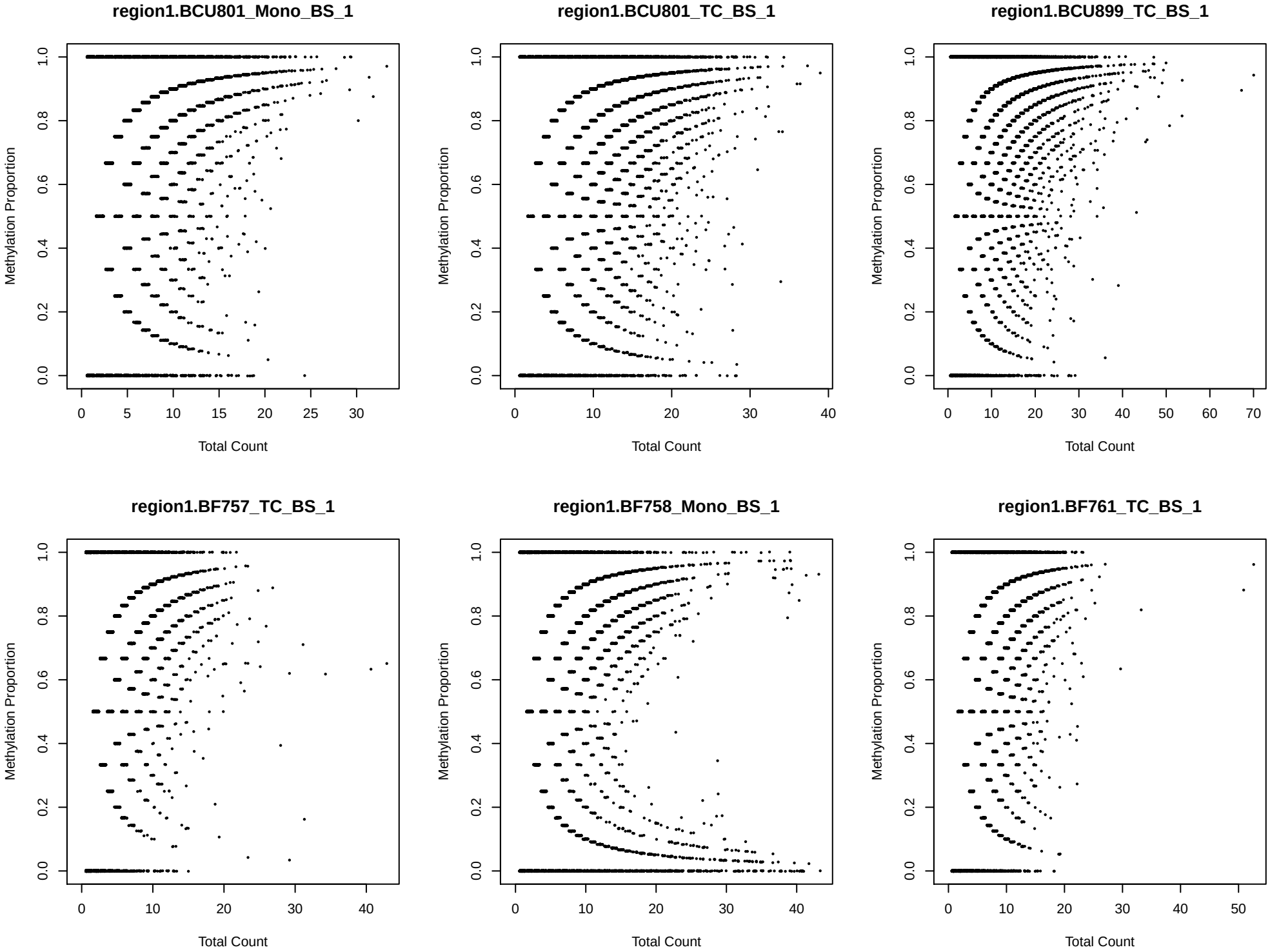

blk.BF775_BC_BS_1
Methylation Proportion
Total Count

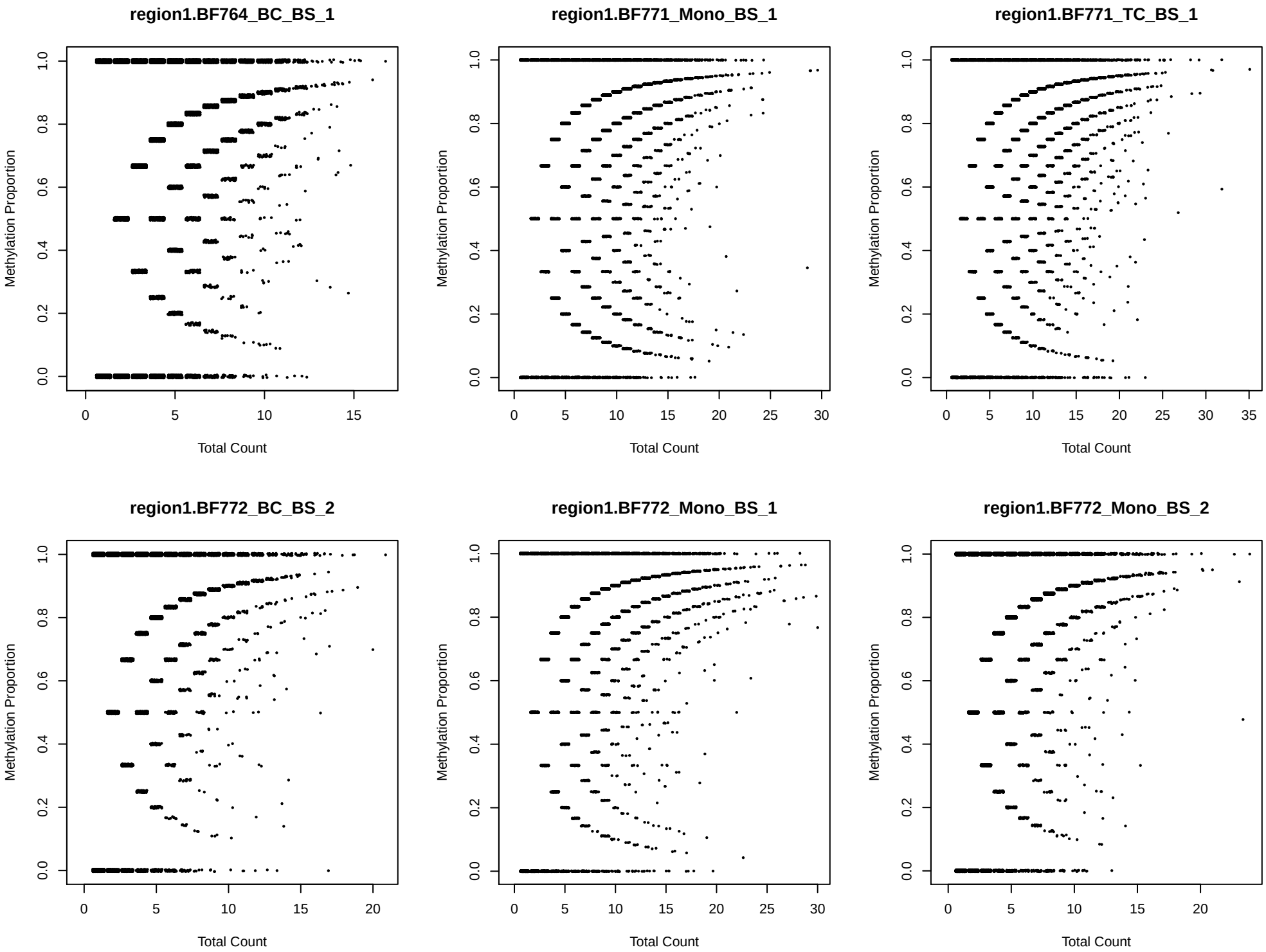

blk.BF775_Mono_BS_1
Methylation Proportion
Total Count

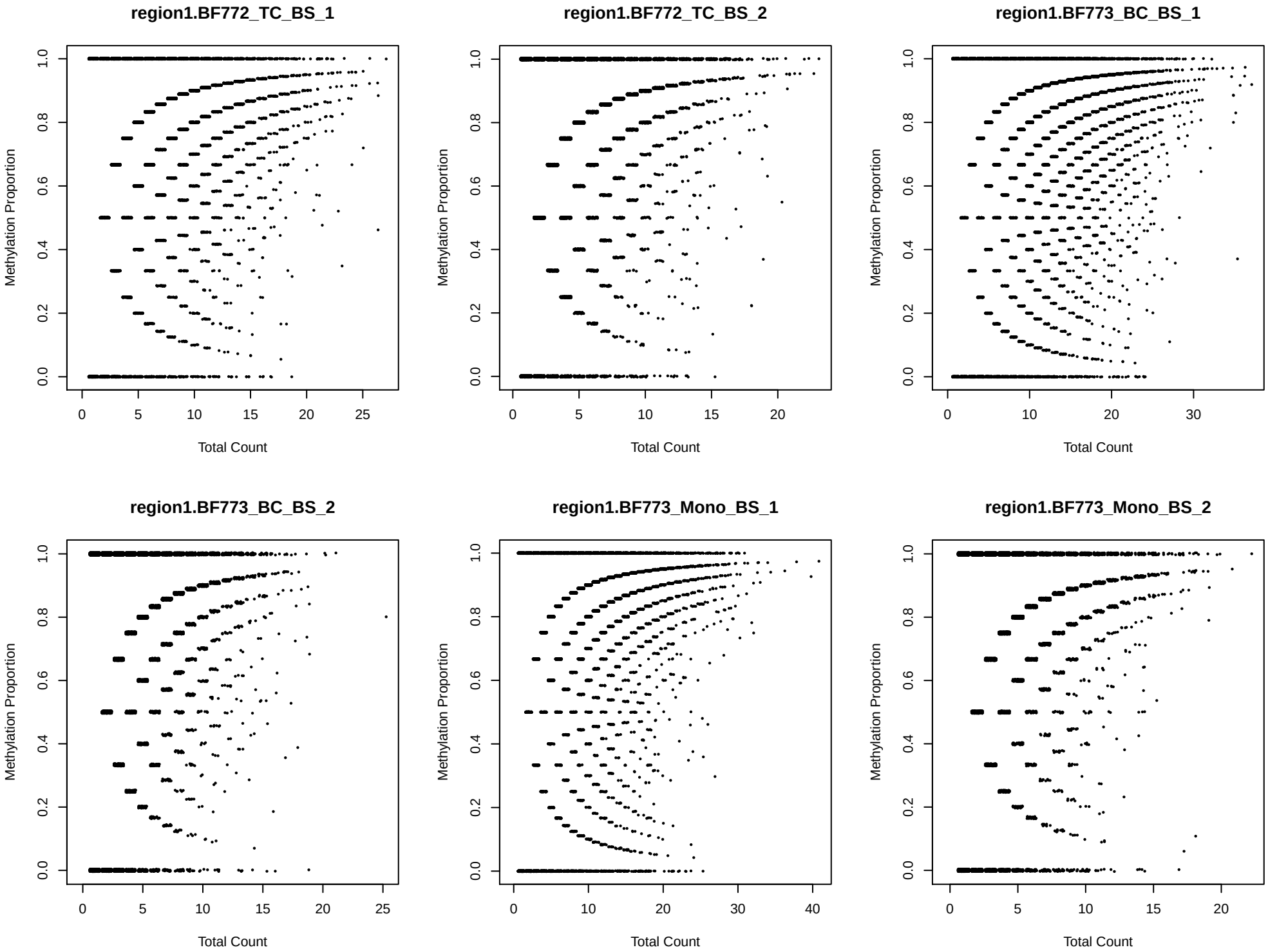

blk.BF775_TC_BS_1
blk.BF776_BC_BS_1
blk.BF776_Mono_BS_1
blk.BF776_TC_BS_1
region1.BCU1053_TC_BS_1
region1.BCU1568_BC_BS_1
Methylation Proportion
Total Count

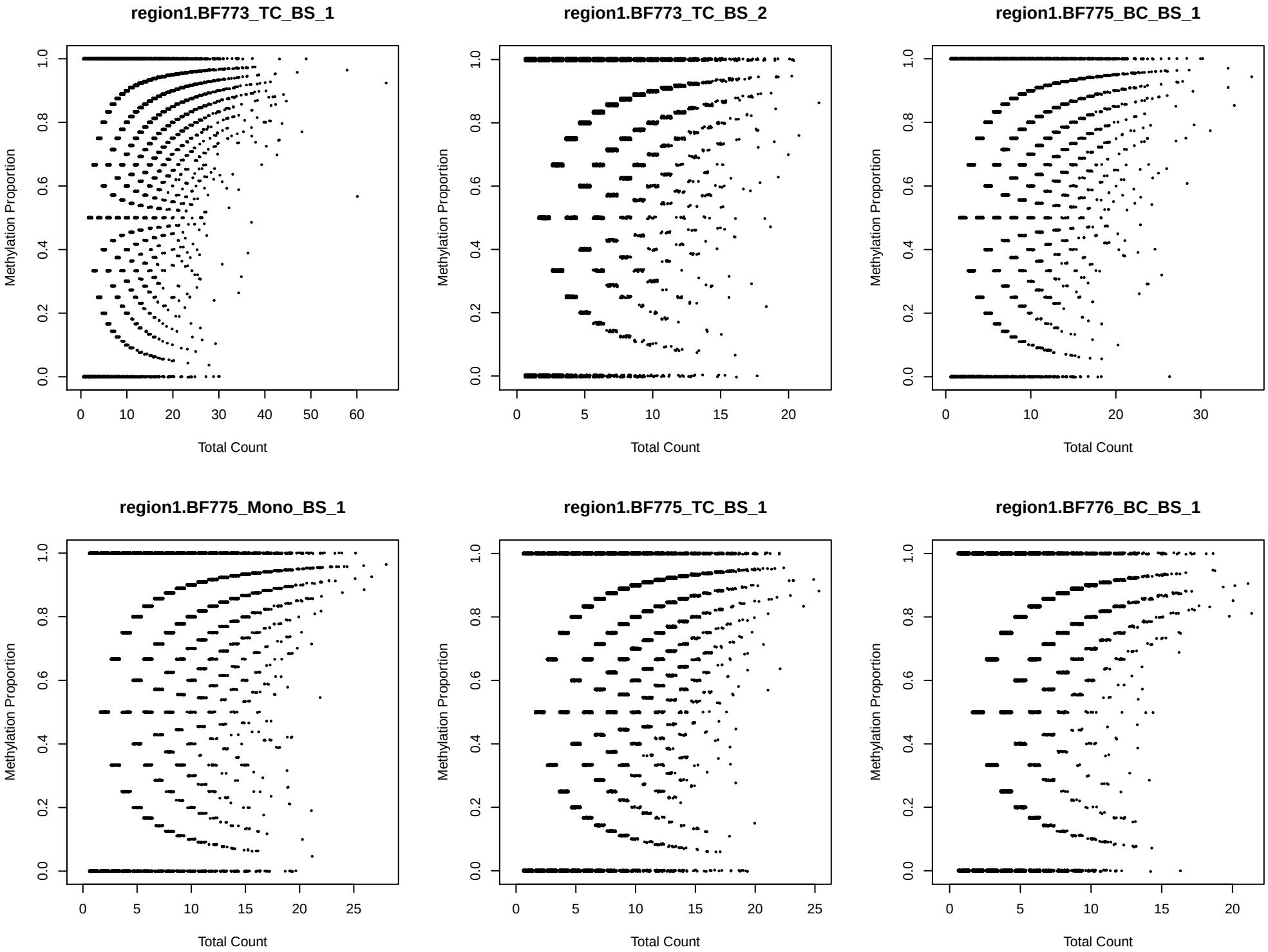

region1.BCU1571_Mono_BS_1
region1.BCU1571_TC_BS_1
region1.BCU1595_TC_BS_1
region1.BCU1653_BC_BS_1
region1.BCU1657_Mono_BS_1
region1.BCU1657_TC_BS_1
Methylation Proportion
Total Count

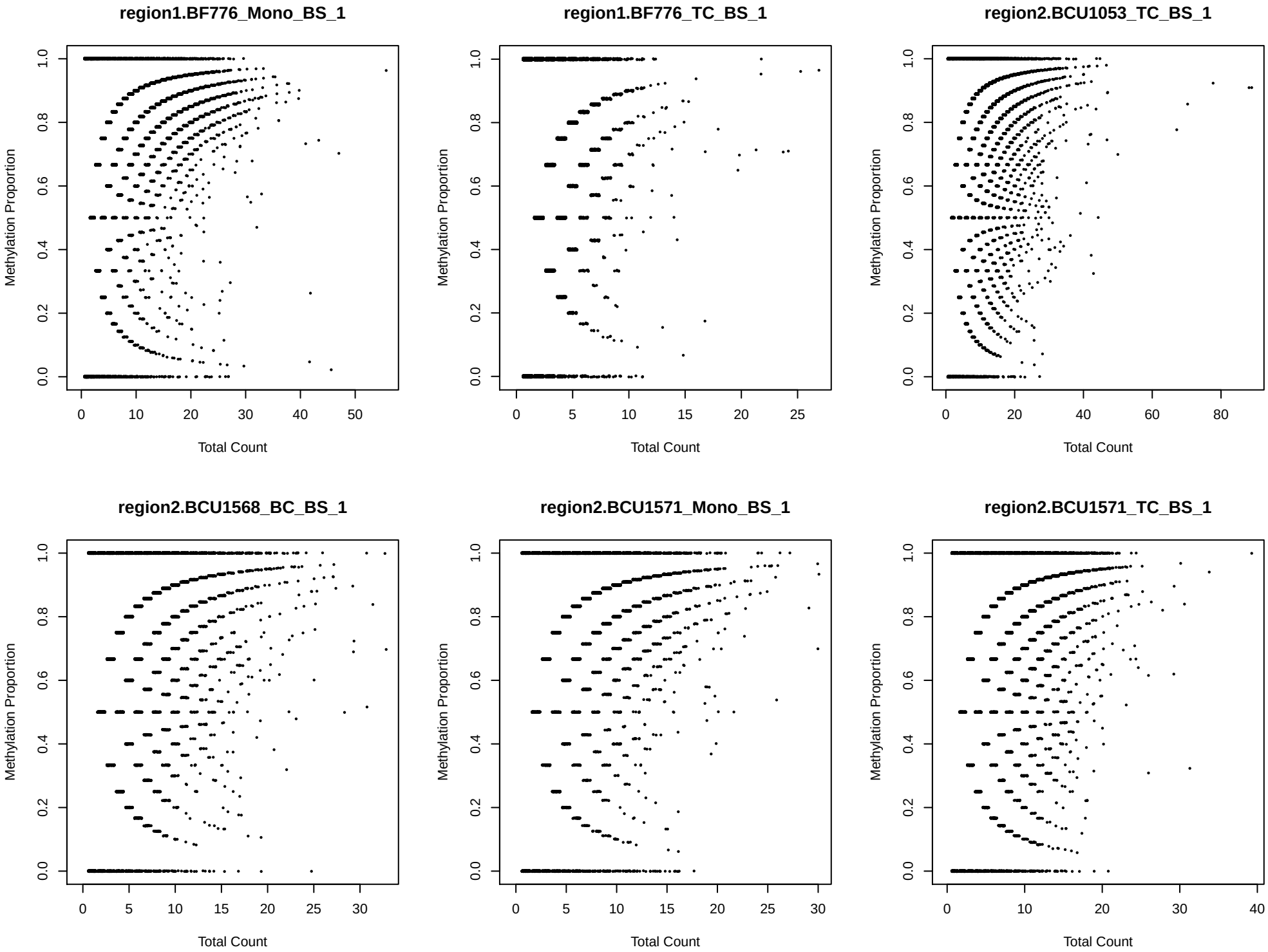

region1.BCU173_TC_BS_1
region1.BCU1799_Mono_BS_1
region1.BCU292_TC_BS_1
region1.BCU551_Mono_BS_1
region1.BCU551_TC_BS_1
region1.BCU582_TC_BS_1
Methylation Proportion
Total Count

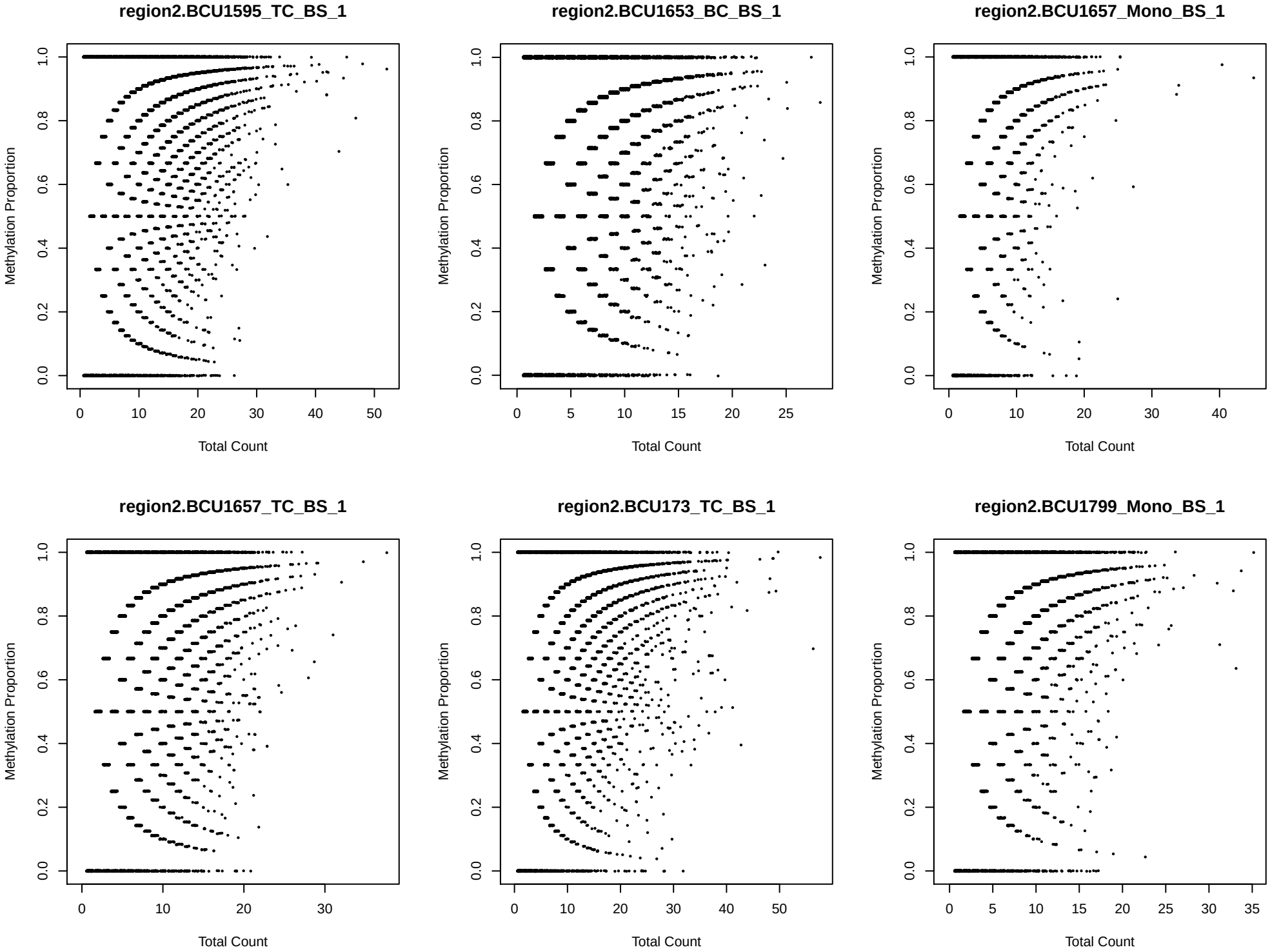

region1.BCU801_Mono_BS_1
region1.BCU801_TC_BS_1
region1.BCU899_TC_BS_1
region1.BF757_TC_BS_1
region1.BF758_Mono_BS_1
region1.BF761_TC_BS_1
Methylation Proportion
Total Count

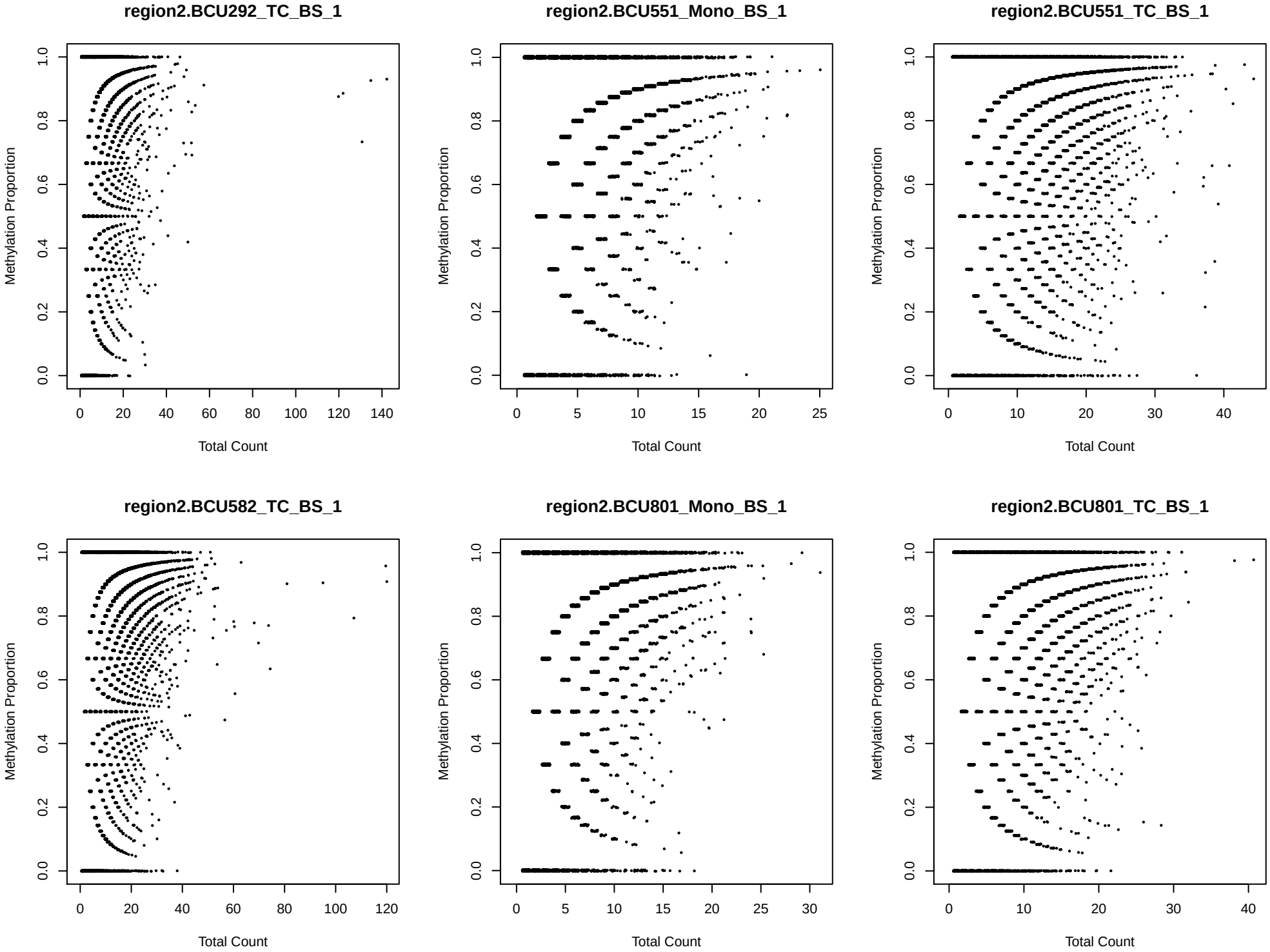

region1.BF764_BC_BS_1
region1.BF771_Mono_BS_1
region1.BF771_TC_BS_1
region1.BF772_BC_BS_2
region1.BF772_Mono_BS_1
region1.BF772_Mono_BS_2
Methylation Proportion
Total Count

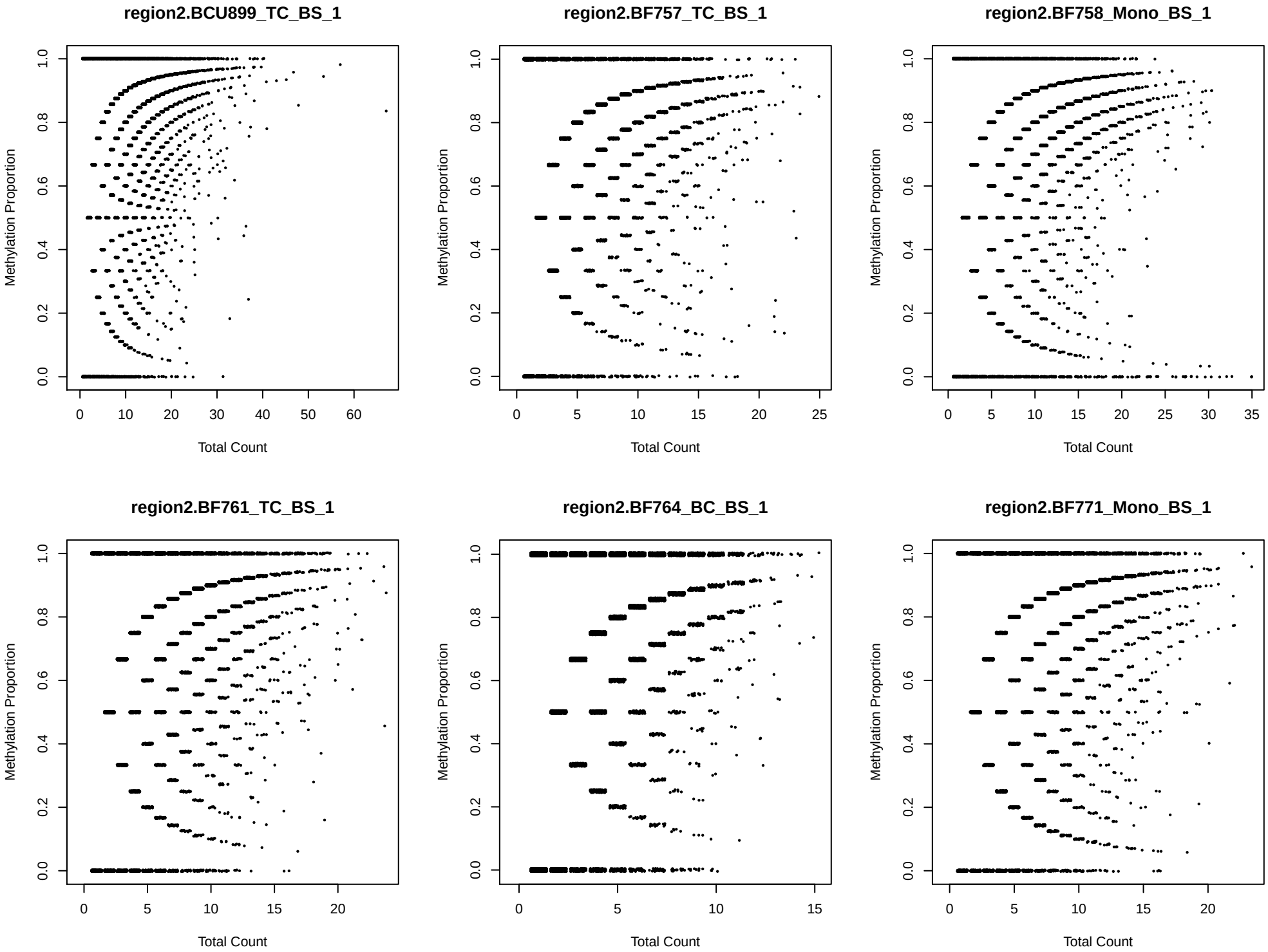

region1.BF772_TC_BS_1
region1.BF772_TC_BS_2
region1.BF773_BC_BS_1
region1.BF773_BC_BS_2
region1.BF773_Mono_BS_1
region1.BF773_Mono_BS_2
Methylation Proportion
Total Count

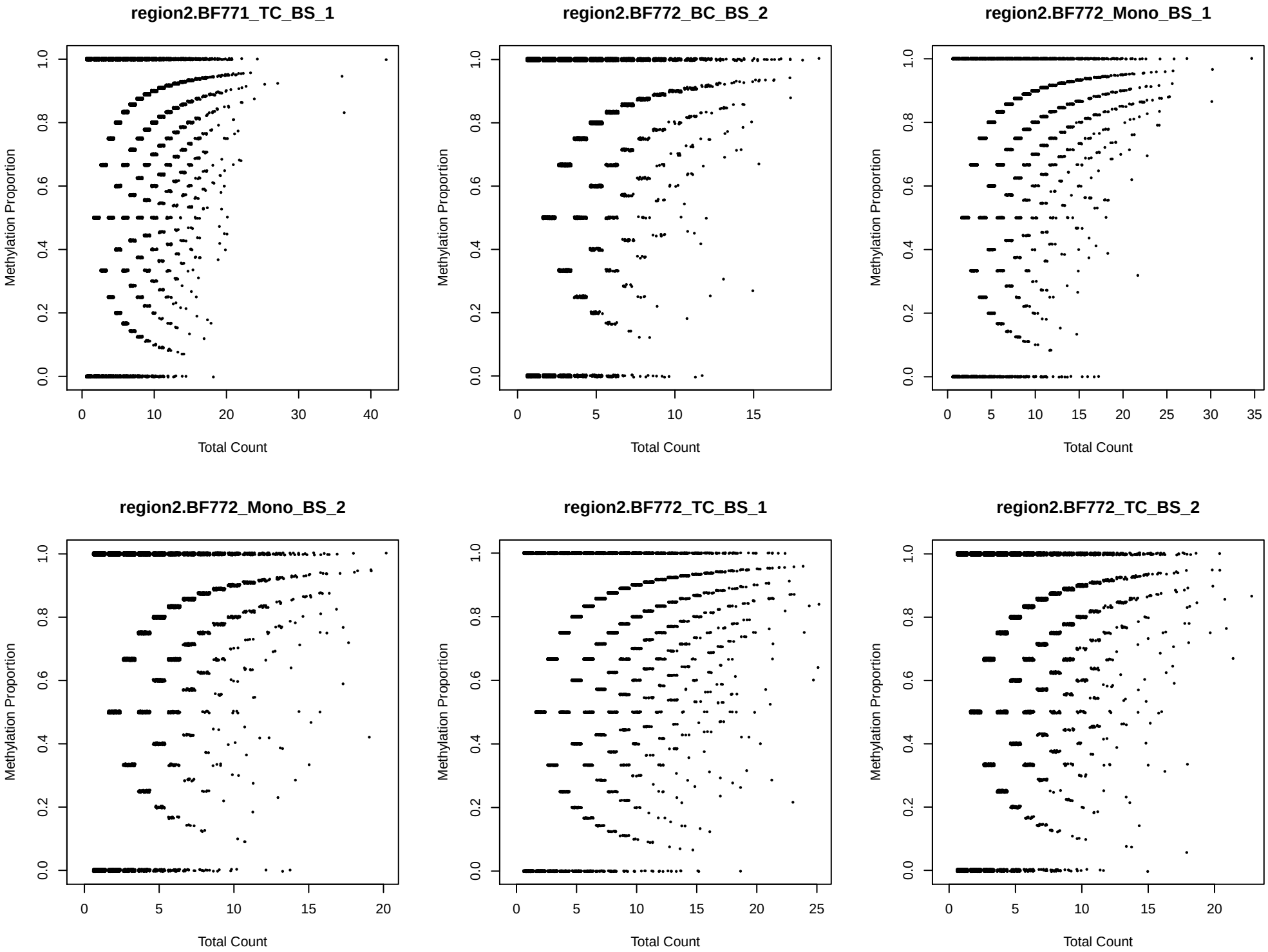

region1.BF773_TC_BS_1
region1.BF773_TC_BS_2
region1.BF775_BC_BS_1
region1.BF775_Mono_BS_1
region1.BF775_TC_BS_1
region1.BF776_BC_BS_1
Methylation Proportion
Total Count

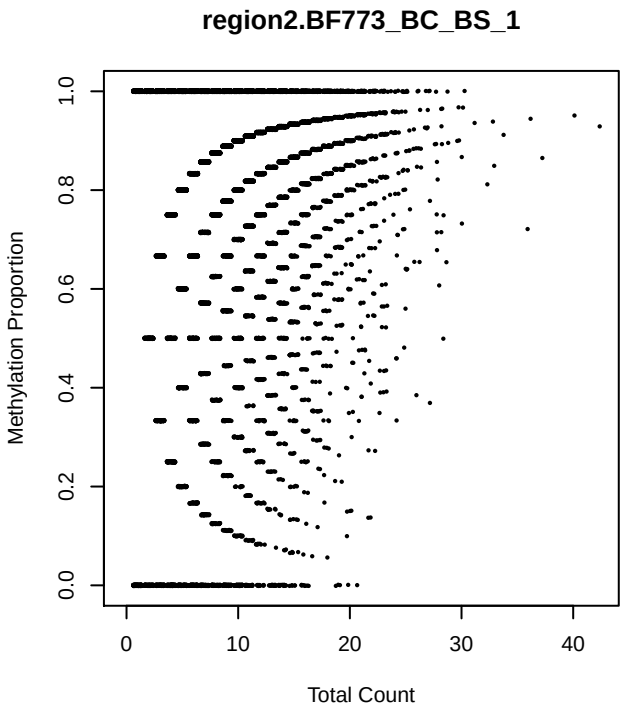

region1.BF776_Mono_BS_1
region1.BF776_TC_BS_1
region2.BCU1053_TC_BS_1
region2.BCU1568_BC_BS_1
region2.BCU1571_Mono_BS_1
region2.BCU1571_TC_BS_1
Methylation Proportion
Total Count

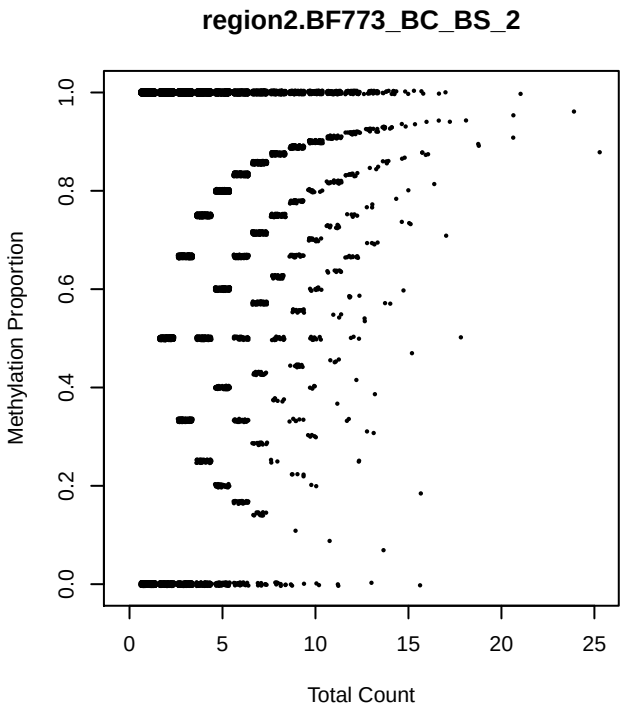

region2.BCU1595_TC_BS_1
region2.BCU1653_BC_BS_1
region2.BCU1657_Mono_BS_1
region2.BCU1657_TC_BS_1
region2.BCU173_TC_BS_1
region2.BCU1799_Mono_BS_1
Methylation Proportion
Total Count

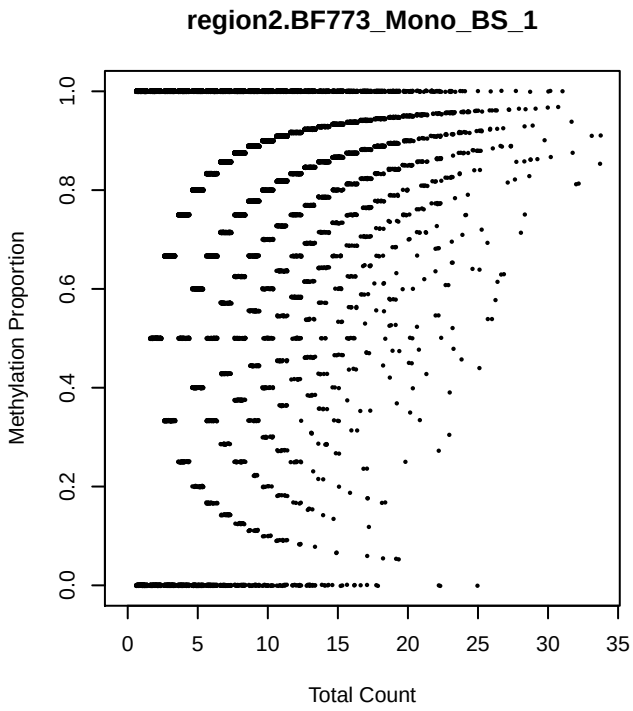

region2.BCU292_TC_BS_1
region2.BCU551_Mono_BS_1
region2.BCU551_TC_BS_1
region2.BCU582_TC_BS_1
region2.BCU801_Mono_BS_1
region2.BCU801_TC_BS_1
Methylation Proportion
Total Count

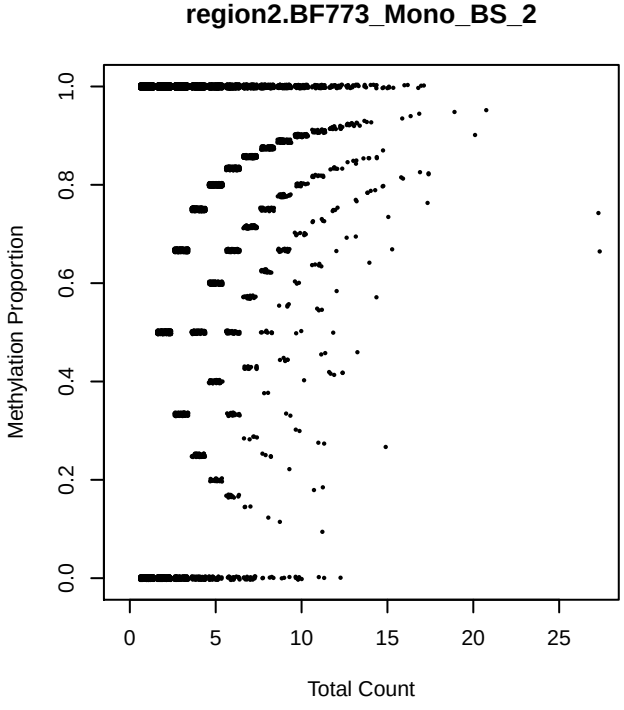

region2.BCU899_TC_BS_1
region2.BF757_TC_BS_1
region2.BF758_Mono_BS_1
region2.BF761_TC_BS_1
region2.BF764_BC_BS_1
region2.BF771_Mono_BS_1
Methylation Proportion
Total Count

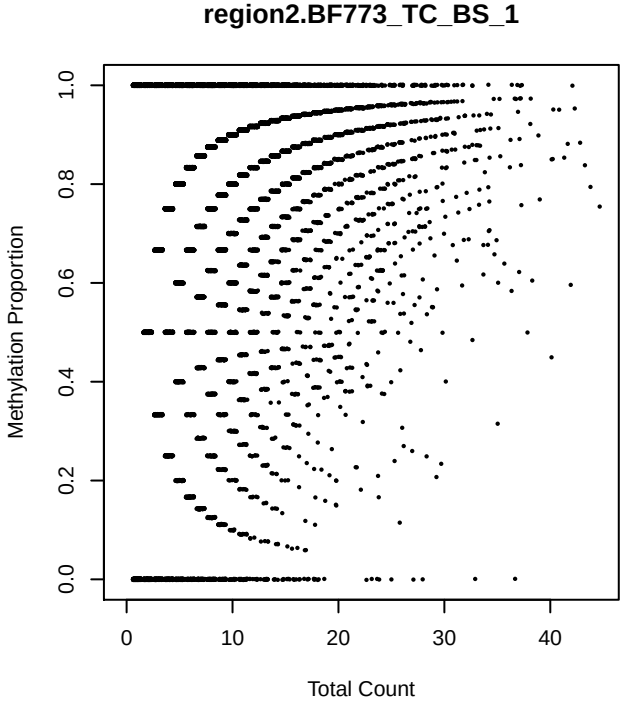

region2.BF771_TC_BS_1
region2.BF772_BC_BS_2
region2.BF772_Mono_BS_1
region2.BF772_Mono_BS_2
region2.BF772_TC_BS_1
region2.BF772_TC_BS_2
Methylation Proportion
Total Count

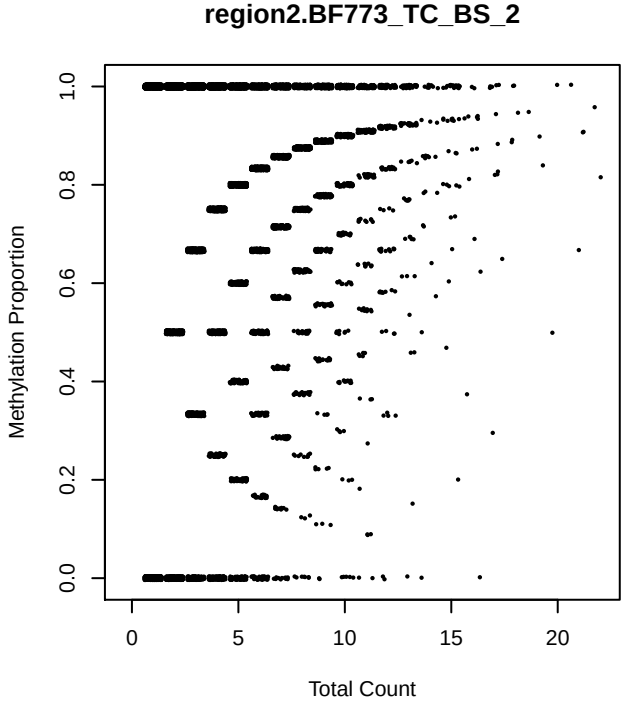

region2.BF773_BC_BS_1
Methylation Proportion
Total Count

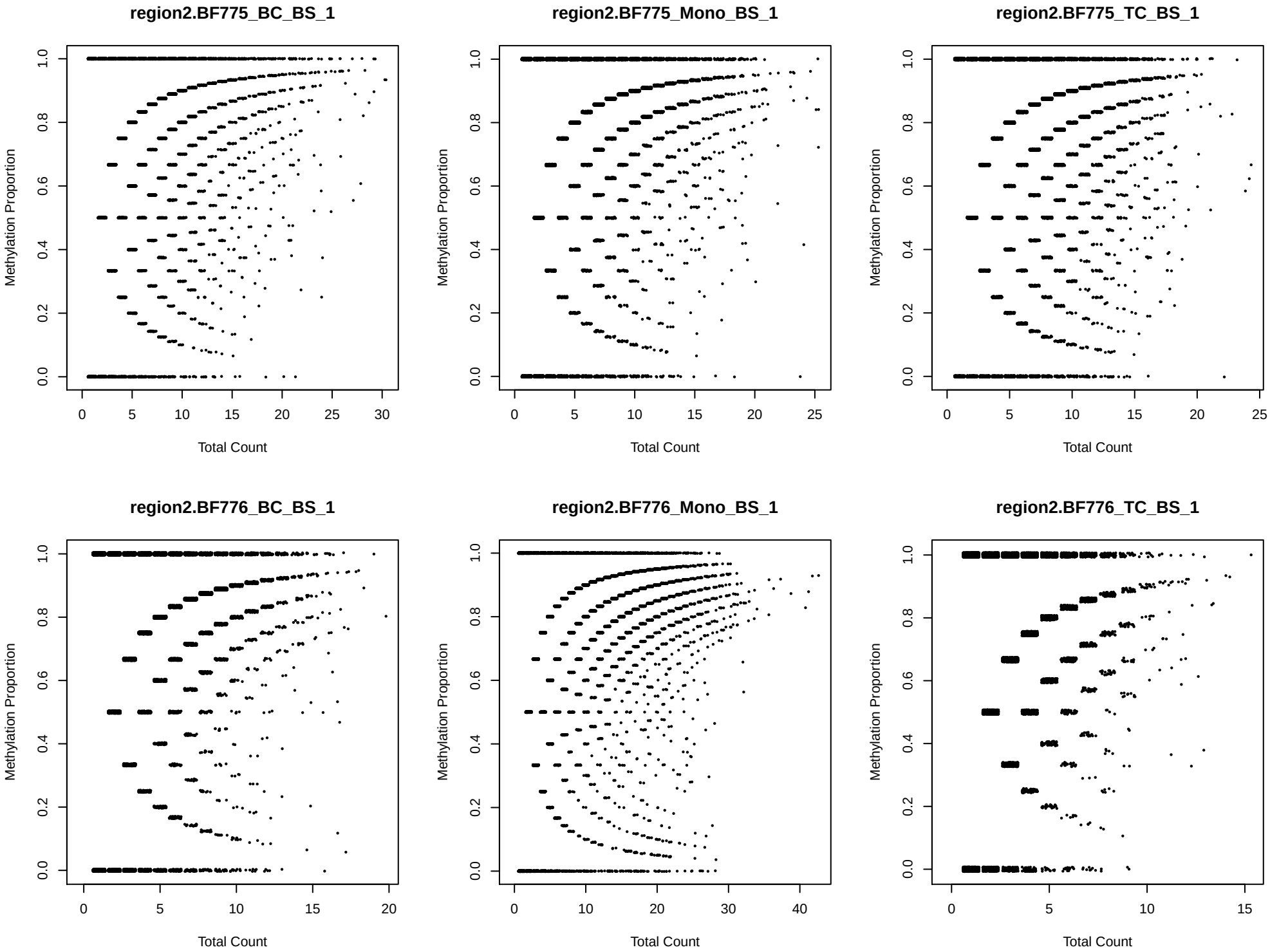

region2.BF773_BC_BS_2
Methylation Proportion
Total Count

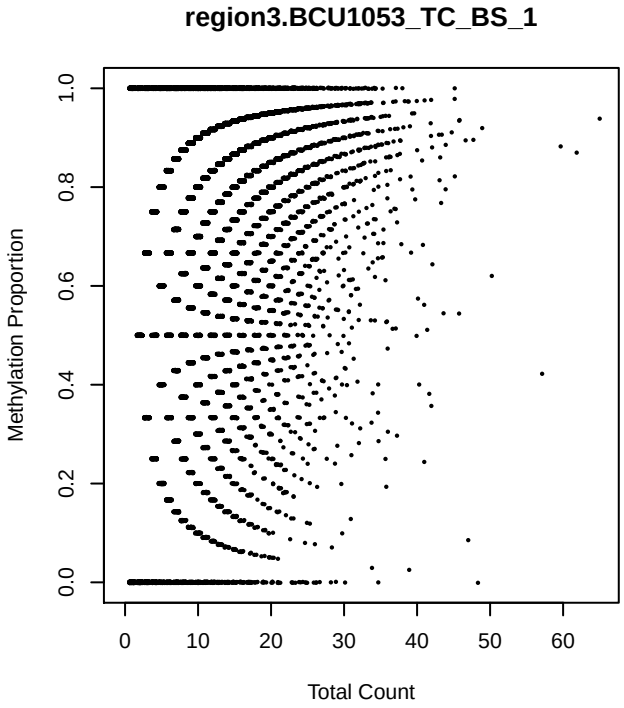

region2.BF773_Mono_BS_1
Methylation Proportion
Total Count

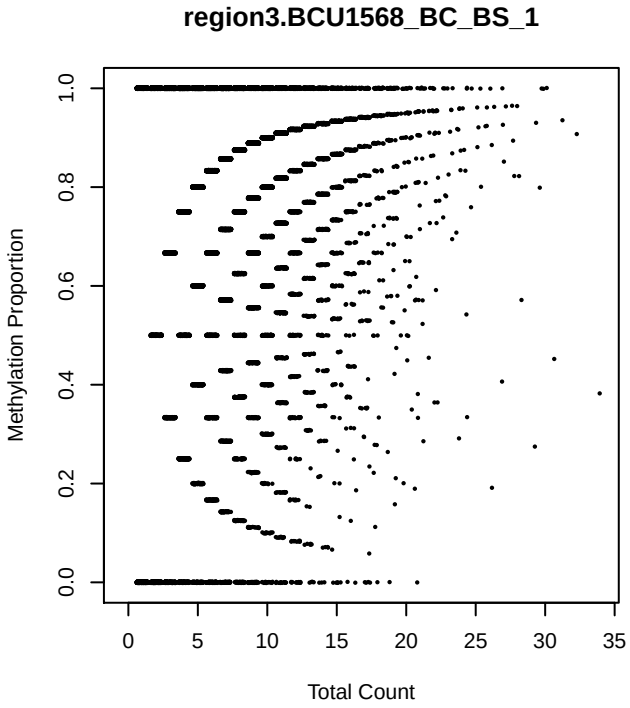

region2.BF773_Mono_BS_2
Methylation Proportion
Total Count

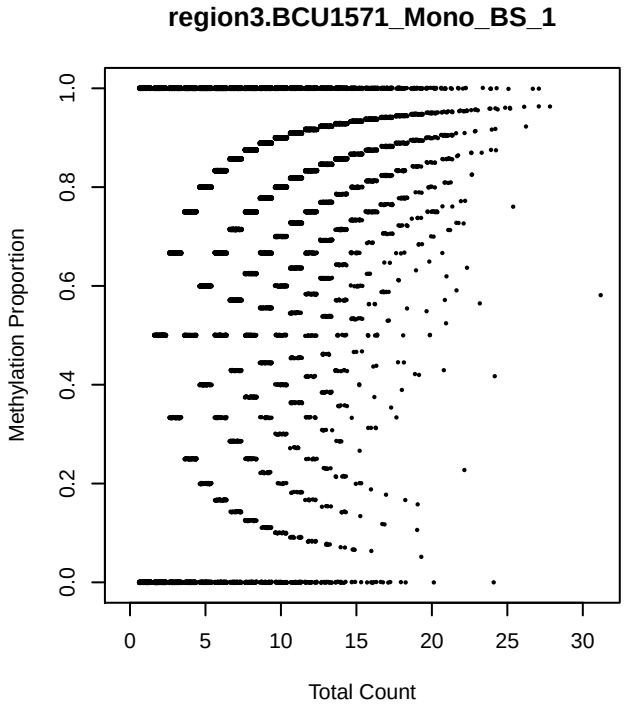

region2.BF773_TC_BS_1
Methylation Proportion
Total Count

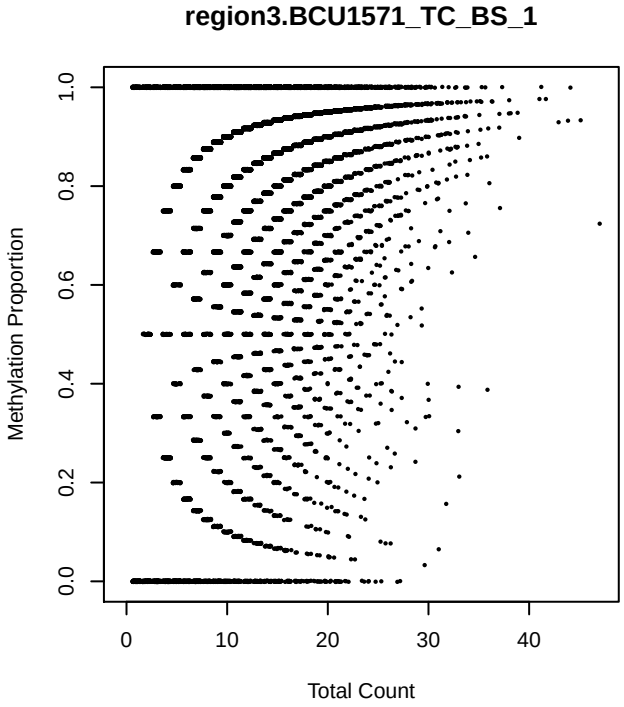

region2.BF773_TC_BS_2
Methylation Proportion
Total Count

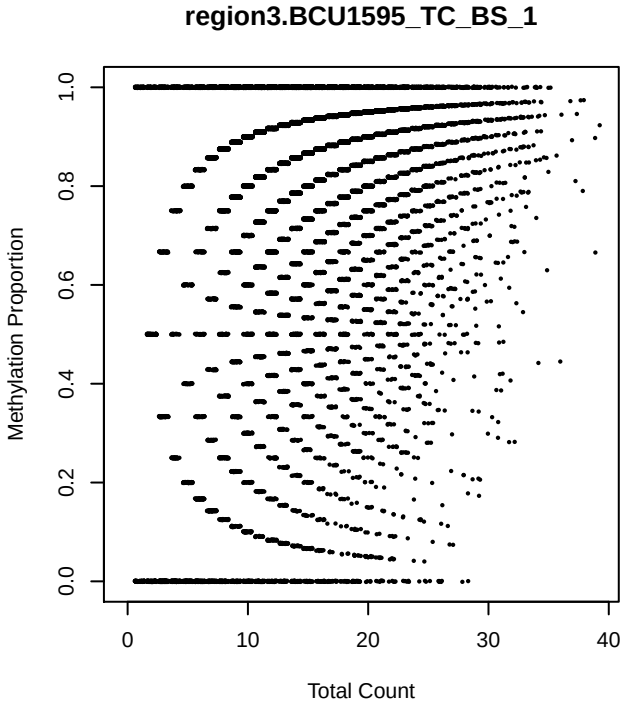

region2.BF775_BC_BS_1
region2.BF775_Mono_BS_1
region2.BF775_TC_BS_1
region2.BF776_BC_BS_1
region2.BF776_Mono_BS_1
region2.BF776_TC_BS_1
Methylation Proportion
Total Count

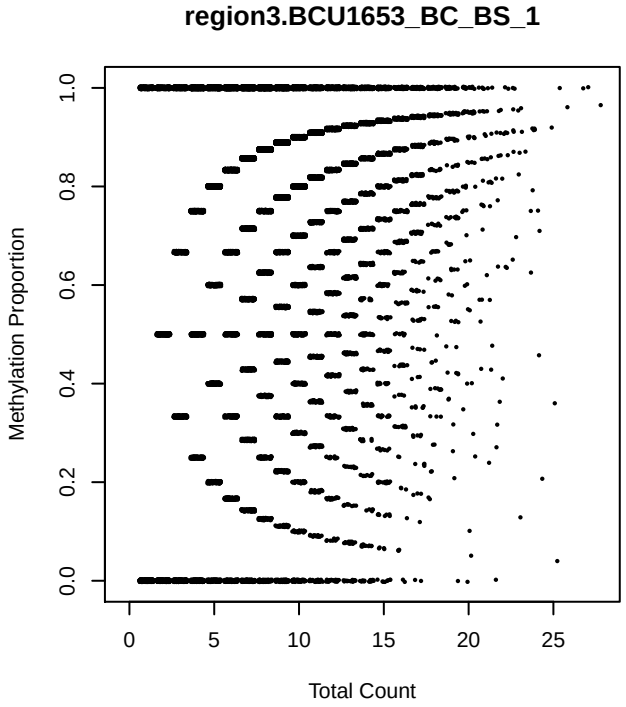

region3.BCU1053_TC_BS_1
Methylation Proportion
Total Count

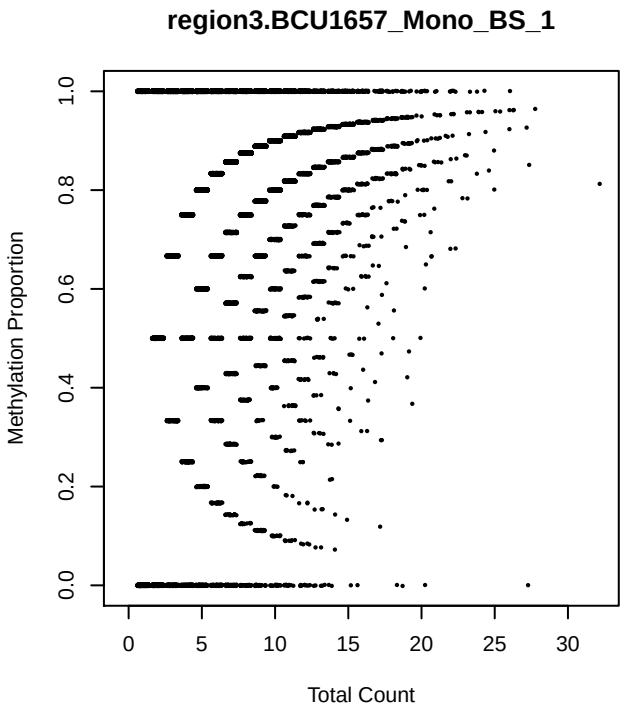

region3.BCU1568_BC_BS_1
Methylation Proportion
Total Count

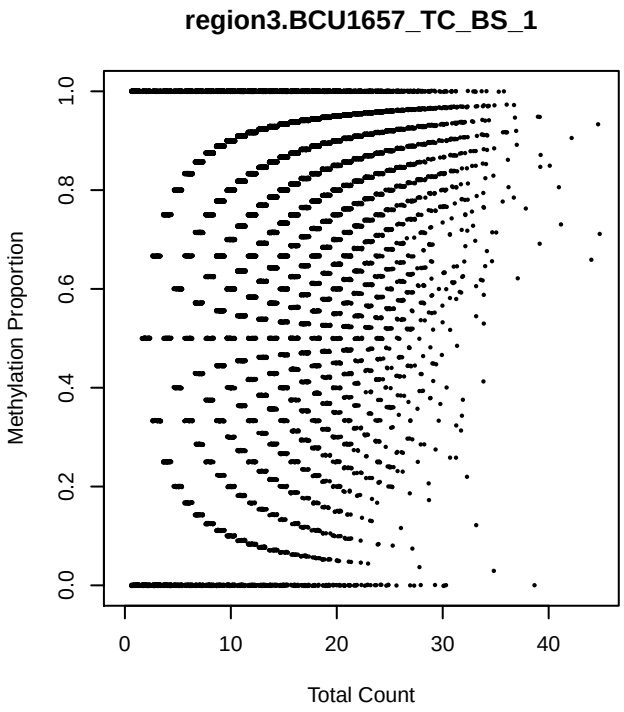

region3.BCU1571_Mono_BS_1
Methylation Proportion
Total Count

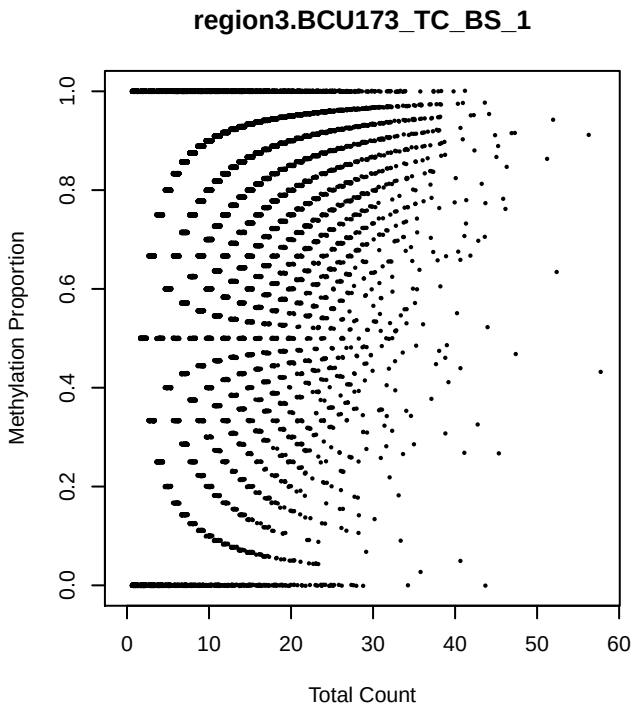

region3.BCU1571_TC_BS_1
Methylation Proportion
Total Count

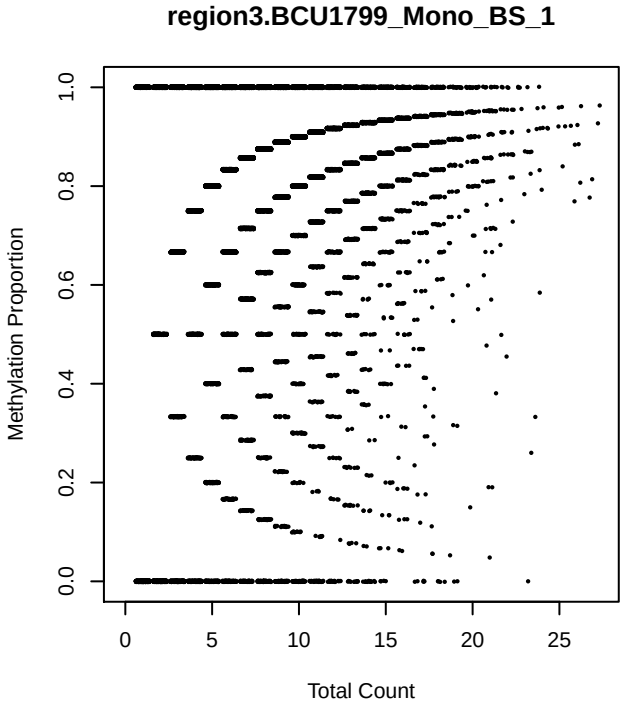

region3.BCU1595_TC_BS_1
Methylation Proportion
Total Count

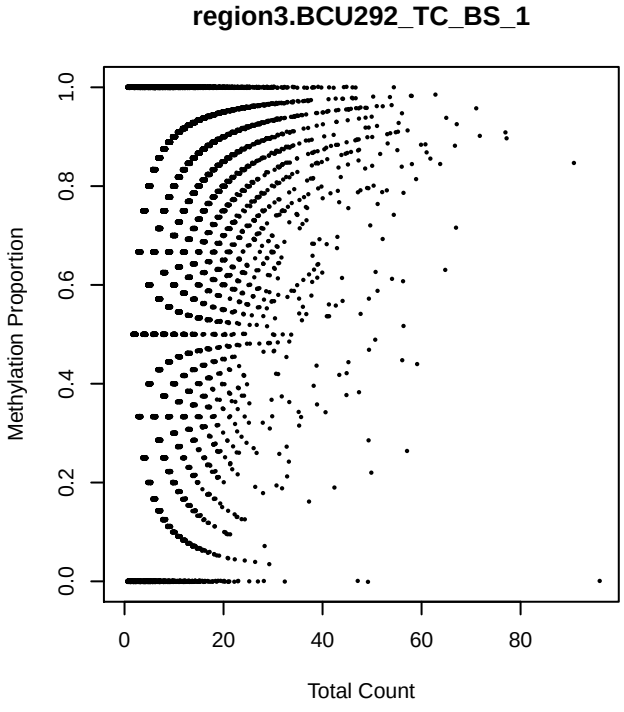

region3.BCU1653_BC_BS_1
Methylation Proportion
Total Count

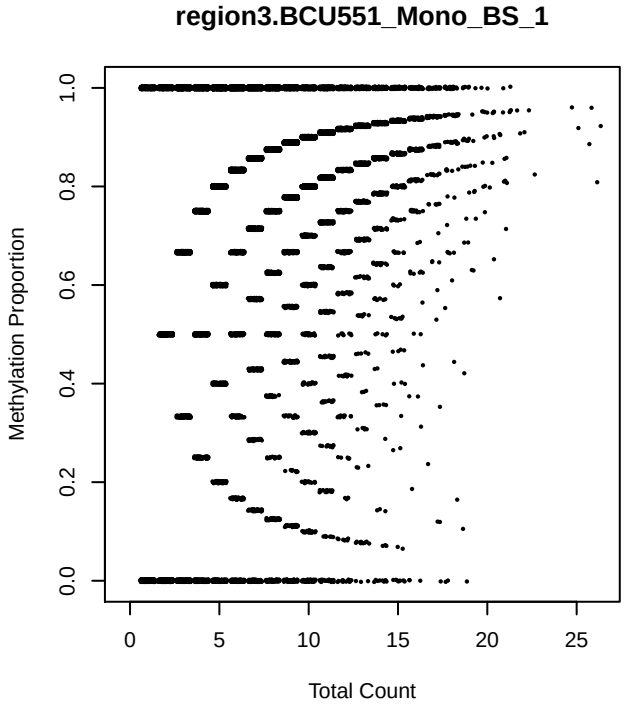

region3.BCU1657_Mono_BS_1
Methylation Proportion
Total Count

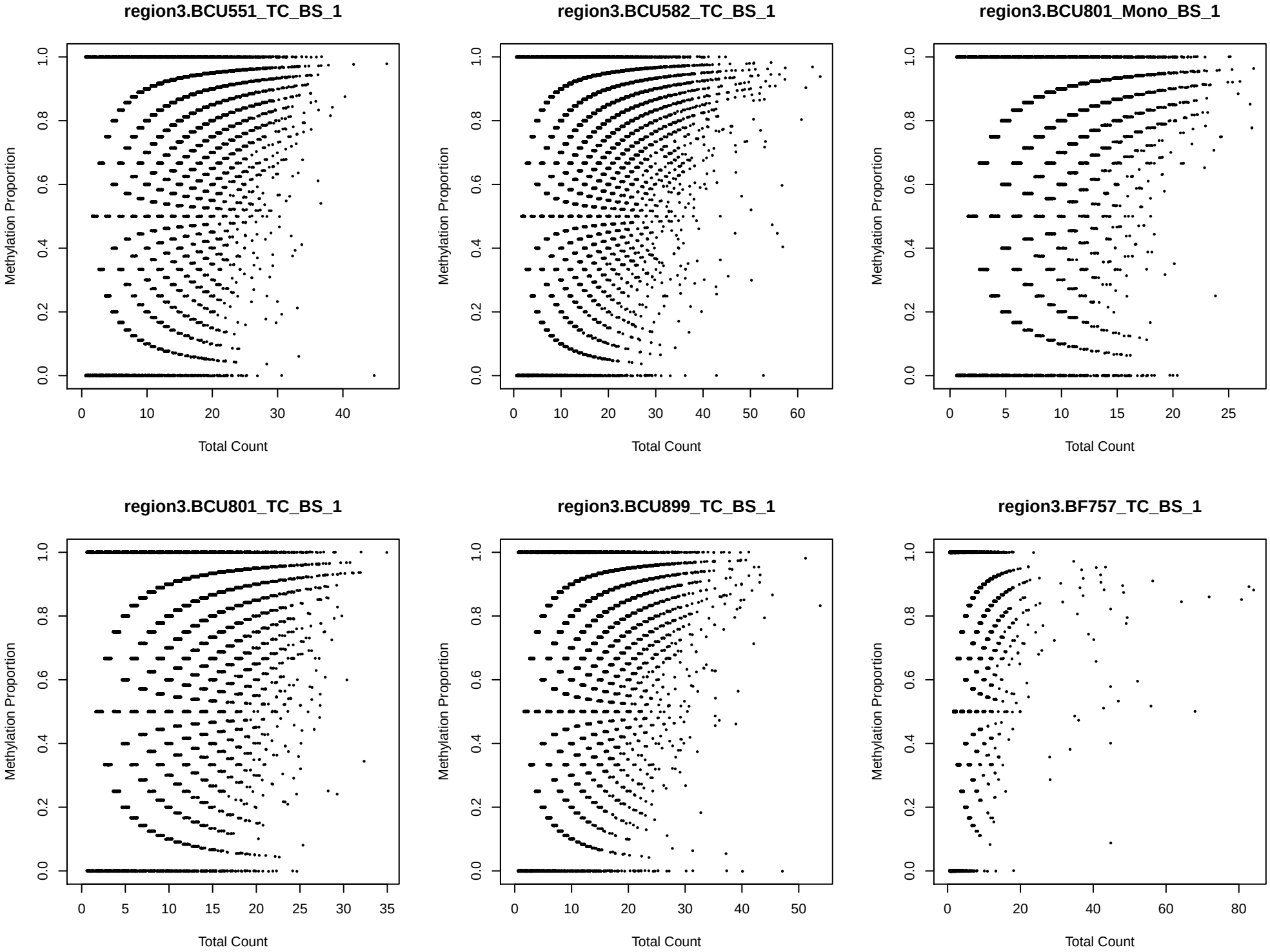

region3.BCU1657_TC_BS_1
Methylation Proportion
Total Count

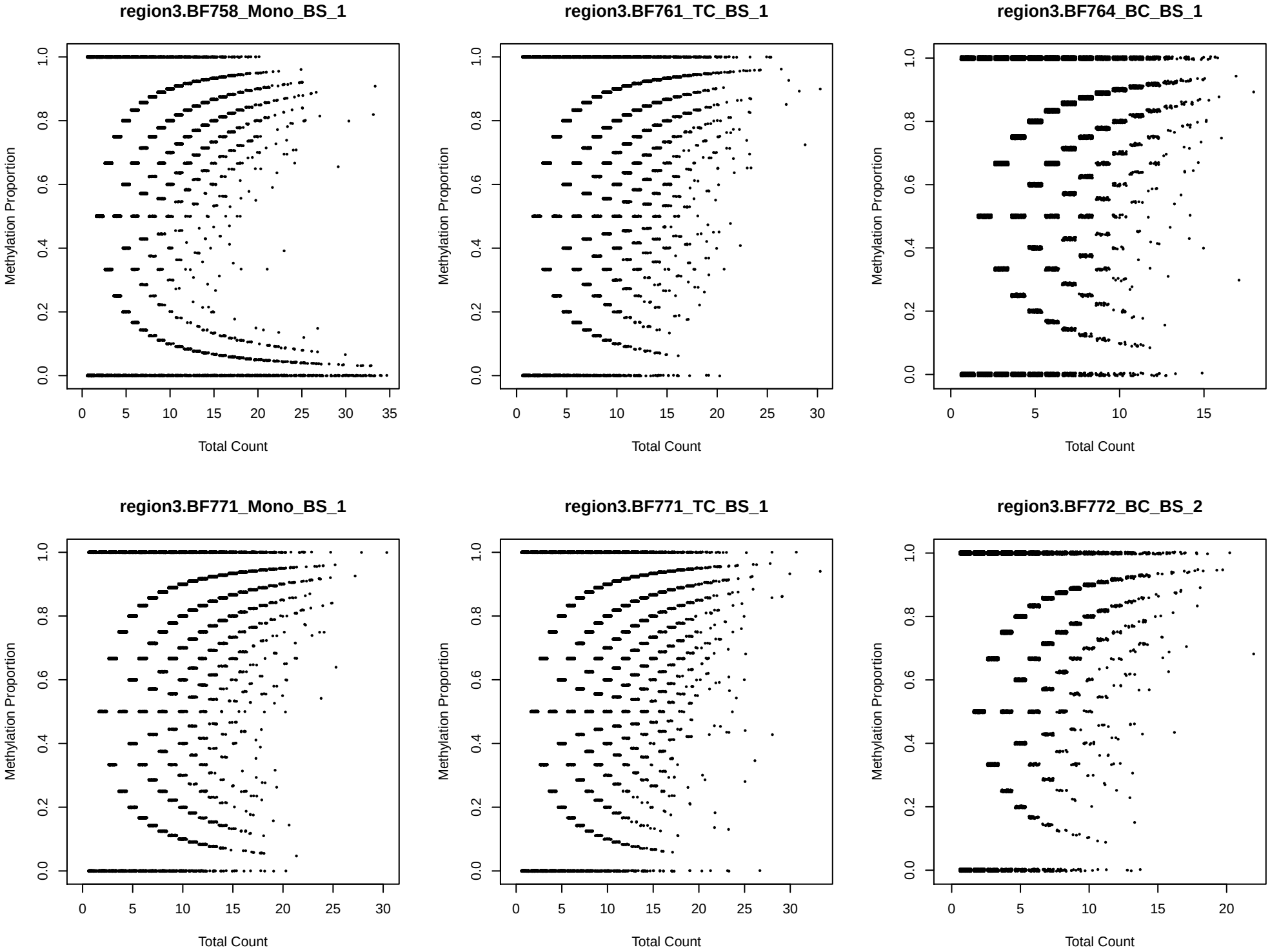

region3.BCU173_TC_BS_1
Methylation Proportion
Total Count

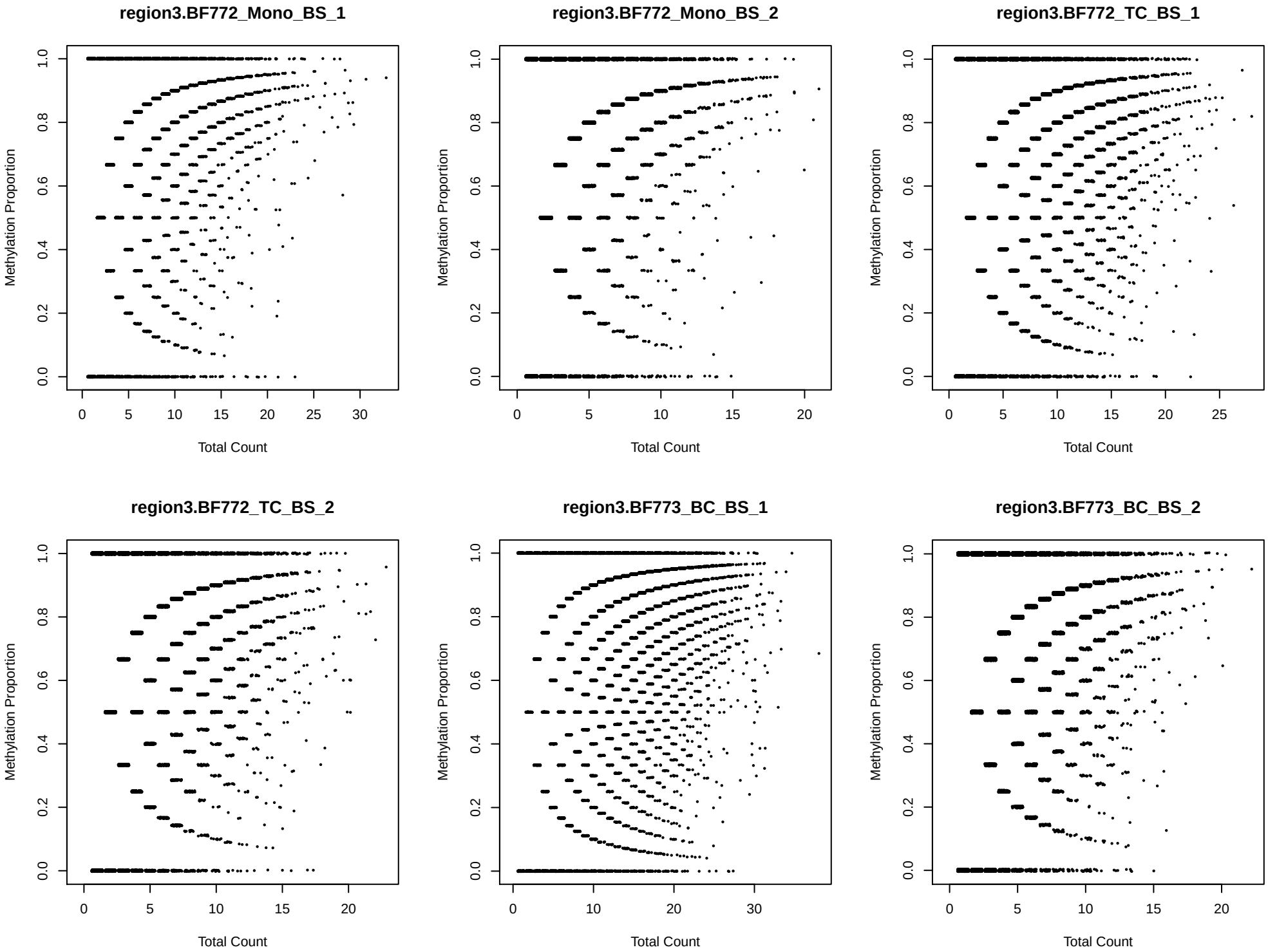

region3.BCU1799_Mono_BS_1
Methylation Proportion
Total Count

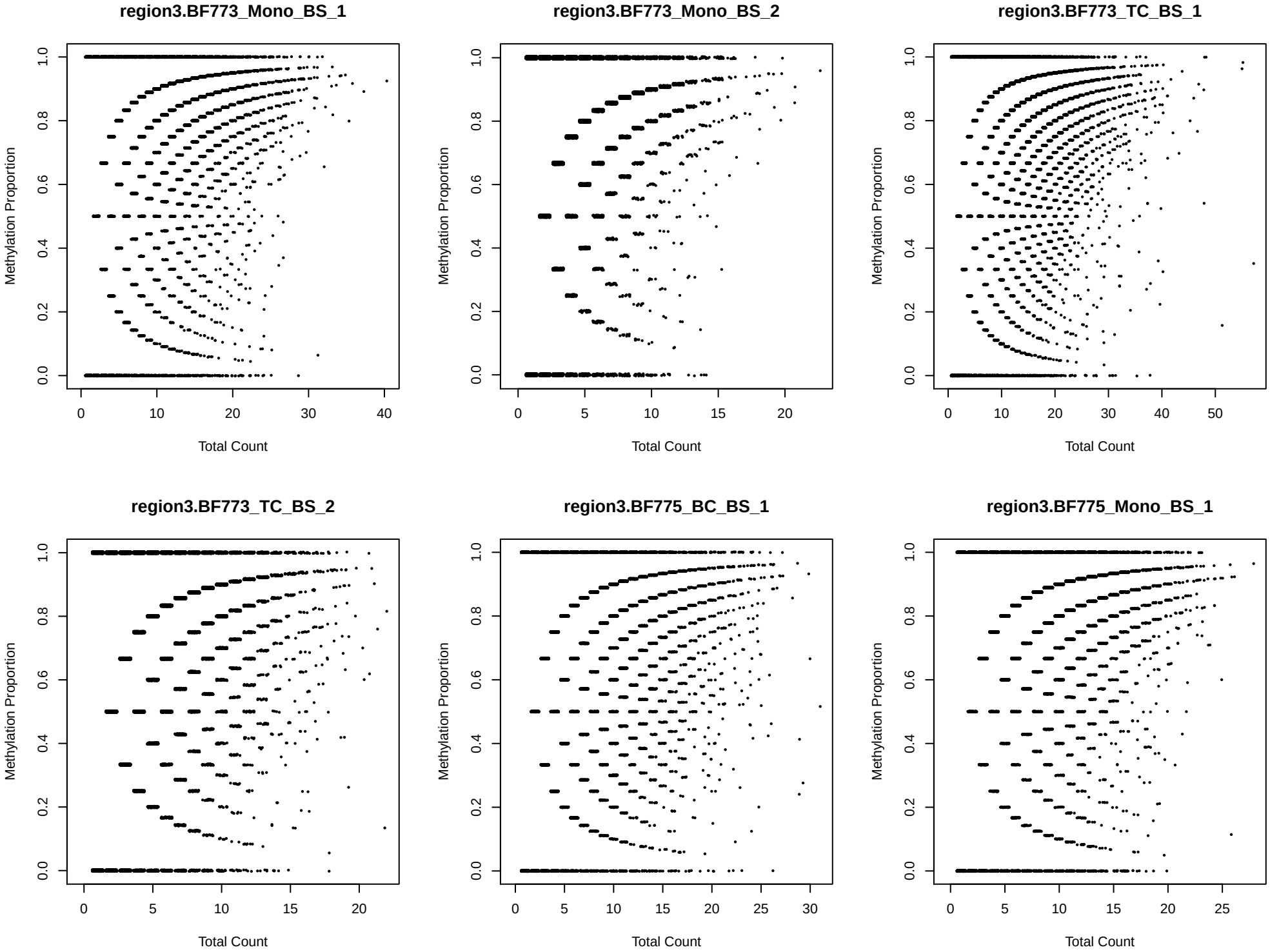

region3.BCU292_TC_BS_1
Methylation Proportion
Total Count

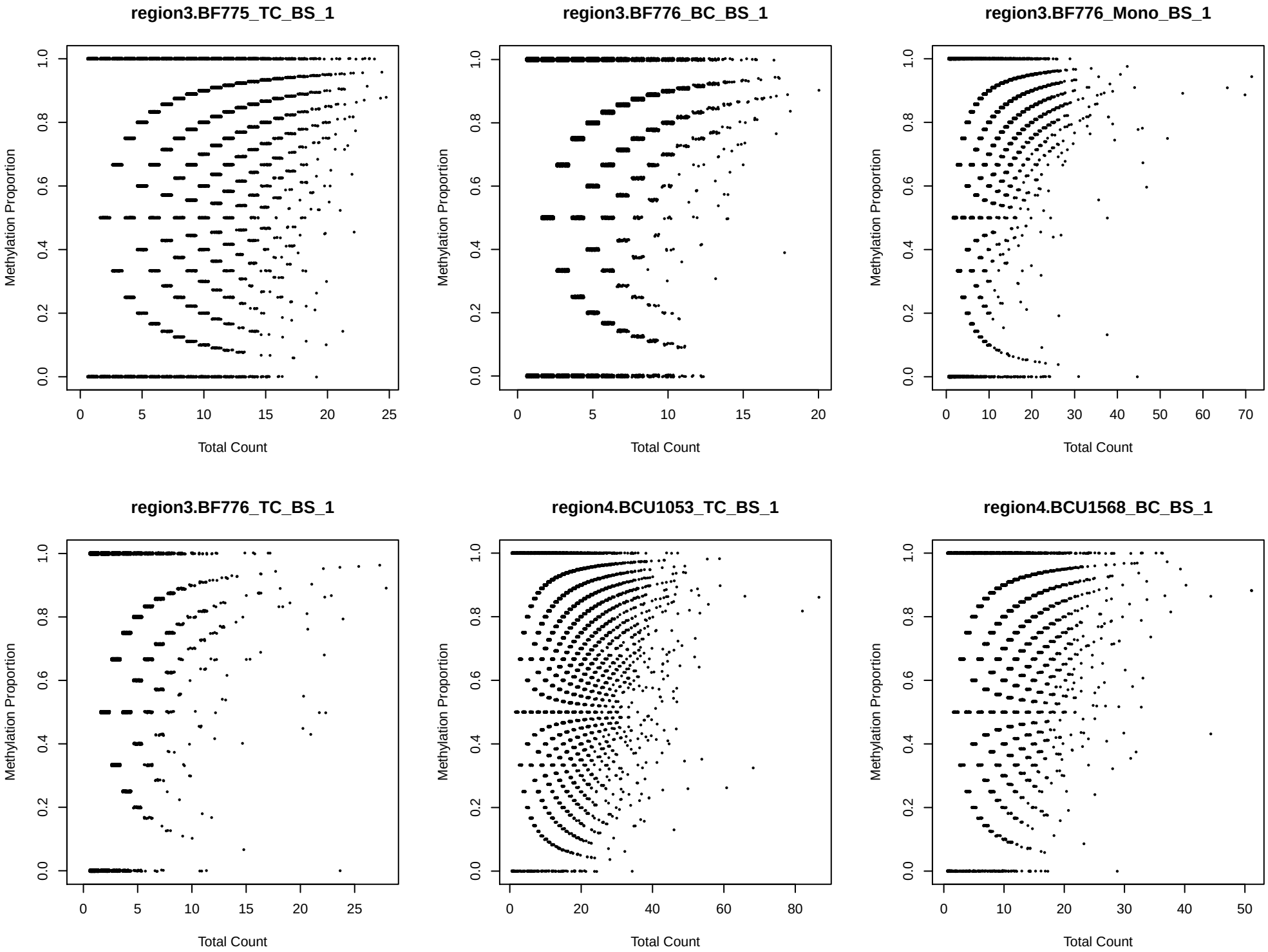

region3.BCU551_Mono_BS_1
Methylation Proportion
Total Count

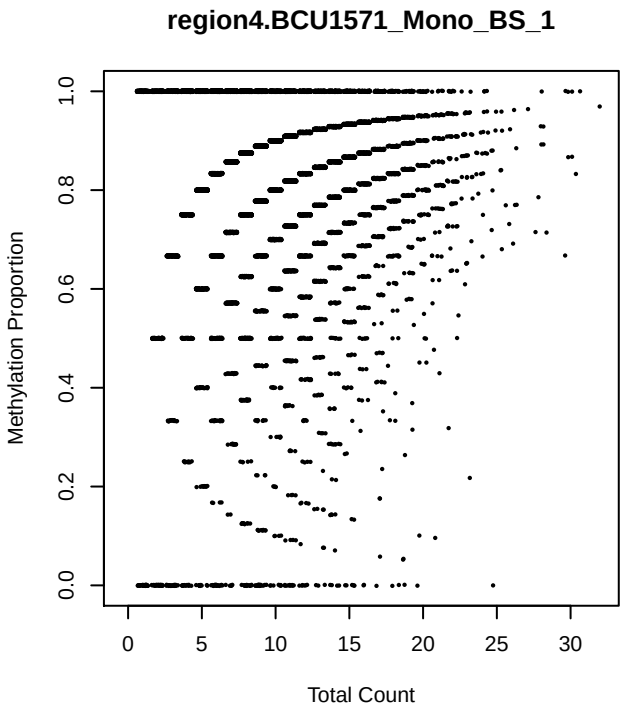

region3.BCU551_TC_BS_1
region3.BCU582_TC_BS_1
region3.BCU801_Mono_BS_1
region3.BCU801_TC_BS_1
region3.BCU899_TC_BS_1
region3.BF757_TC_BS_1
Methylation Proportion
Total Count

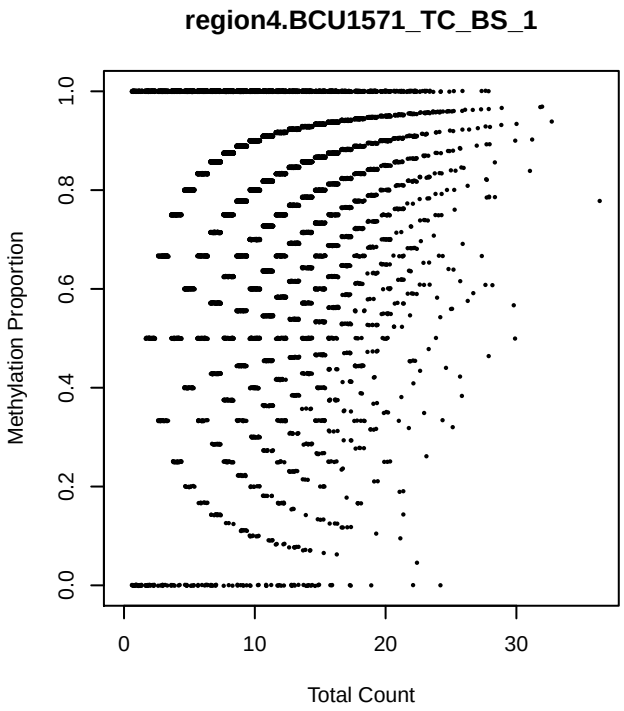

region3.BF758_Mono_BS_1
region3.BF761_TC_BS_1
region3.BF764_BC_BS_1
region3.BF771_Mono_BS_1
region3.BF771_TC_BS_1
region3.BF772_BC_BS_2
Methylation Proportion
Total Count

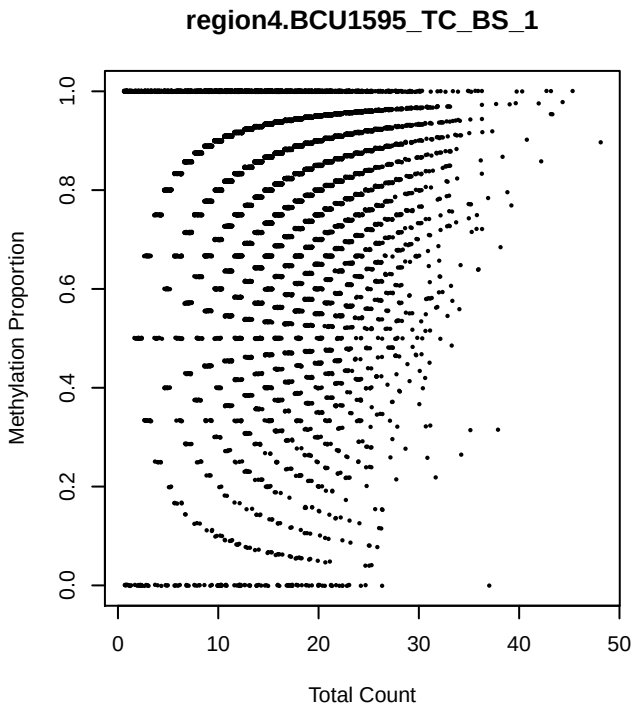

region3.BF772_Mono_BS_1
region3.BF772_Mono_BS_2
region3.BF772_TC_BS_1
region3.BF772_TC_BS_2
region3.BF773_BC_BS_1
region3.BF773_BC_BS_2
Methylation Proportion
Total Count

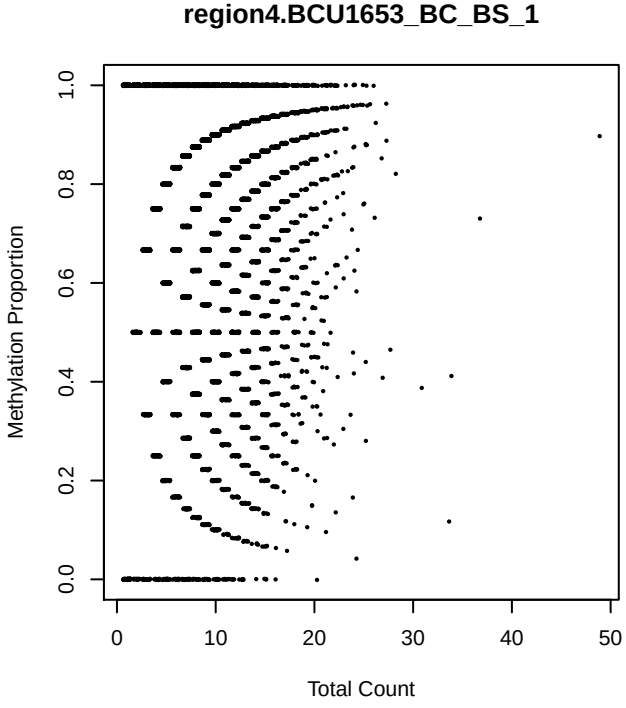

region3.BF773_Mono_BS_1
region3.BF773_Mono_BS_2
region3.BF773_TC_BS_1
region3.BF773_TC_BS_2
region3.BF775_BC_BS_1
region3.BF775_Mono_BS_1
Methylation Proportion
Total Count

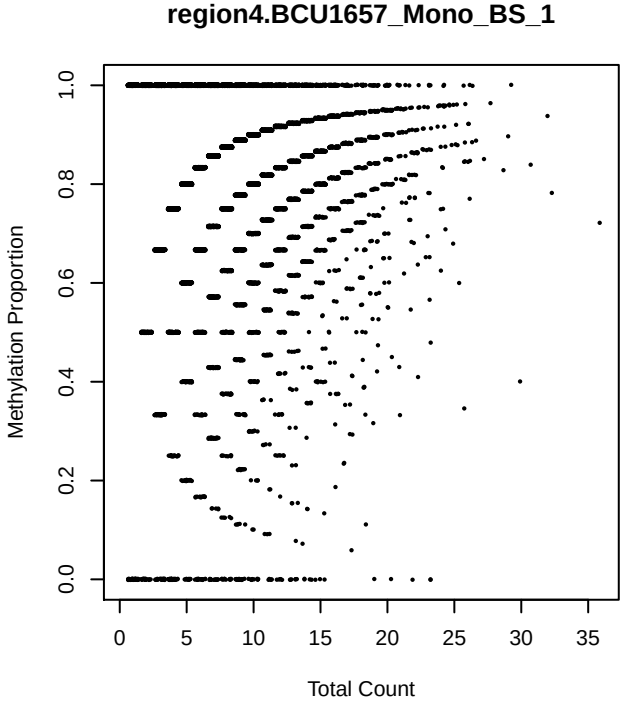

region3.BF775_TC_BS_1
region3.BF776_BC_BS_1
region3.BF776_Mono_BS_1
region3.BF776_TC_BS_1
region4.BCU1053_TC_BS_1
region4.BCU1568_BC_BS_1
Methylation Proportion
Total Count

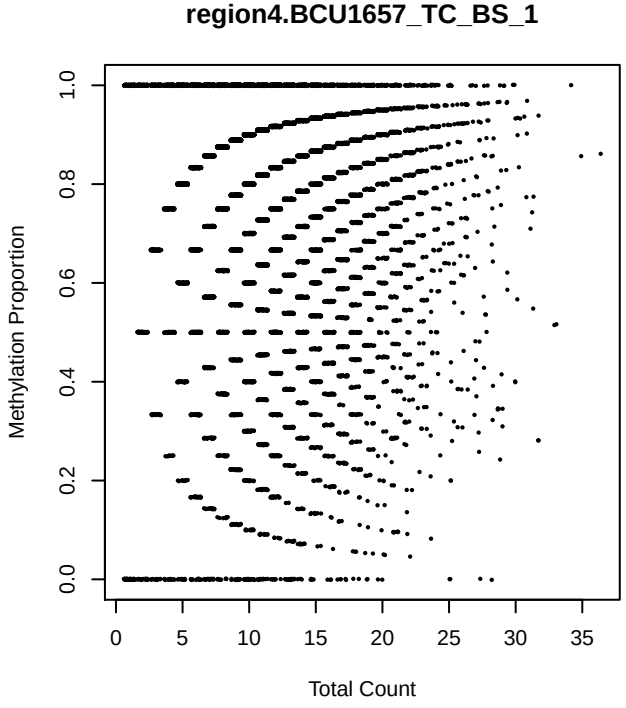

region4.BCU1571_Mono_BS_1
Methylation Proportion
Total Count

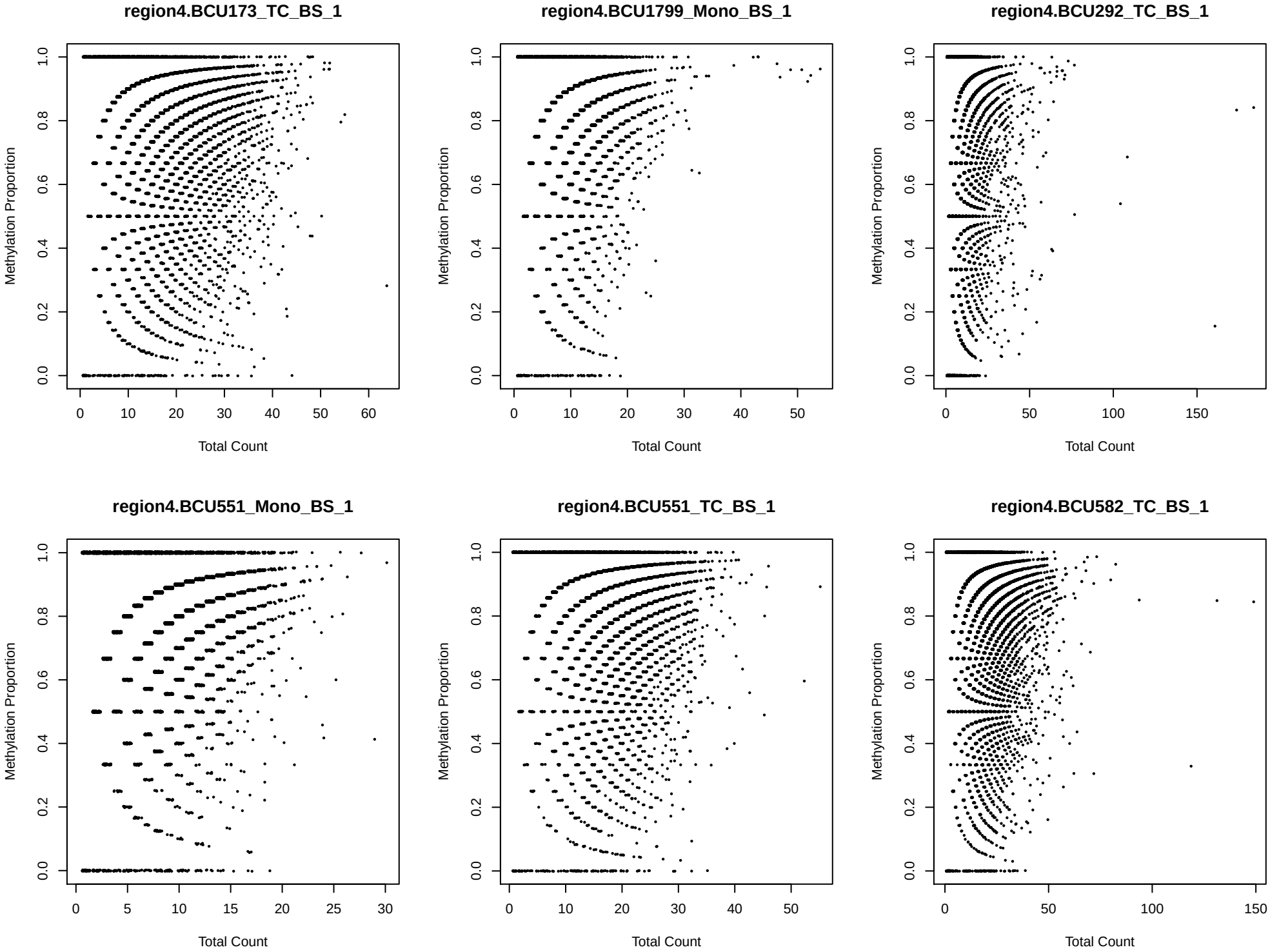

region4.BCU1571_TC_BS_1
Methylation Proportion
Total Count

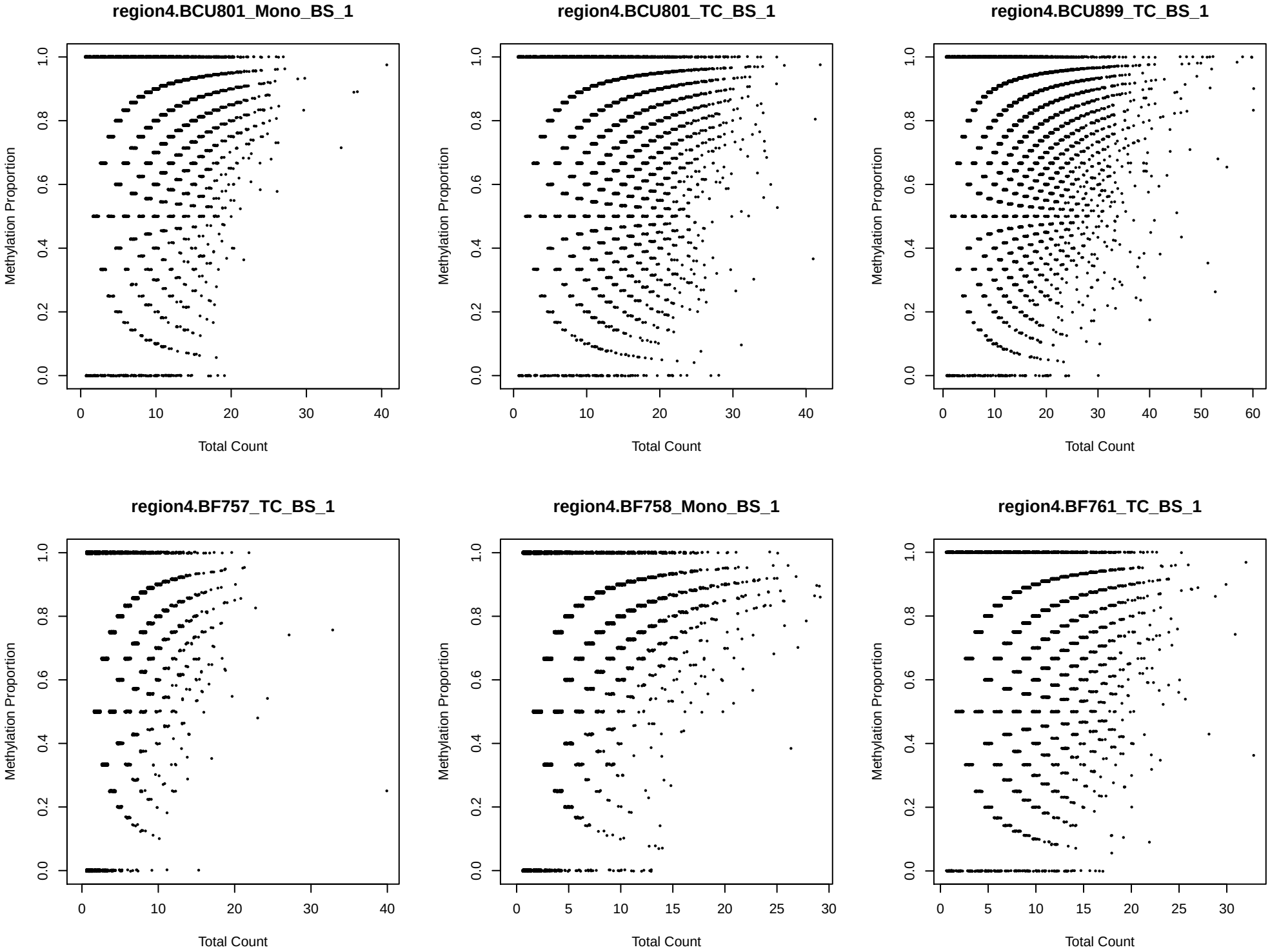

region4.BCU1595_TC_BS_1
Methylation Proportion
Total Count

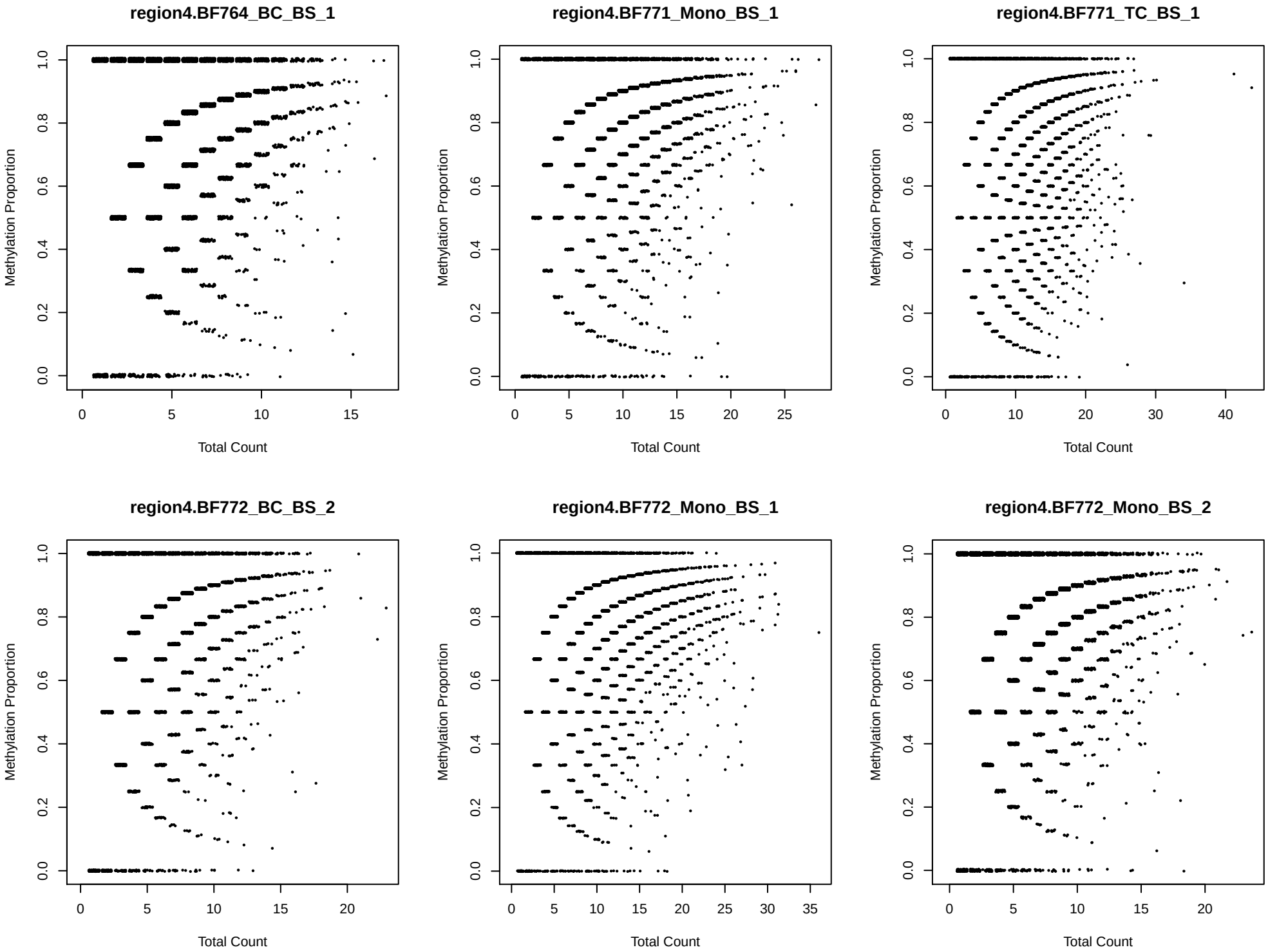

region4.BCU1653_BC_BS_1
Methylation Proportion
Total Count

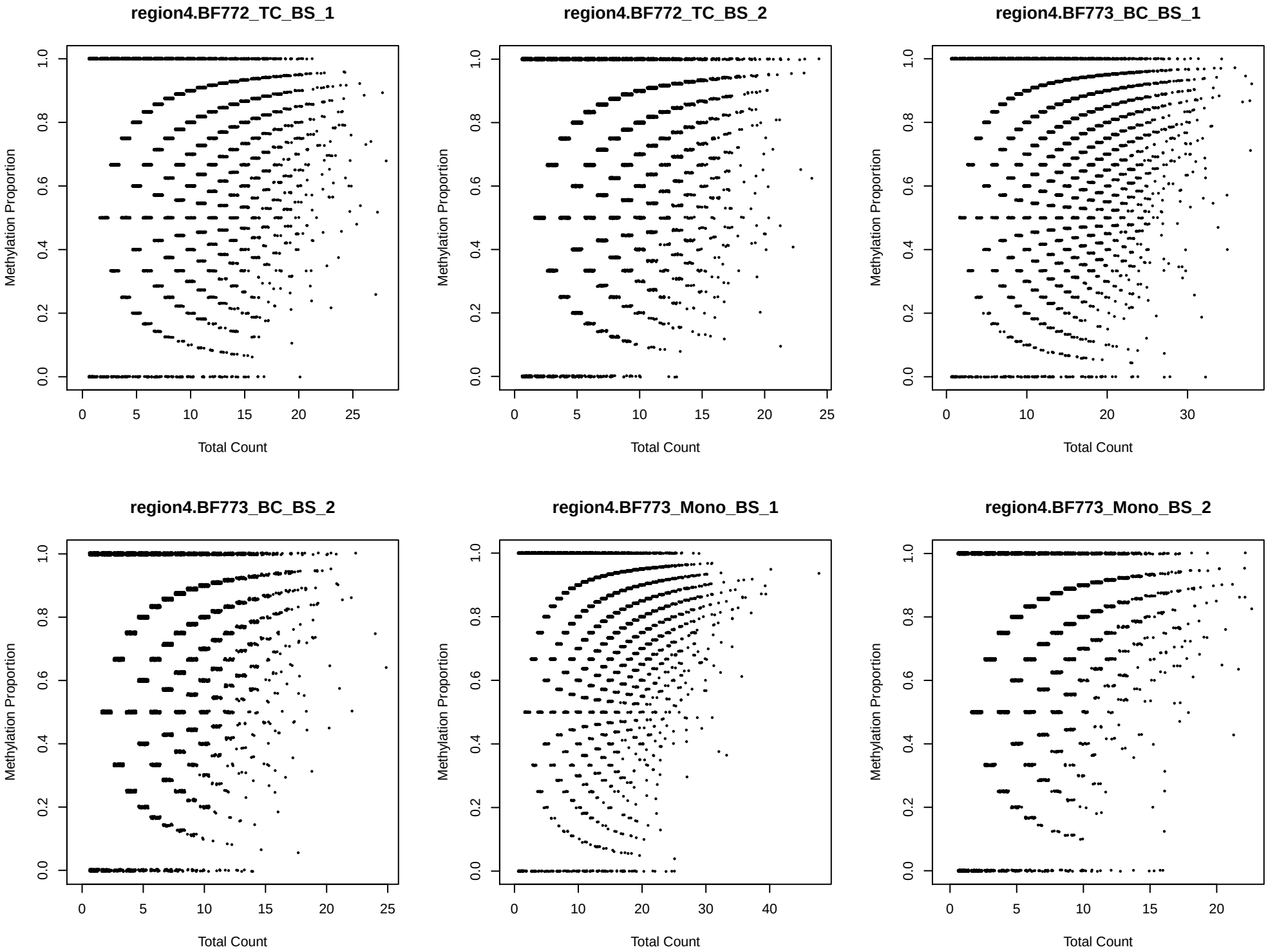

region4.BCU1657_Mono_BS_1
Methylation Proportion
Total Count

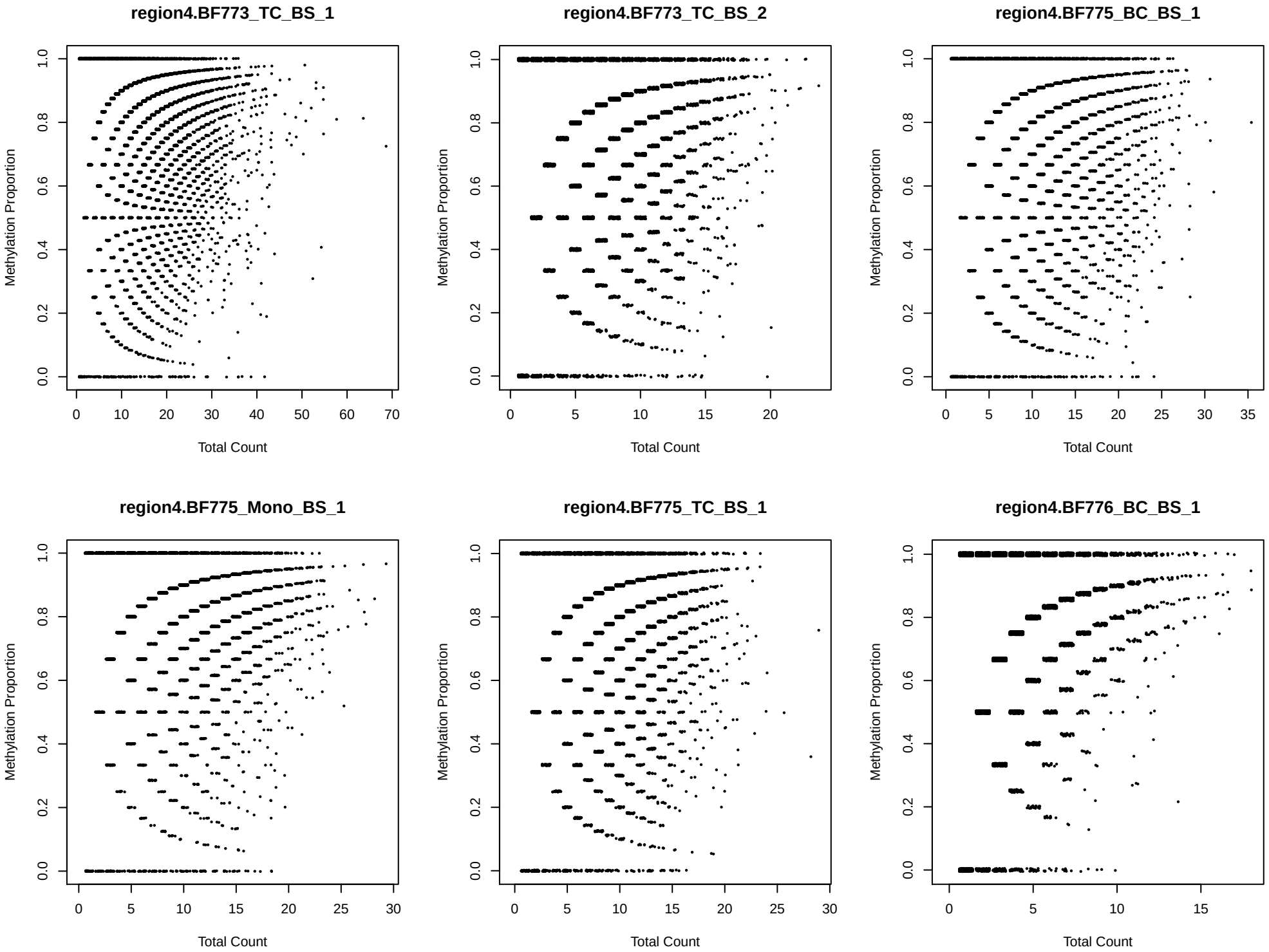

region4.BCU1657_TC_BS_1
Methylation Proportion
Total Count

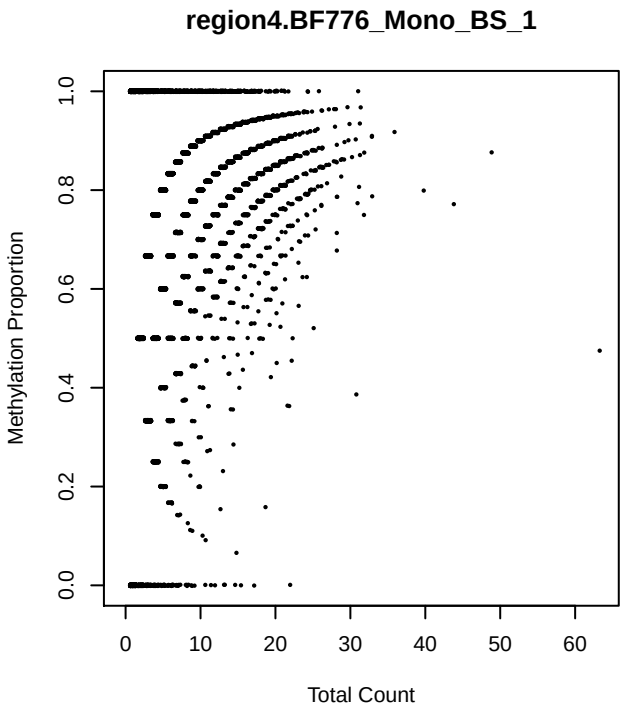

region4.BCU173_TC_BS_1
region4.BCU1799_Mono_BS_1
region4.BCU292_TC_BS_1
region4.BCU551_Mono_BS_1
region4.BCU551_TC_BS_1
region4.BCU582_TC_BS_1
Methylation Proportion
Total Count

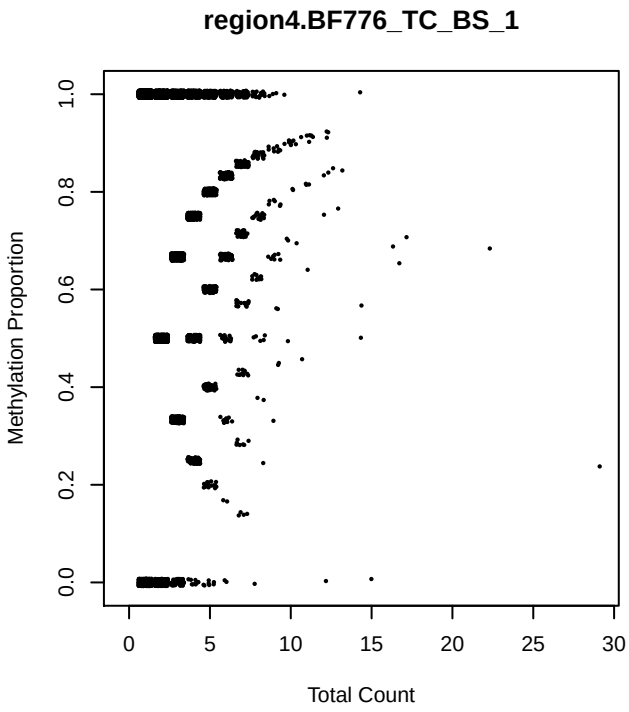

region4.BCU801_Mono_BS_1
region4.BCU801_TC_BS_1
region4.BCU899_TC_BS_1
region4.BF757_TC_BS_1
region4.BF758_Mono_BS_1
region4.BF761_TC_BS_1
Methylation Proportion
Total Count

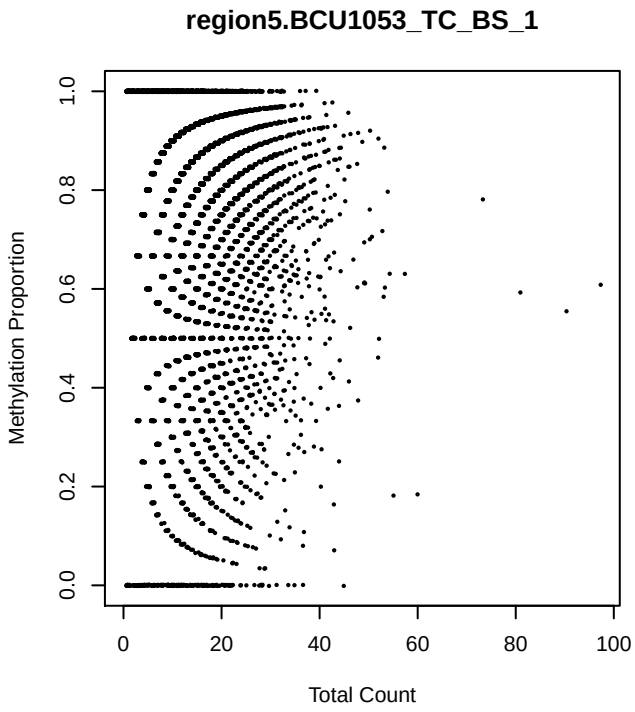

region4.BF764_BC_BS_1
region4.BF771_Mono_BS_1
region4.BF771_TC_BS_1
region4.BF772_BC_BS_2
region4.BF772_Mono_BS_1
region4.BF772_Mono_BS_2
Methylation Proportion
Total Count

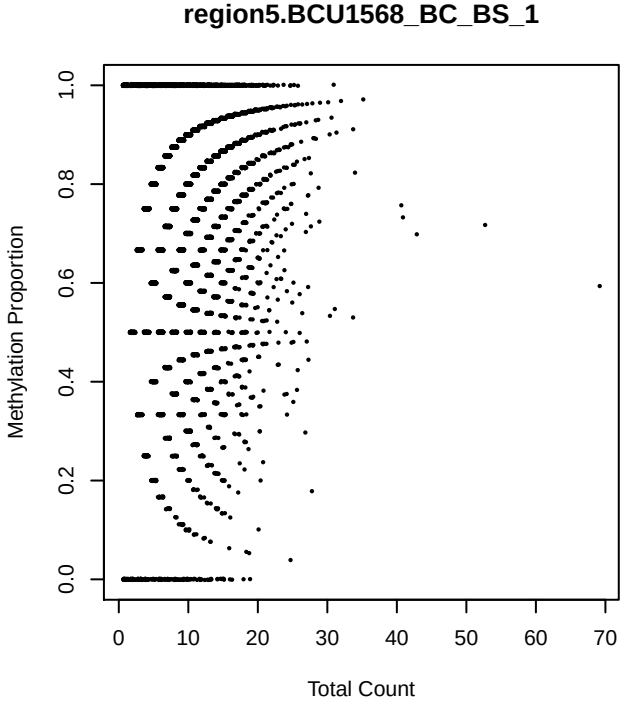

region4.BF772_TC_BS_1
region4.BF772_TC_BS_2
region4.BF773_BC_BS_1
region4.BF773_BC_BS_2
region4.BF773_Mono_BS_1
region4.BF773_Mono_BS_2
Methylation Proportion
Total Count

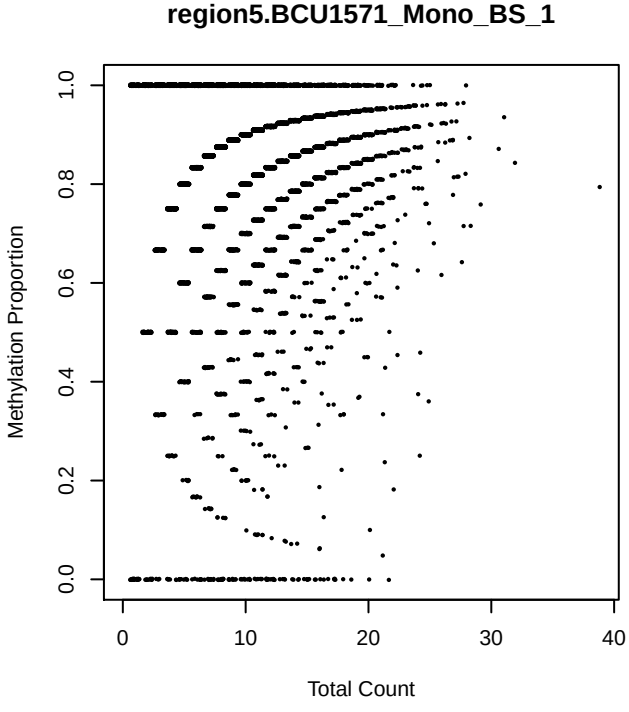

region4.BF773_TC_BS_1
region4.BF773_TC_BS_2
region4.BF775_BC_BS_1
region4.BF775_Mono_BS_1
region4.BF775_TC_BS_1
region4.BF776_BC_BS_1
Methylation Proportion
Total Count

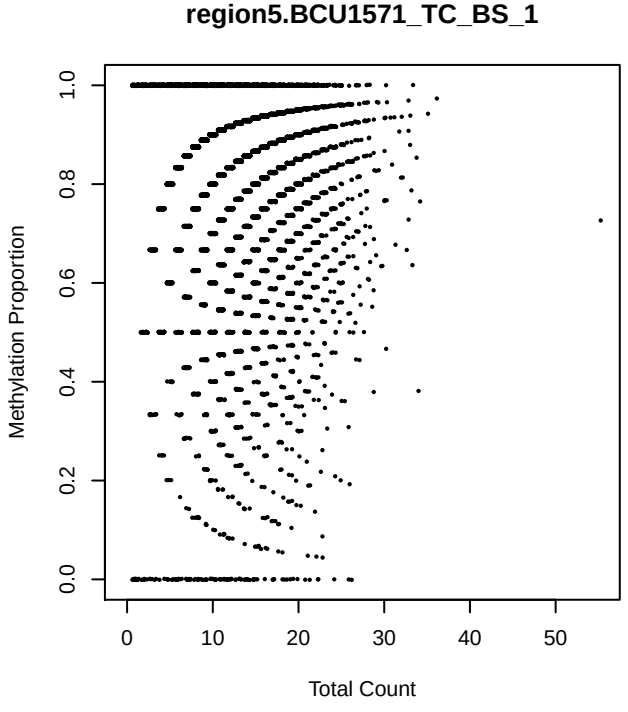

region4.BF776_Mono_BS_1
Methylation Proportion
Total Count

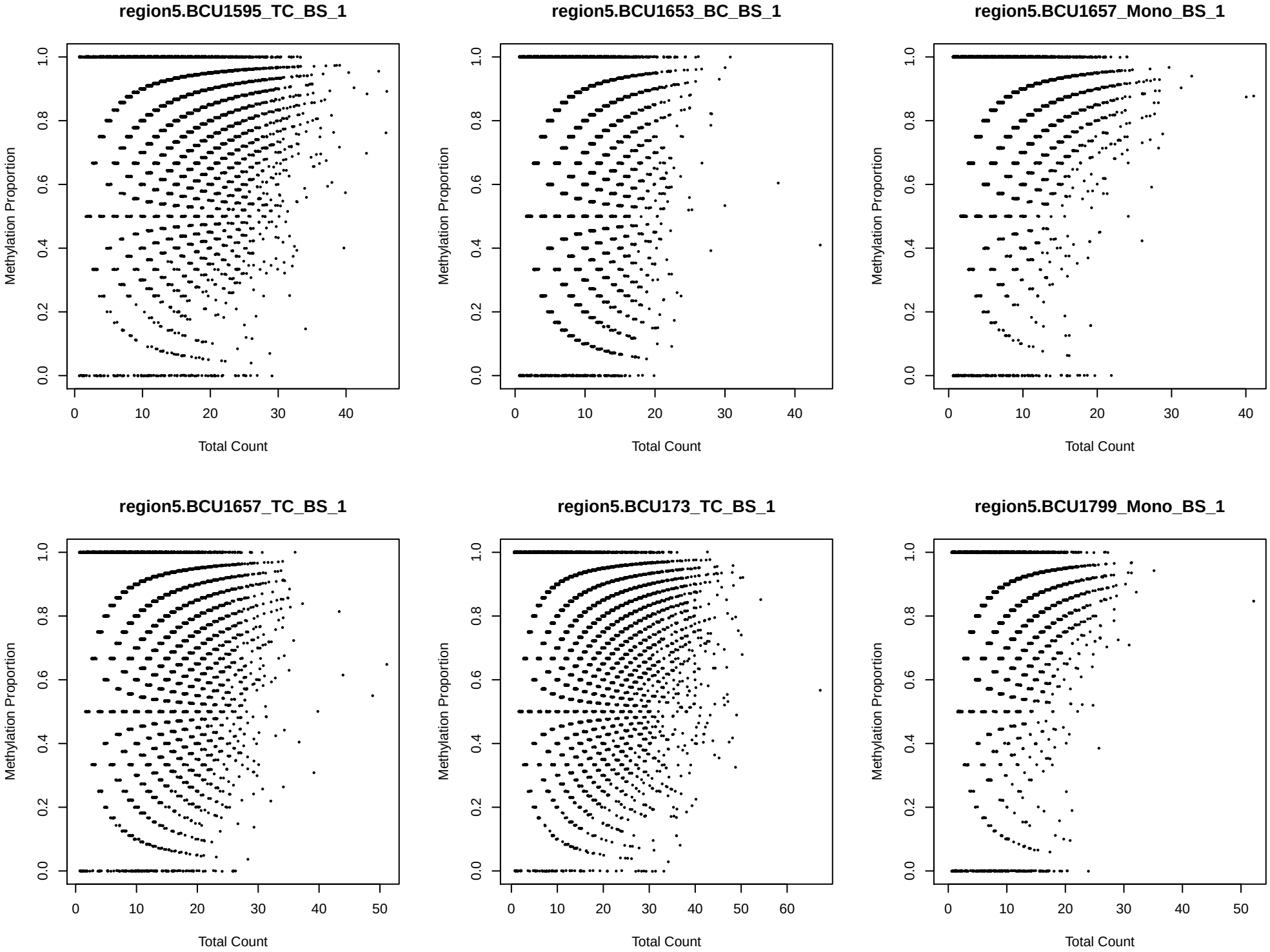

region4.BF776_TC_BS_1
Methylation Proportion
Total Count

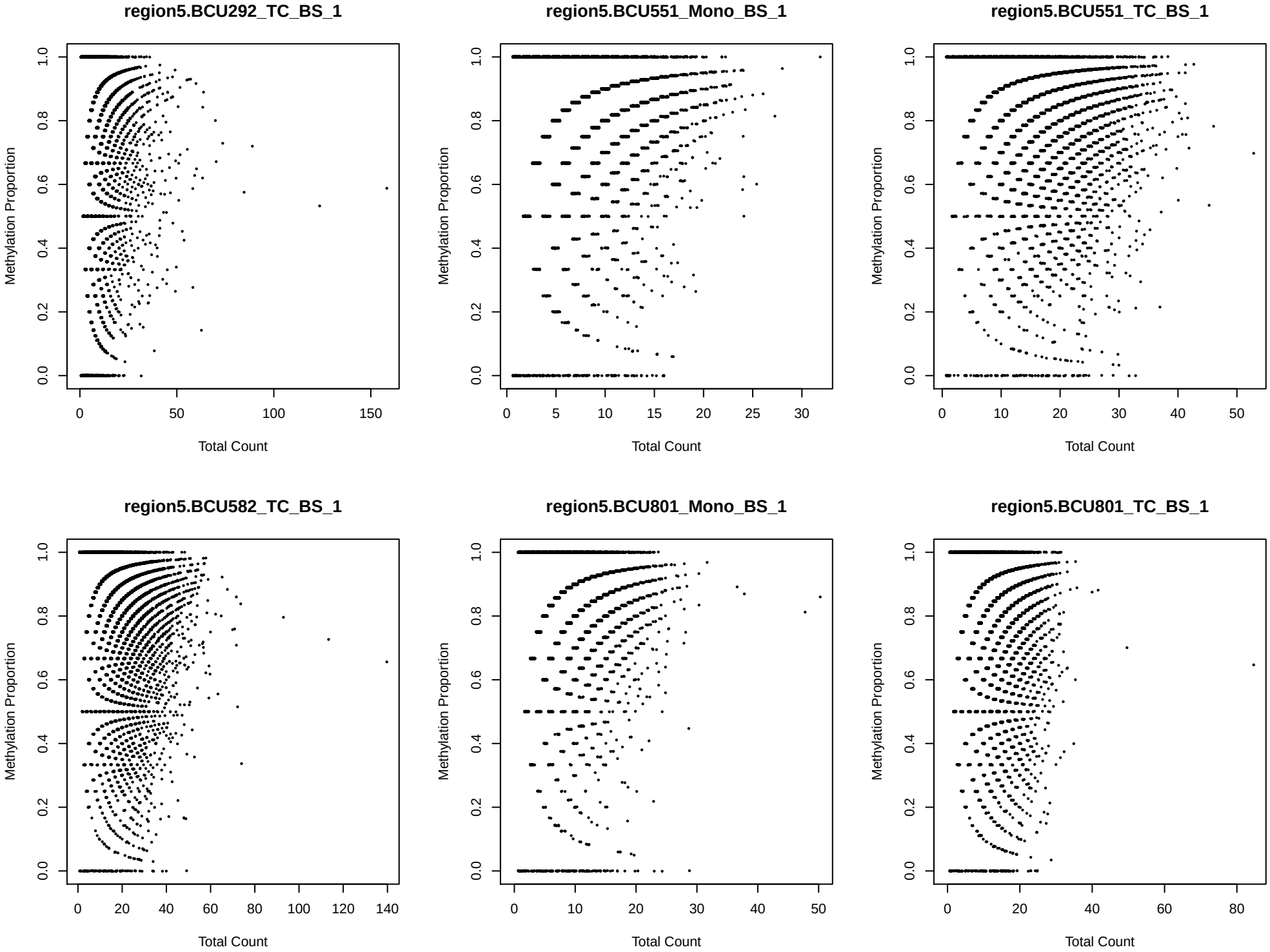

region5.BCU1053_TC_BS_1
Methylation Proportion
Total Count

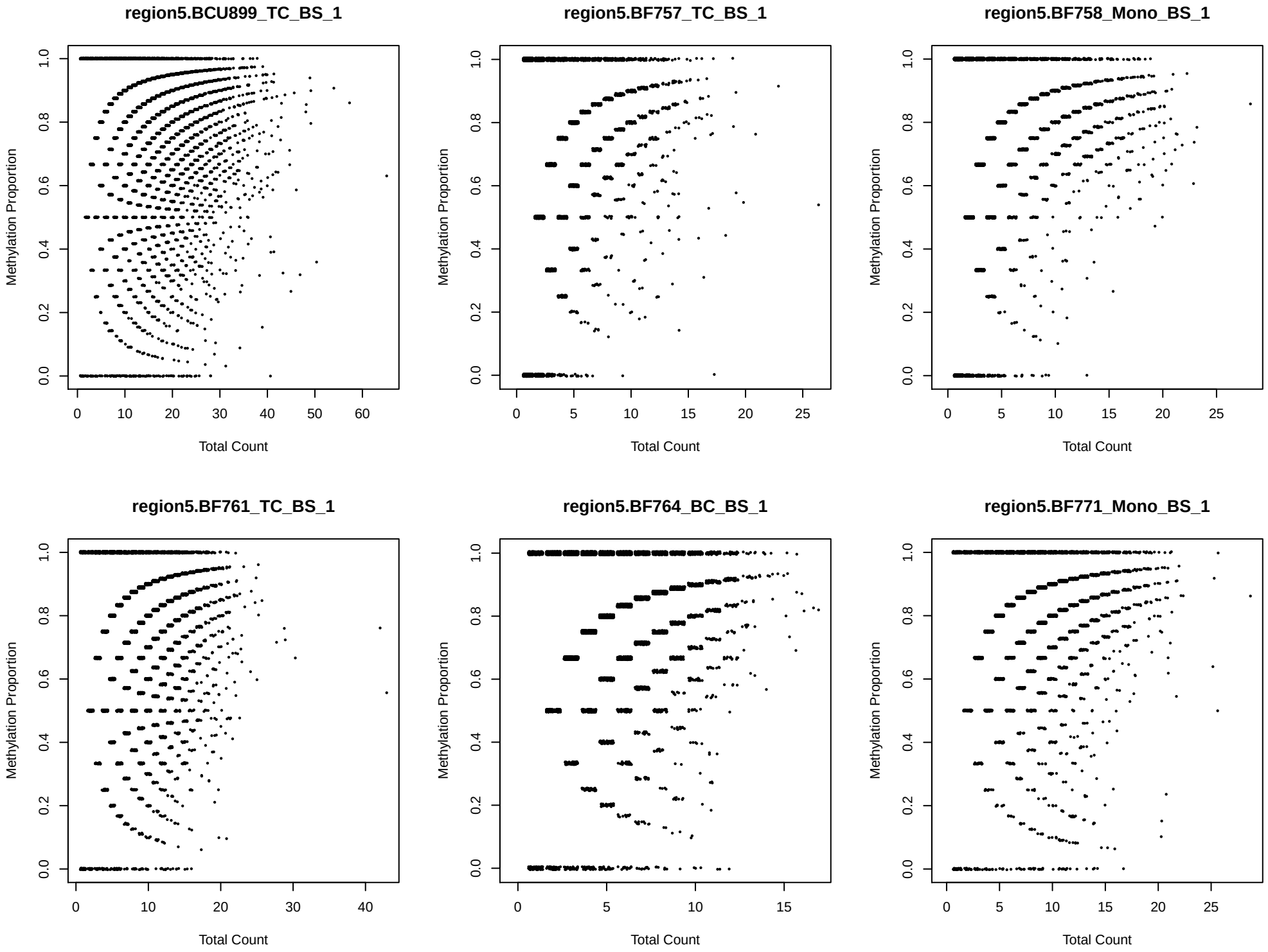

region5.BCU1568_BC_BS_1
Methylation Proportion
Total Count

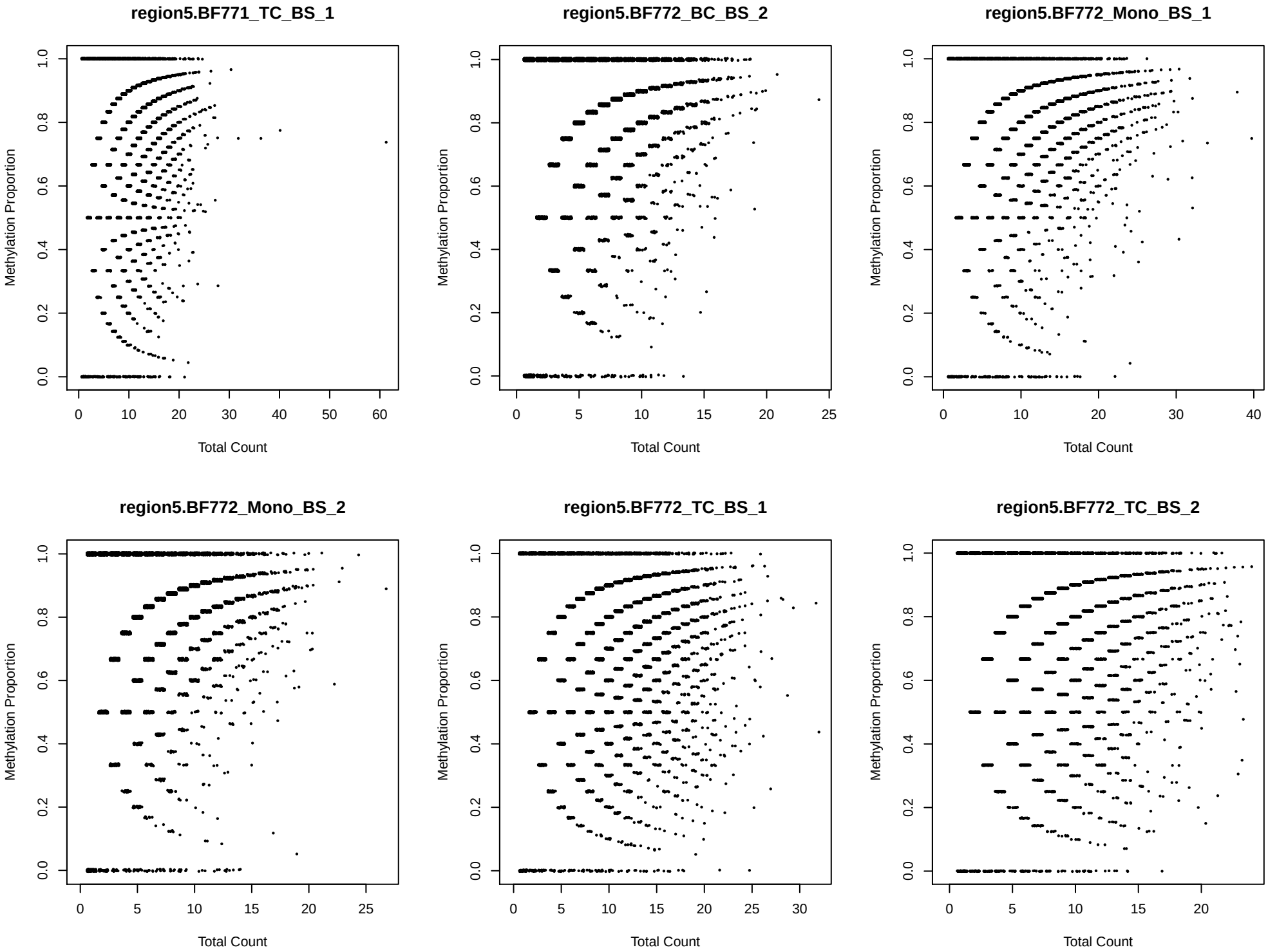

region5.BCU1571_Mono_BS_1
Methylation Proportion
Total Count

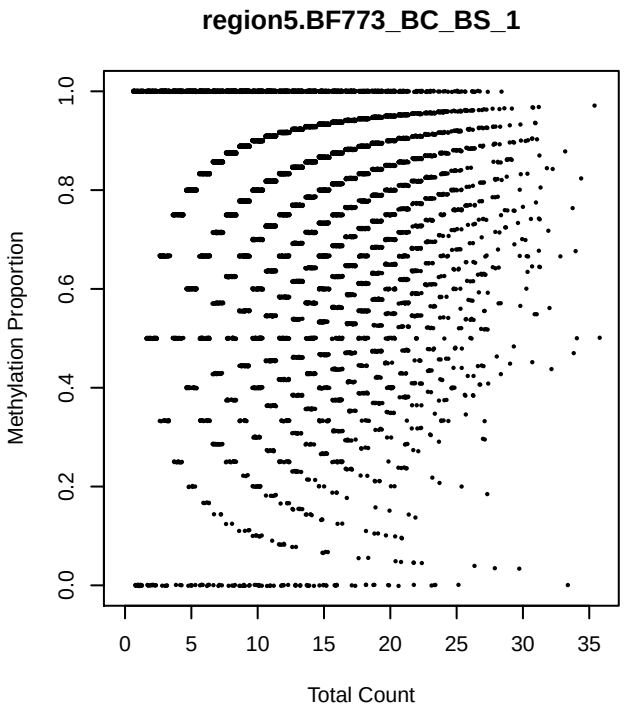

region5.BCU1571_TC_BS_1
Methylation Proportion
Total Count

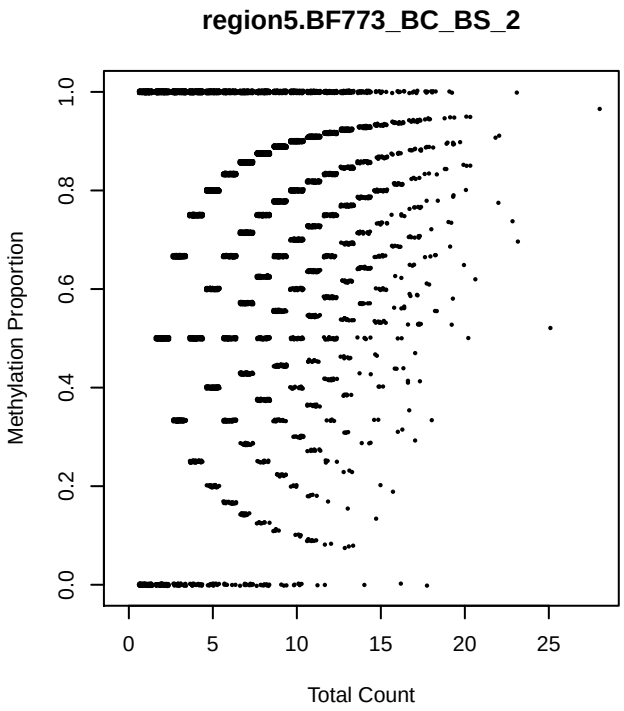

region5.BCU1595_TC_BS_1
region5.BCU1653_BC_BS_1
region5.BCU1657_Mono_BS_1
region5.BCU1657_TC_BS_1
region5.BCU173_TC_BS_1
region5.BCU1799_Mono_BS_1
Methylation Proportion
Total Count

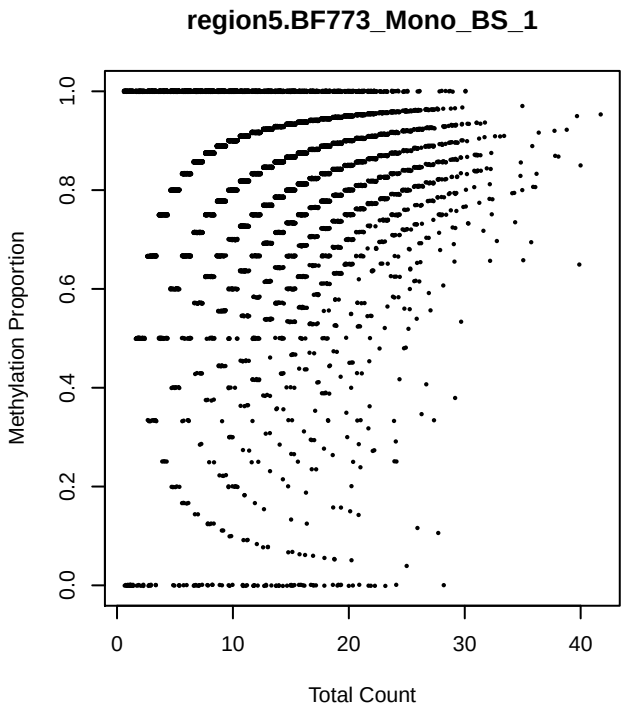

region5.BCU292_TC_BS_1
region5.BCU551_Mono_BS_1
region5.BCU551_TC_BS_1
region5.BCU582_TC_BS_1
region5.BCU801_Mono_BS_1
region5.BCU801_TC_BS_1
Methylation Proportion
Total Count

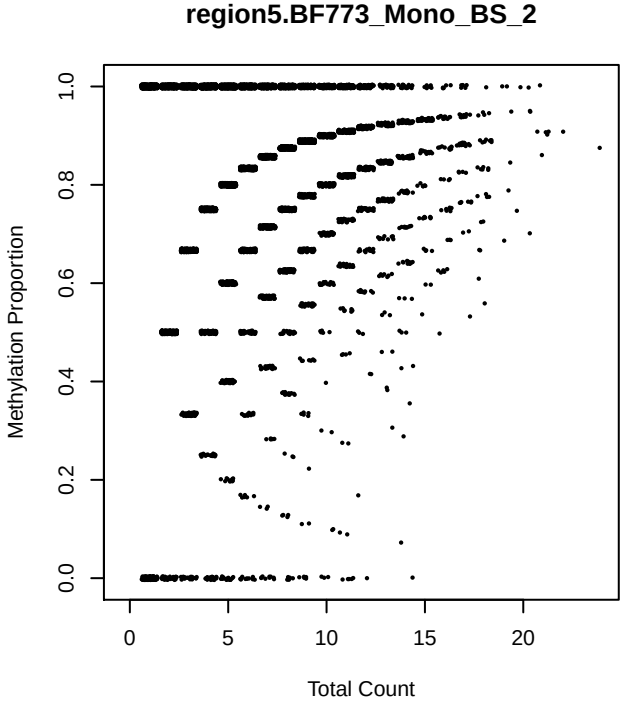

region5.BCU899_TC_BS_1
region5.BF757_TC_BS_1
region5.BF758_Mono_BS_1
region5.BF761_TC_BS_1
region5.BF764_BC_BS_1
region5.BF771_Mono_BS_1
Methylation Proportion
Total Count

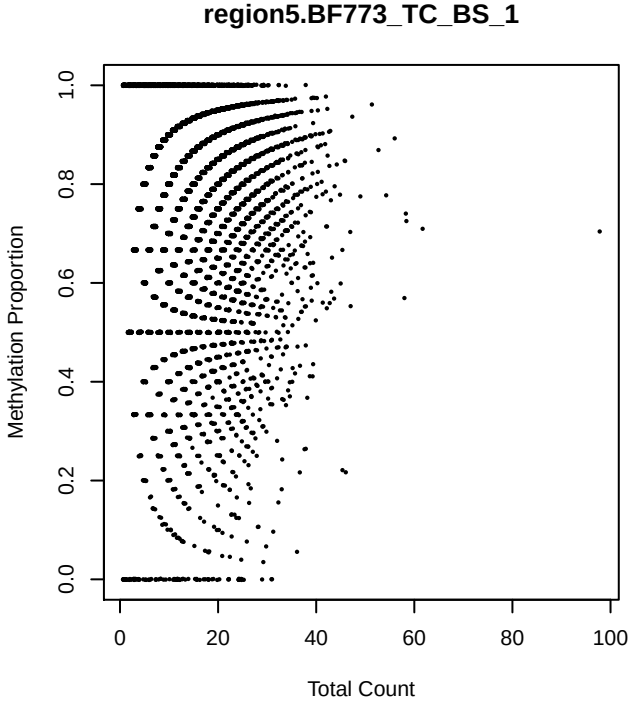

region5.BF771_TC_BS_1
region5.BF772_BC_BS_2
region5.BF772_Mono_BS_1
region5.BF772_Mono_BS_2
region5.BF772_TC_BS_1
region5.BF772_TC_BS_2
Methylation Proportion
Total Count

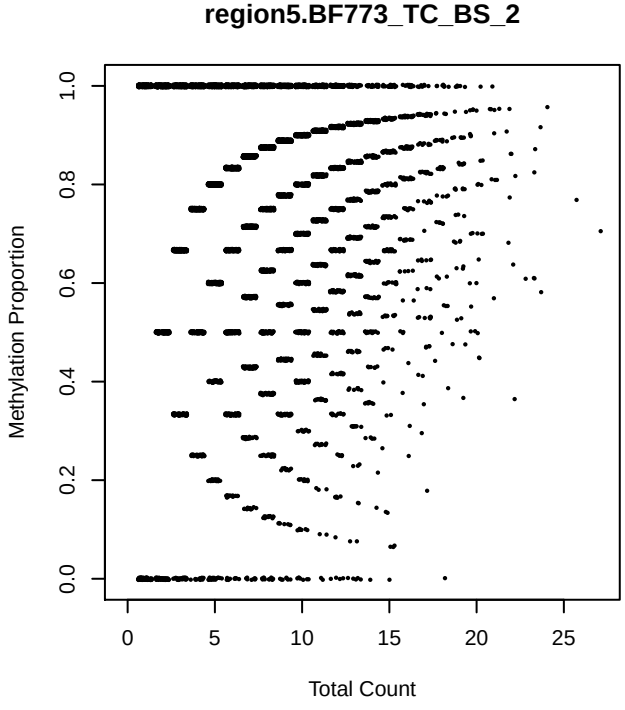

region5.BF773_BC_BS_1
Methylation Proportion
Total Count

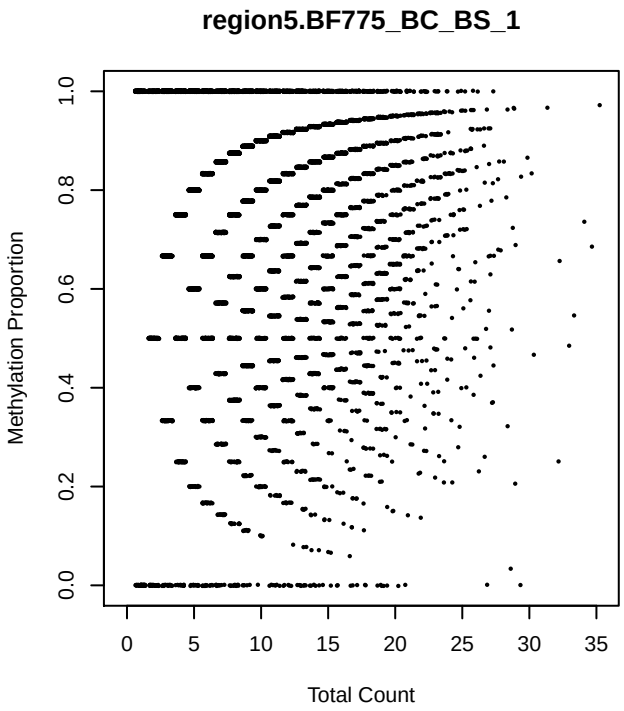

region5.BF773_BC_BS_2
Methylation Proportion
Total Count

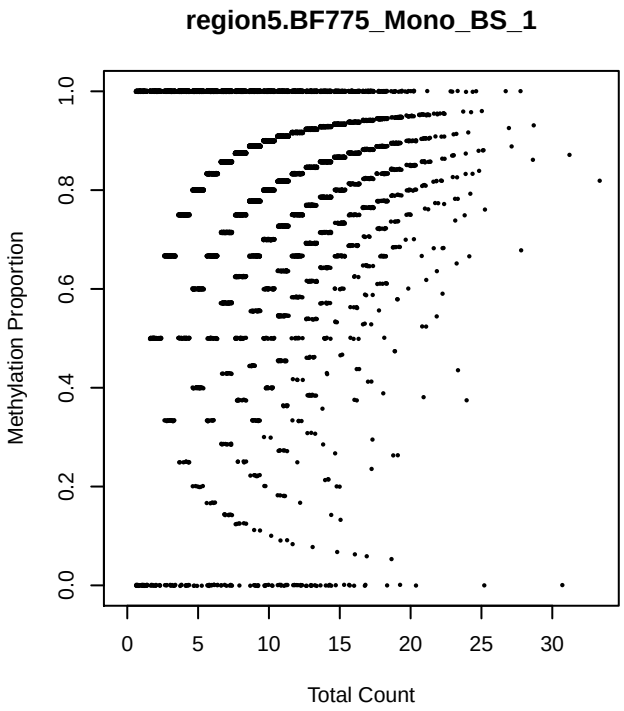

region5.BF773_Mono_BS_1
Methylation Proportion
Total Count

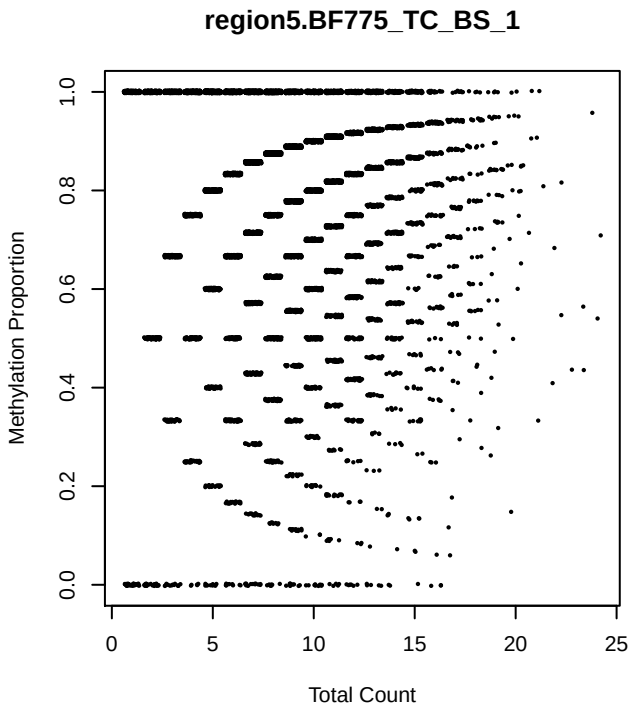

region5.BF773_Mono_BS_2
Methylation Proportion
Total Count

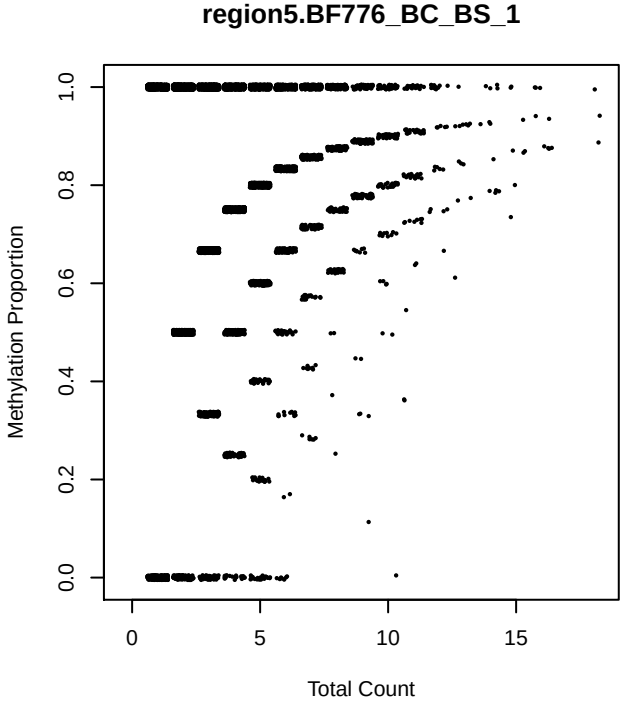

region5.BF773_TC_BS_1
Methylation Proportion
Total Count

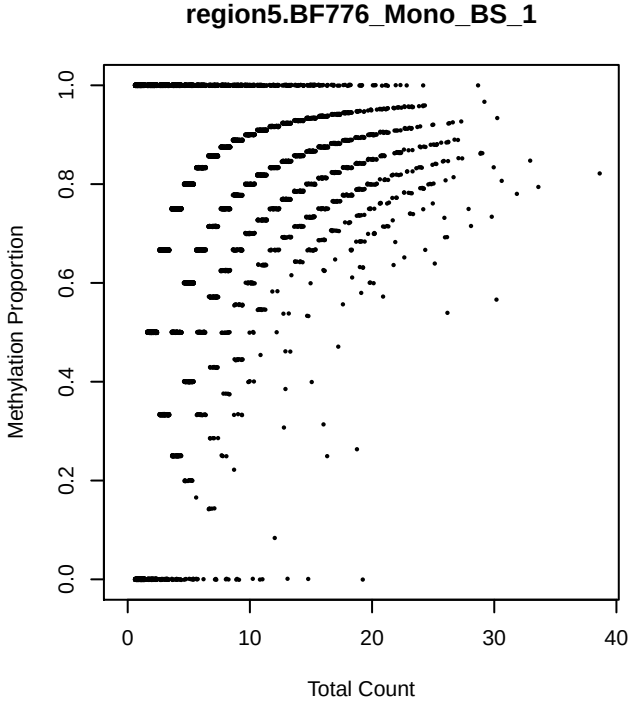

region5.BF773_TC_BS_2
Methylation Proportion
Total Count

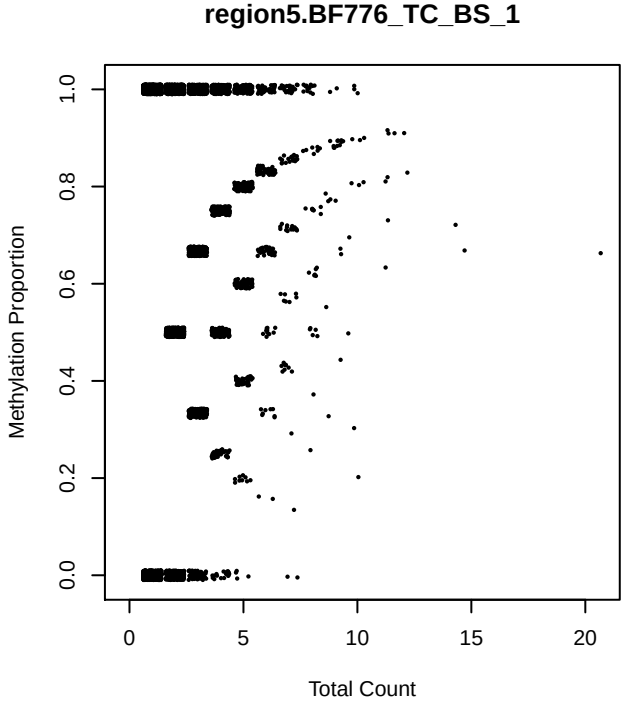

region5.BF775_BC_BS_1
Methylation Proportion
Total Count

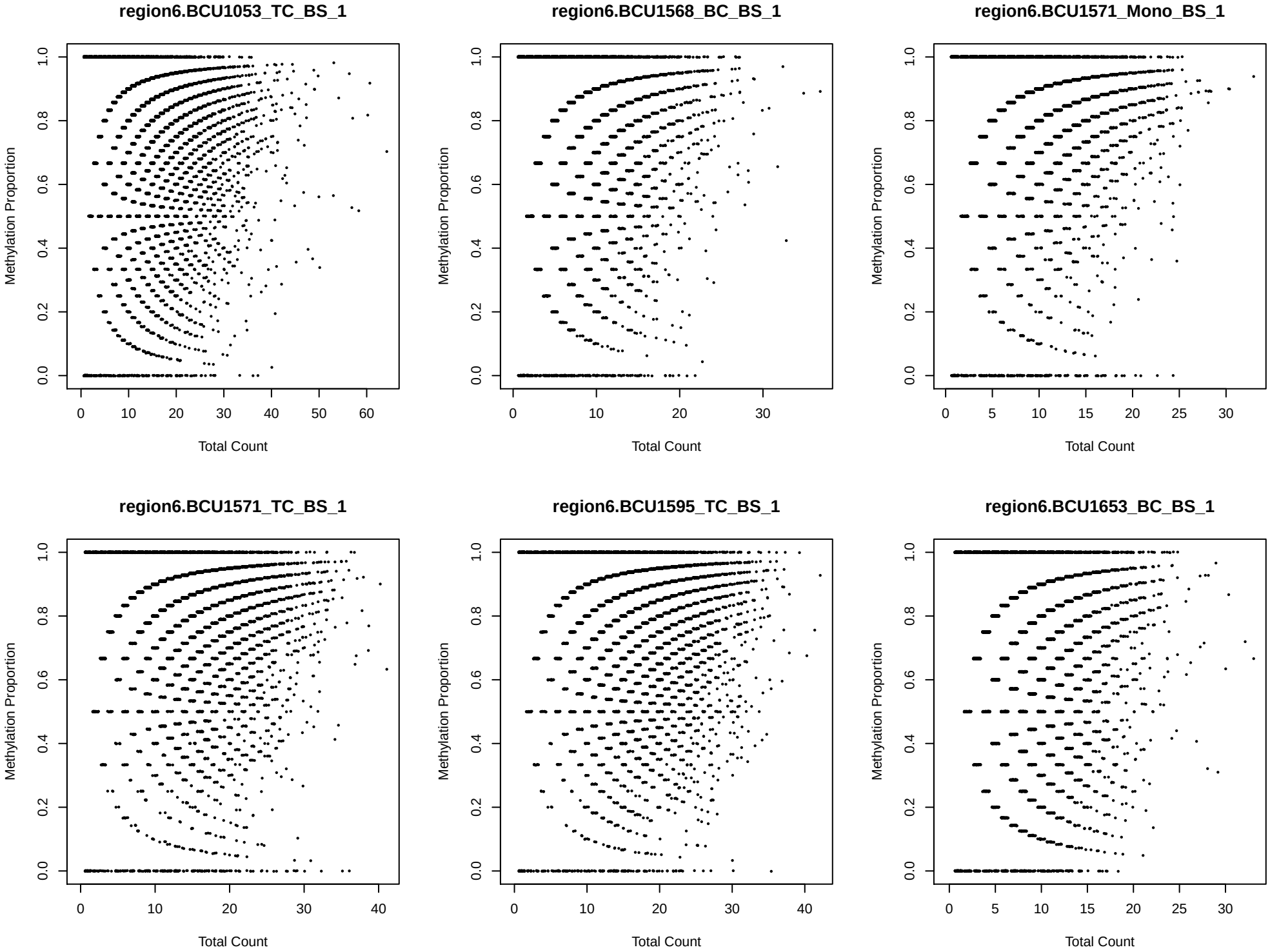

region5.BF775_Mono_BS_1
Methylation Proportion
Total Count

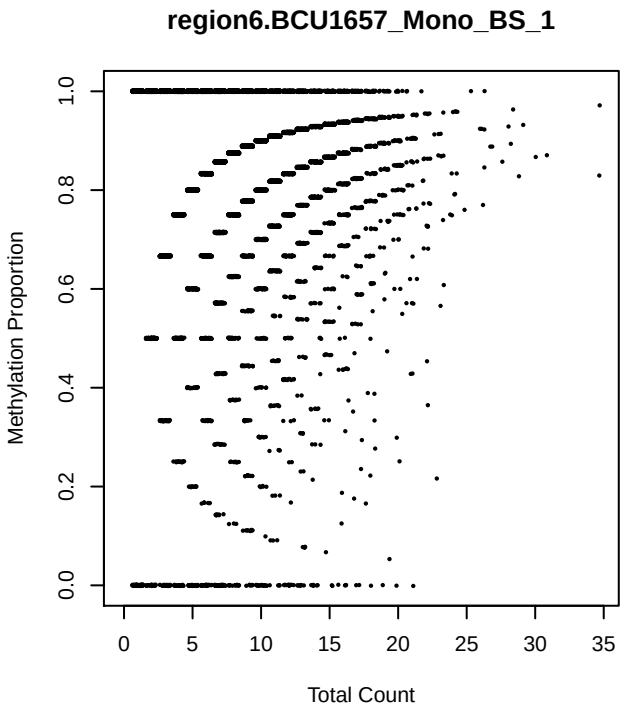

region5.BF775_TC_BS_1
Methylation Proportion
Total Count

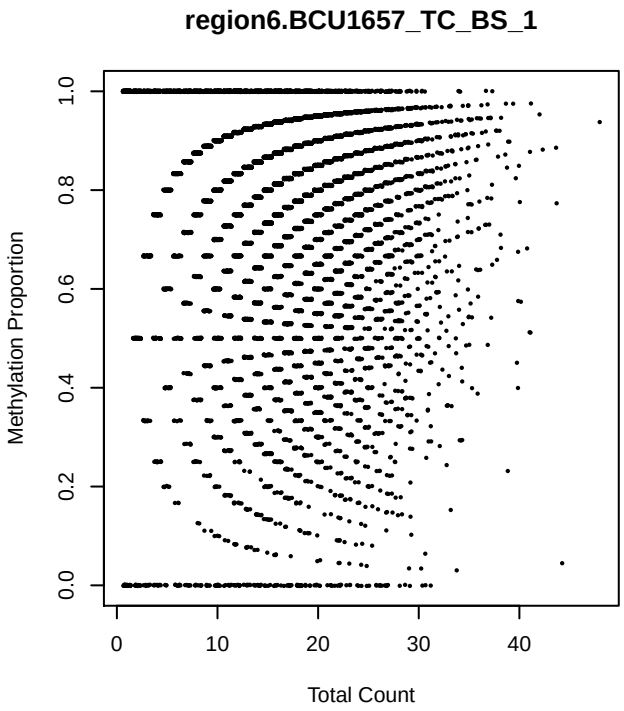

region5.BF776_BC_BS_1
Methylation Proportion
Total Count

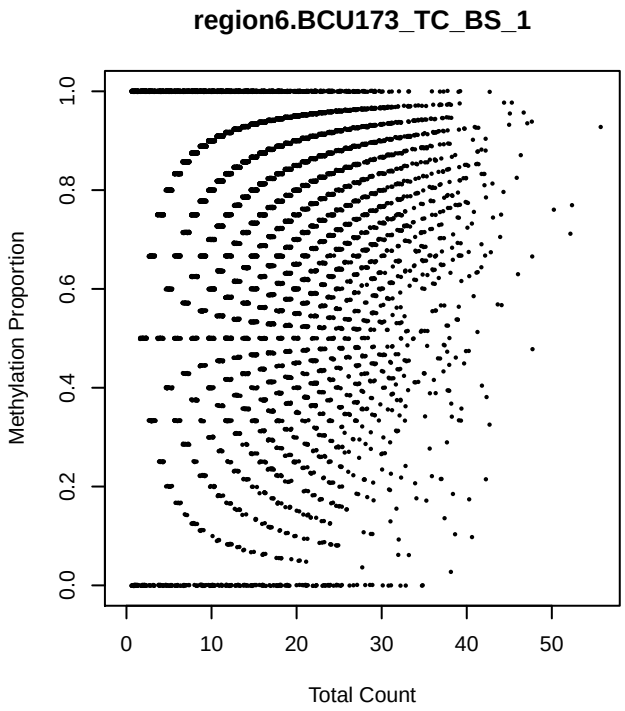

region5.BF776_Mono_BS_1
Methylation Proportion
Total Count

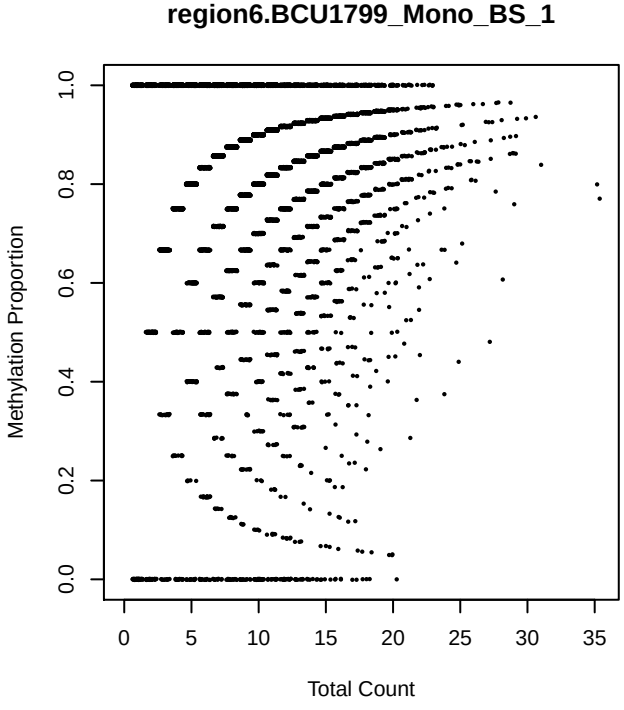

region5.BF776_TC_BS_1
Methylation Proportion
Total Count

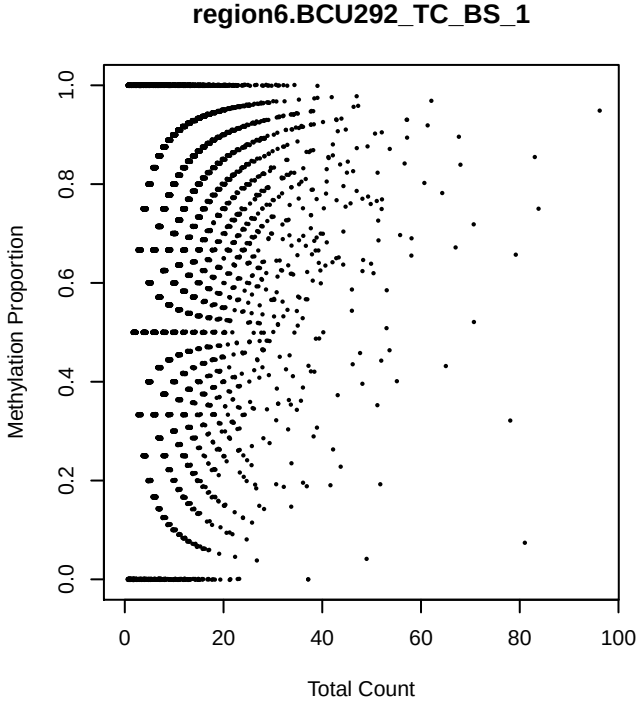

region6.BCU1053_TC_BS_1
region6.BCU1568_BC_BS_1
region6.BCU1571_Mono_BS_1
region6.BCU1571_TC_BS_1
region6.BCU1595_TC_BS_1
region6.BCU1653_BC_BS_1
Methylation Proportion
Total Count

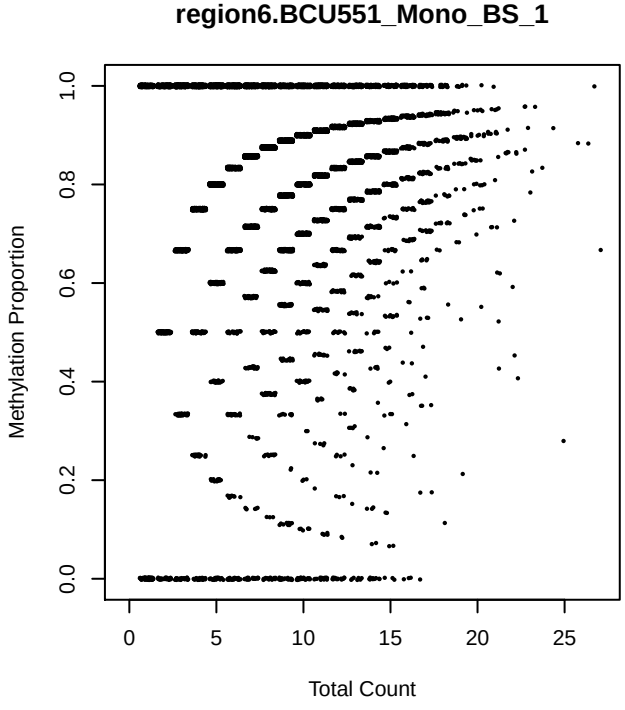

region6.BCU1657_Mono_BS_1
Methylation Proportion
Total Count

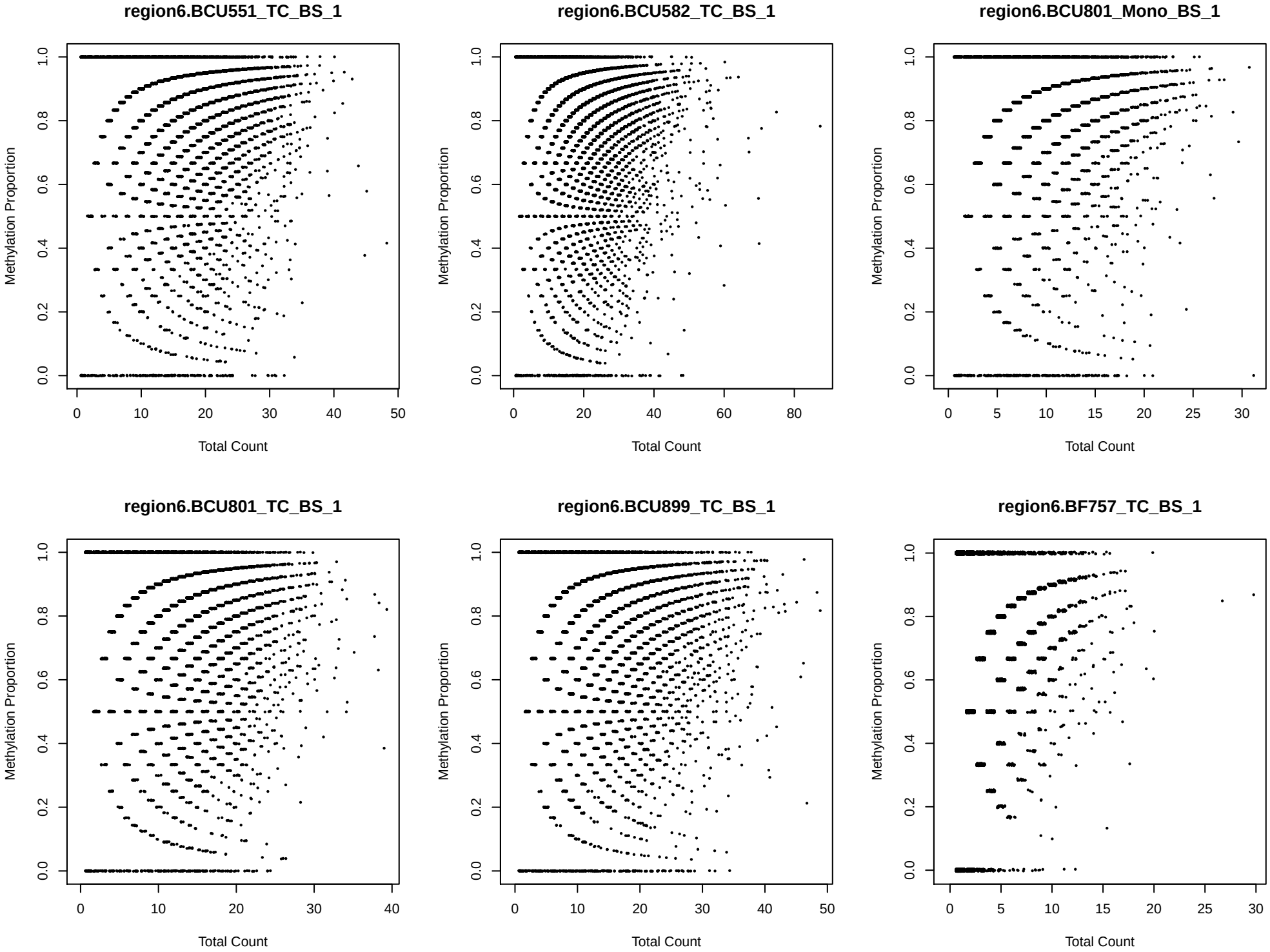

region6.BCU1657_TC_BS_1
Methylation Proportion
Total Count

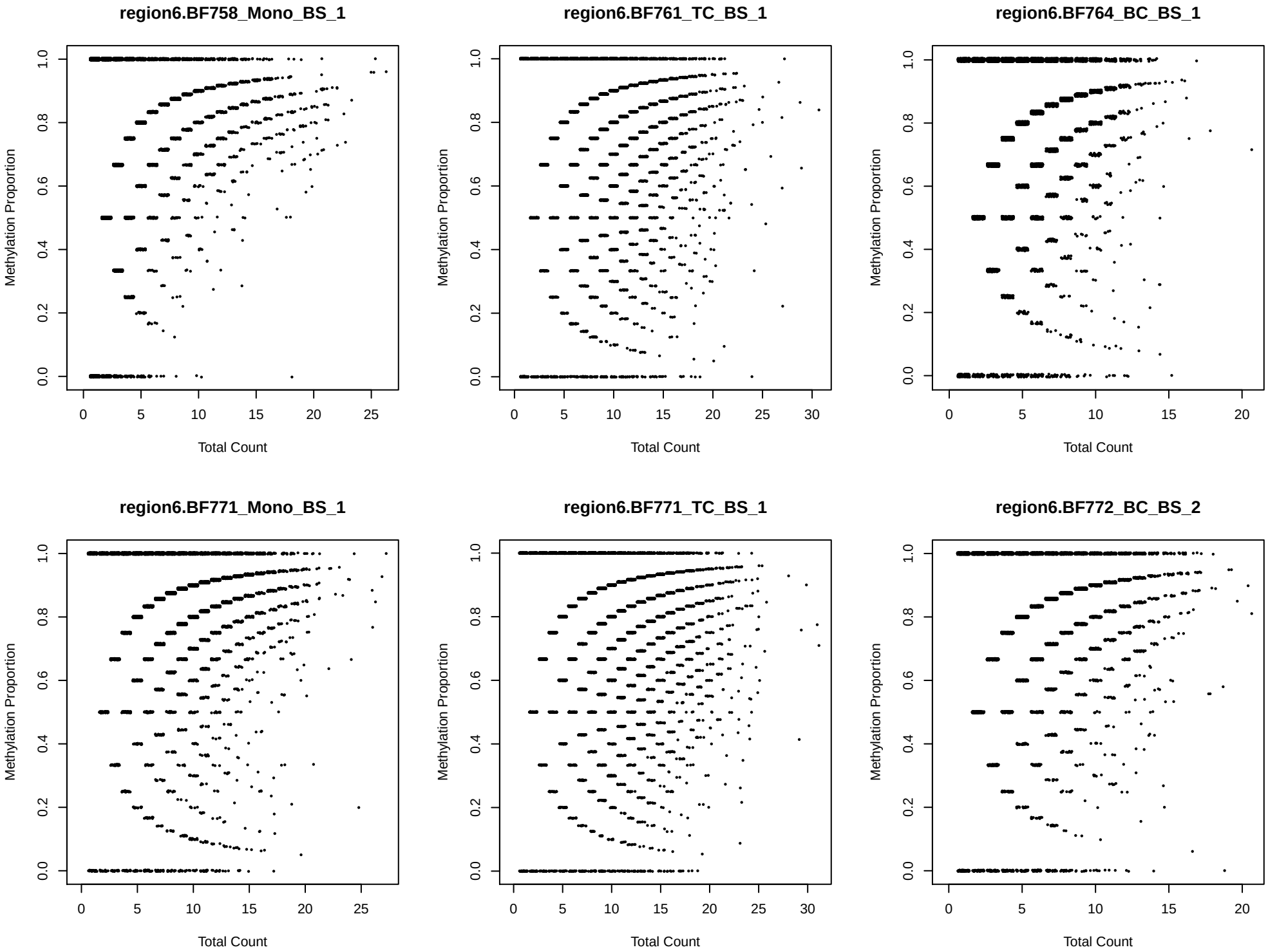

region6.BCU173_TC_BS_1
Methylation Proportion
Total Count

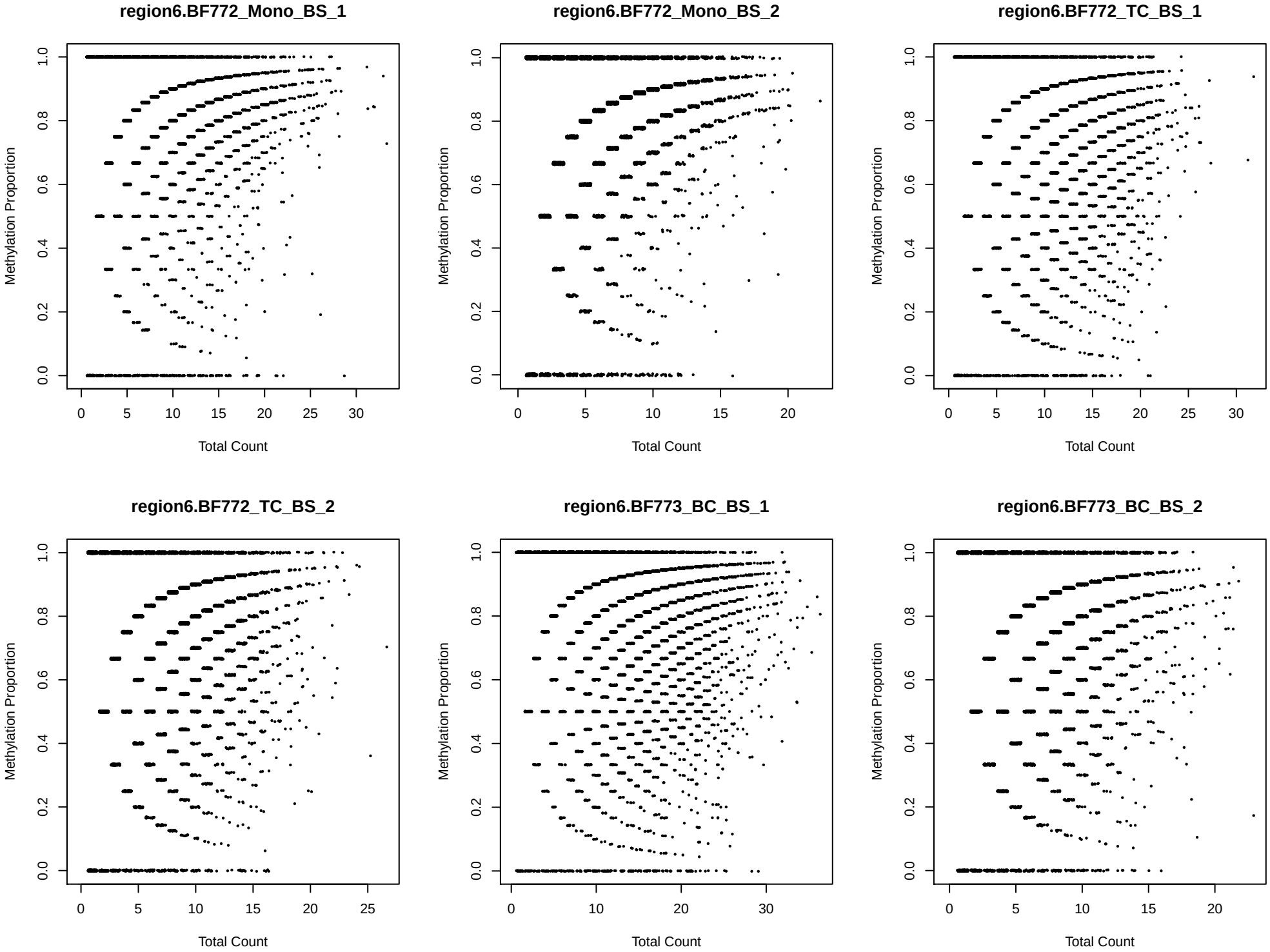

region6.BCU1799_Mono_BS_1
Methylation Proportion
Total Count

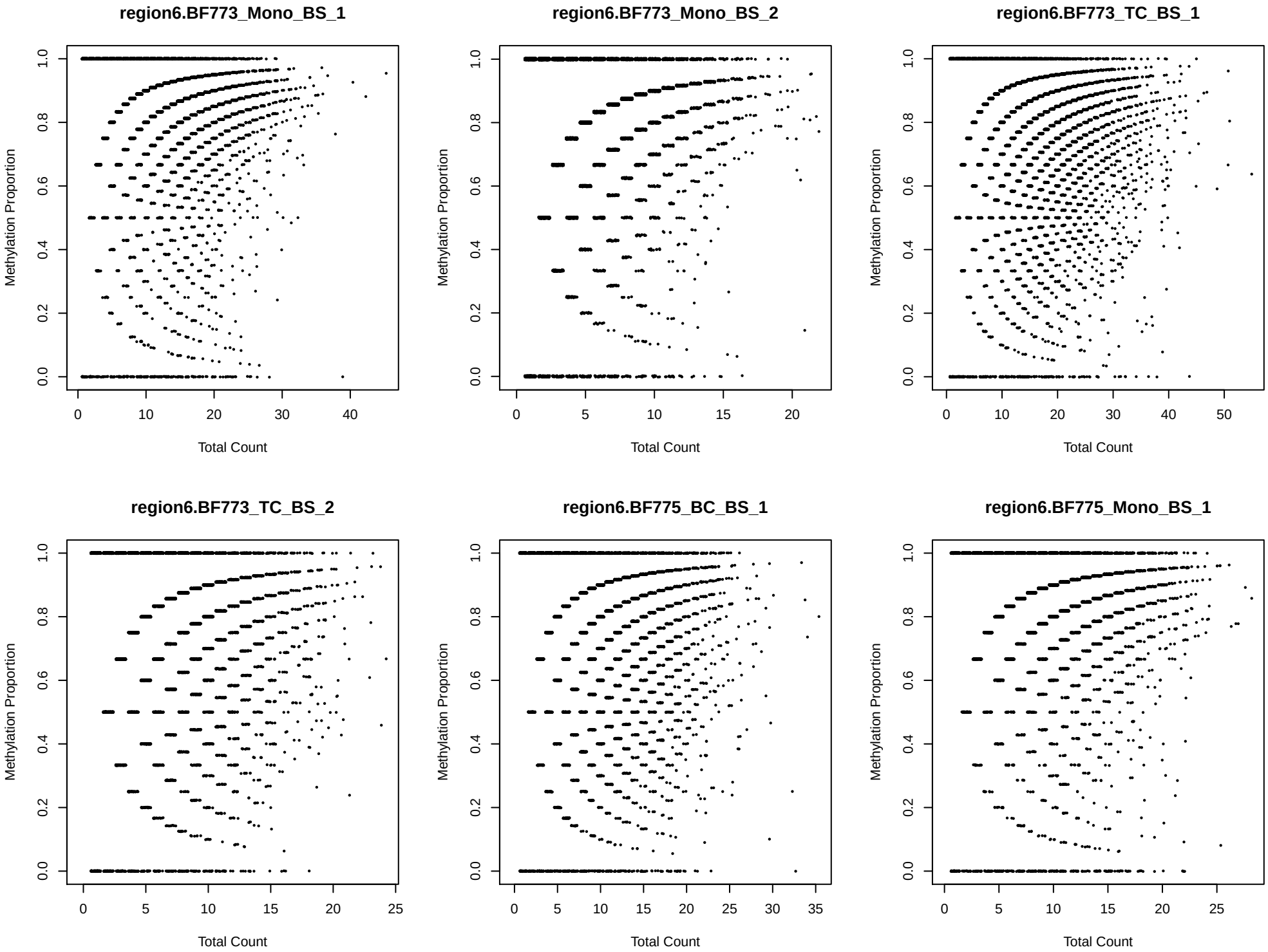

region6.BCU292_TC_BS_1
Methylation Proportion
Total Count

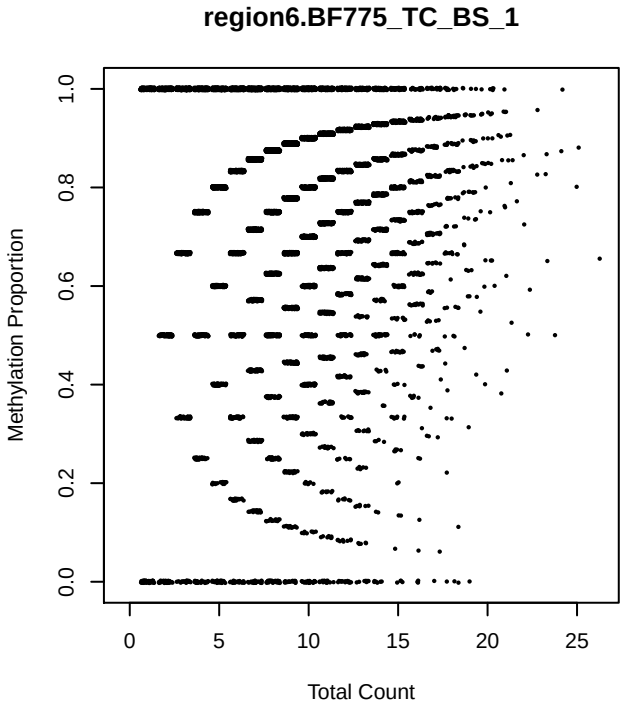

region6.BCU551_Mono_BS_1
Methylation Proportion
Total Count

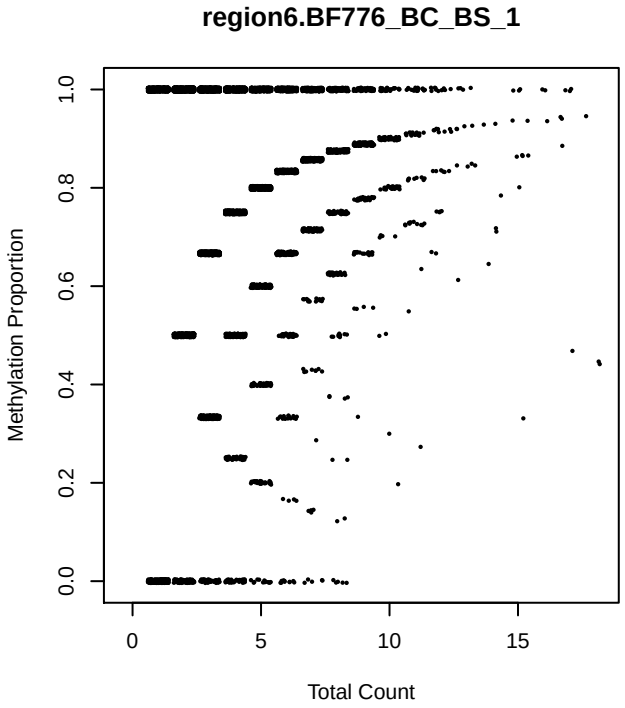

region6.BCU551_TC_BS_1
region6.BCU582_TC_BS_1
region6.BCU801_Mono_BS_1
region6.BCU801_TC_BS_1
region6.BCU899_TC_BS_1
region6.BF757_TC_BS_1
Methylation Proportion
Total Count

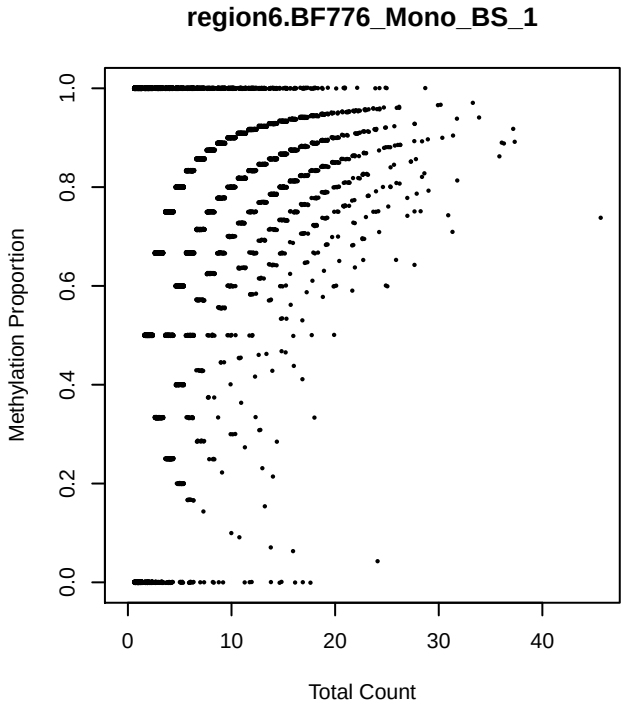

region6.BF758_Mono_BS_1
region6.BF761_TC_BS_1
region6.BF764_BC_BS_1
region6.BF771_Mono_BS_1
region6.BF771_TC_BS_1
region6.BF772_BC_BS_2
Methylation Proportion
Total Count
0.0
0.2
0.4
0.6
0.8
1.0
0
5
10
15
20
25
30

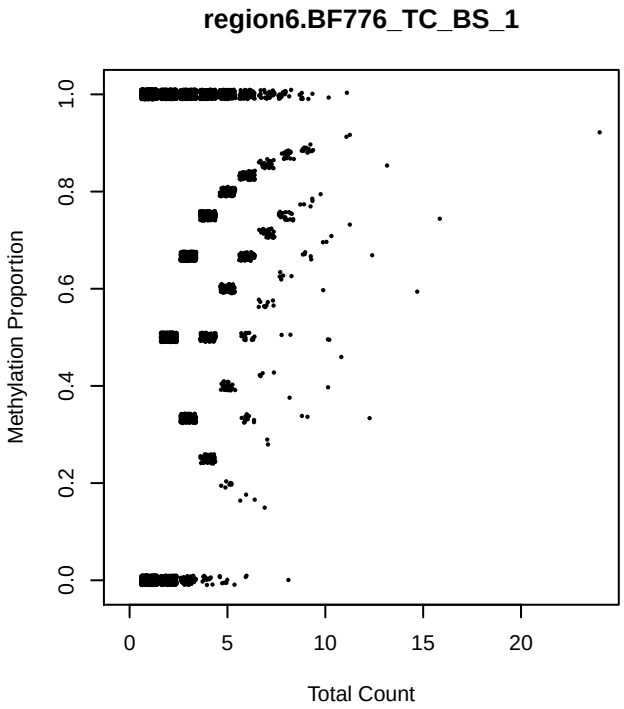

region6.BF772_Mono_BS_1
region6.BF772_Mono_BS_2
region6.BF772_TC_BS_1
region6.BF772_TC_BS_2
region6.BF773_BC_BS_1
region6.BF773_BC_BS_2
Methylation Proportion
Total Count


region6.BF773_Mono_BS_1
region6.BF773_Mono_BS_2
region6.BF773_TC_BS_1
region6.BF773_TC_BS_2
region6.BF775_BC_BS_1
region6.BF775_Mono_BS_1
Methylation Proportion
Total Count


region6.BF775_TC_BS_1
Methylation Proportion
Total Count
0.0
0.2
0.4
0.6
0.8
1.0
0
5
10
15
20
25


region6.BF776_BC_BS_1
Methylation Proportion
Total Count
0.0
0.2
0.4
0.6
0.8
1.0
0
5
10
15


region6.BF776_Mono_BS_1
Methylation Proportion
Total Count
0.0
0.2
0.4
0.6
0.8
1.0
0
10
20
30
40


region6.BF776_TC_BS_1
Methylation Proportion
Total Count
0.0
0.2
0.4
0.6
0.8
1.0
0
5
10
15
20