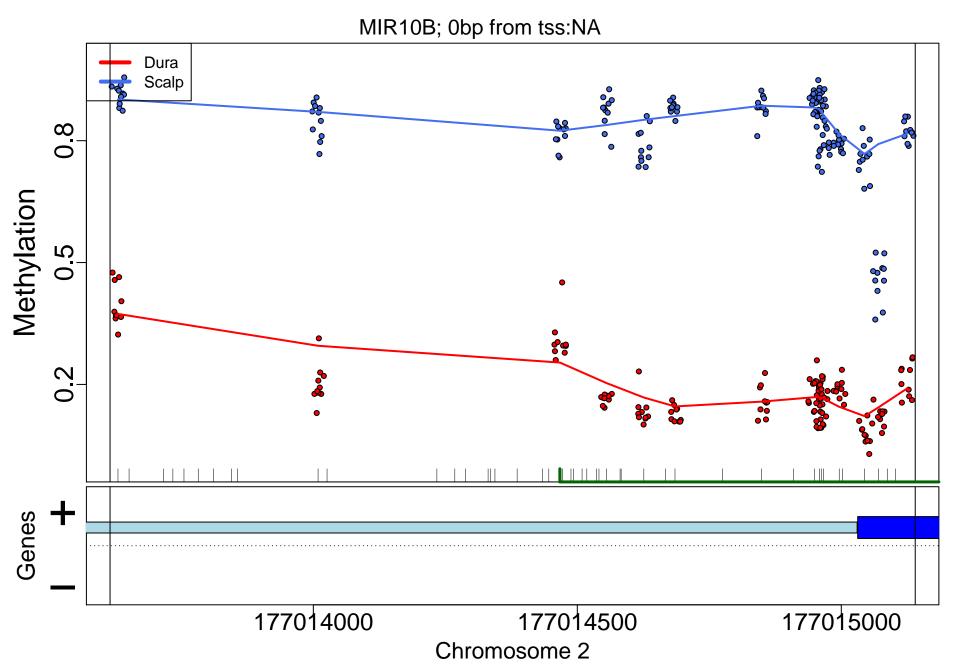
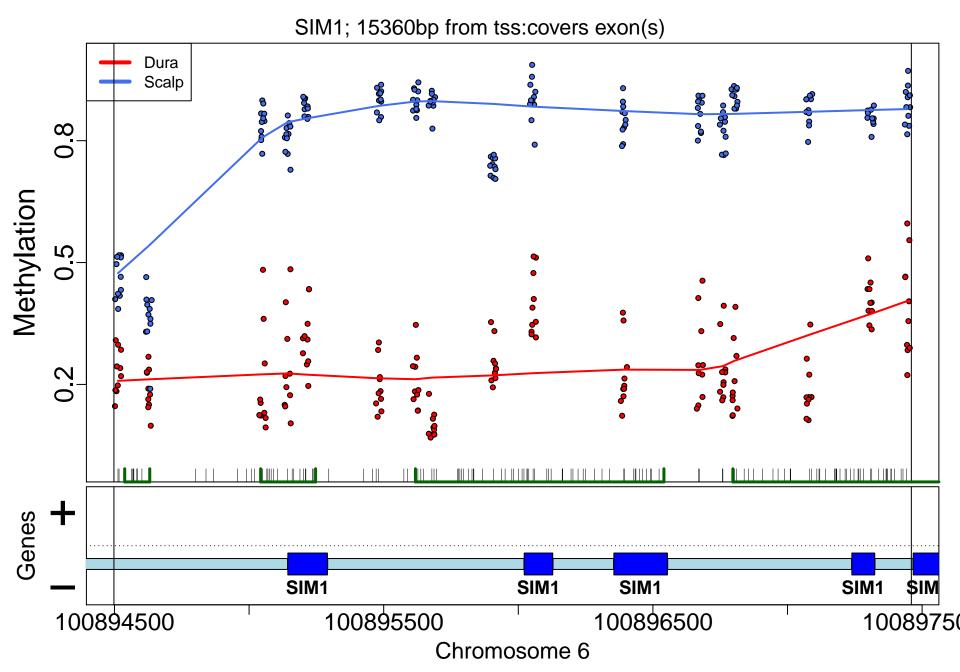
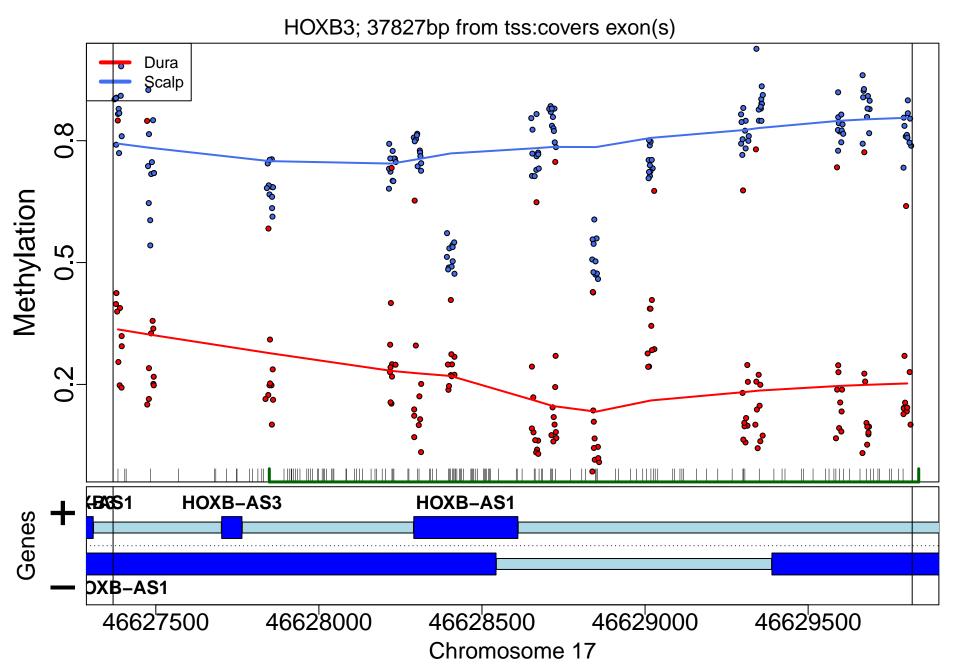
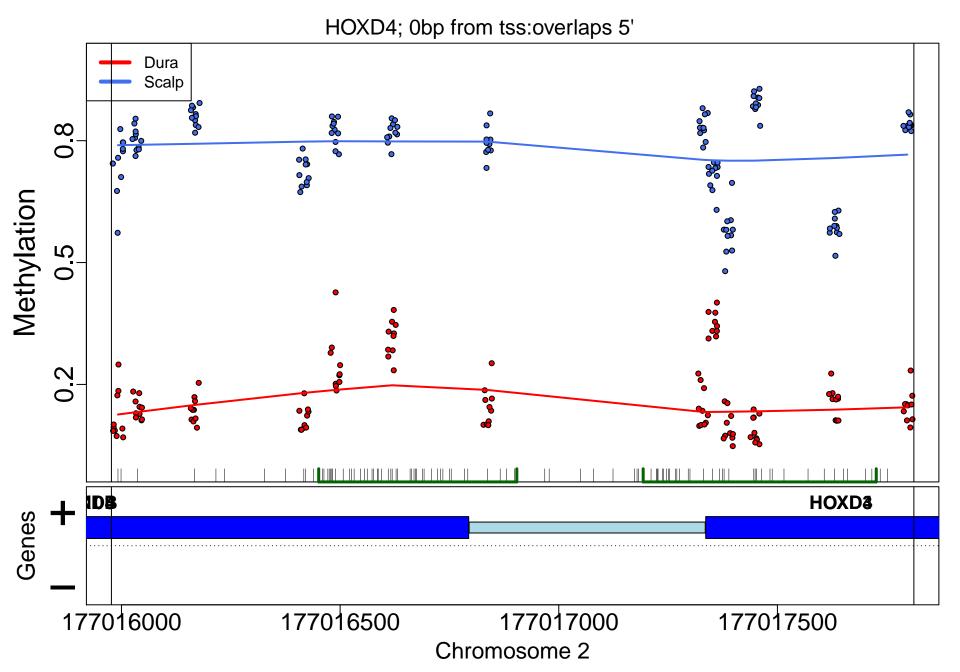


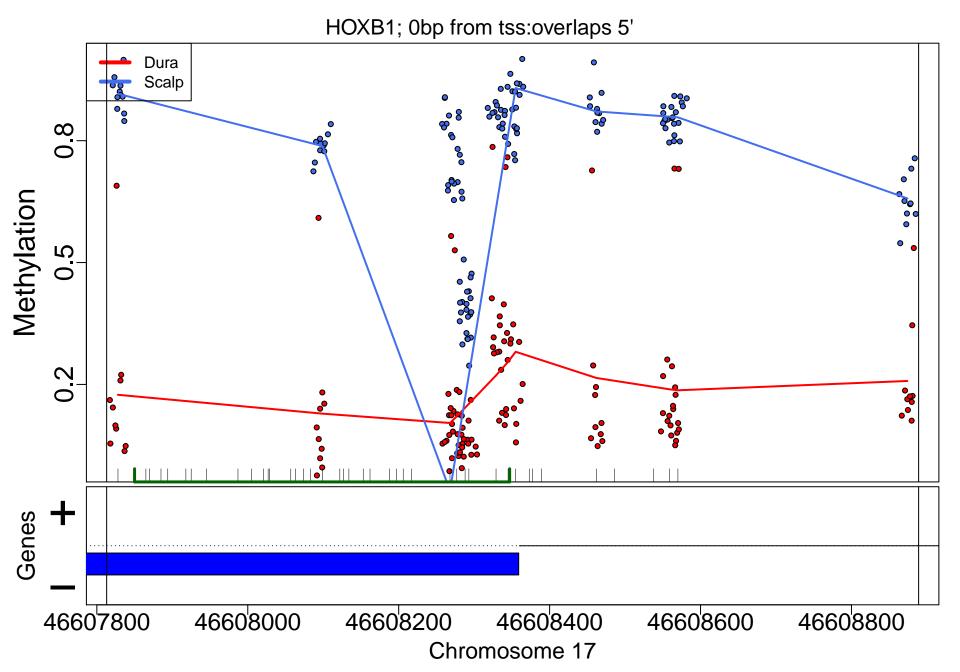
RUNX3; 32933bp from tss:inside intron Dura Scalp 80. Methylation 0.2 Genes **RUNX3** 25256000 25257000 25258000 25259000 Chromosome 1

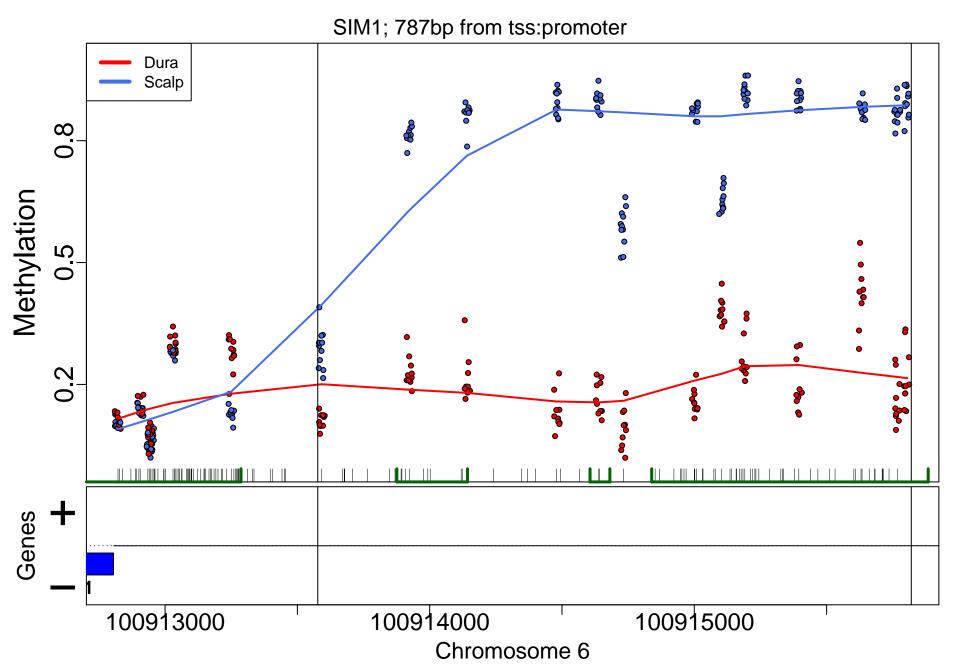


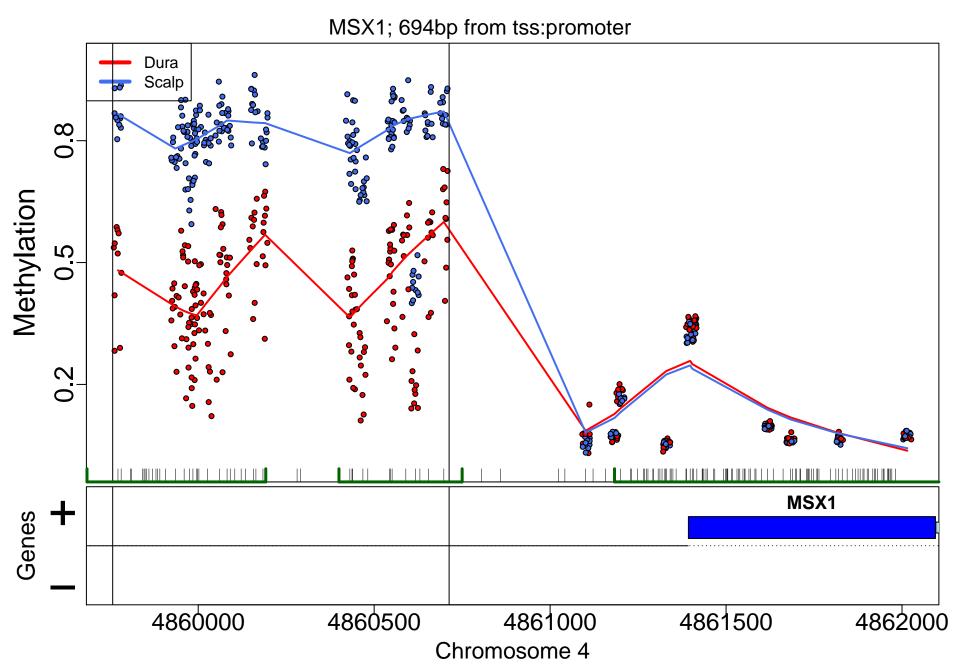


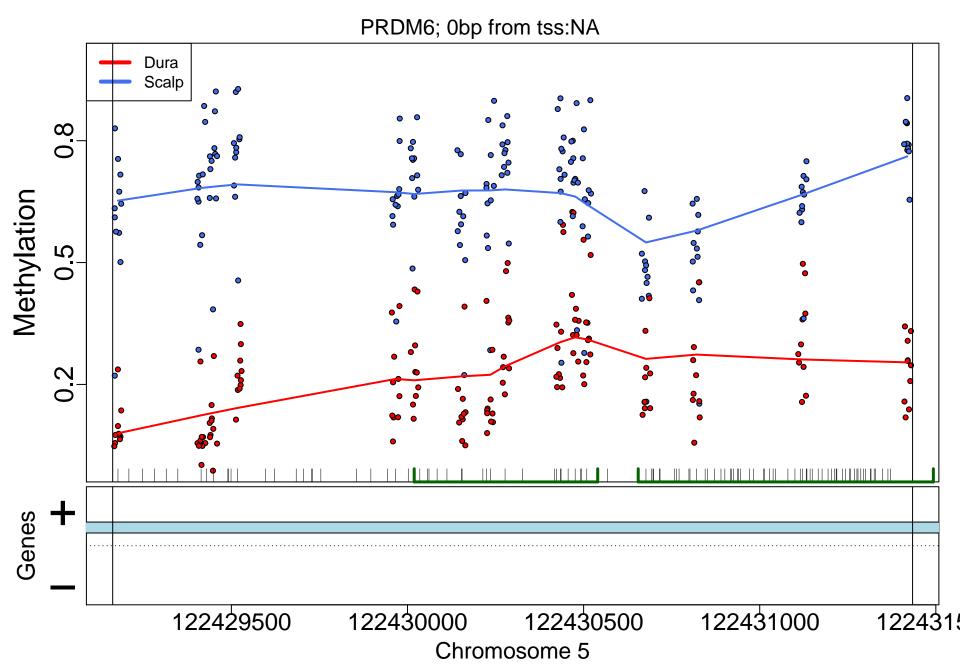


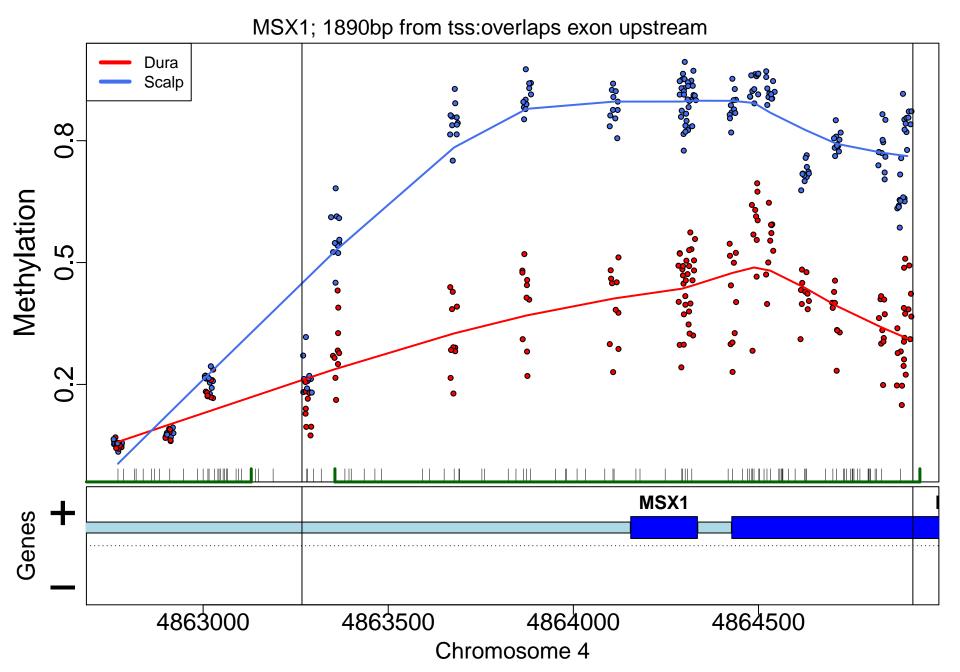


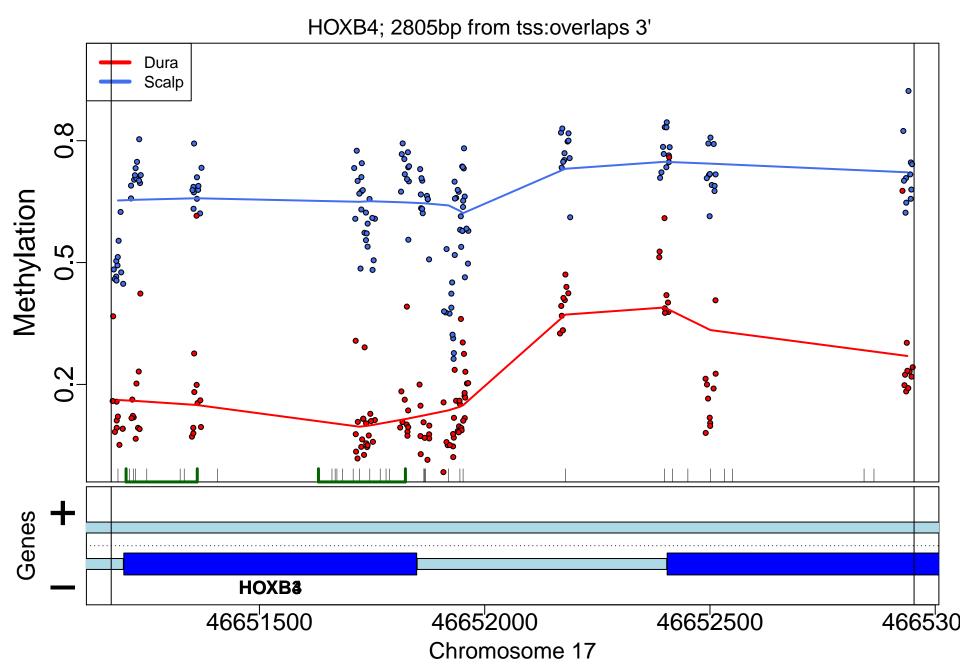




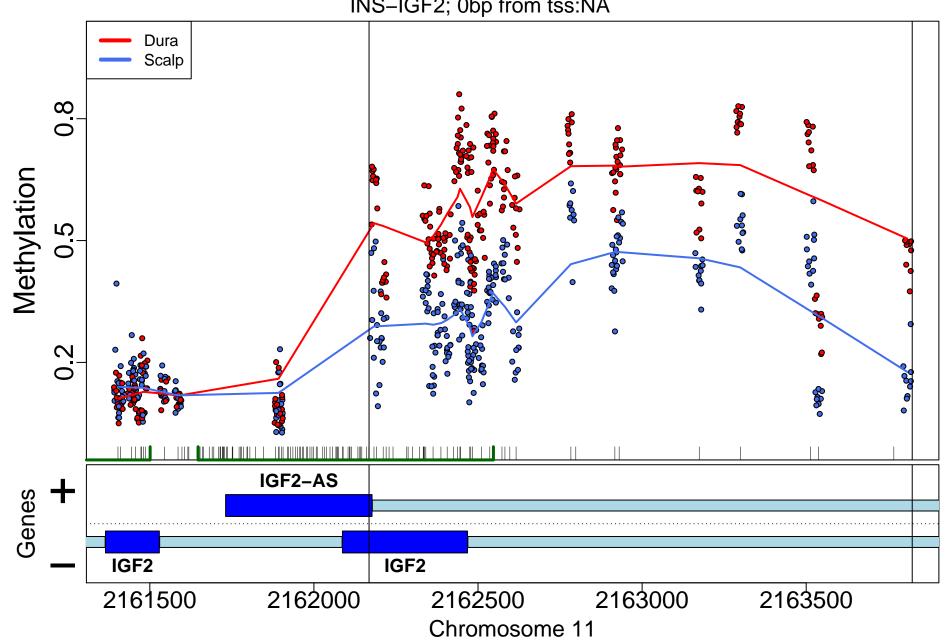


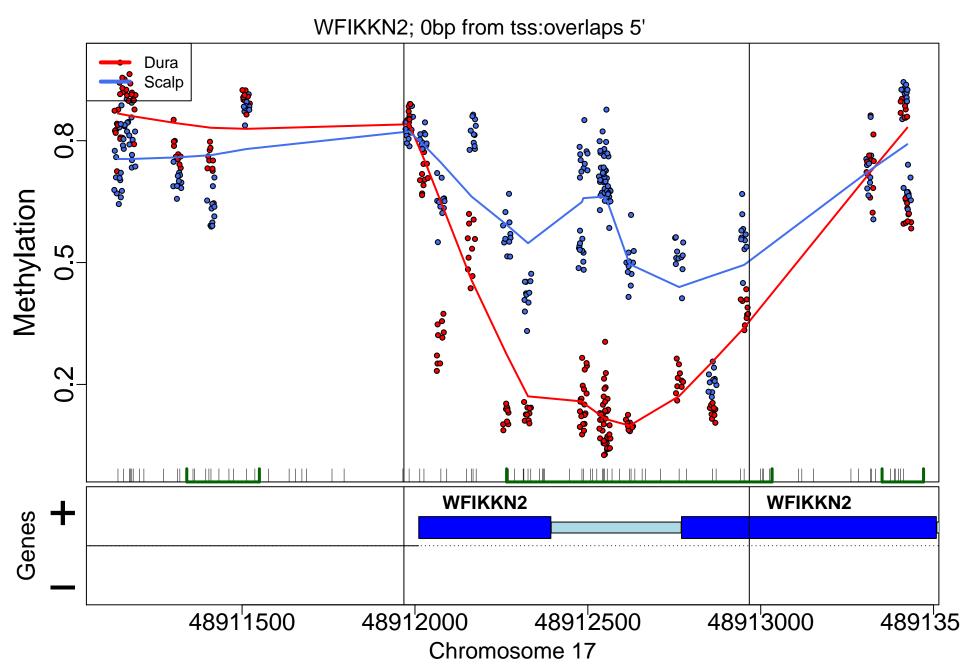


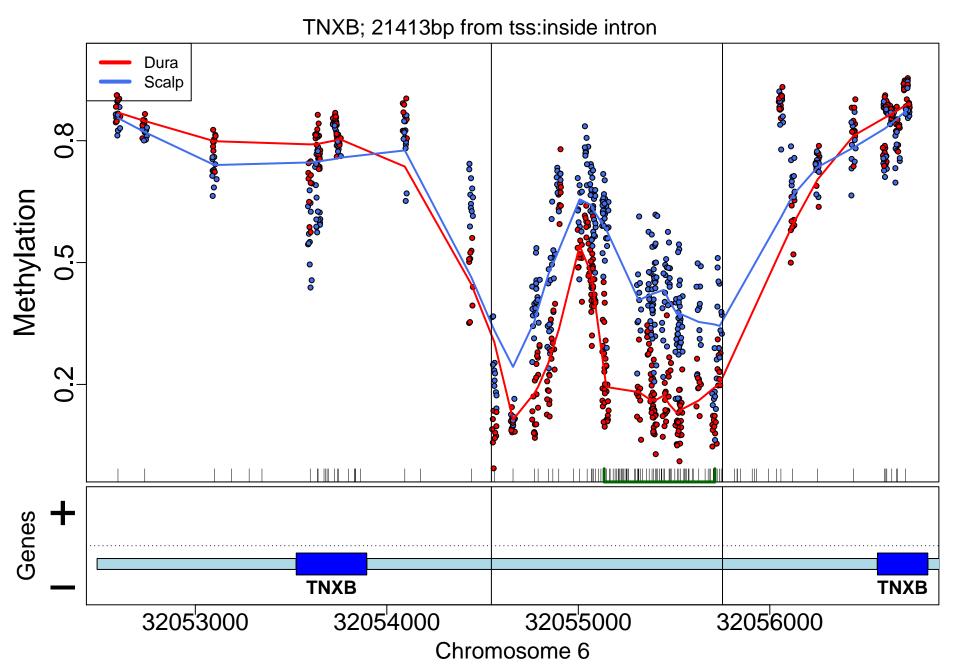




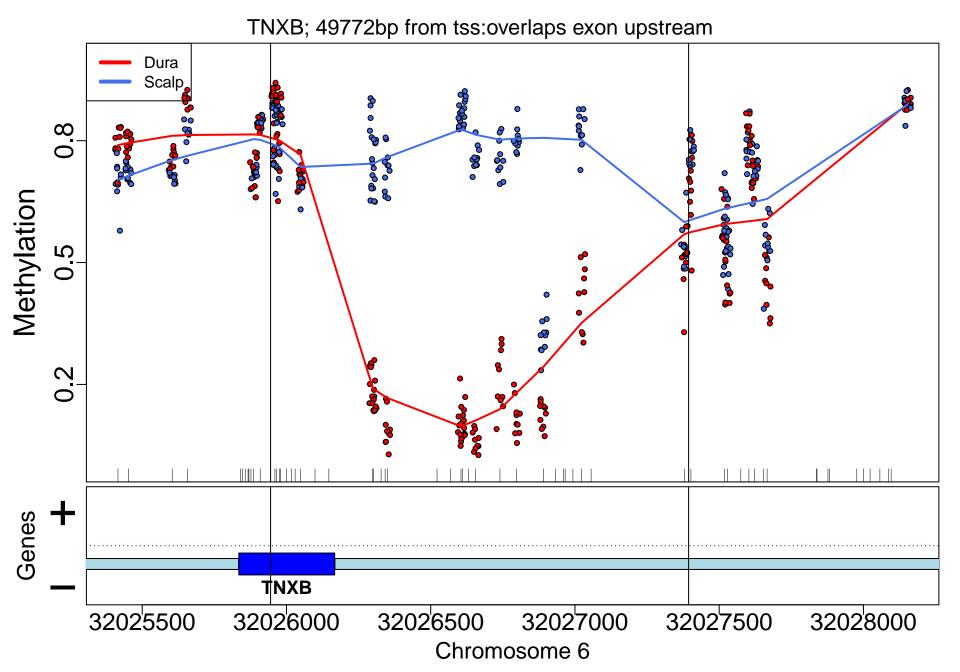
INS-IGF2; 0bp from tss:NA







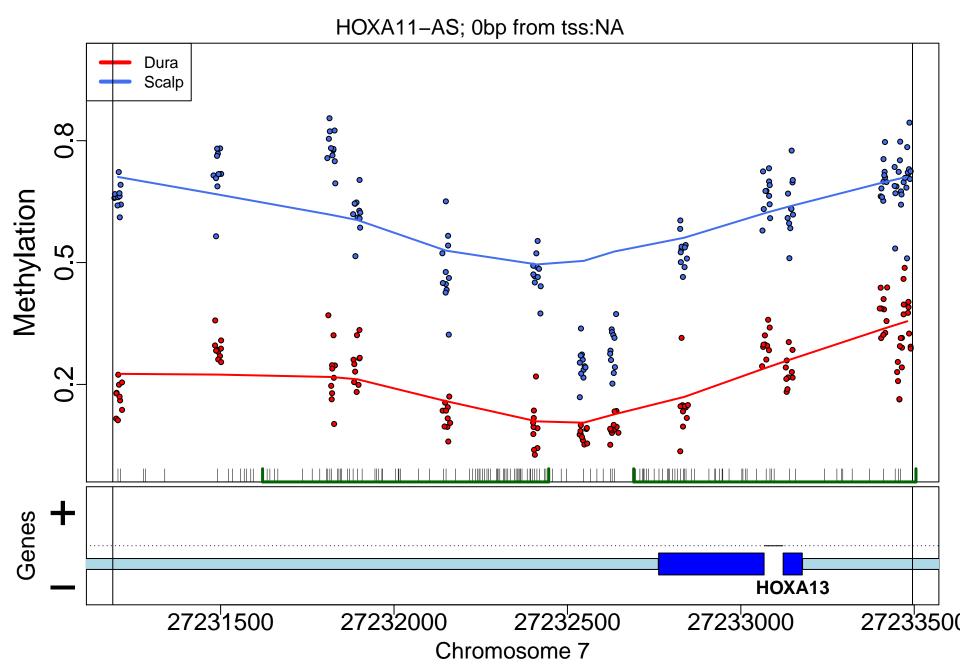
ESR1; 0bp from tss:overlaps 5' Dura Scalp <u>∞</u>. Methylation 0.2 ESR1 ESR1 ESR1 Genes 152128500 152129500 152130000 152128000 152129000 Chromosome 6

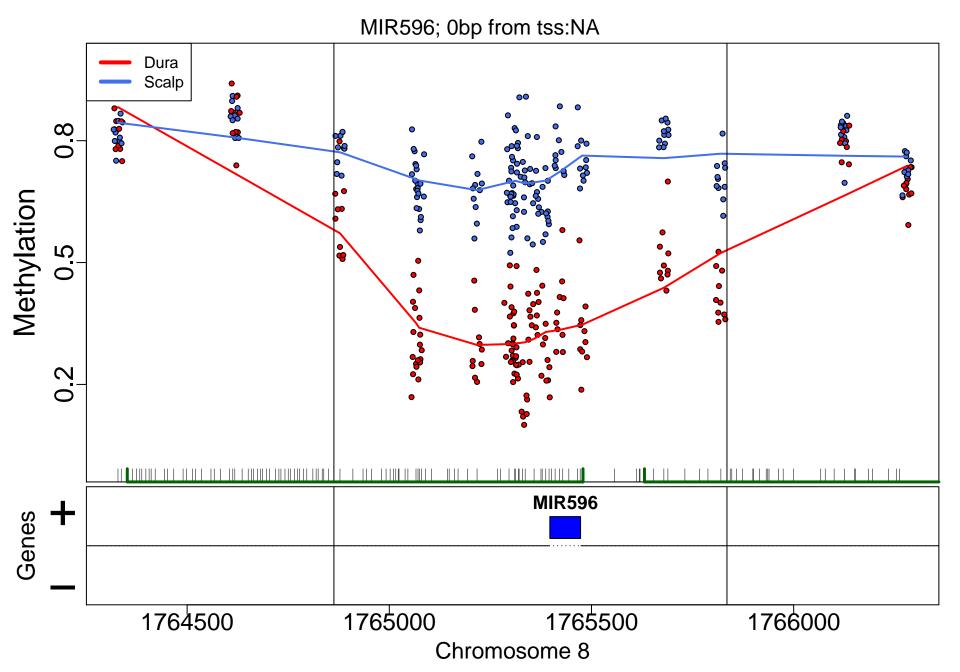


MIR10B; 0bp from tss:NA Dura Scap **%** <u>ω</u>_ Methylation Genes 177012400 177012800 177012200 177012600 177013000 Chromosome 2

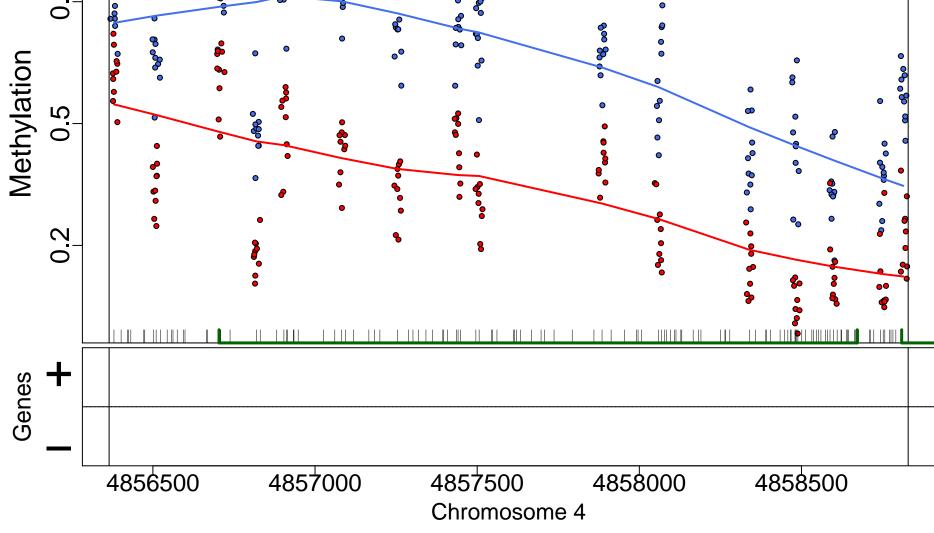
NBEA; 0bp from tss:overlaps 5' Dura Scalp <u>ω</u>_ Methylation 0.2 **NBEA** Genes MAB21L1 36049000 36050500 3605 1000 36049500 36050000 Chromosome 13

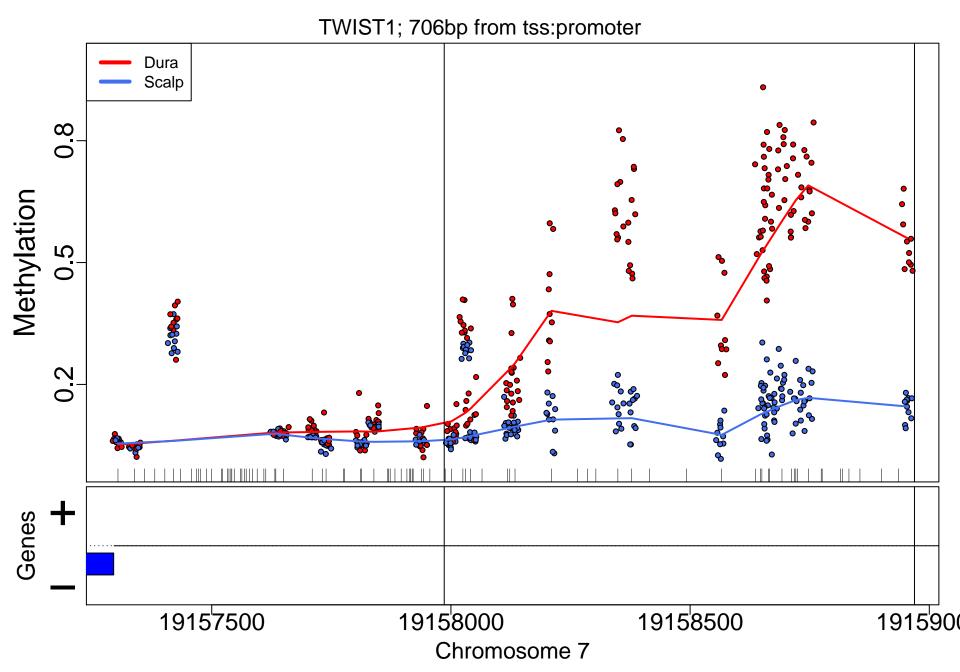
SIX1; 5361bp from tss:upstream Dura Scalp <u>ω</u>_ Methylation Genes SIX1 61122500 6112 1500 61123500 61124500 Chromosome 14





MSX1; 2578bp from tss:upstream Dura Scalp <u>ω</u>_ 0.5 0.2





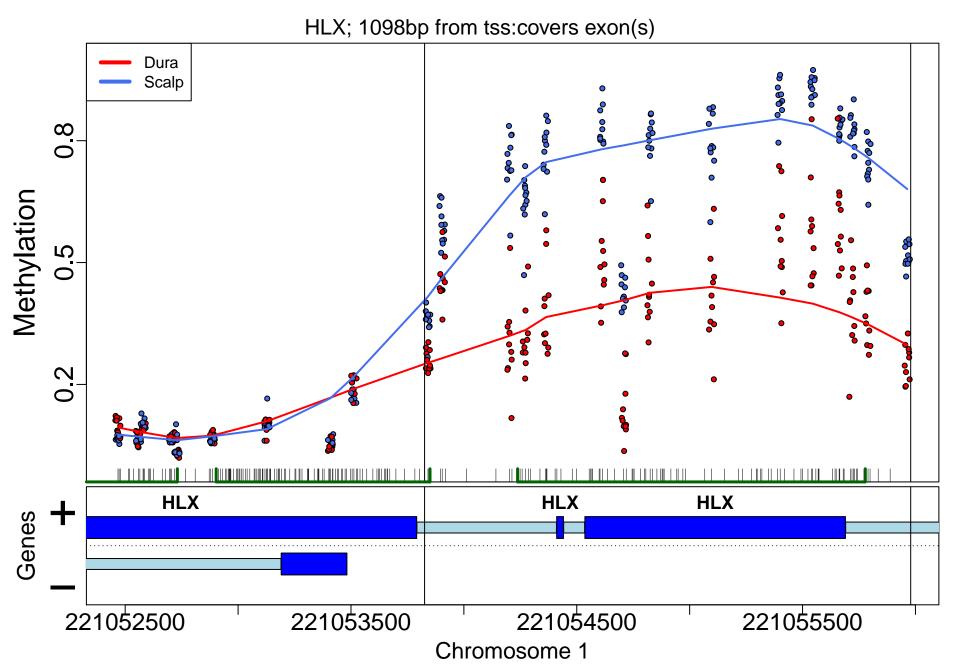
LINC00311; 0bp from tss:NA Dura Scalp <u>∞</u>. Methylation 0.2 Genes 85320000 85320200 85320600 85320800 85320400 Chromosome 16

ZIC2; 6888bp from tss:downstream Dura Scalp 0 Methylation 0.2 Genes 100641000 100643000 100642000 100644000 Chromosome 13

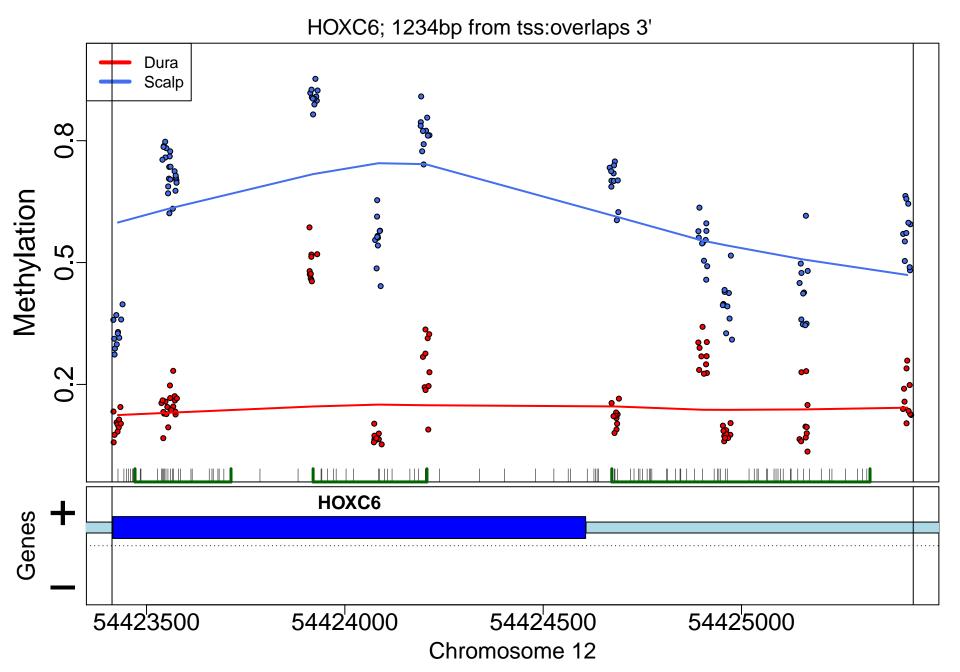
SPON2; 35036bp from tss:inside intron Dura Scalp <u>∞</u>. Methylation 0.2 Genes SPON2 SPON2 1167500 1166500 1167000

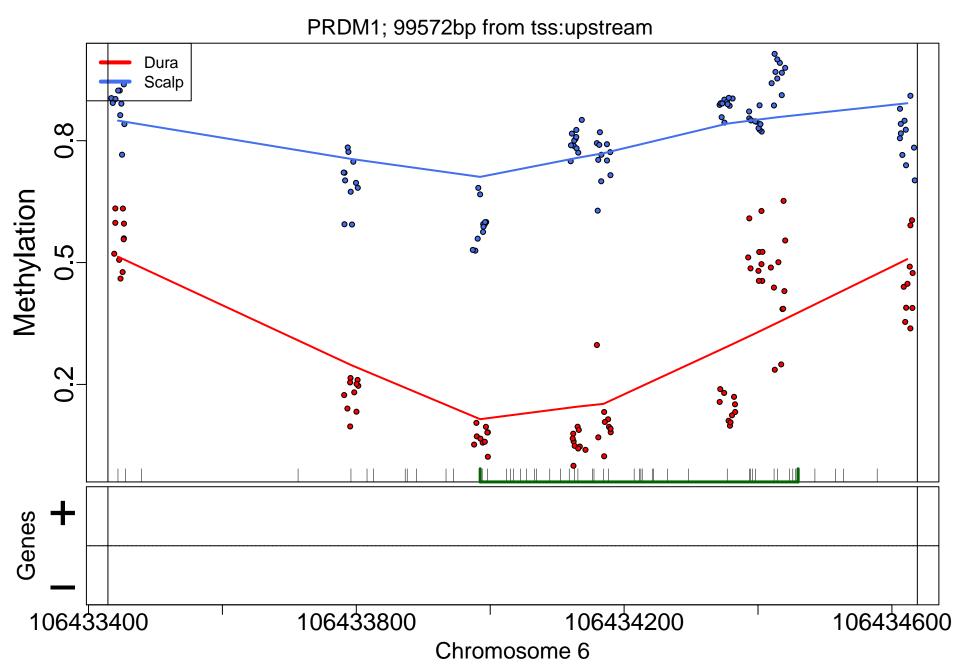
Chromosome 4

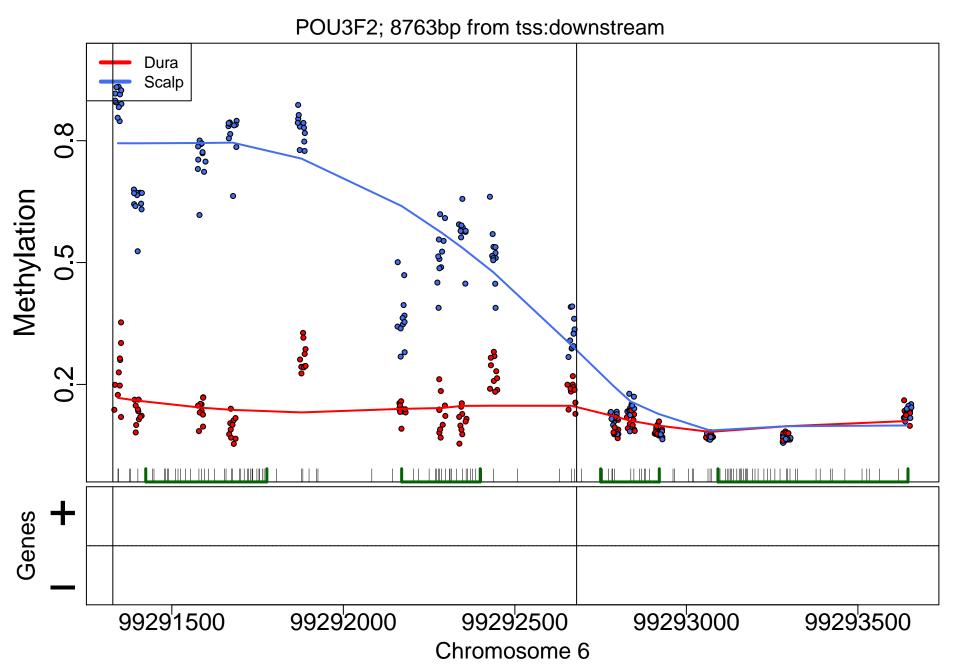
MSX1; 4655bp from tss:downstream Dura Scalp ω_{_} Methylation 0.5 0.2 Genes 4867000 4866000 4867500 4866500 4868000 4868500 4869000 Chromosome 4

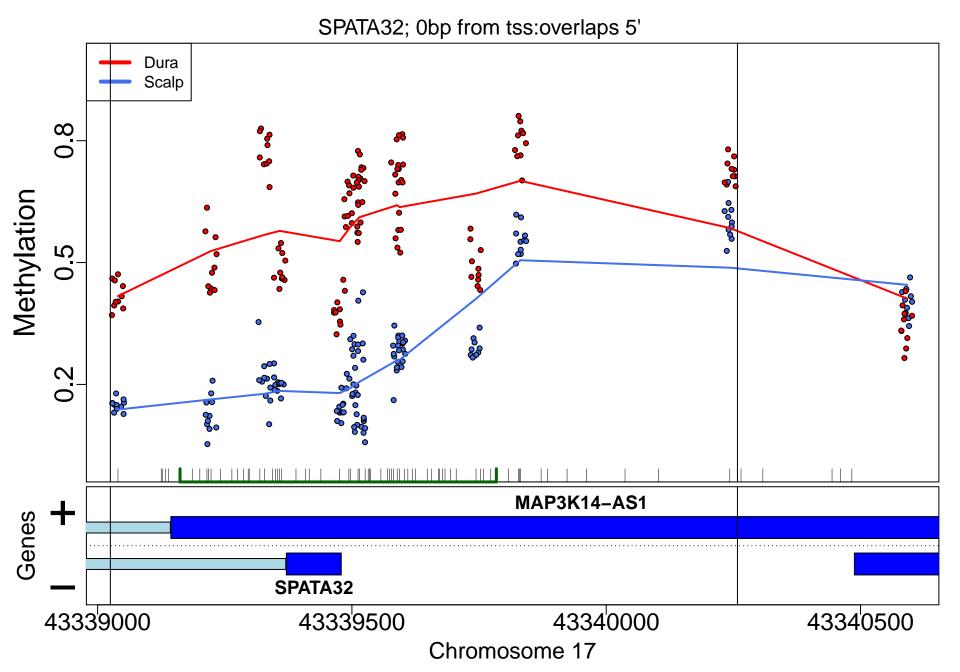


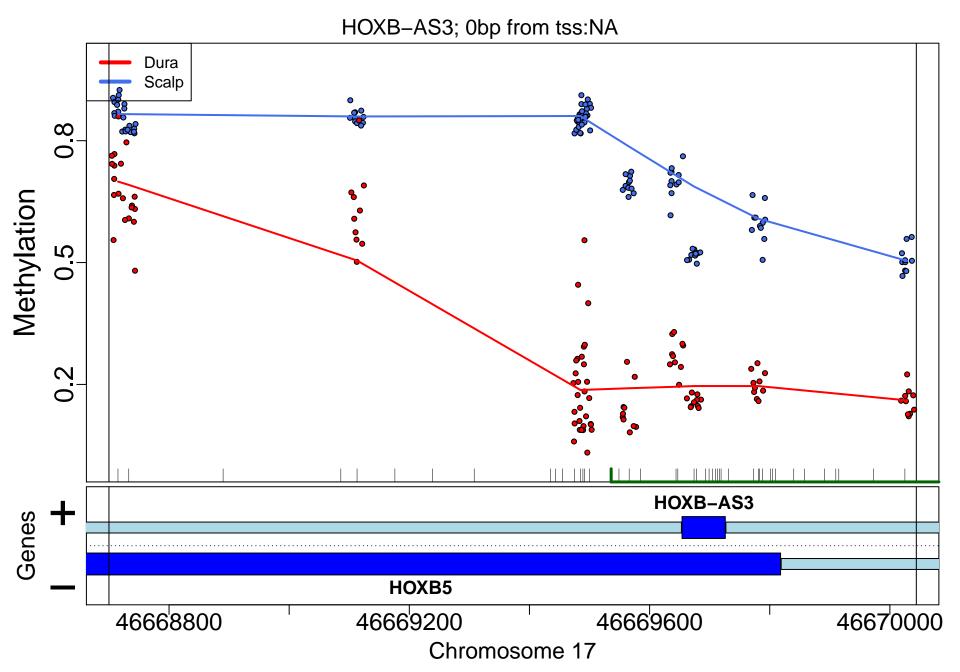
LSP1; 0bp from tss:overlaps 5' Dura Scalp <u>ω</u>_ Methylation 0.2 LSP1 Genes 1892600 1892800 1891800 1892000 1892200 1892400 Chromosome 11

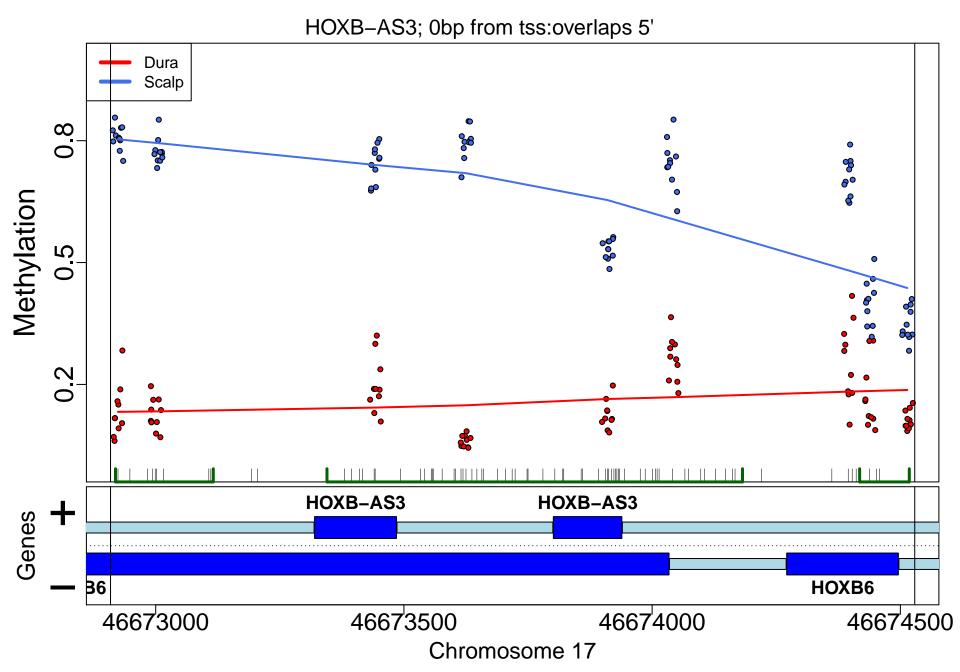


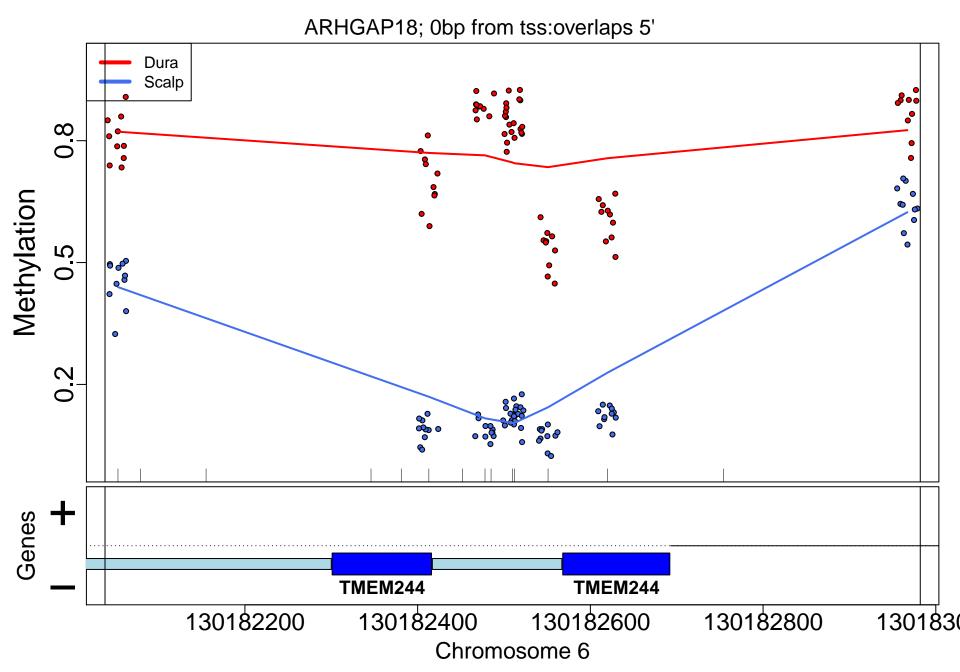


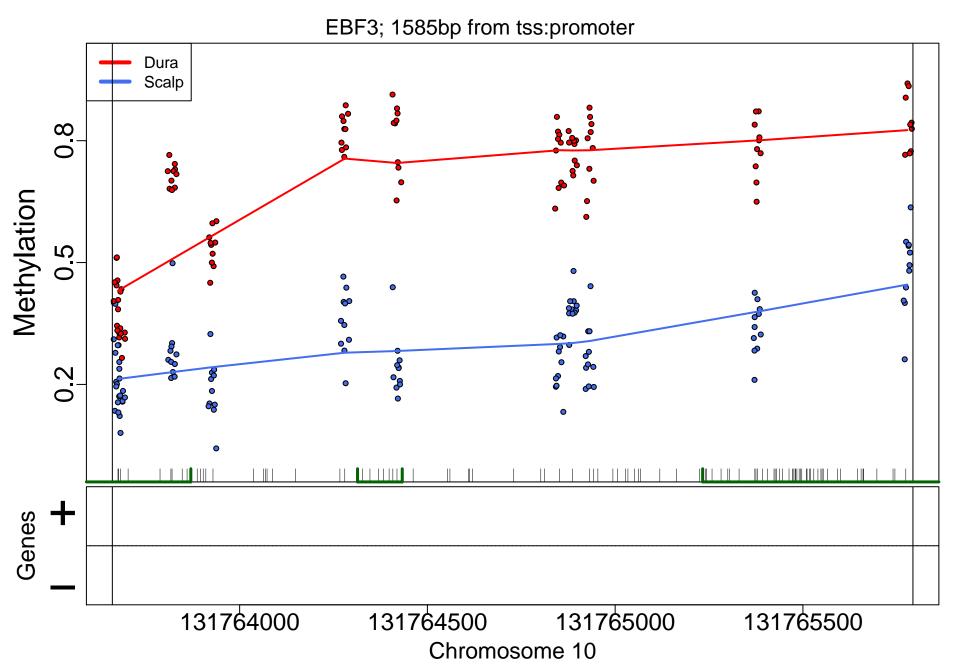


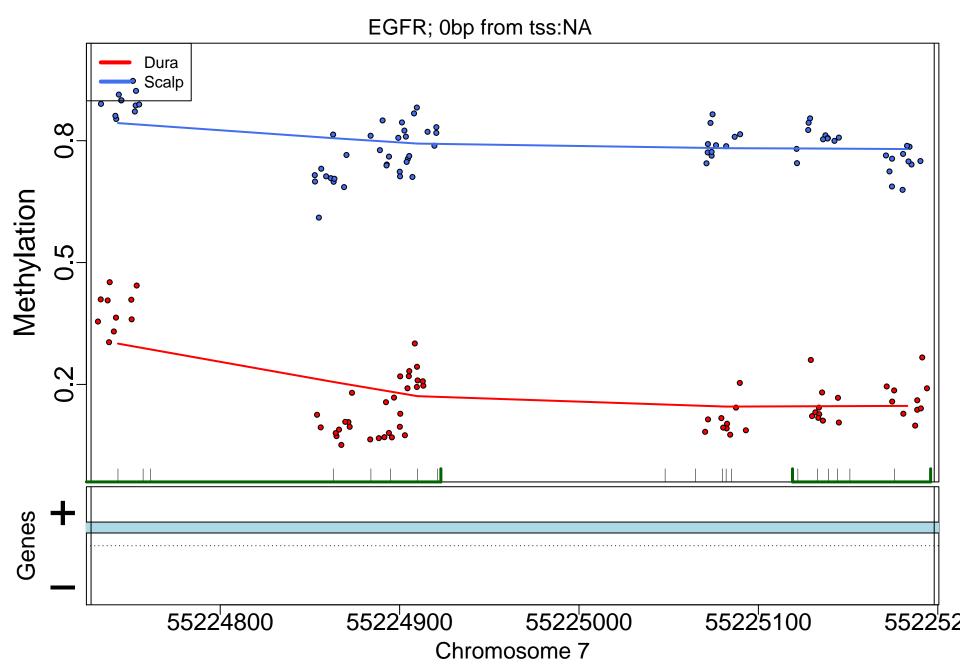




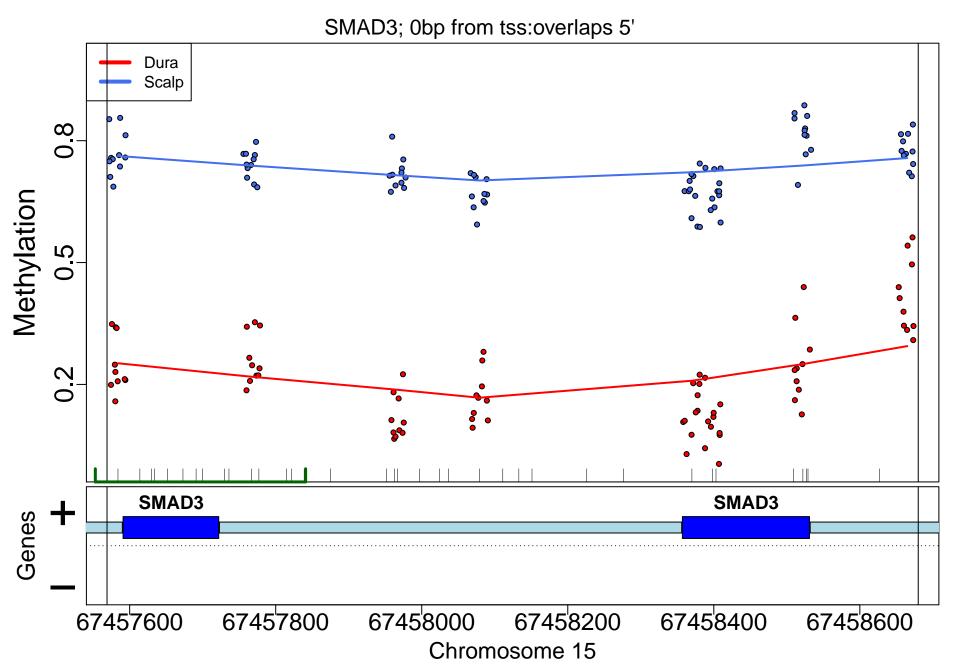


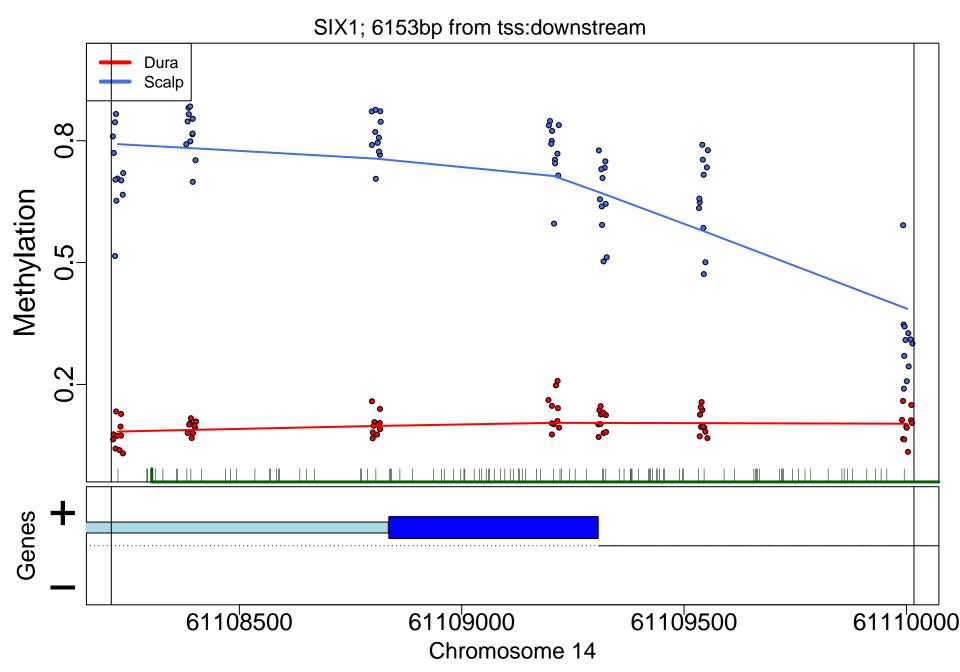


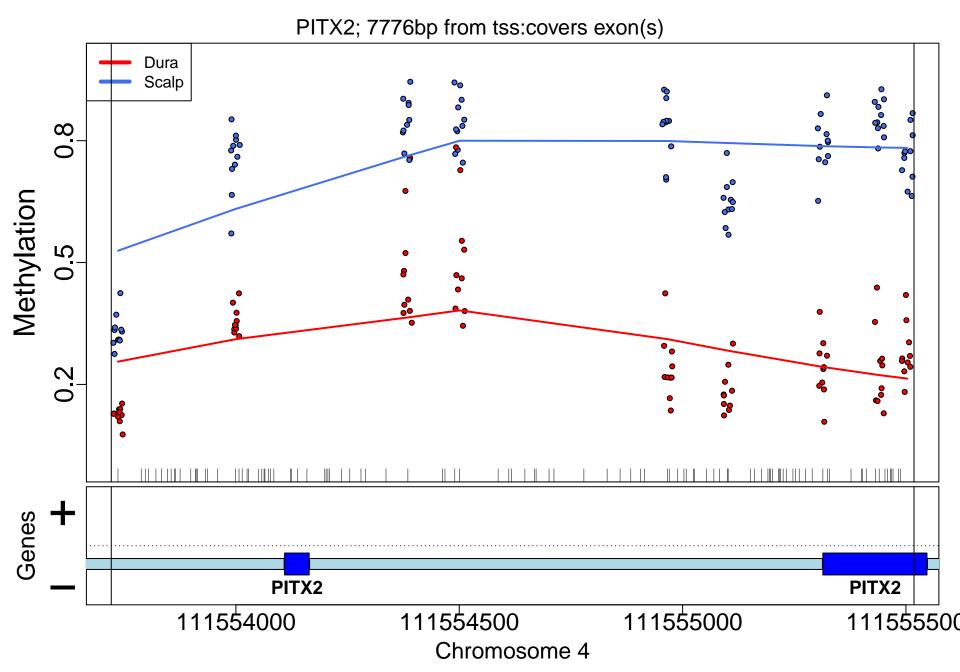


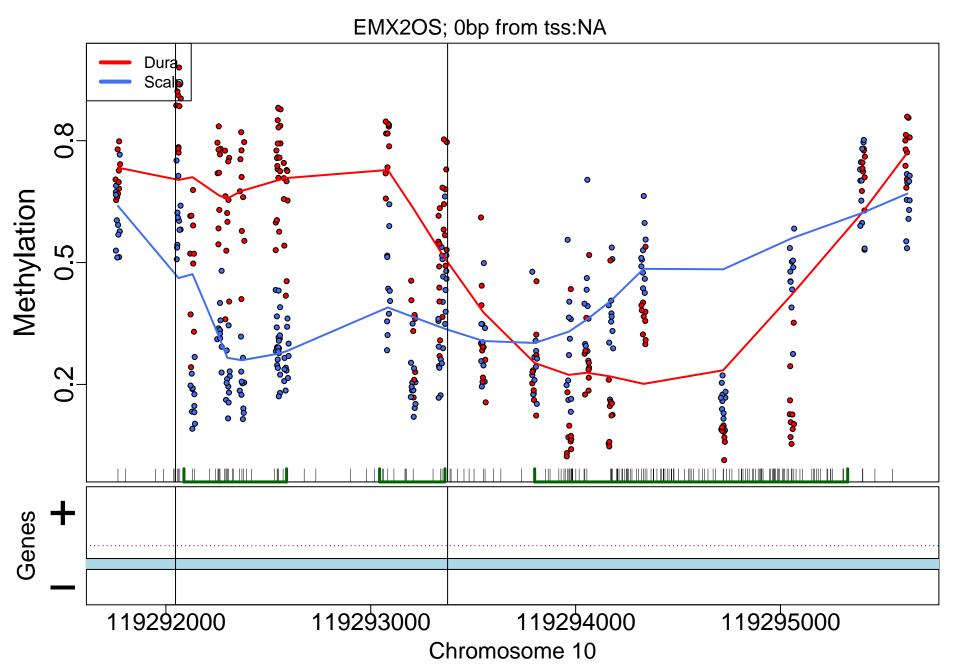


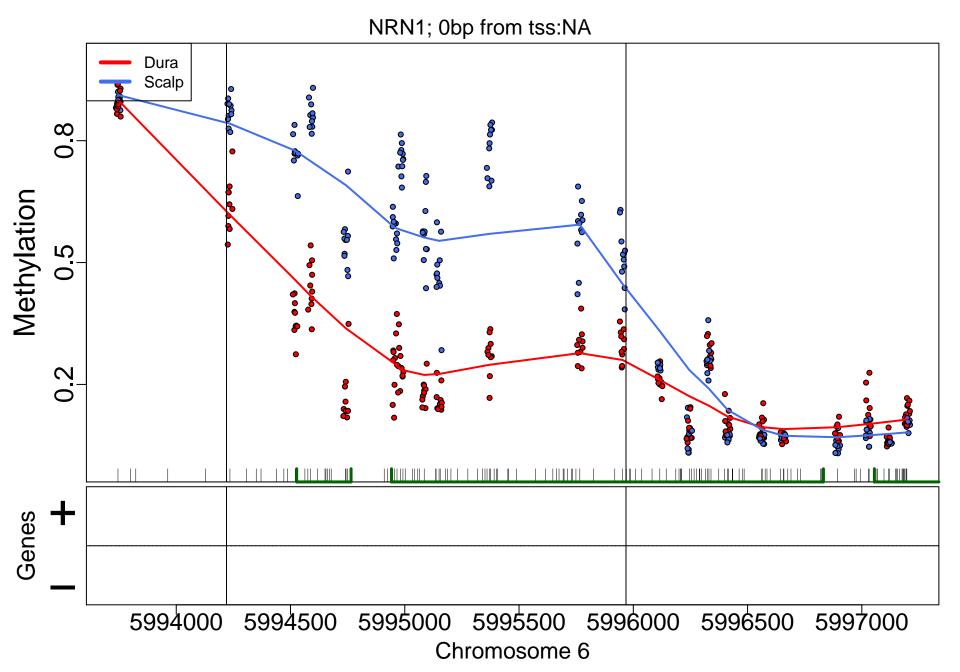
EN1; 8730bp from tss:upstream Dura Scalp 0 。 。 Methylation 0.2 Genes 119614500 119615000 119615500 119616000 119616500 119617000 Chromosome 2



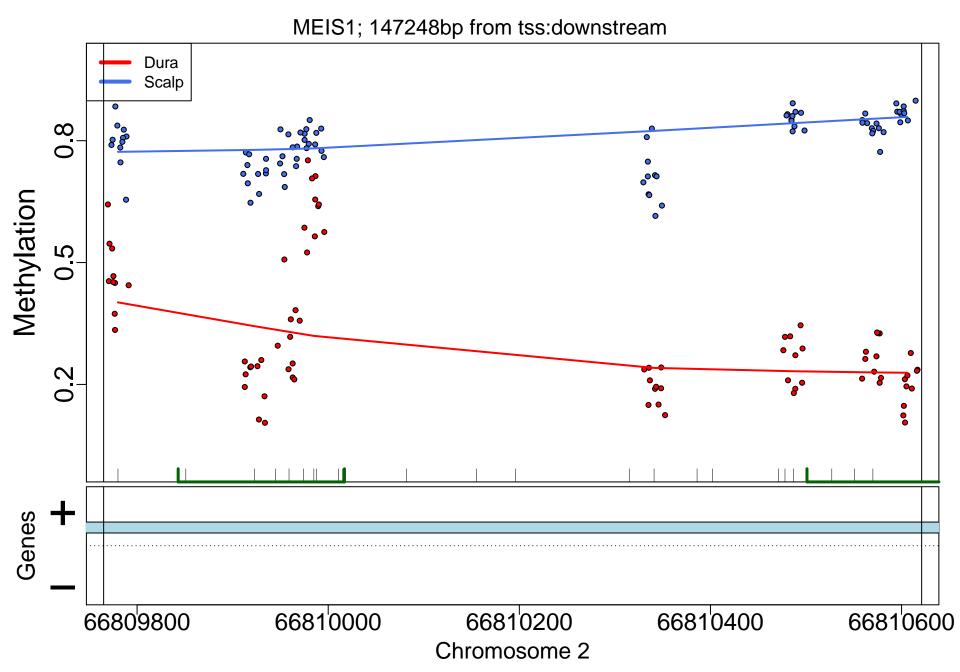


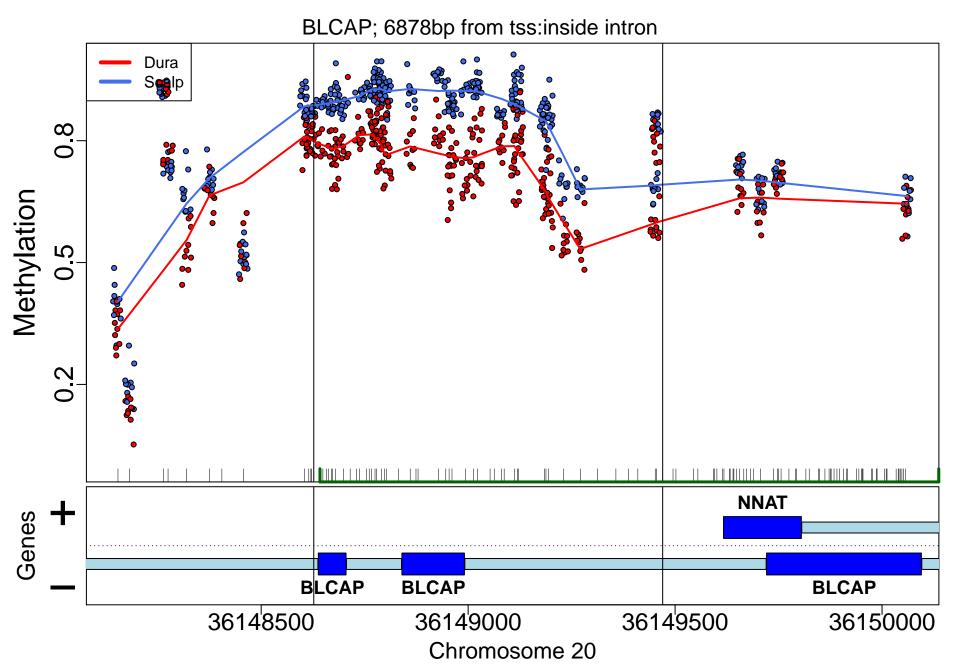


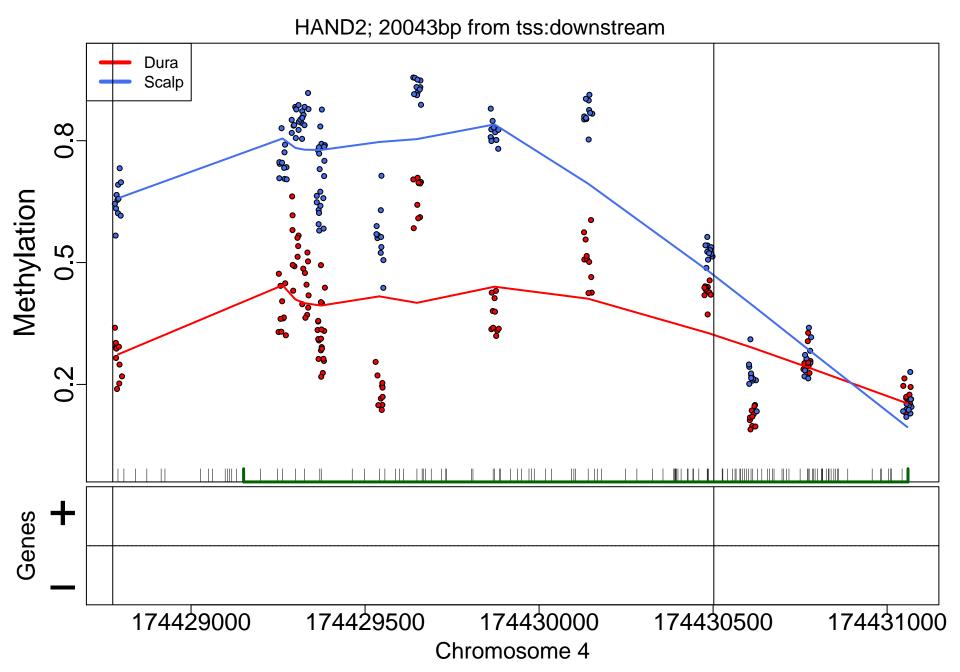


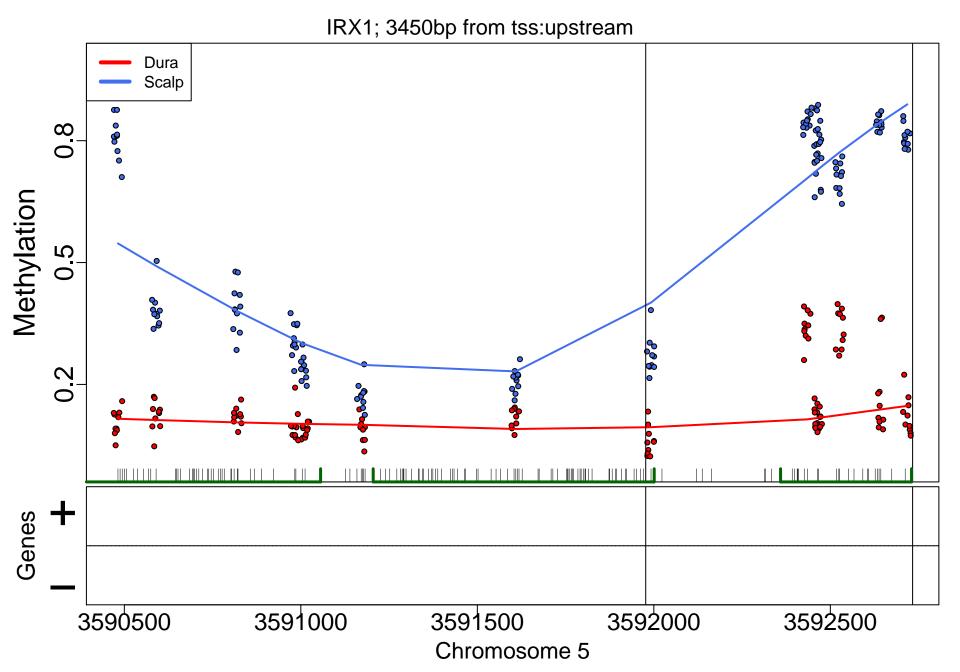


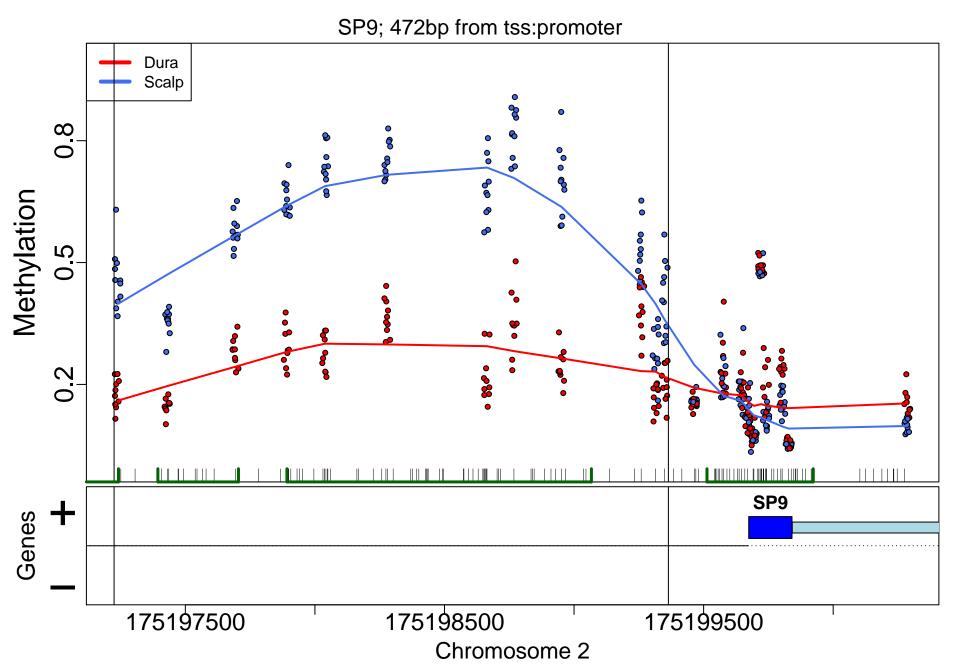
SIM1; 4111bp from tss:upstream Dura Scalp <u>∞</u>_ Methylation 0.2 Genes 100917000 100917200 100917400 100917600 100917800 Chromosome 6

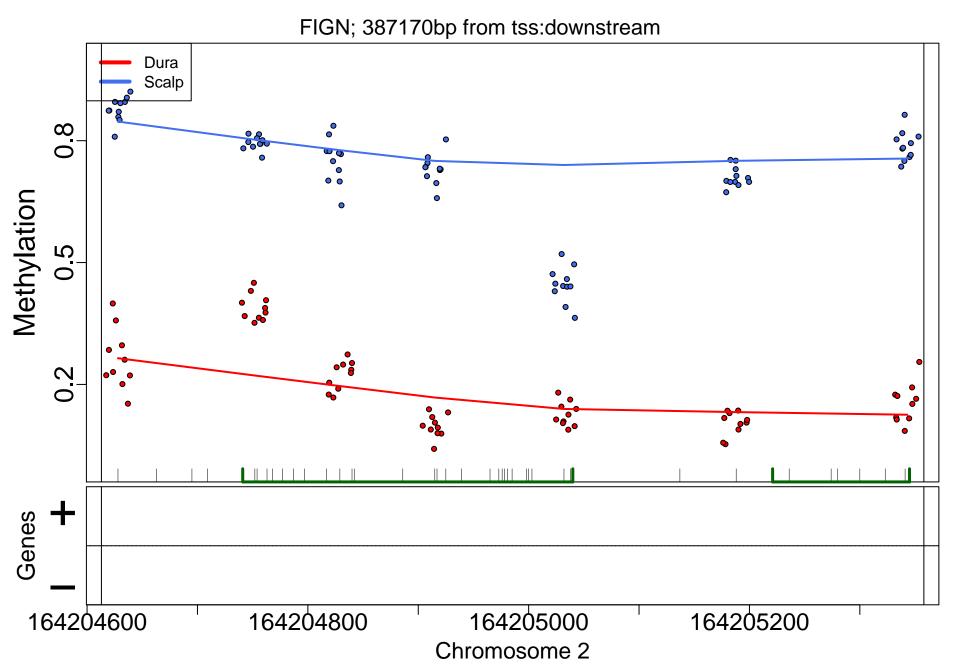








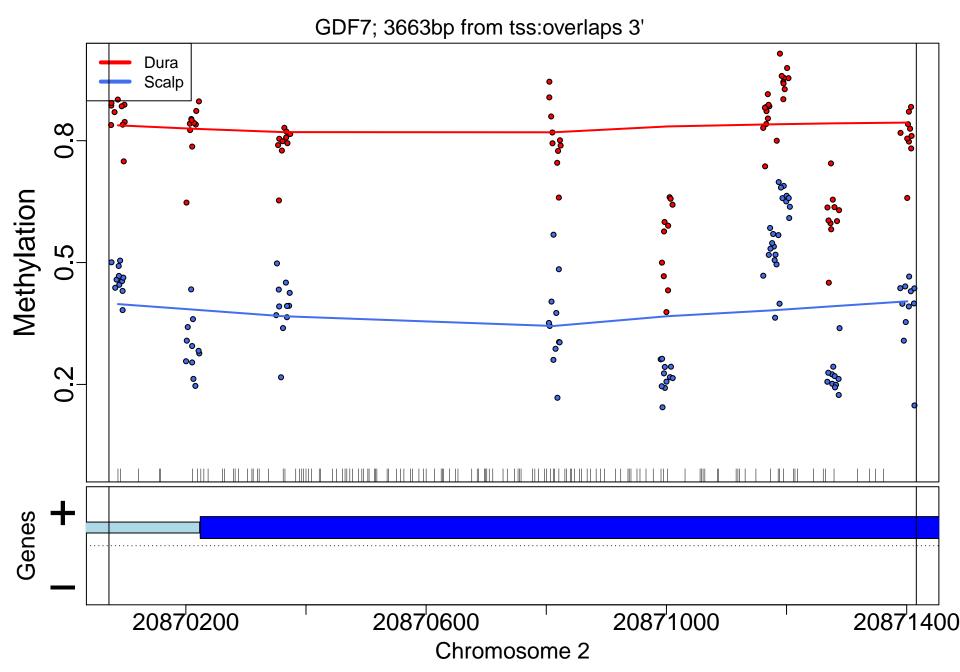


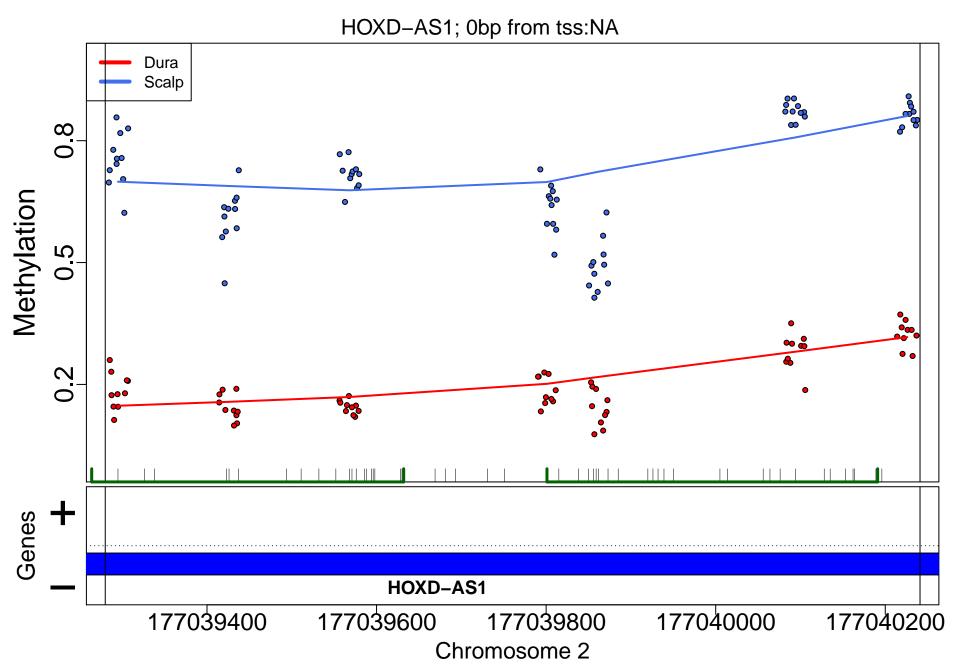


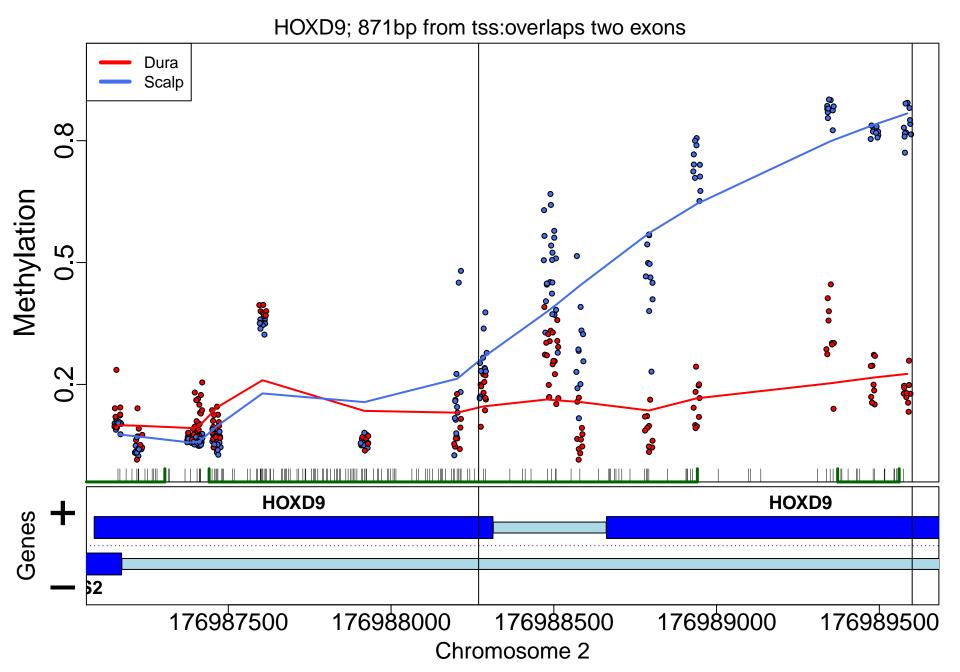
HLX; 15153bp from tss:downstream Dura Scalp 80. Methylation 0.2 Genes 221068000 221069500 221070000 221068500 221069000

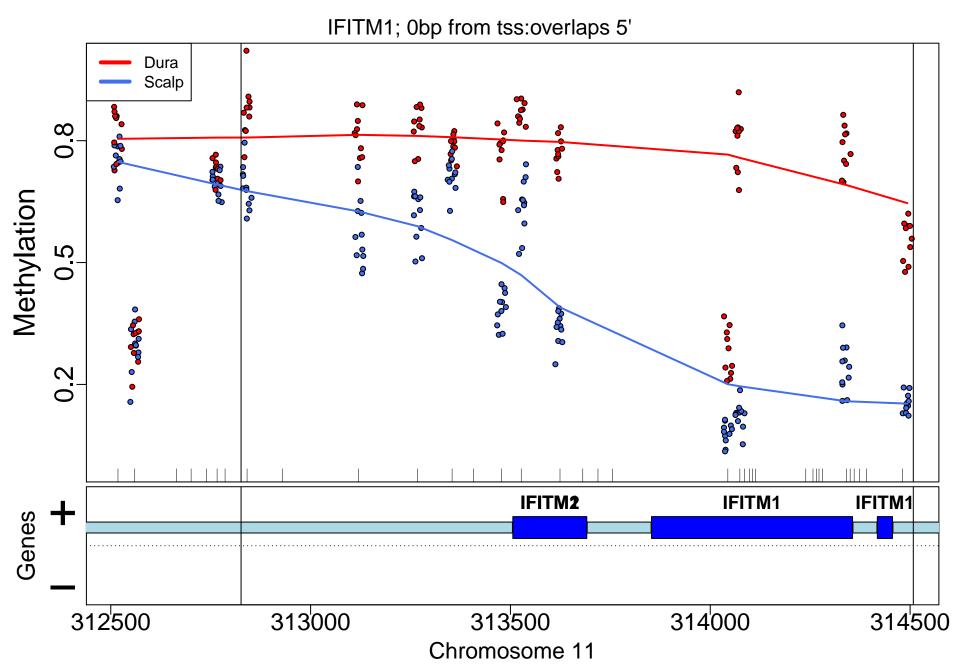
Chromosome 1

SFRP2; 733bp from tss:promoter Dura Scalp 80. Methylation 0.2 Genes 154711000 154713000 154712000 154714000 Chromosome 4





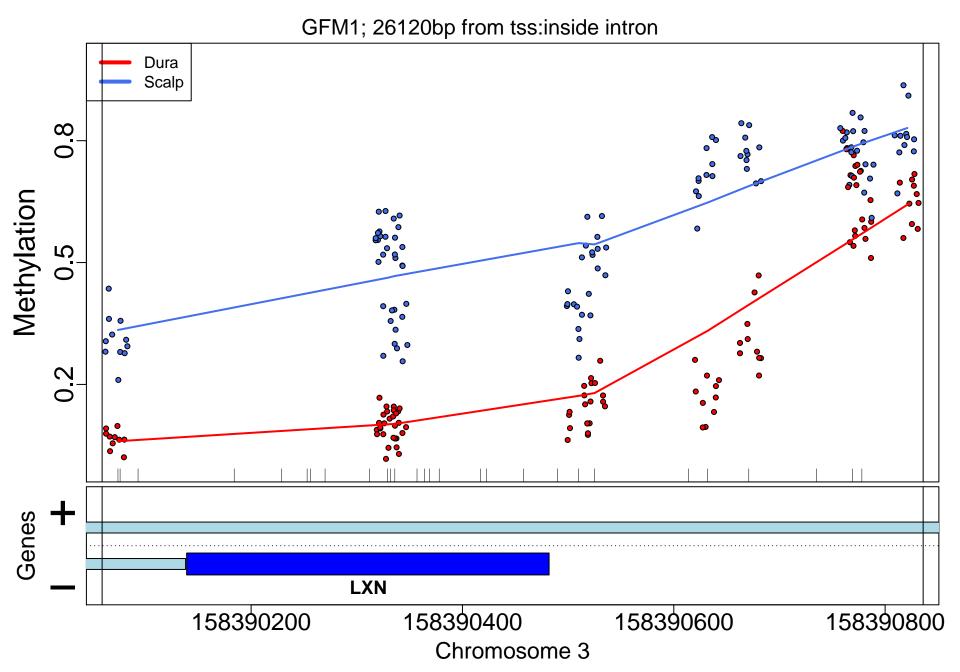


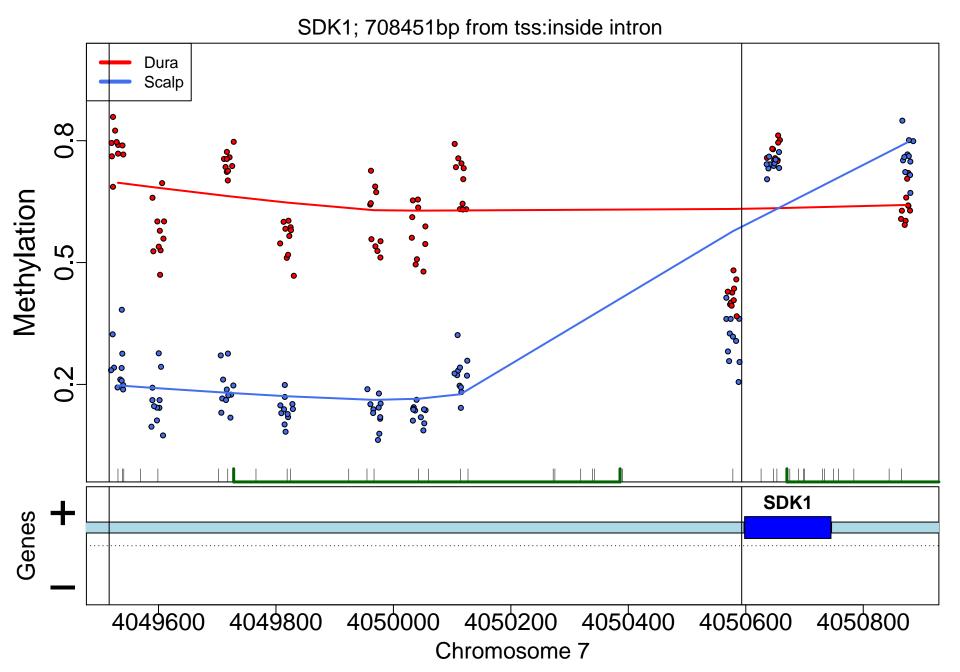


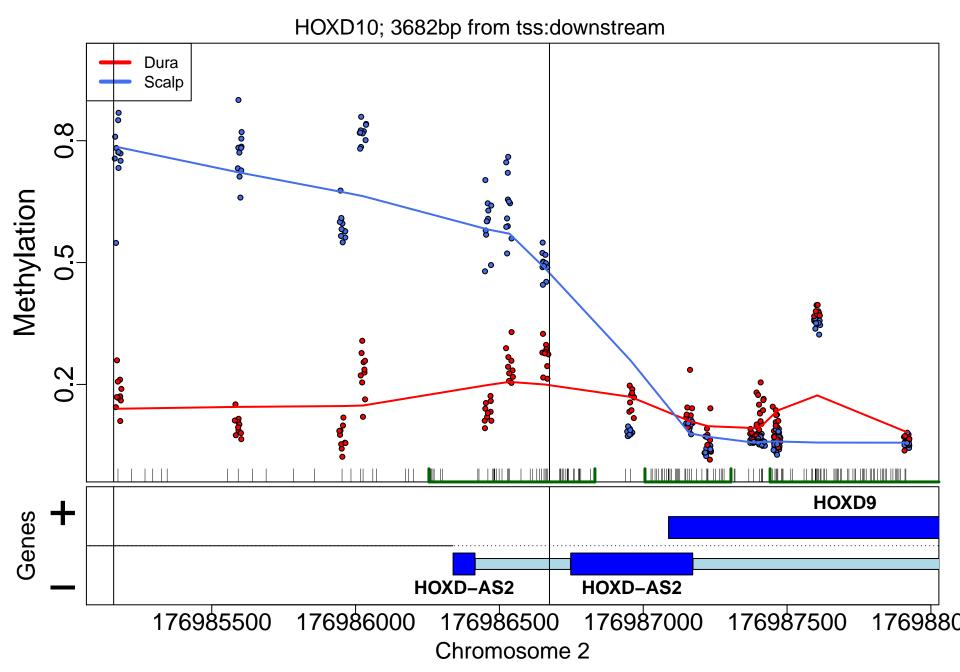
SLC9A3; 21629bp from tss:inside intron Dura Scalp <u>∞</u>. Methylation Genes 502200 502600 502800 502400 502000 Chromosome 5

RGS22; 0bp from tss:NA Dura Scalp <u>∞</u>. Methylation 0.2 Genes RGS22 RGS22 101118000 101118800 101118200 101118400 101118600 Chromosome 8

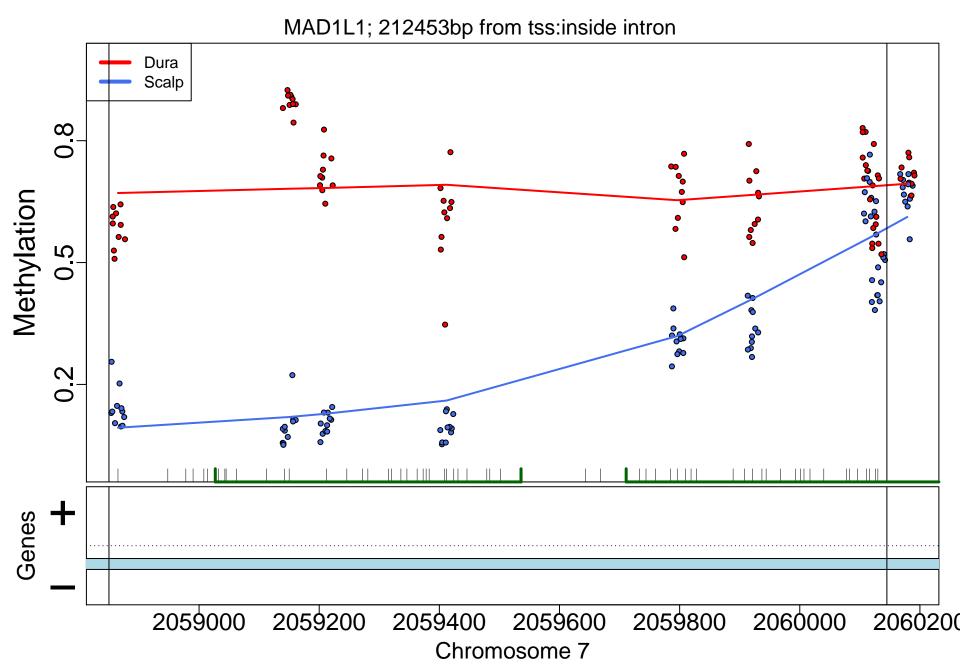
SIX6; 1413bp from tss:promoter Dura Scalp <u>∞</u>. Methylation Genes 60973600 60973800 60974200 60974000 60974400 Chromosome 14

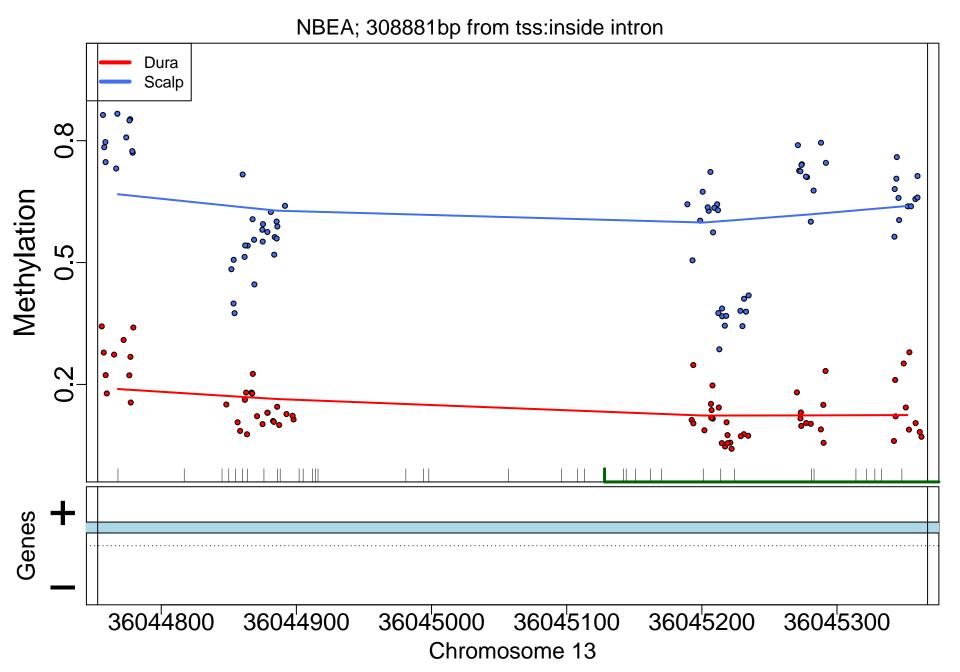


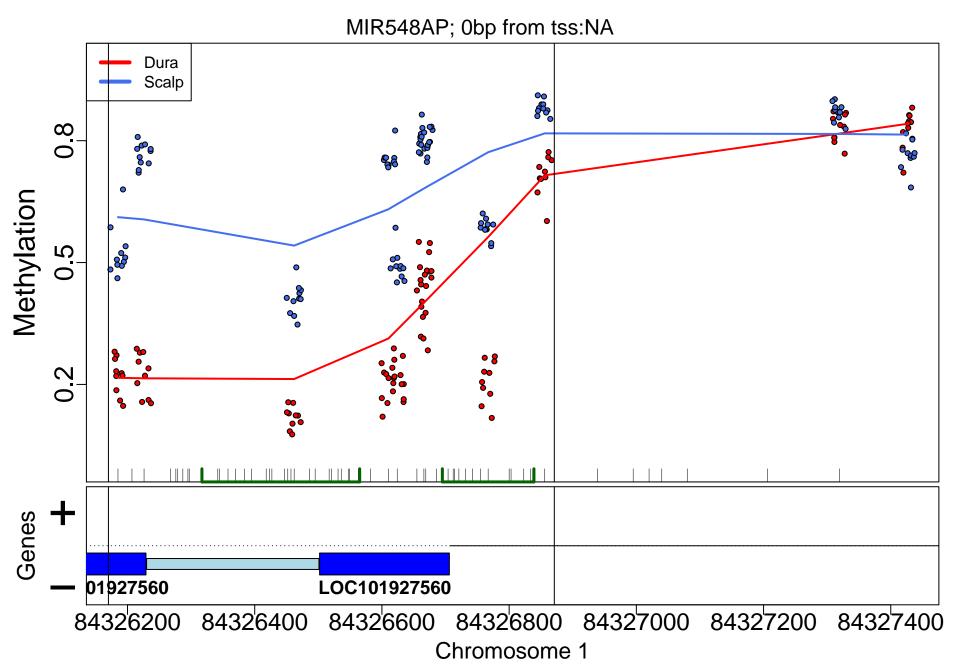




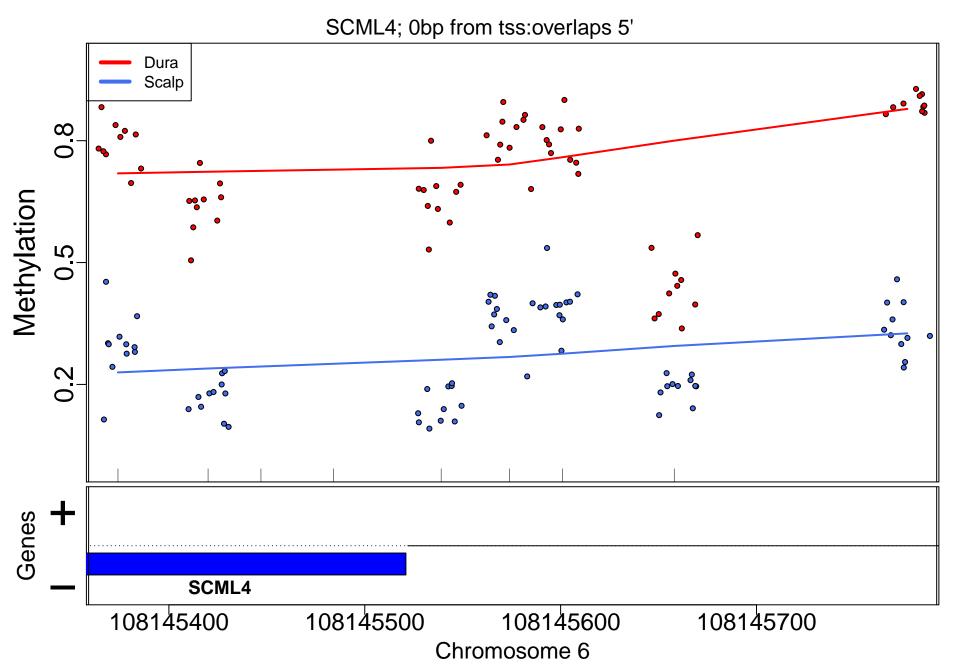
SIX2; 3654bp from tss:overlaps 3' Dura Scalp <u>∞</u>. Methylation 0.5 0.2 Genes SI 4523 1500 45232500 45232000 Chromosome 2

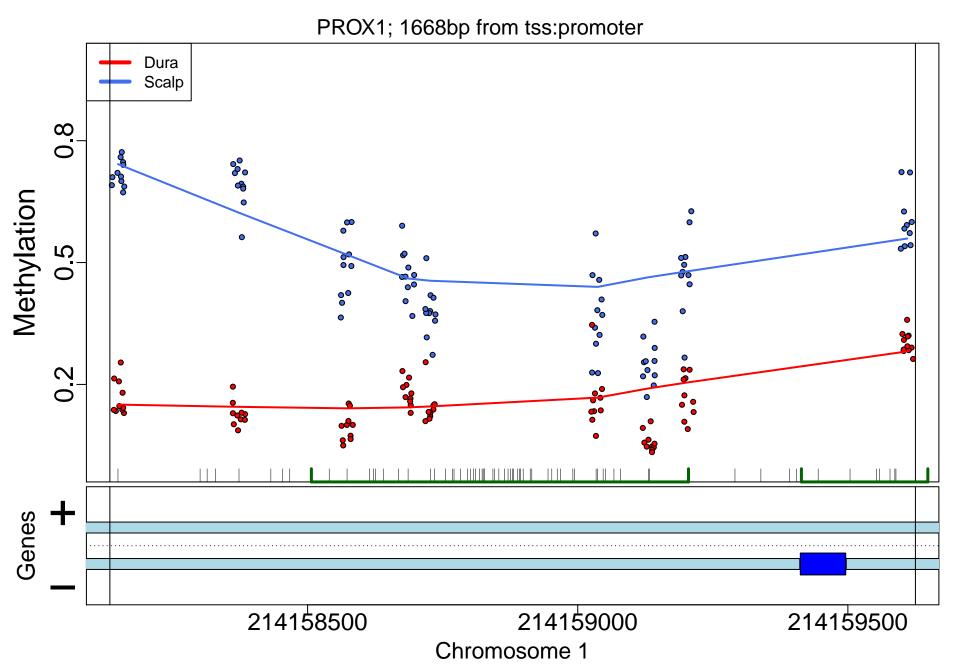


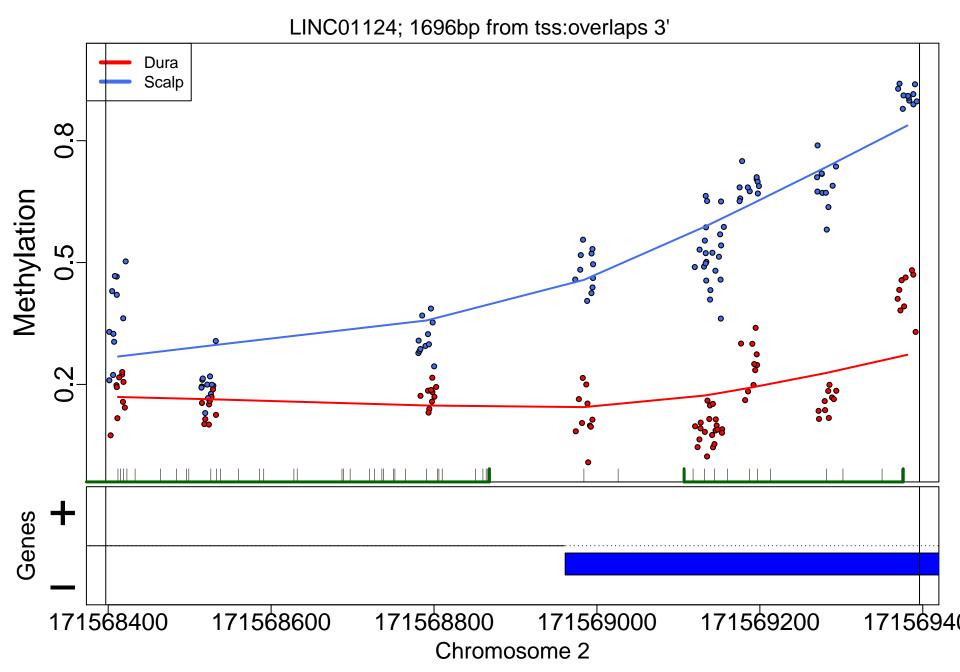


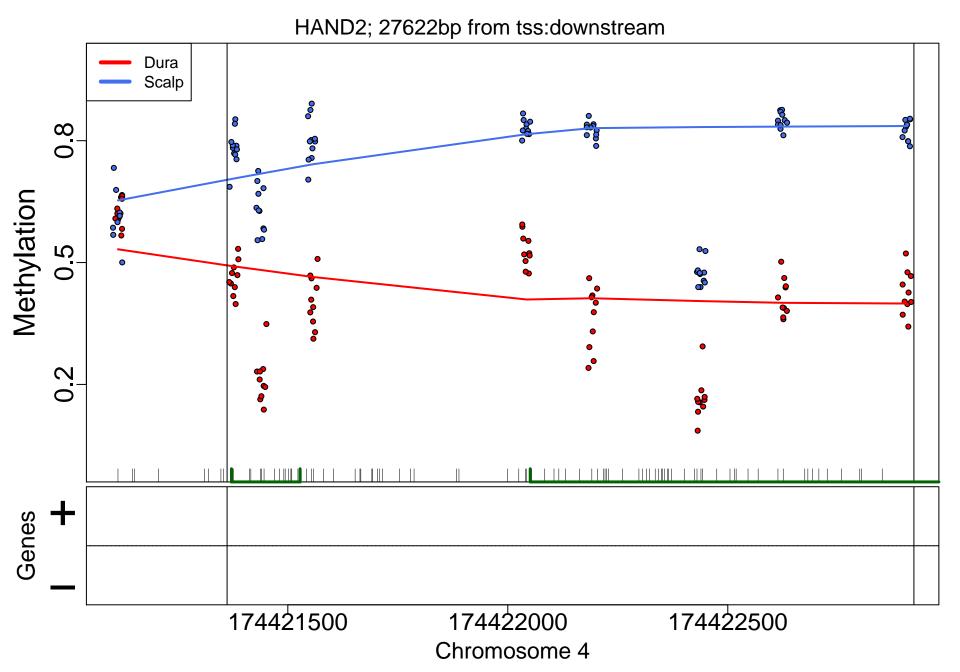


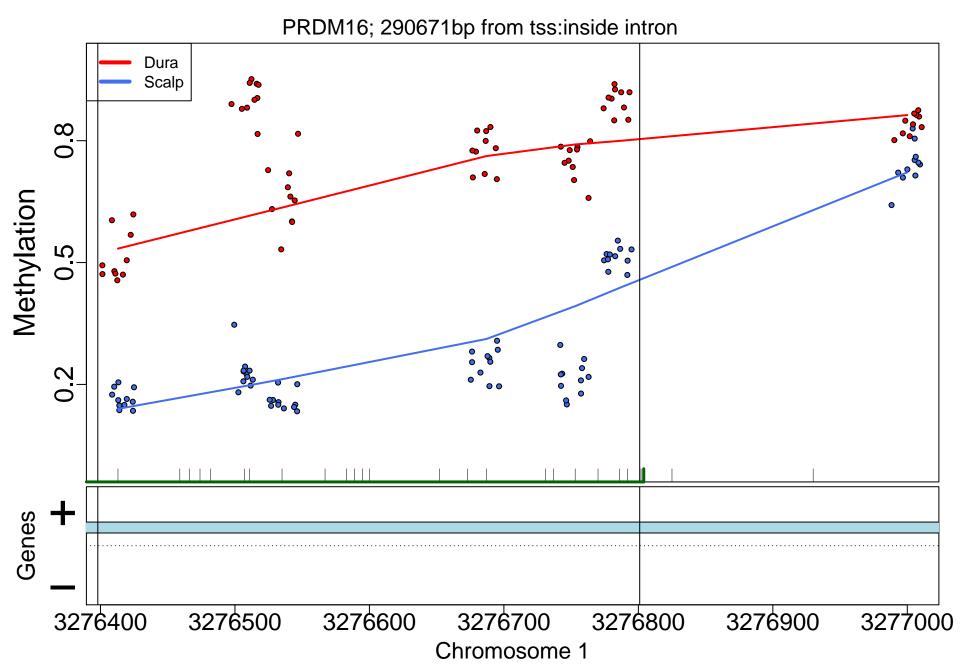
ZFHX3; 4308bp from tss:upstream Dura Scalp <u>∞</u>_ Methylation 0.5 0.2 Genes 73096900 73097000 73097200 73097300 73097100 Chromosome 16

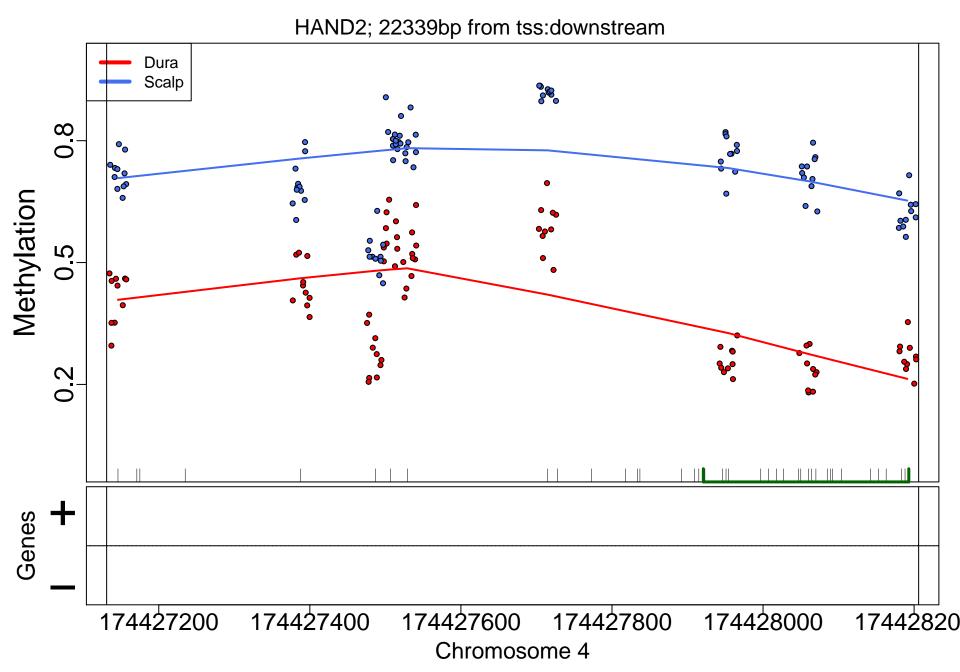


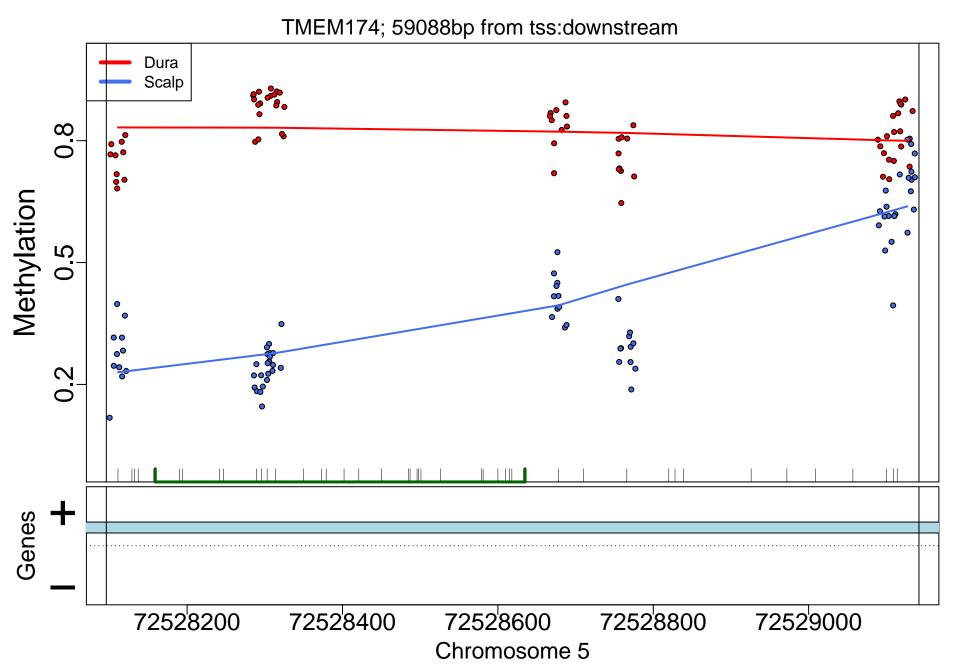


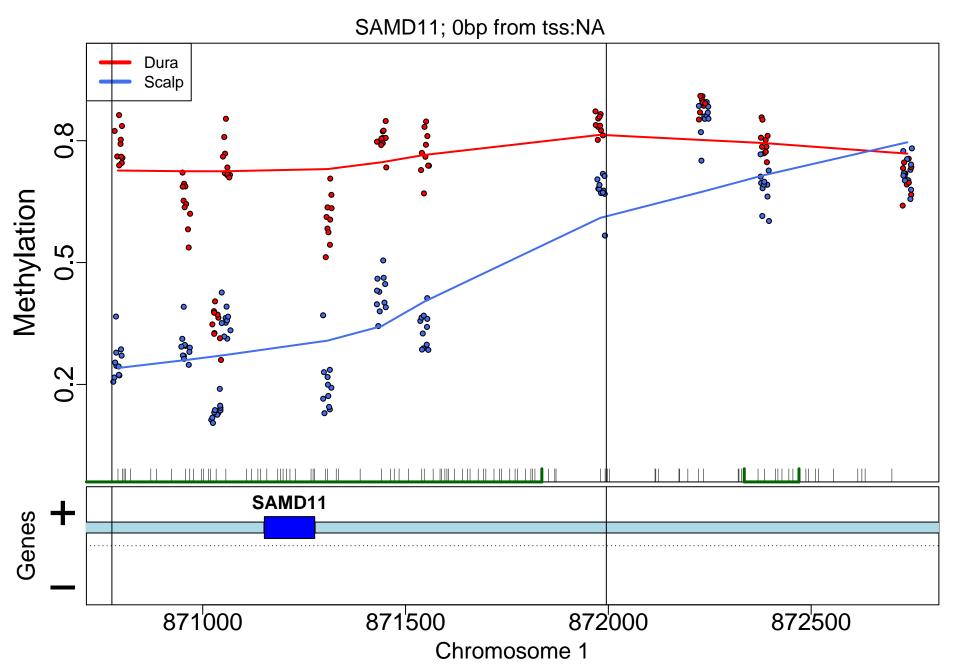




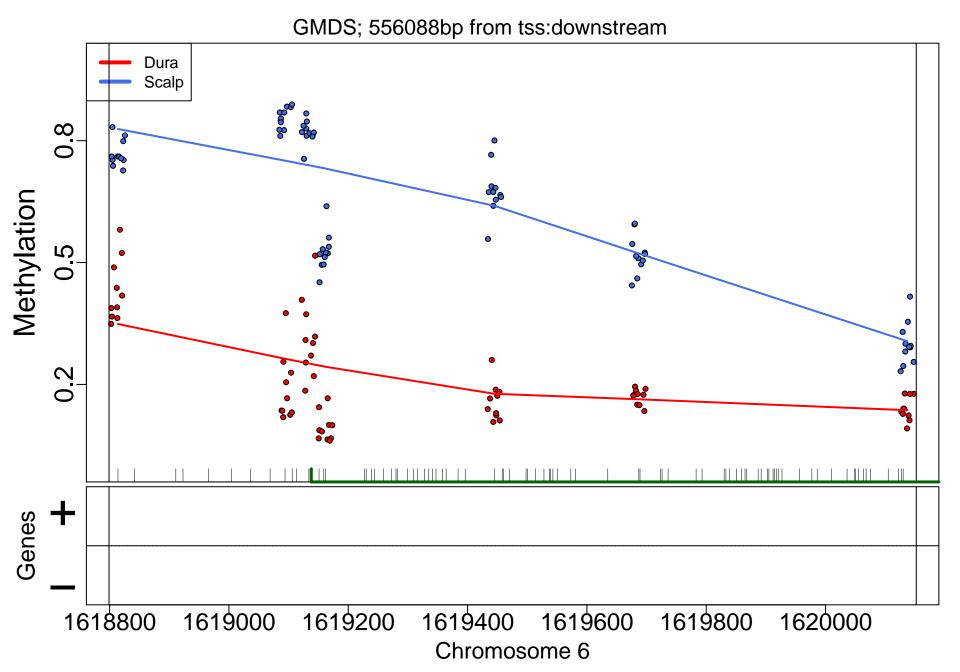


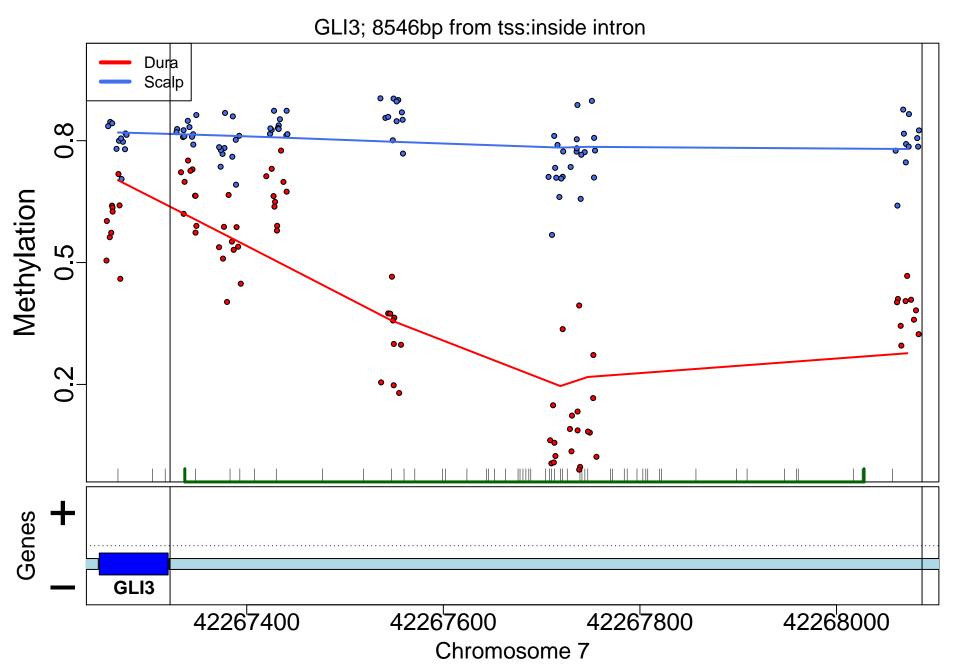


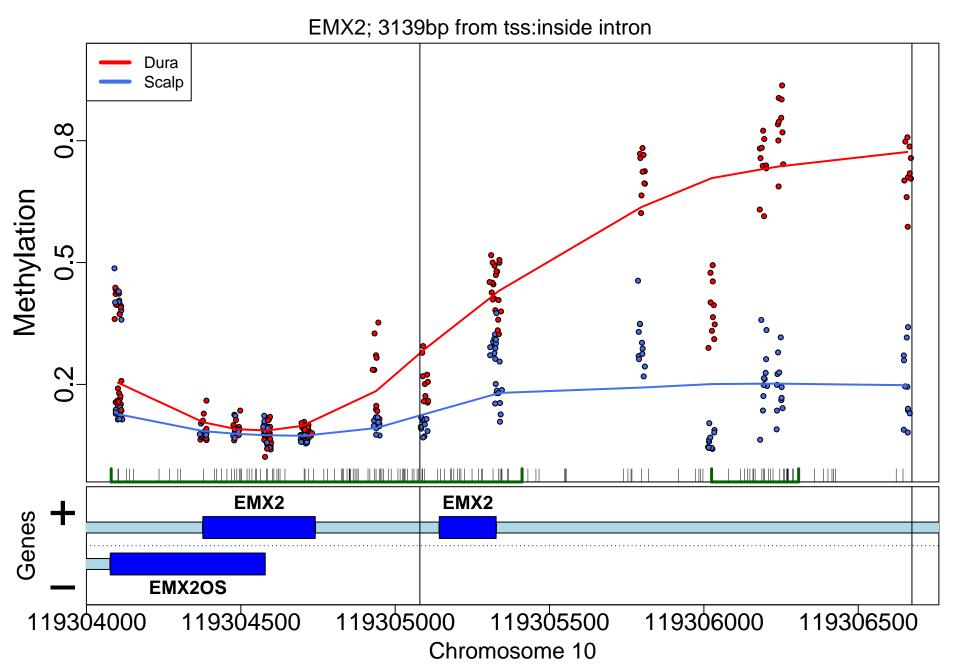


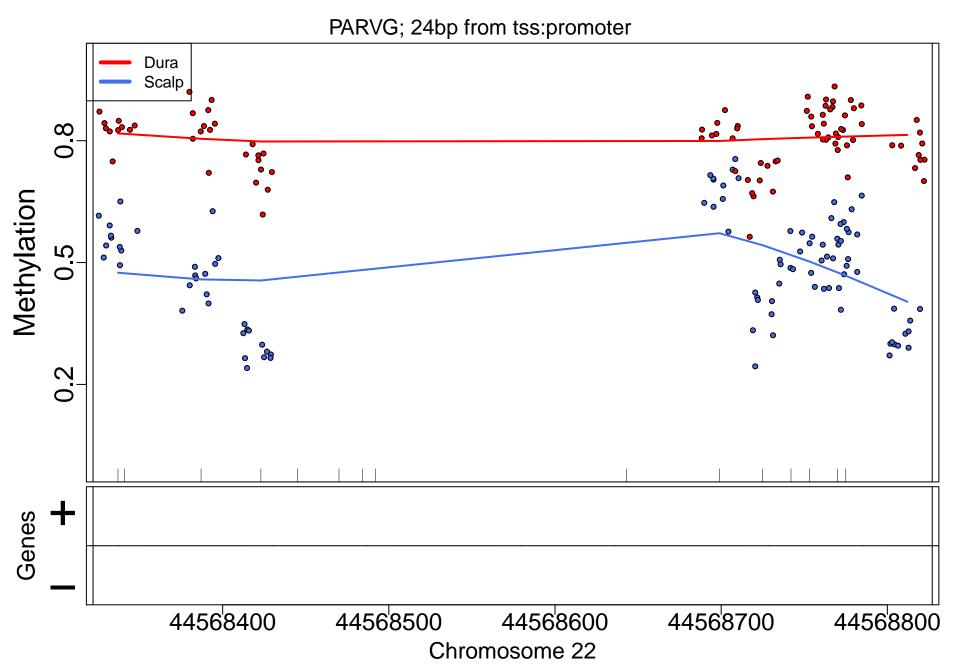


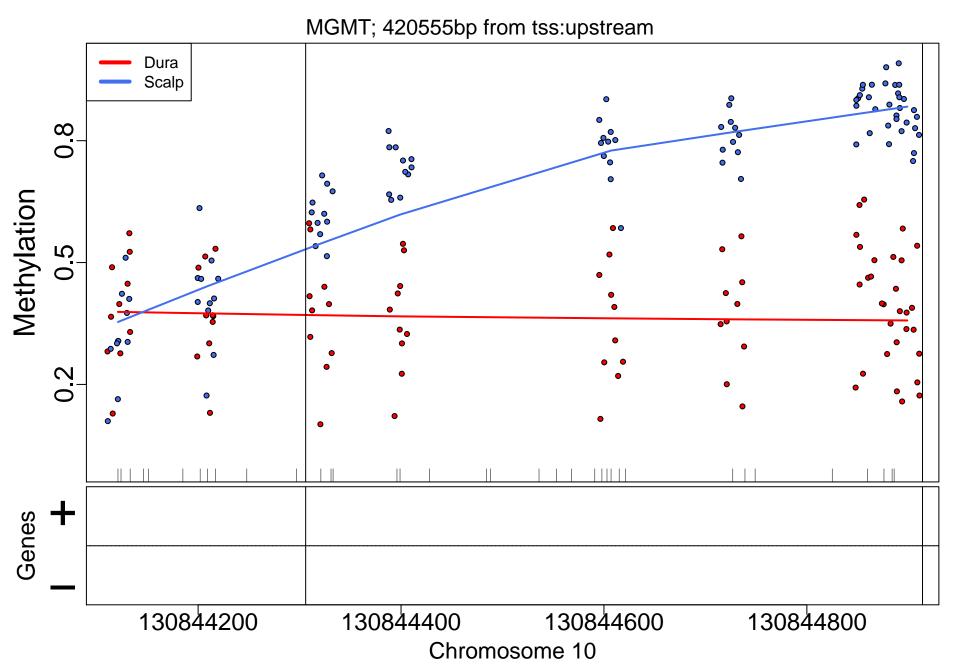
LOC145845; 0bp from tss:NA Dura Scalp 800 <u>ω</u>_ Methylation Genes 37170000 37171000 37170500 Chromosome 15







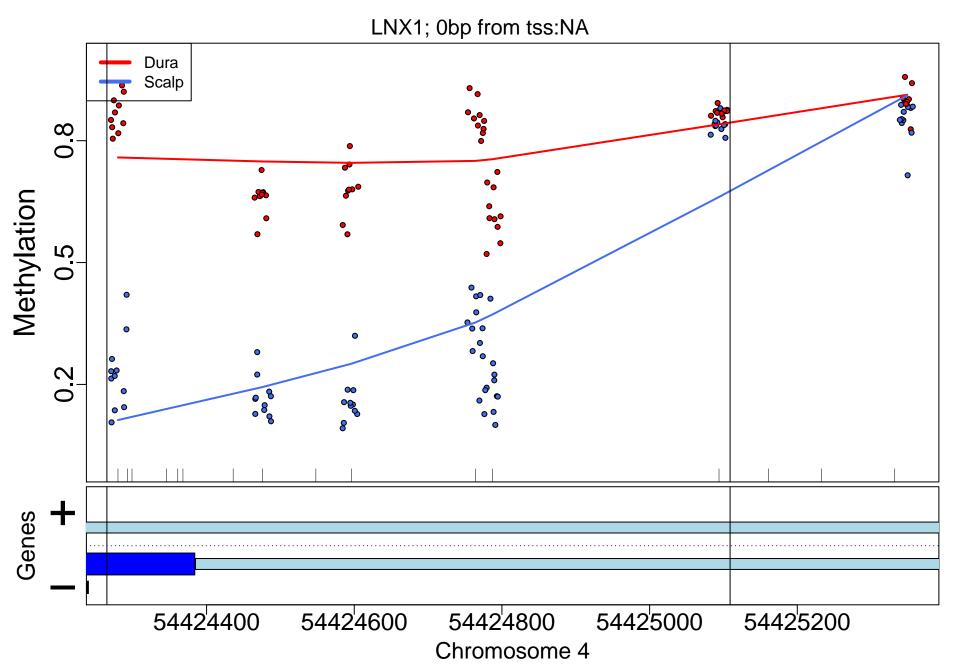


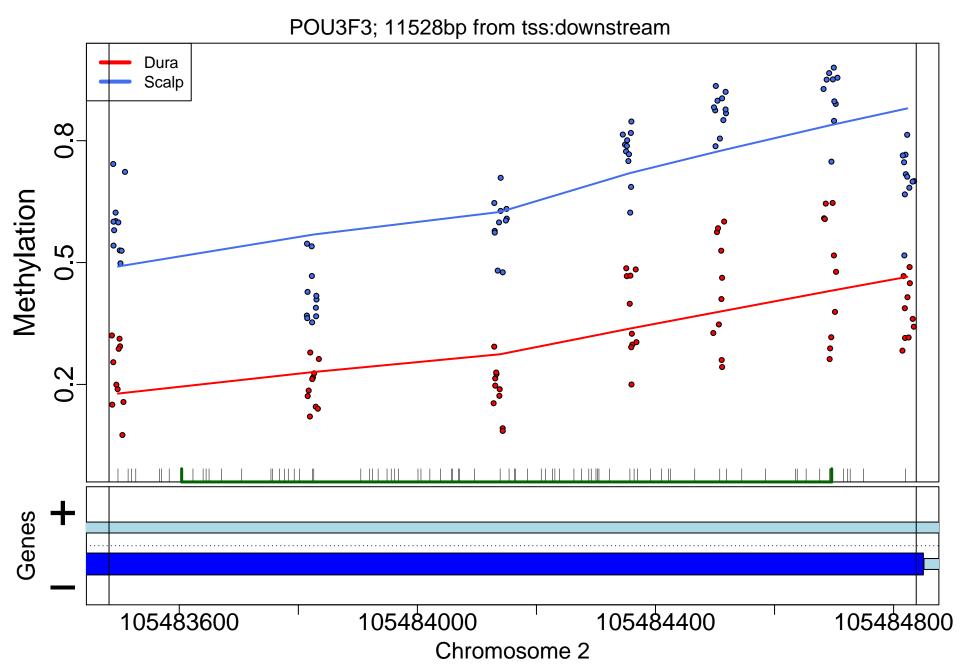


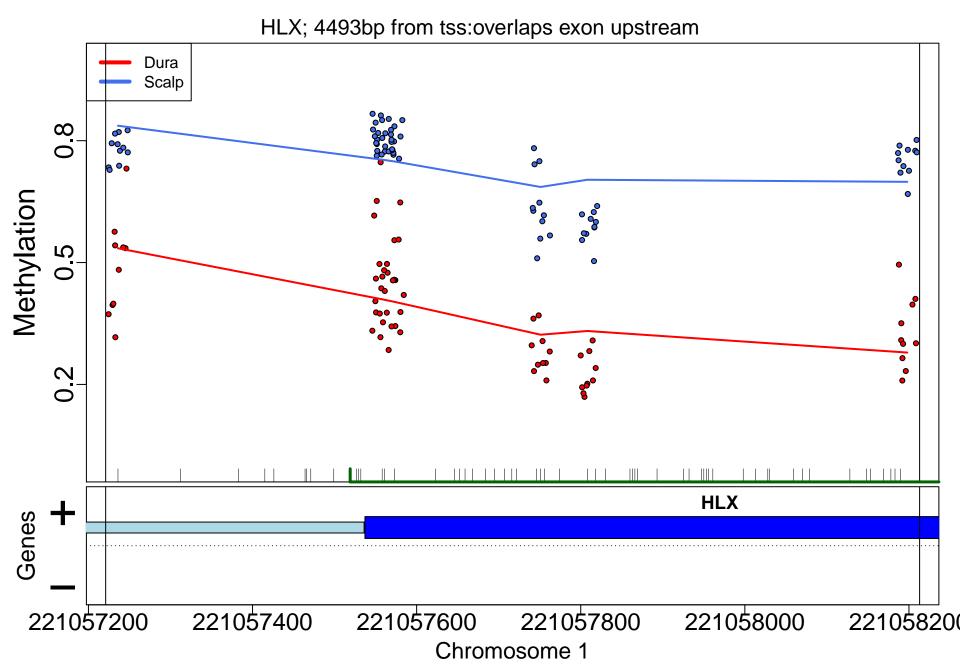
ZFHX3; 5369bp from tss:upstream Dura Scalp <u>∞</u>. Methylation 8 00 0.2 Genes 73098000 73098500 73099000 73099500 Chromosome 16

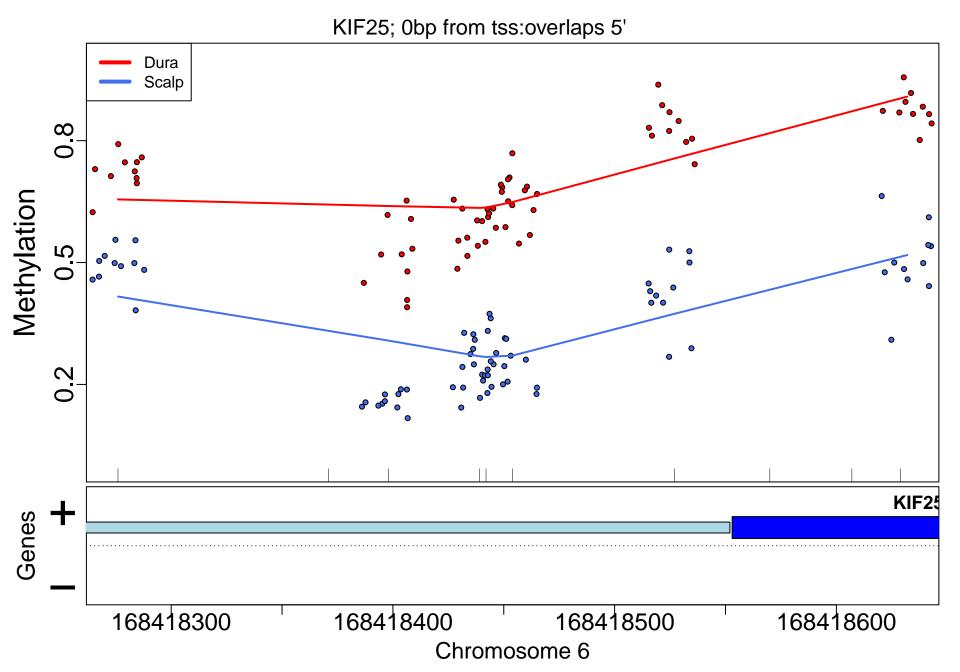
ABR; 3413bp from tss:upstream Dura Scalp 0 Methylation Genes 1094100 1094200 1094300 1094500 1094400 Chromosome 17

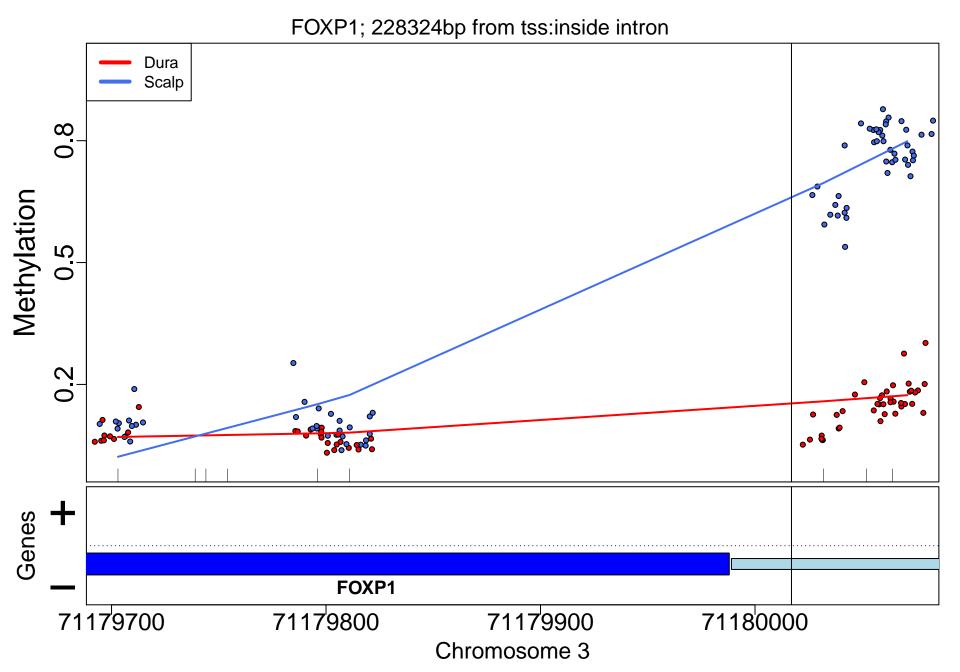
TNXB; 12387bp from tss:inside exon Dura Scalp 0 Methylation 0.2 Genes **TNXB** 32063600 32064800 3206 4000 32064400 Chromosome 6

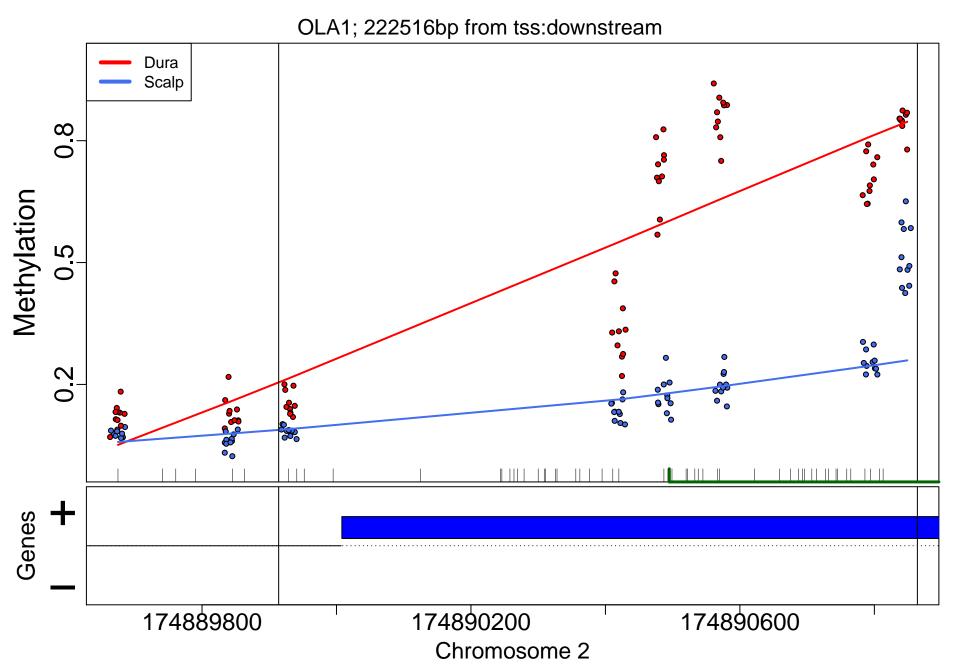


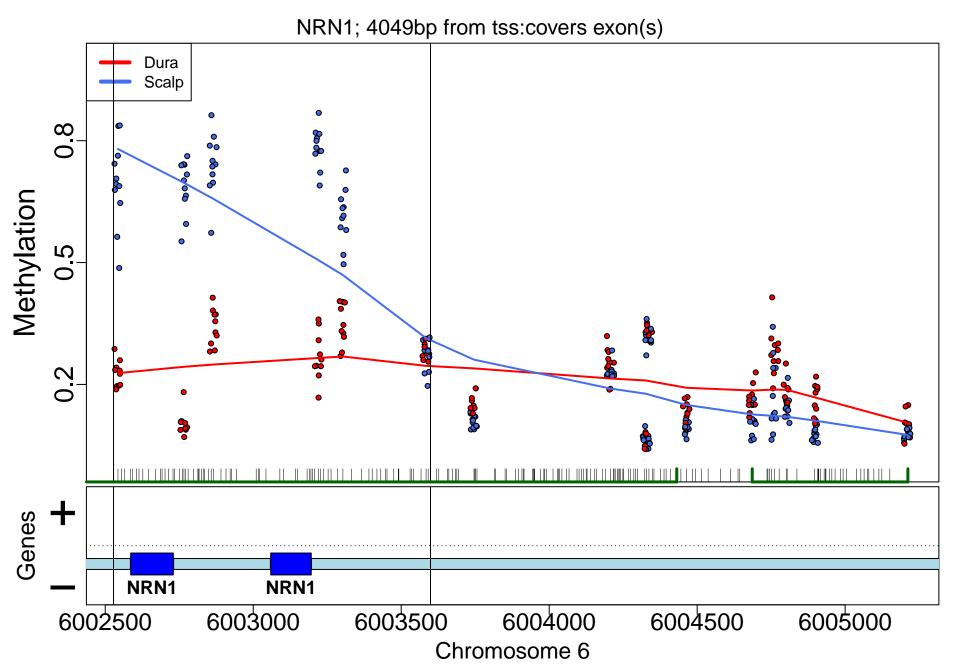


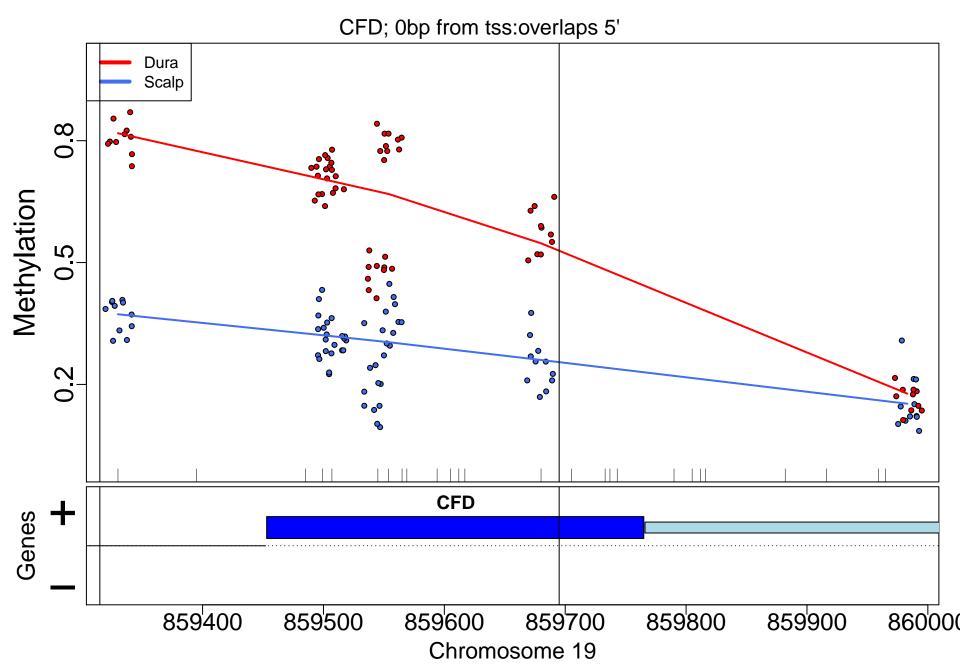


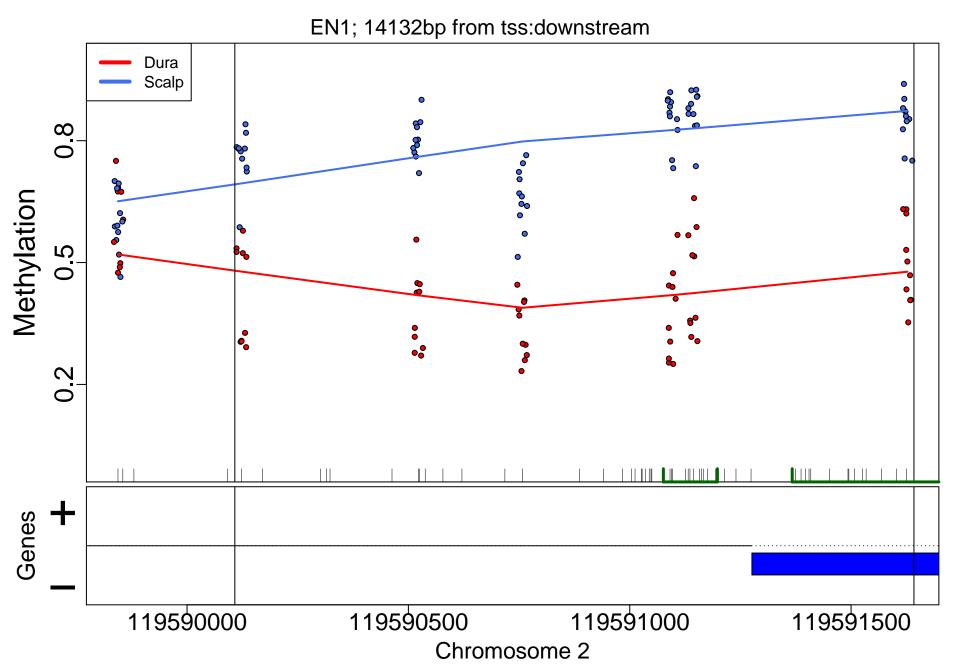


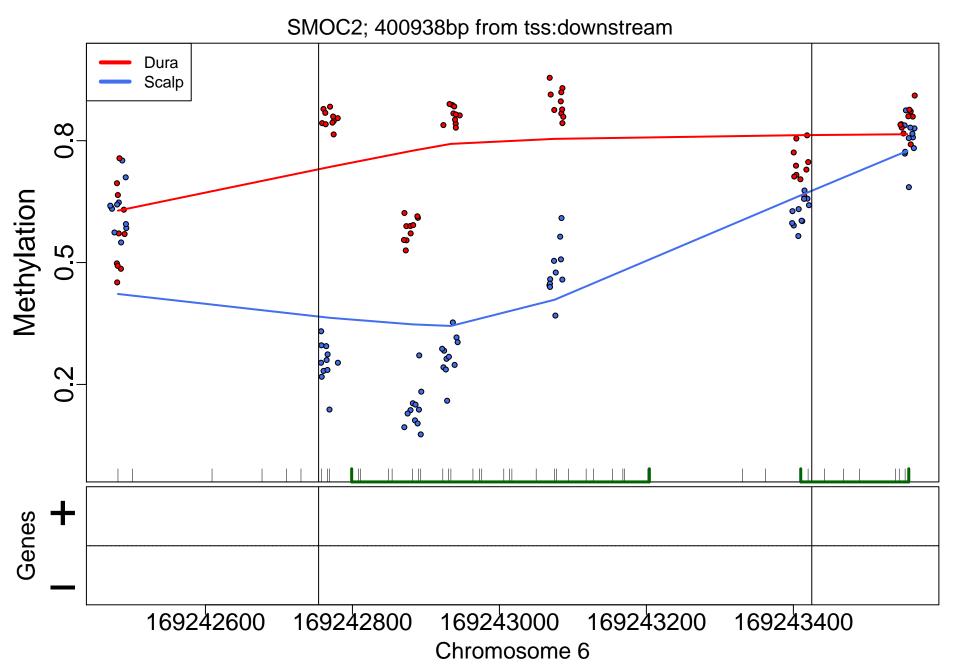


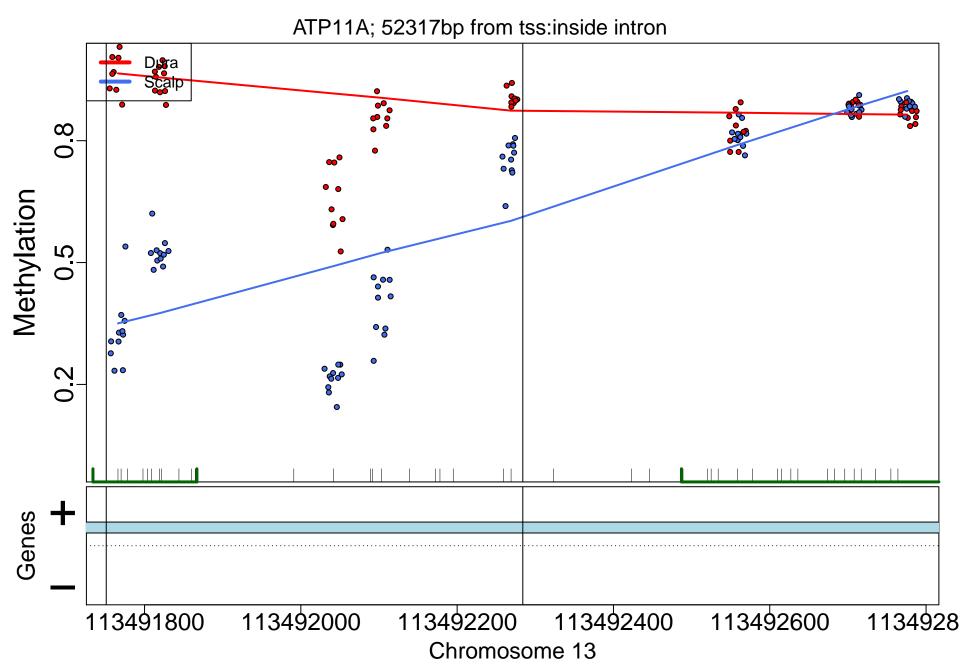


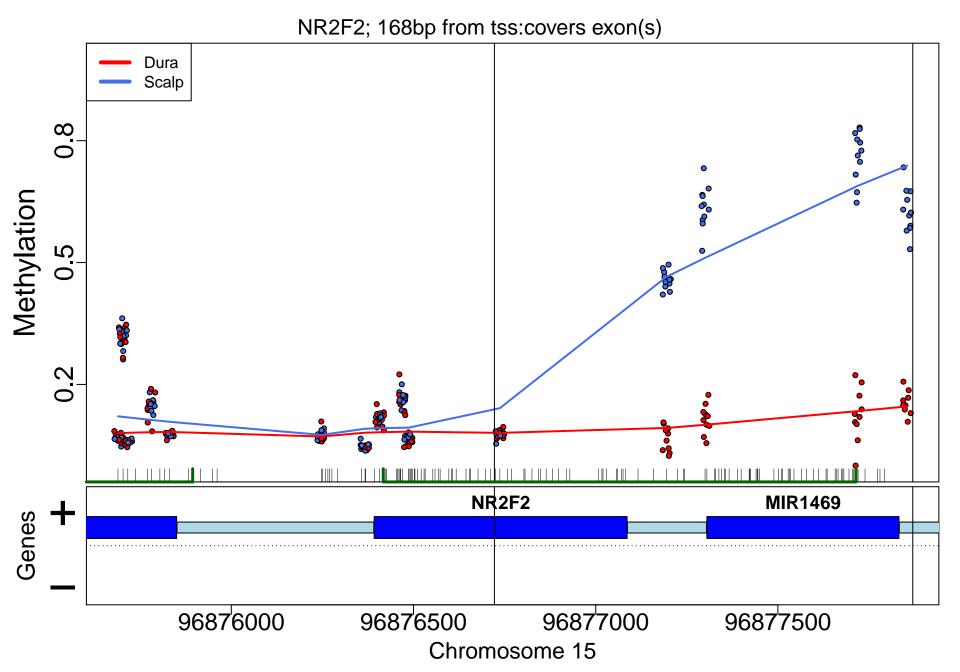


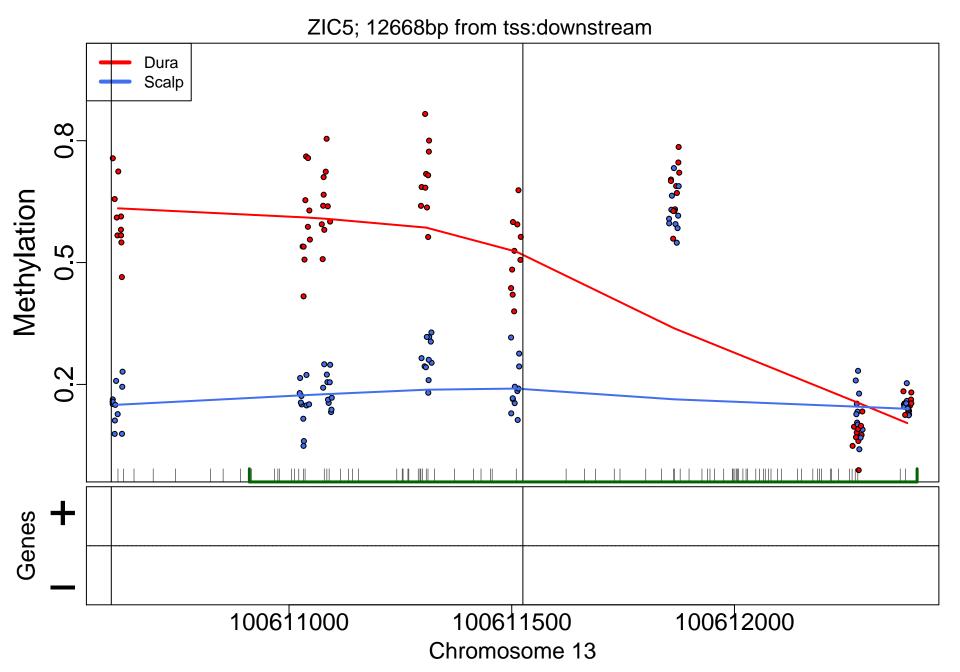


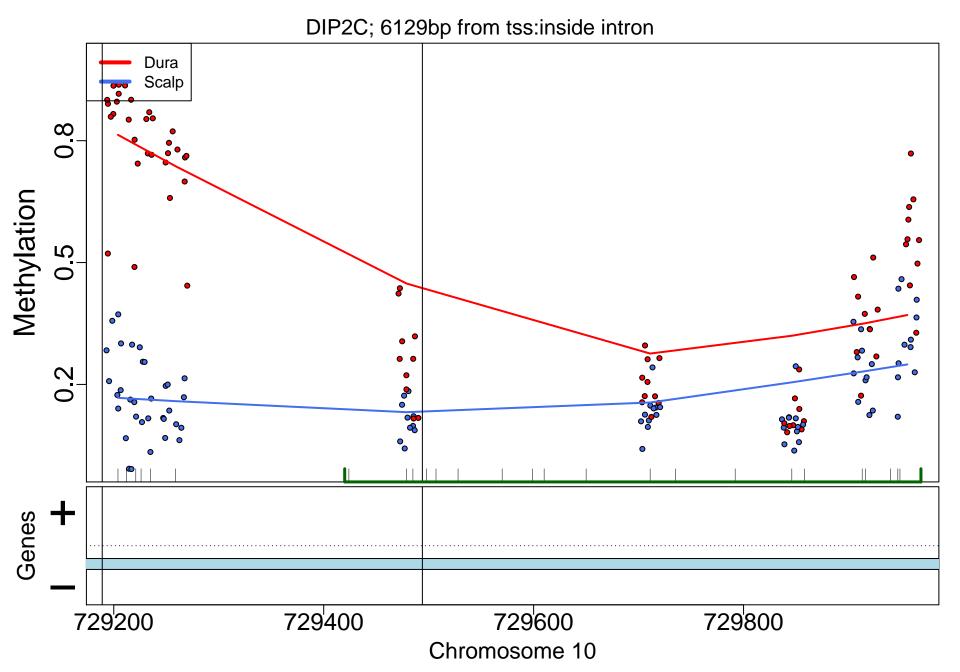


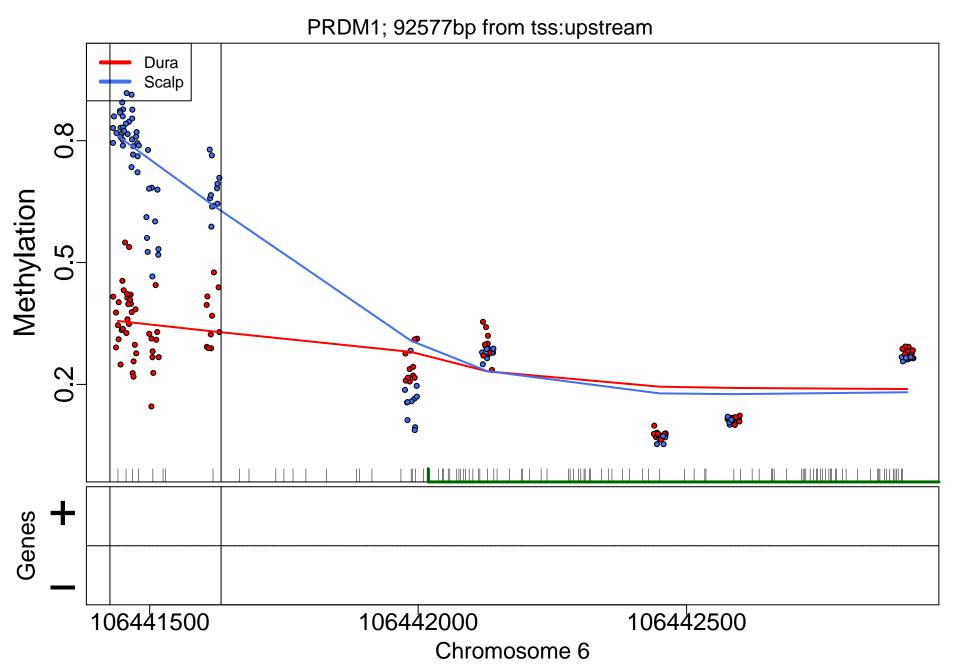


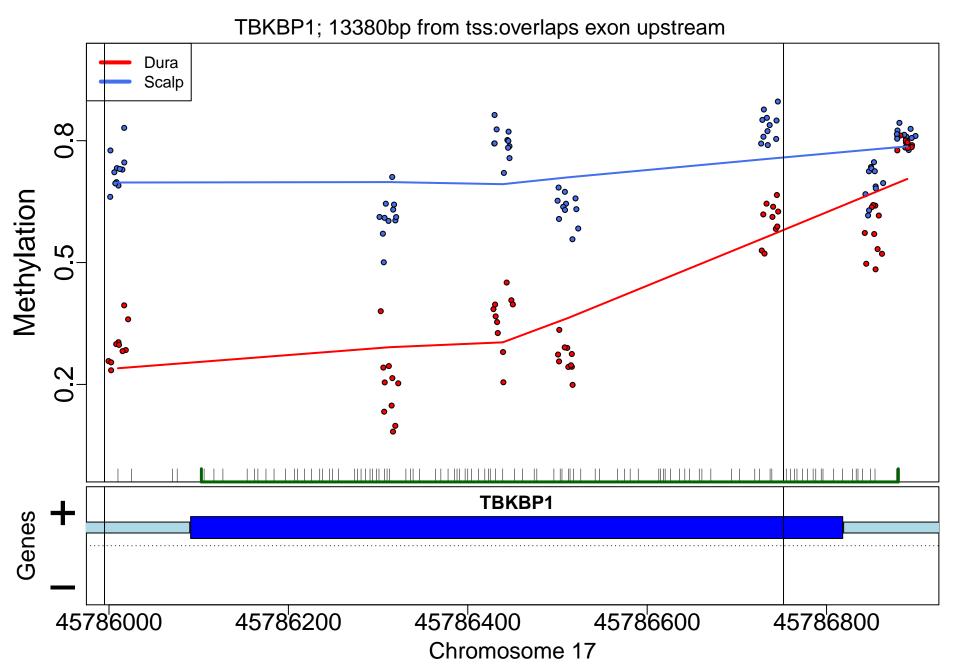


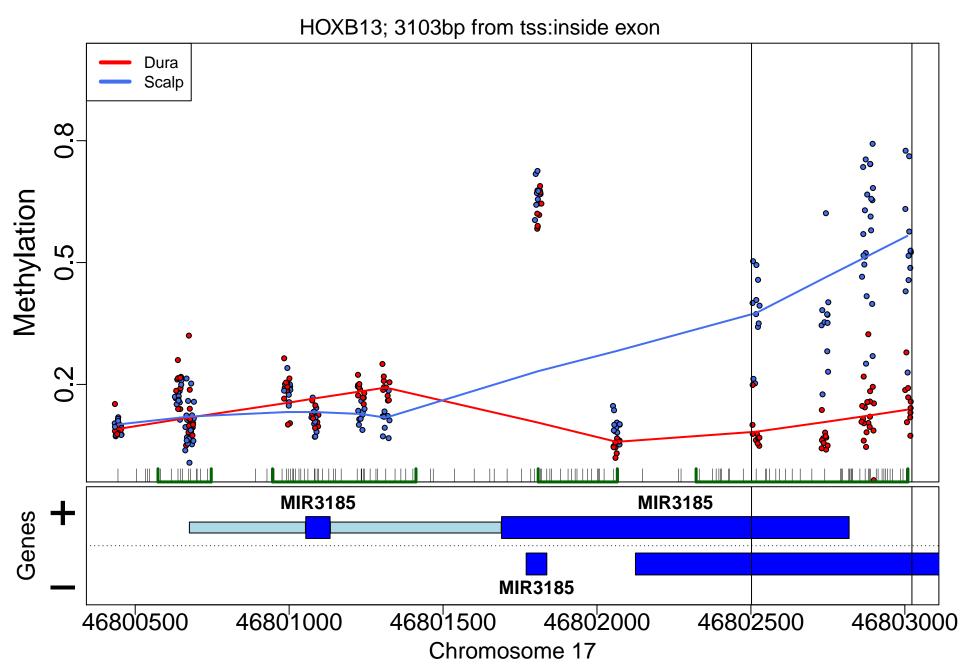


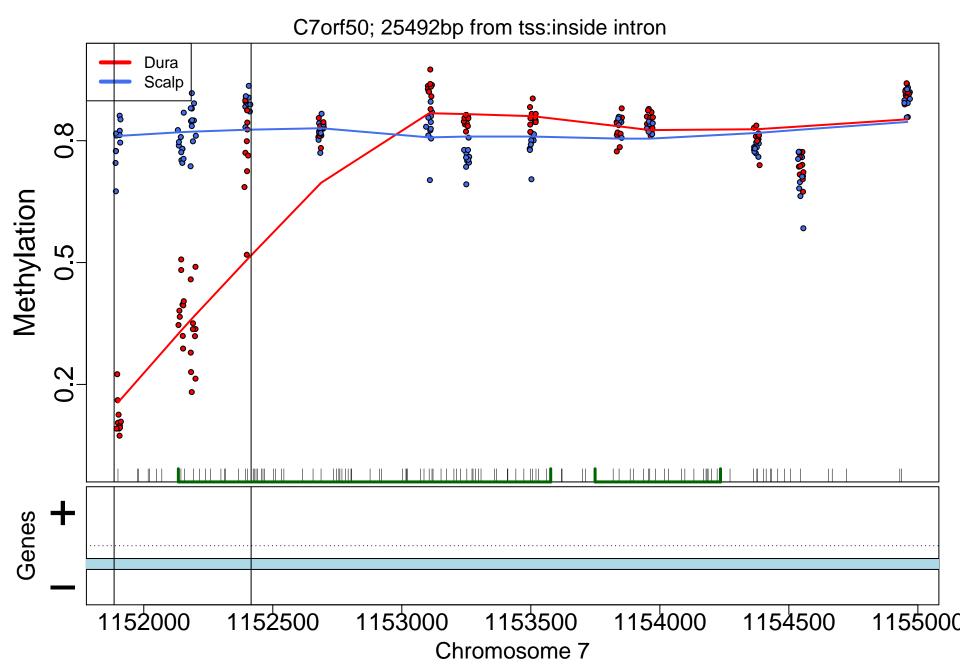


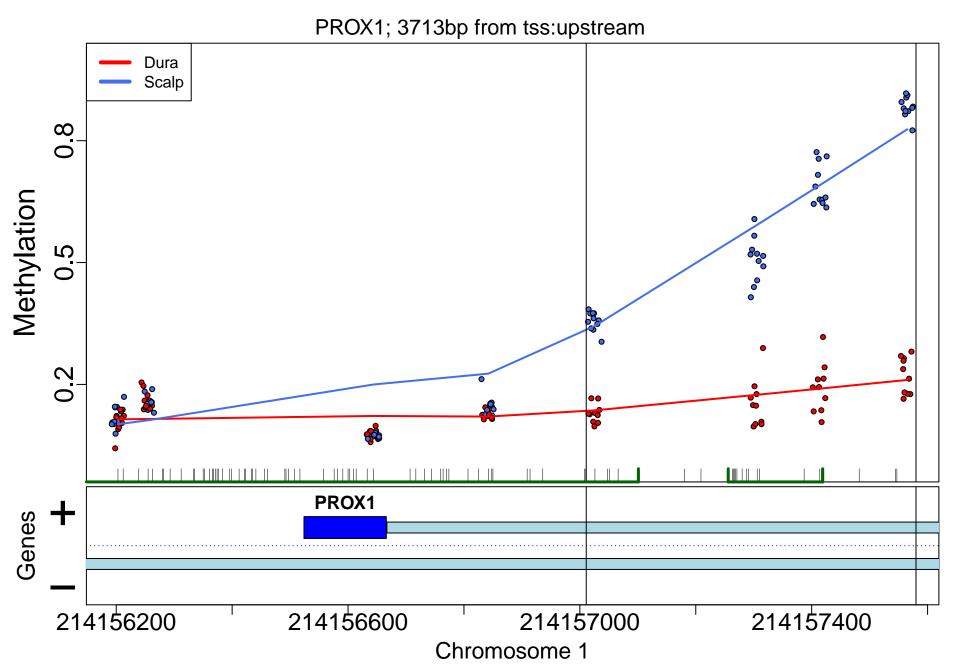


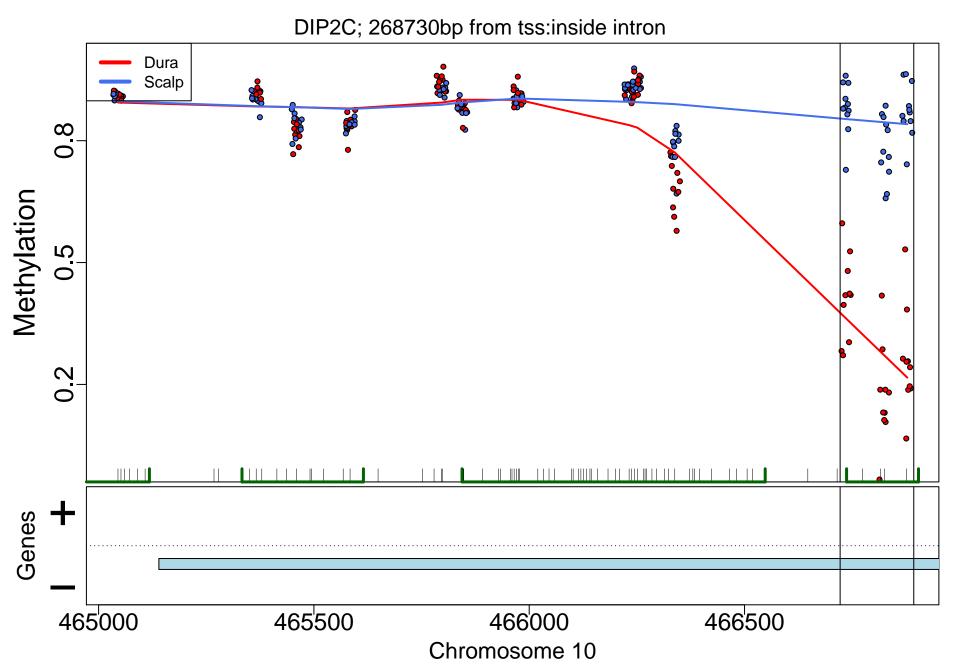


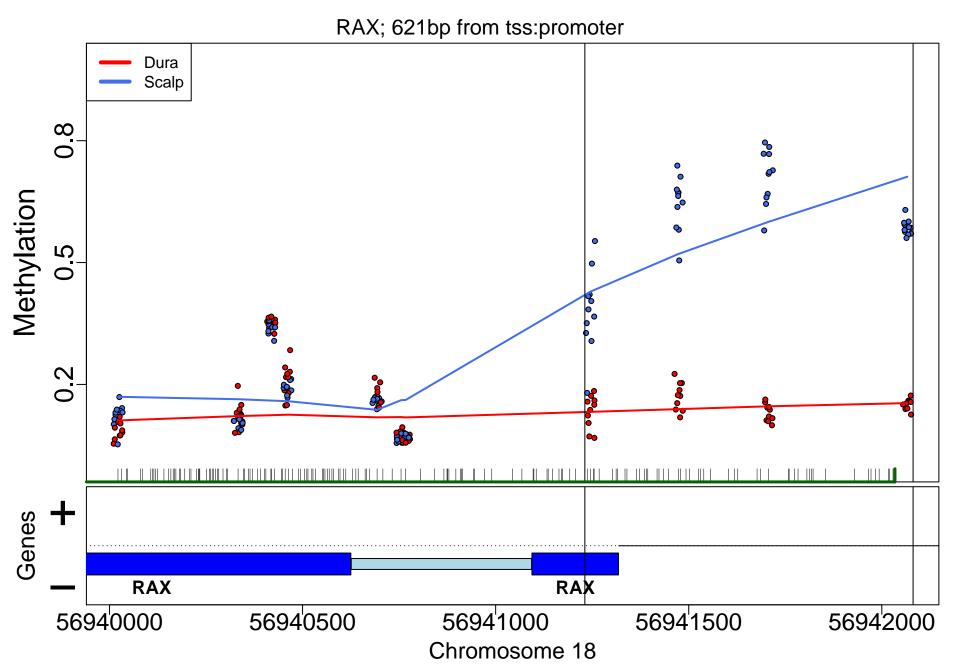


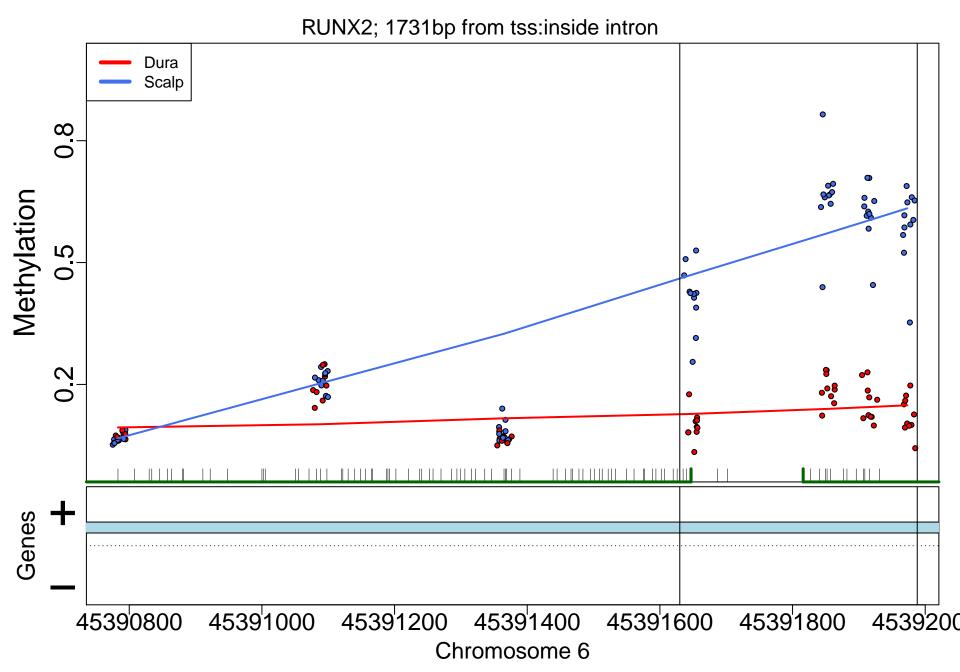


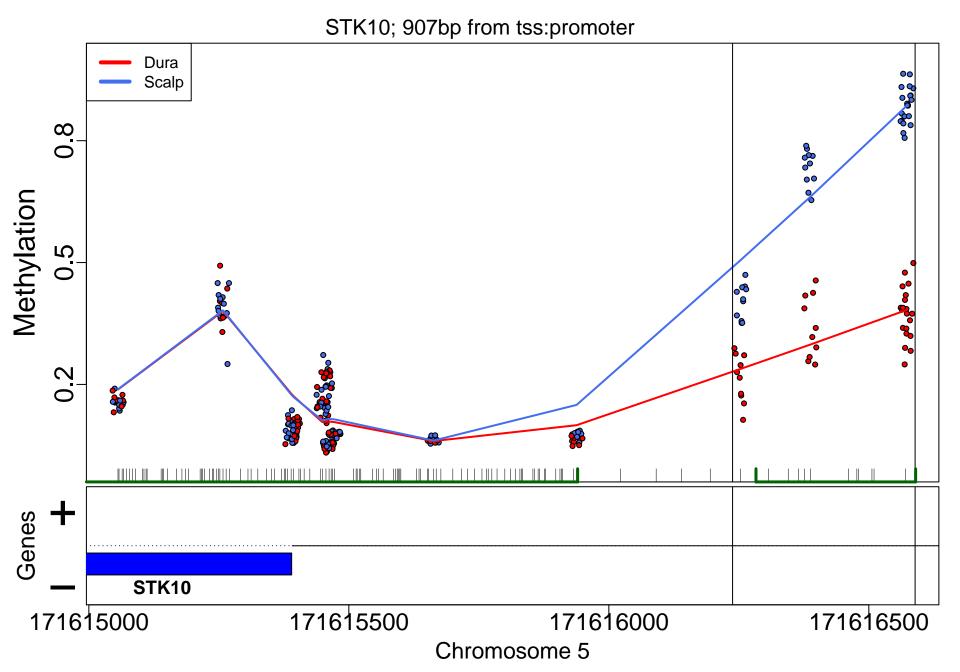


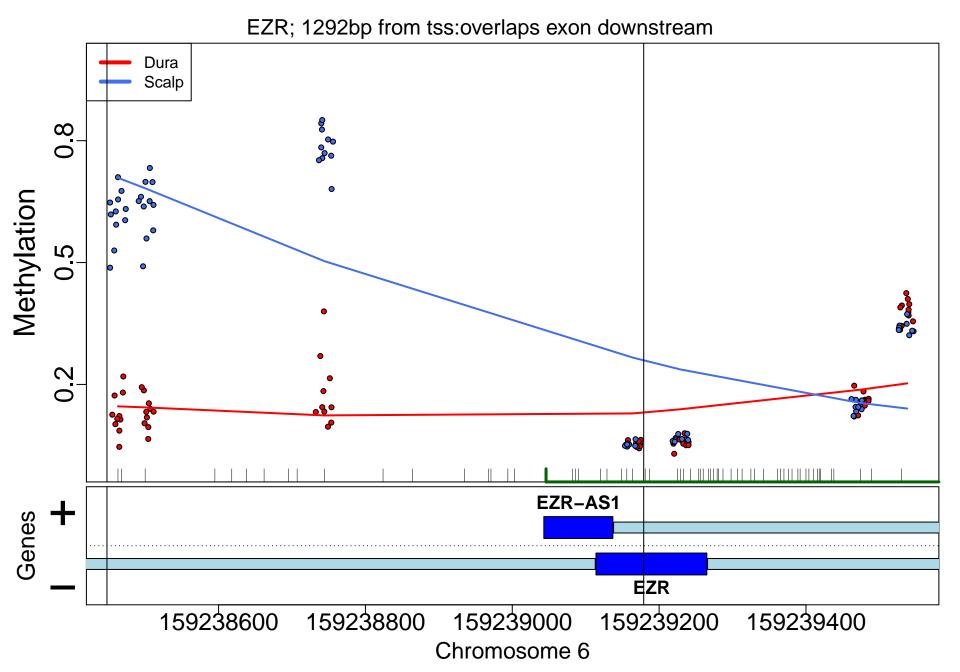


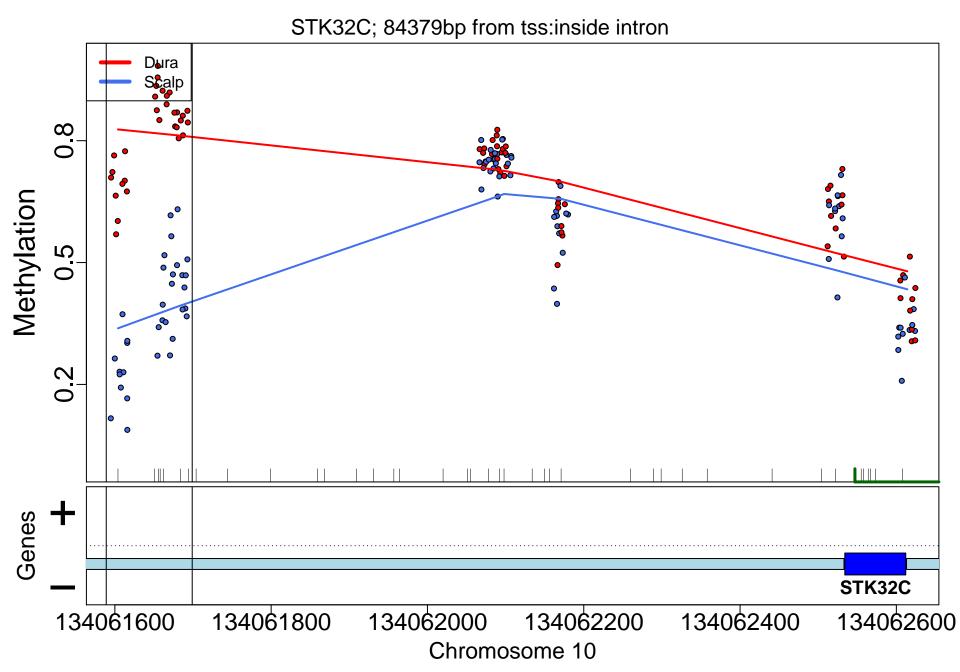


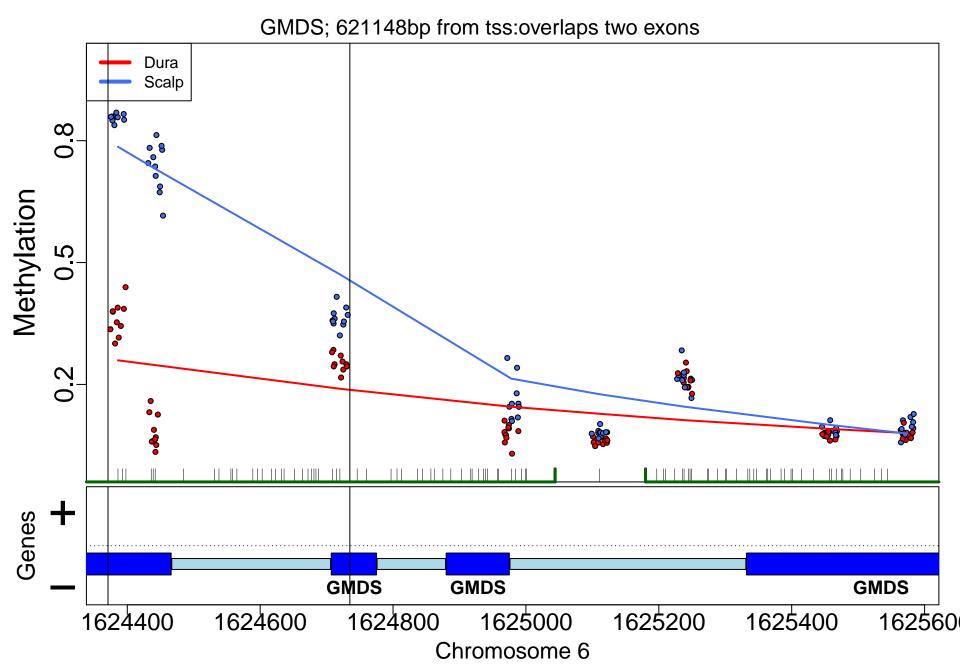


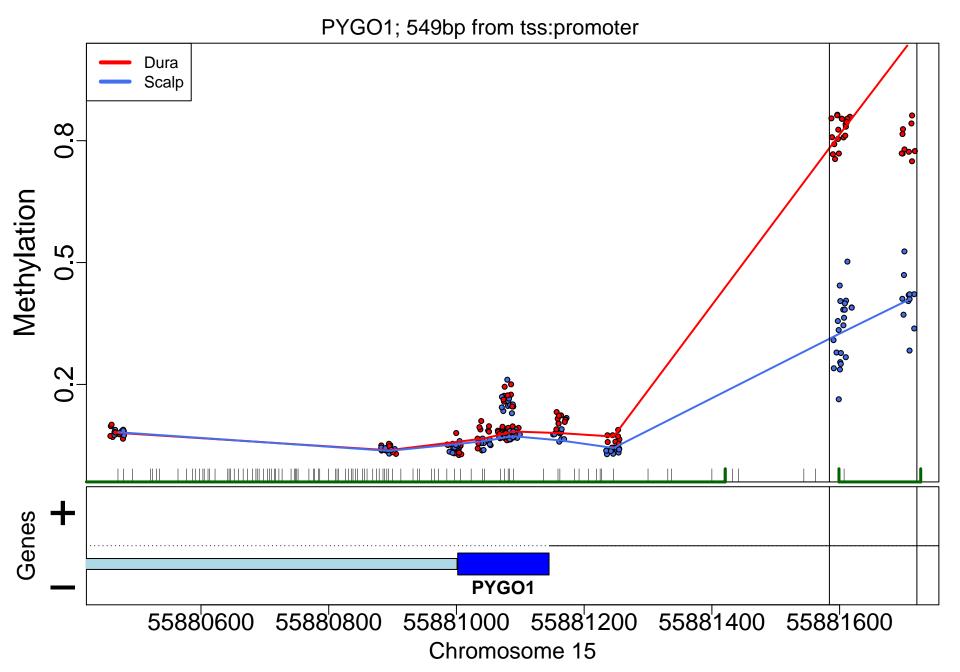


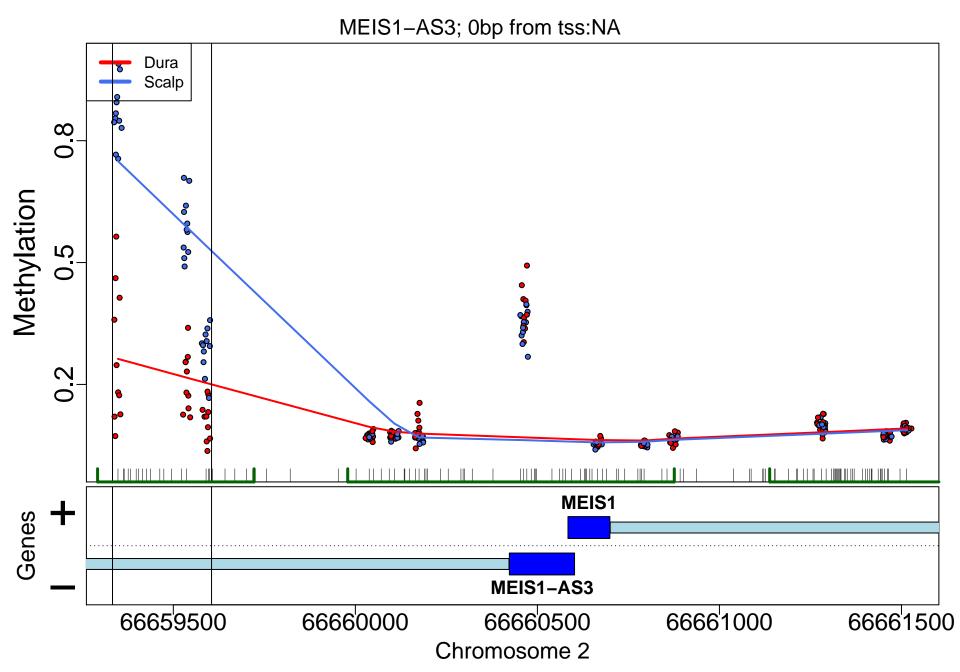


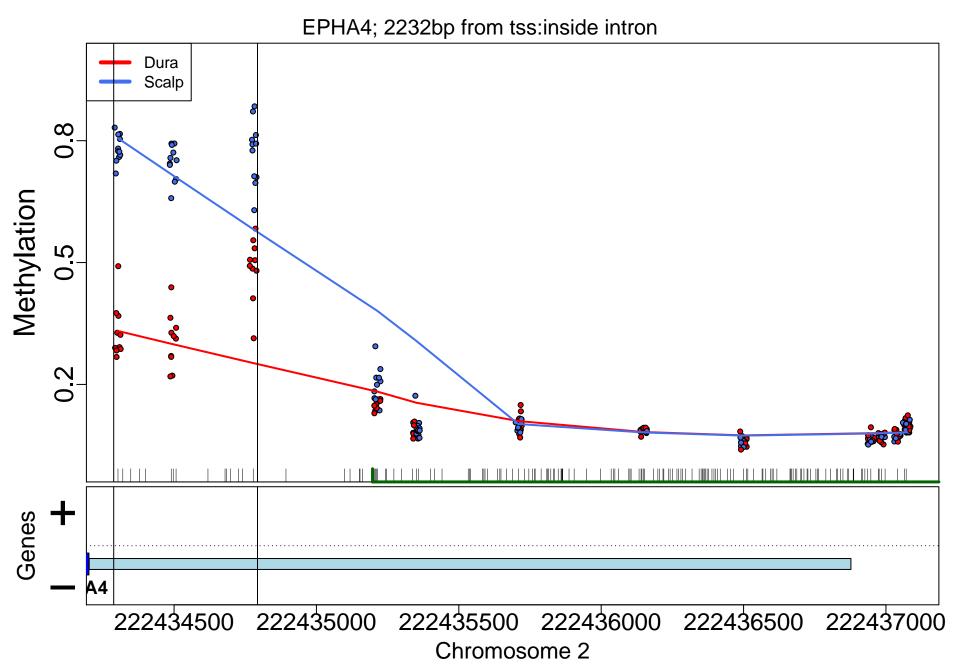


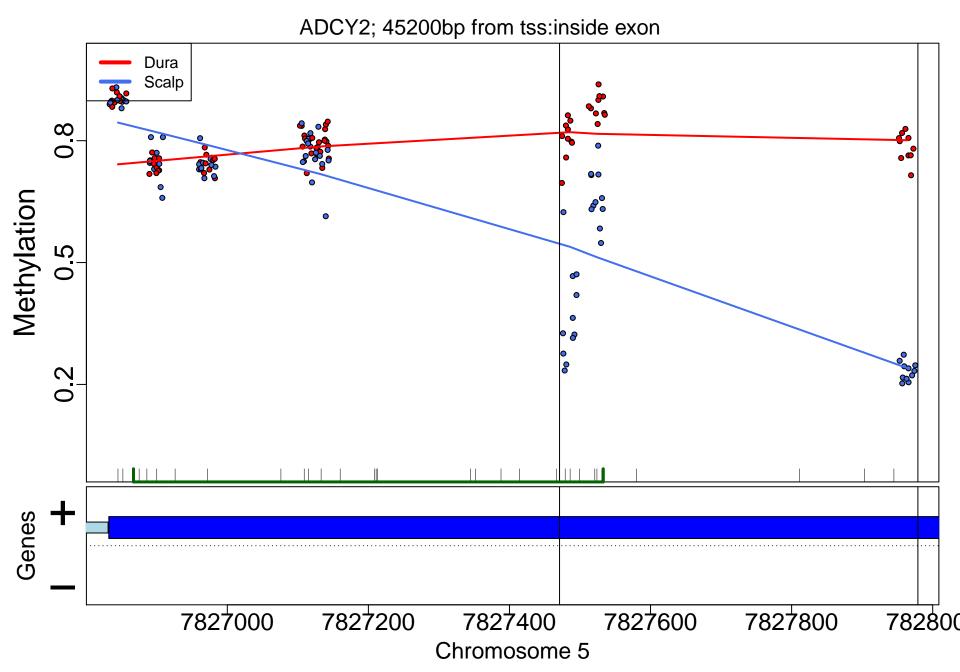


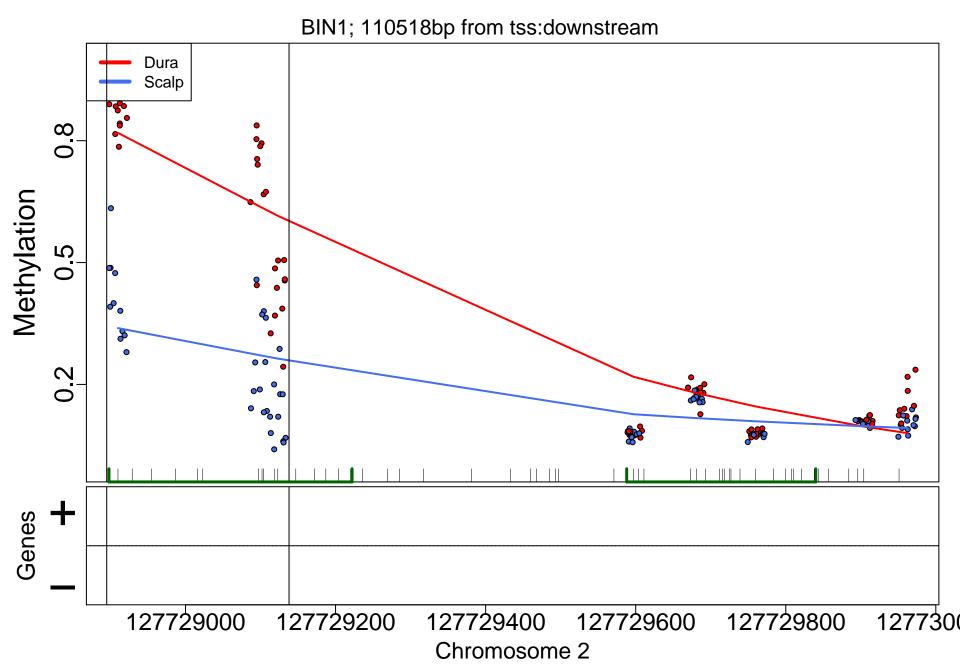


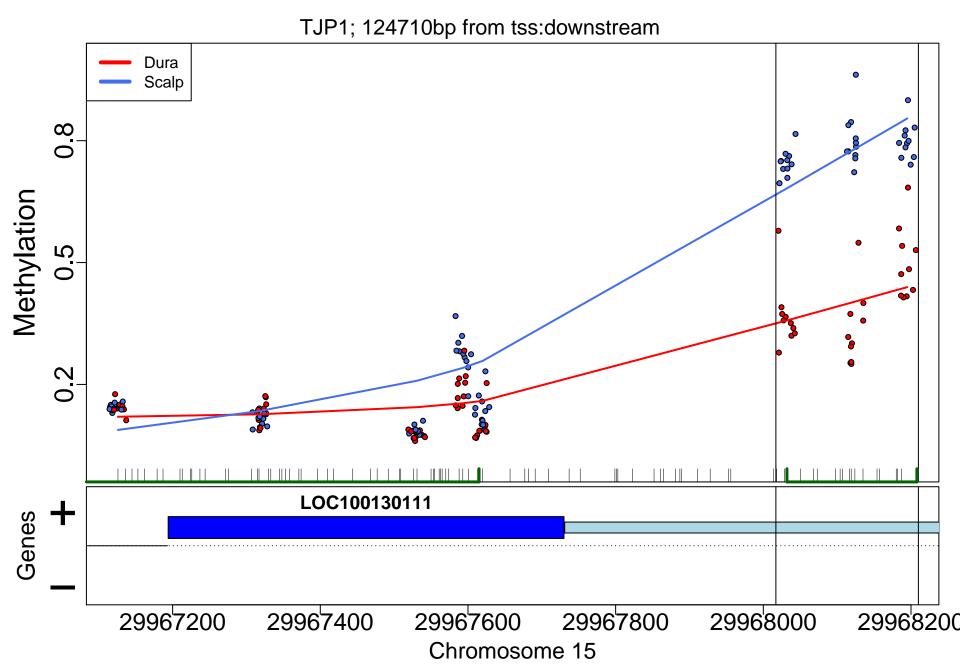




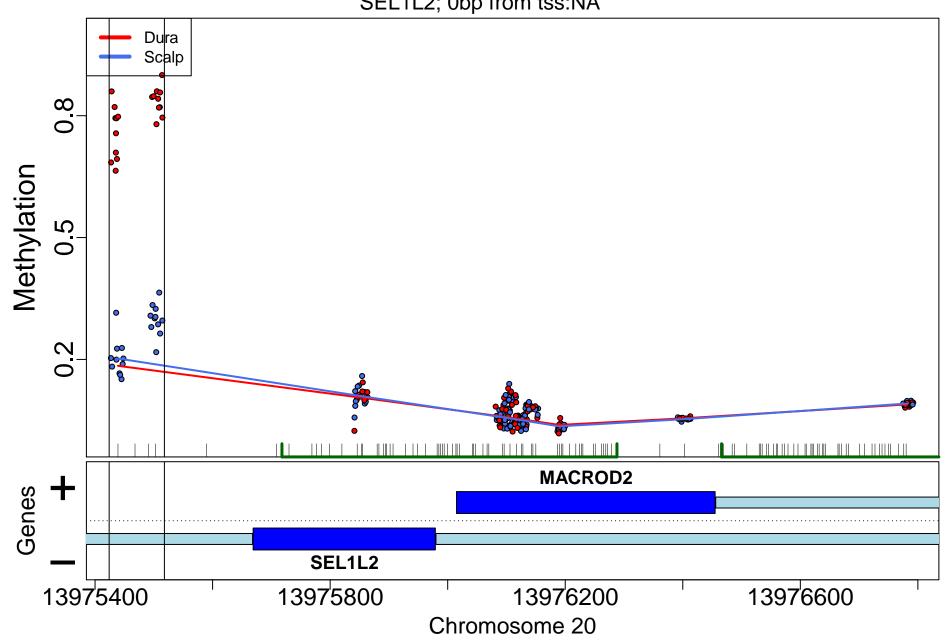






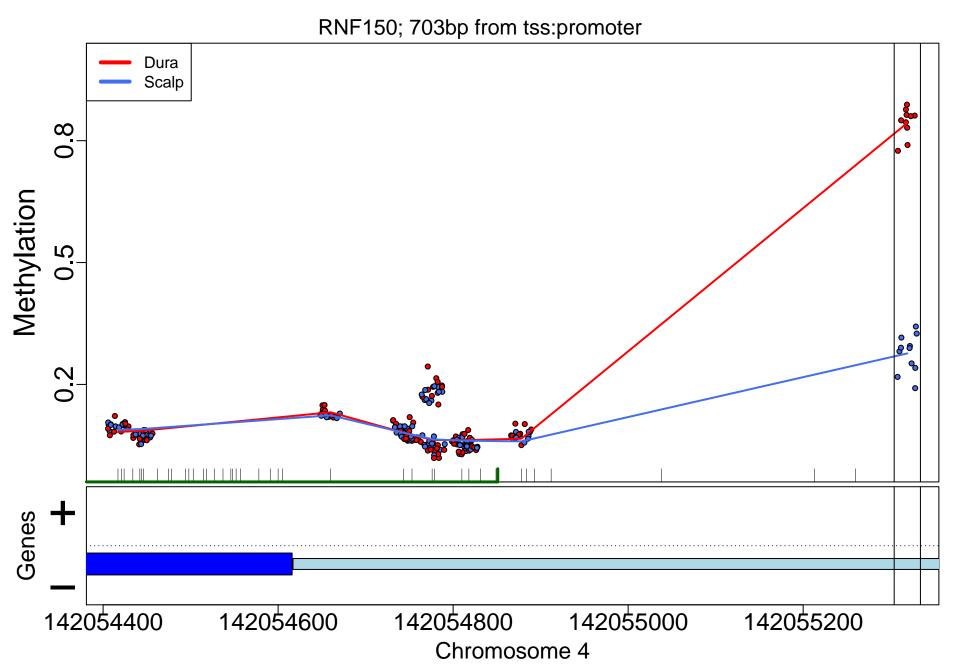


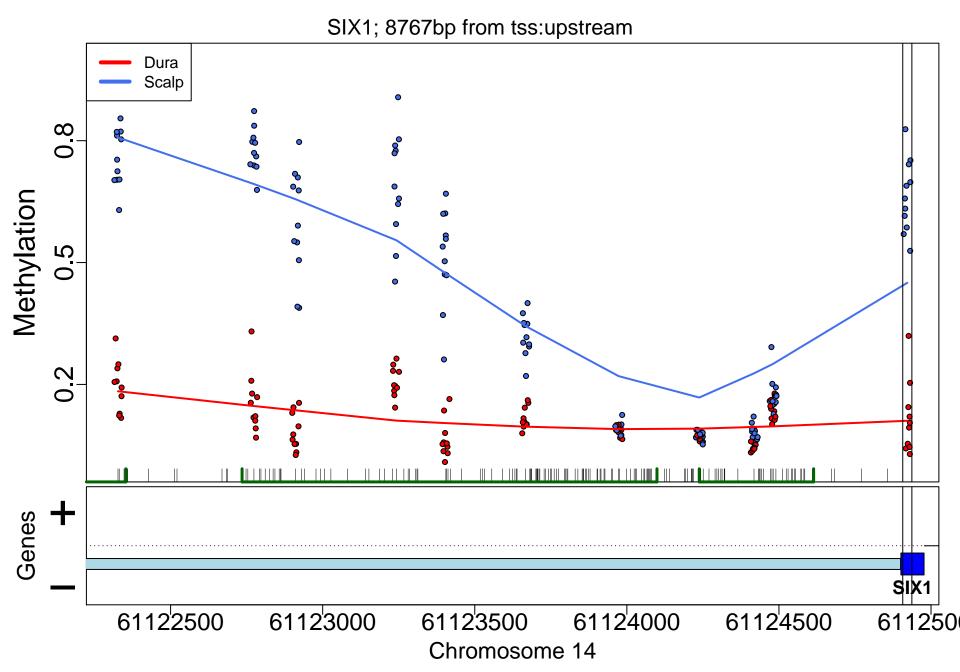
SEL1L2; 0bp from tss:NA

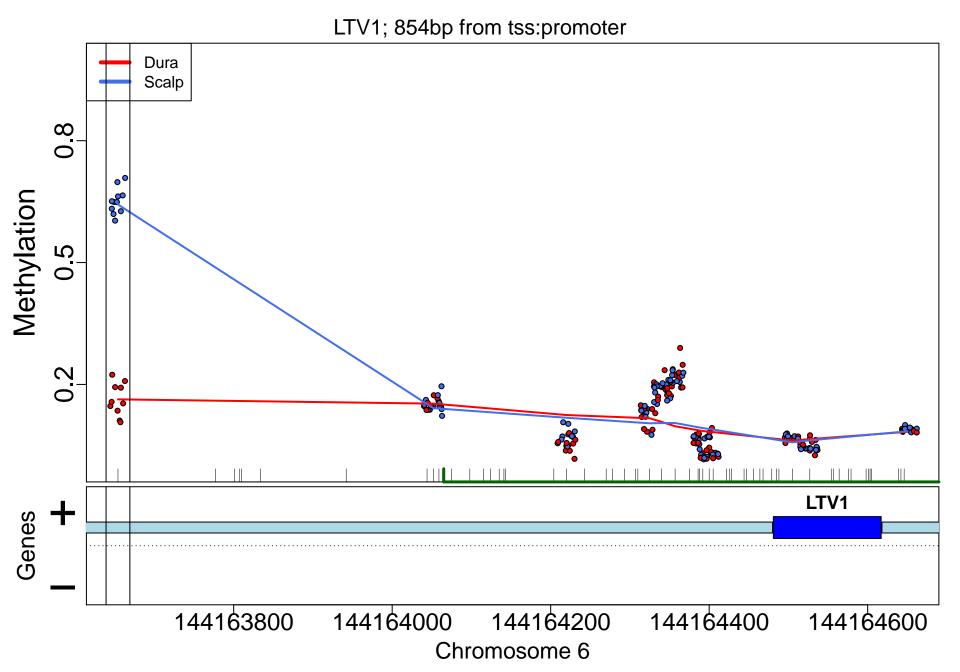


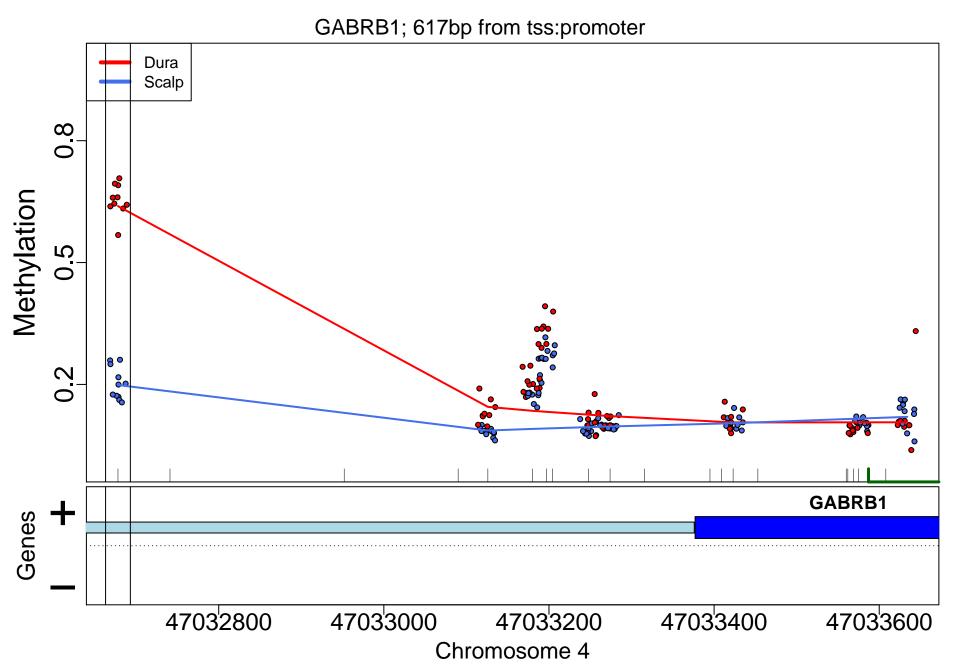
IFITM1; 1111bp from tss:inside exon Dura Scalp <u>ω</u>_ Methylation • ° 08 % 0.2 IFITM1 Genes 315200 315400 316400 315600 315800 316000 316200 Chromosome 11

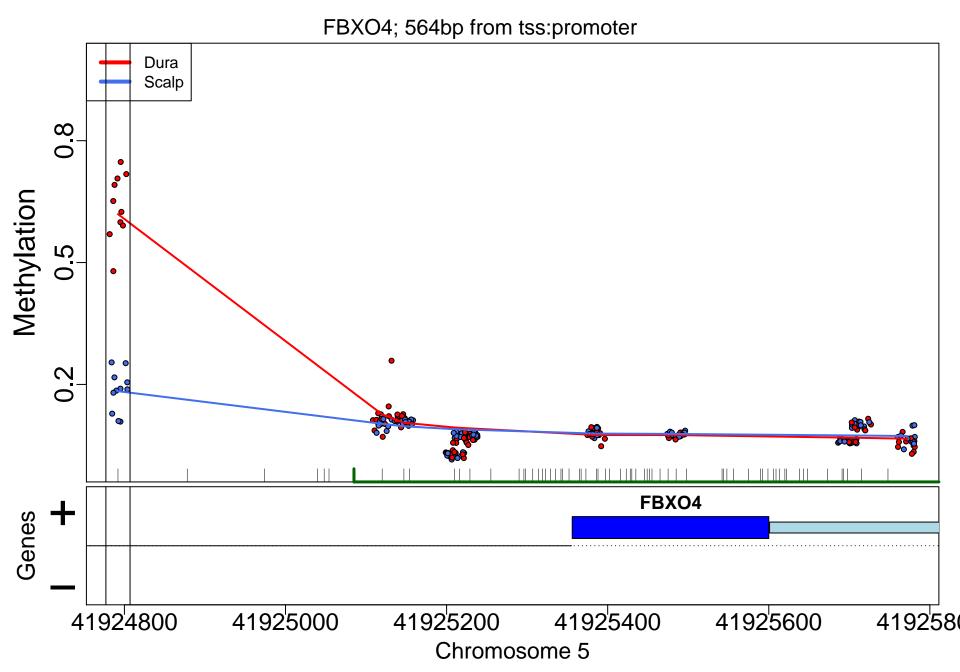
C2CD2; 863bp from tss:inside intron Dura Scalp <u>∞</u>. Methylation 0.2 Genes C2CD2 C2CD2 43373500 43374500 43374000 Chromosome 21

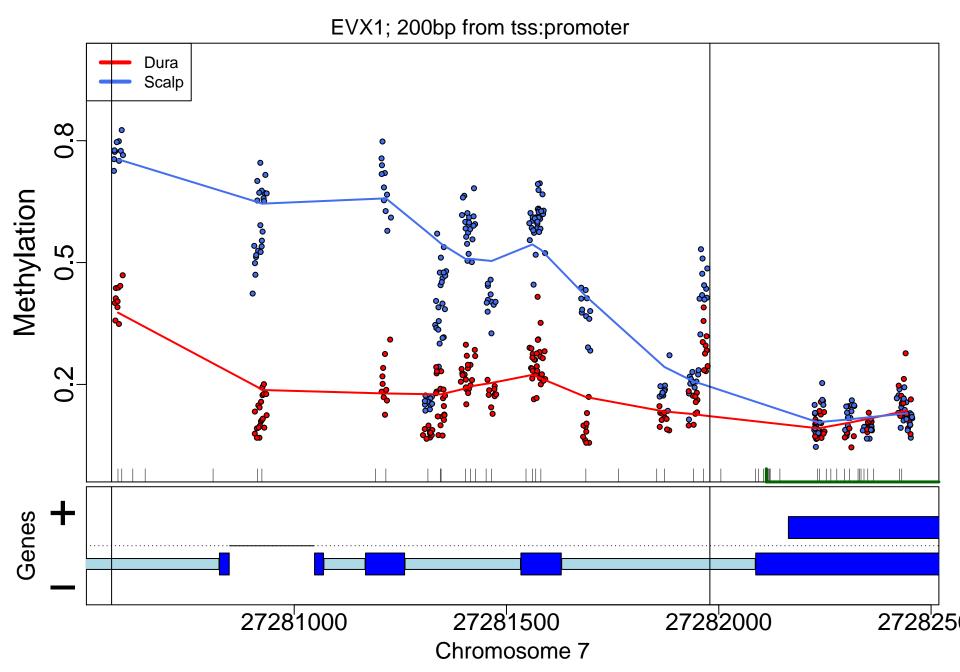




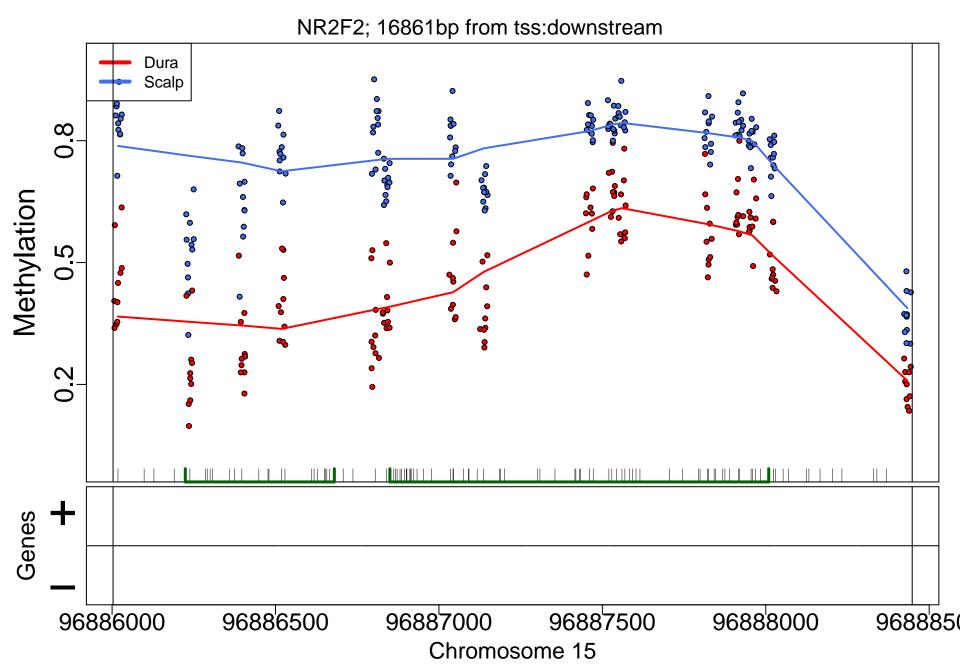


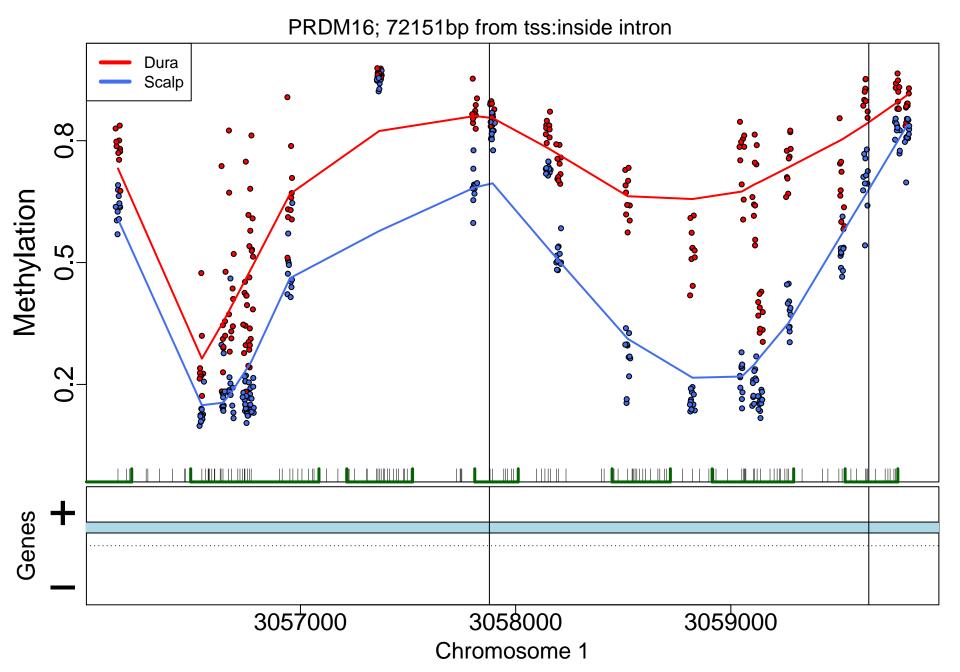


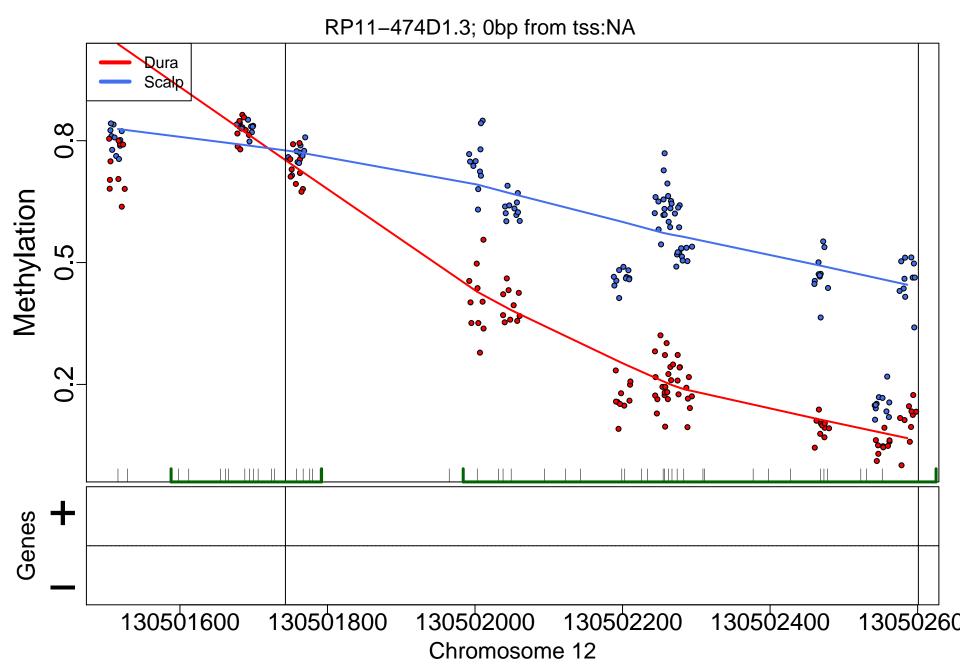


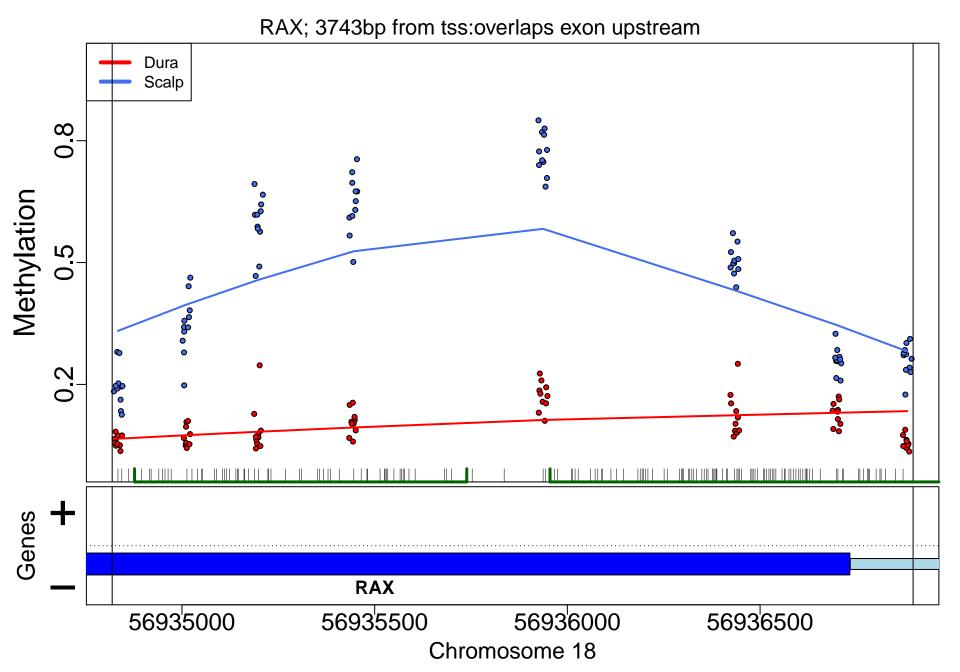


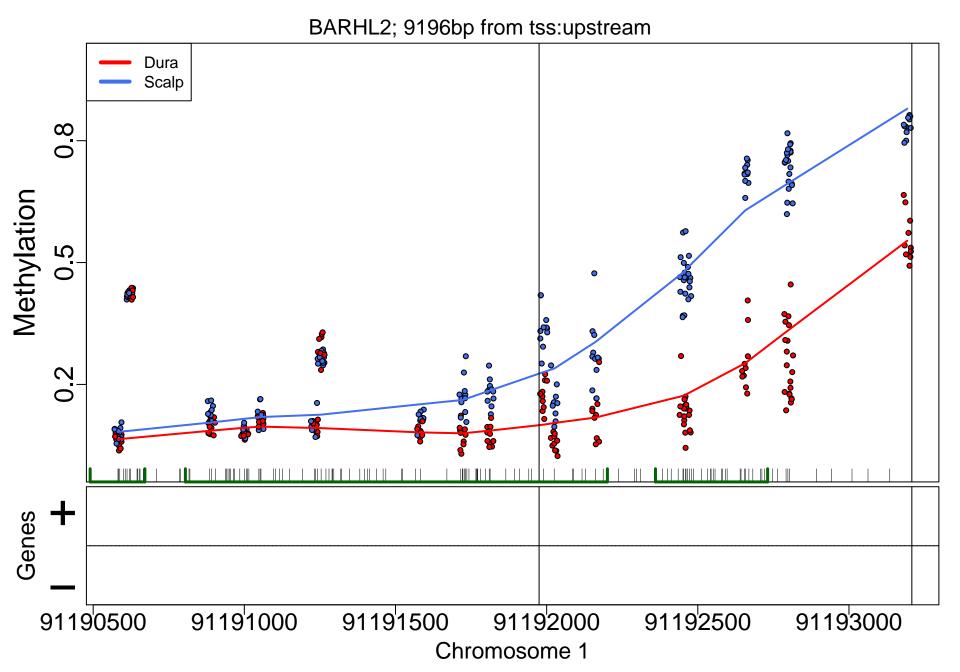
TNXB; 27198bp from tss:overlaps two exons Dura Scalp <u>ω</u>_ Methylation 0.2 Genes TNXB **TNXB** 32048500 32049000 32049500 3205 0000 Chromosome 6

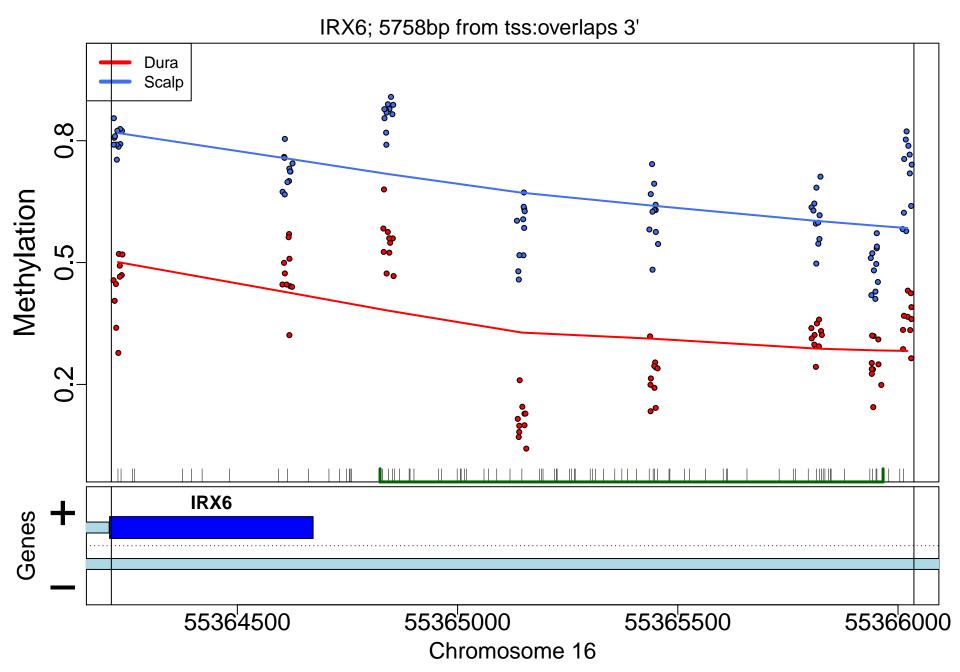


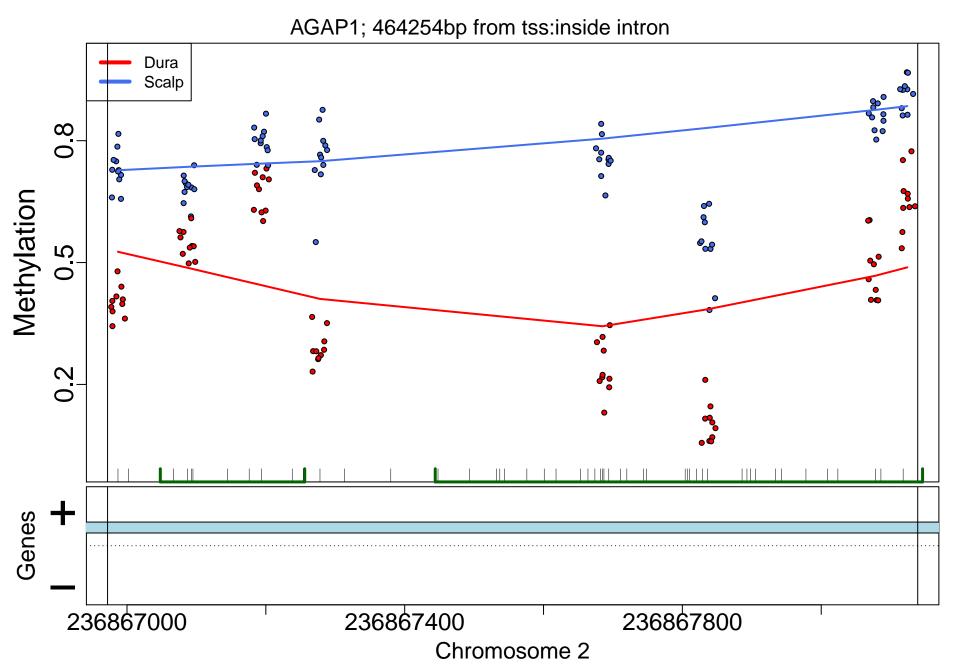


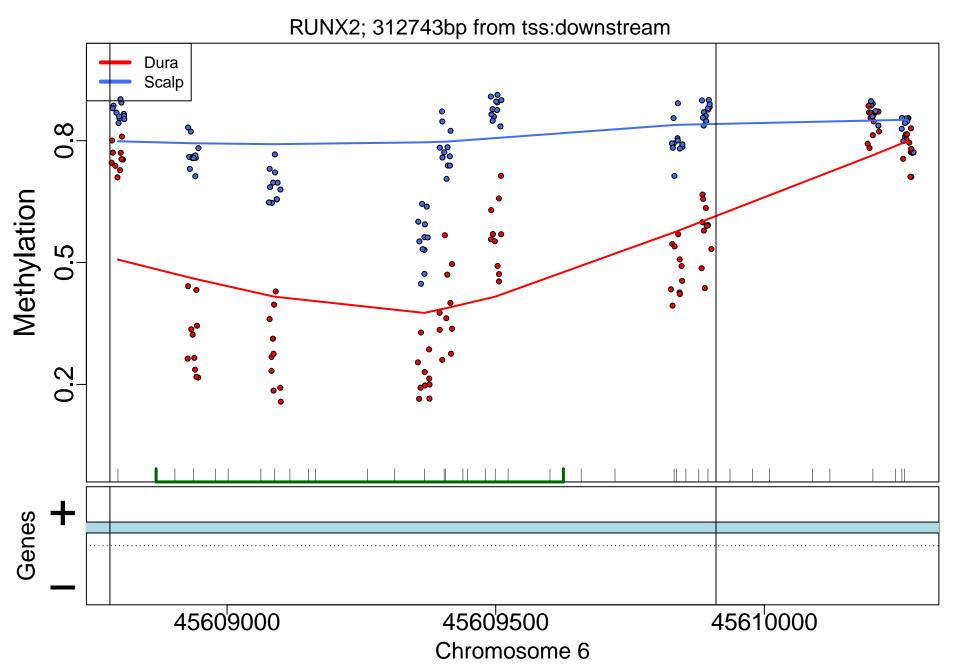


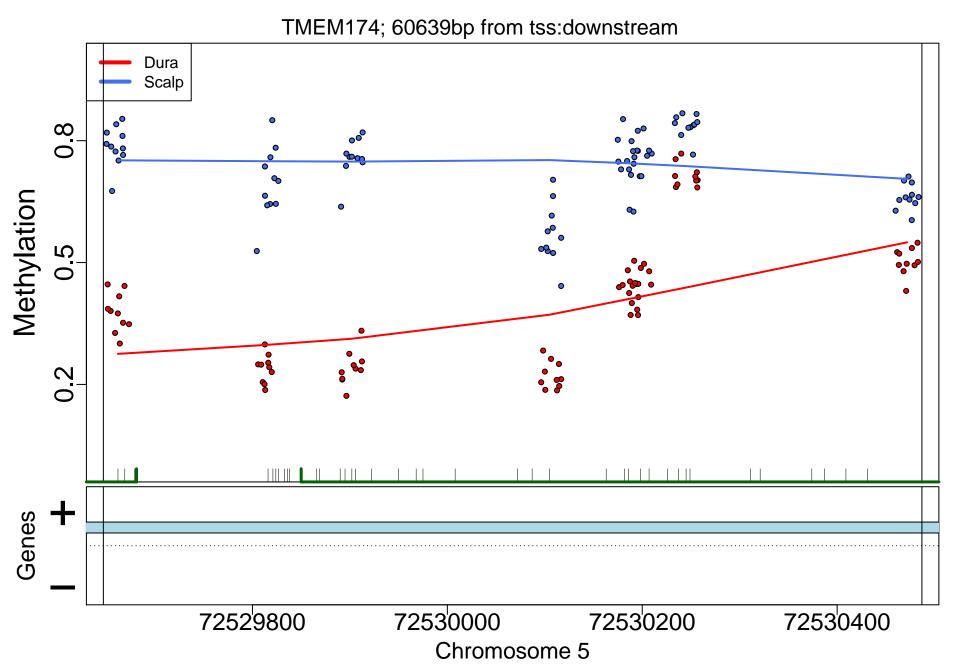


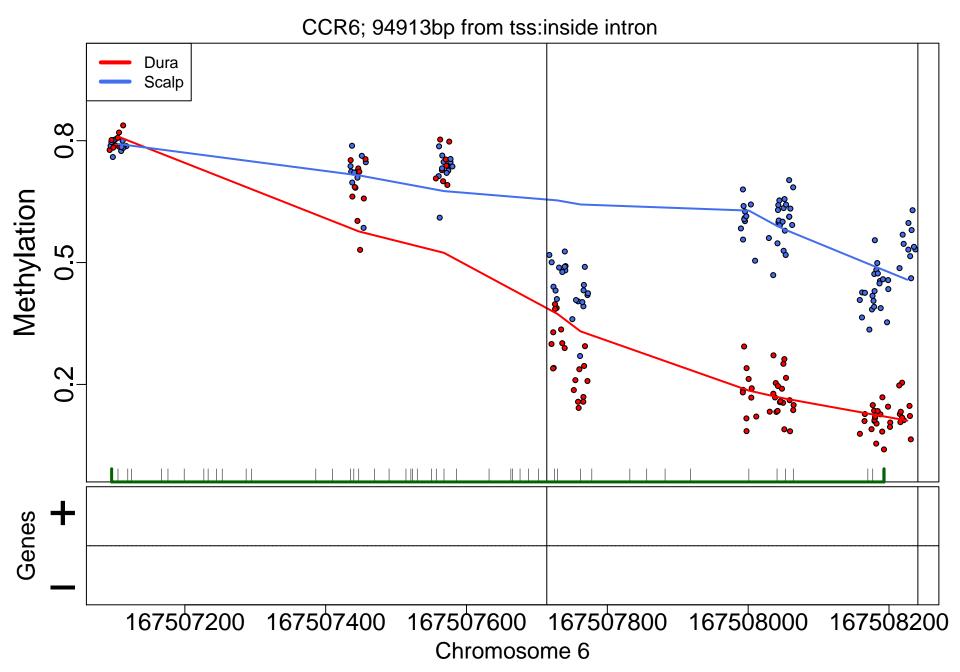


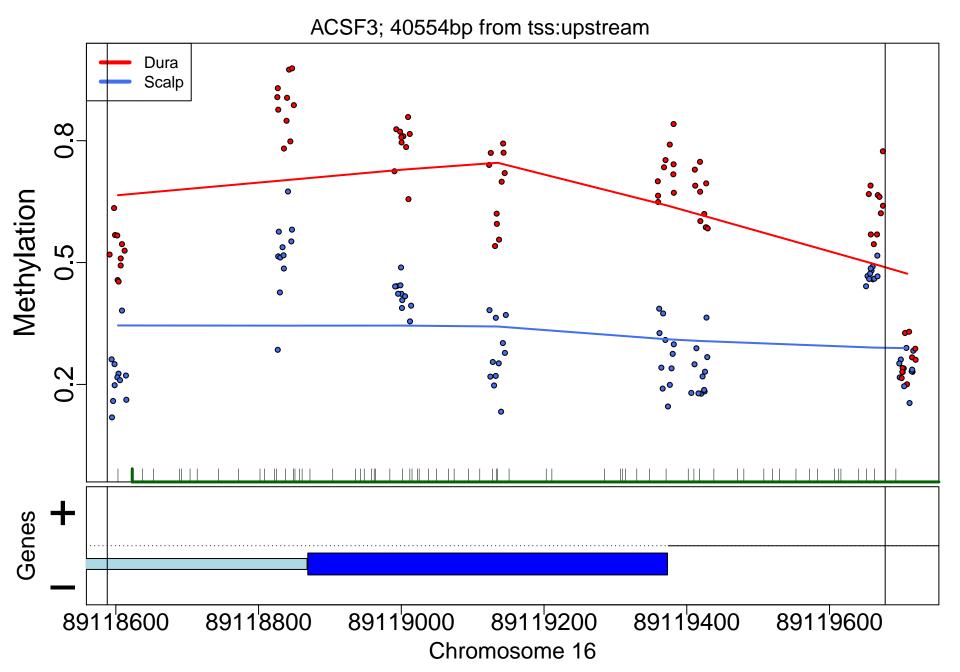


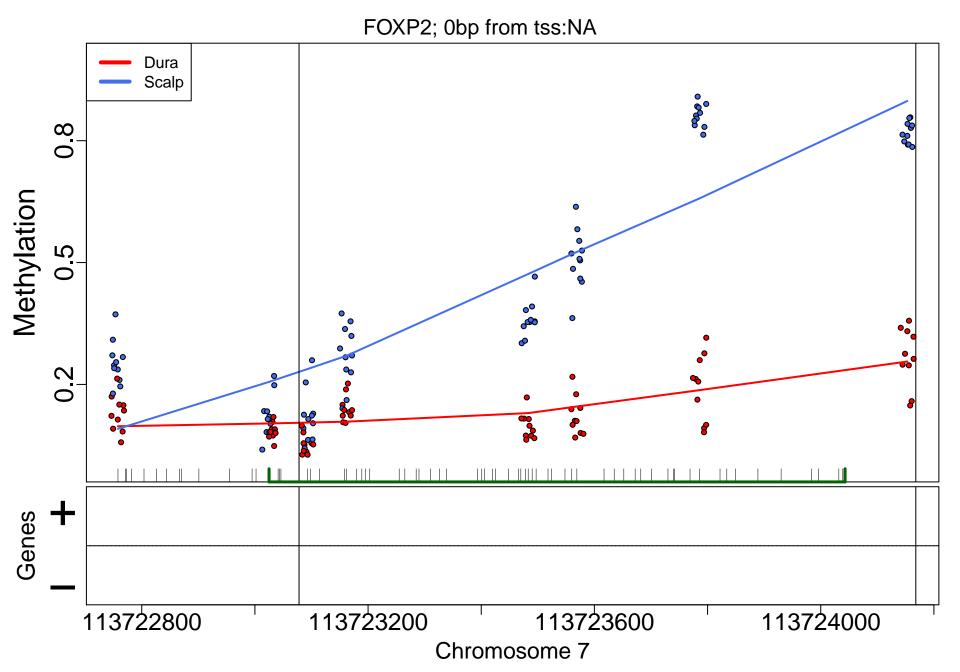


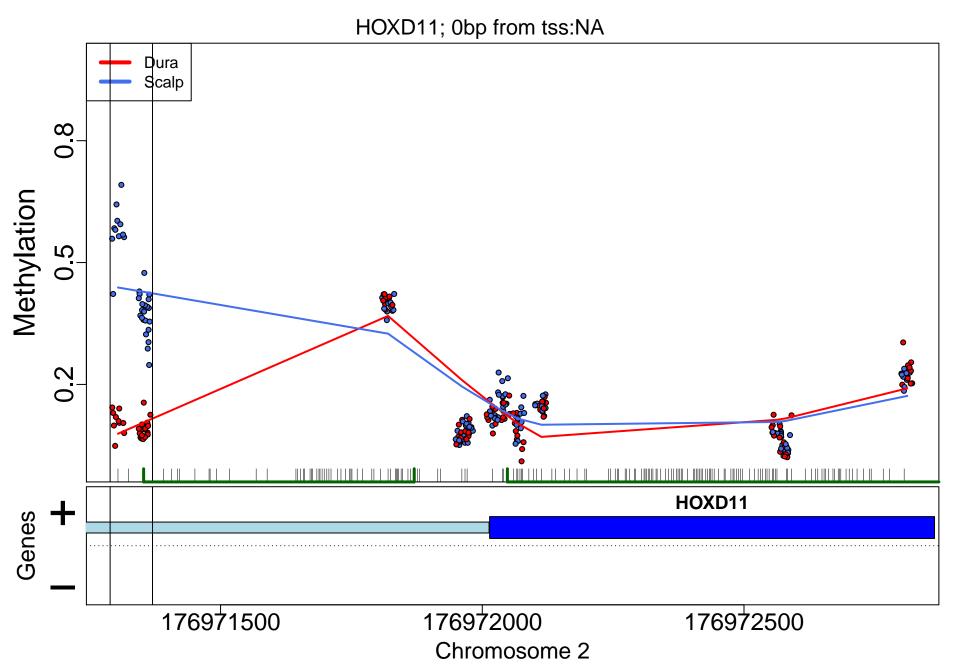


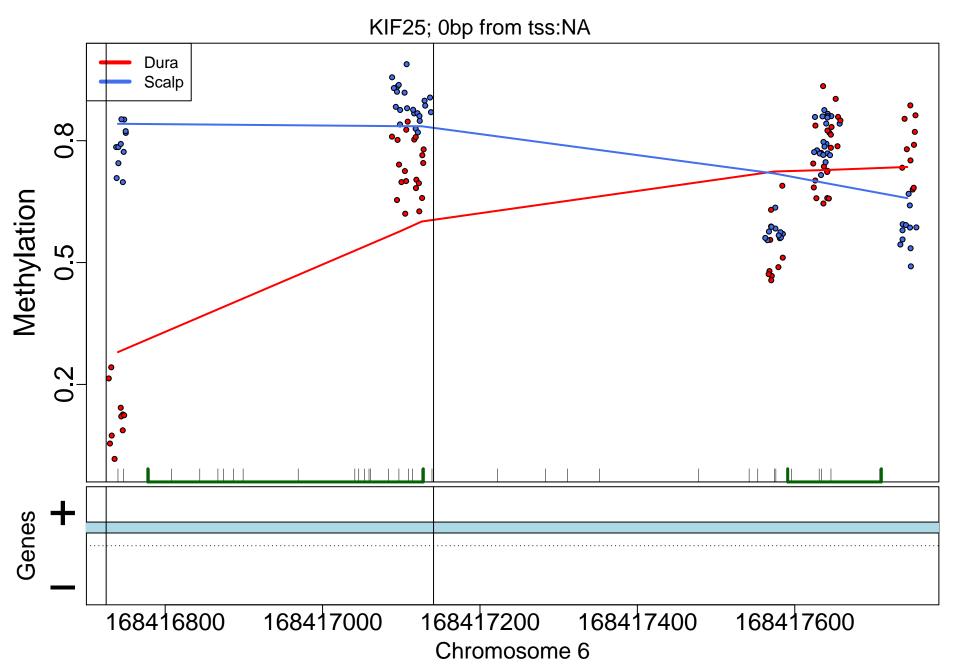


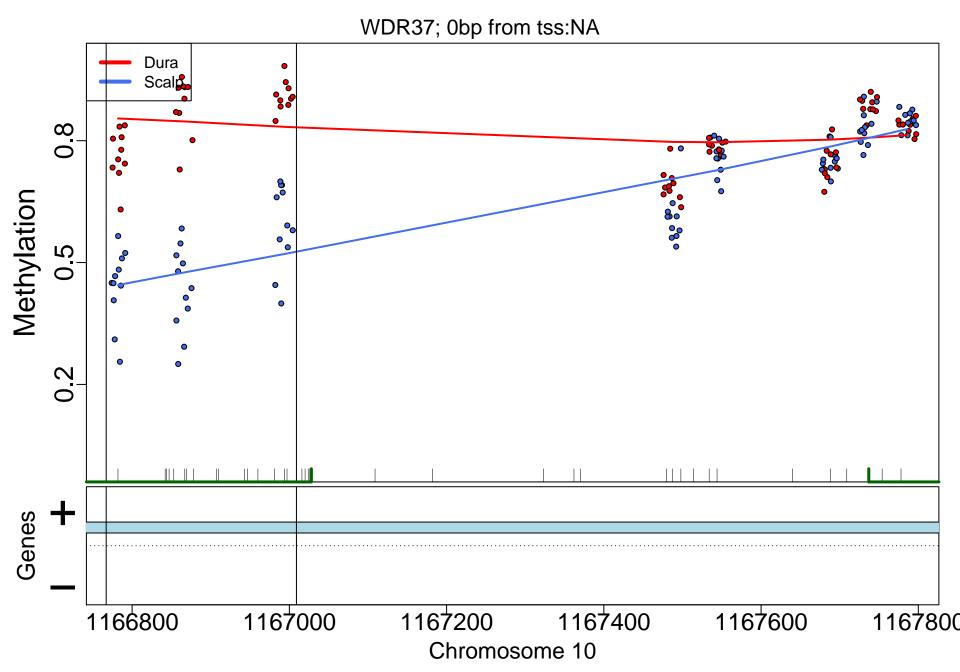


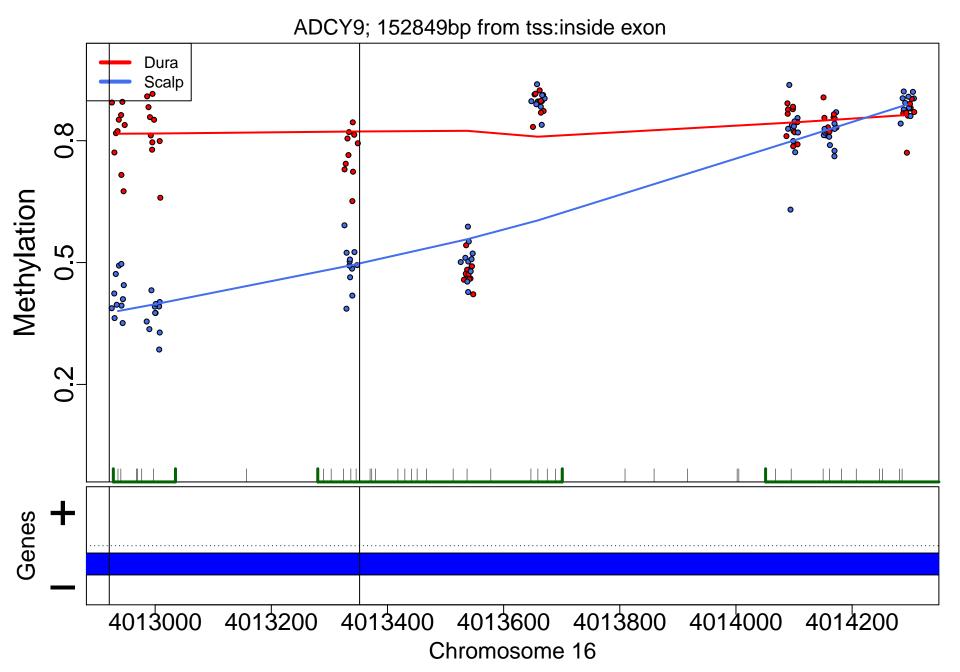


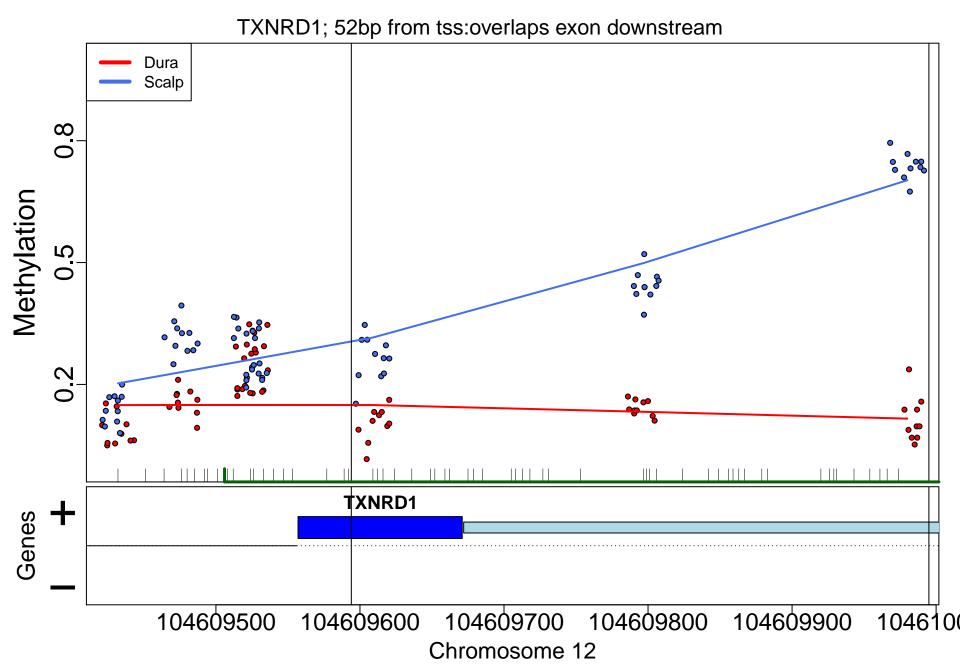


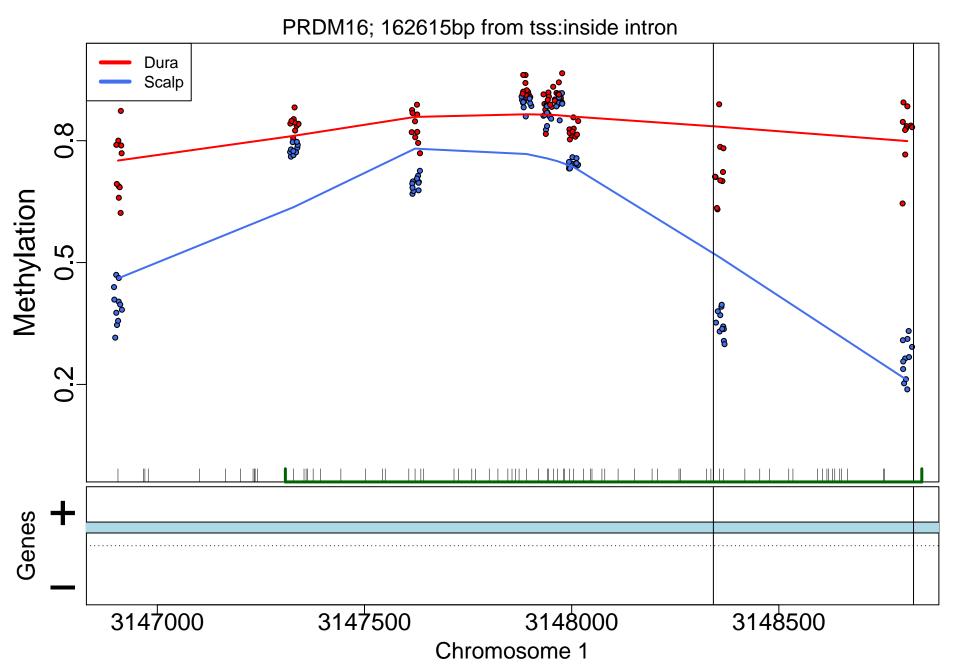


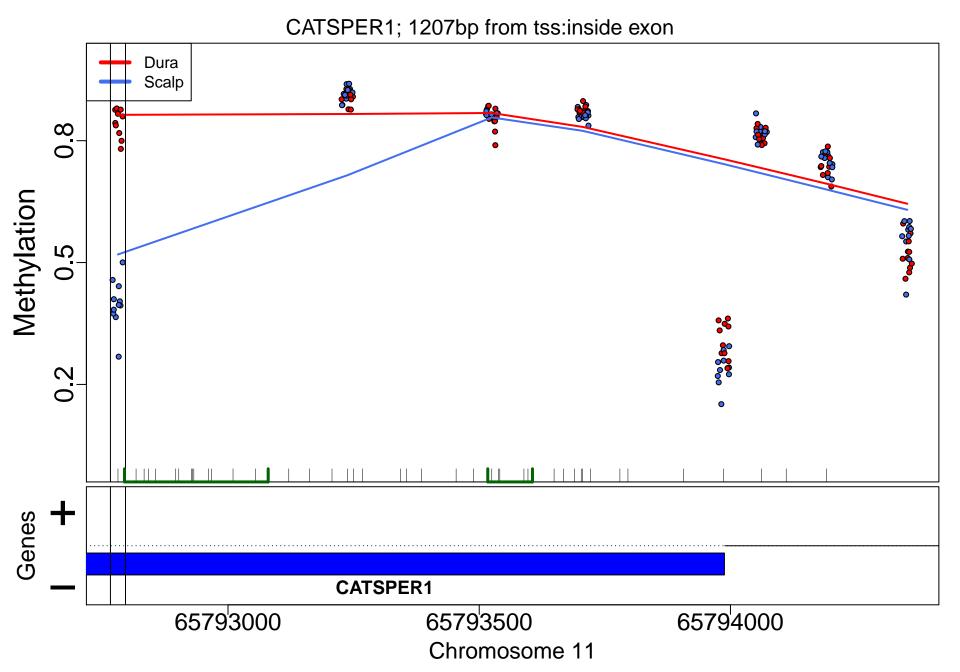


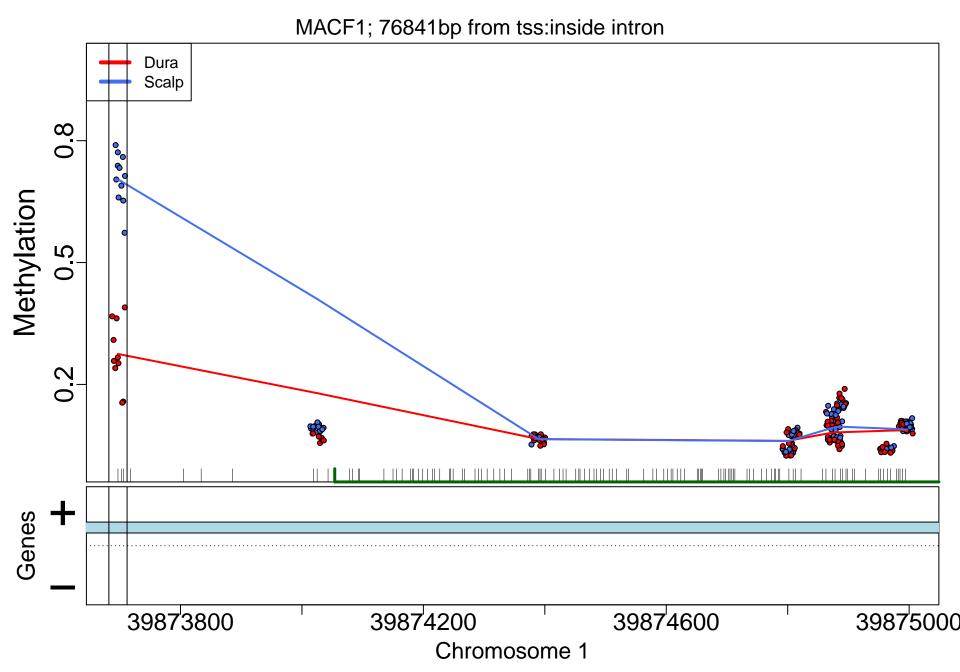


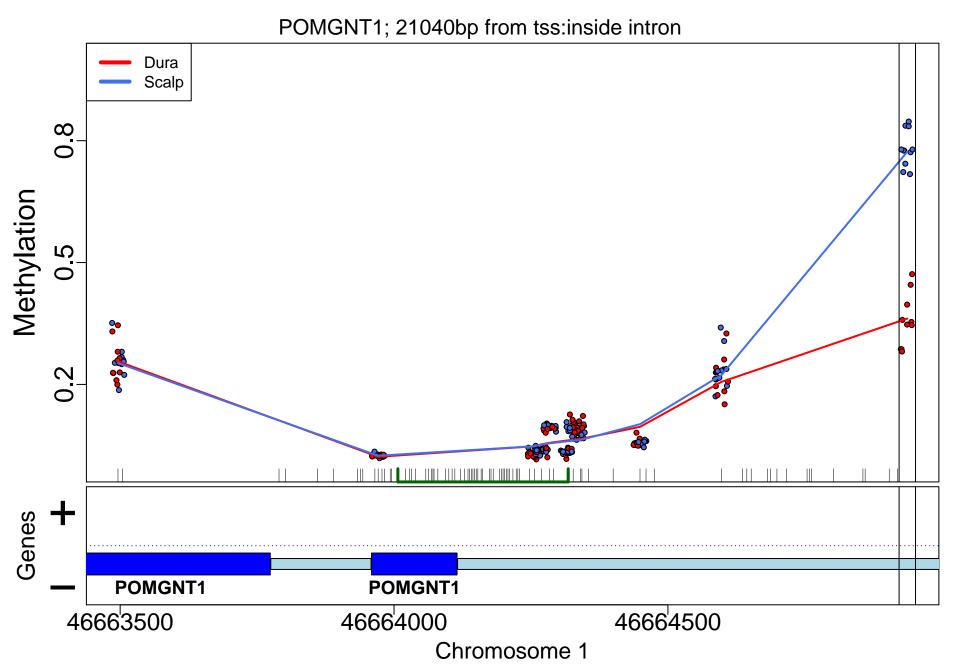


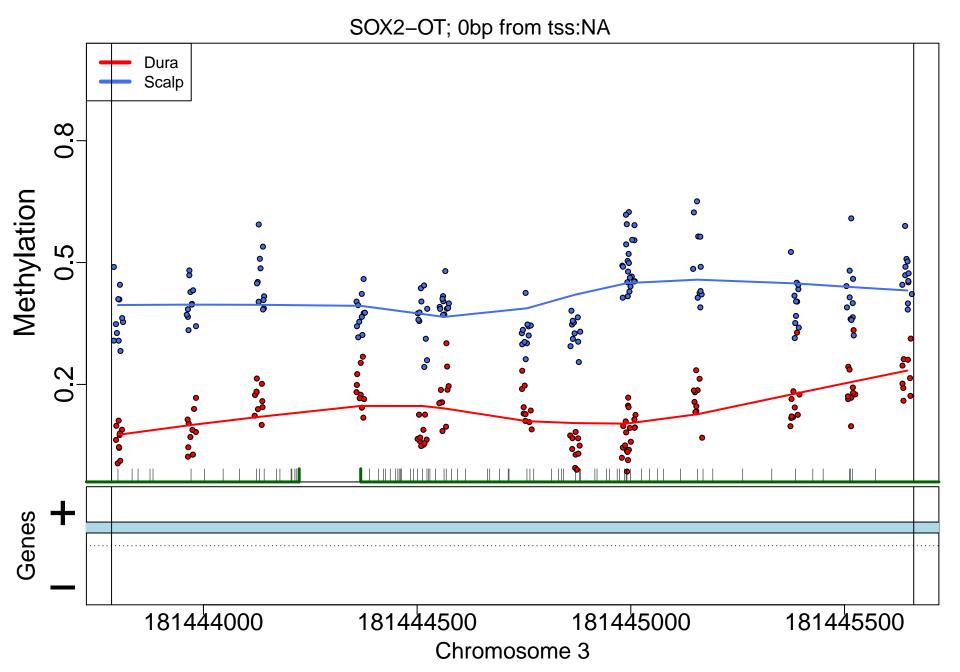




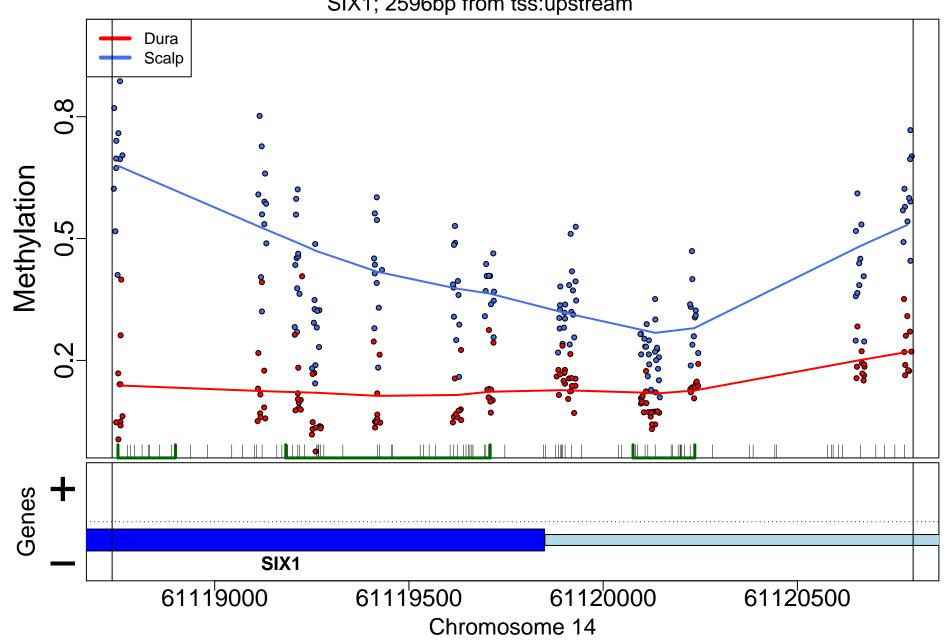


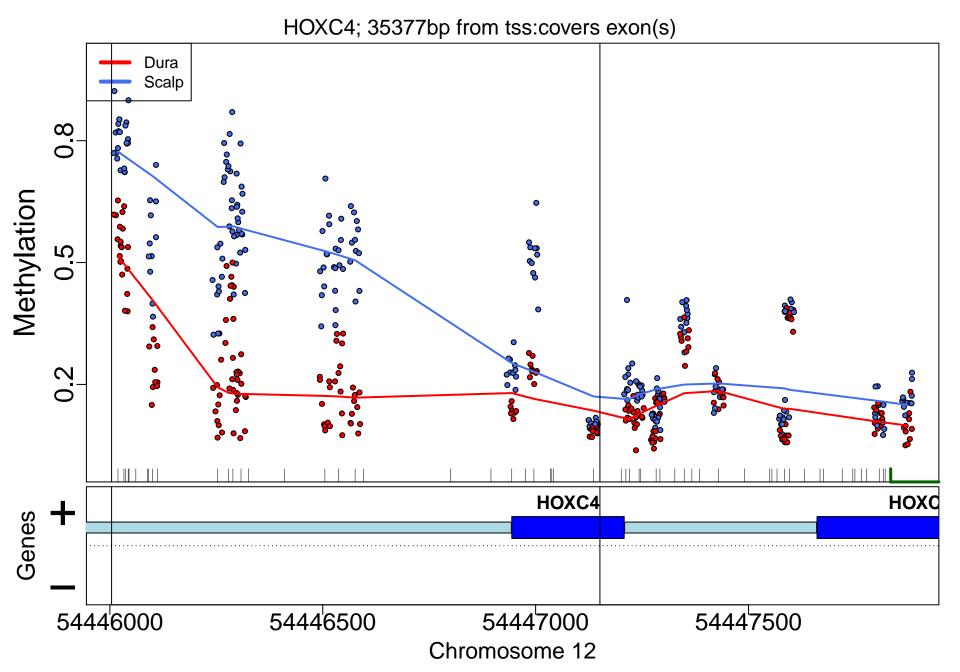


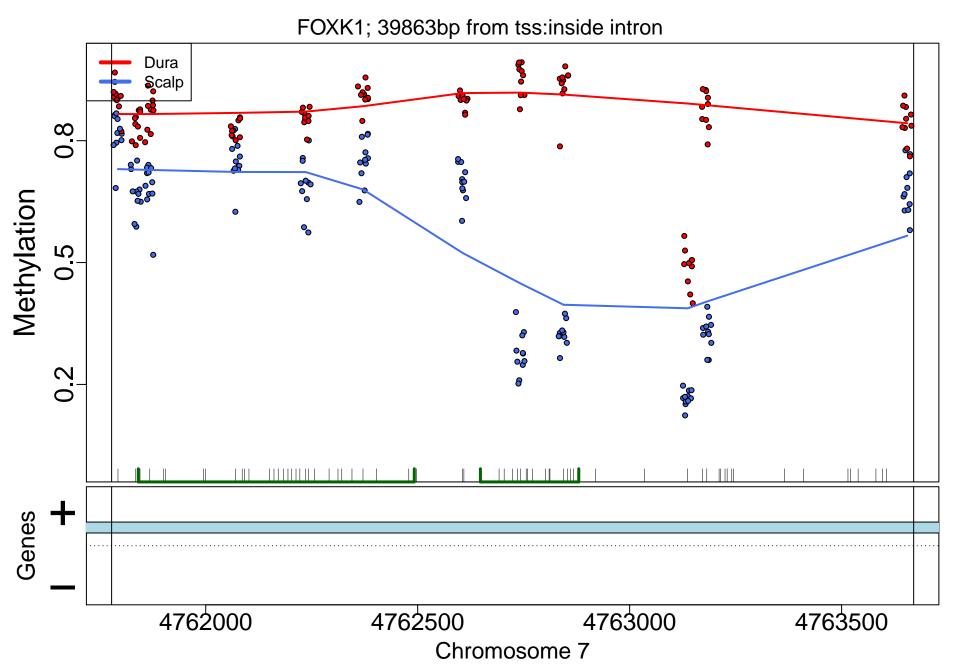


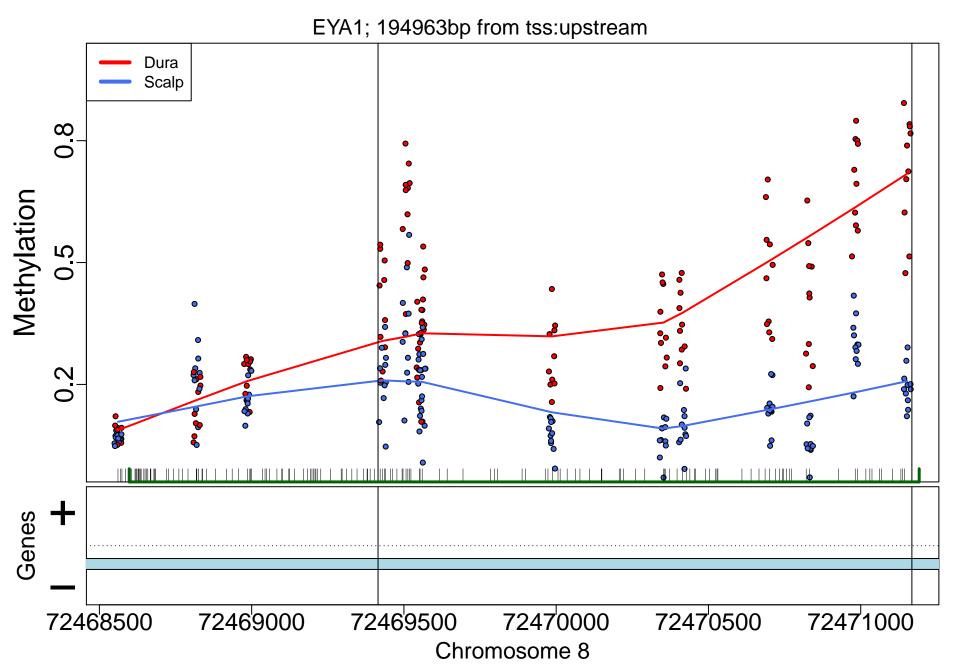


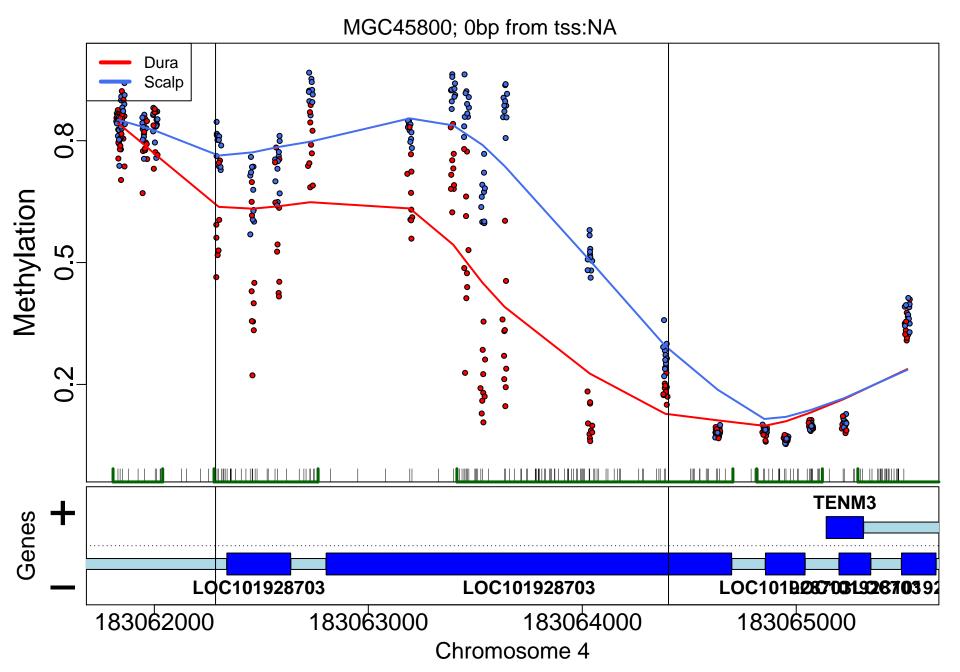
SIX1; 2596bp from tss:upstream

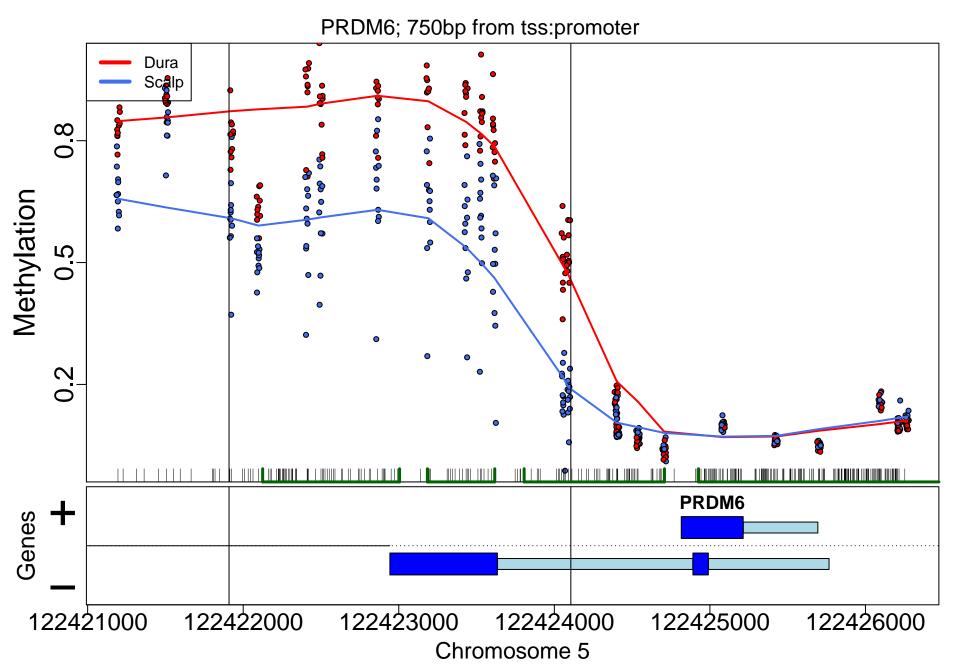


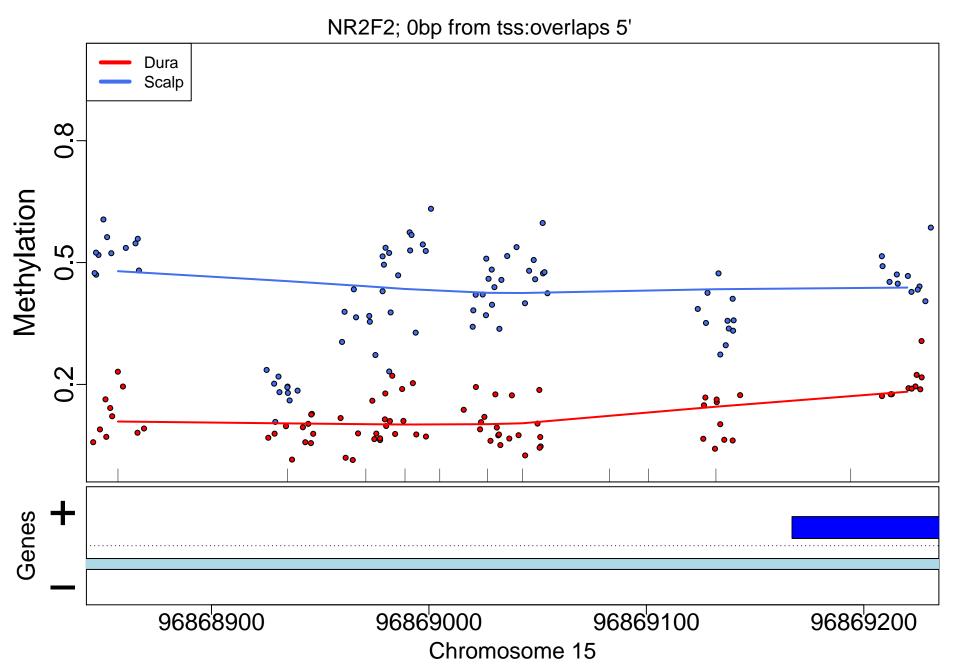


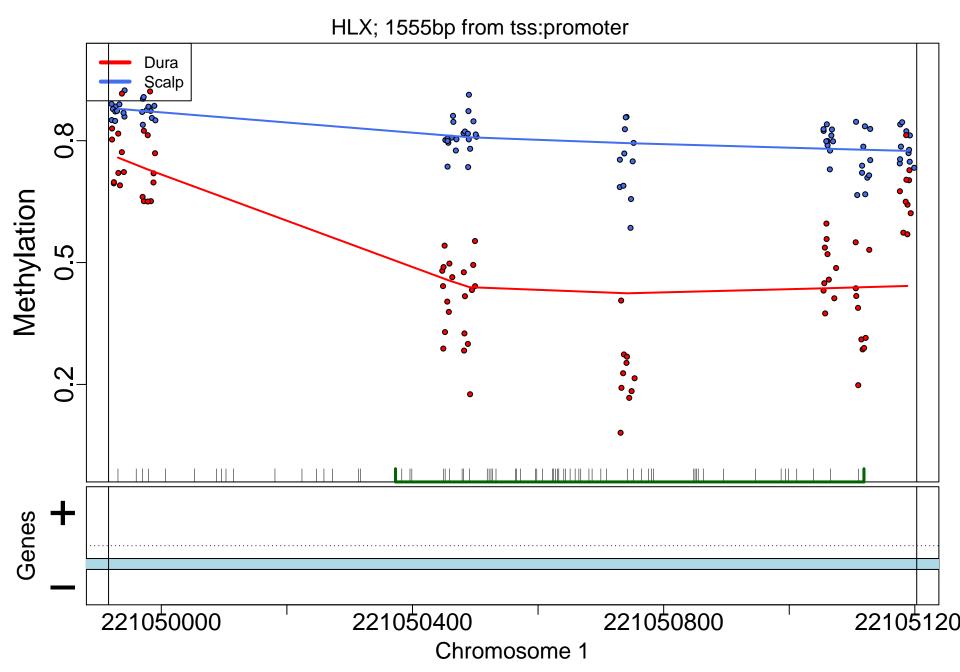


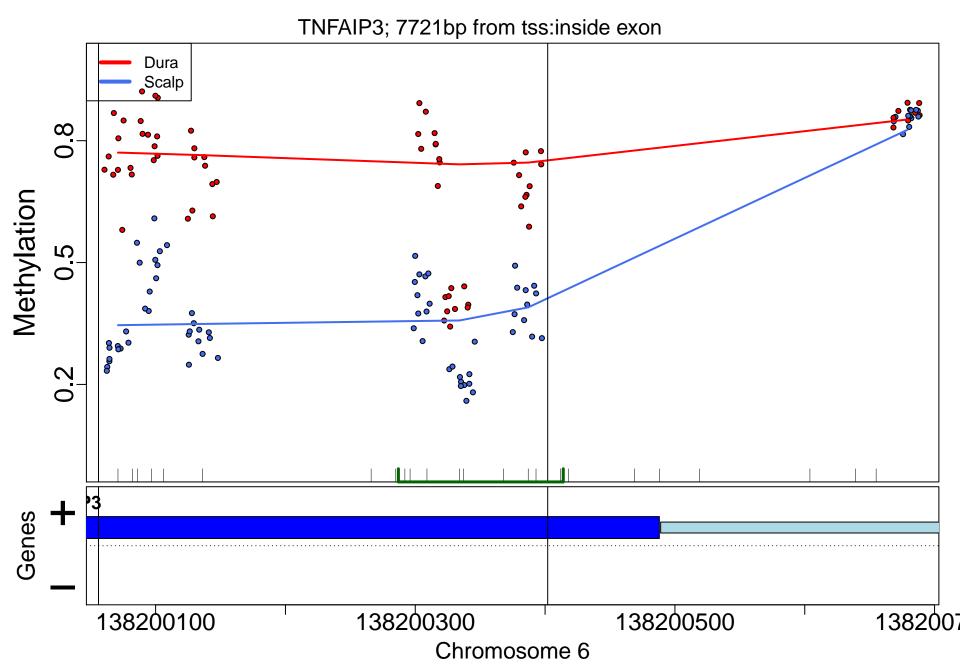


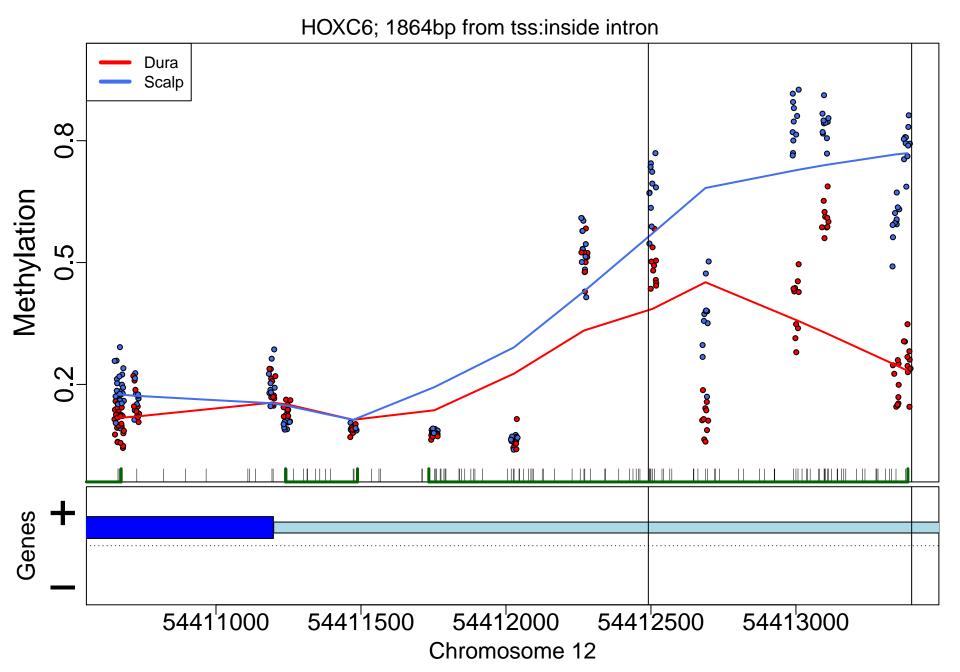


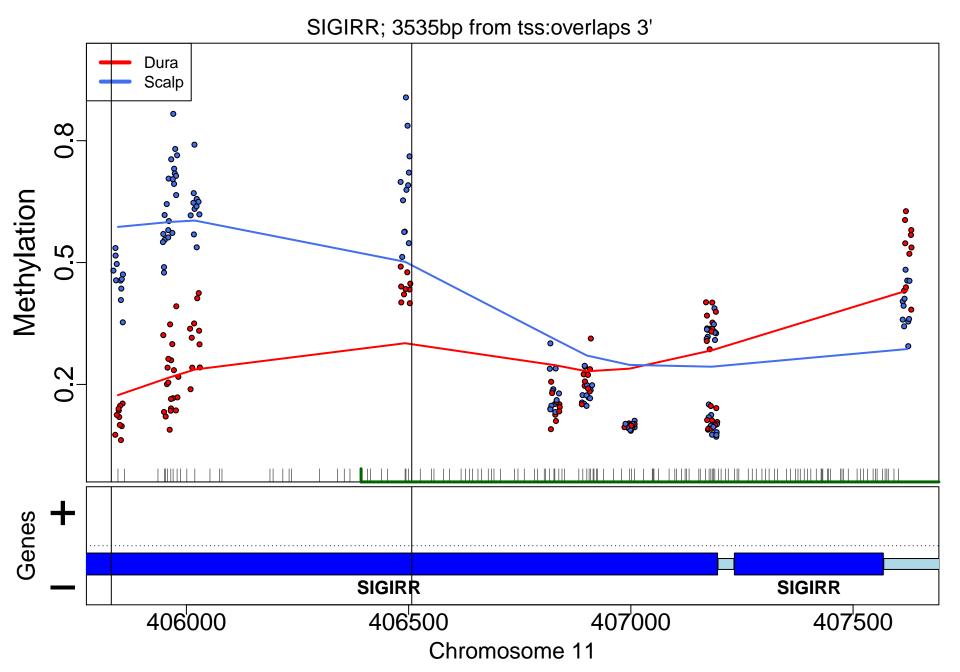


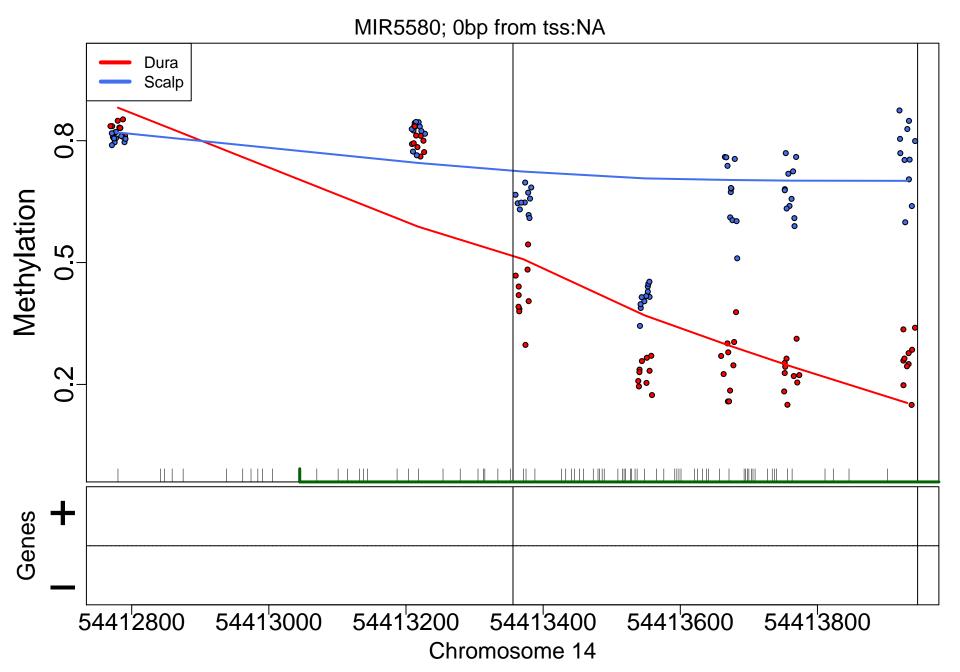


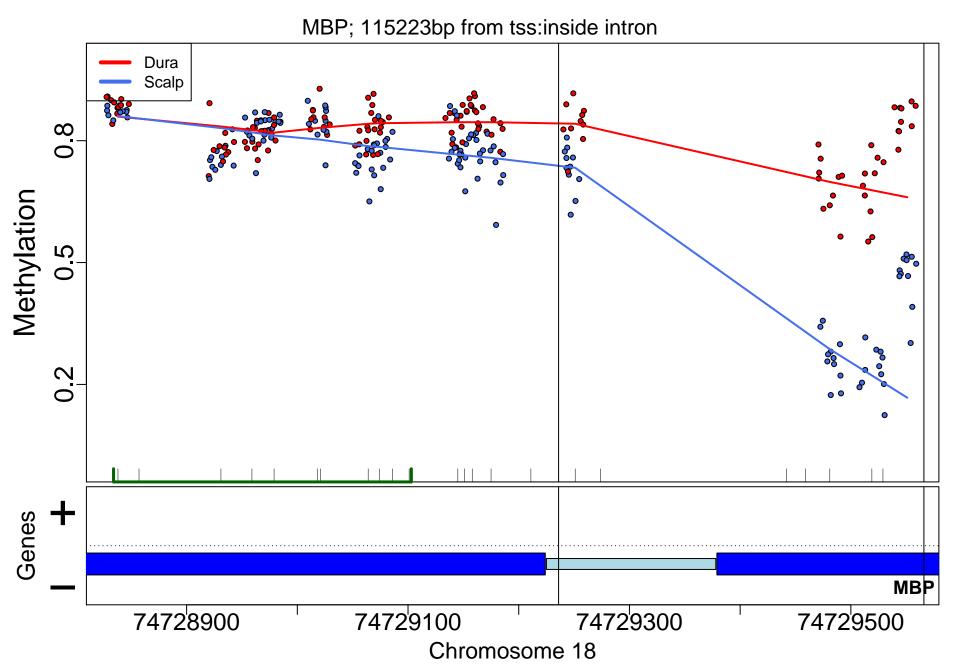


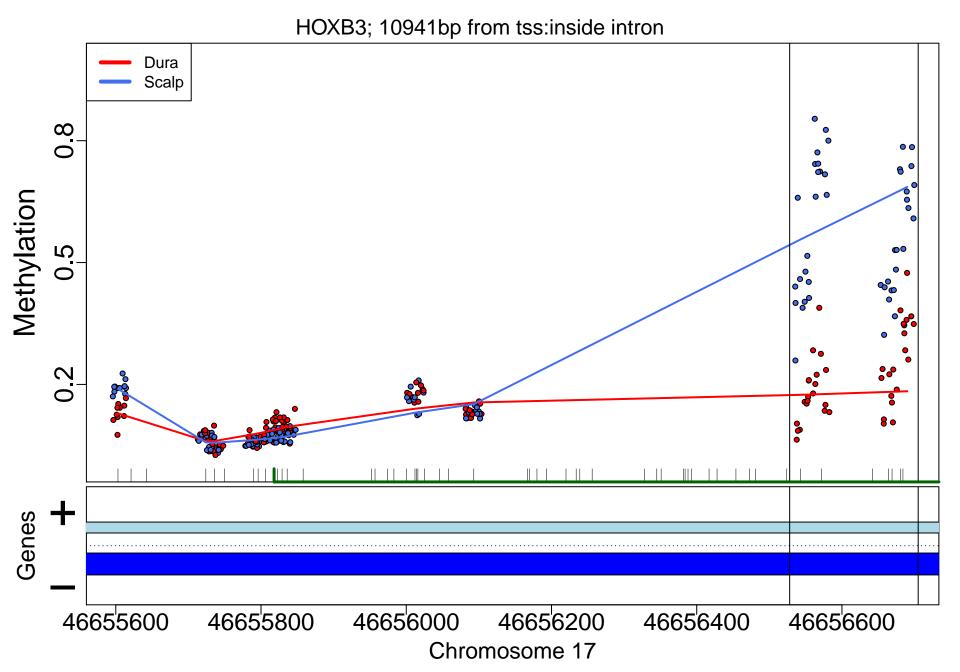


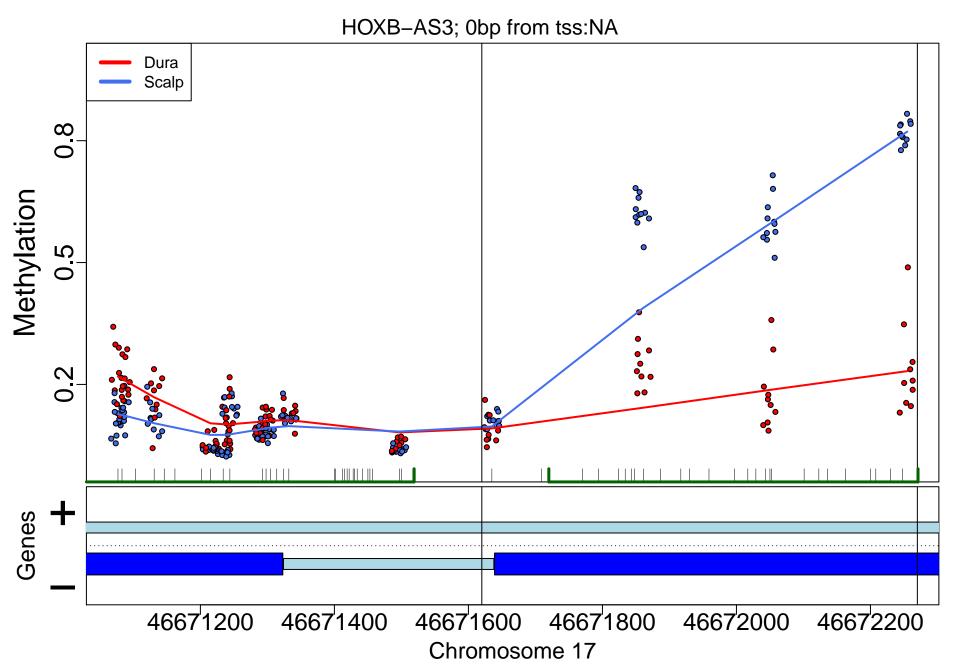


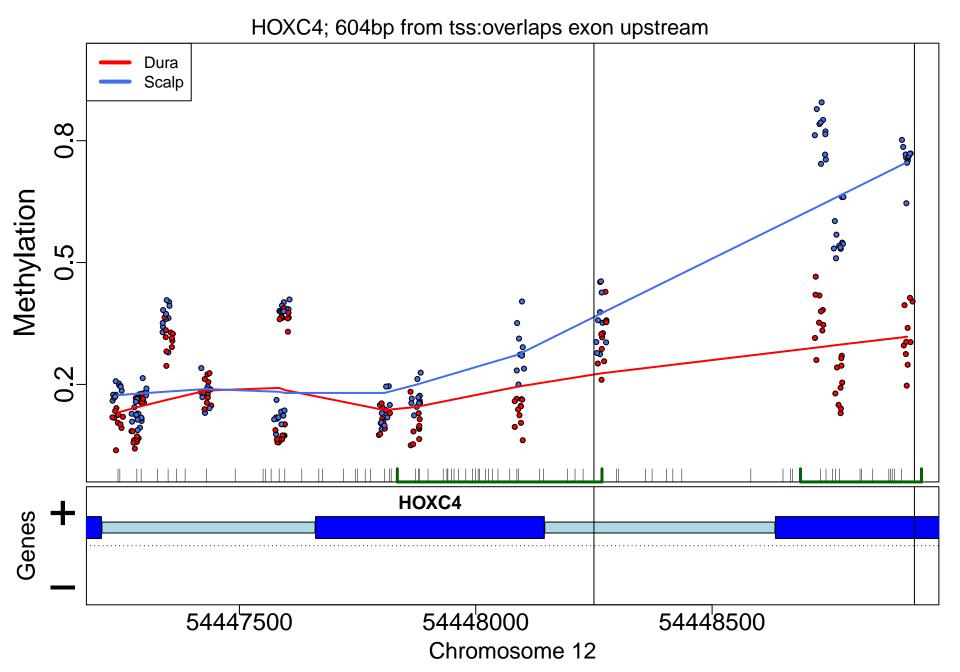


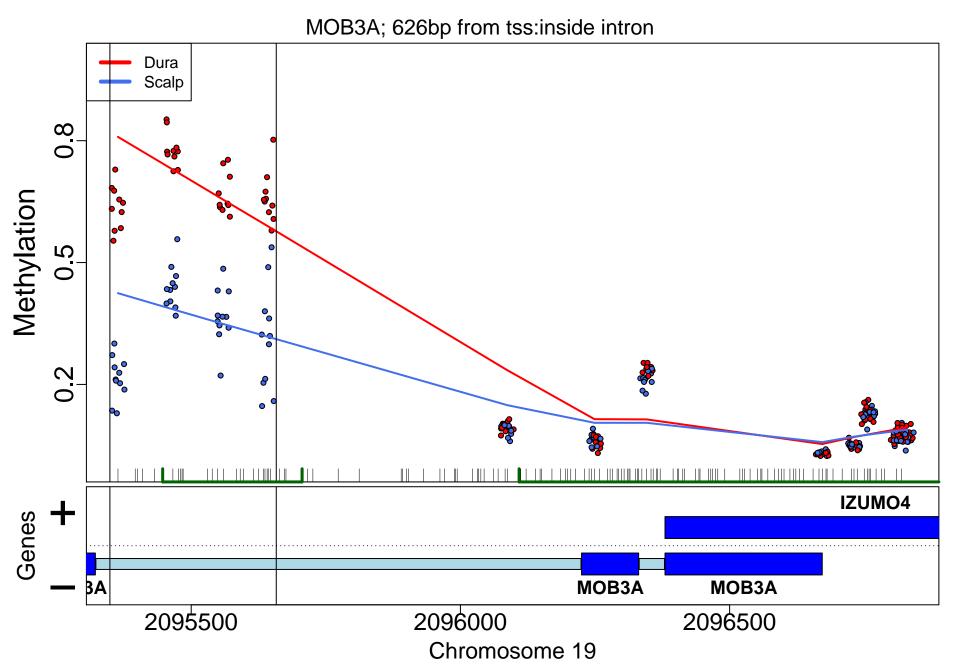


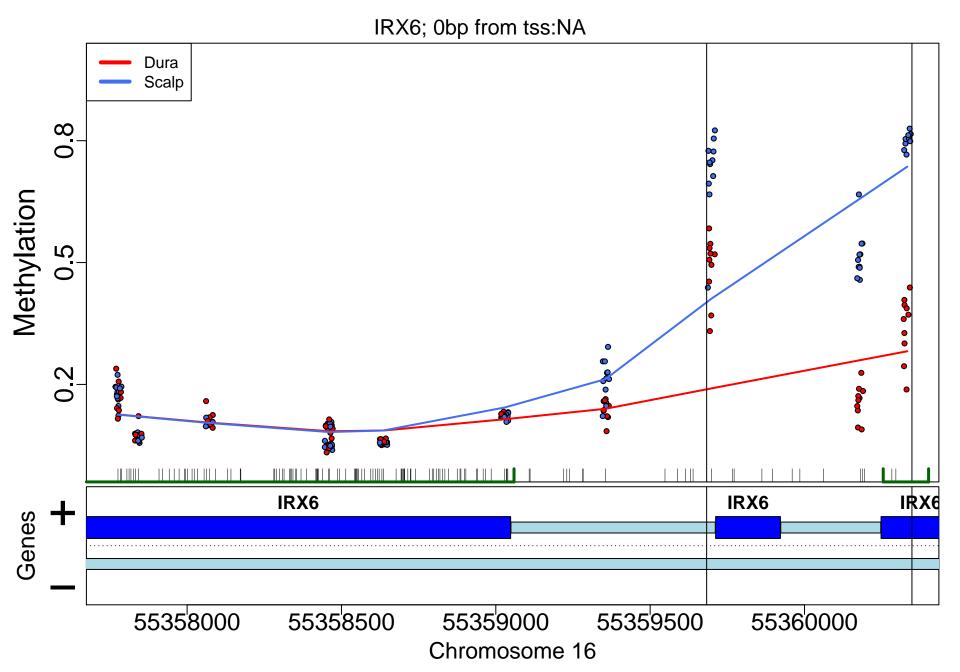


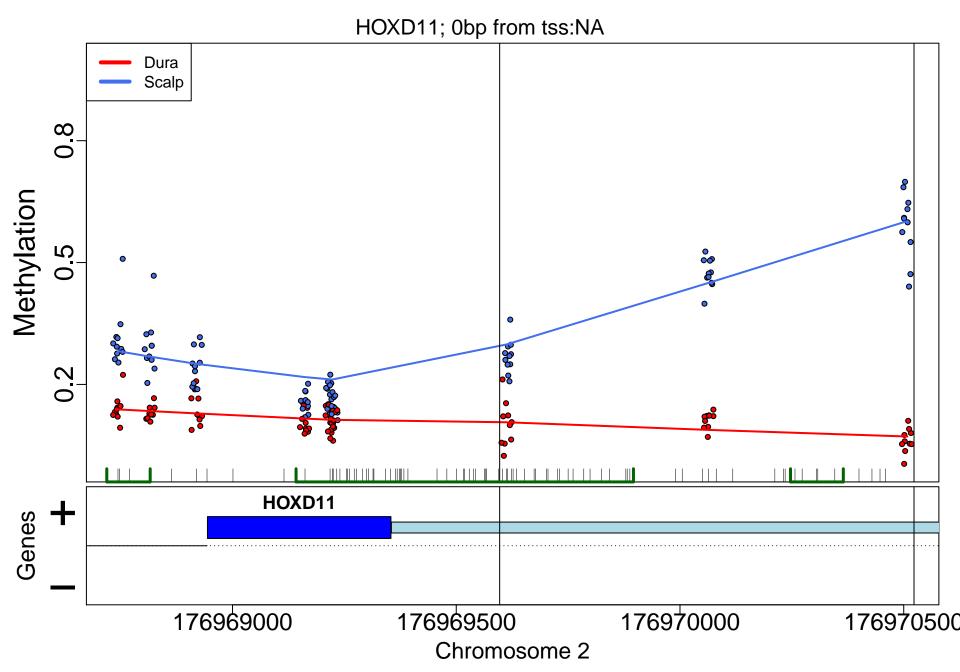


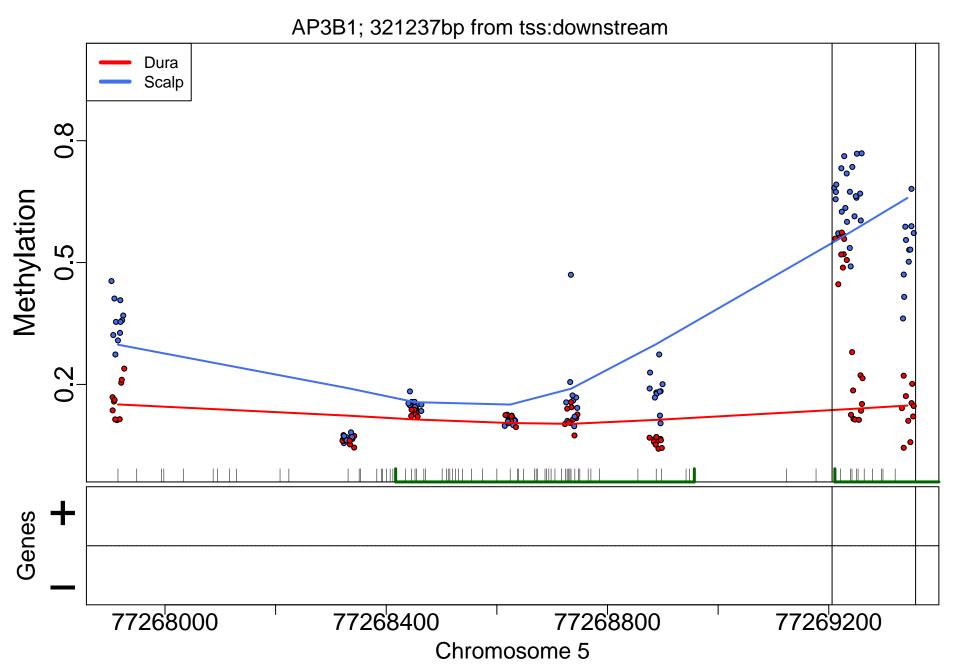


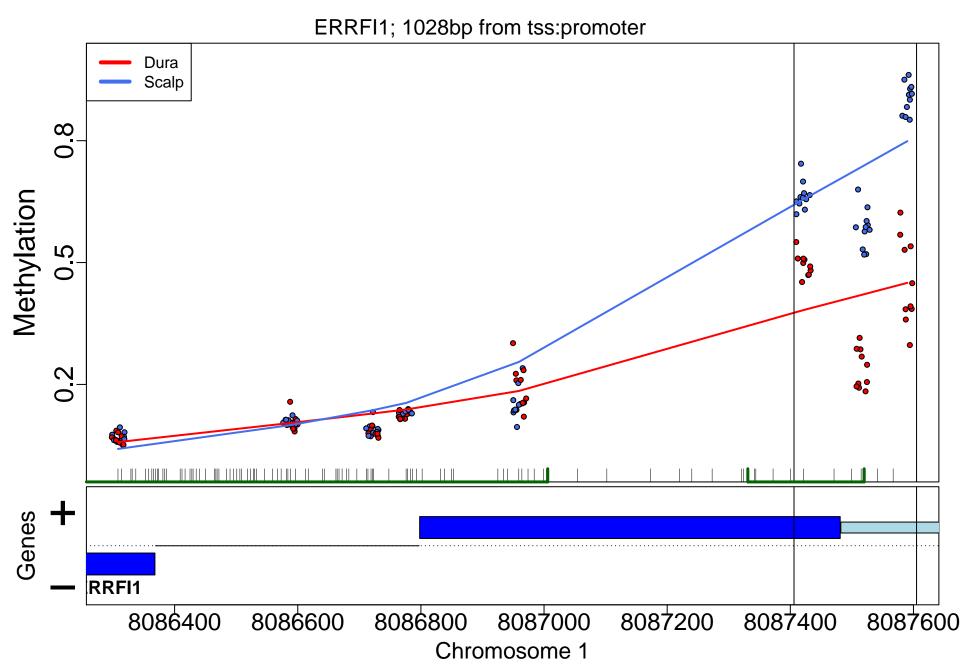


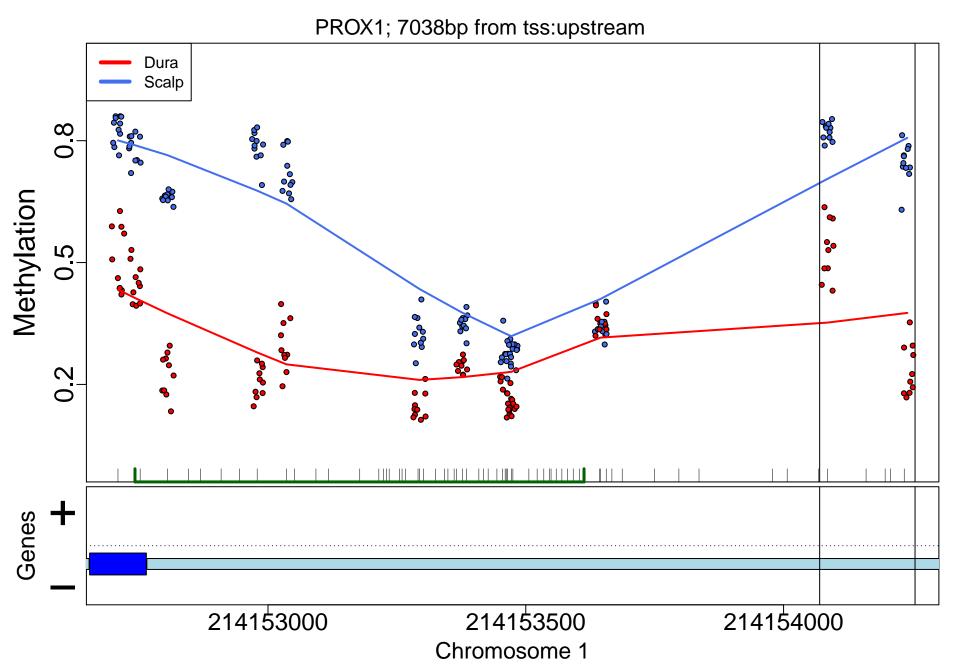


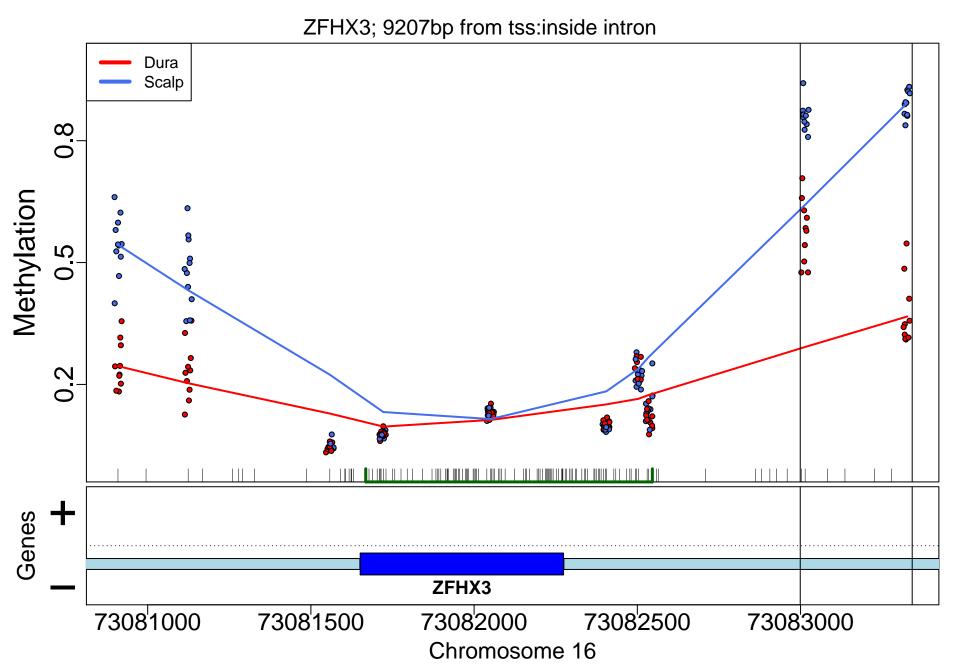


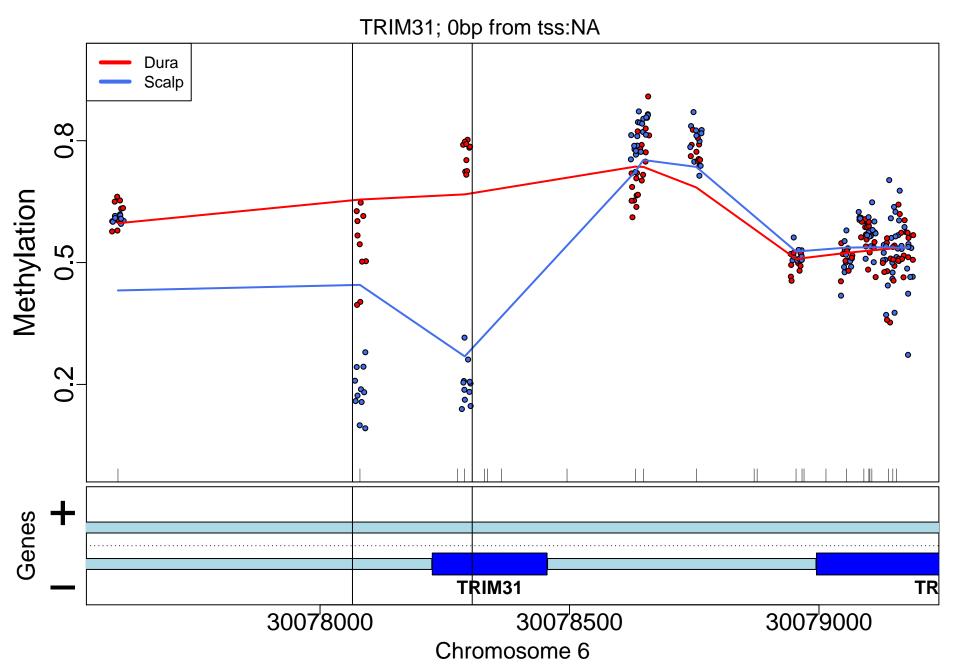


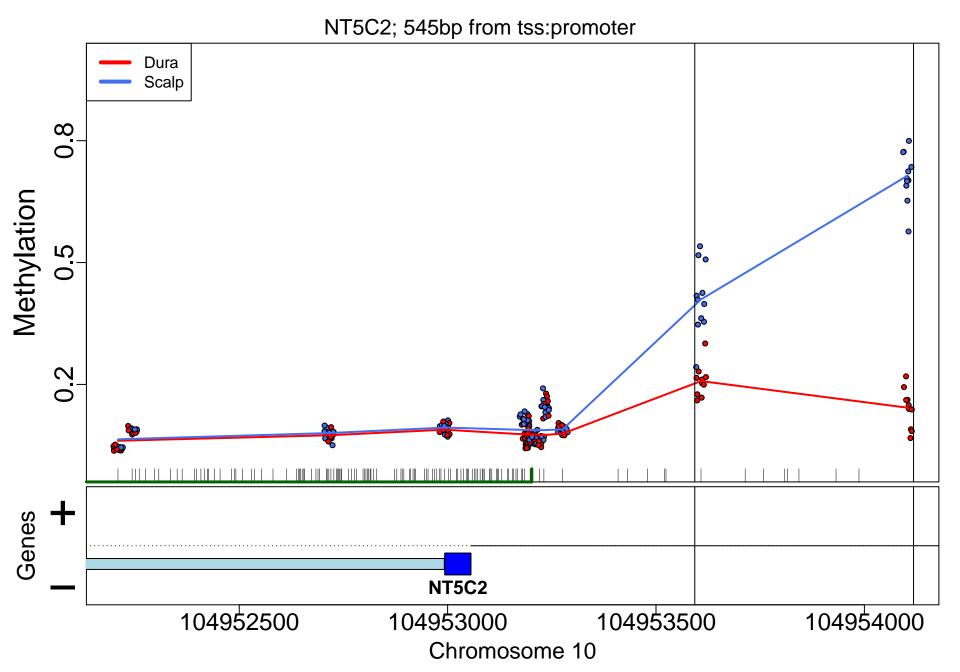


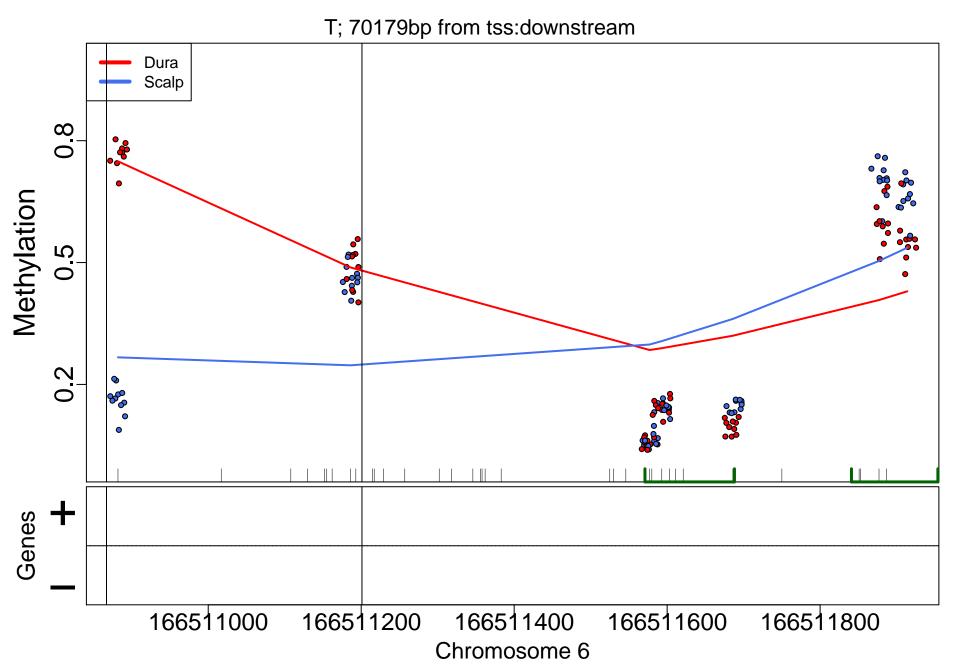


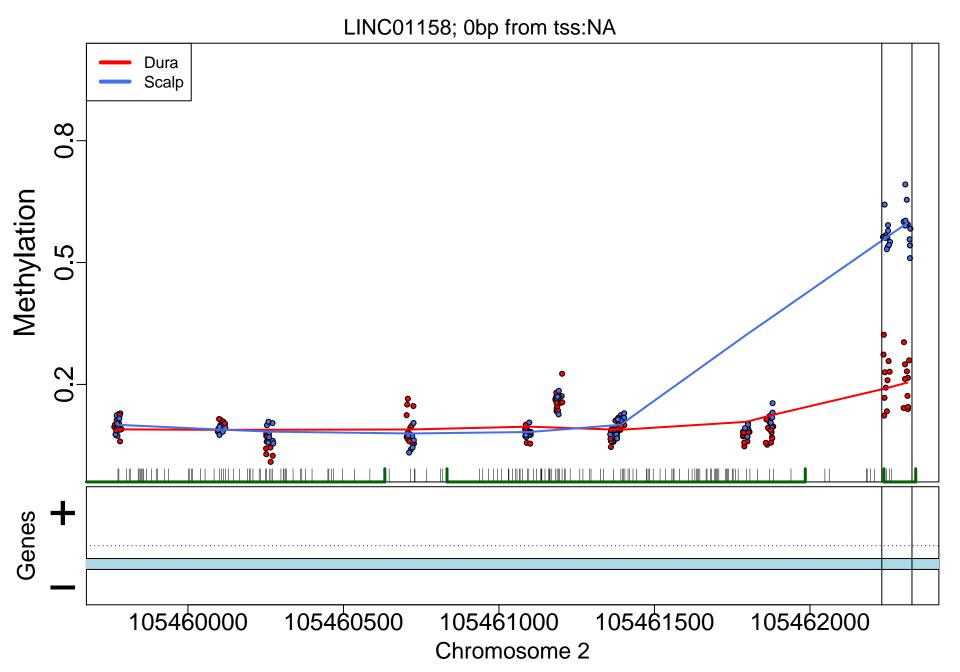




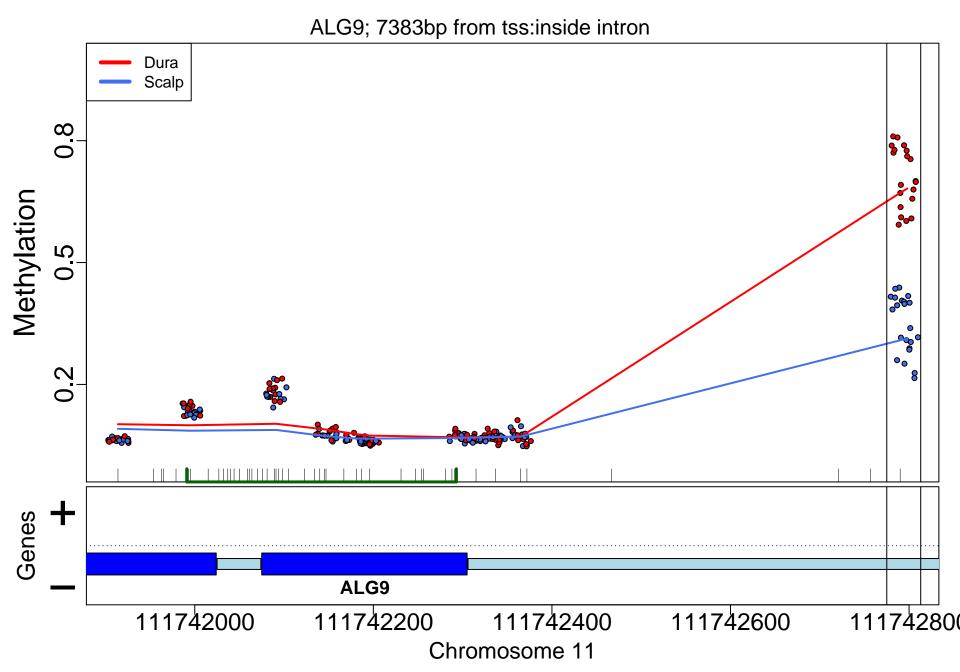


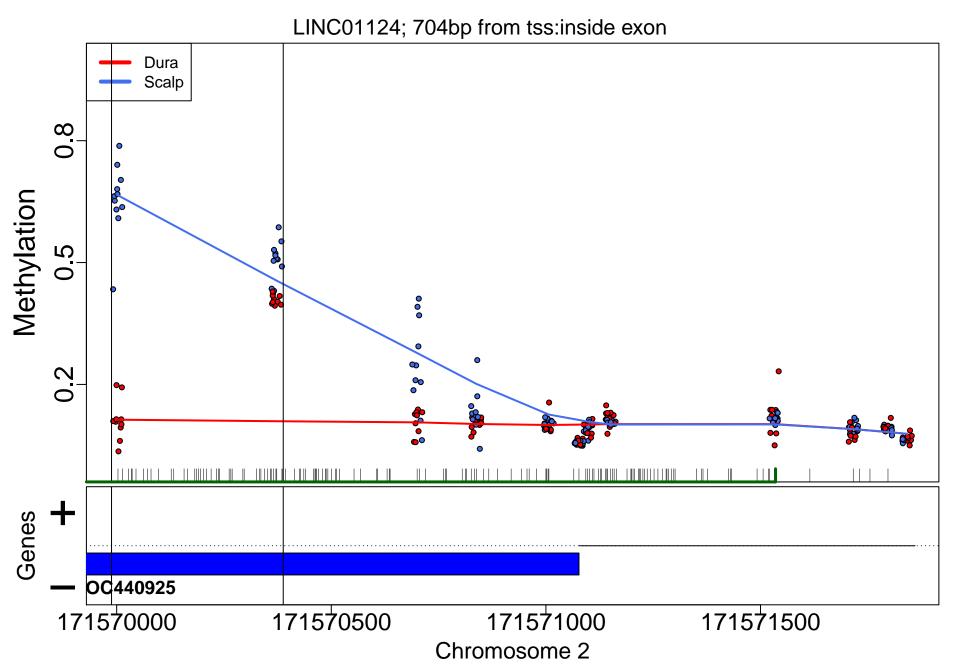


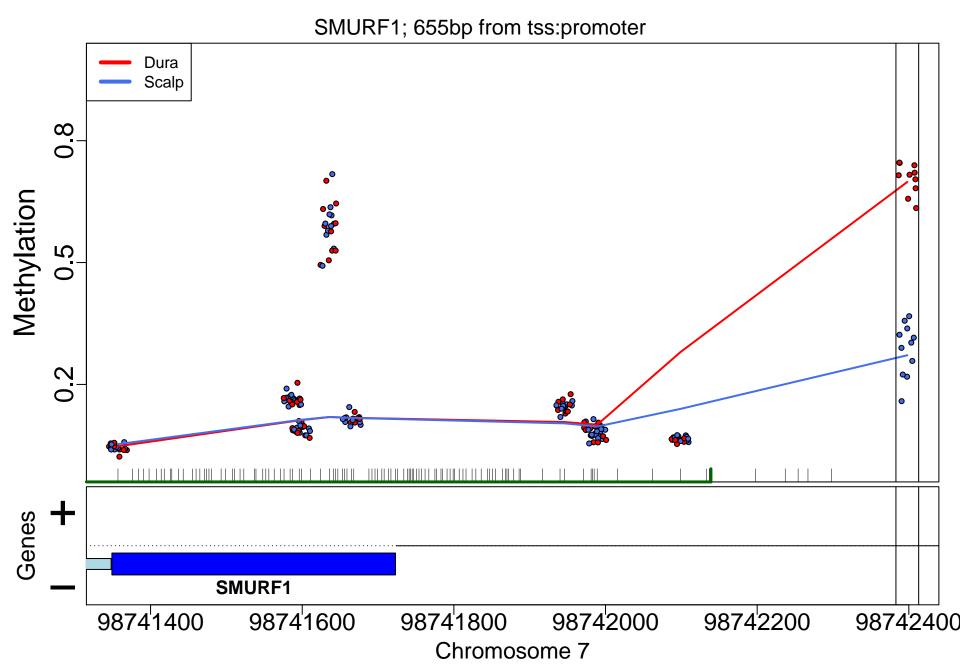


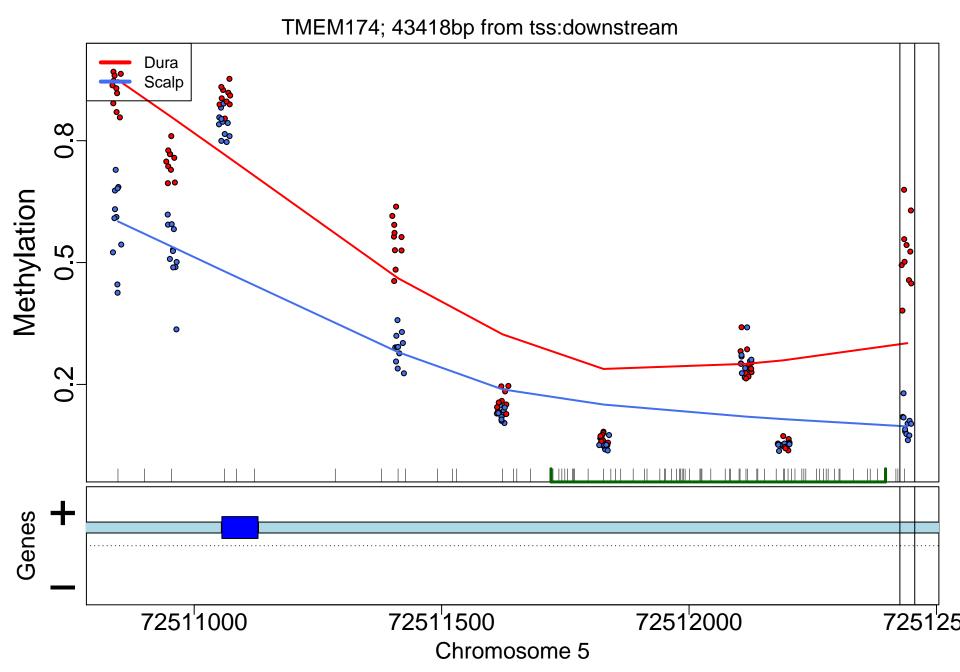


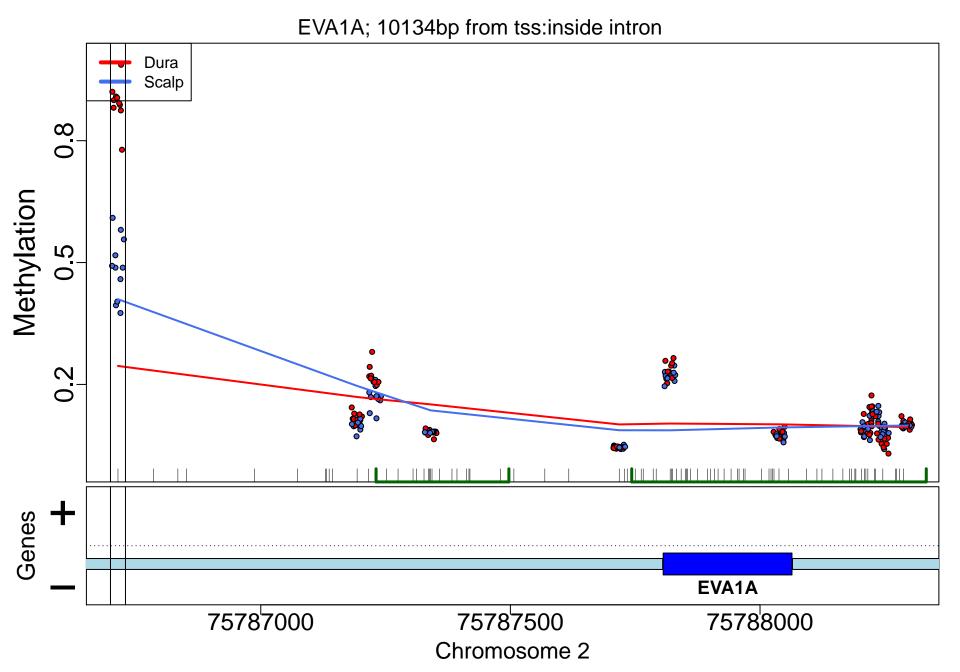
ZFHX4-AS1; 0bp from tss:NA Dura Scalp <u>ω</u>_ **%**% Methylation 0.2 ZF Genes 77593000 77593400 77593600 77593200 Chromosome 8

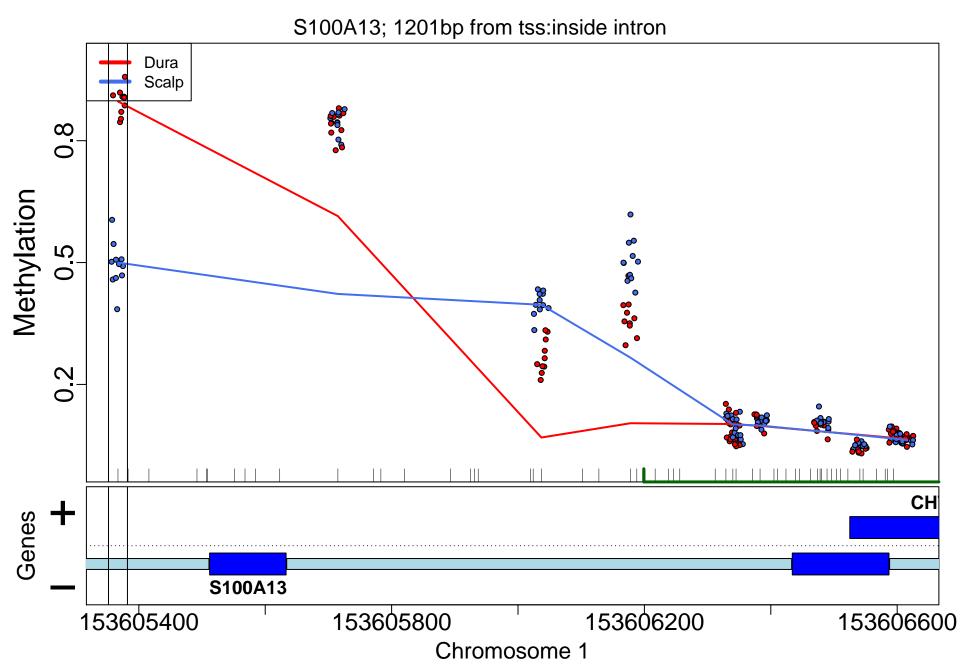


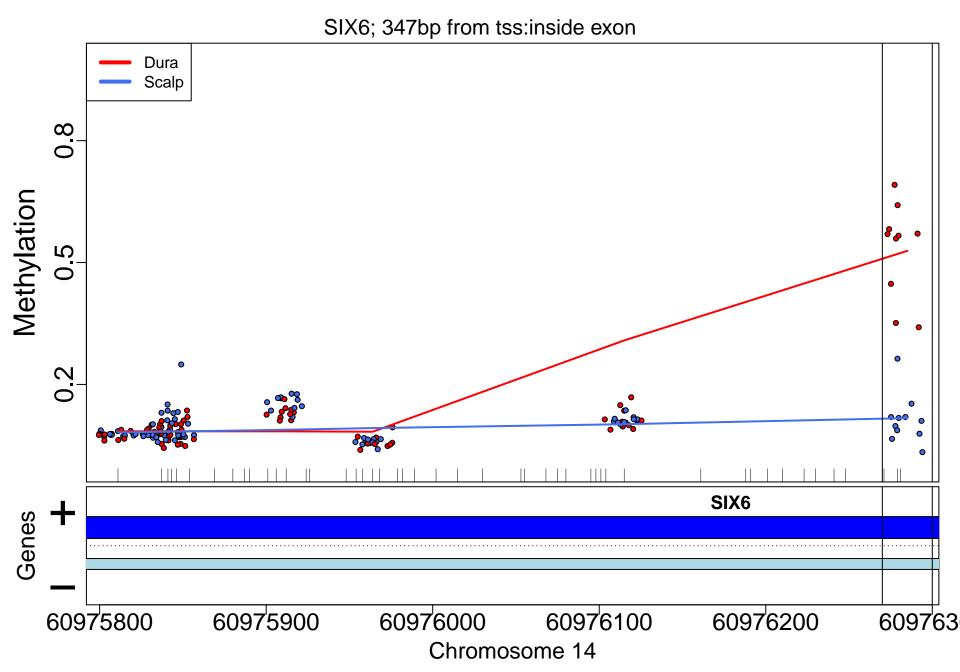


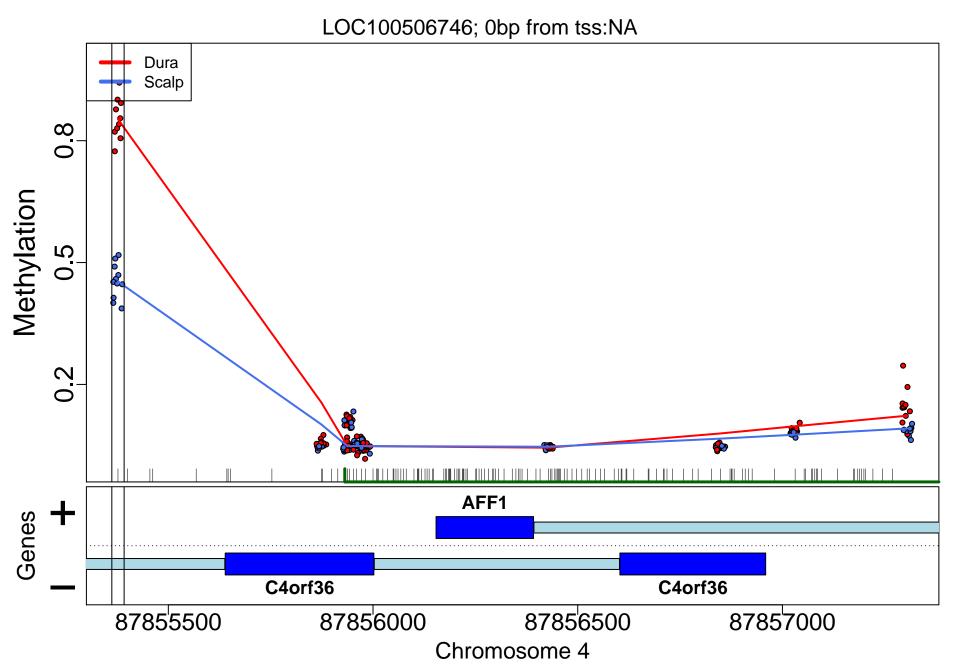


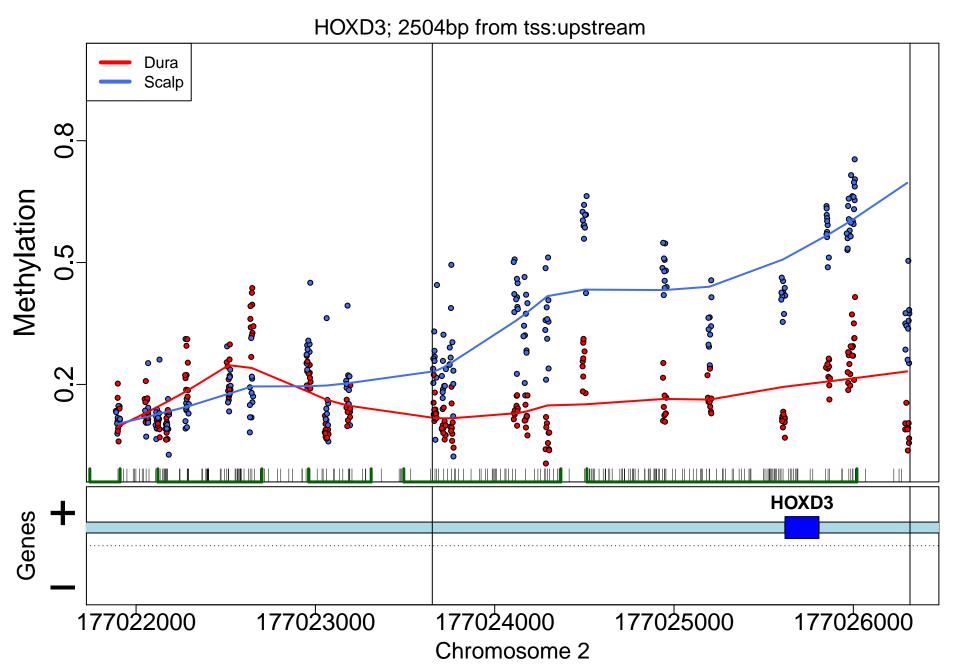




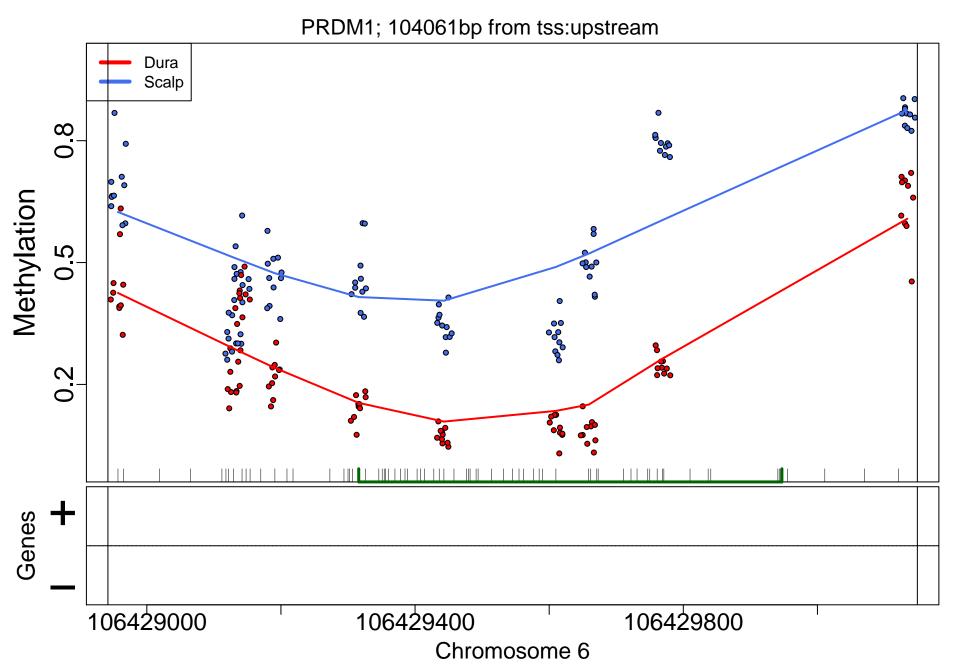


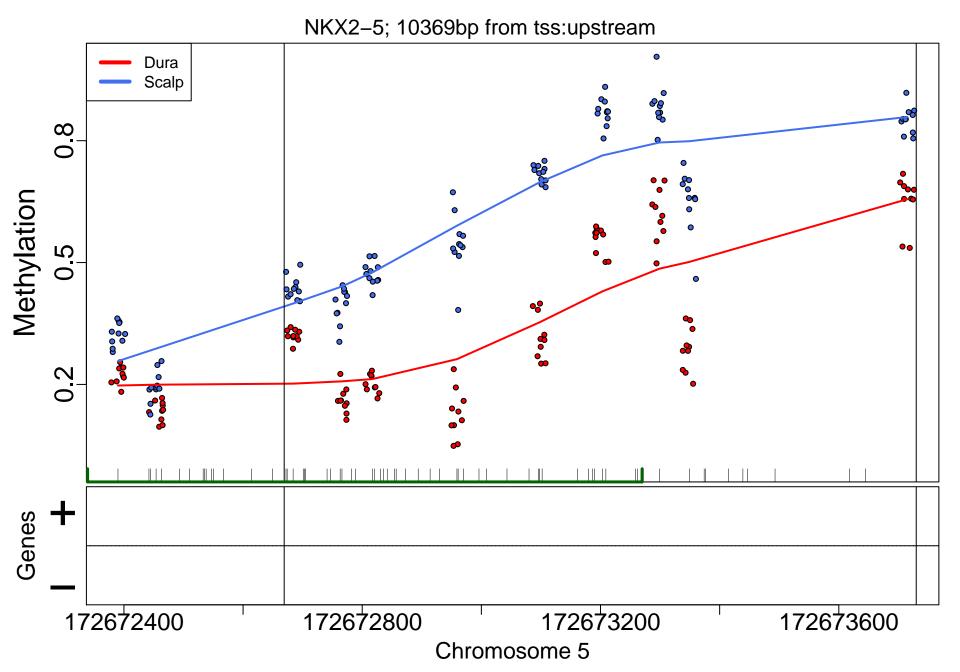


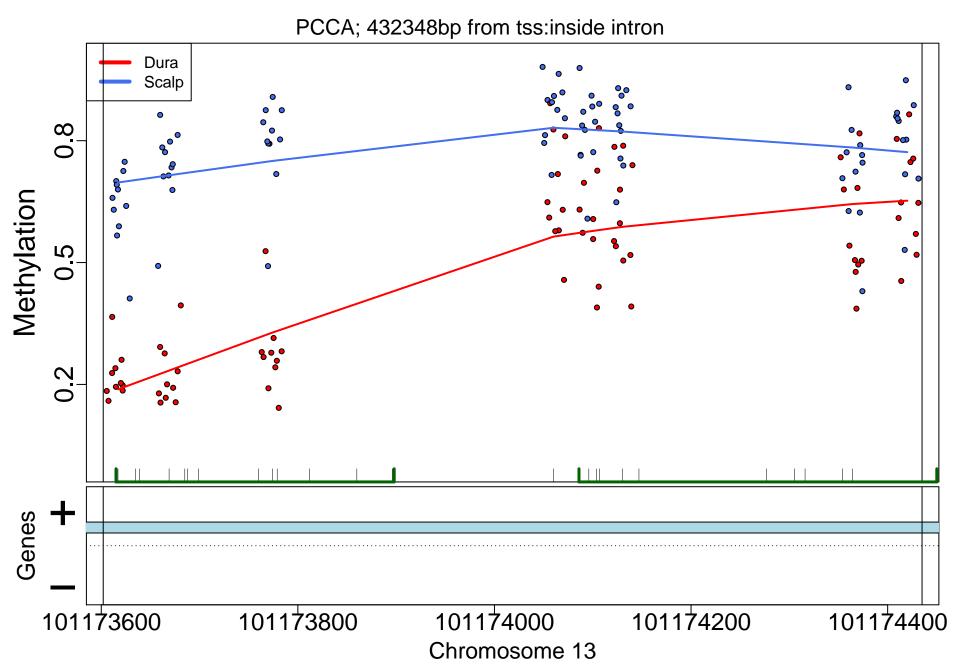


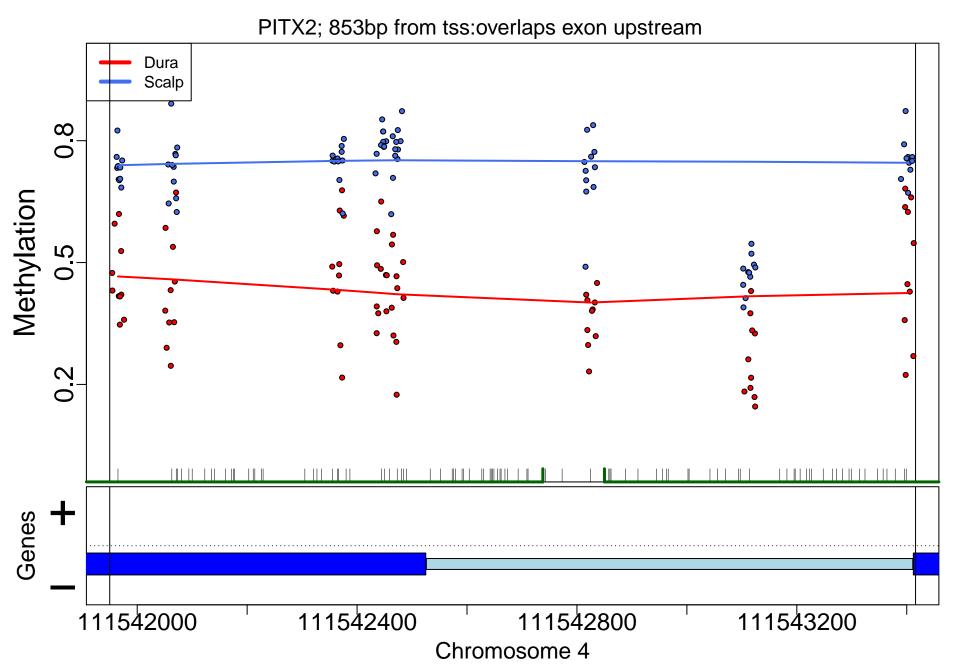


CLEC14A; 0bp from tss:overlaps 5' Dura Scalp <u>∞</u>. Methylation 0.2 Genes CLEC14A 38725500 38724500 38725000 38726000 Chromosome 14





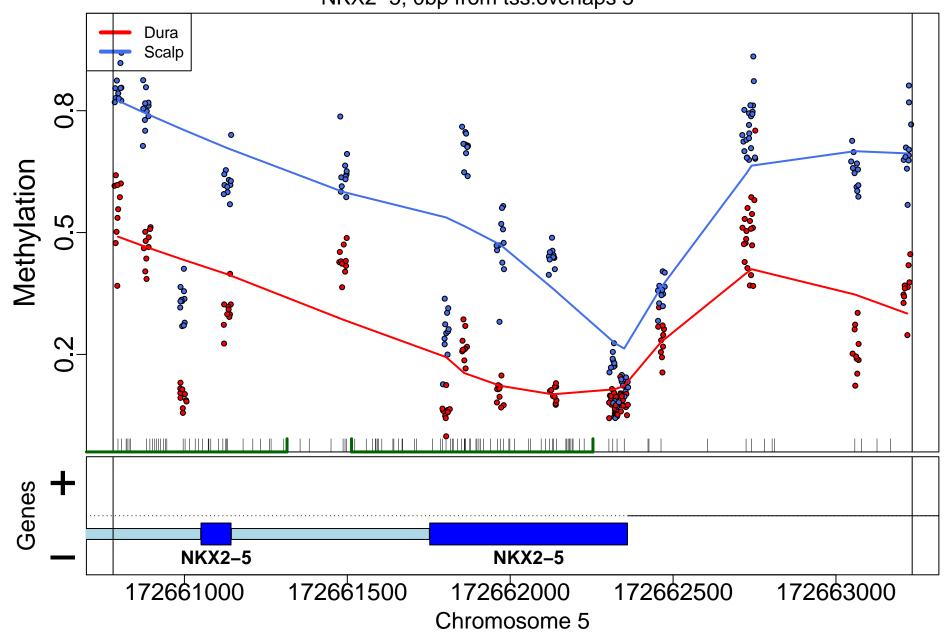


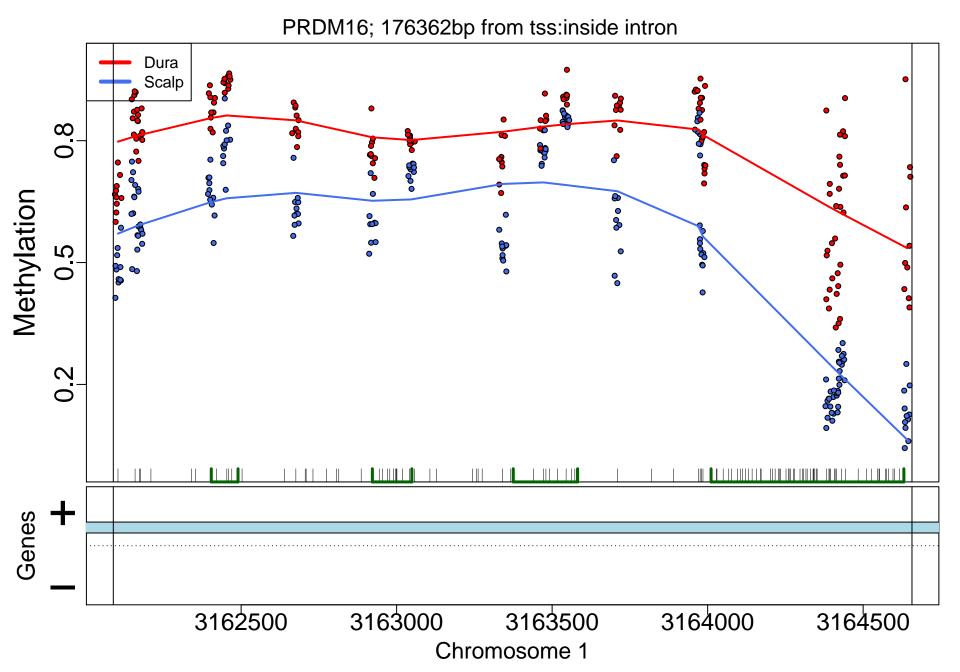


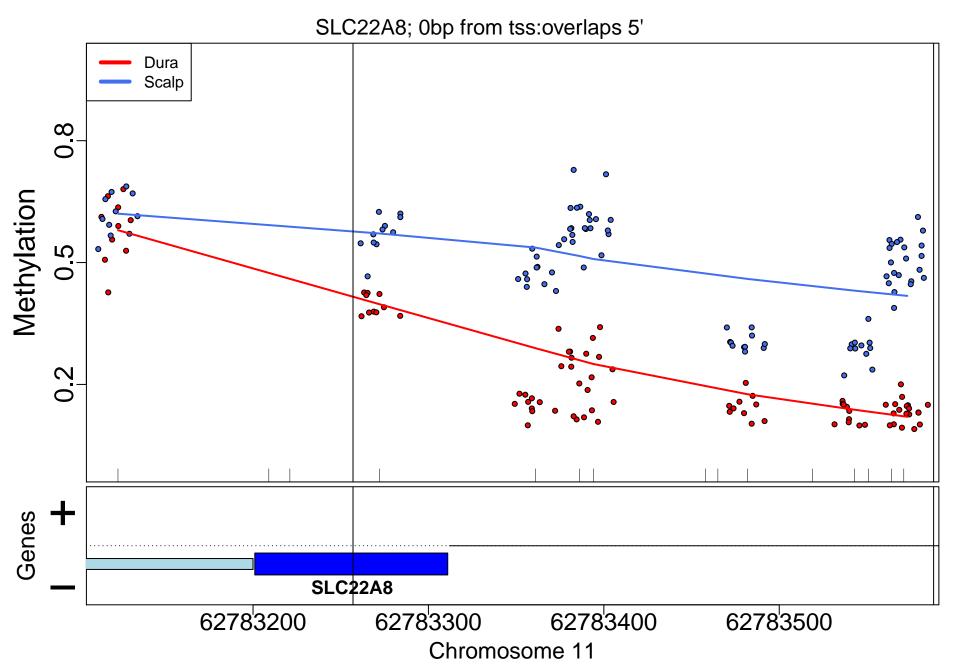
SENP7; 1082bp from tss:inside intron Dura Scalp <u>∞</u>. Methylation Genes SENP7 101231000 101231400 101231800 101232200 Chromosome 3

ALX4; 3293bp from tss:inside intron Dura Scalp <u>∞</u>. Methylation Genes 44325000 44326000 44327000 44328000 Chromosome 11

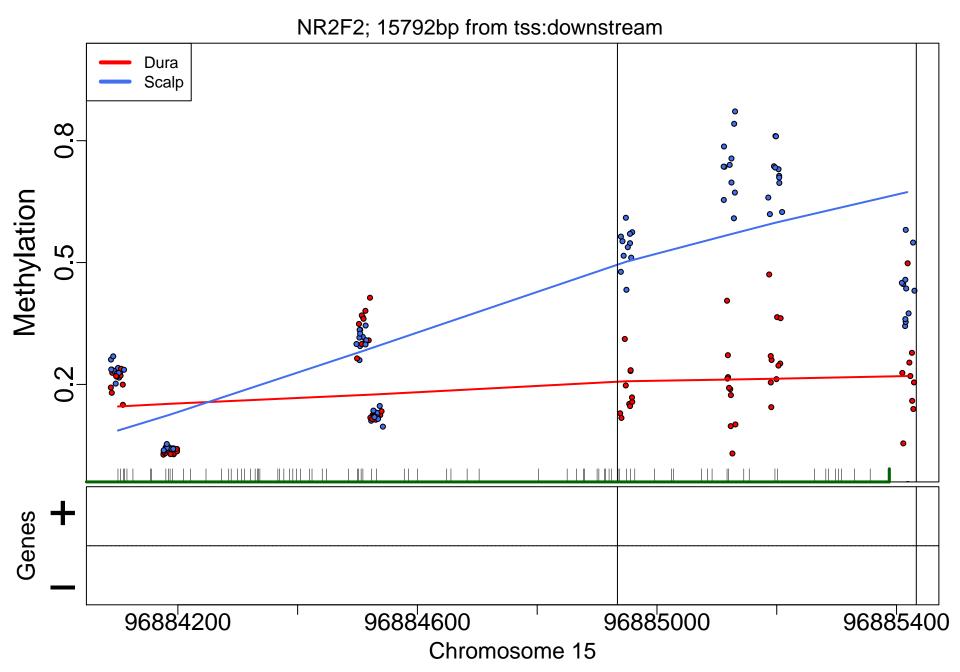
NKX2-5; 0bp from tss:overlaps 5'

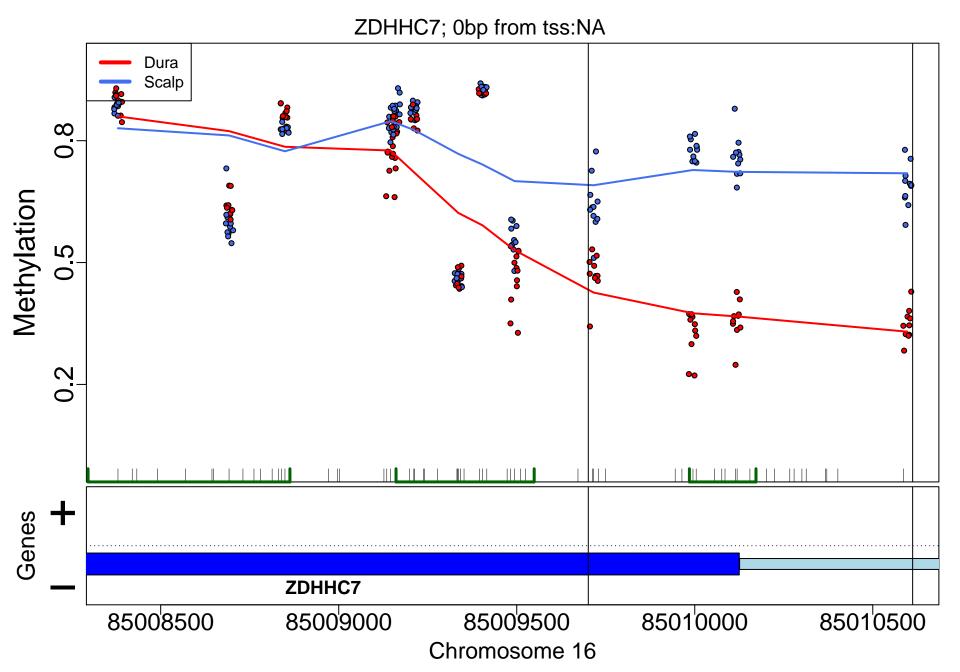


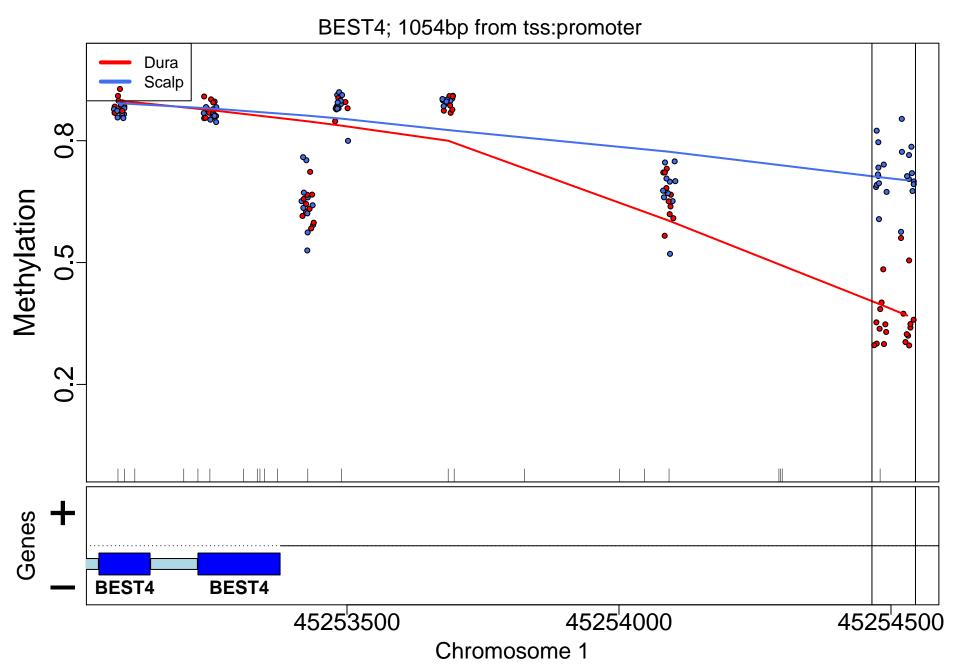




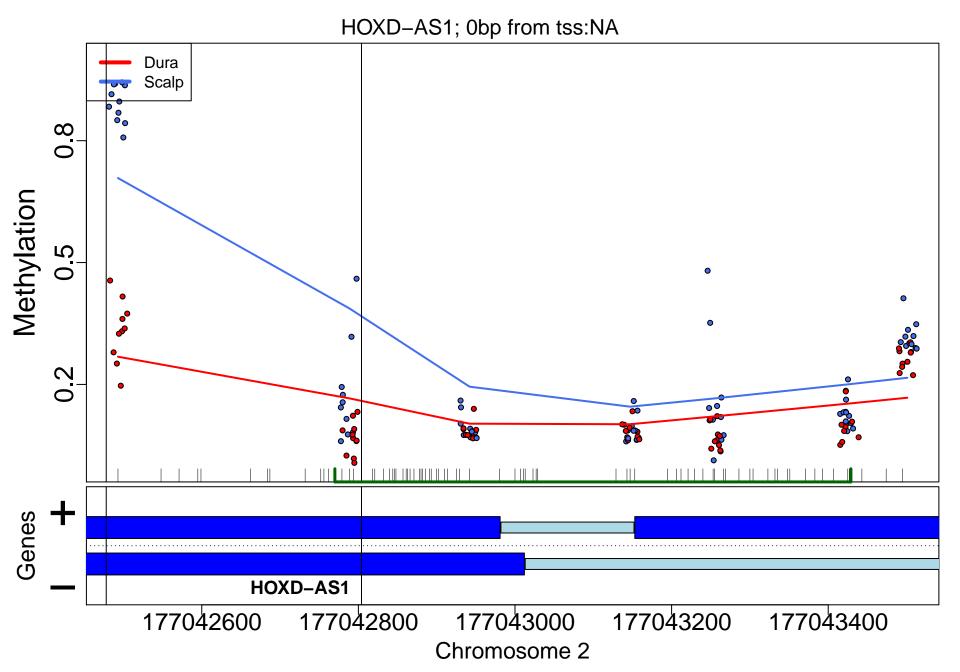
PITX2; 12140bp from tss:inside intron Dura Scalp Methylation Genes 111550000 111550400 111550800 Chromosome 4



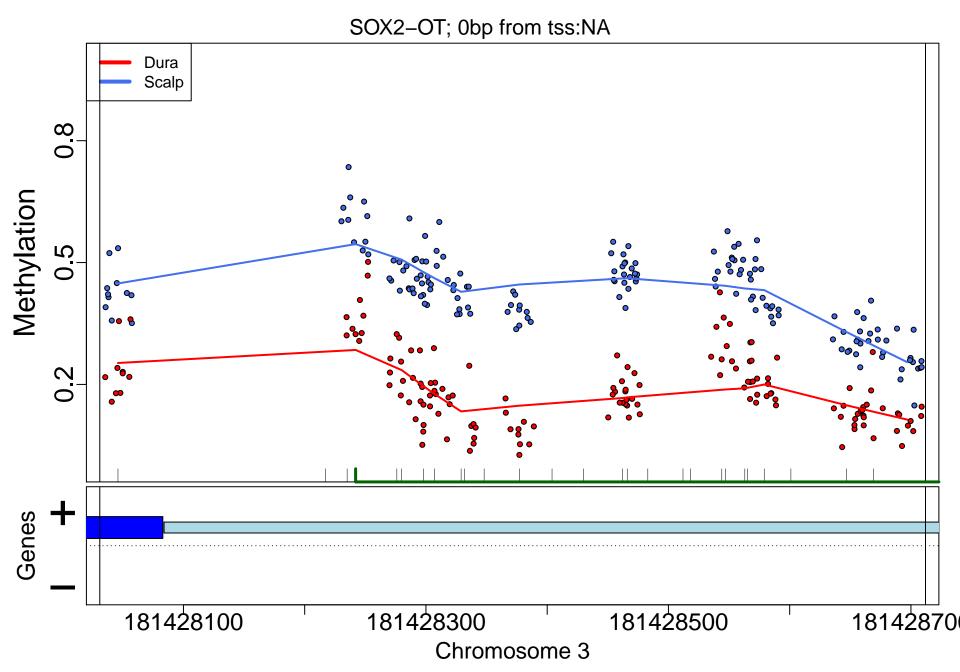


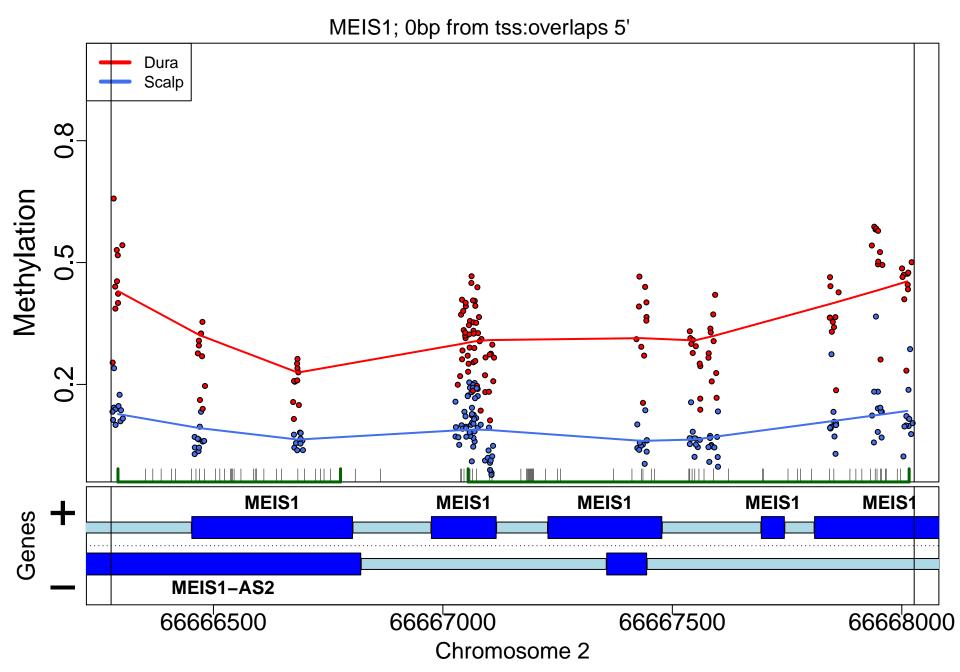


DEPTOR; 351bp from tss:promoter Dura \$calp <u>ω</u>_ Methylation 0 0.2 **3 DEPTOR** Genes 120885600 120886000 120886400 120886800 Chromosome 8

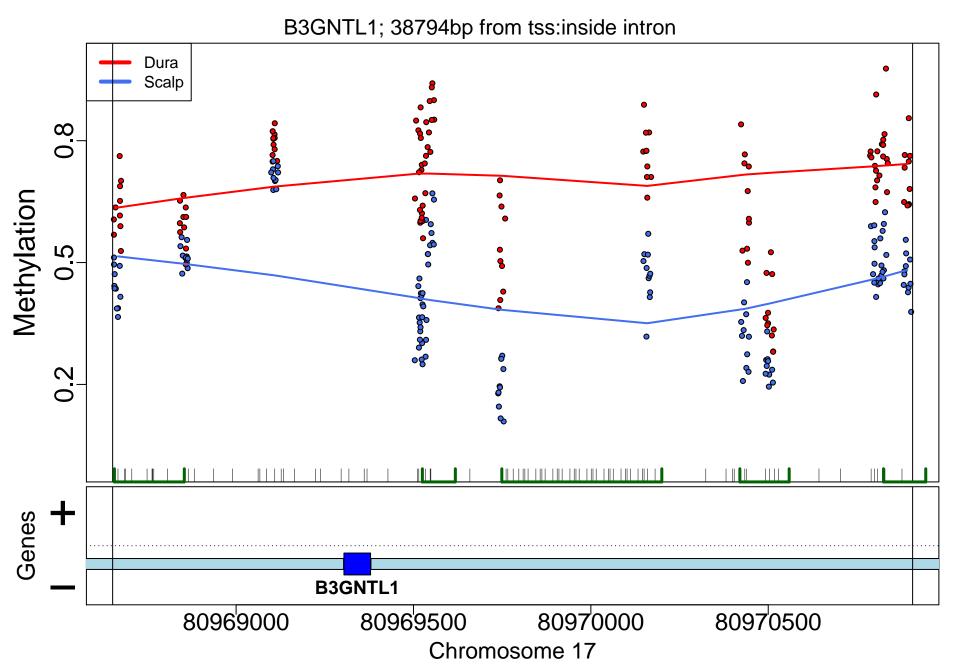


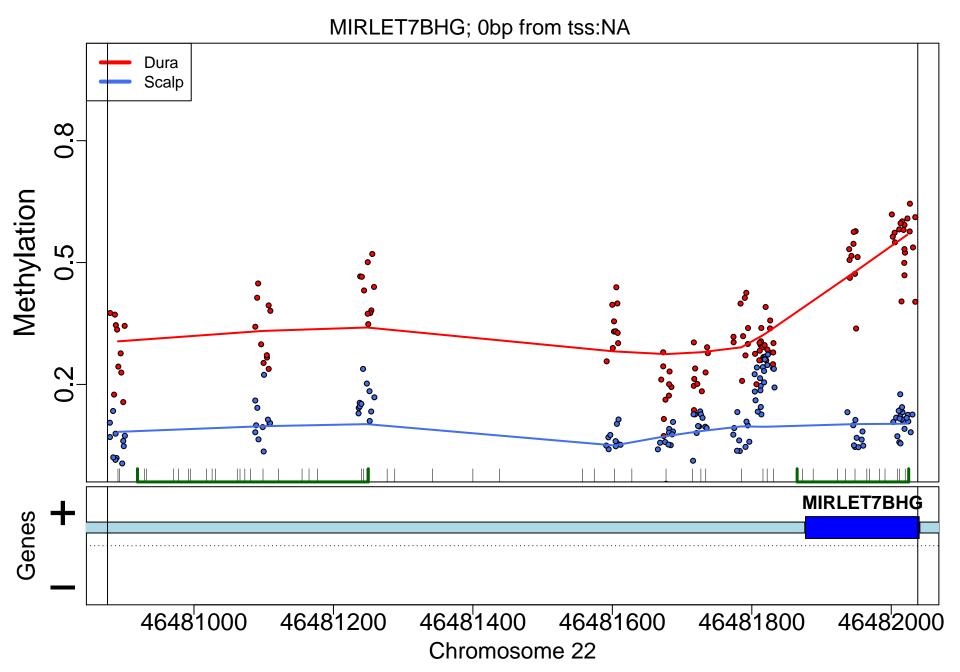
MCF2L; 0bp from tss:overlaps 5' Dura Scalp Methylation MCF2L Genes 113698400 113698600 113699200 113698800 113699000 113699400 Chromosome 13

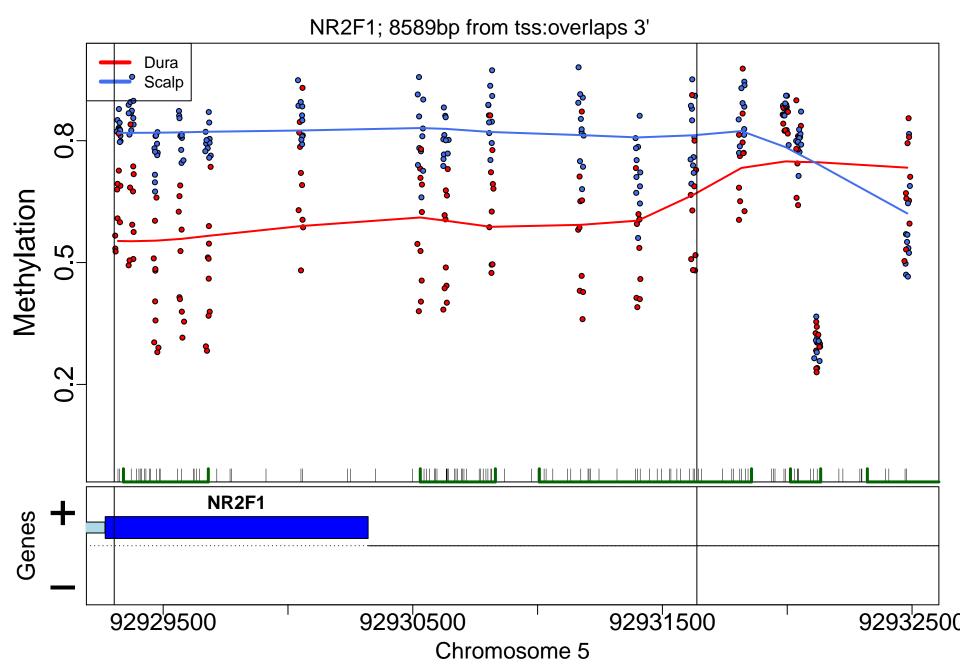


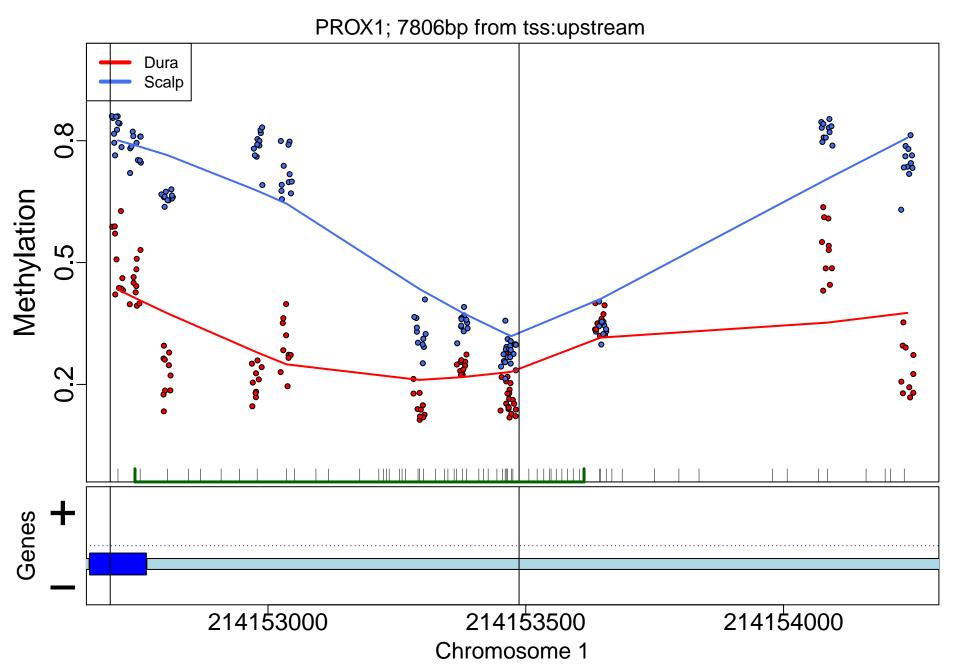


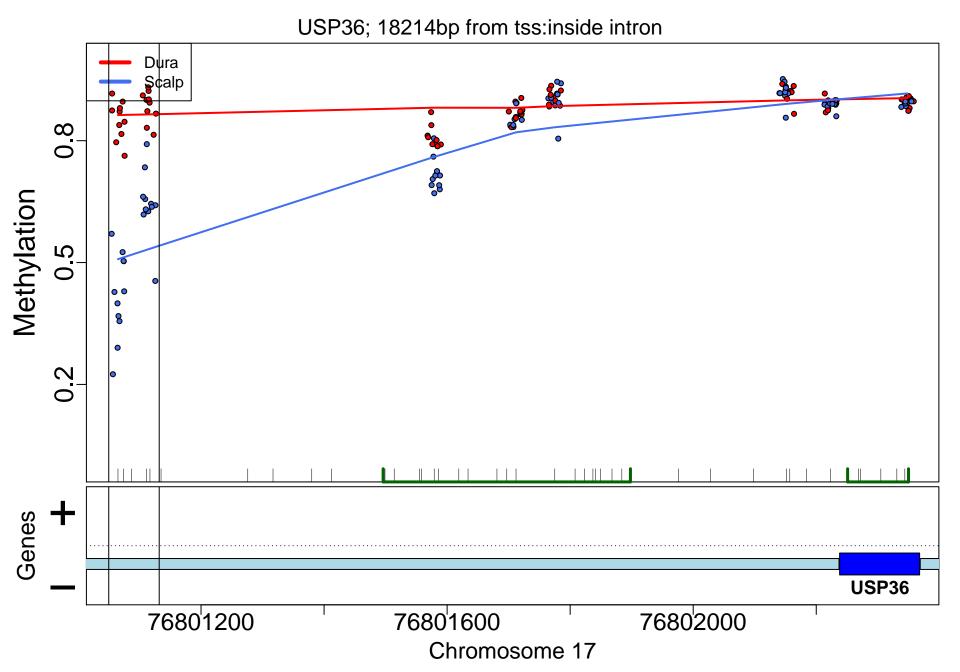
GSC; 2172bp from tss:promoter Dura Scalp <u>∞</u>. Methylation 0.2 Genes 95239000 95240000 95240500 95239500 Chromosome 14

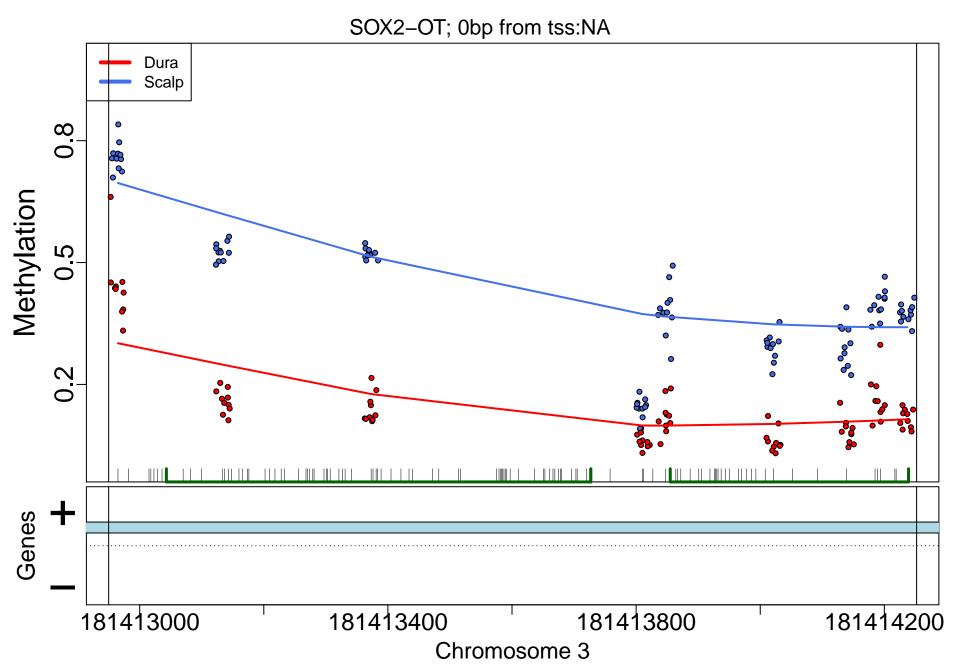




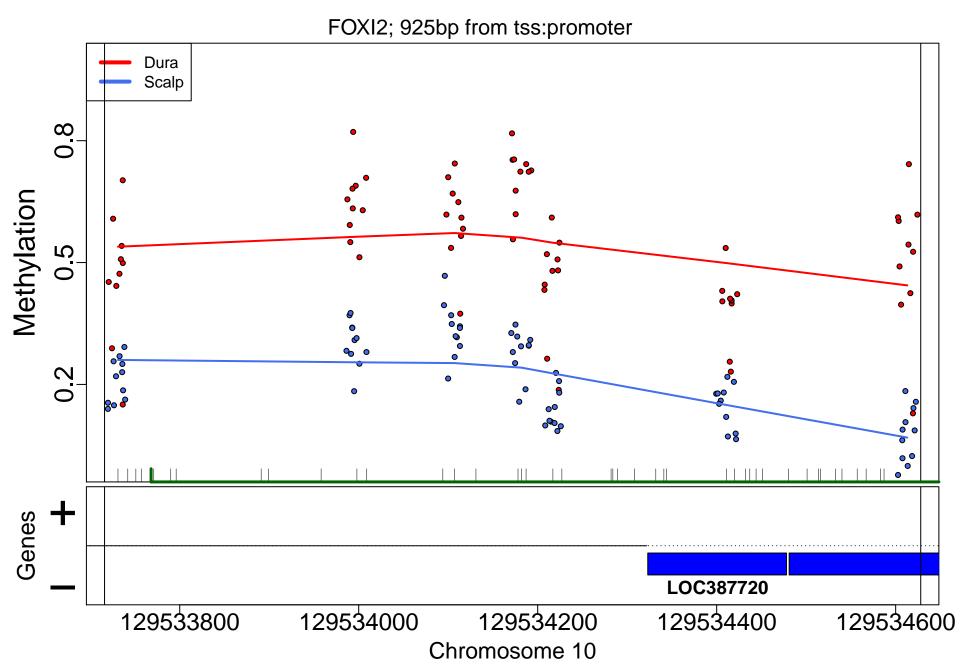


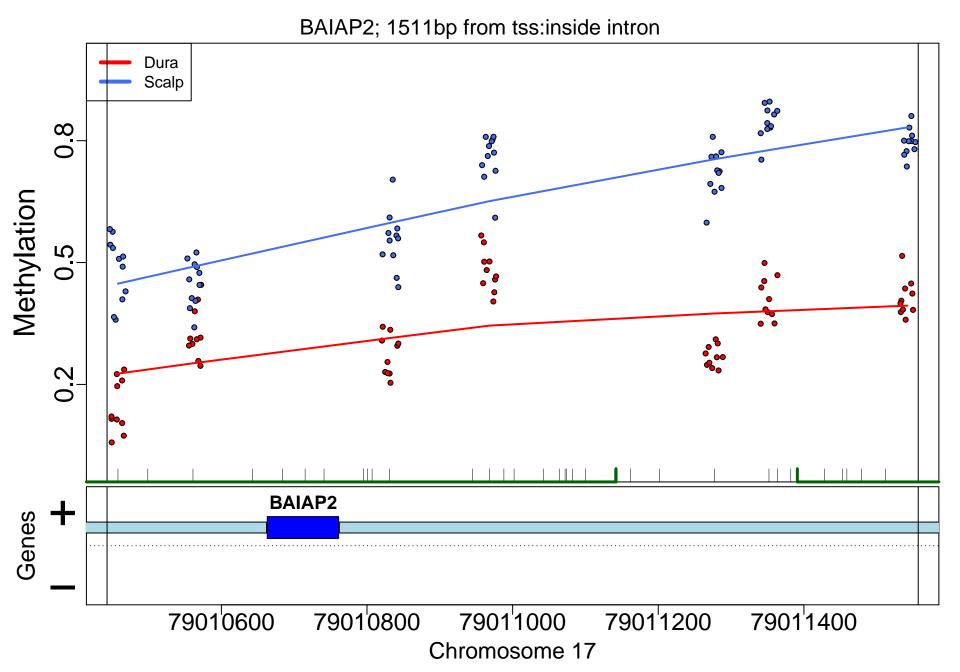


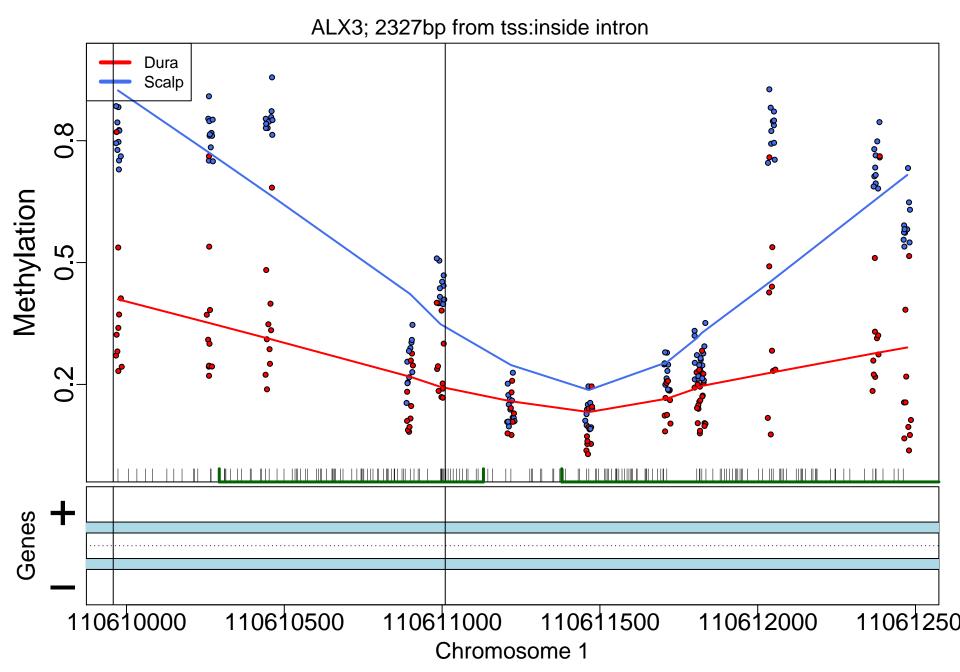


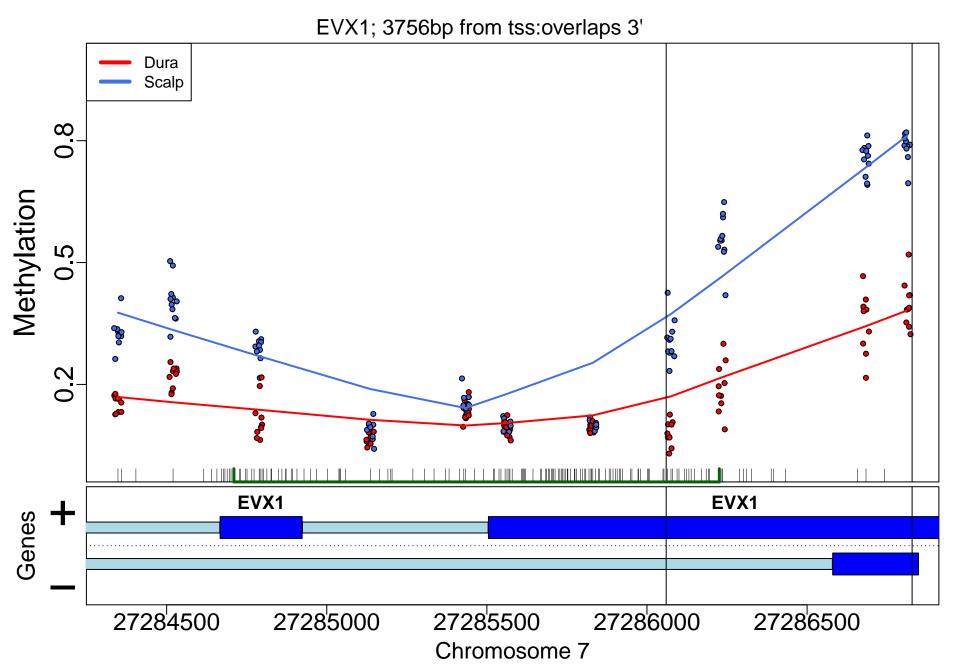


SOX2-OT; 0bp from tss:NA Dura Scalp <u>ω</u>_ Methylation **%**° 0.2 Genes 181441400 181441600 181441800 181442000 Chromosome 3

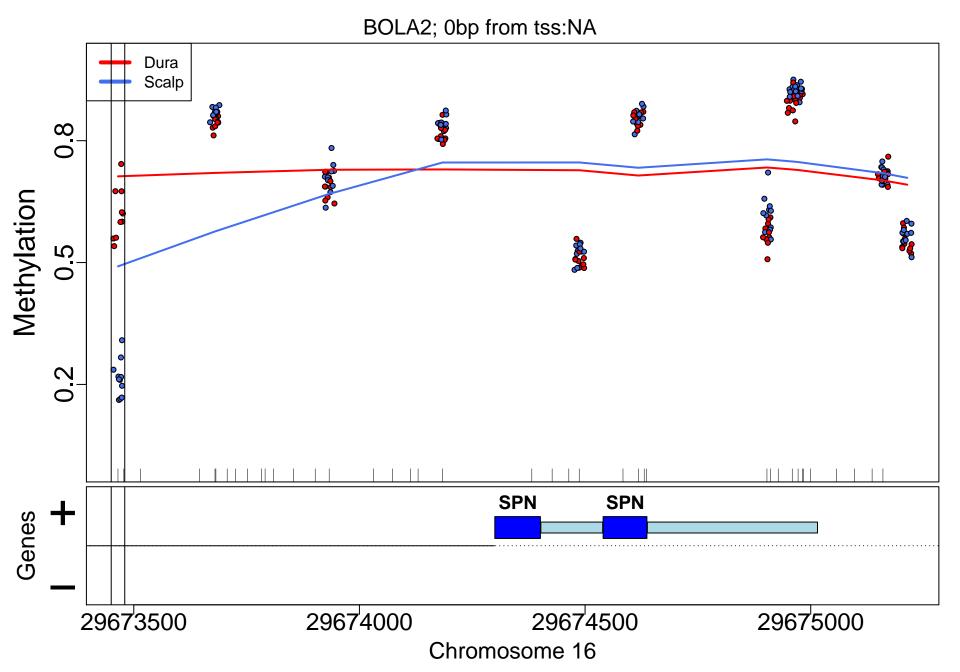


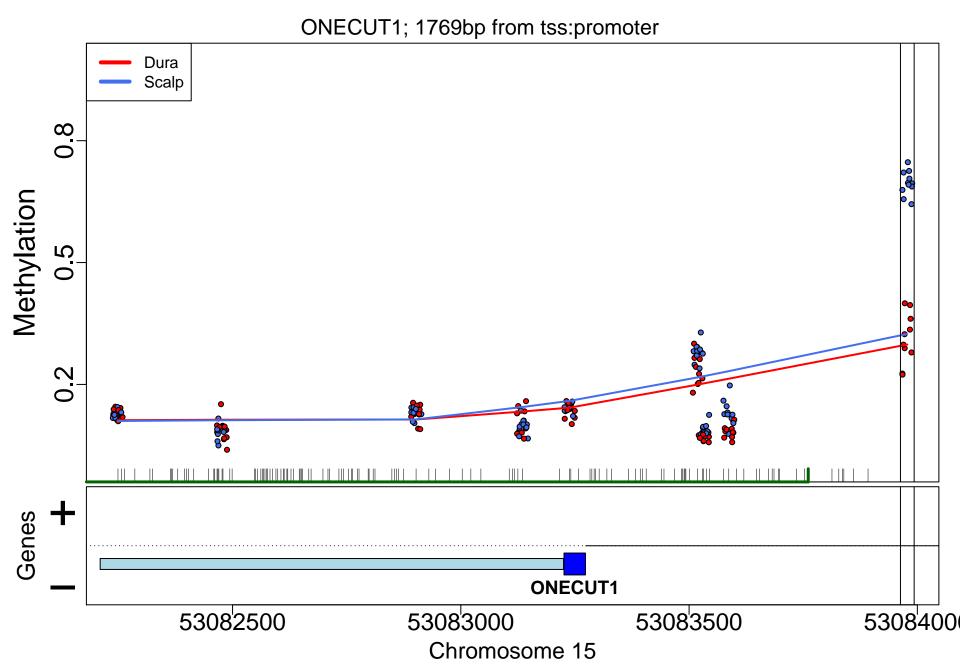






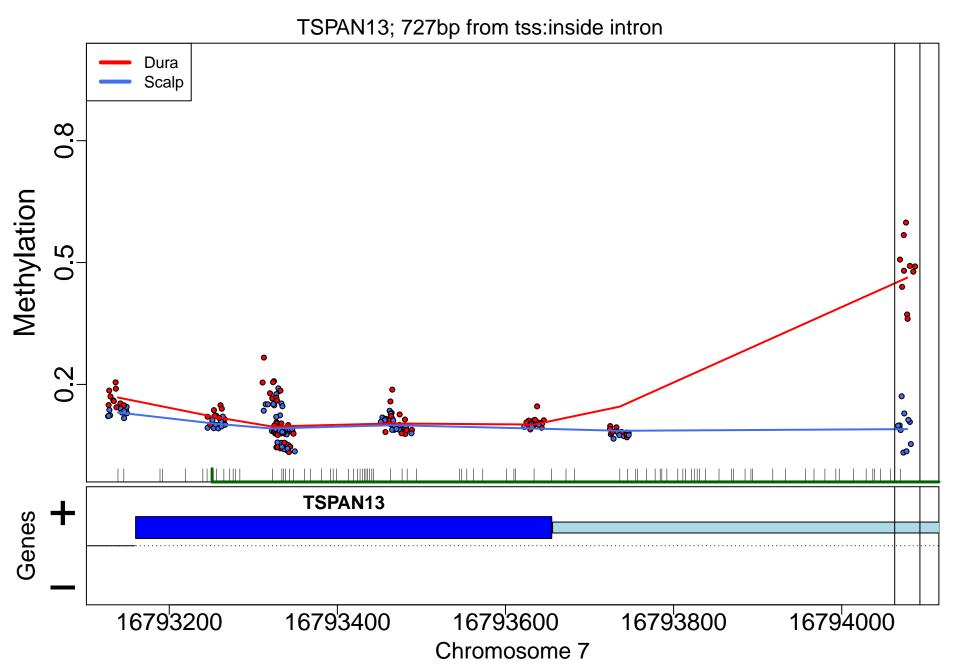
LINC01019; 0bp from tss:NA Dura Scalp <u>ω</u>_ Methylation **8** 0.2 Genes LINC01019 3535600 3535800 3536200 3536400 3535400 3536000 Chromosome 5

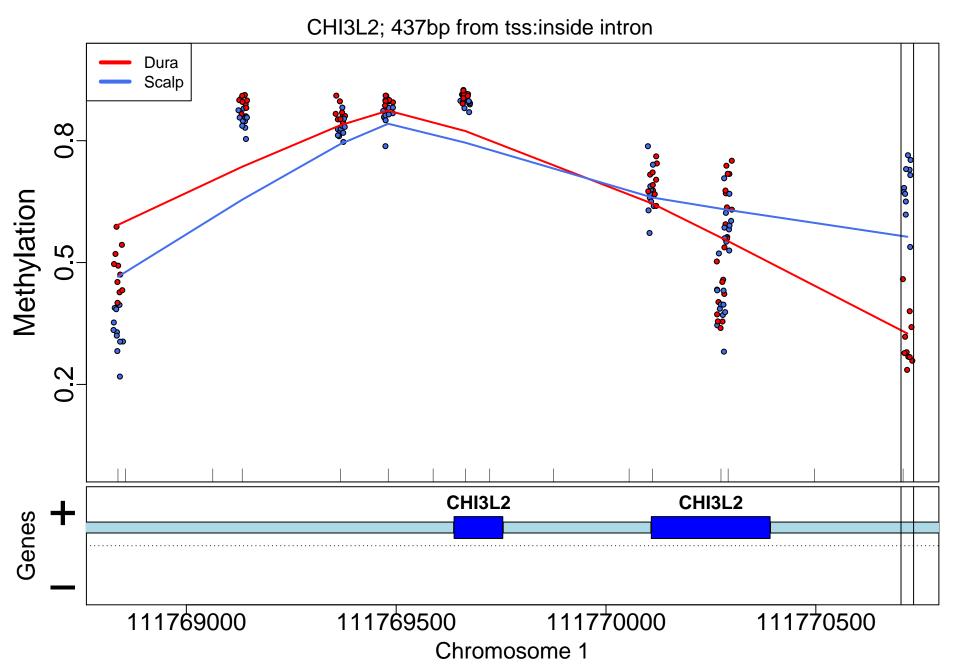




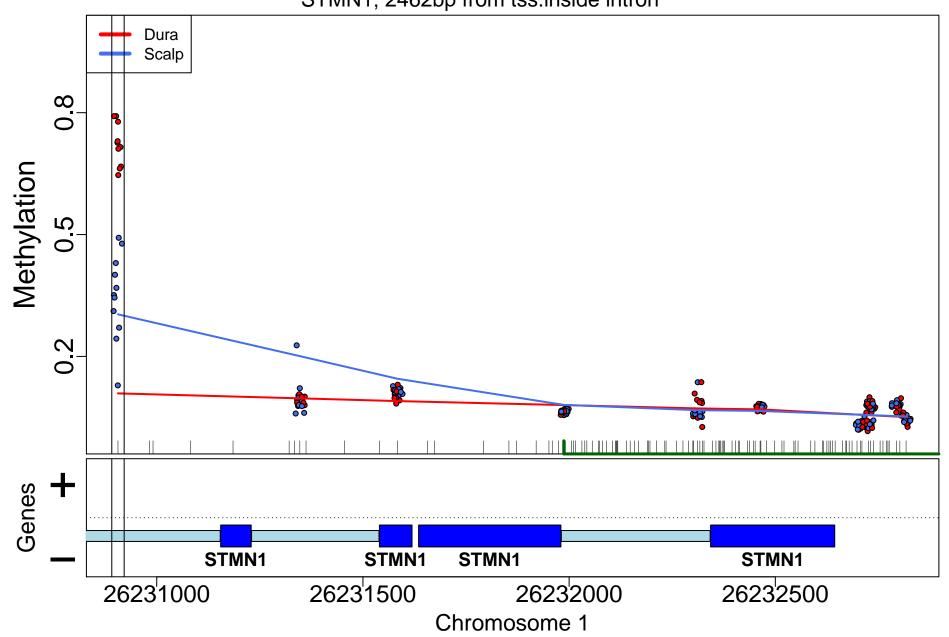
ZNF192P1; 0bp from tss:NA Dura Scalp <u>ω</u>_ Methylation 0.5 0.2 **ZNF192P1** Genes 28129100 28129200 28129400 28129500 28129600 28129300

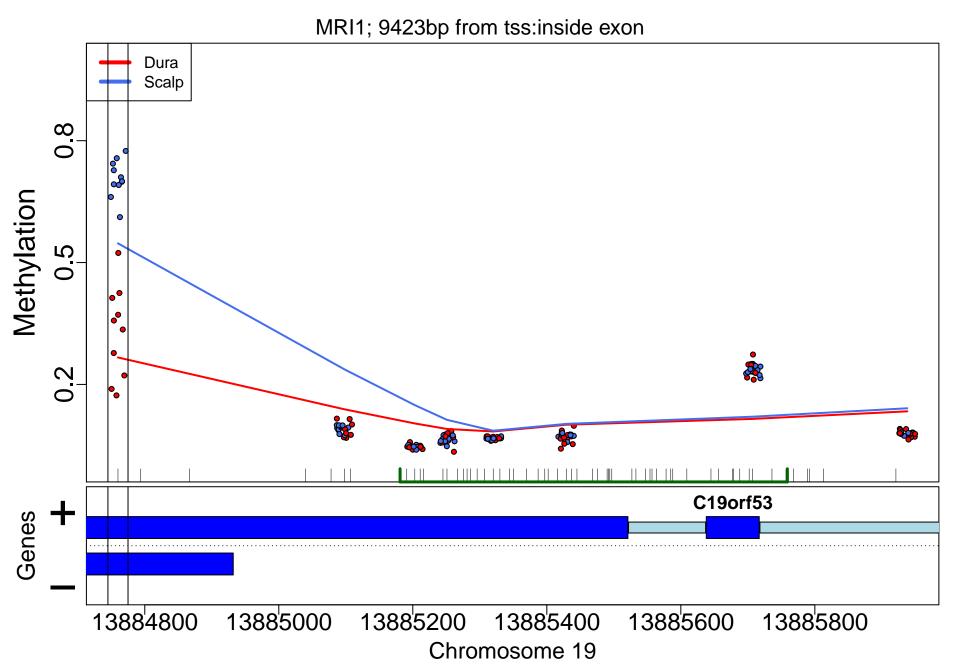
Chromosome 6

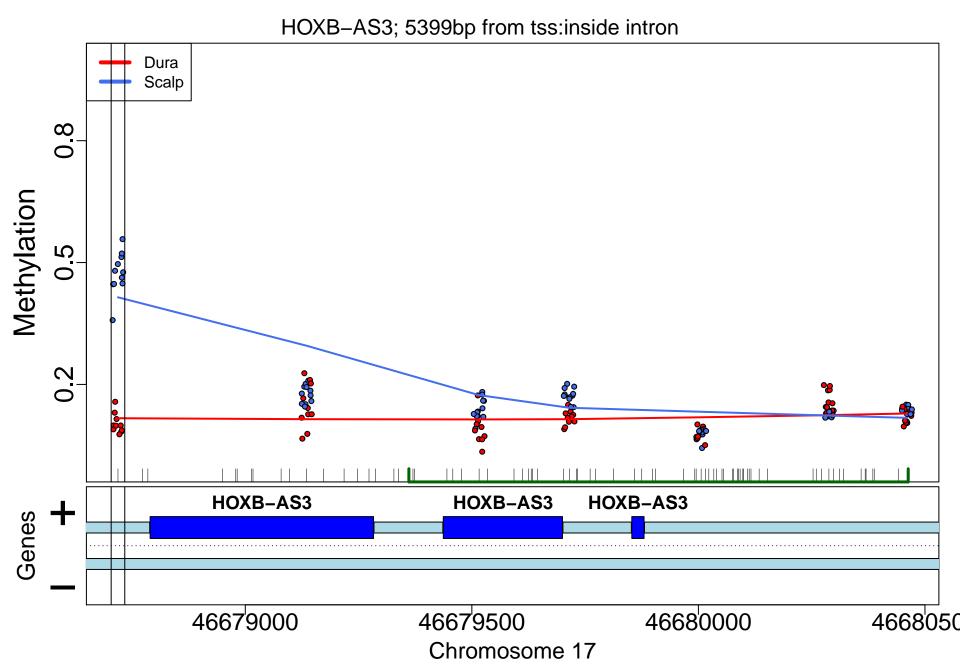


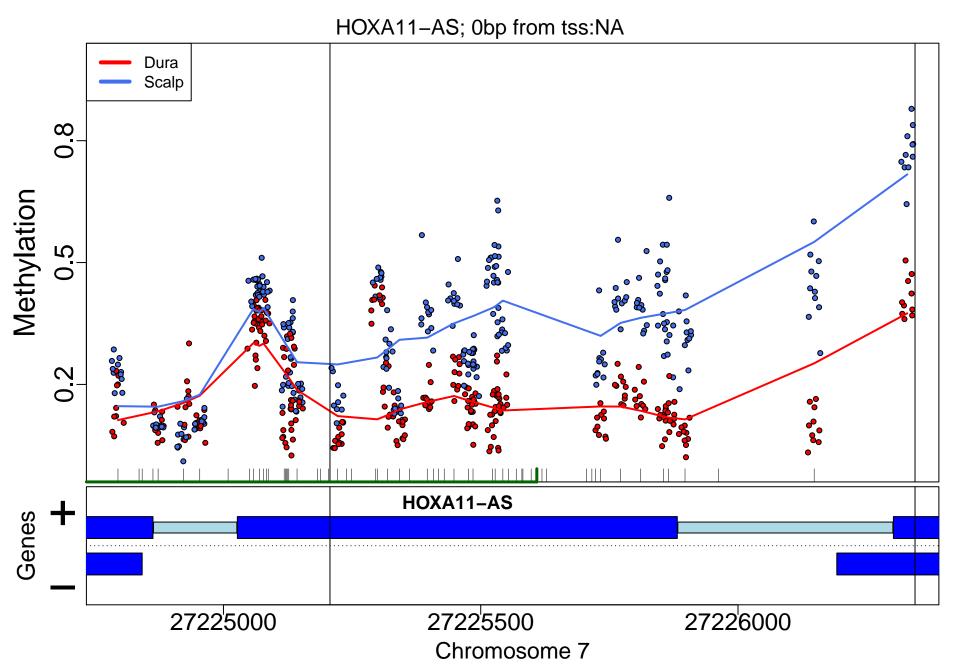


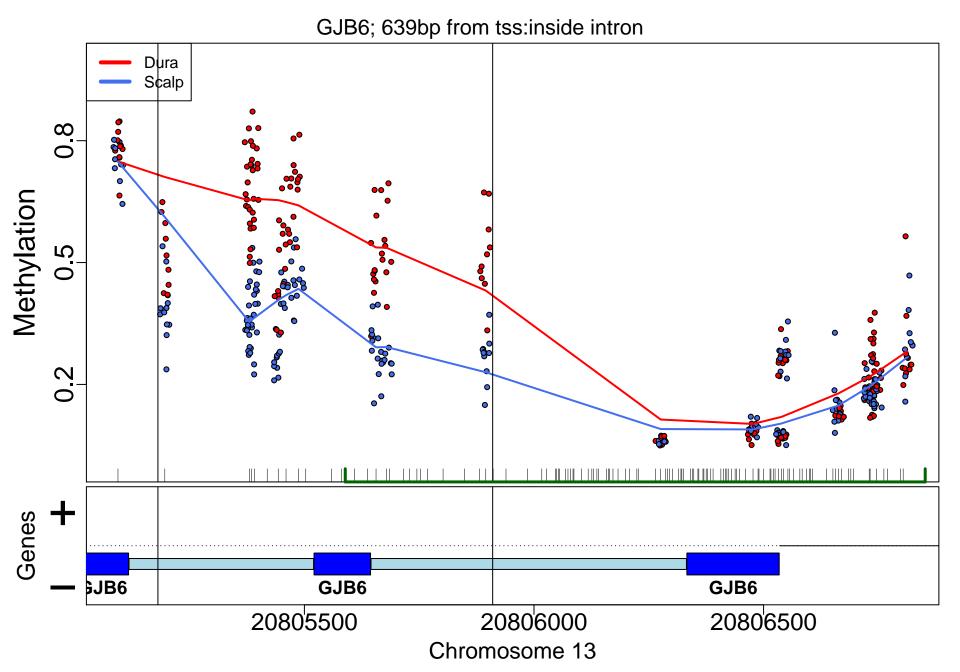
STMN1; 2462bp from tss:inside intron

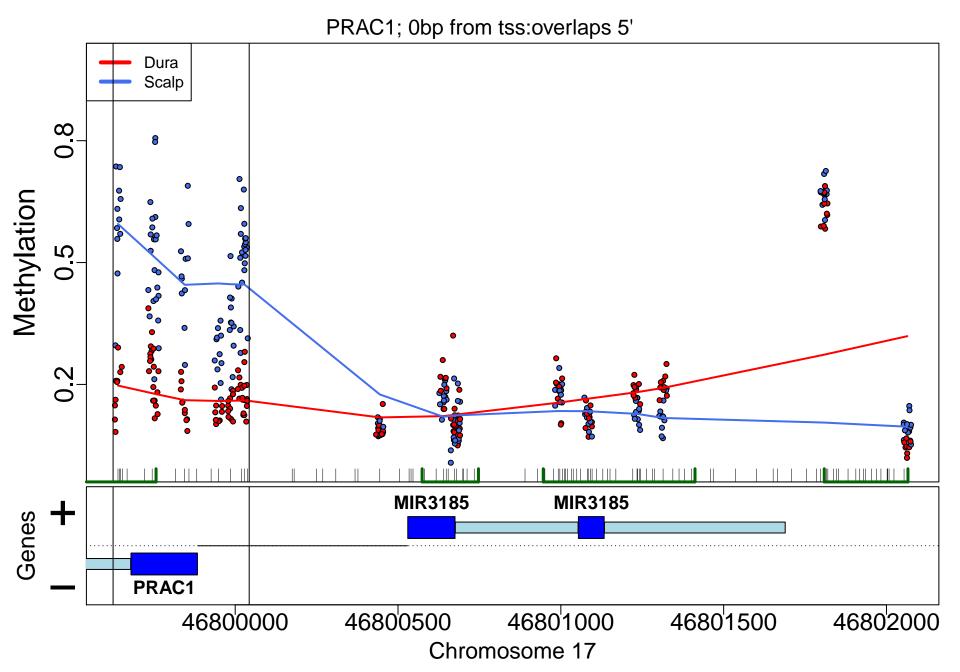


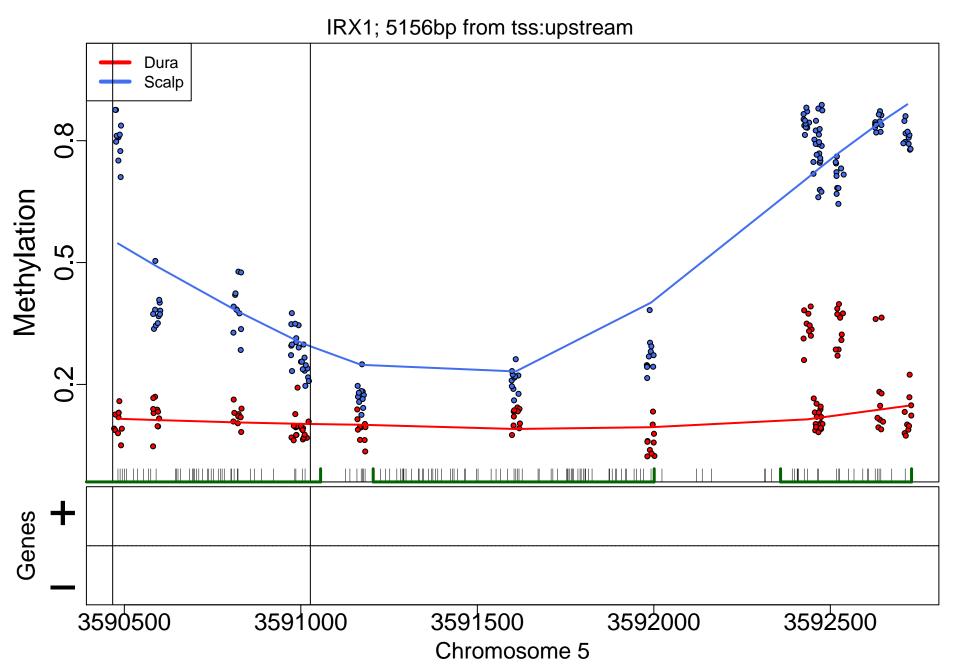


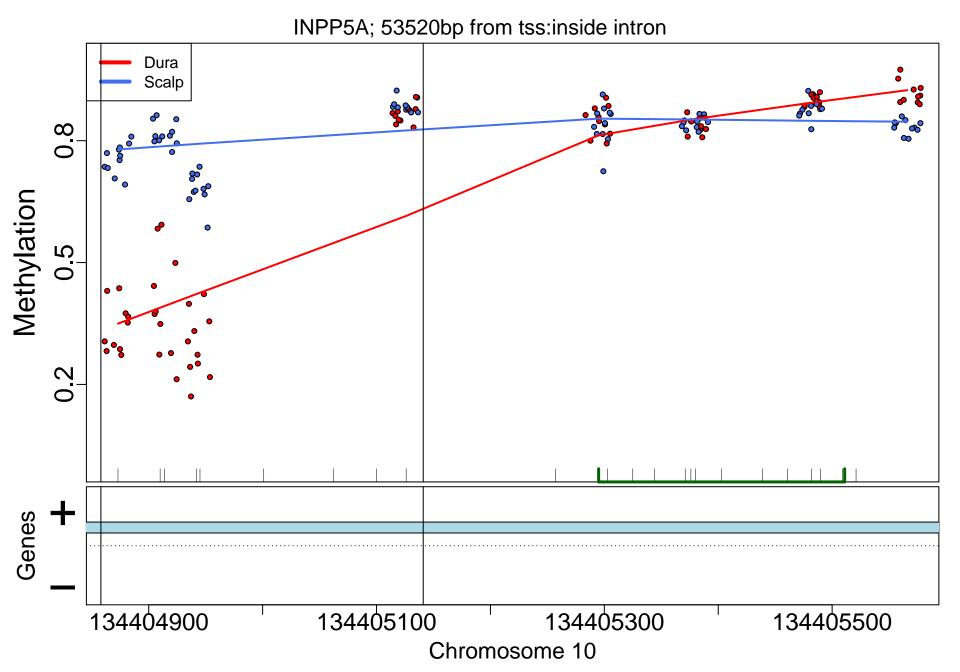


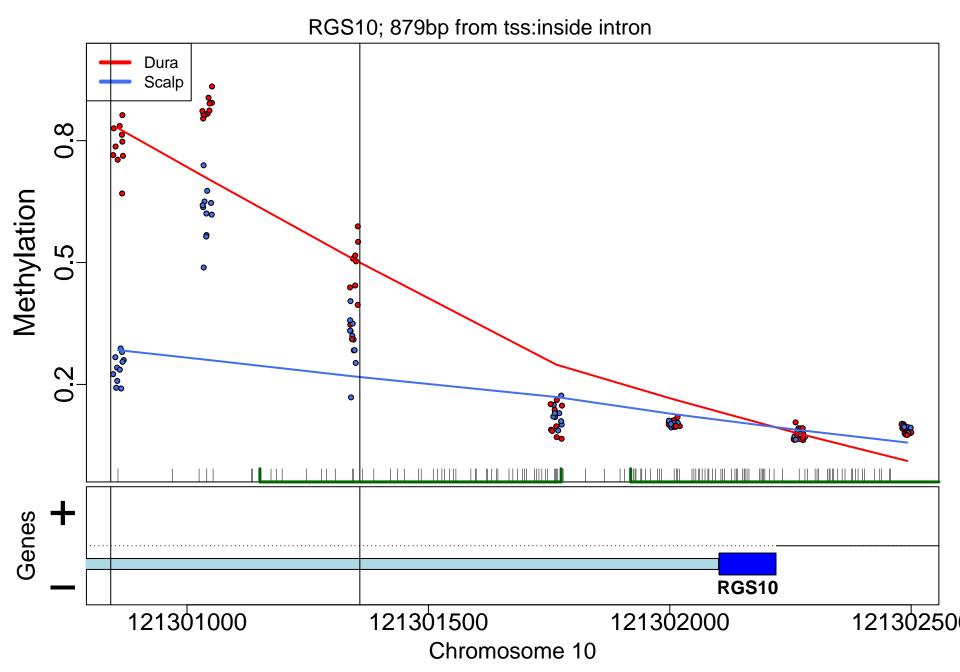


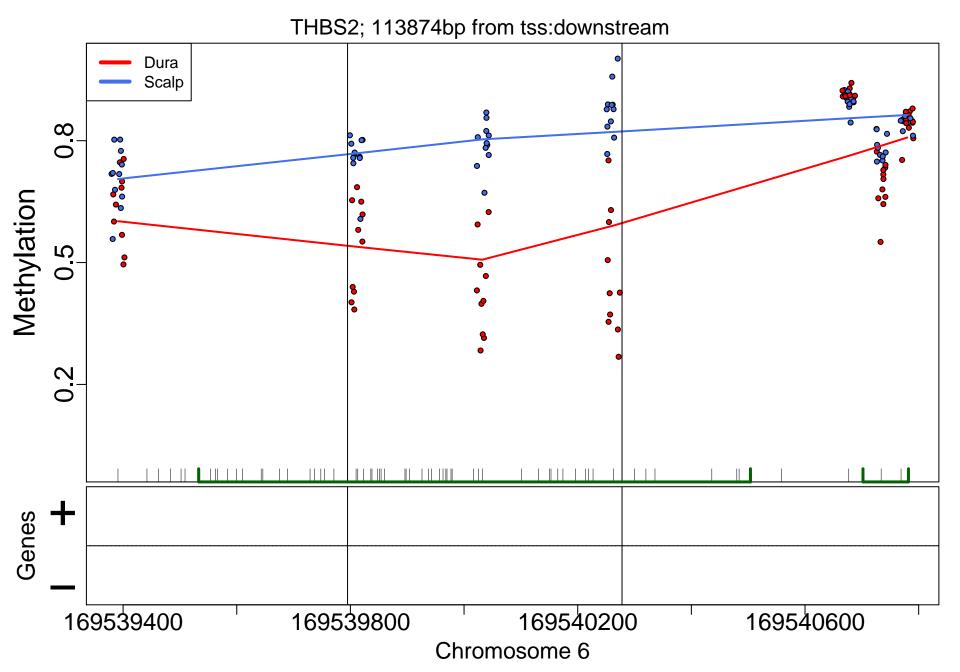


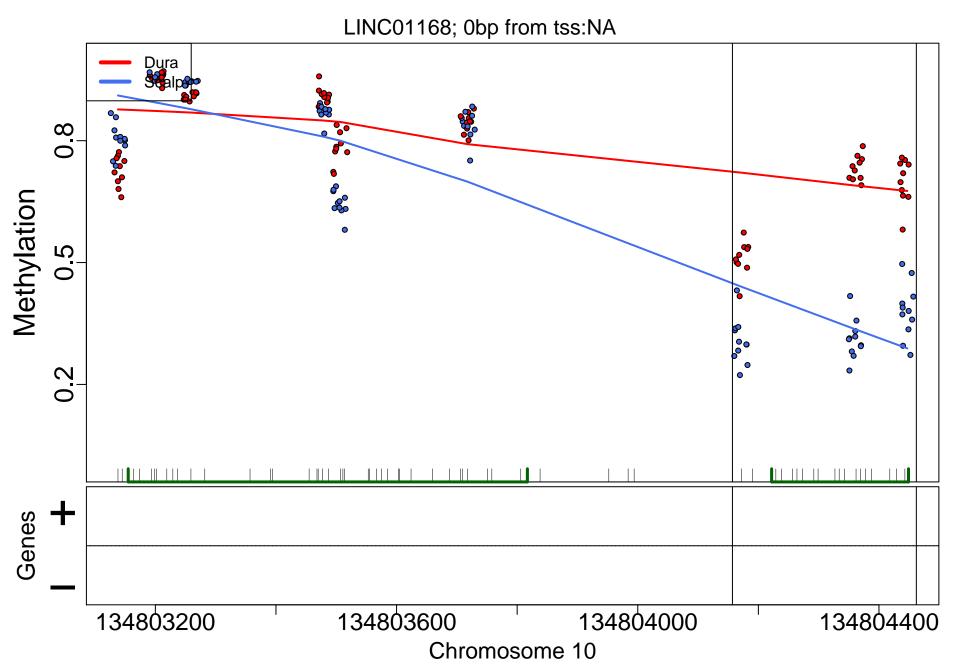


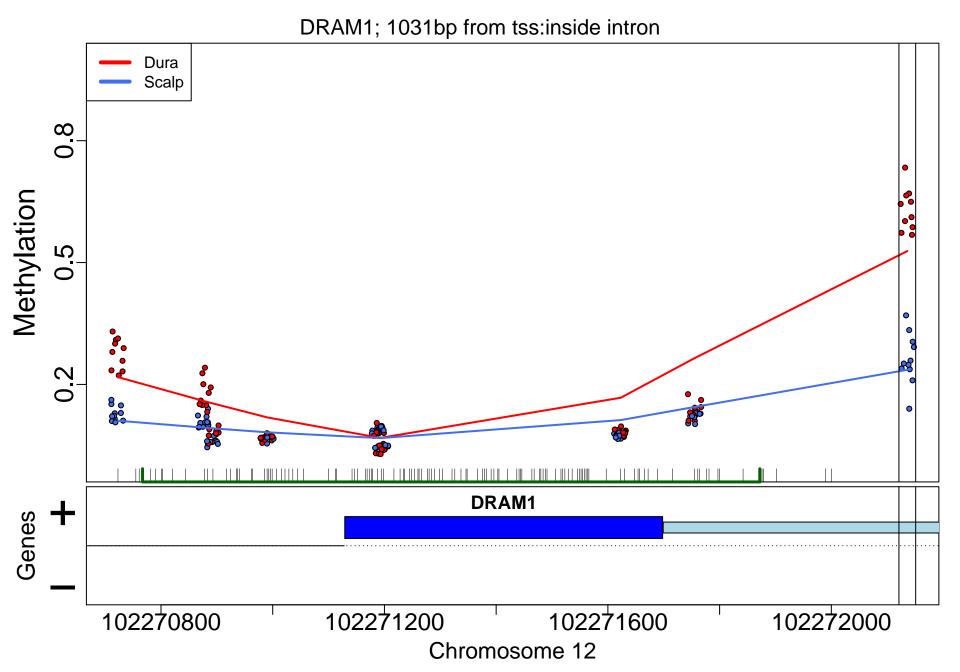




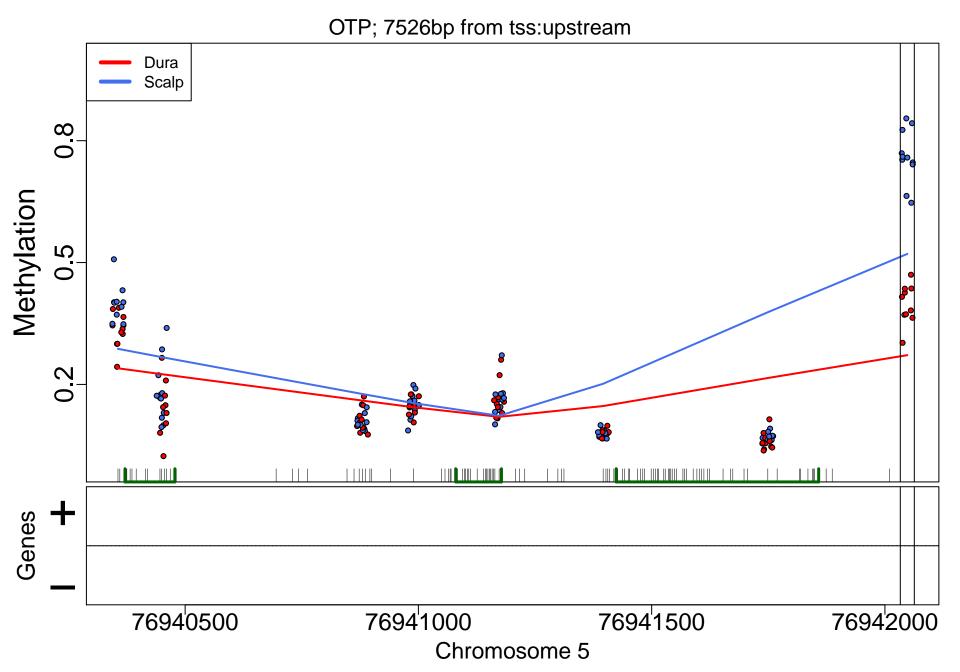


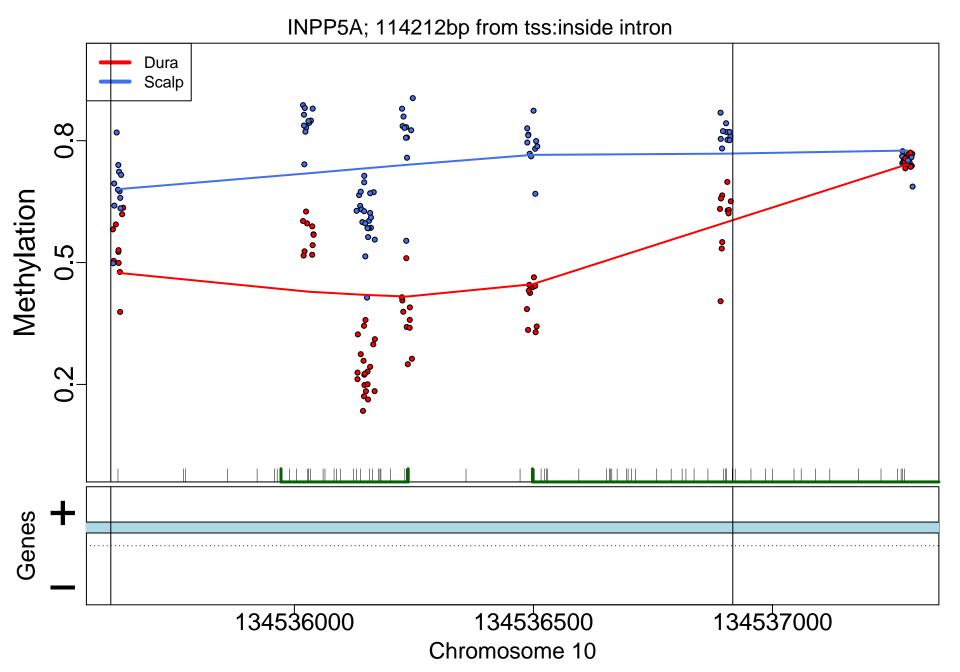


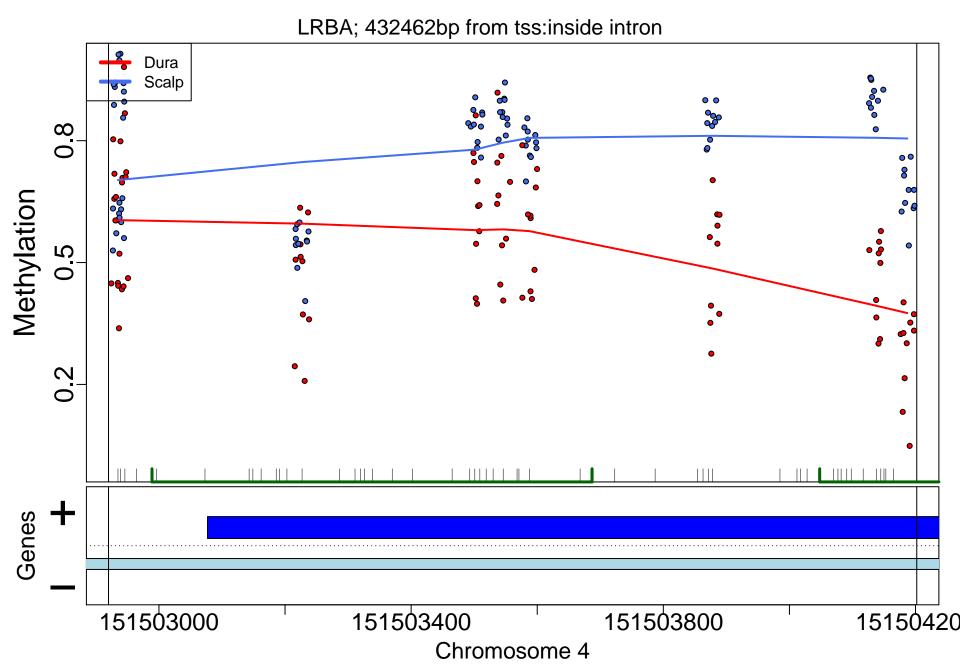


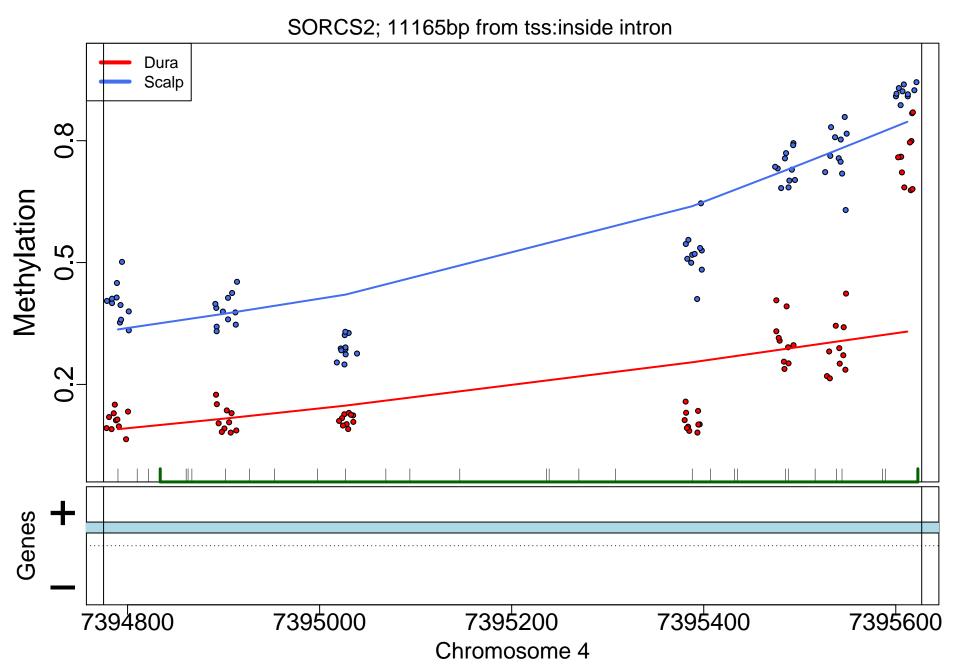


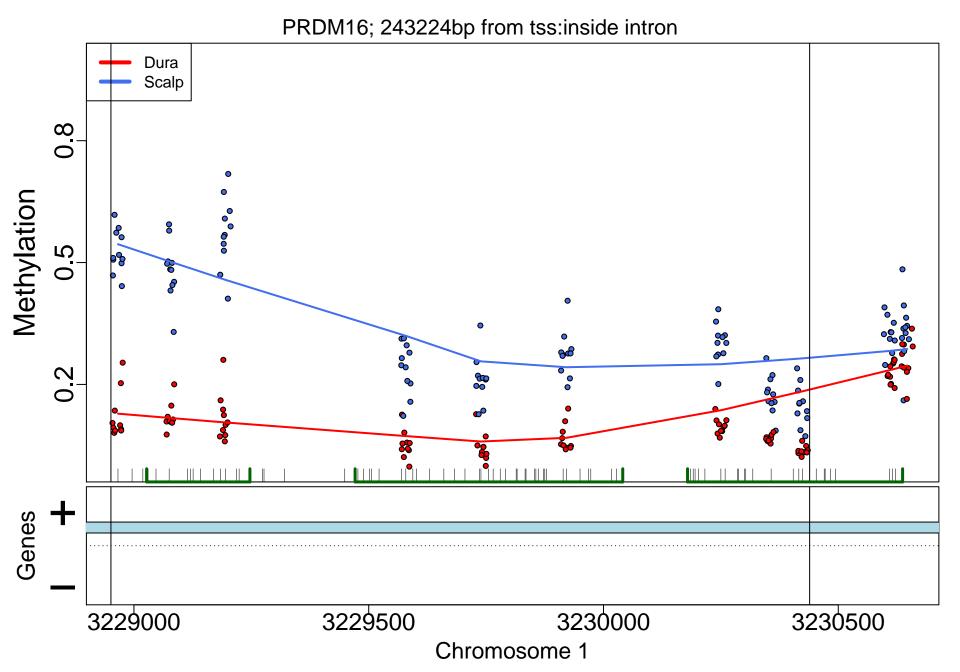
CSGALNACT1; 75391bp from tss:inside intron Dura Scalp <u>ω</u>_ Methylation 0.2 8 Genes CSGALNACT1 CSGALNACT1 CSGALNACT1 19540600 19540000 19540200 19540400 Chromosome 8

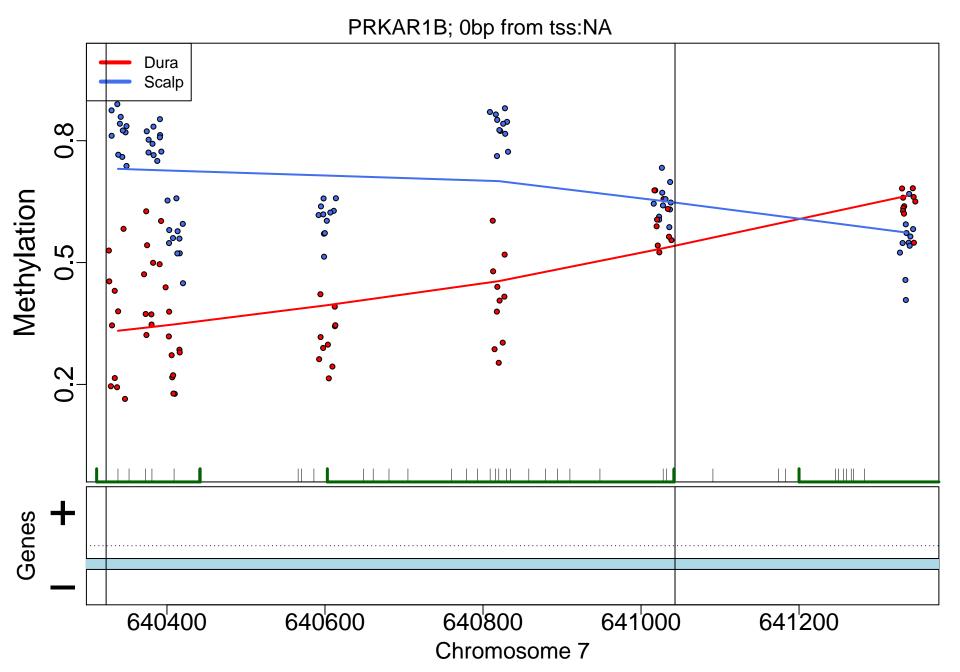


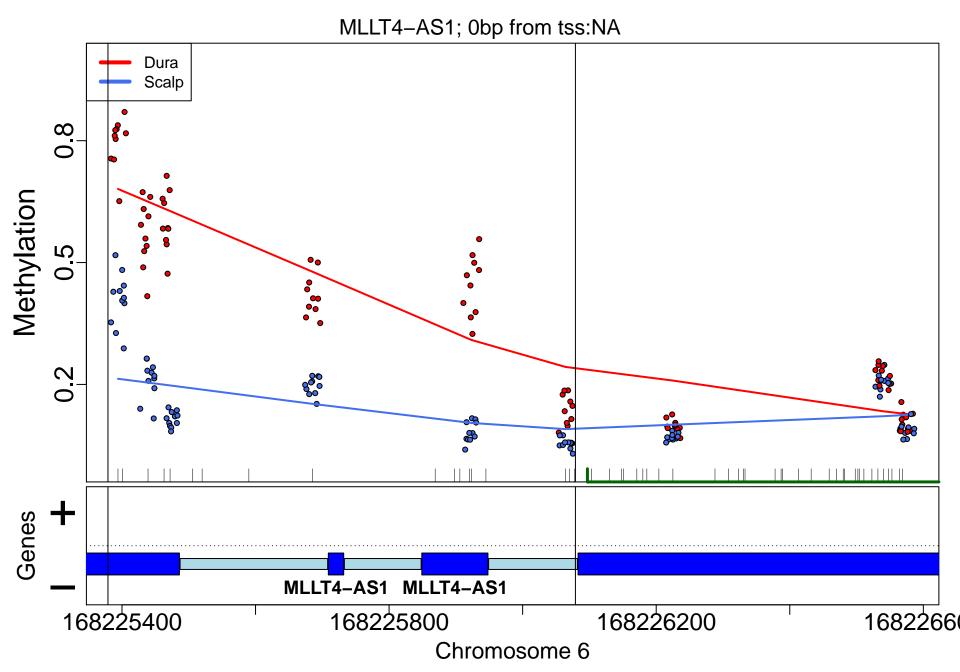


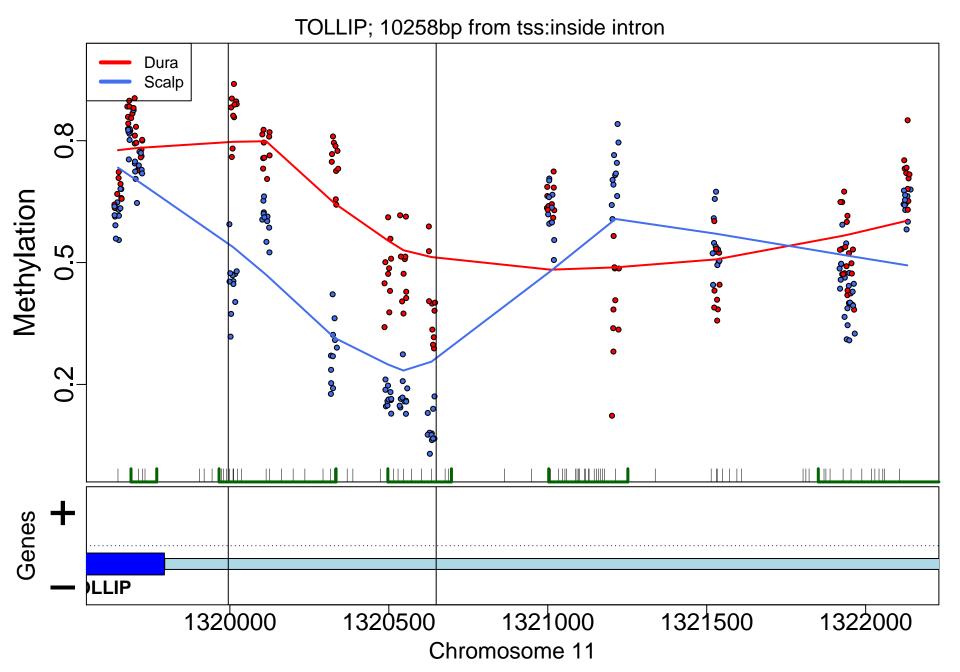


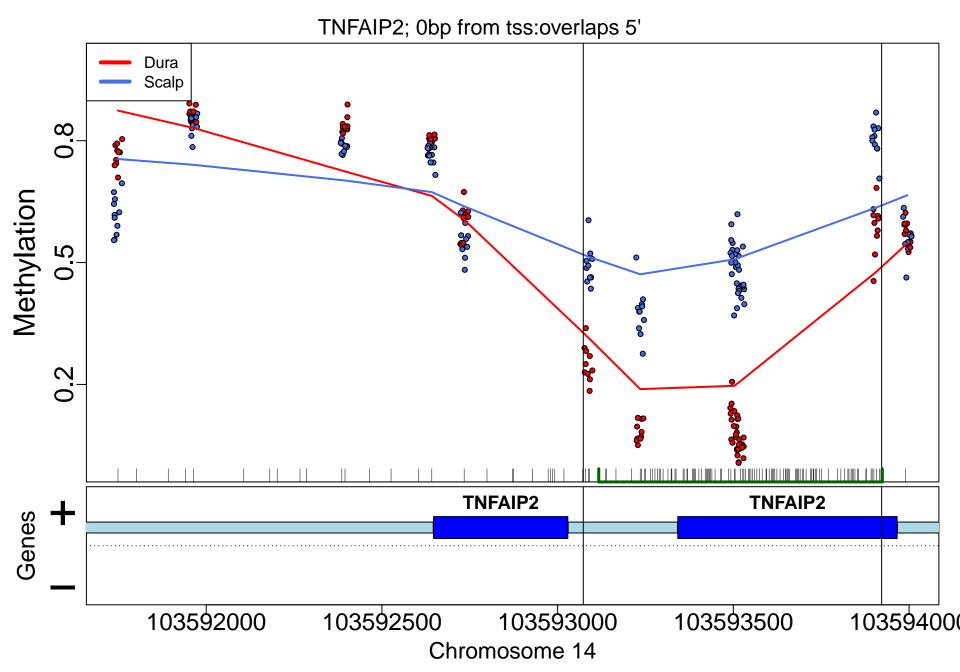


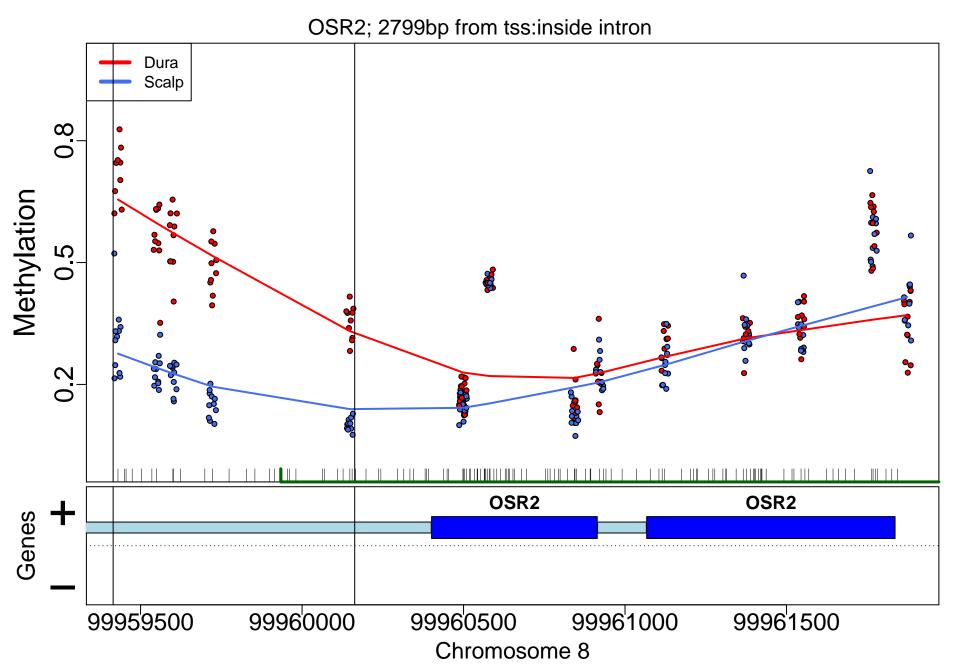


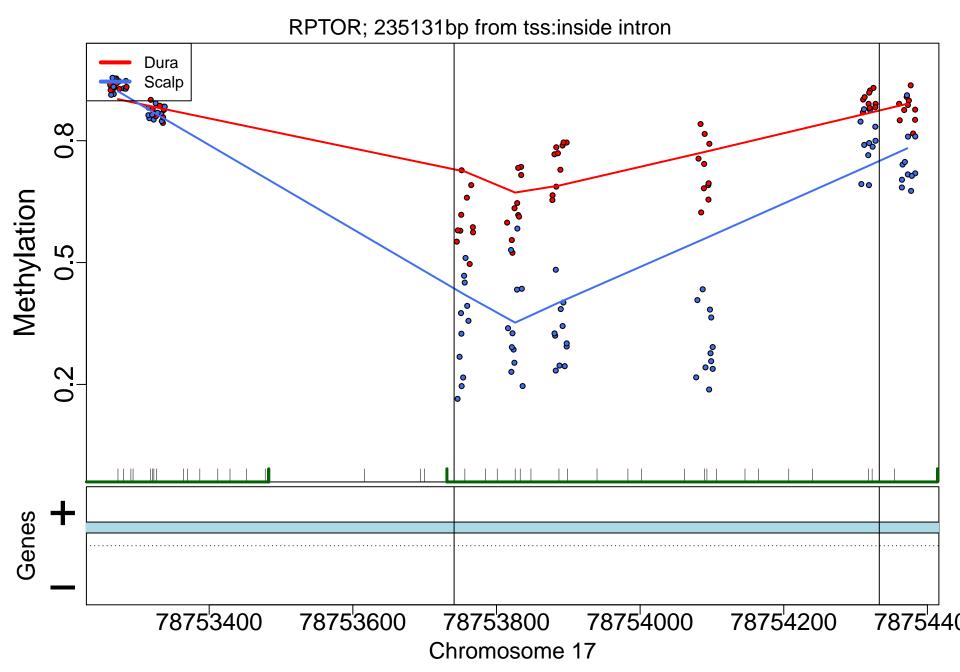


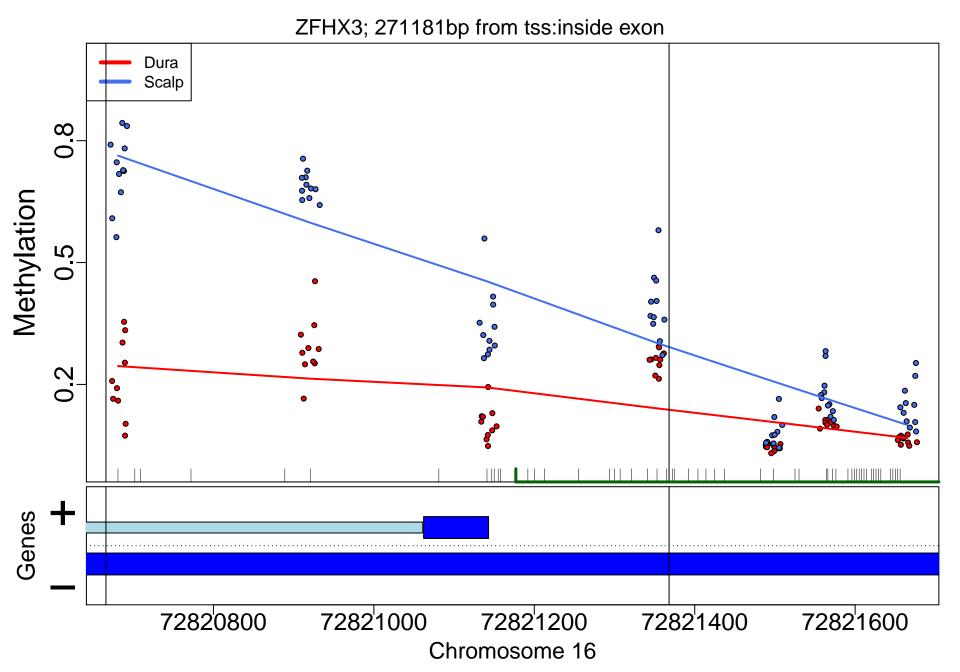


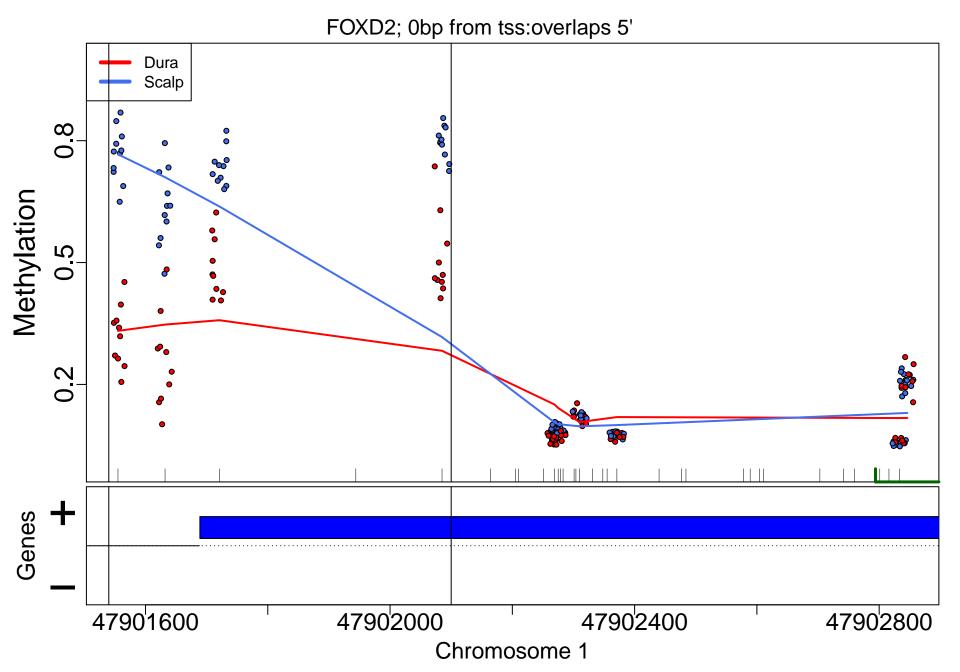


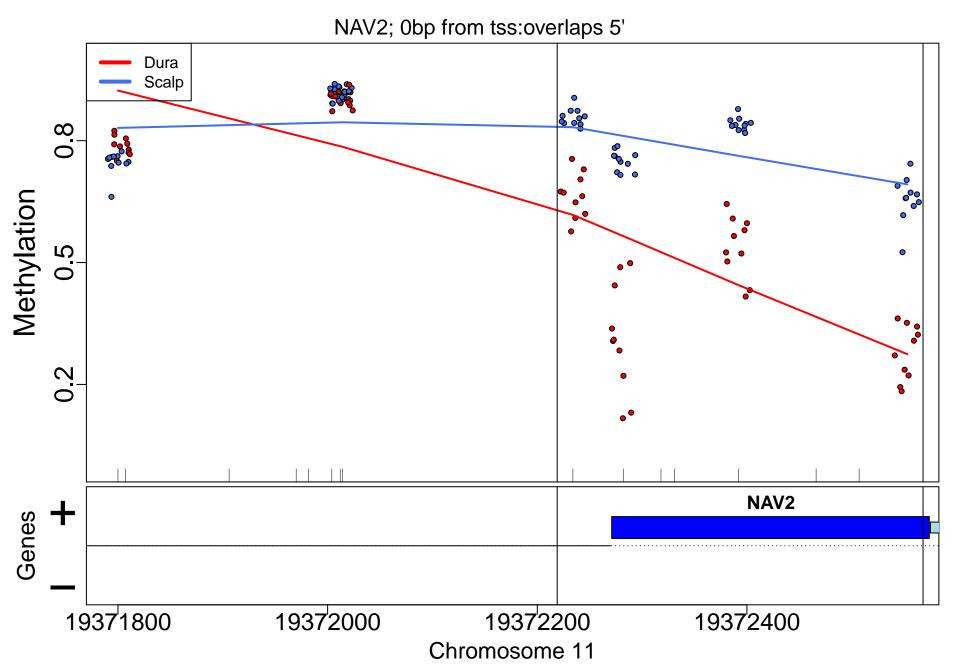


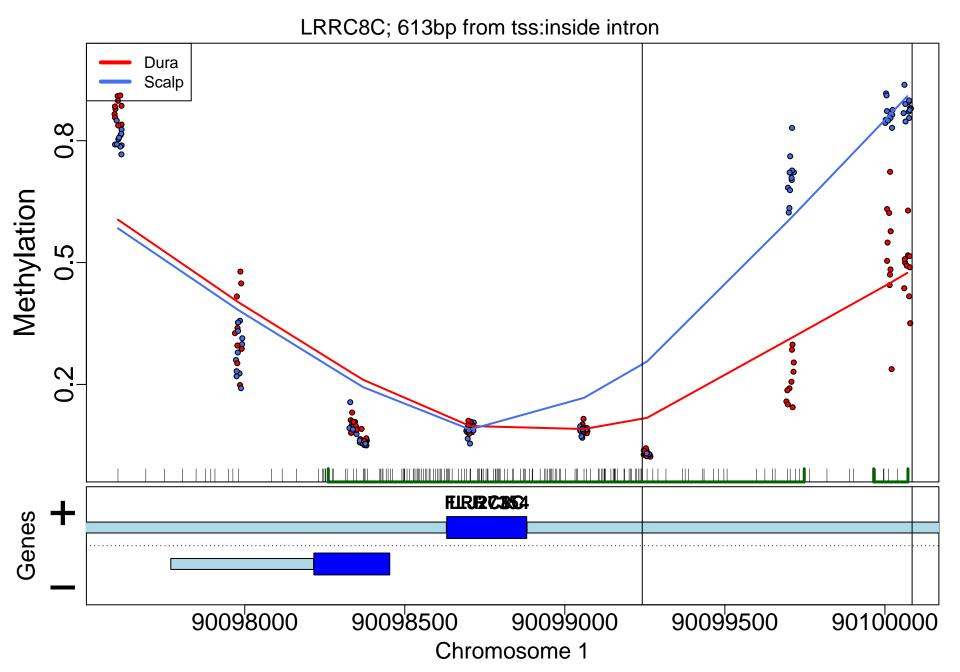


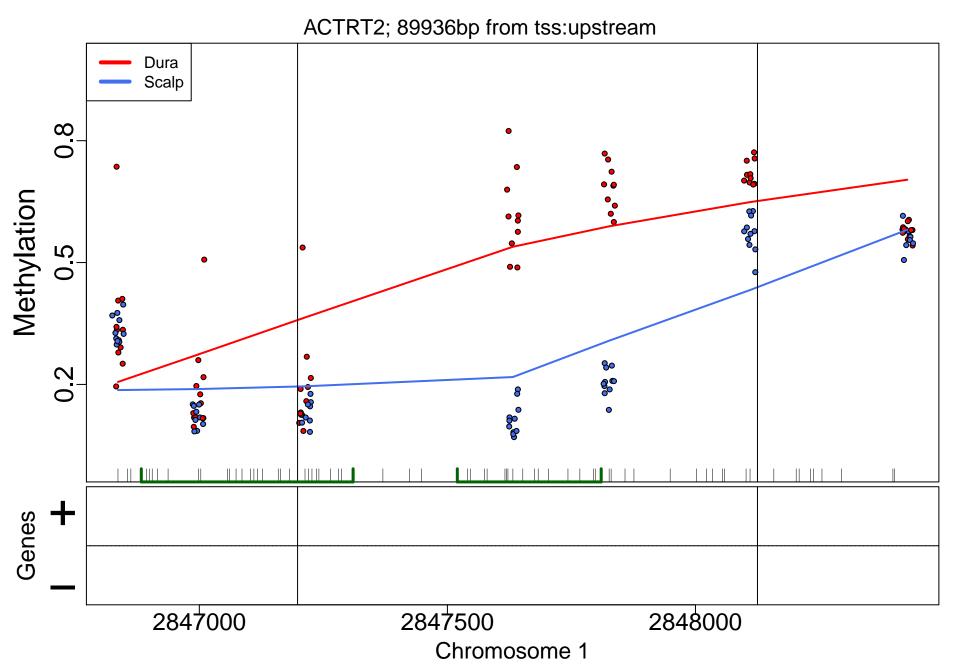


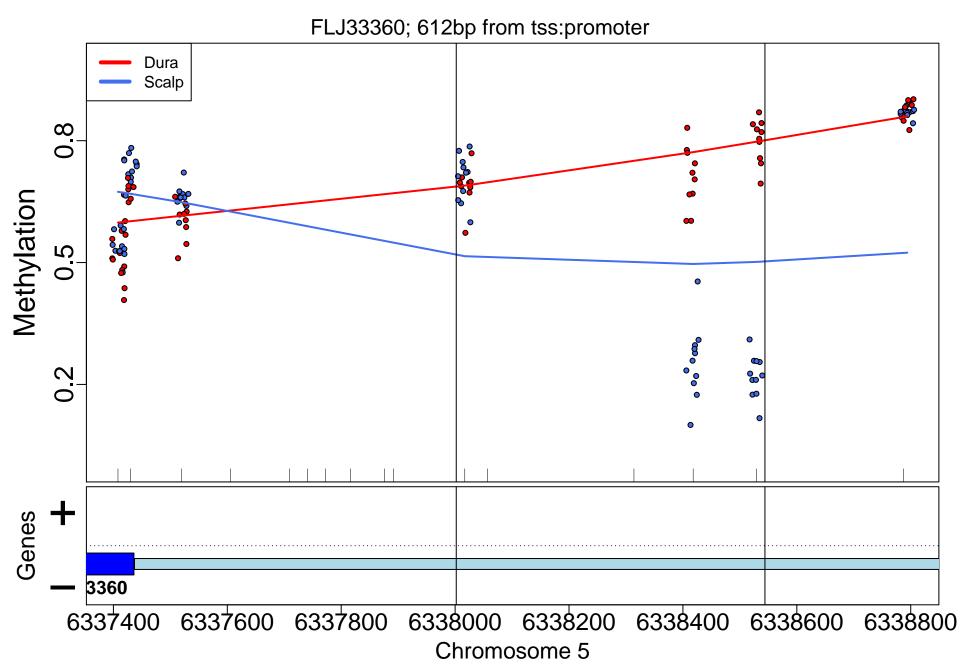


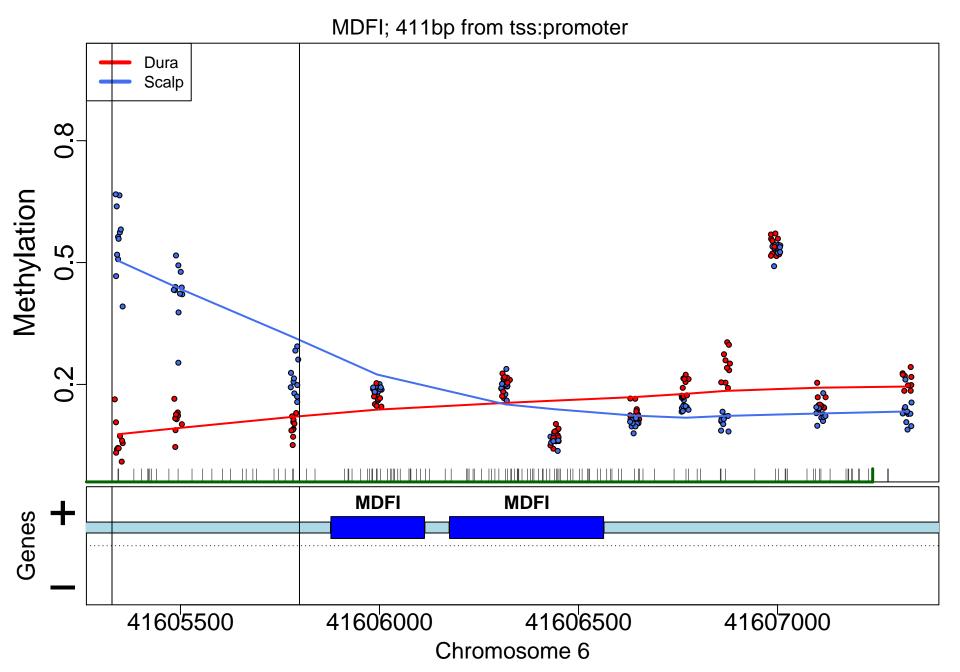


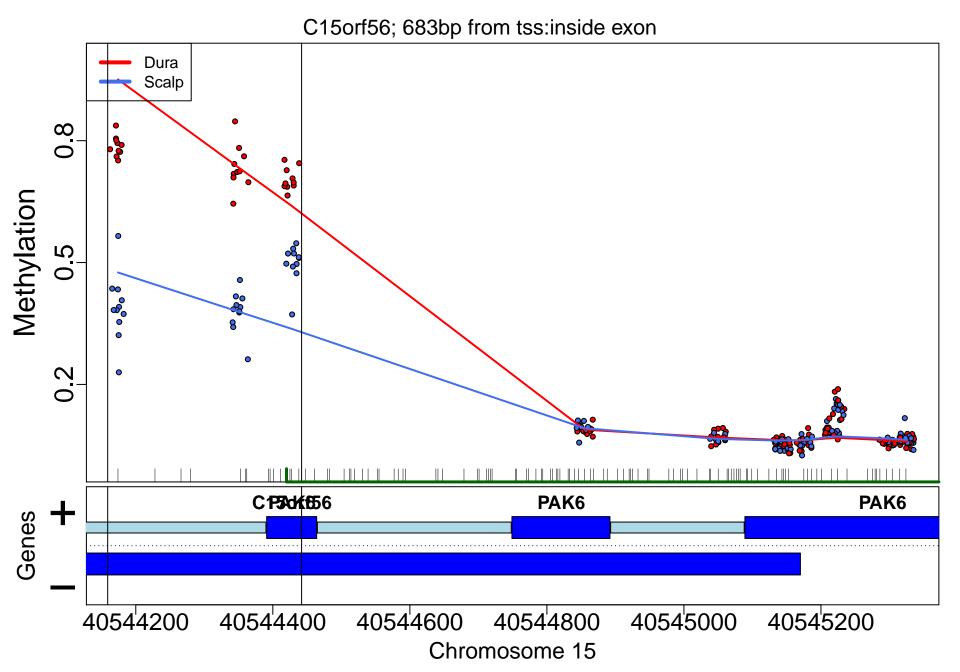


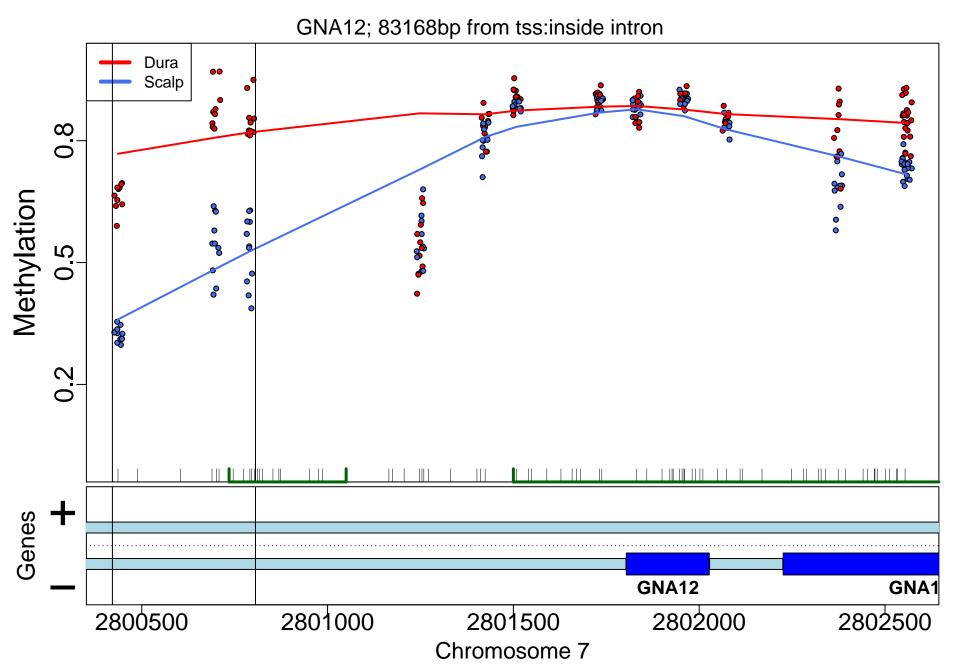


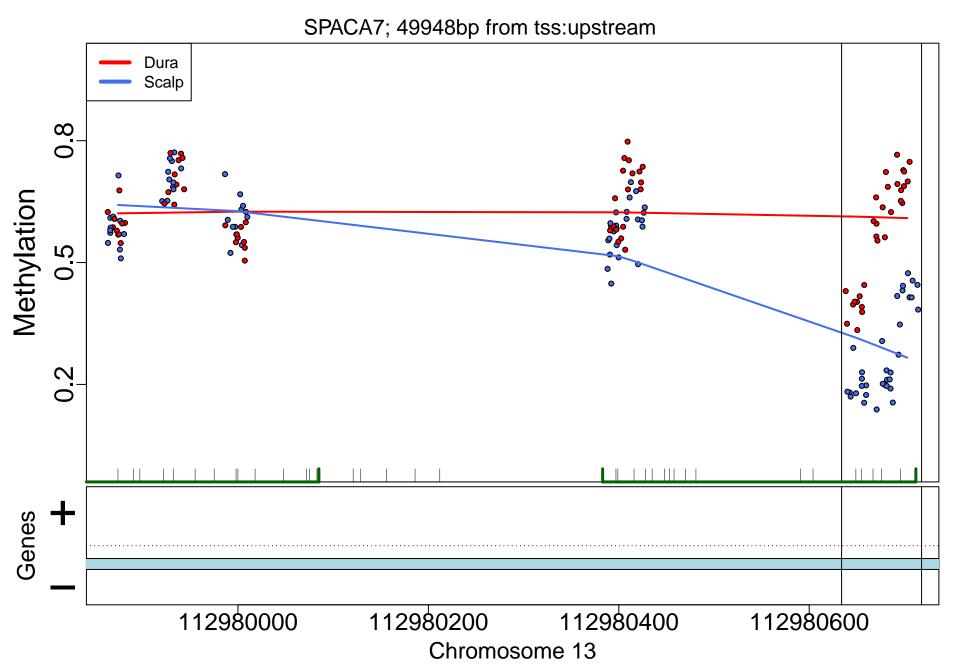




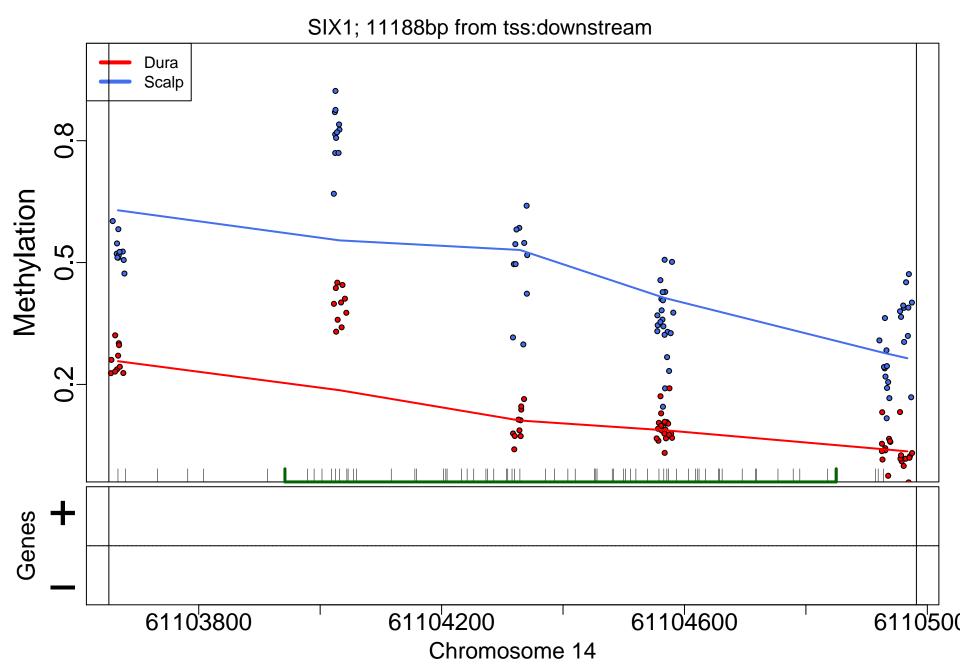


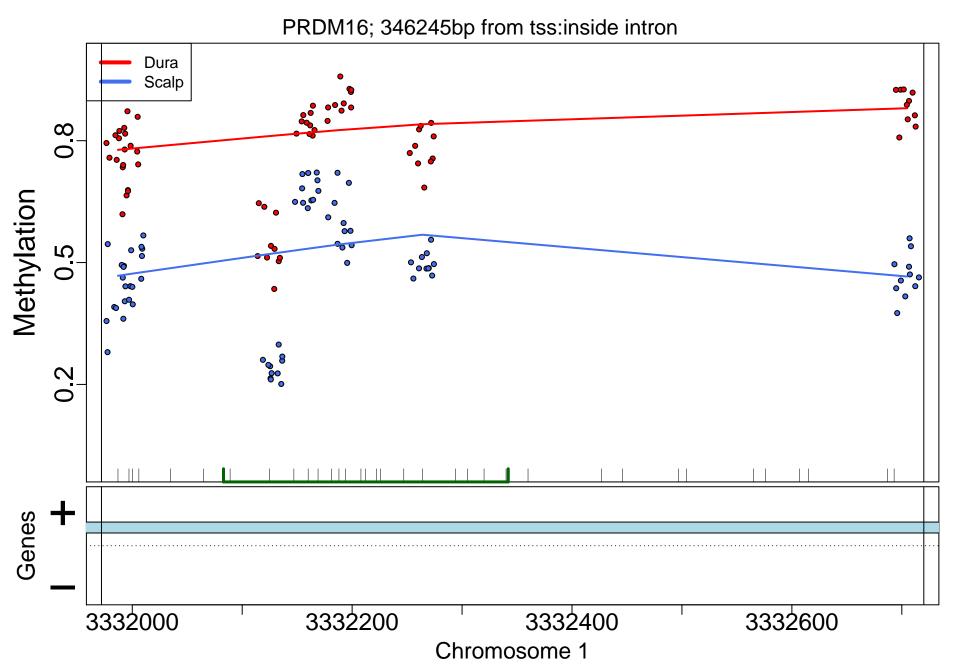


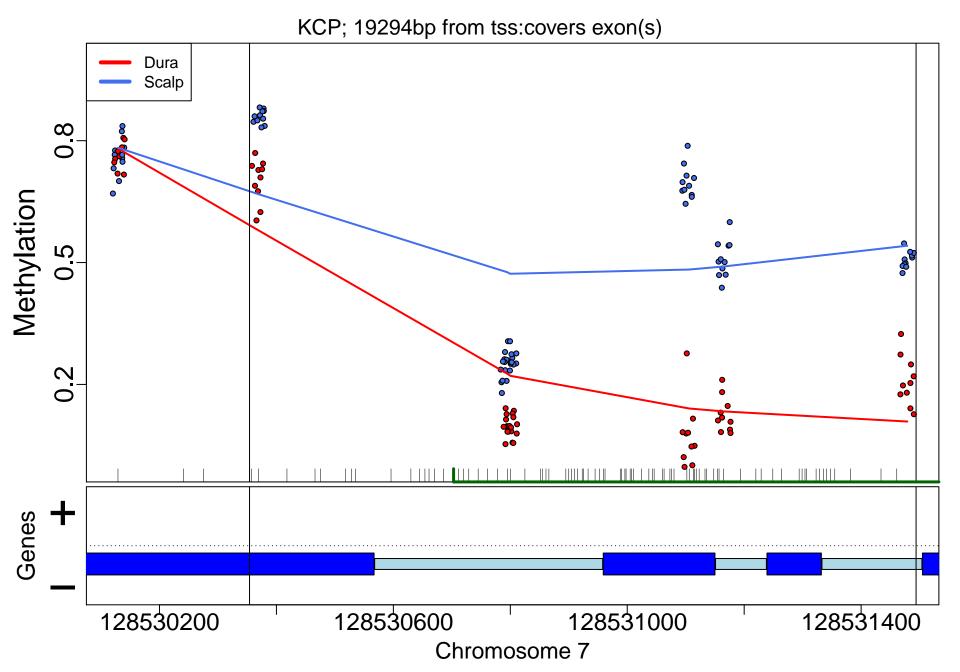


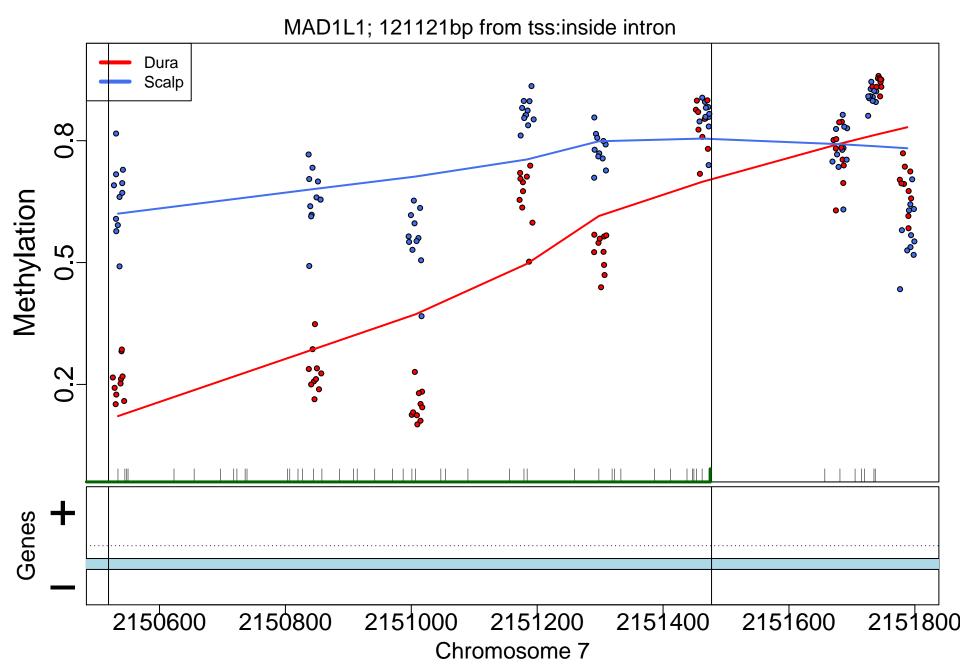


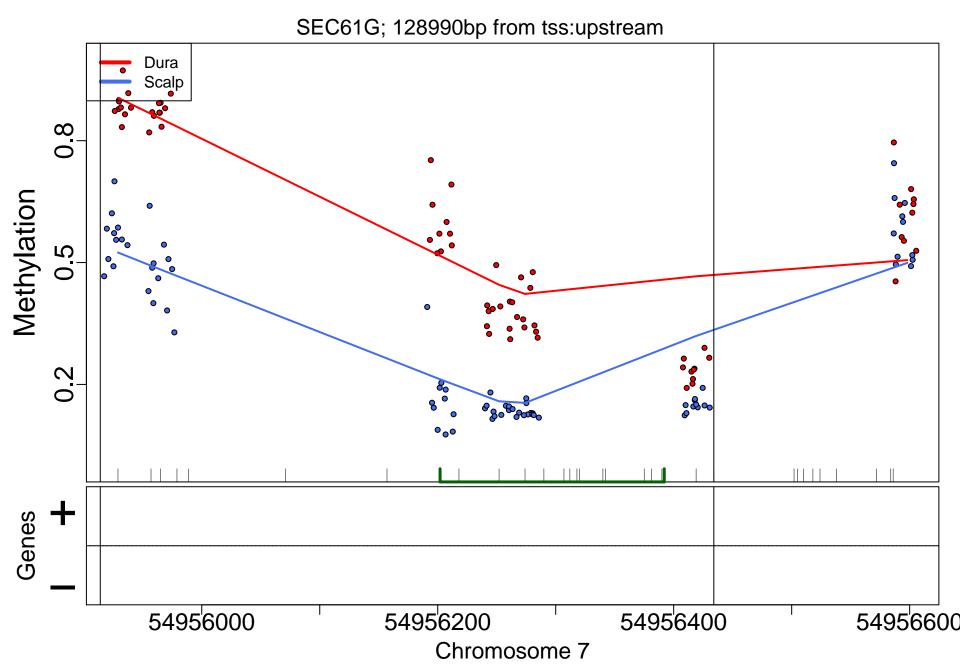
KRTAP17-1; 0bp from tss:overlaps 5' Dura Scalp <u>∞</u>. Methylation 0.2 Genes KRTAP17-1 39471400 39472200 39471800 39472000 39472400 39471600 Chromosome 17

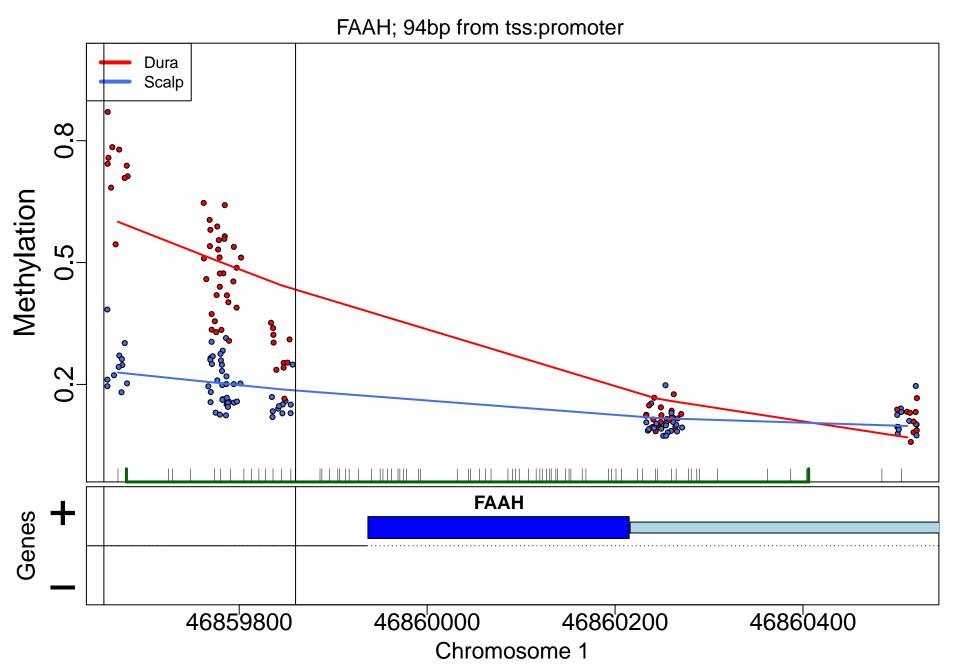


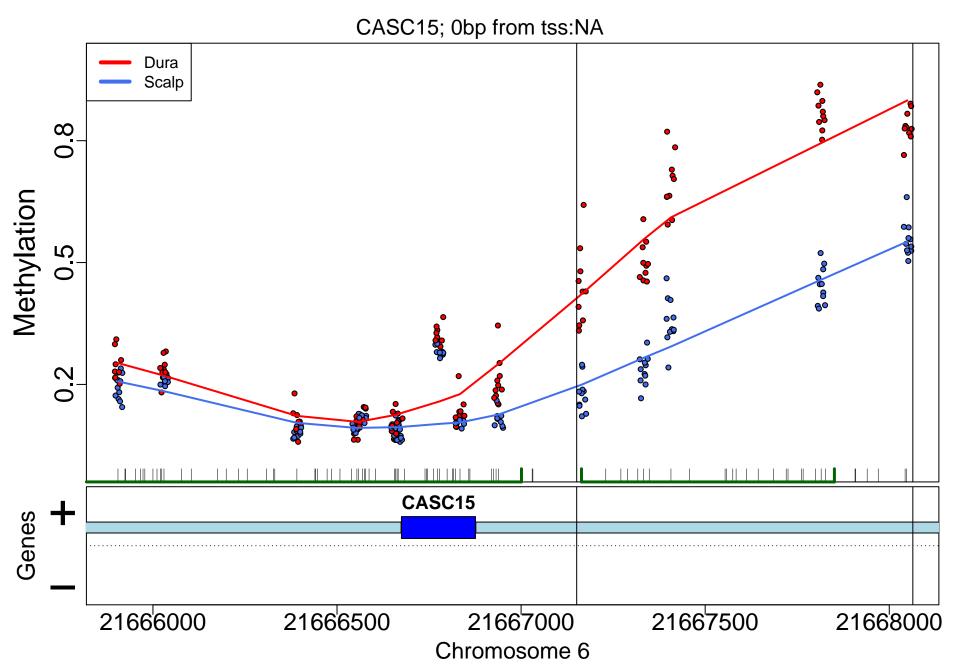


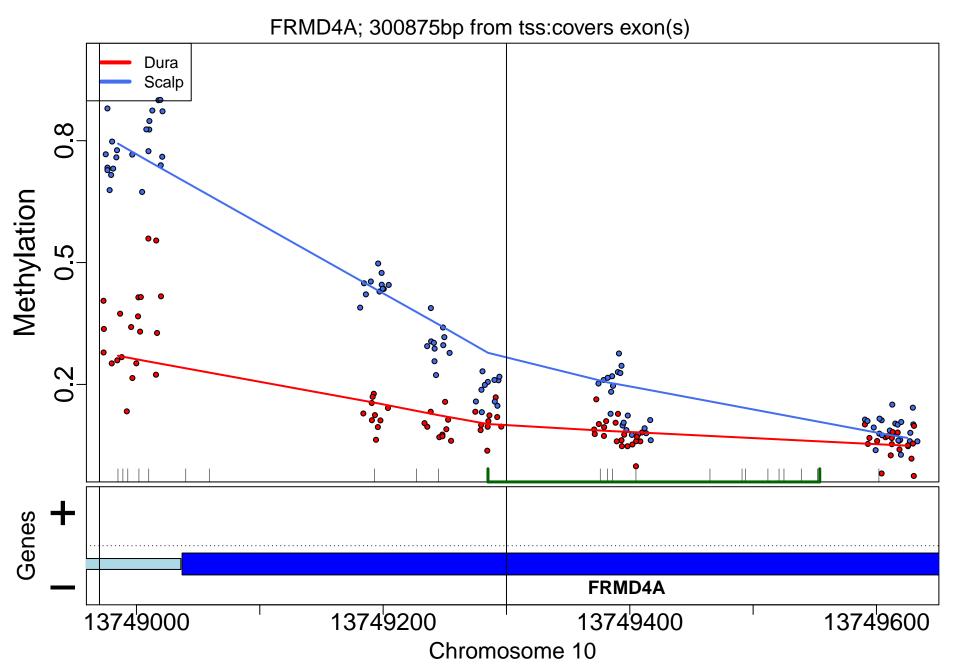


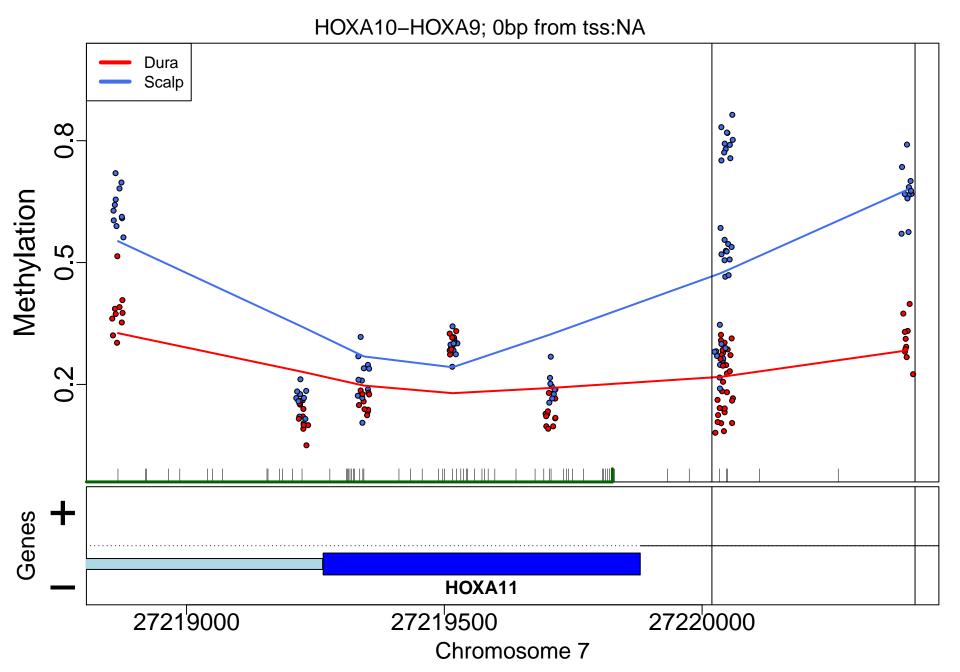


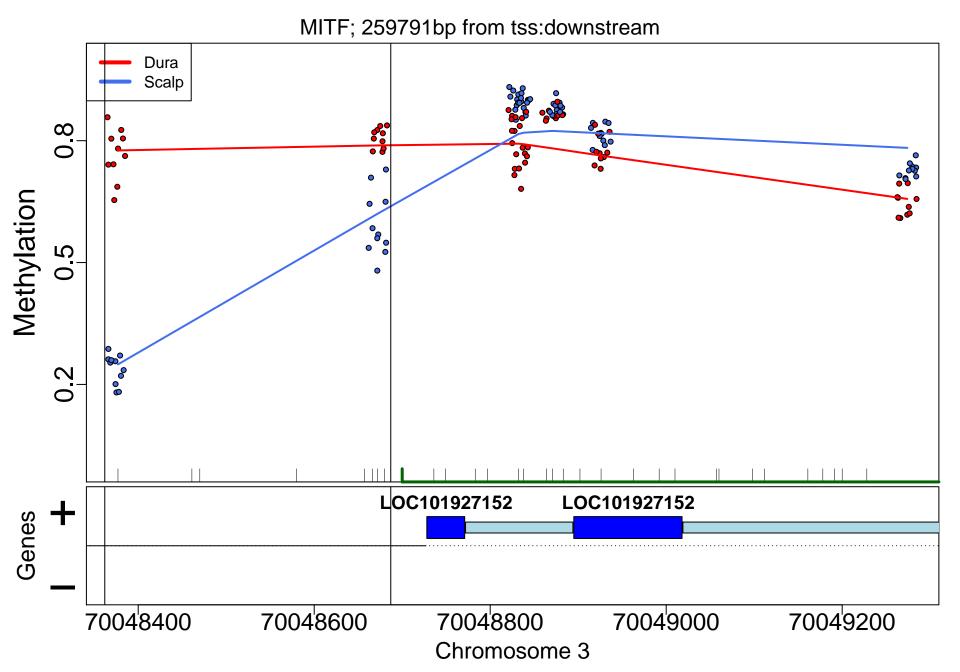


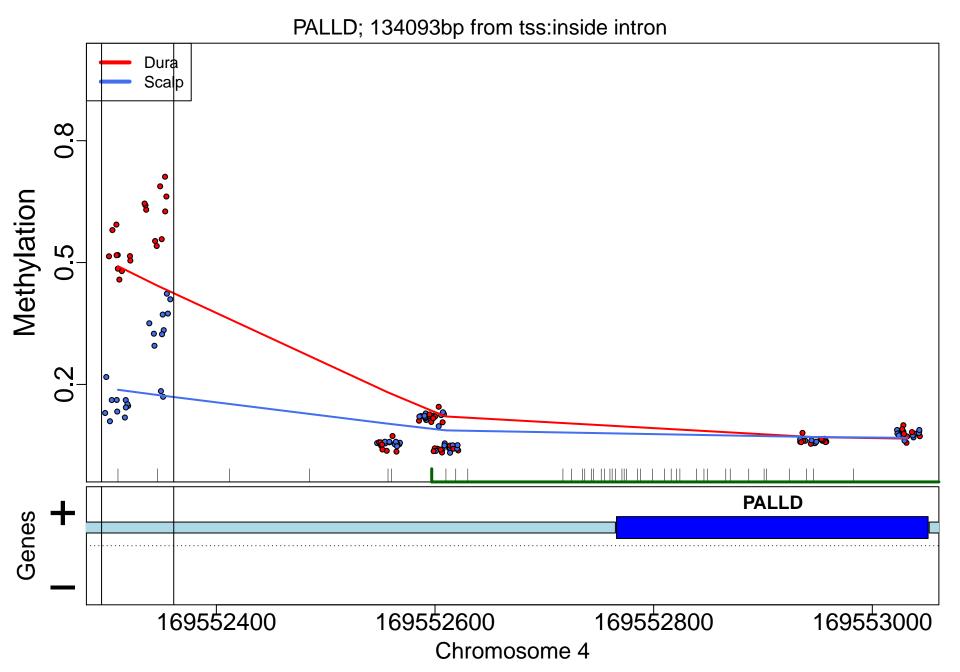


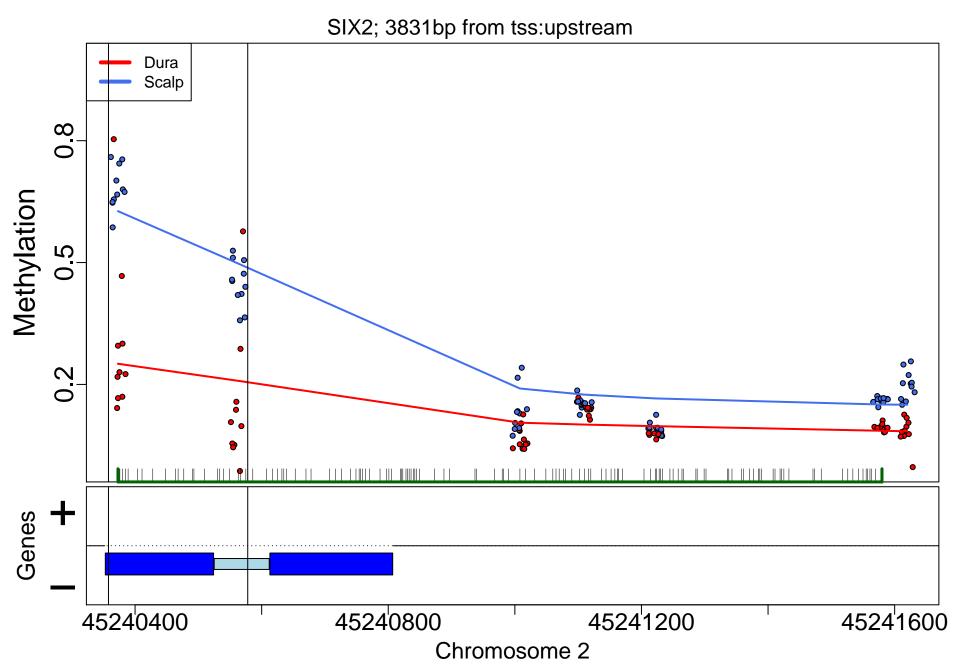


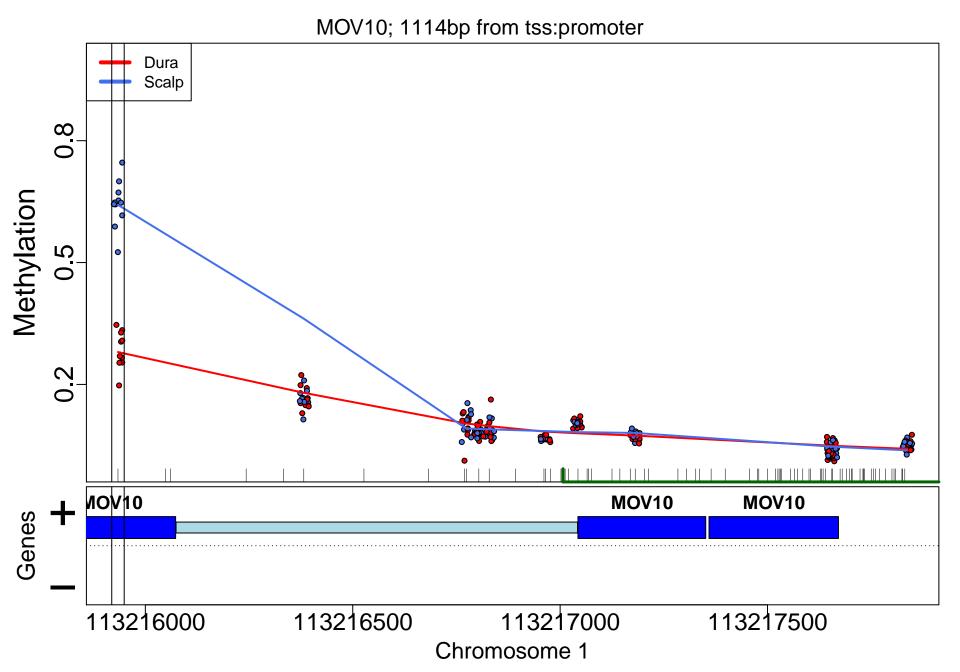


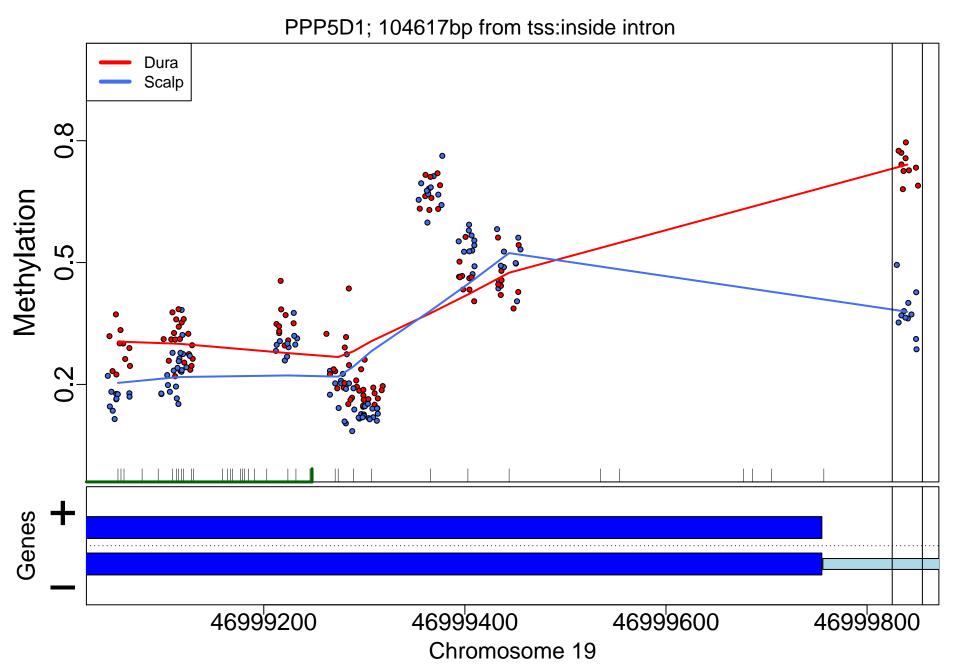




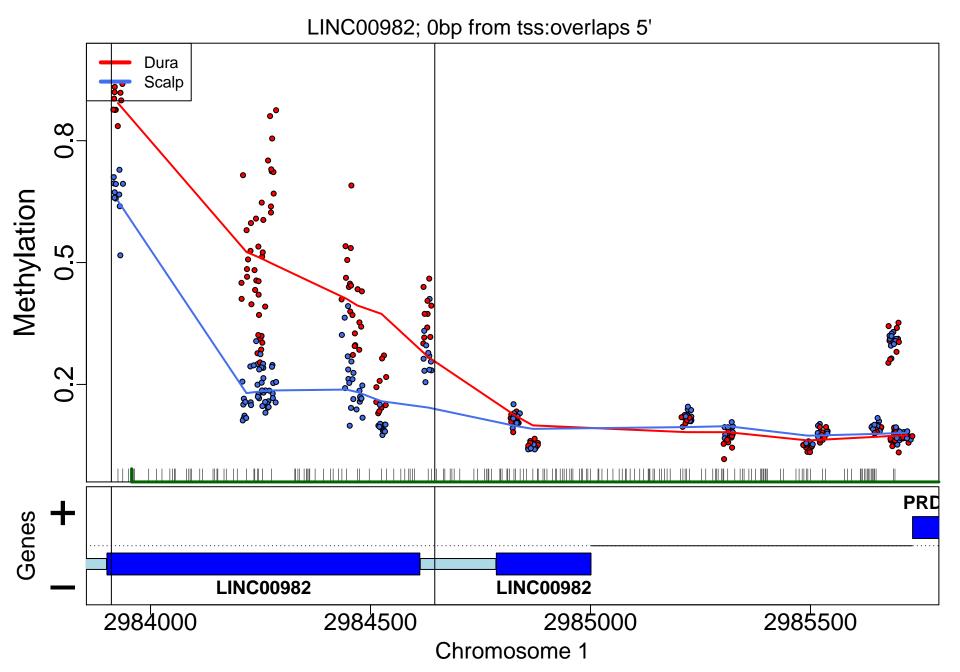


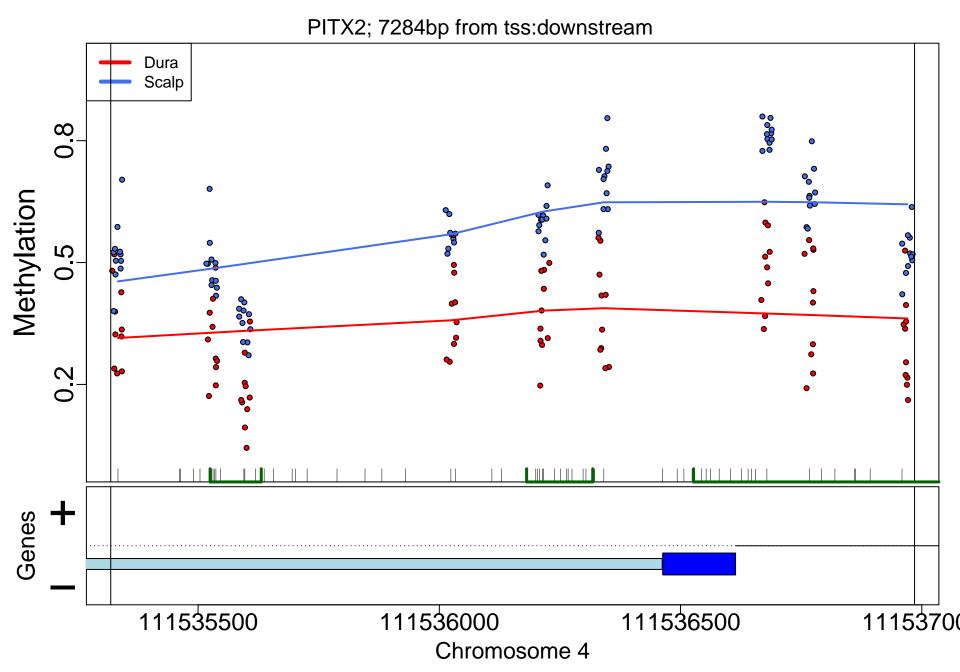


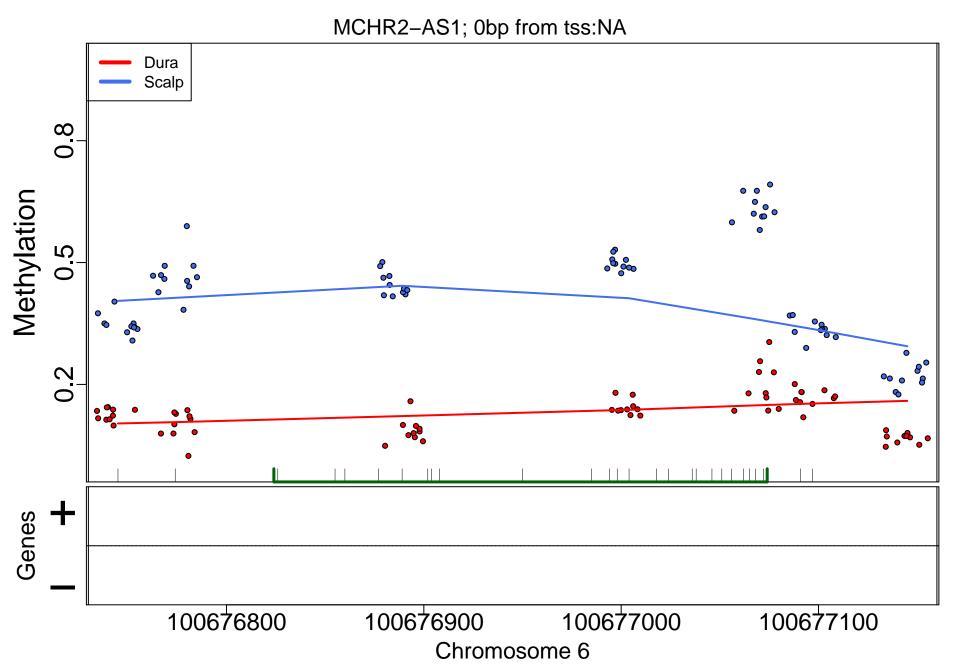


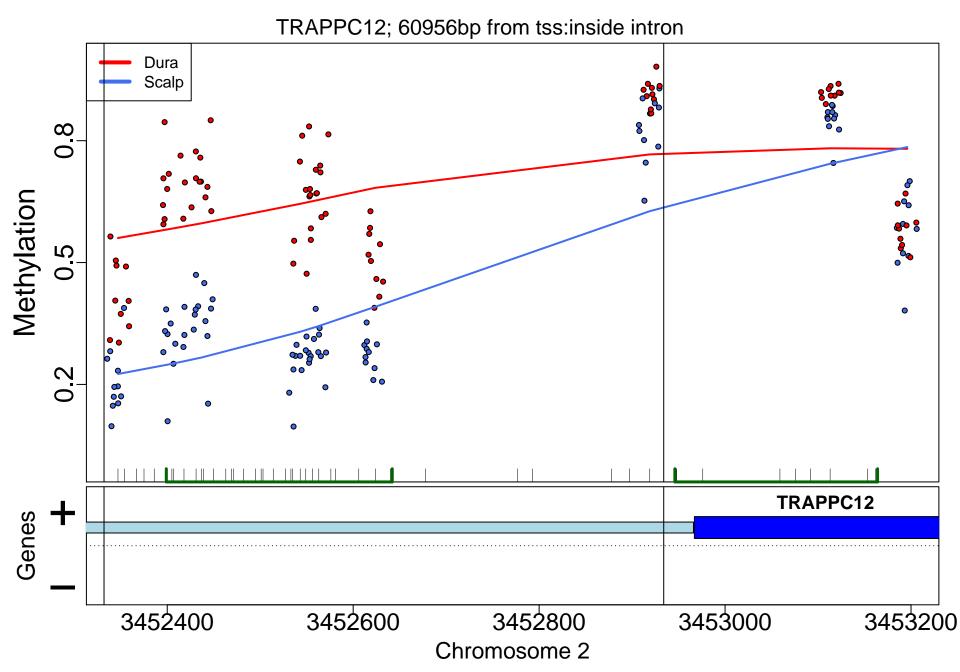


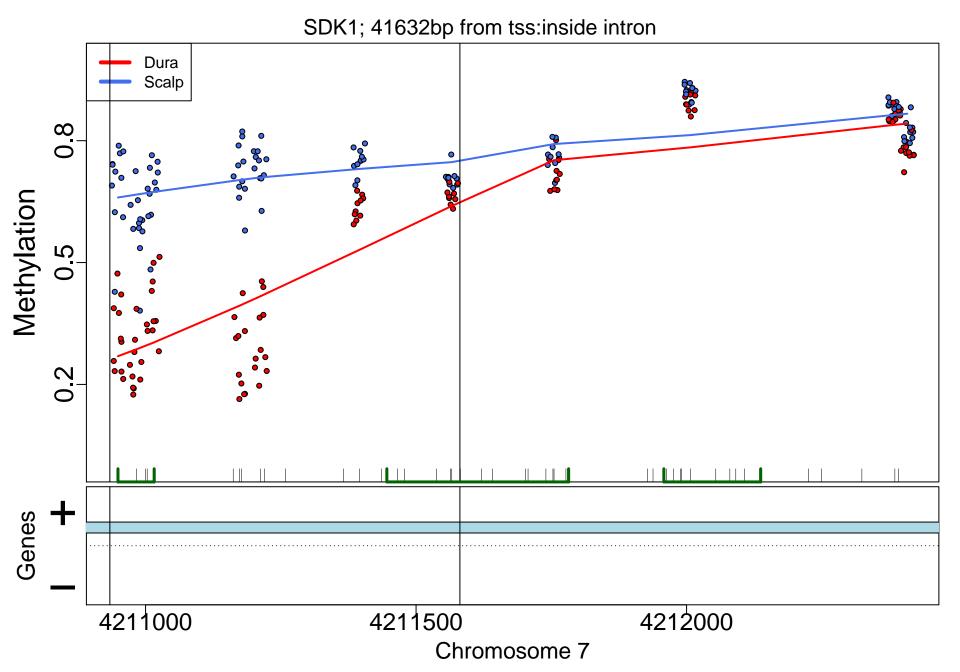
TBX20; 929bp from tss:promoter Dura Scalp 80. Methylation 0.2 Genes 35294500 35295500 35296500 35297500 Chromosome 7

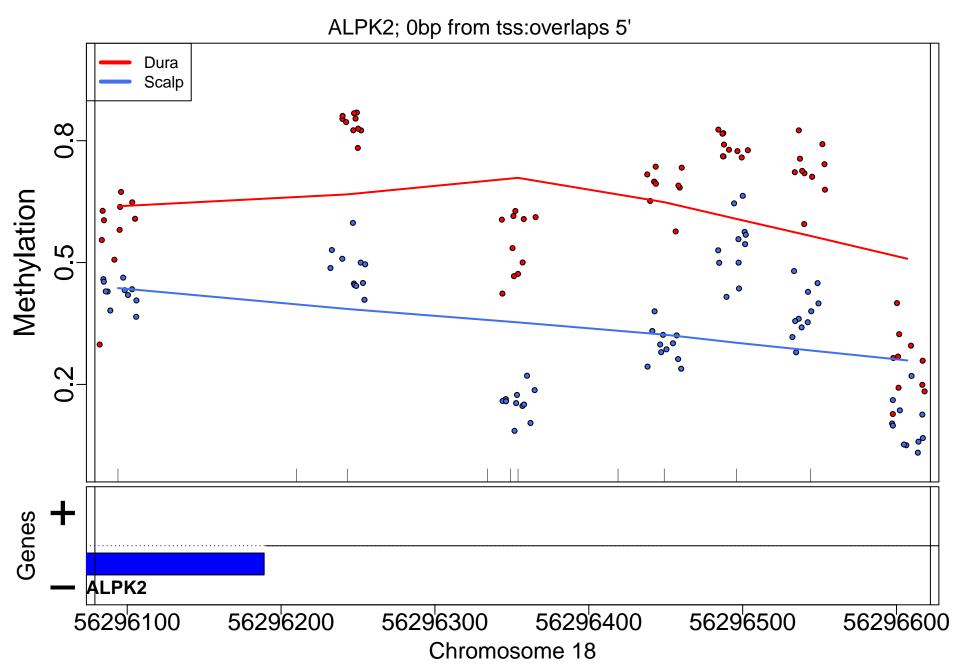




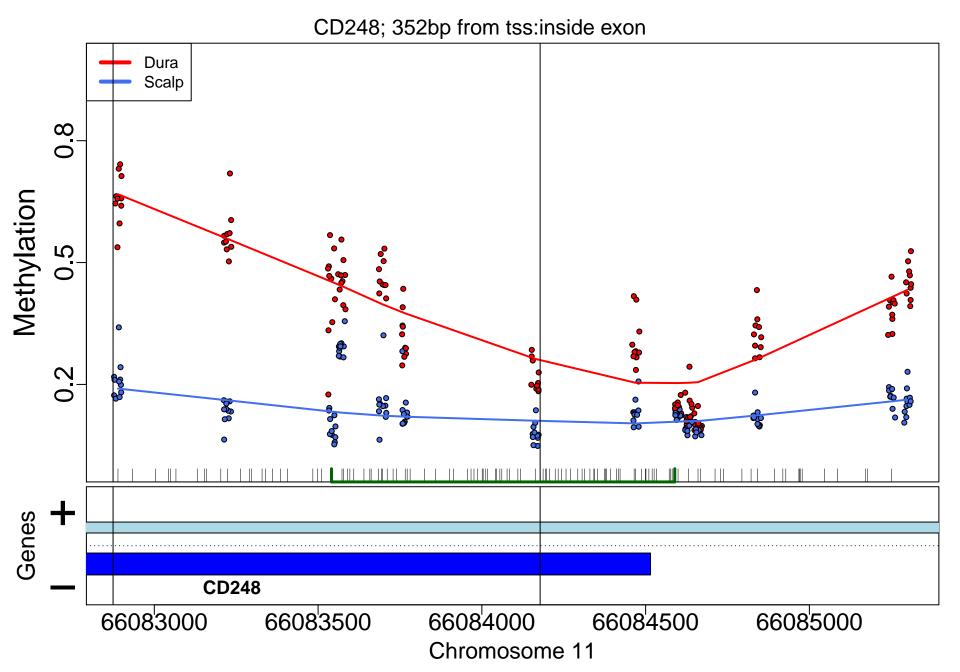


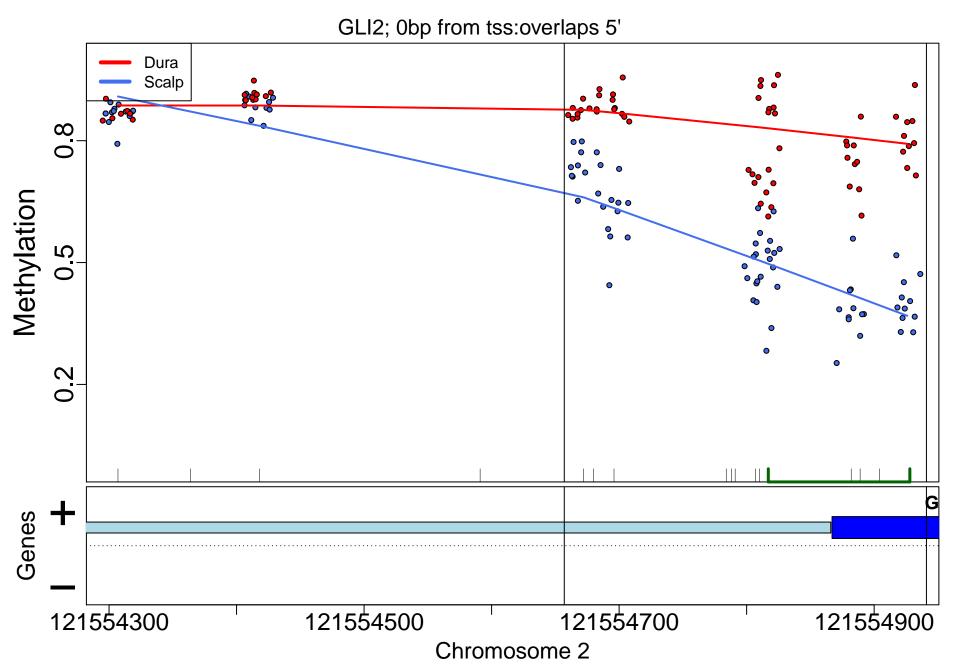


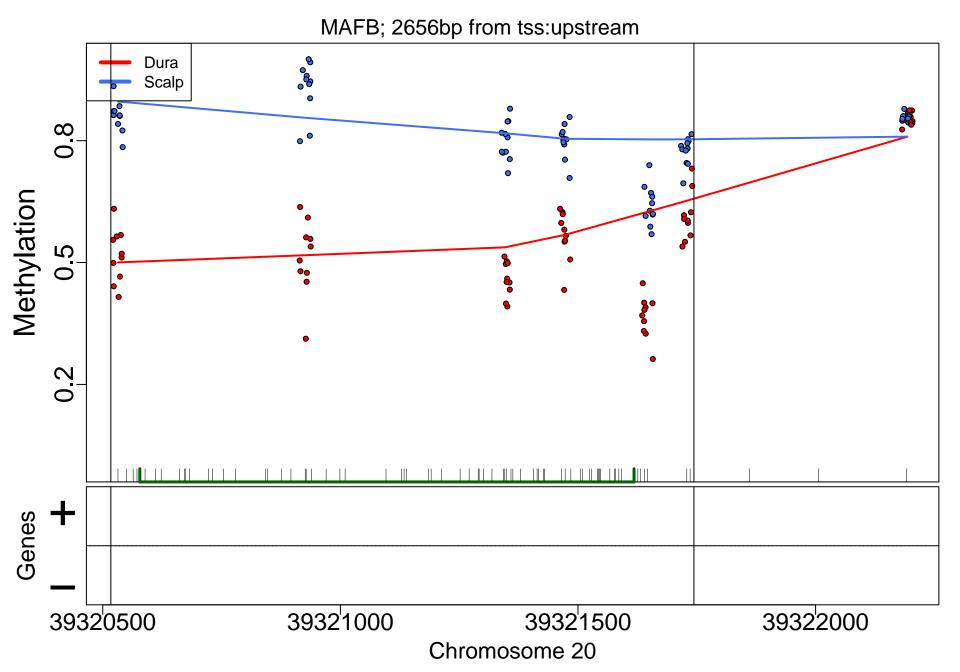


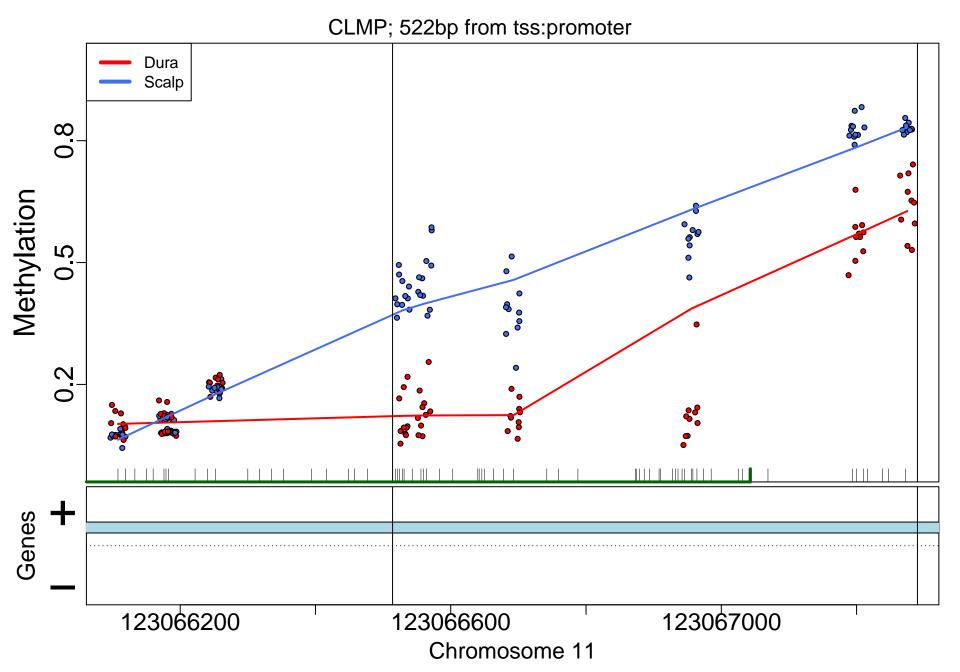


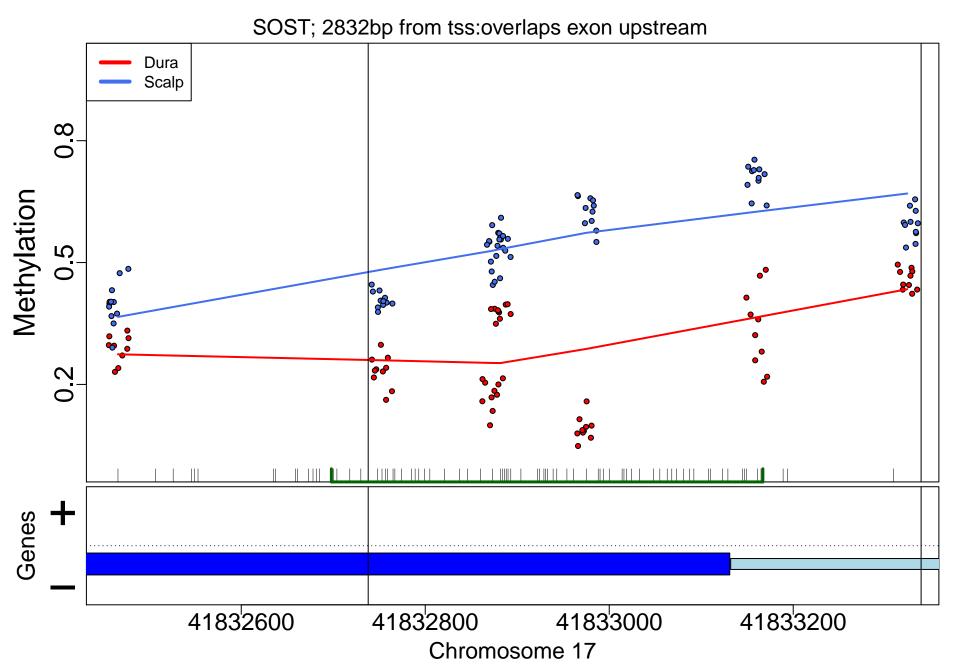
CIR1; 51188bp from tss:downstream Dura Scalp <u>∞</u>. Methylation 0.5 0.2 Genes 175208400 175208200 175208600 175208800 175209000 175209200 Chromosome 2

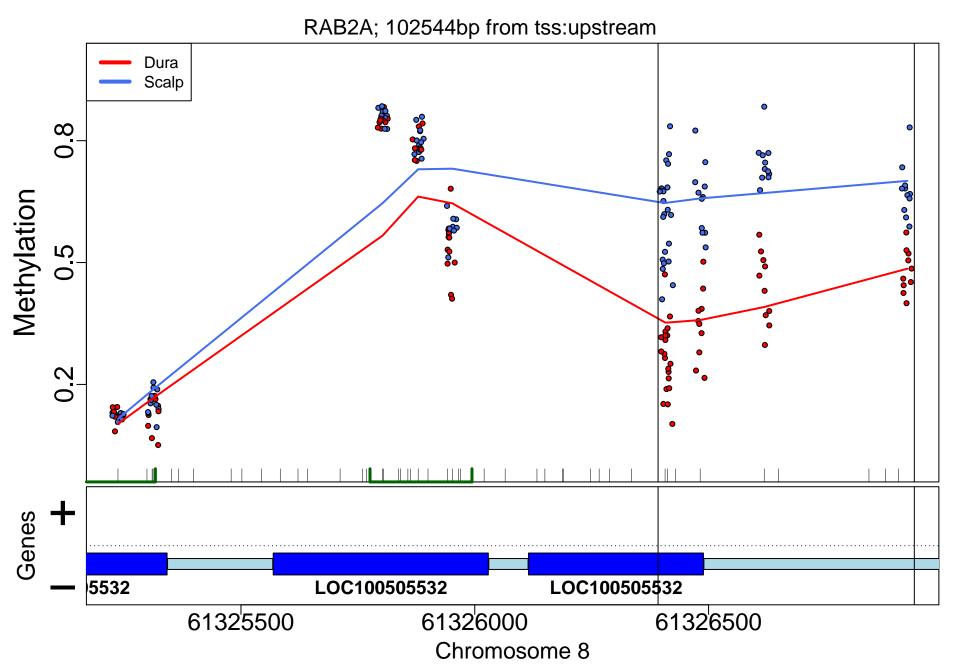


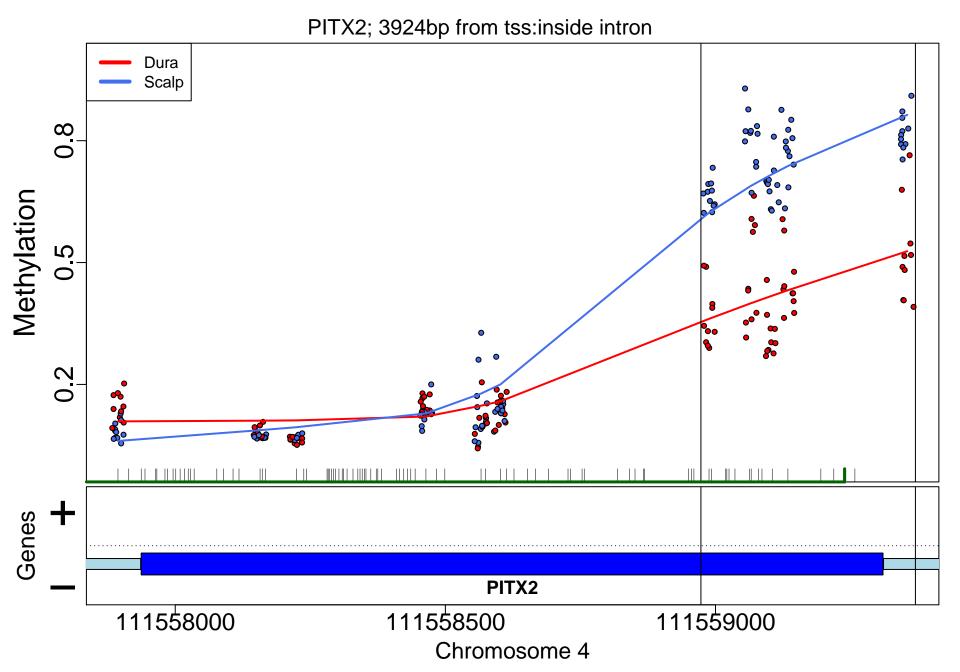


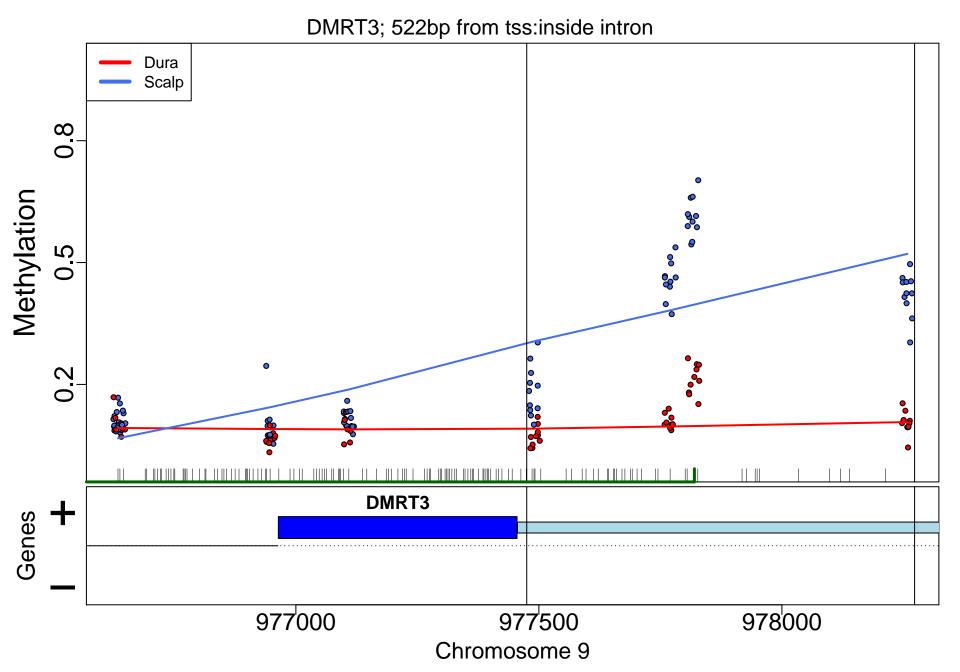


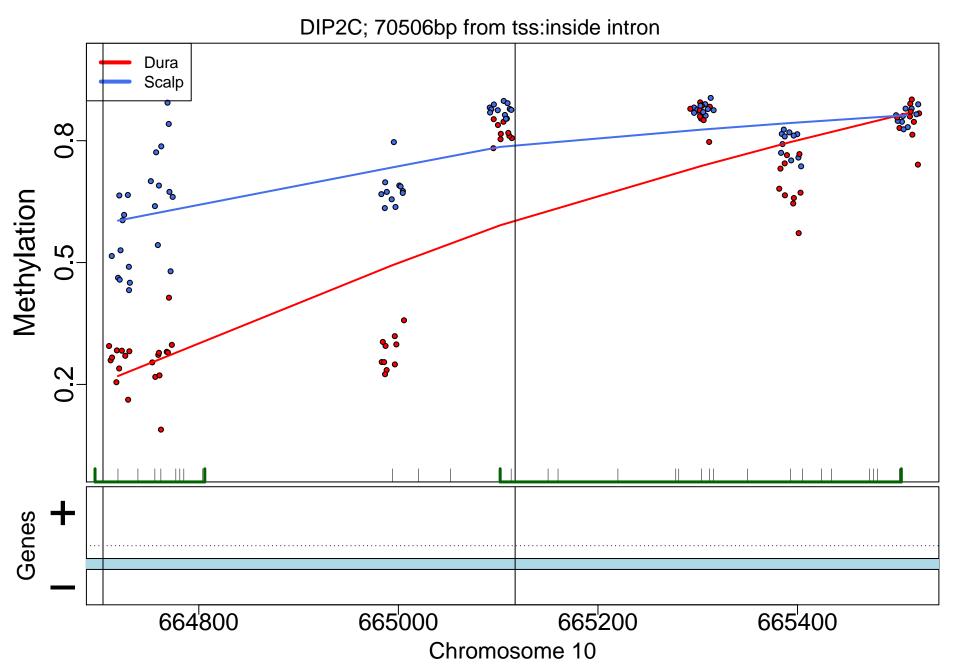


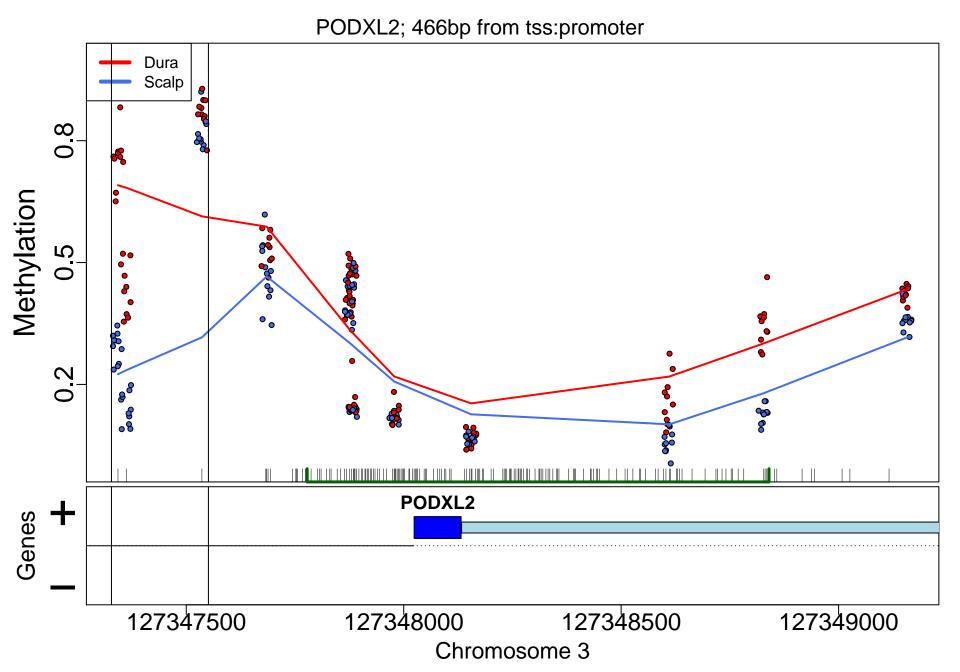


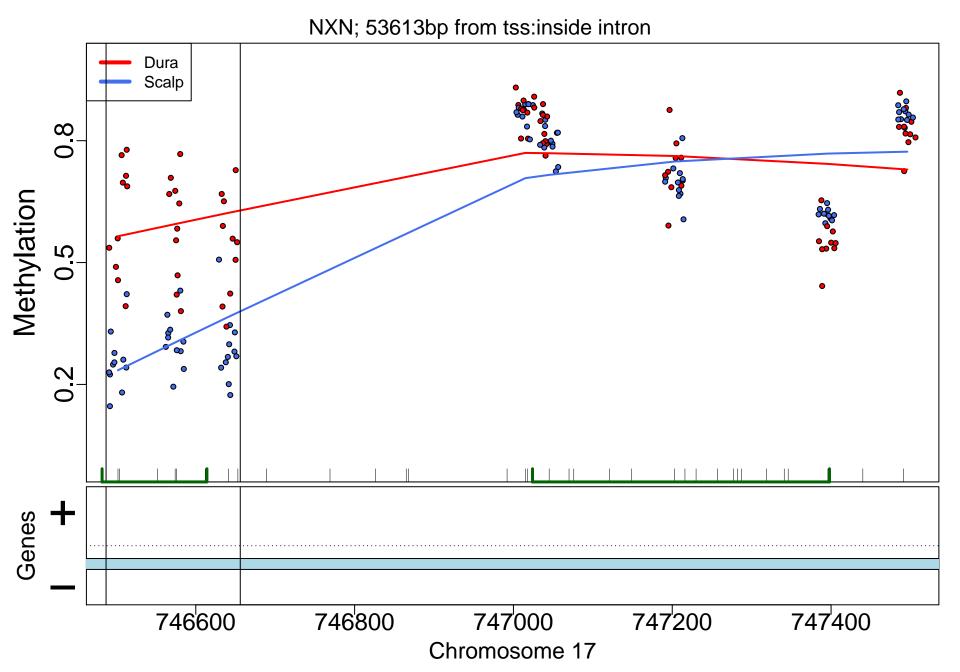


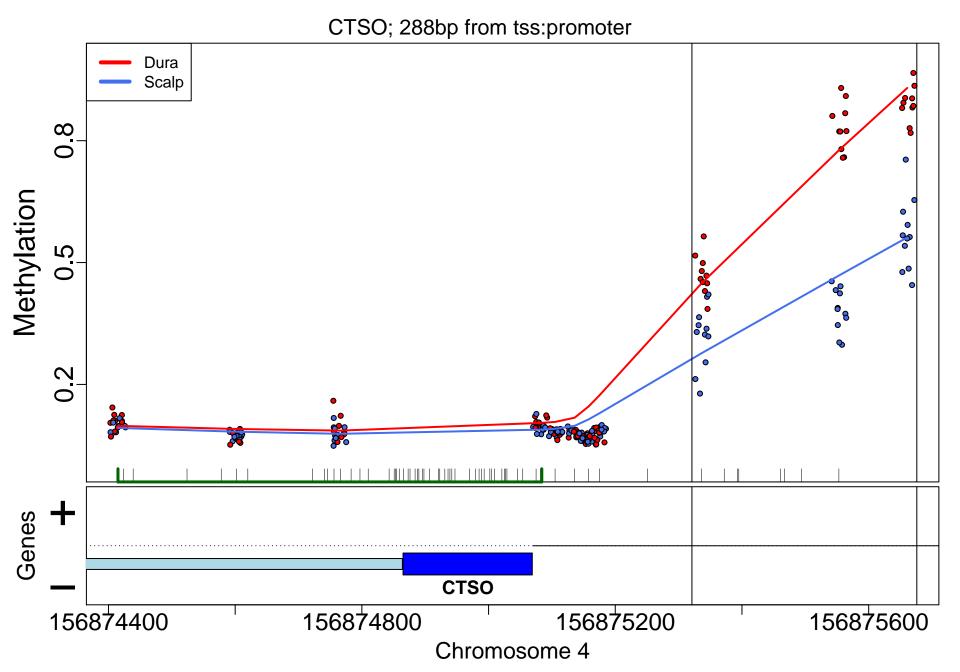


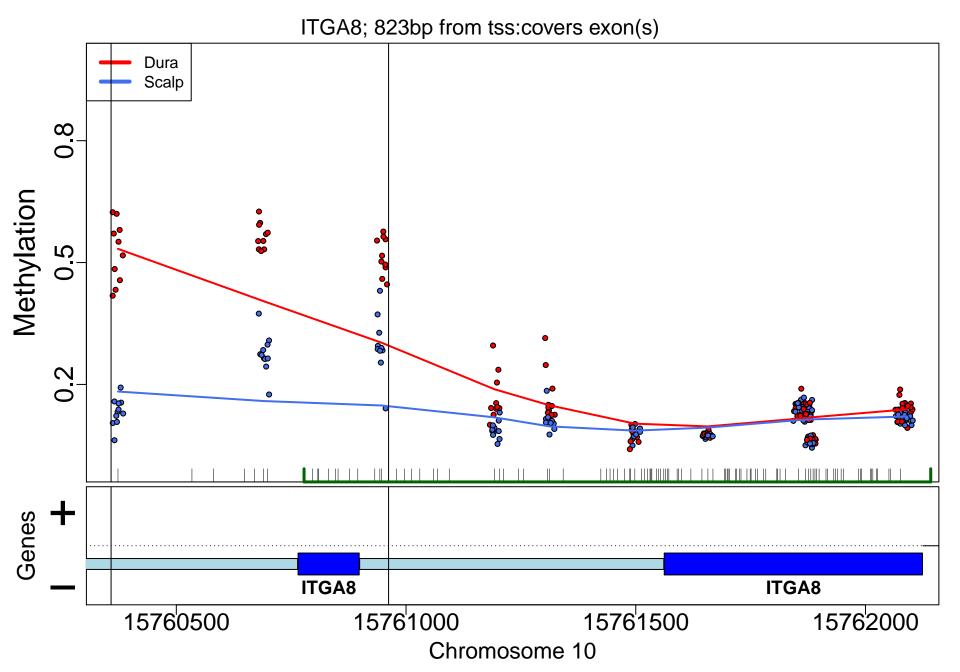


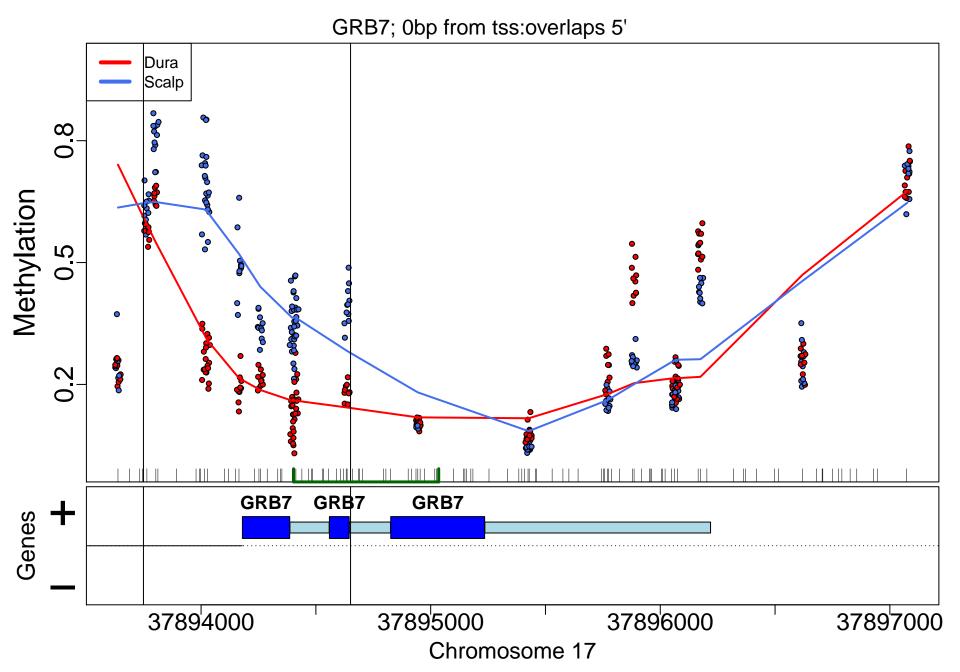


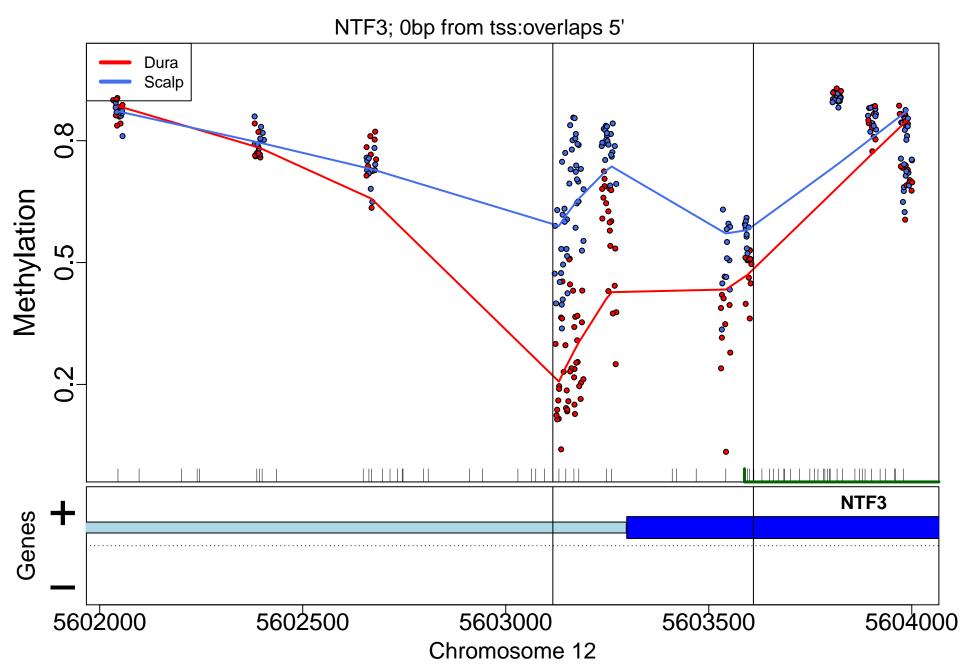




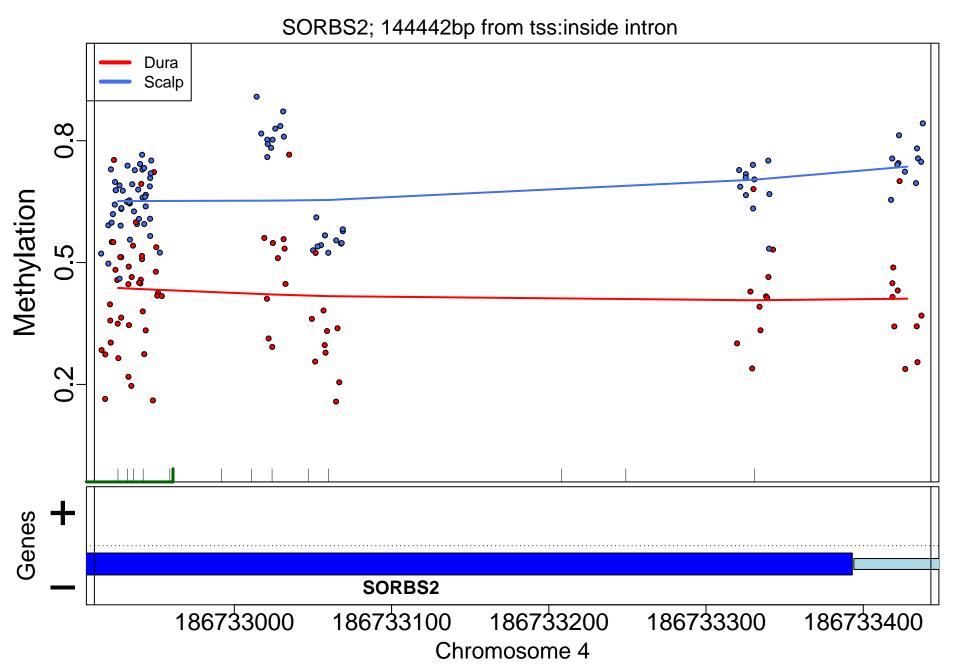


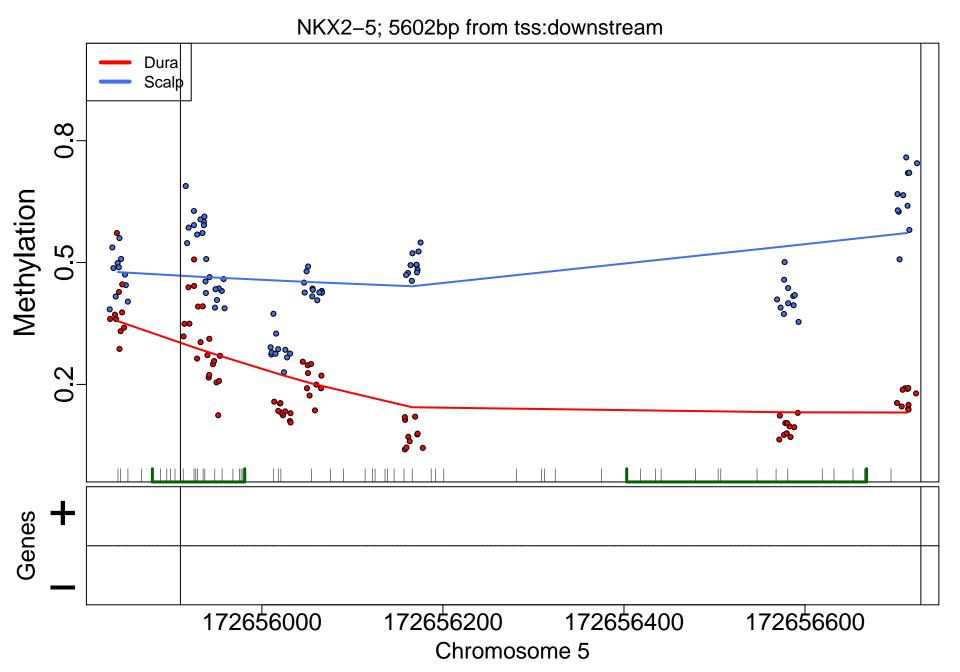


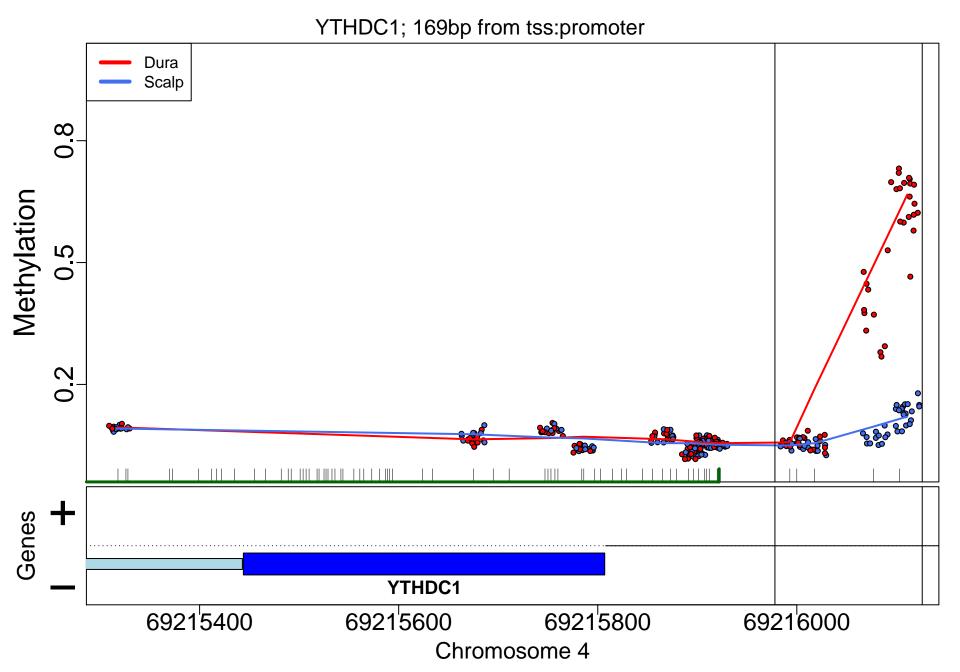


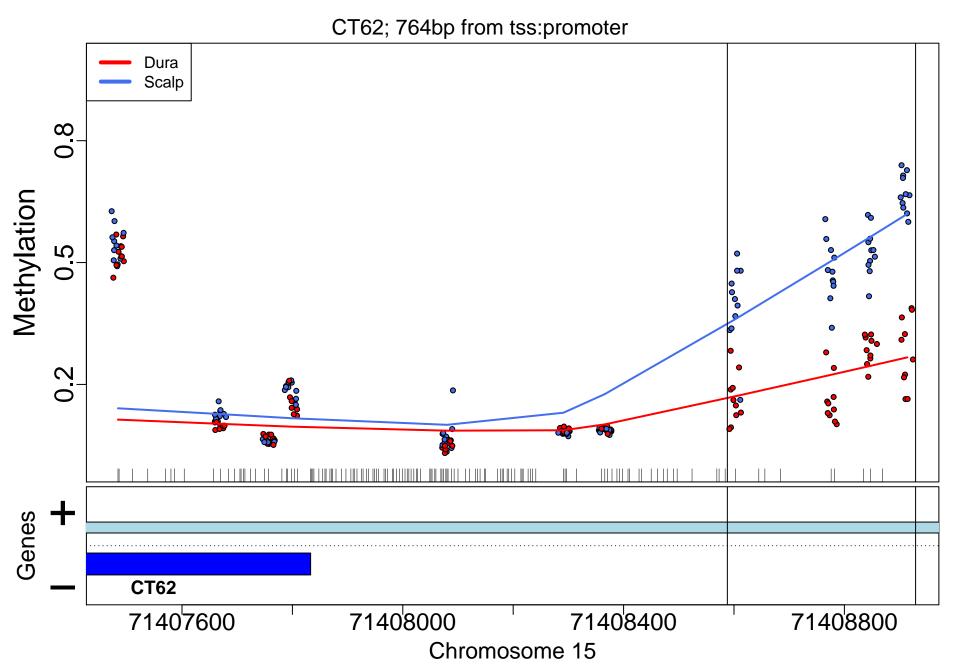


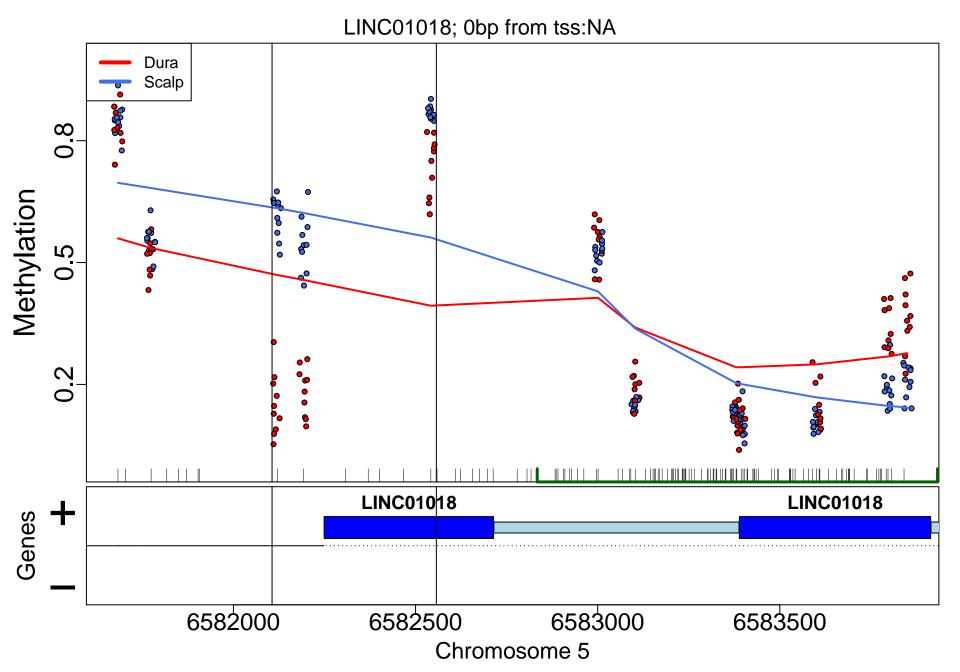
SMOC2; 55861bp from tss:upstream Dura Scalp <u>ω</u>_ Methylation 0.2 Genes 168785300 168785500 168785700 168785900 Chromosome 6

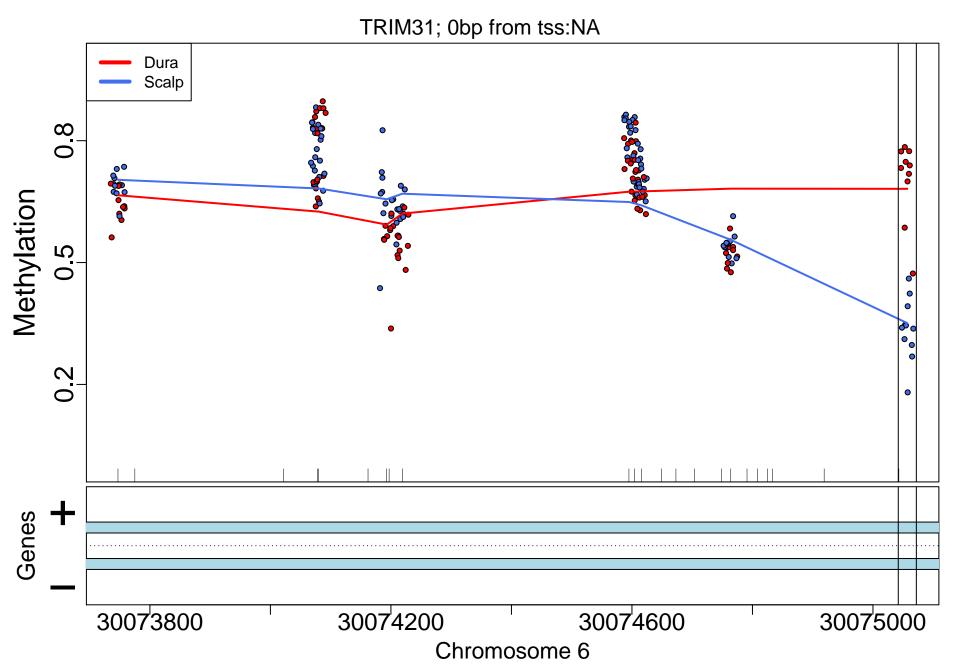


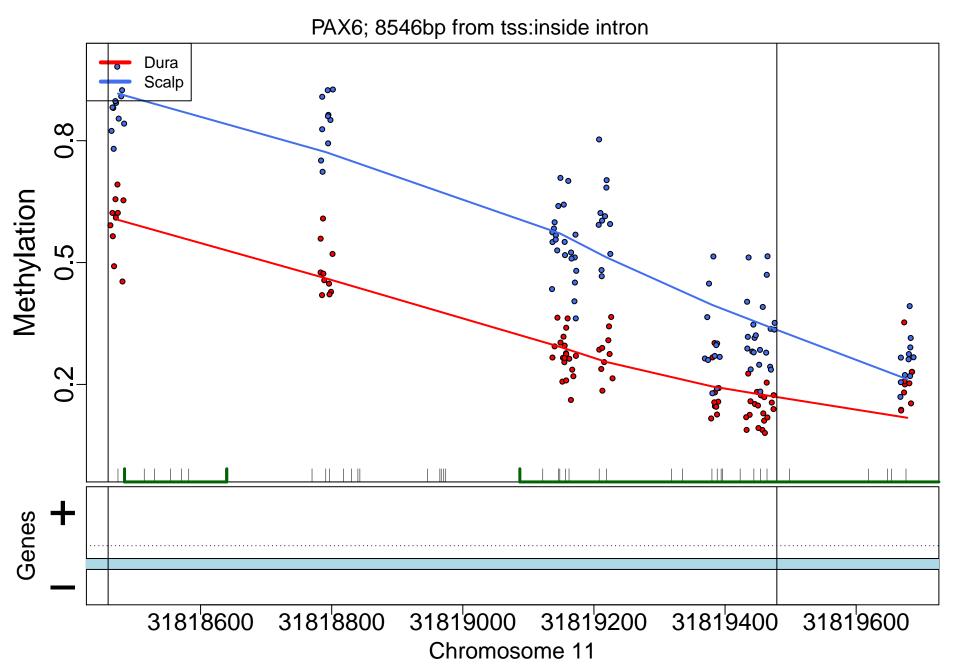


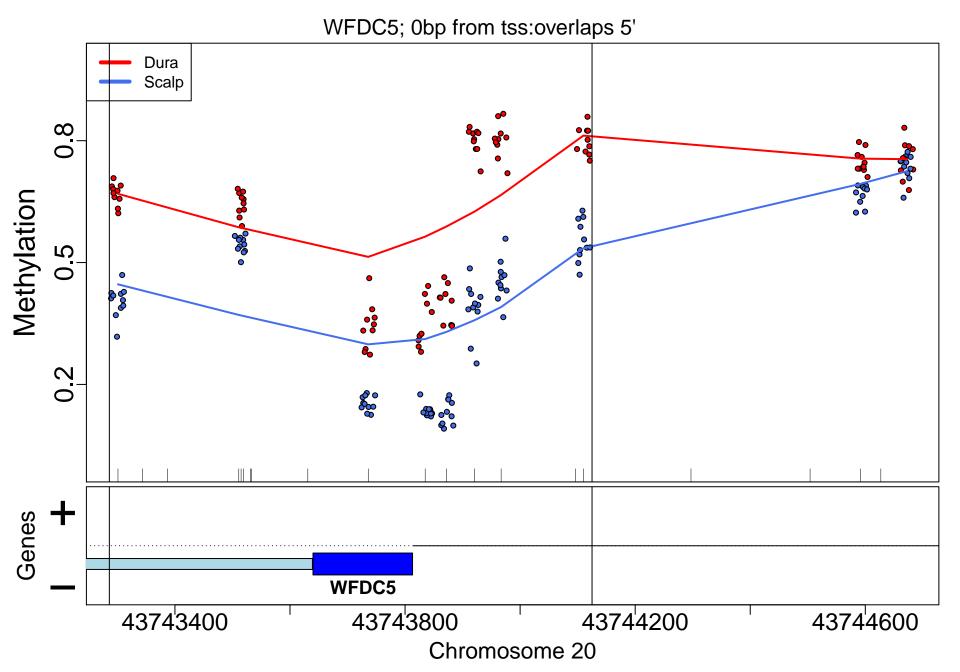


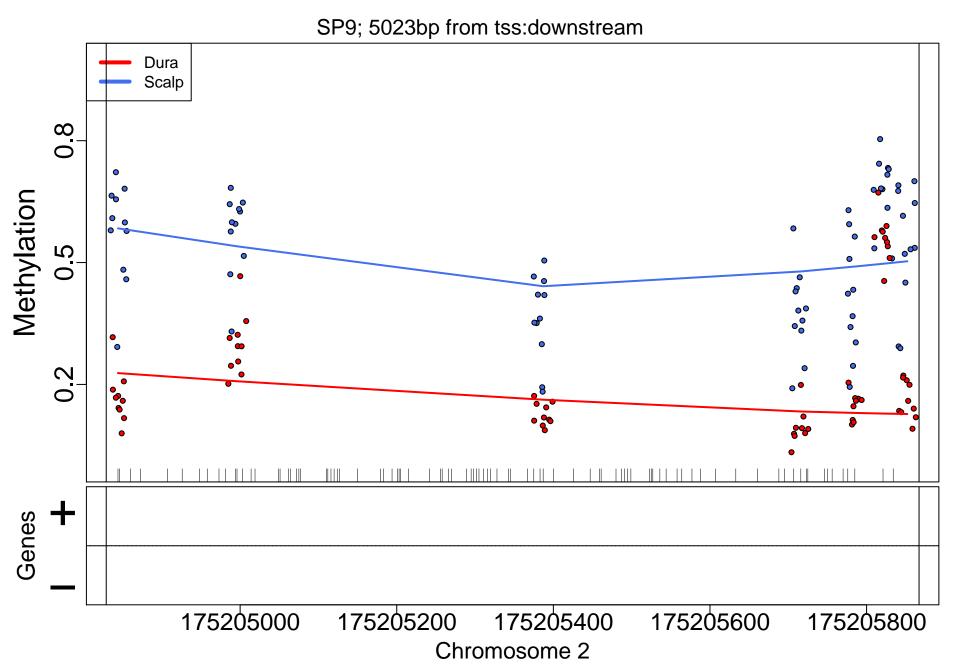


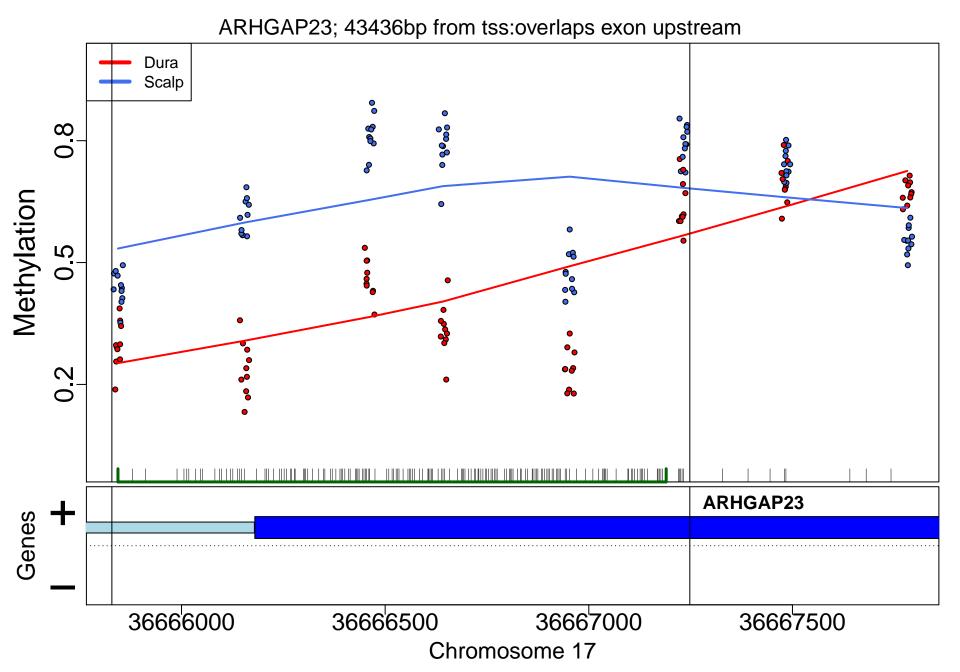


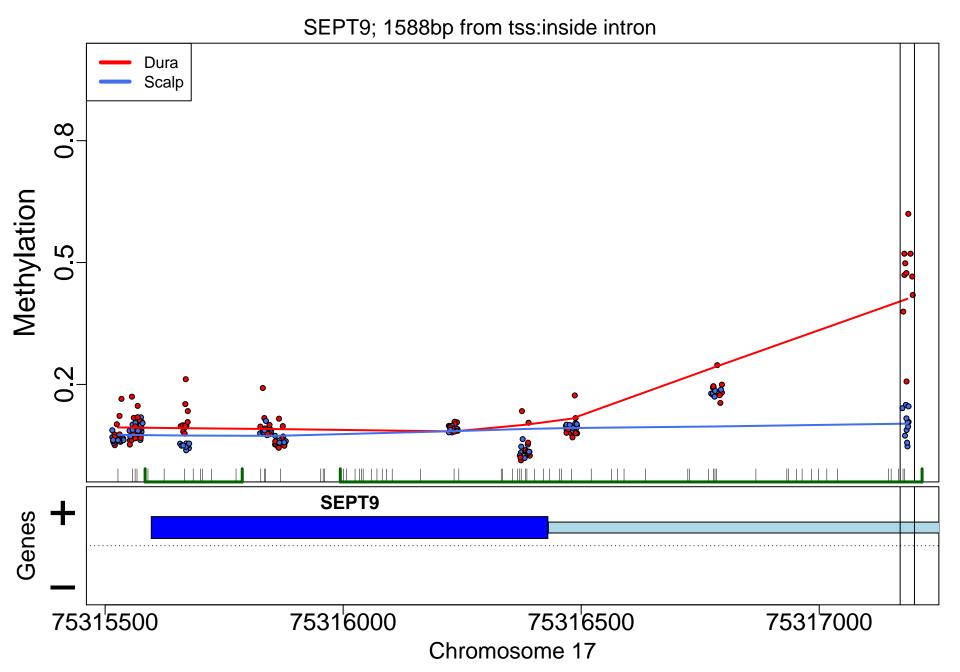


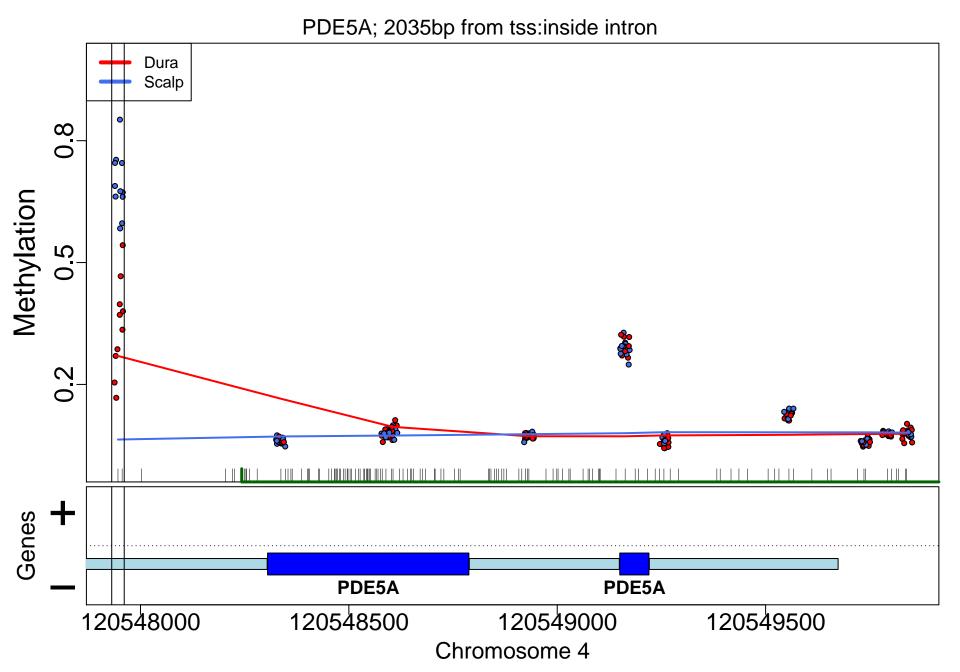


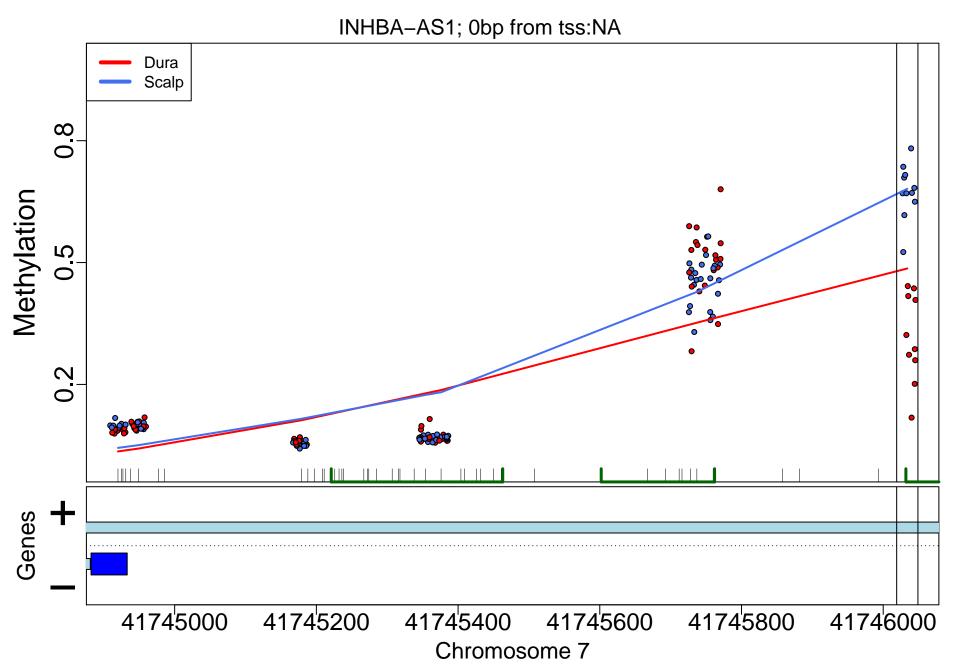




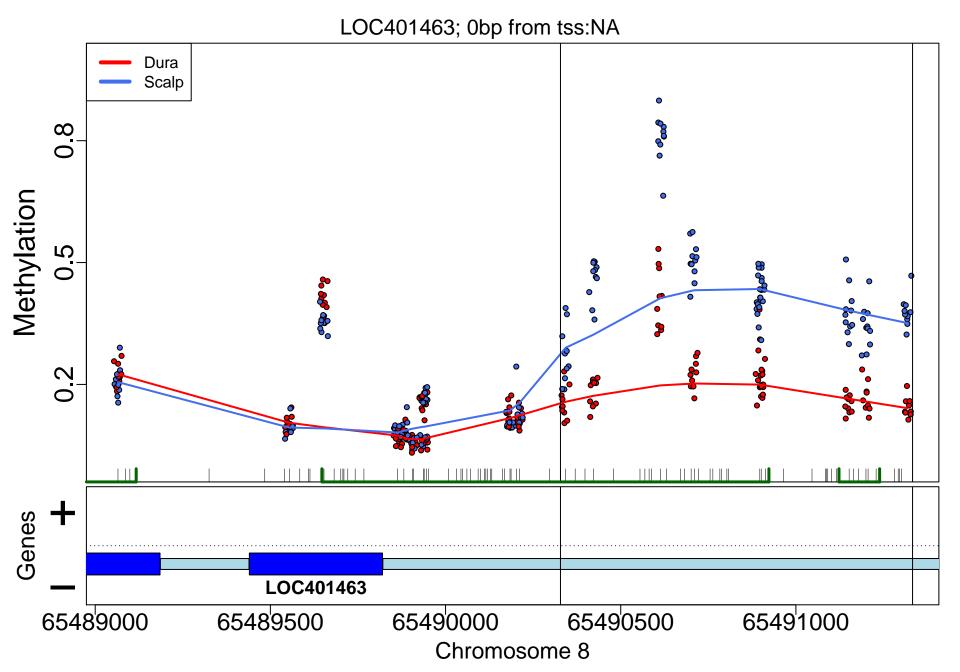




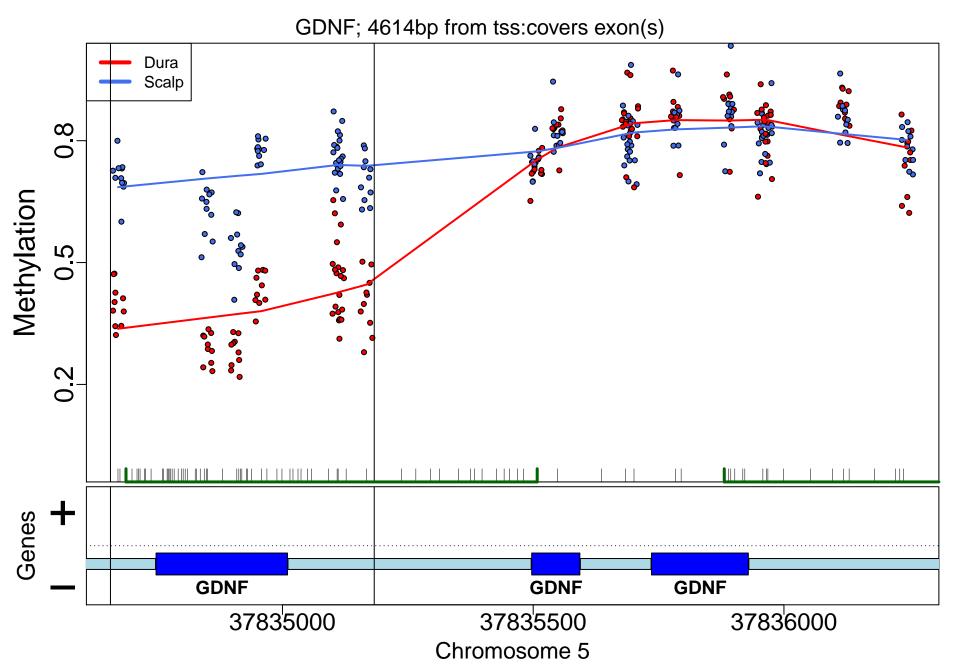


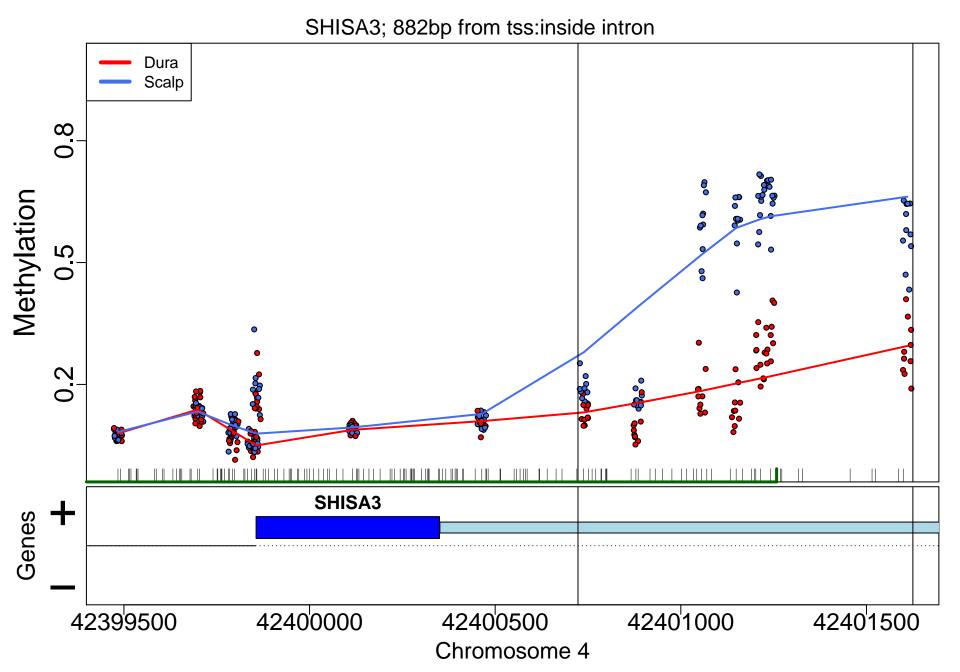


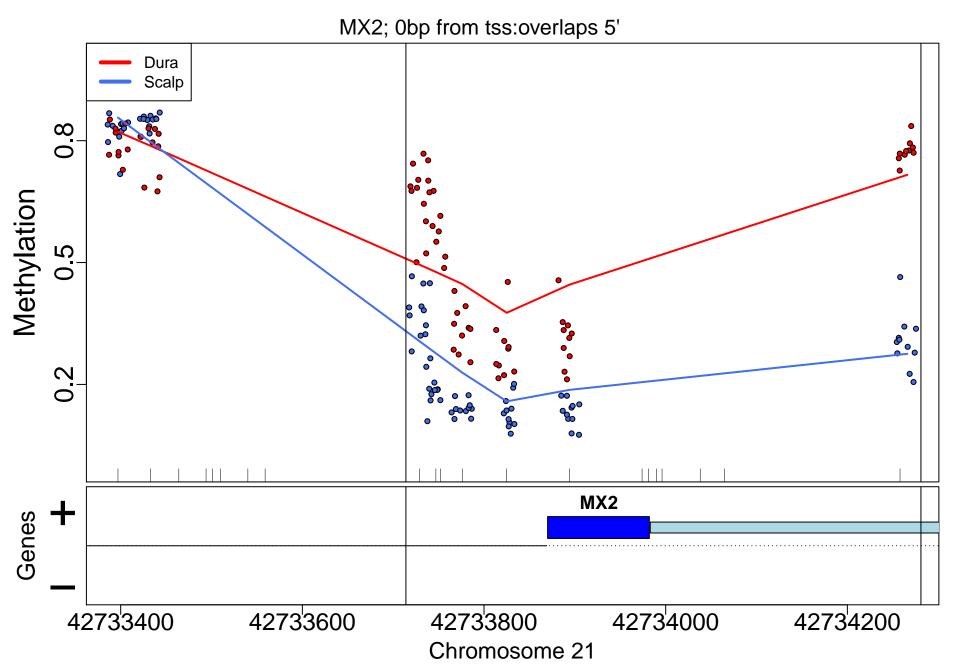
LMX1A; 1273bp from tss:covers exon(s) Dura Scalp 80. Methylation Genes LMX1A LMX1A LMX1A LMX1A 165322000 165323000 165324000 165325000 Chromosome 1

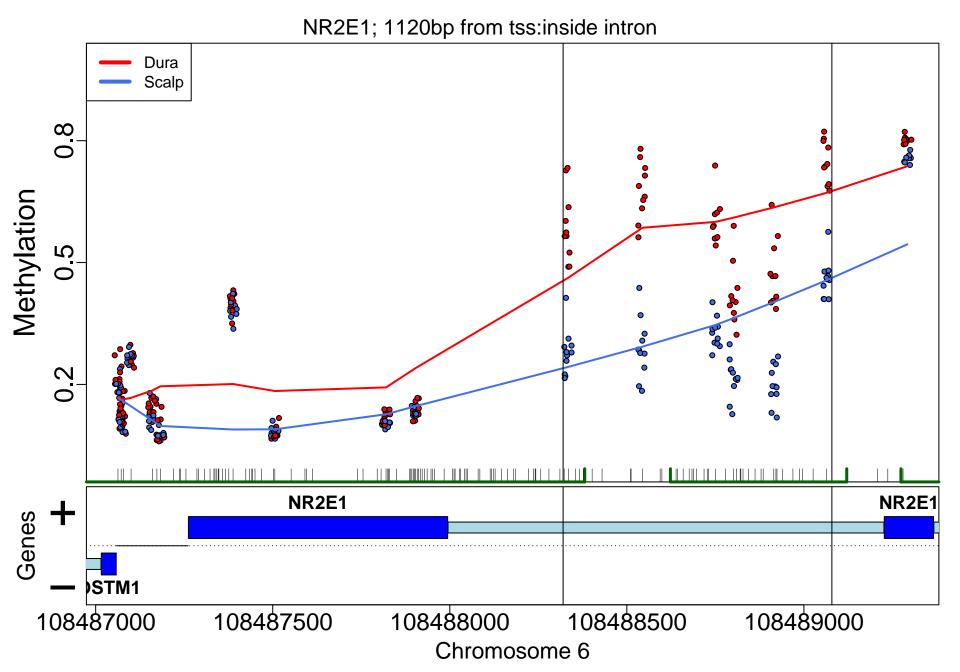


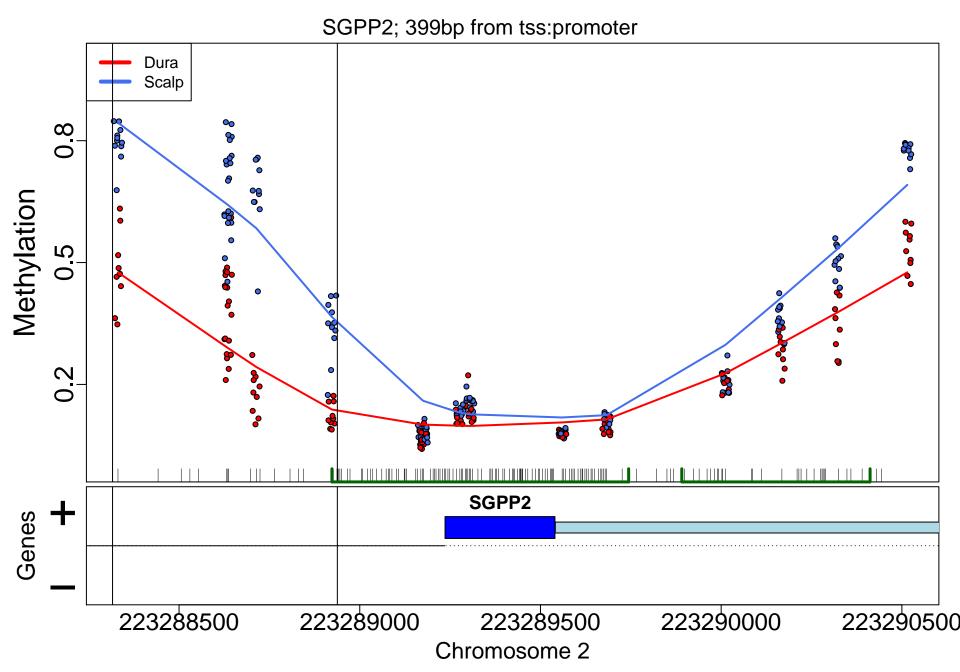
TWIST2; 42767bp from tss:inside intron Dura Scalp 80. 88 Methylation 0.5 0.2 Genes 239800000 239800200 239799600 239799800 Chromosome 2

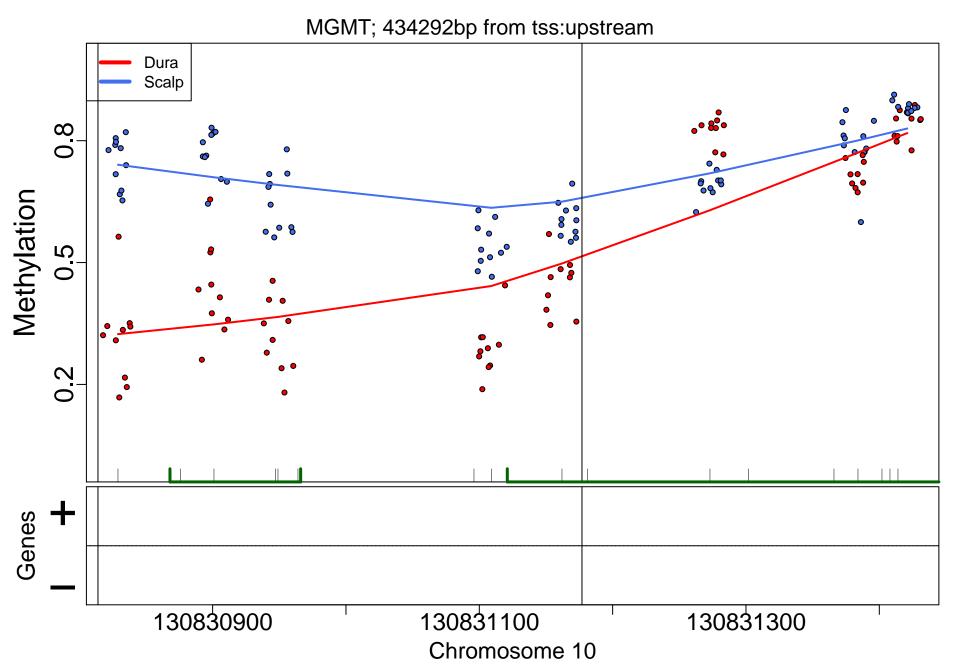


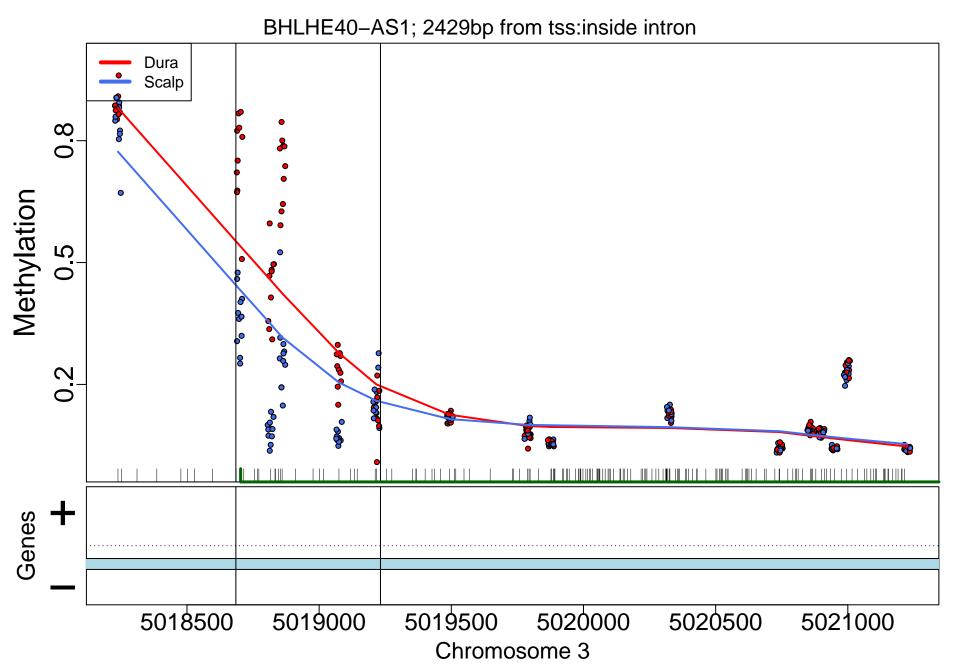


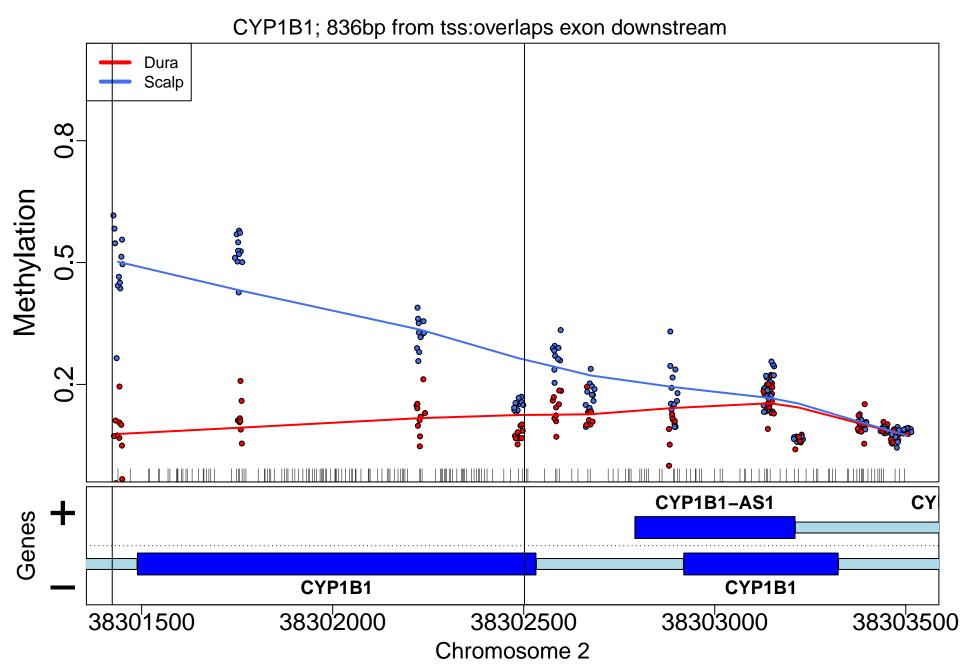


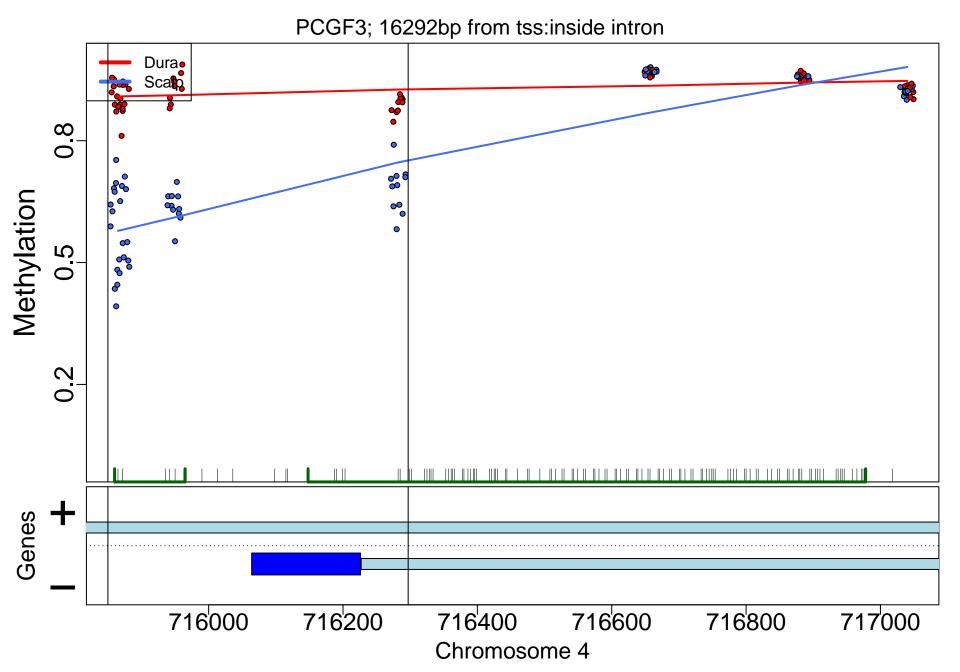


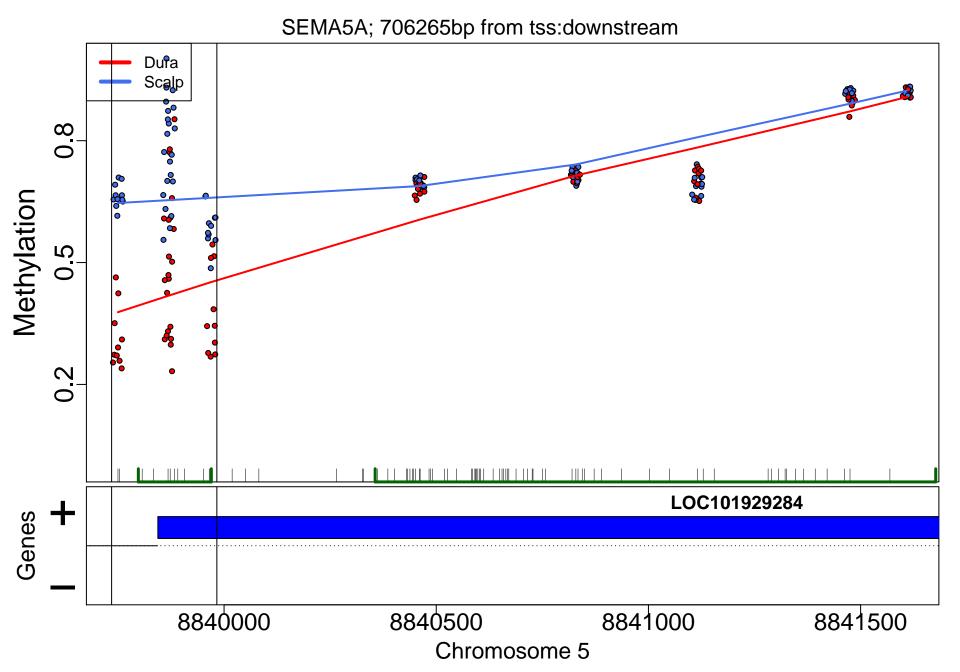


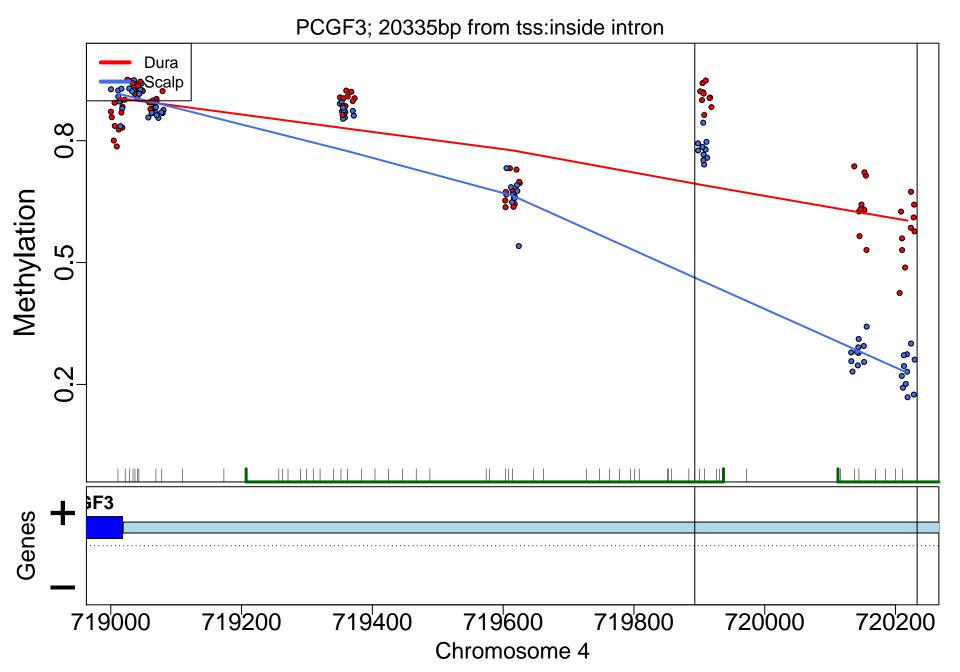


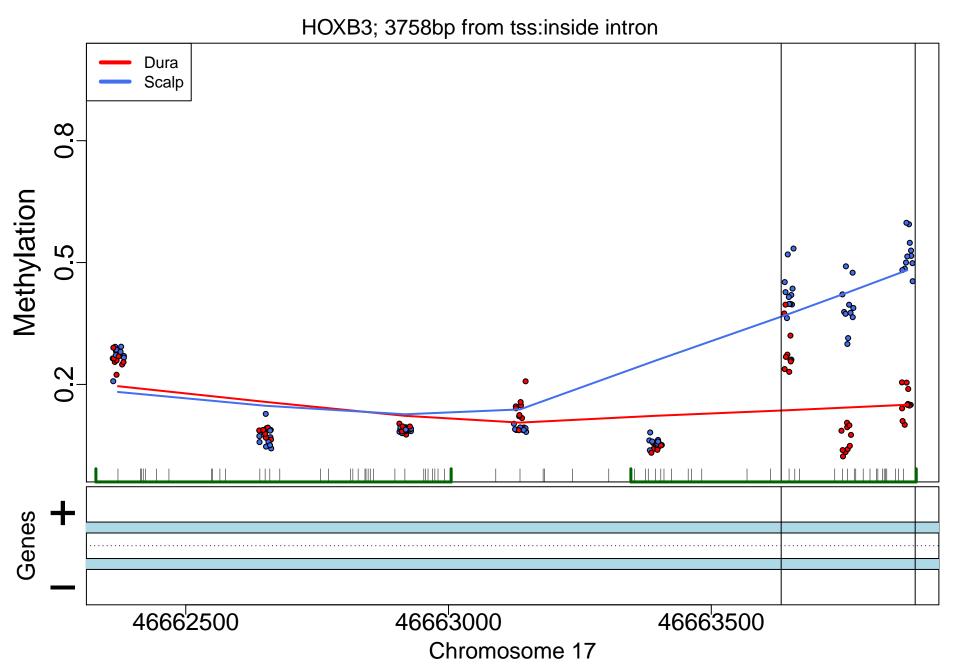


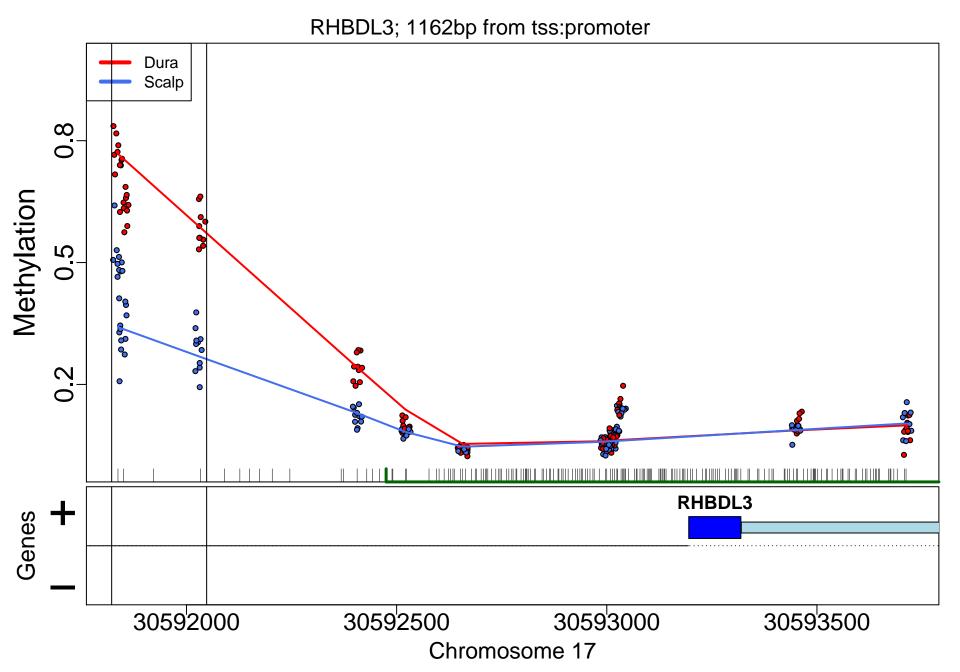


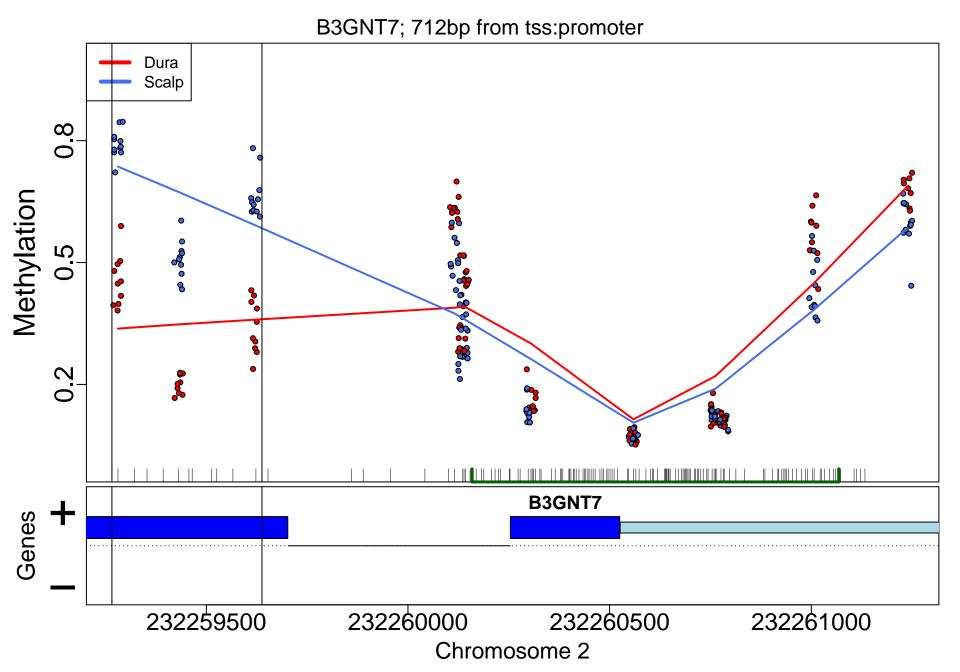


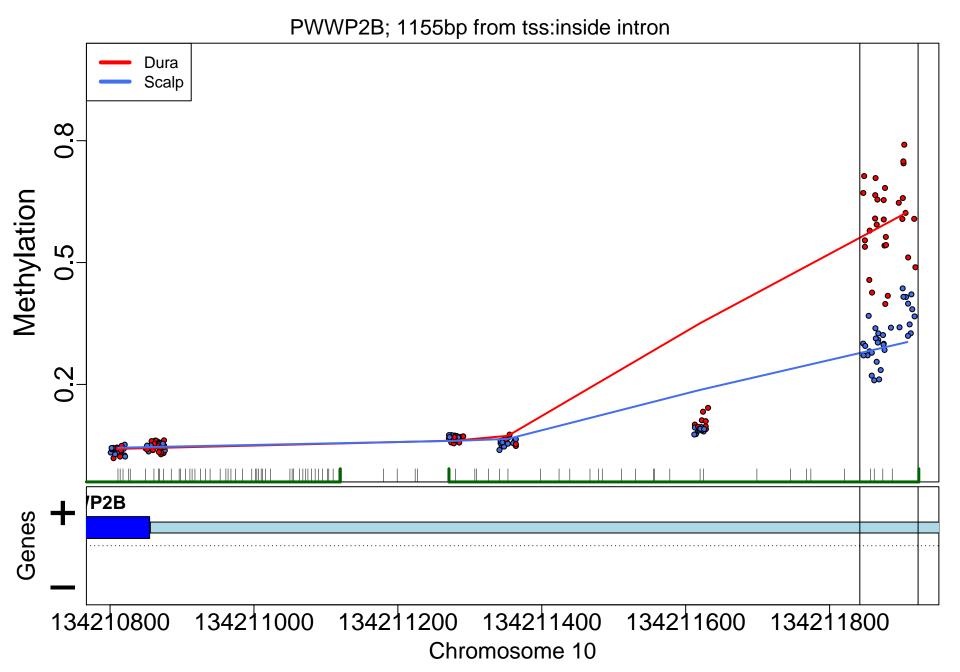


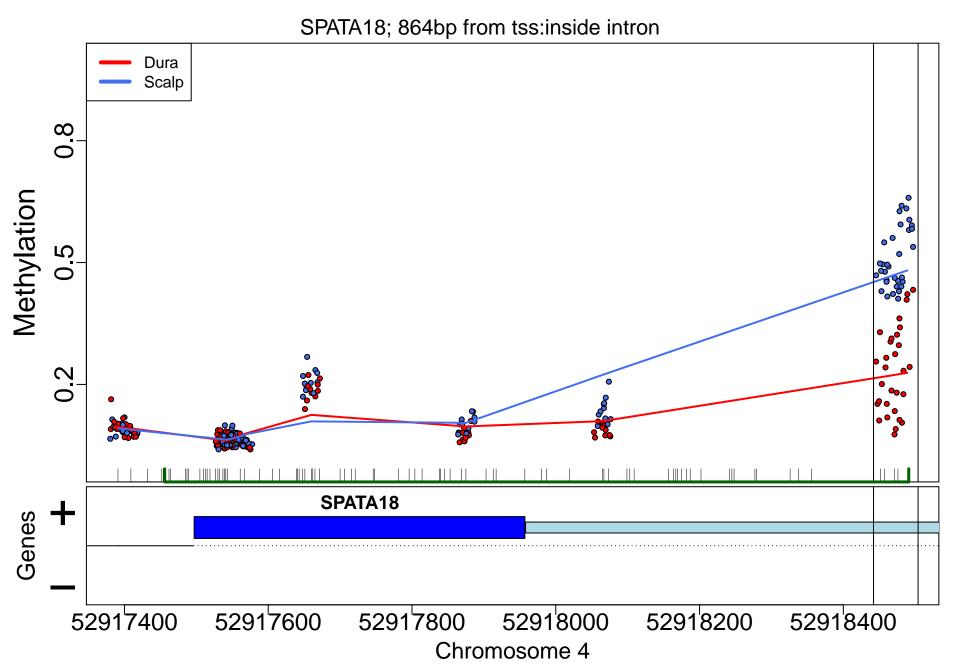


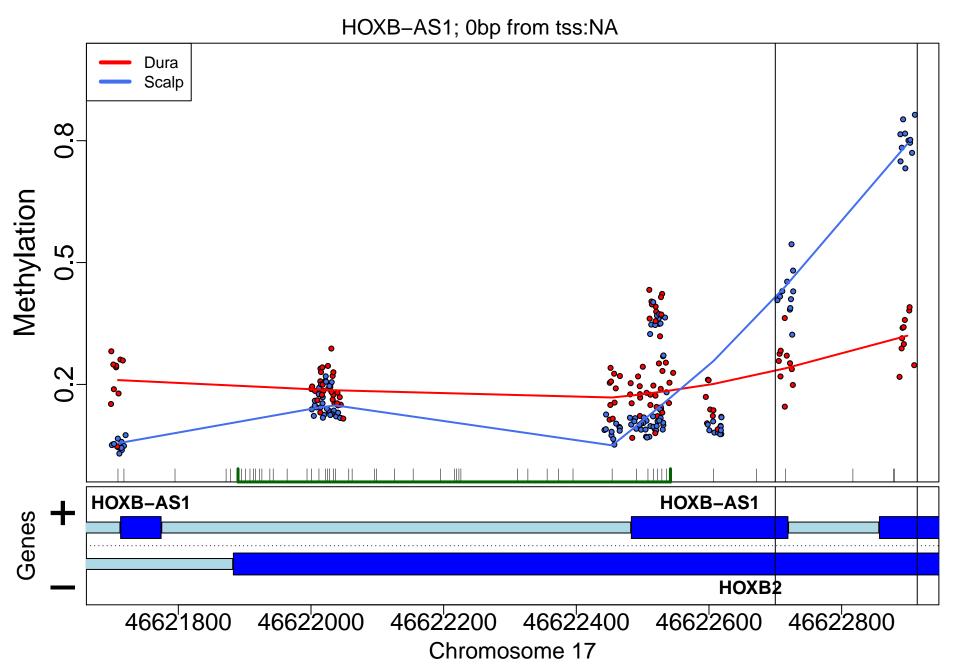


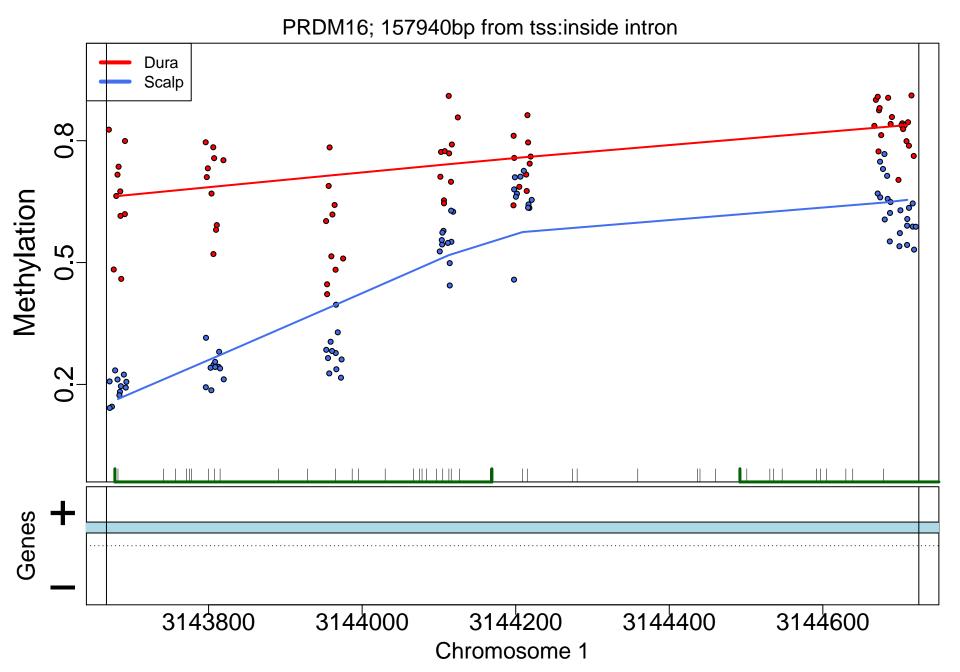


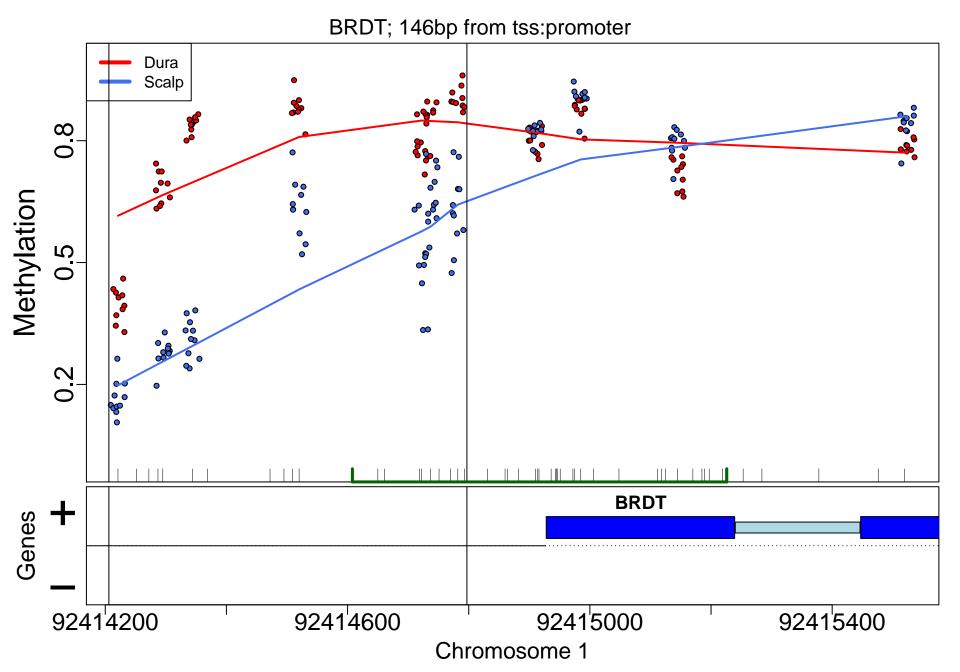


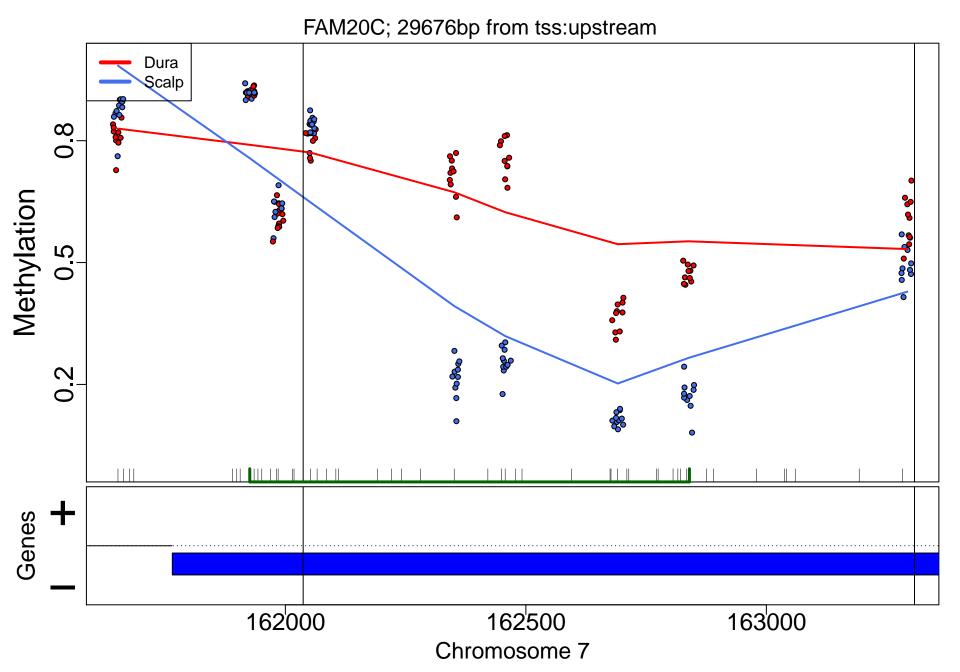


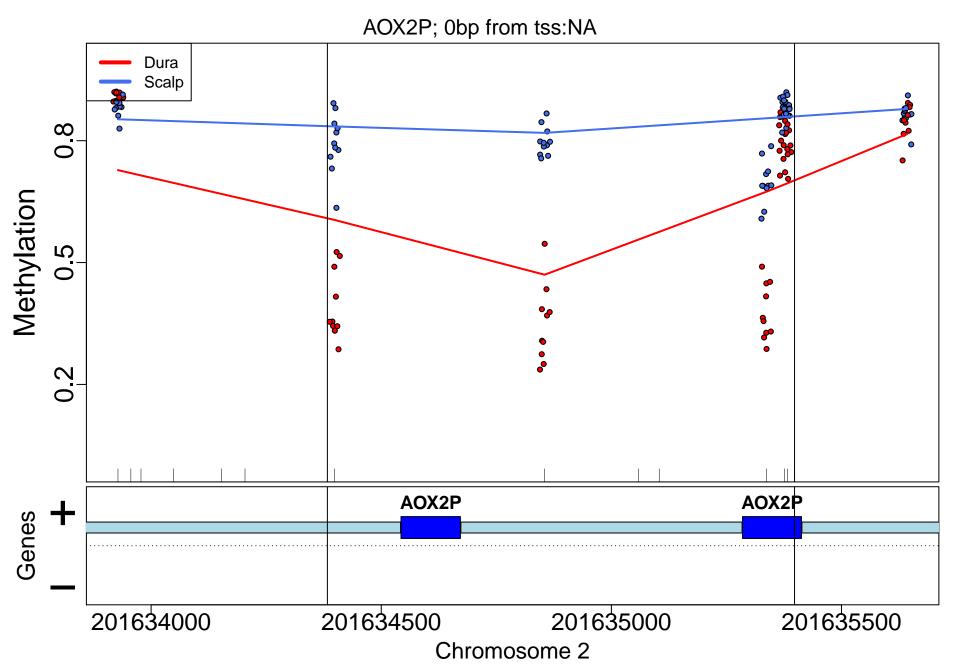




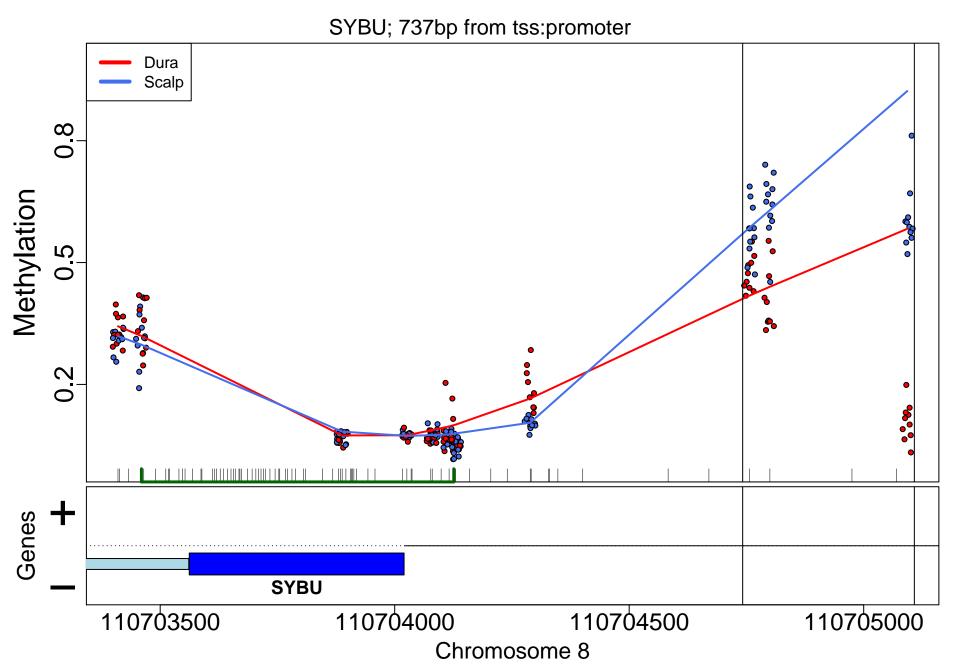


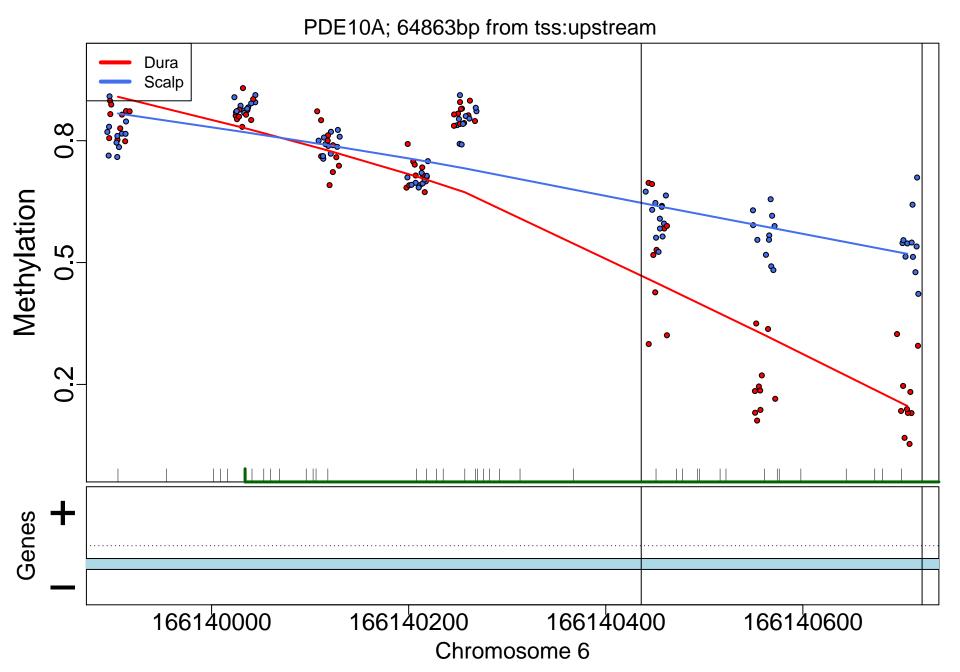


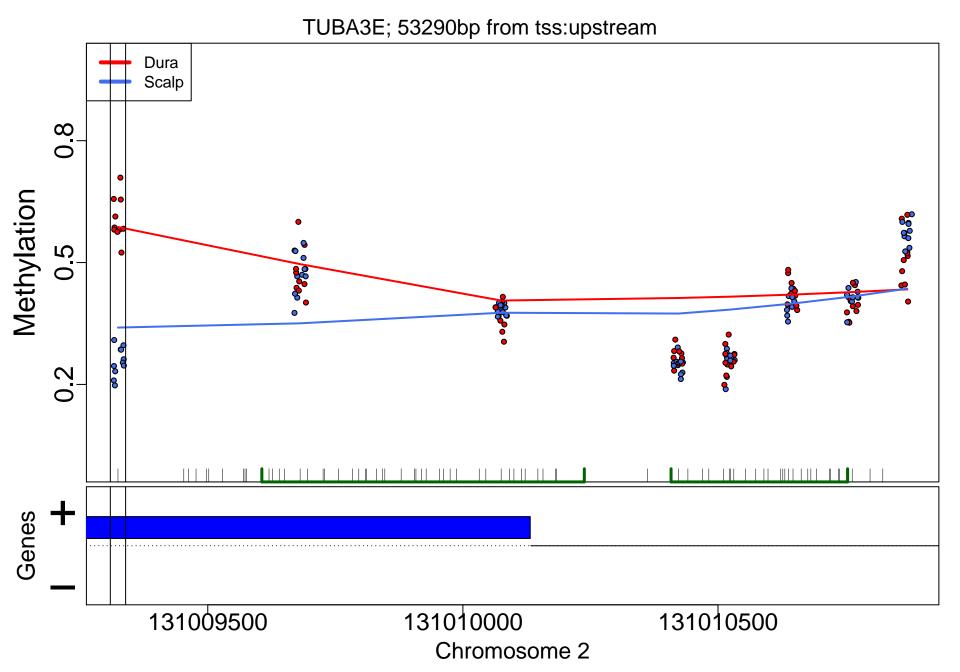




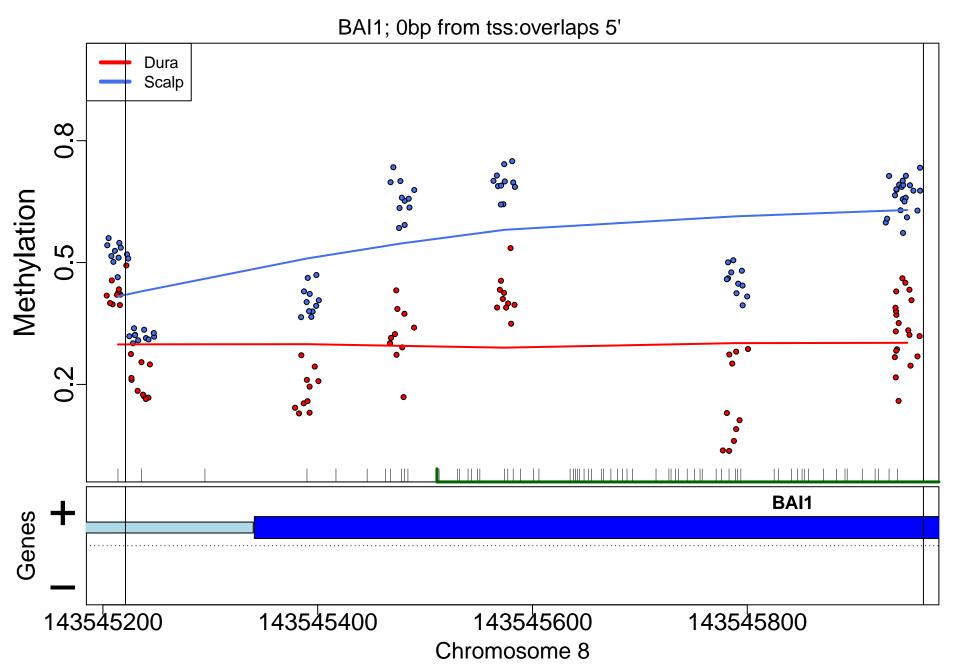
TRPM2; 50bp from tss:inside exon Dura Scalp <u>∞</u>. Methylation 0.2 TRPM2 TRPM2 Genes 45774000 45773500 45774500 Chromosome 21

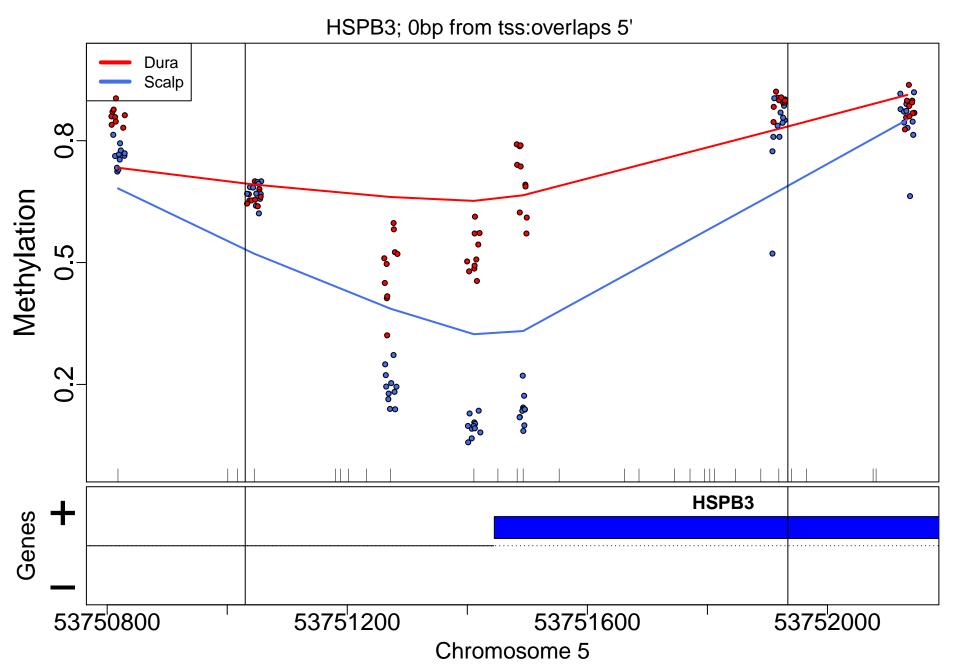


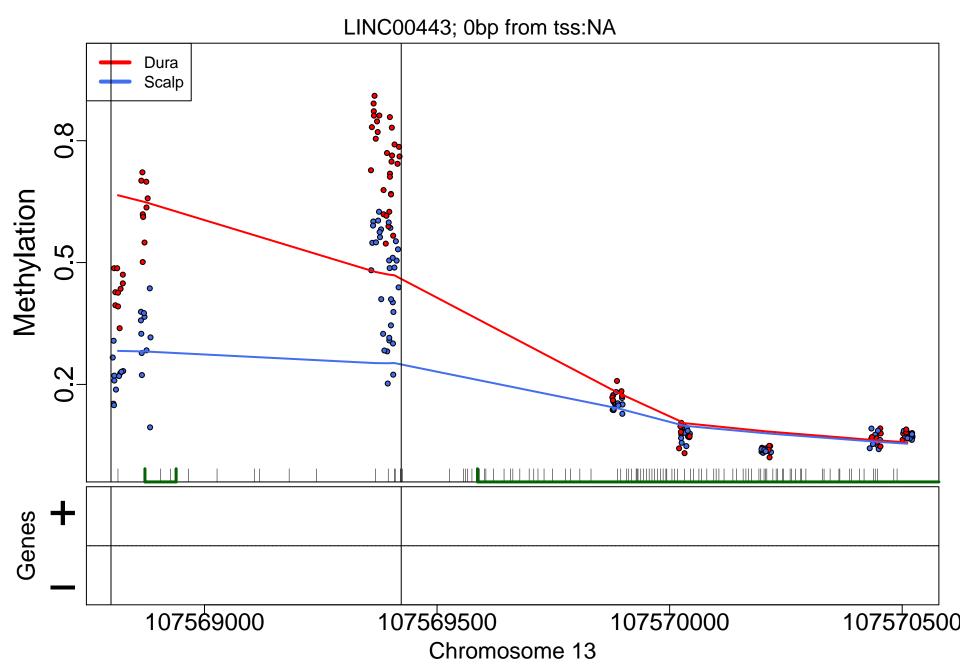


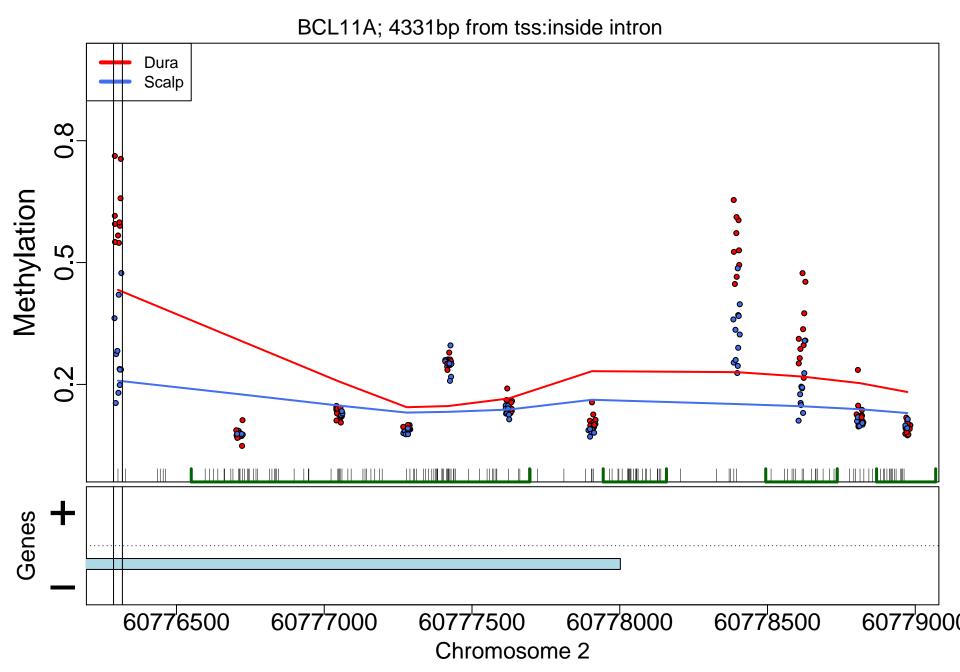


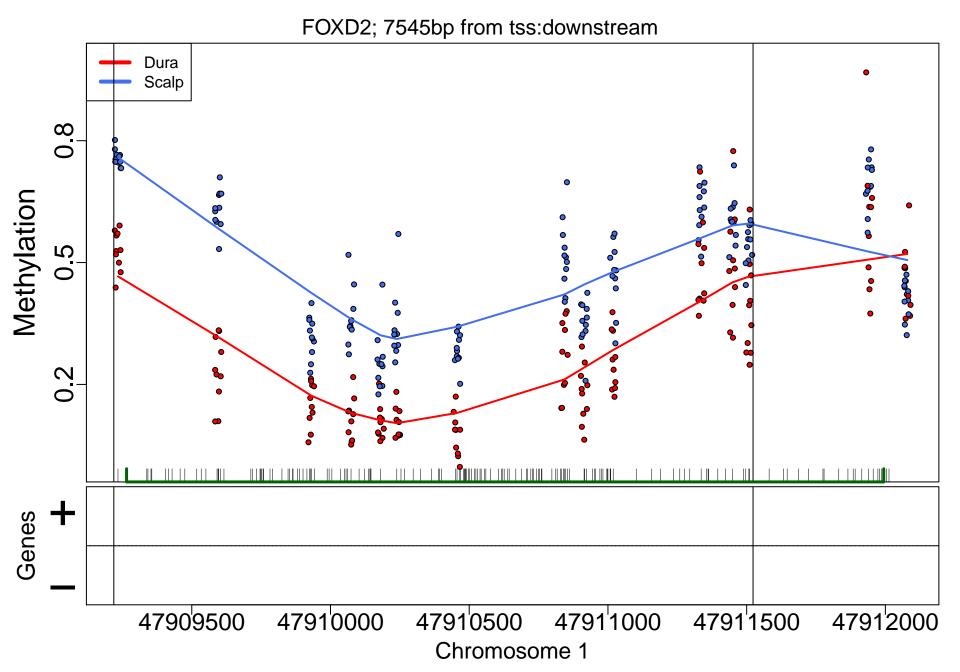
TBX15; 9654bp from tss:upstream Dura Scalp 0 Methylation 0.2 Genes 119542000 119543000 119544000 Chromosome 1

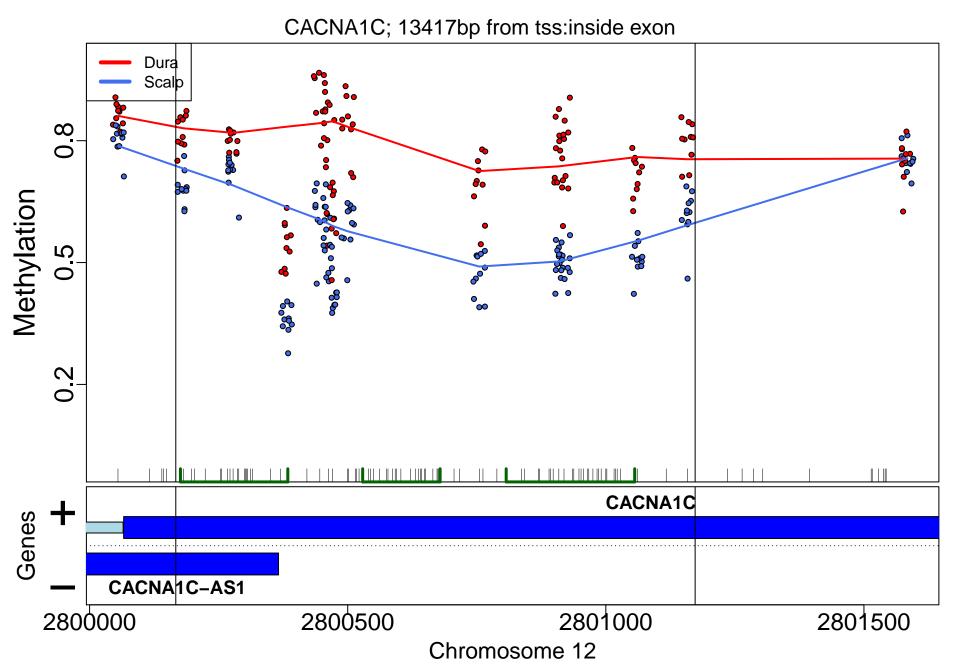




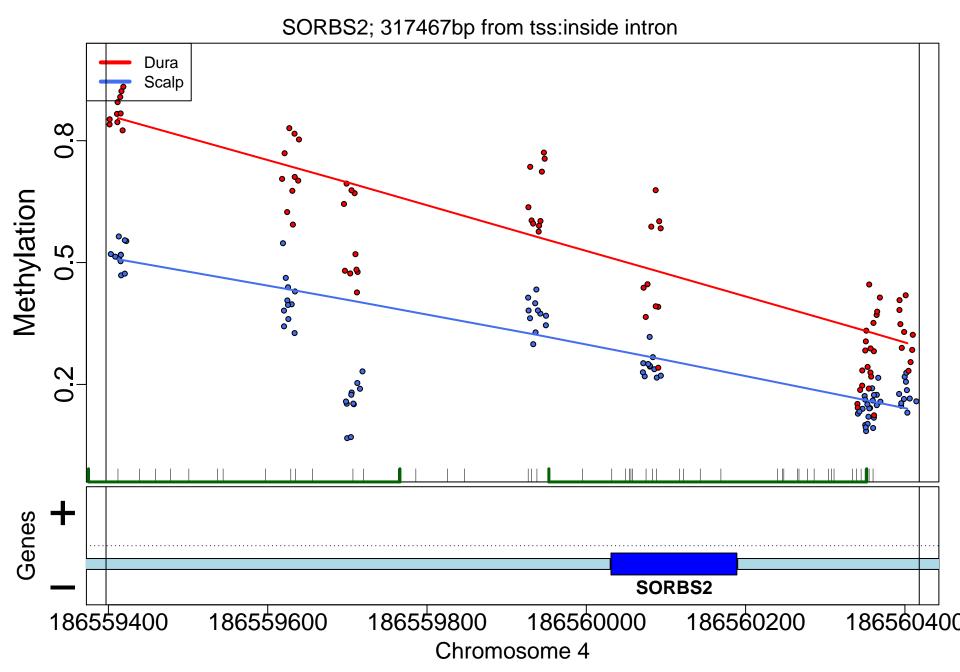




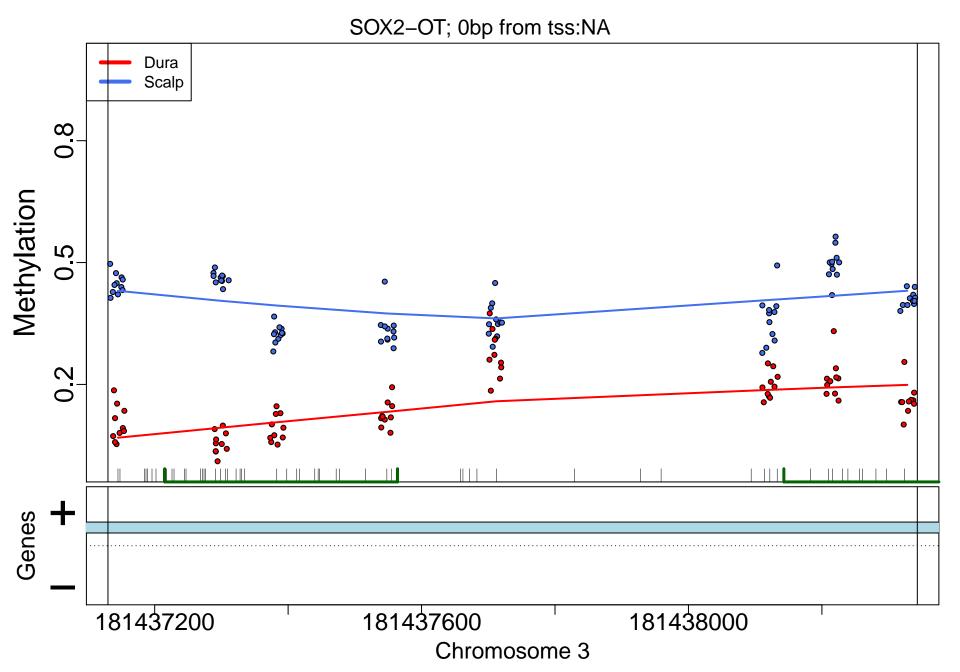


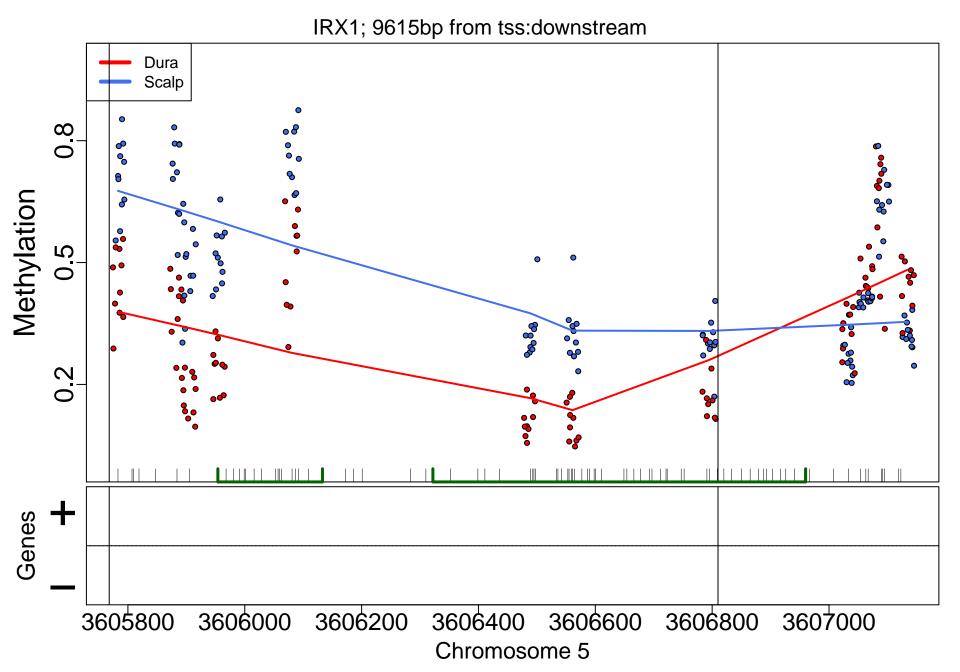


DLX6-AS1; 0bp from tss:NA Dura Scalp 80. Methylation 0.5 0.2 Genes DLX6-AS1 96632000 9663 1000 9663 1500 96633000 96632500 9663 3500 Chromosome 7

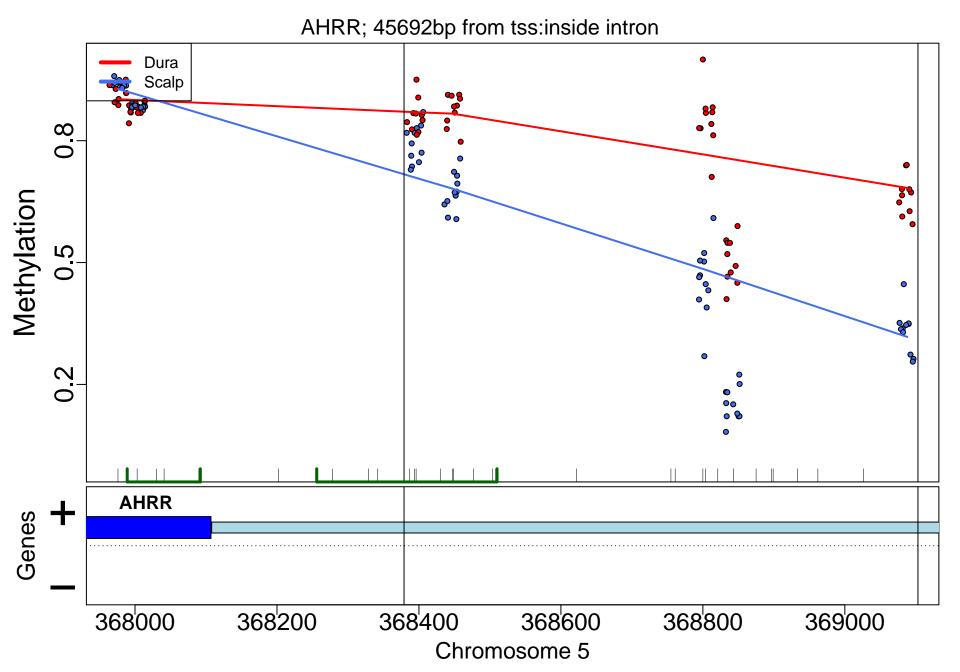


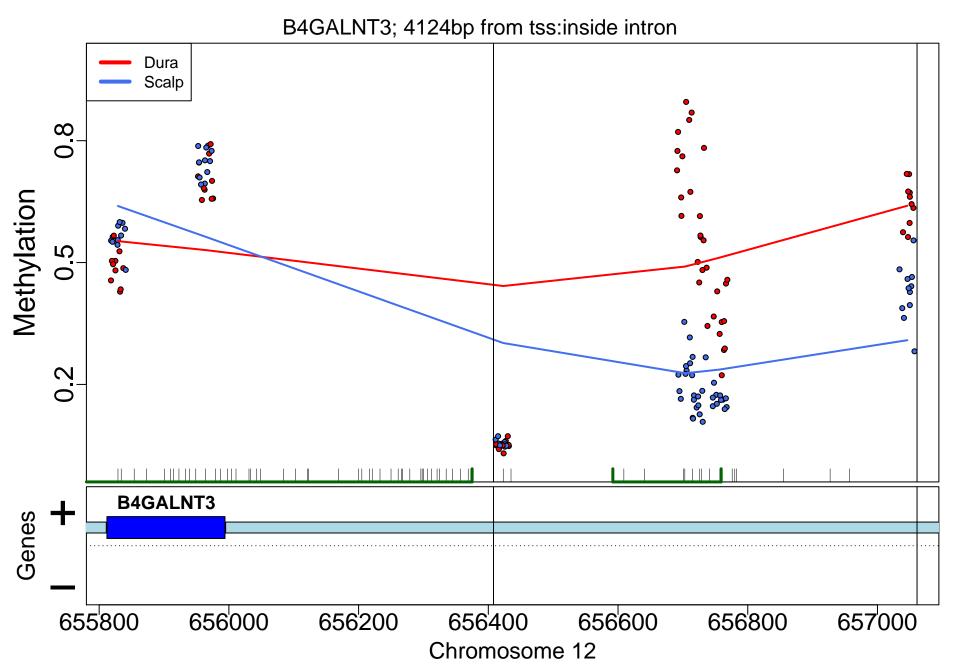
COMMD3-BMI1; 17147bp from tss:downstream Dura Scalp <u>ω</u>_ Methylation 0.2 Genes 22623000 22622000 22625000 22624000 Chromosome 10

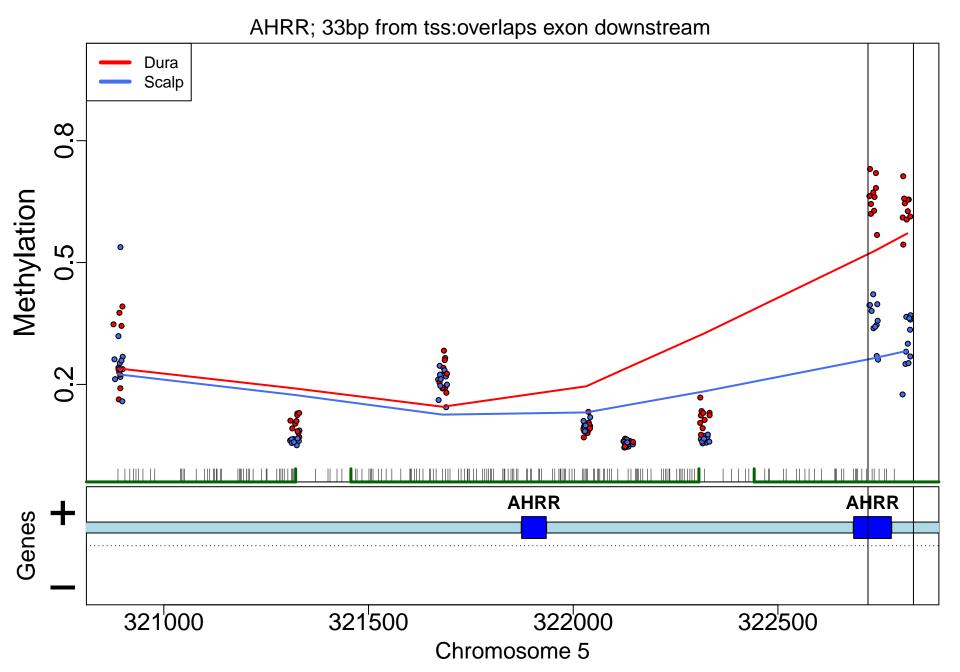


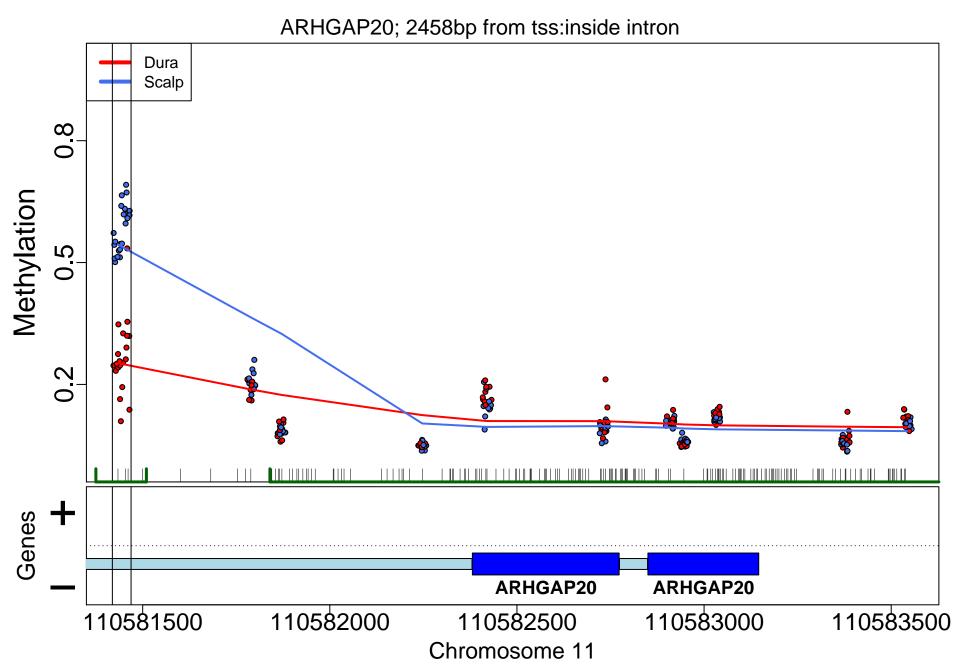


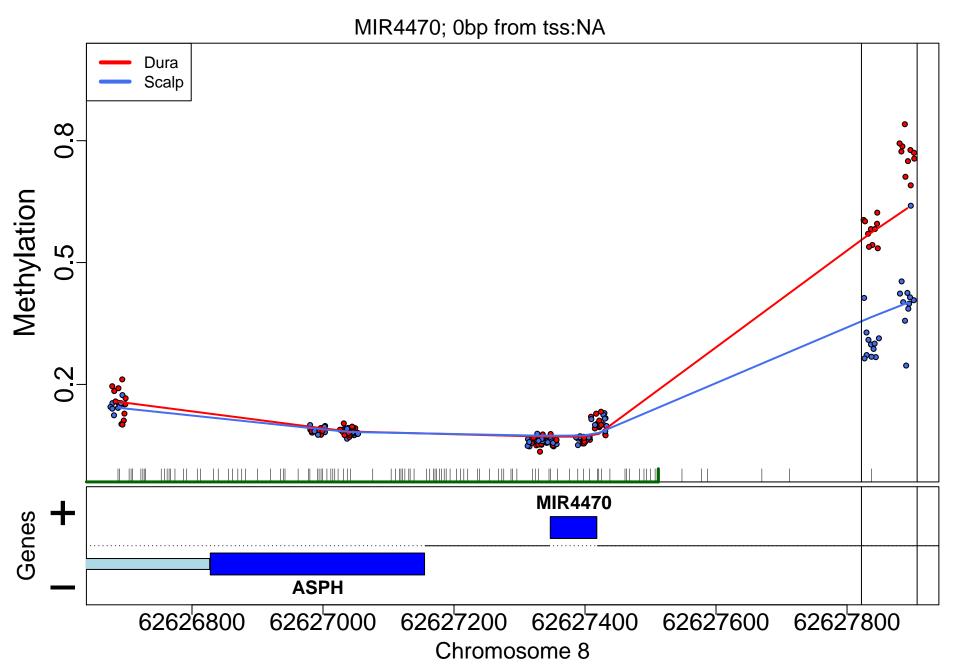
TBX15; 5023bp from tss:inside intron Dura Scalp 80. Methylation Genes 119526000 119527000 119528000 119529000 Chromosome 1

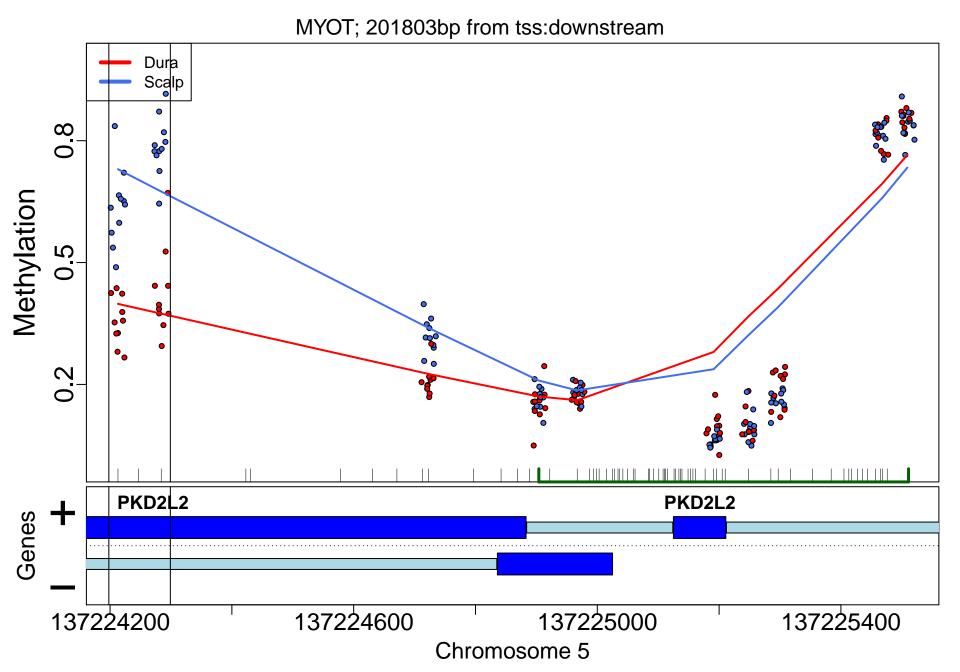


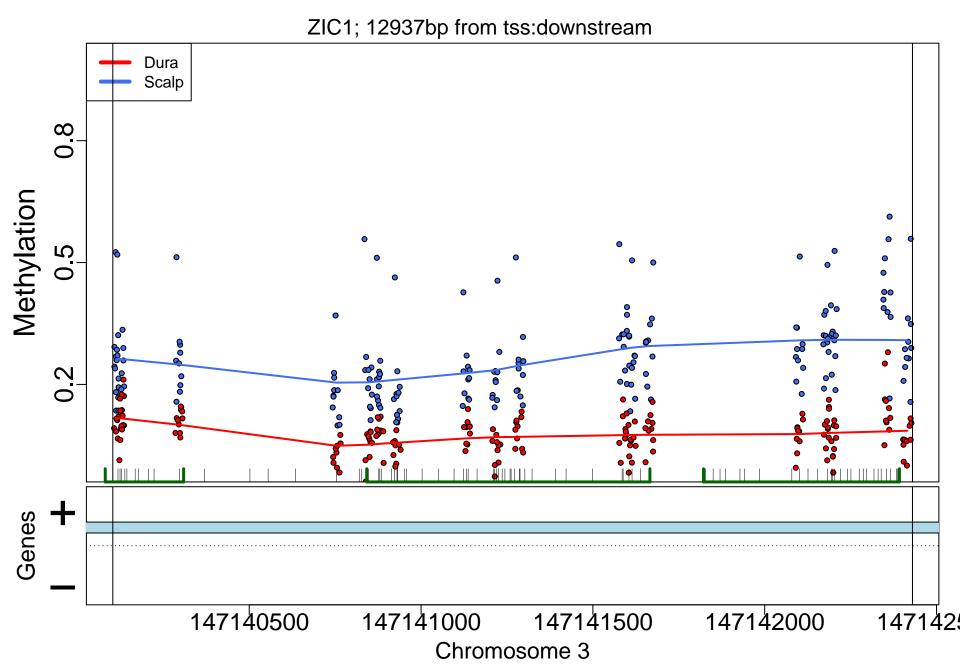


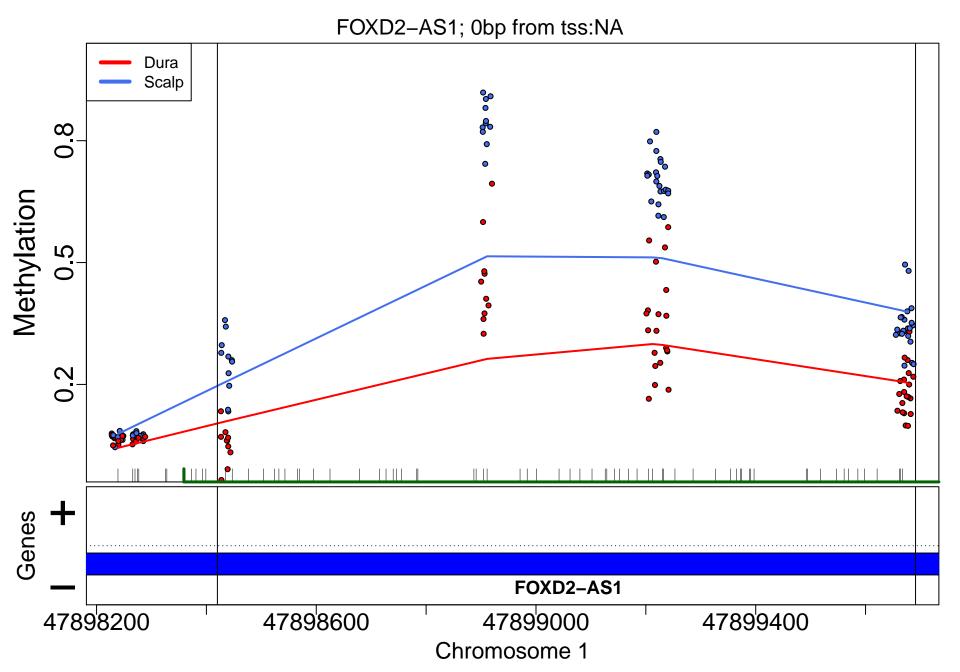


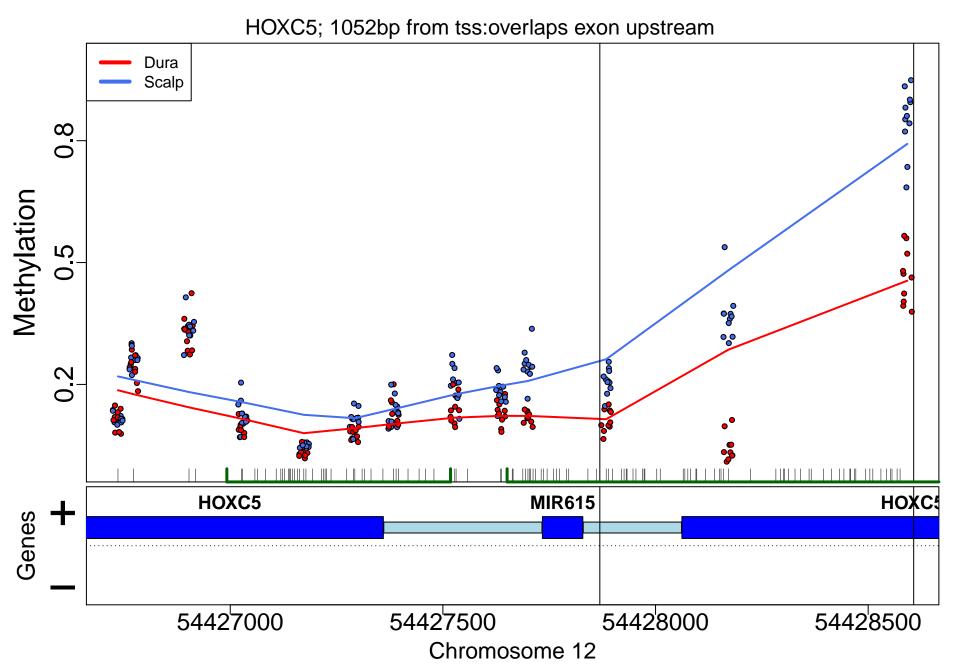


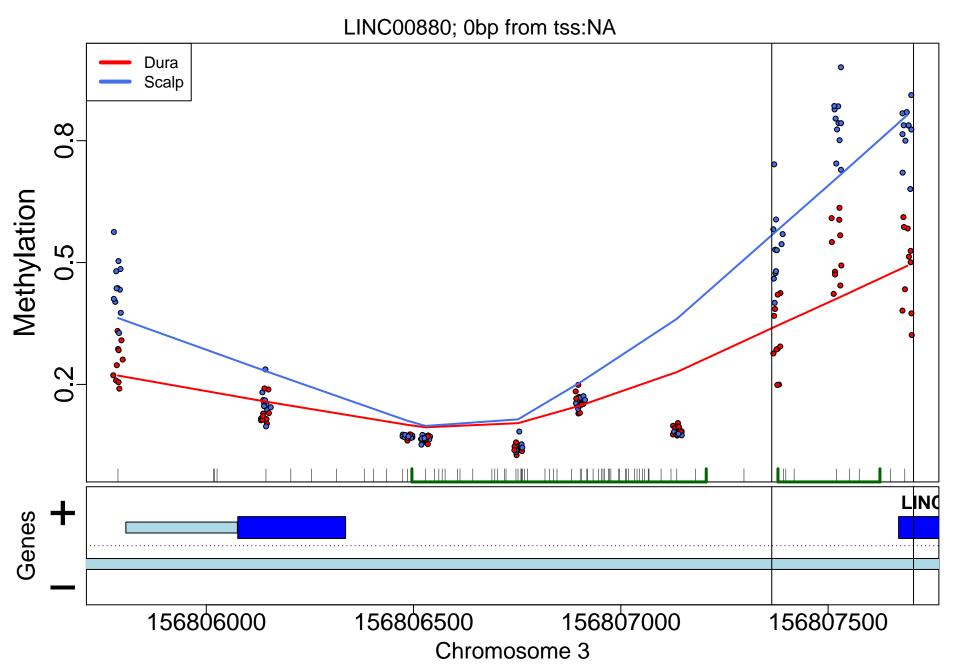


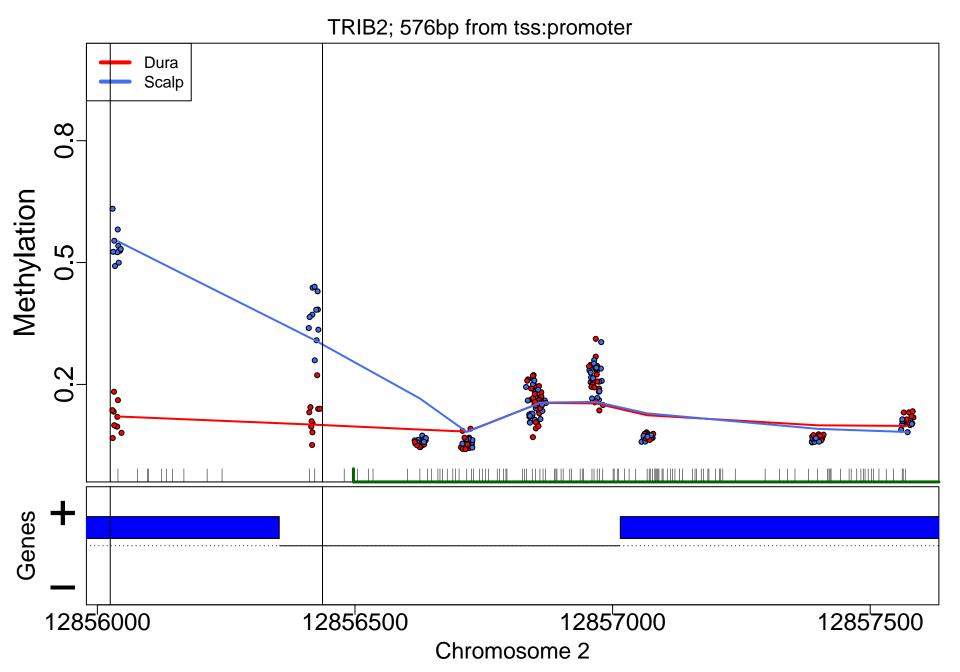


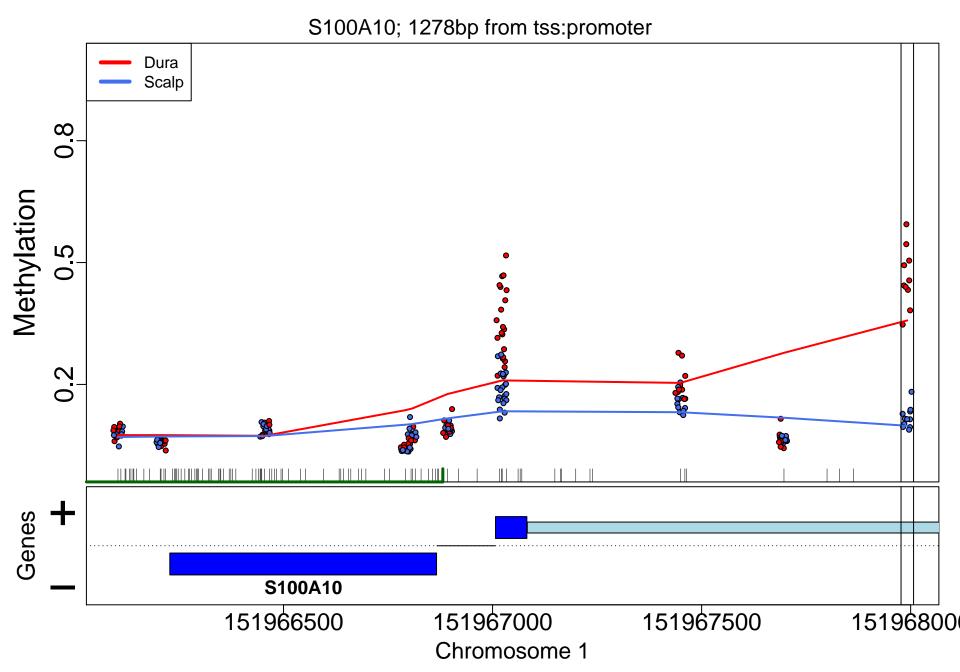


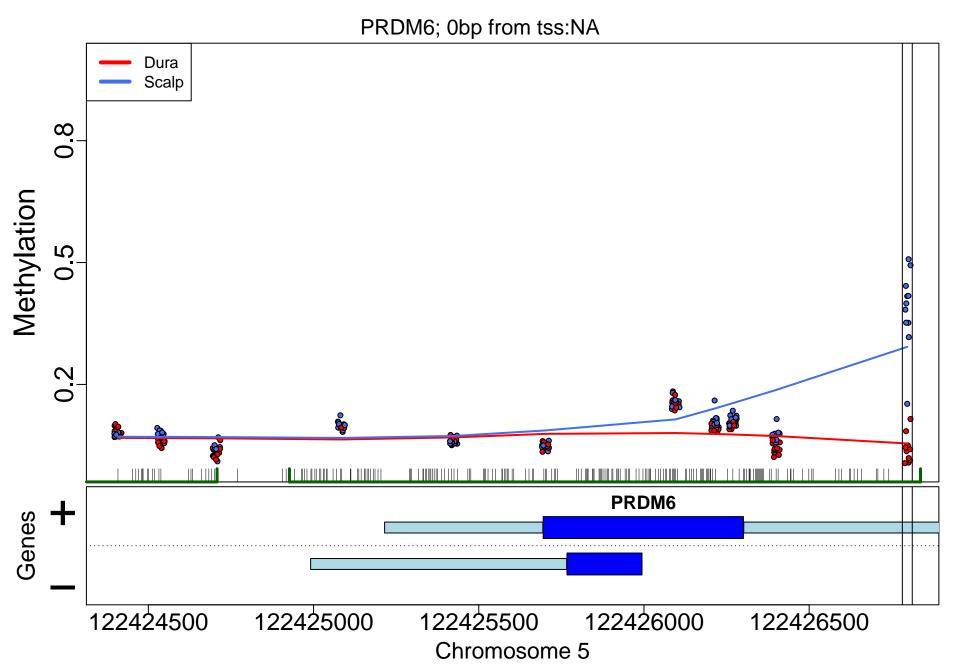


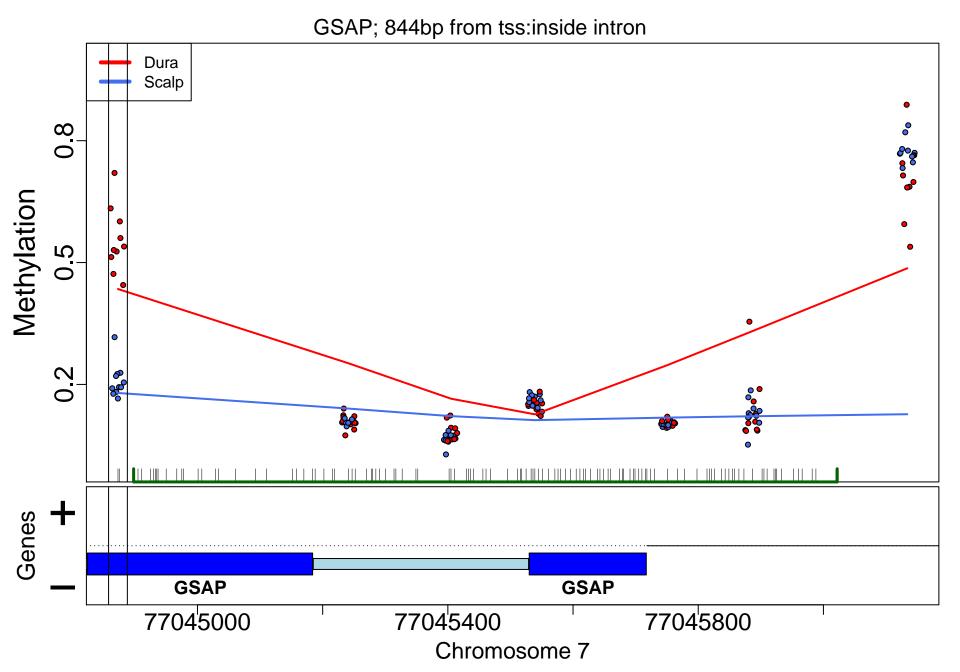




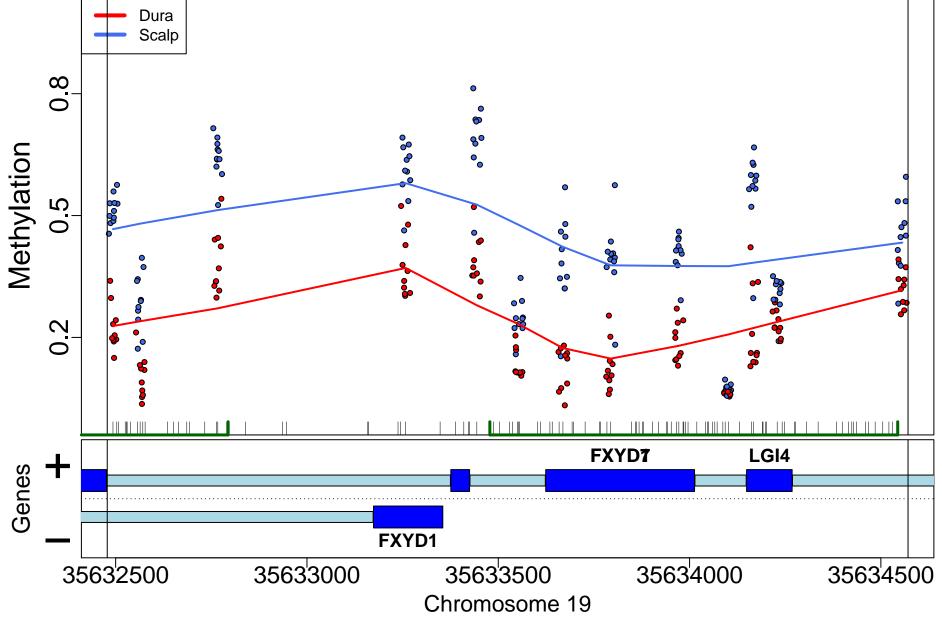


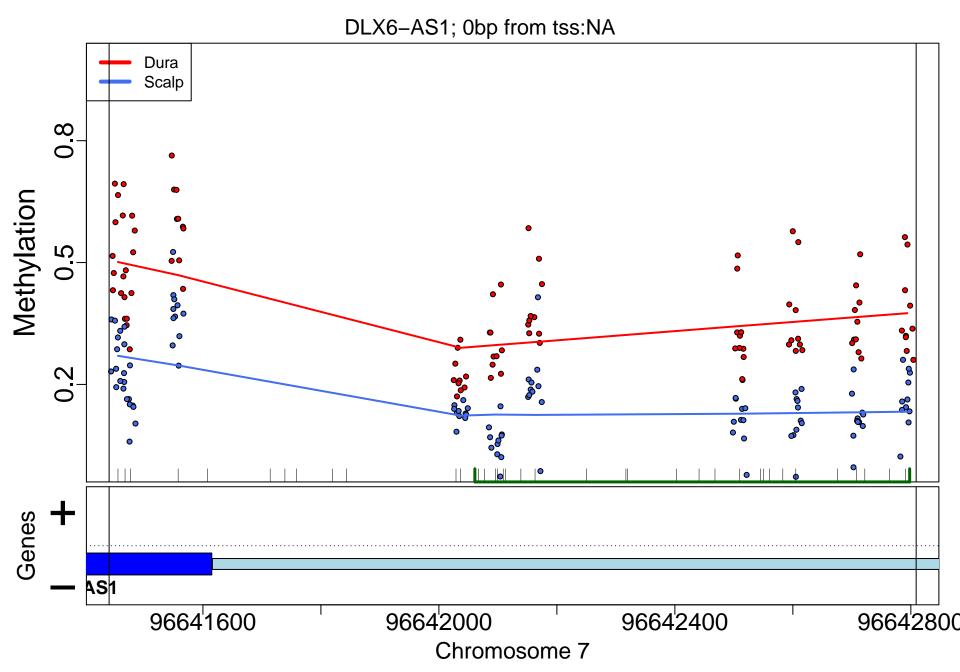


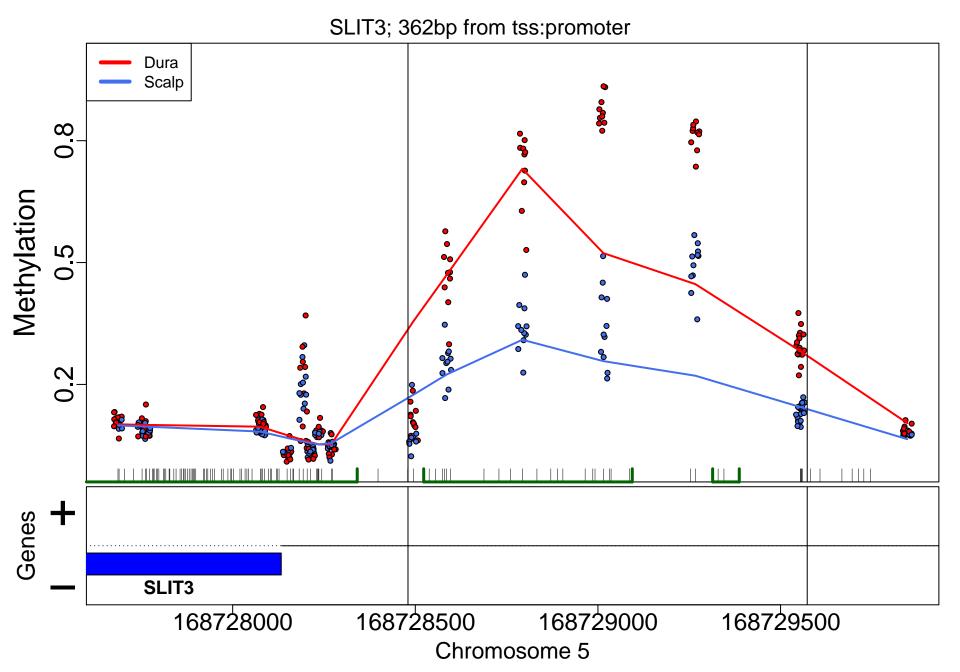


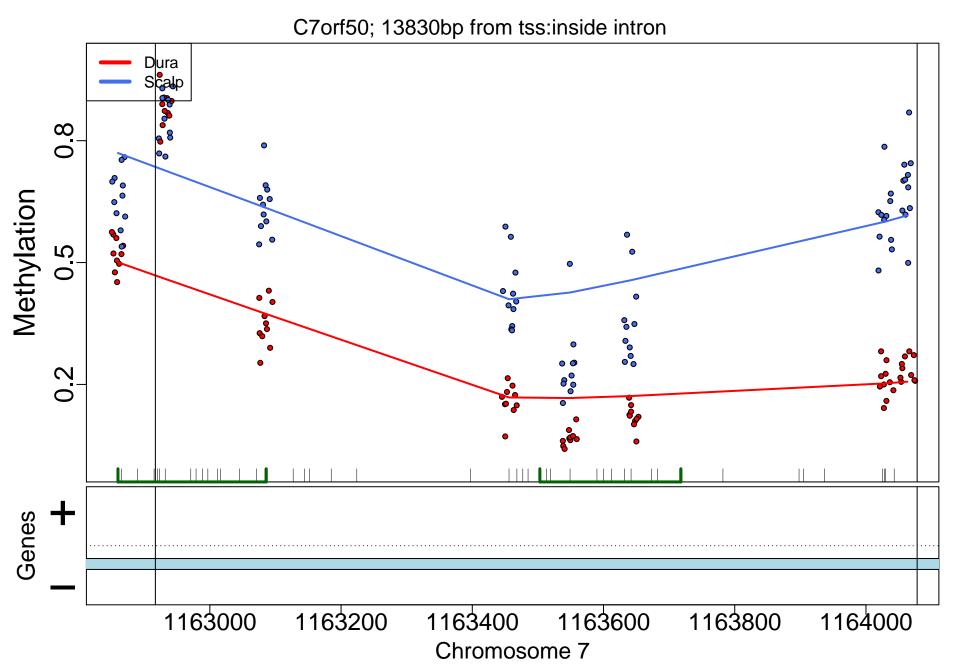


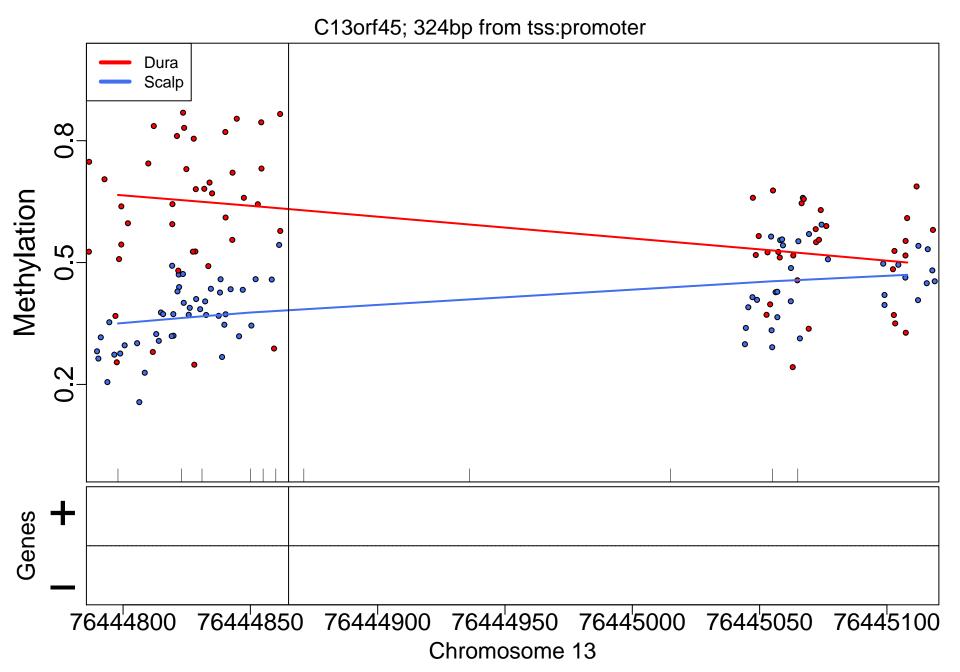
FXYD7; 0bp from tss:overlaps 5' Dura Scalp FXYD7 LGI4

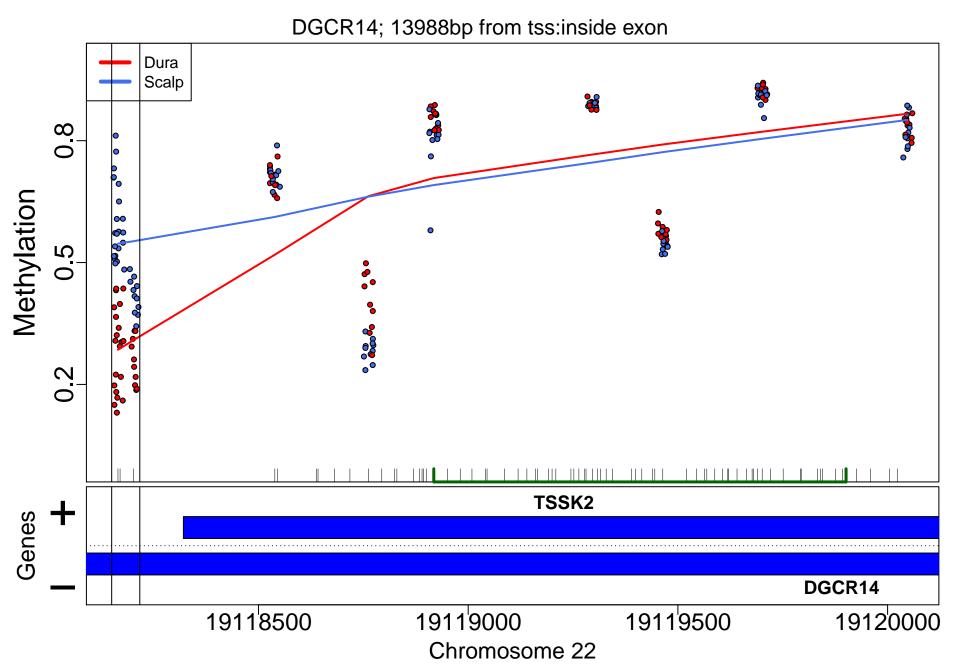


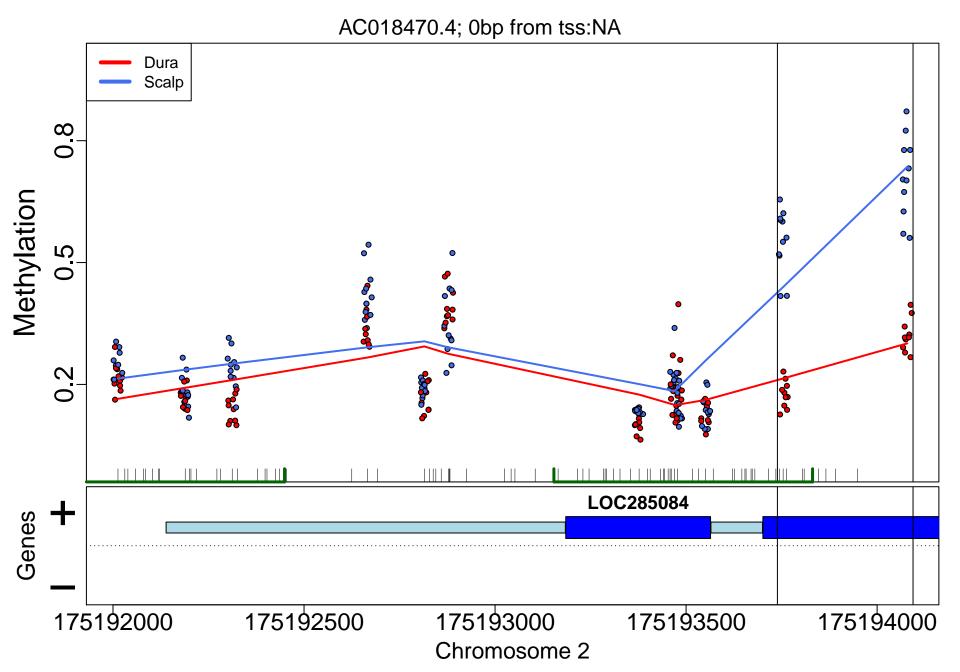


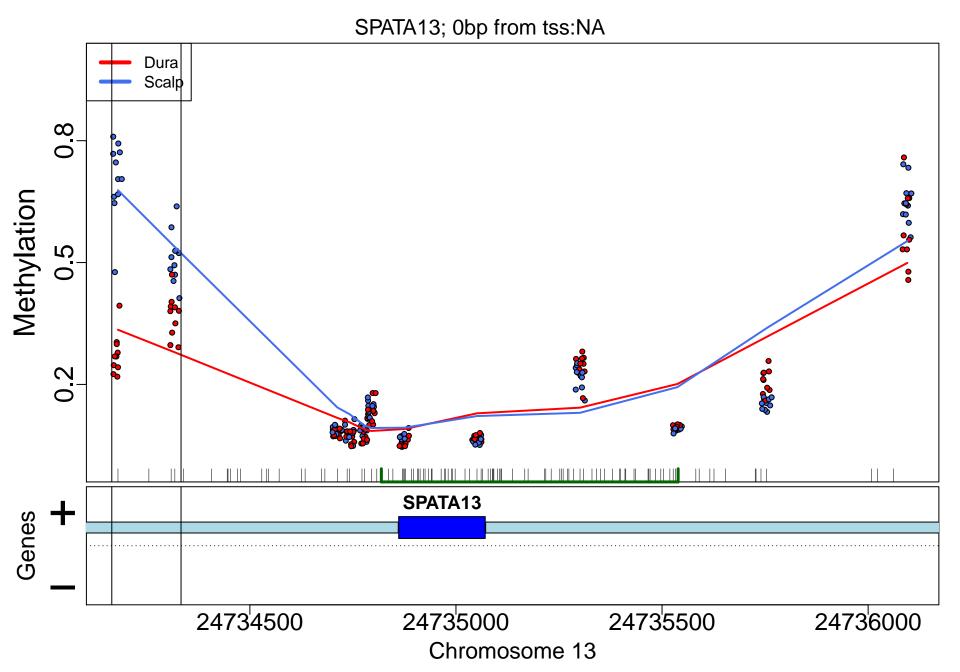


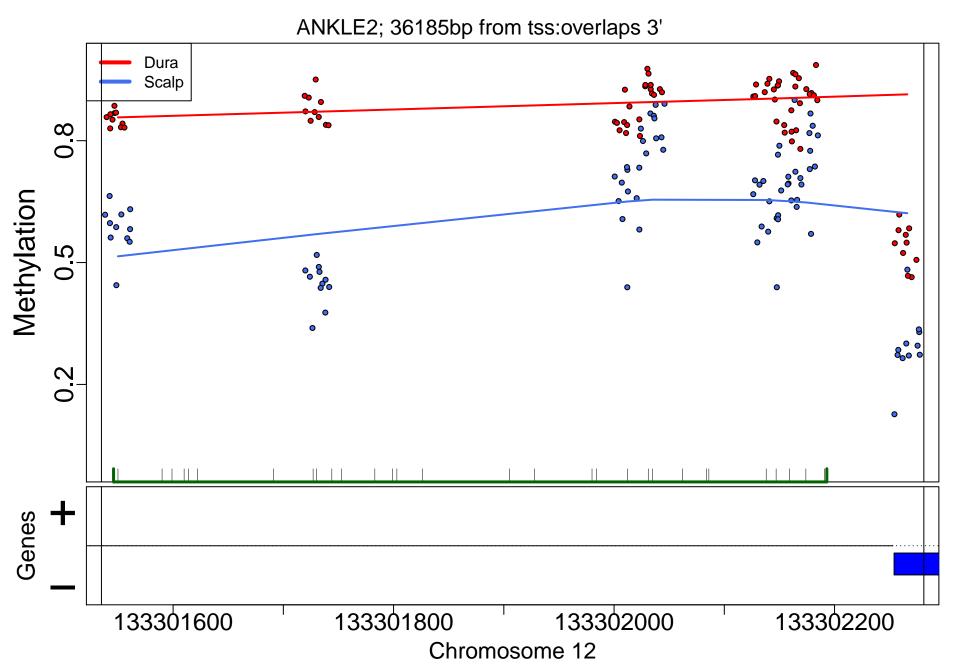


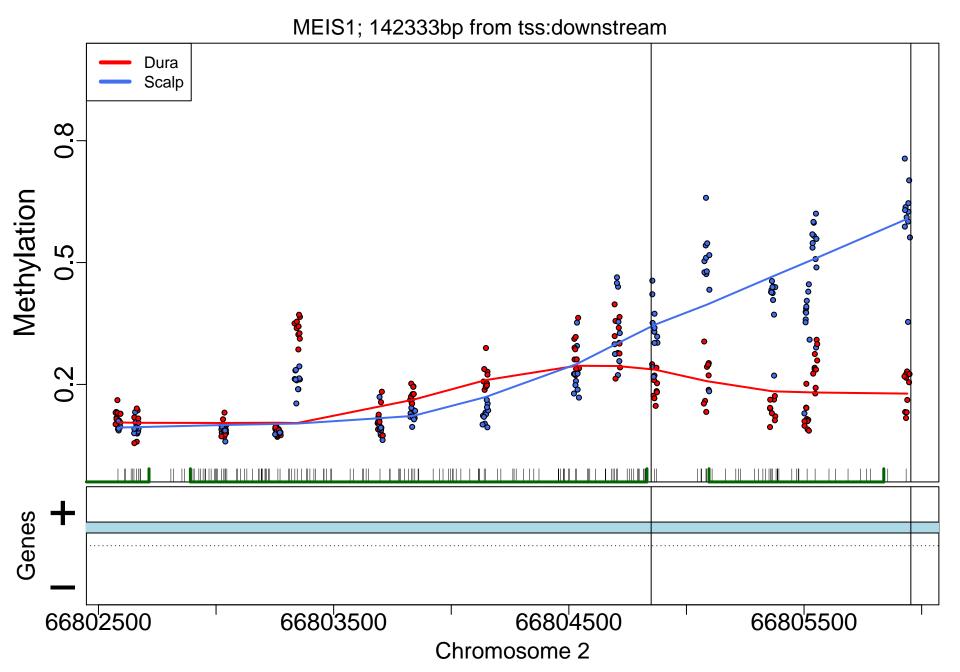


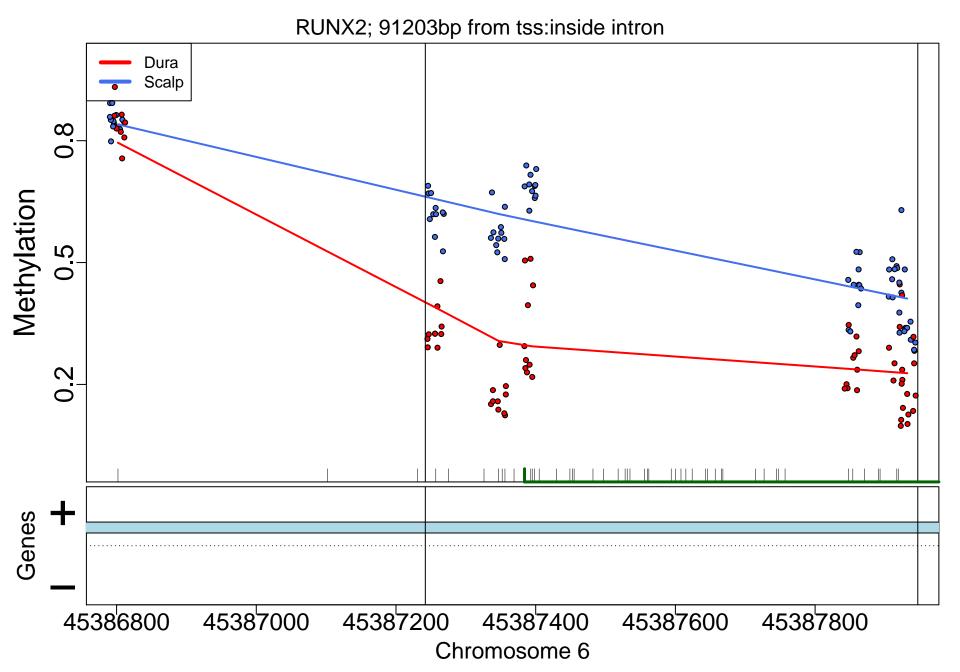




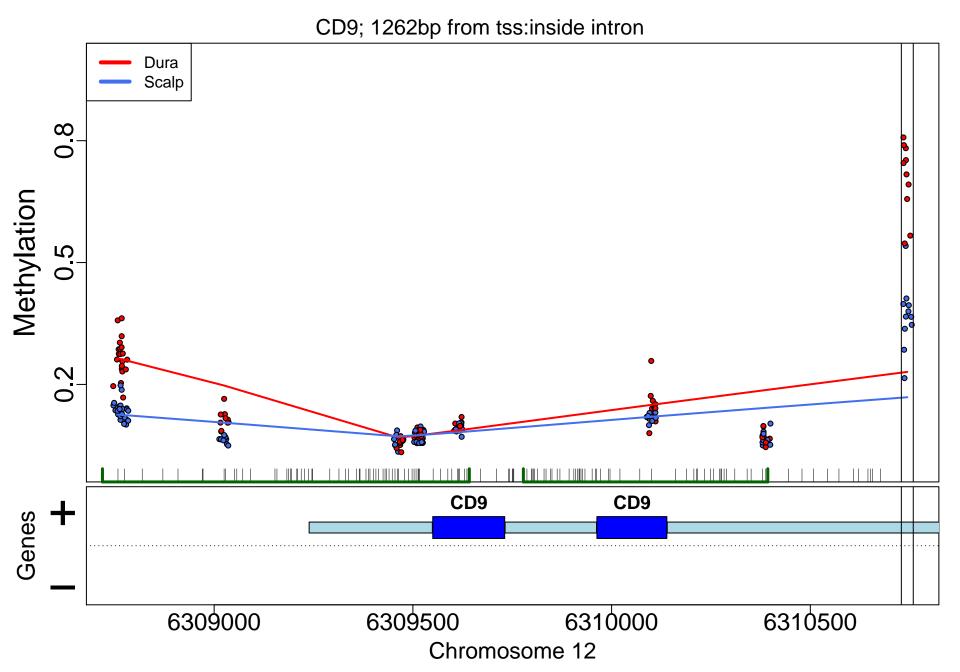


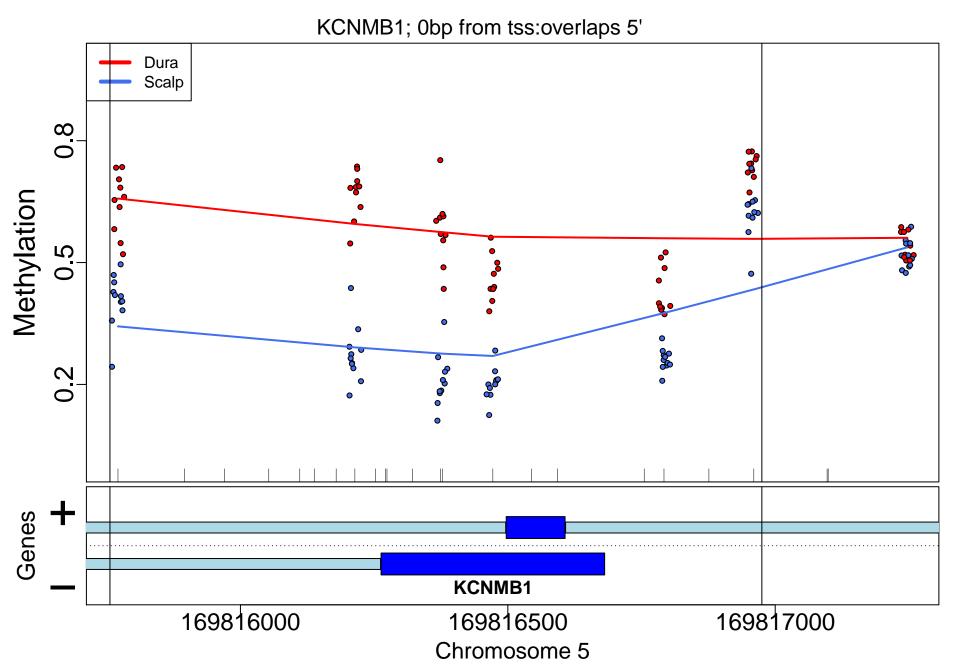


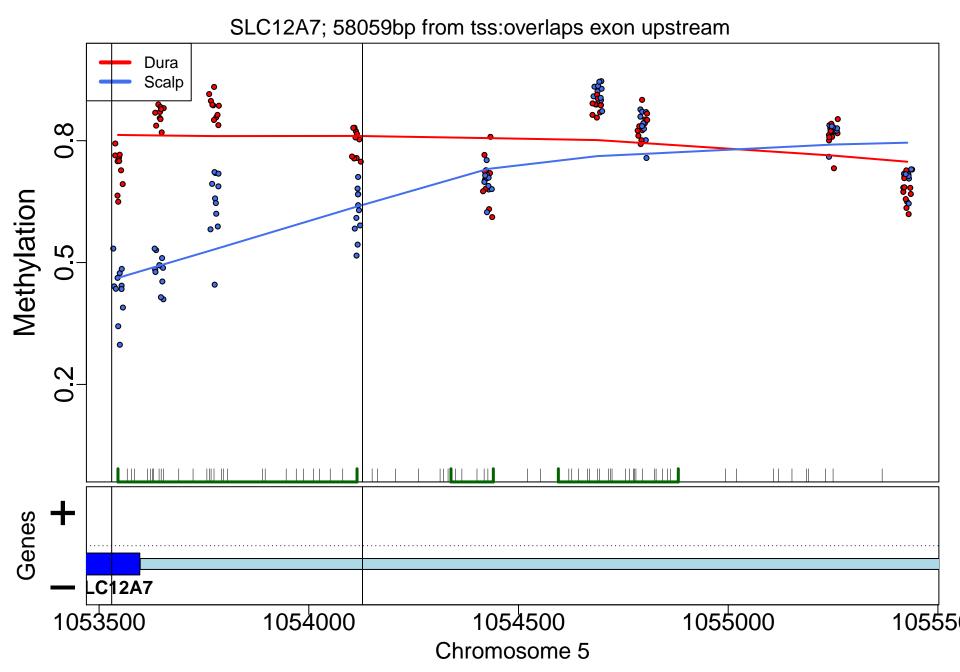


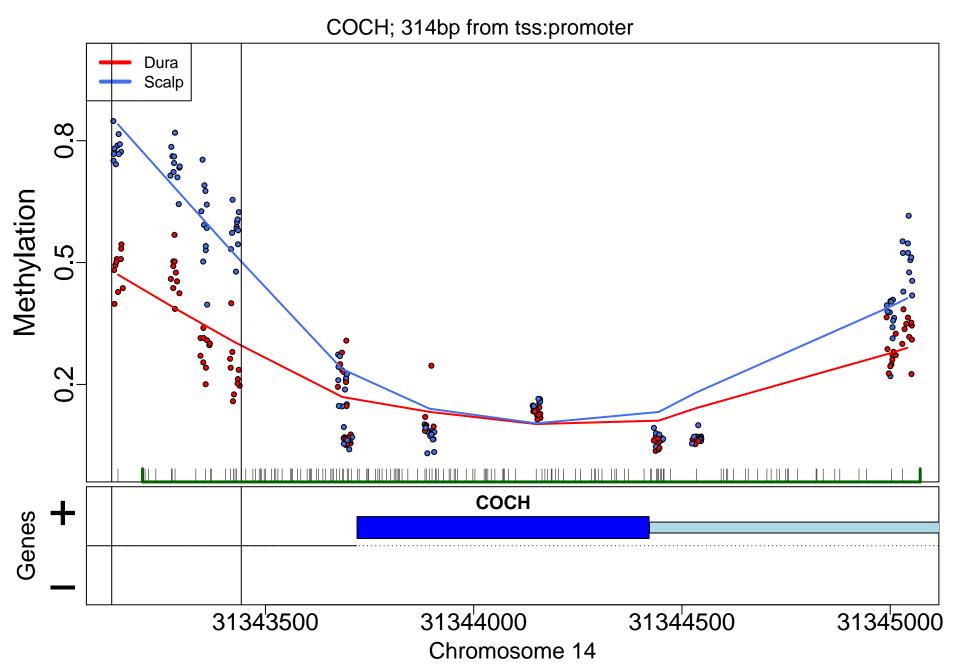


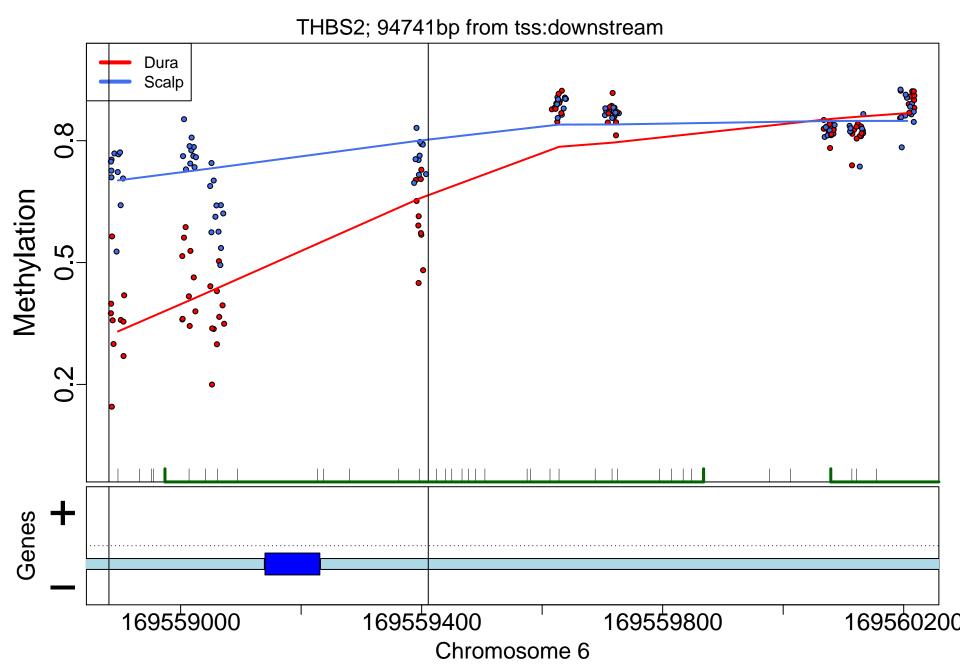
RNF8; 411bp from tss:promoter Dura Scalp <u>ω</u>_ Methylation 0.2 RNF8 Genes 37321400 3732 1800 37322200 37322600 Chromosome 6

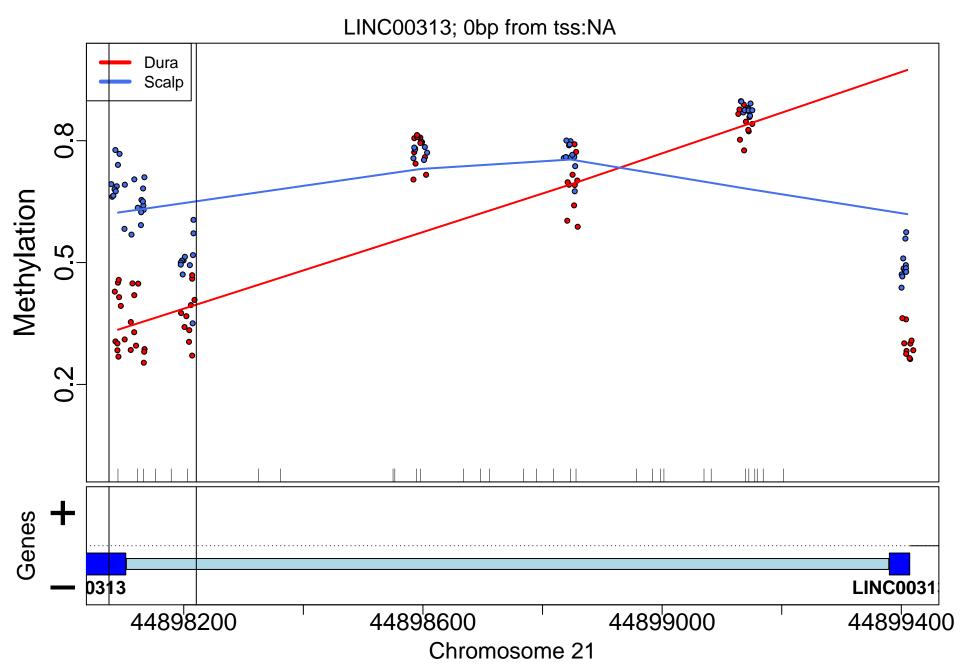


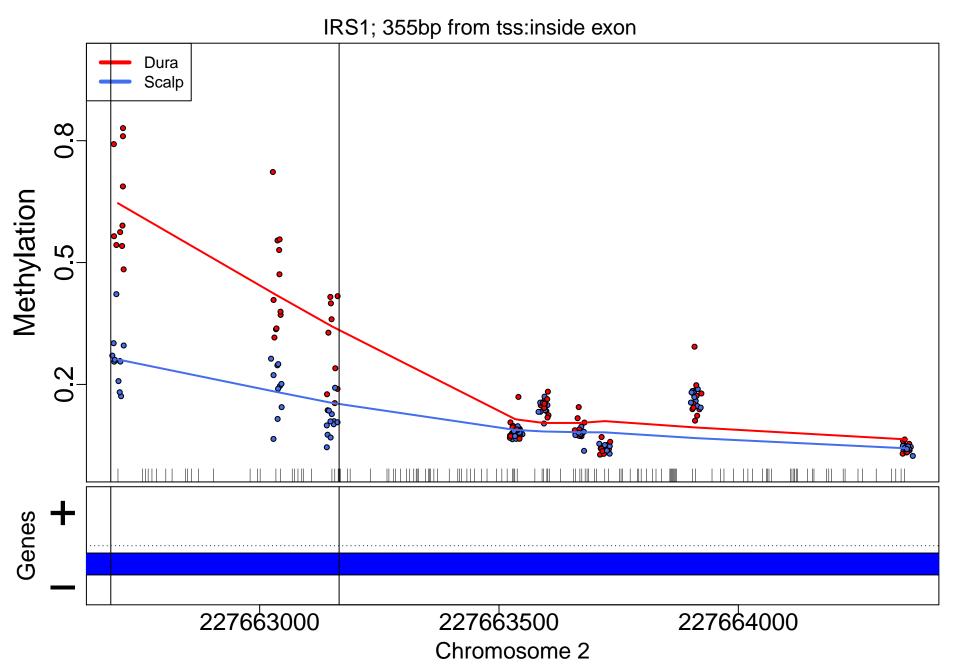


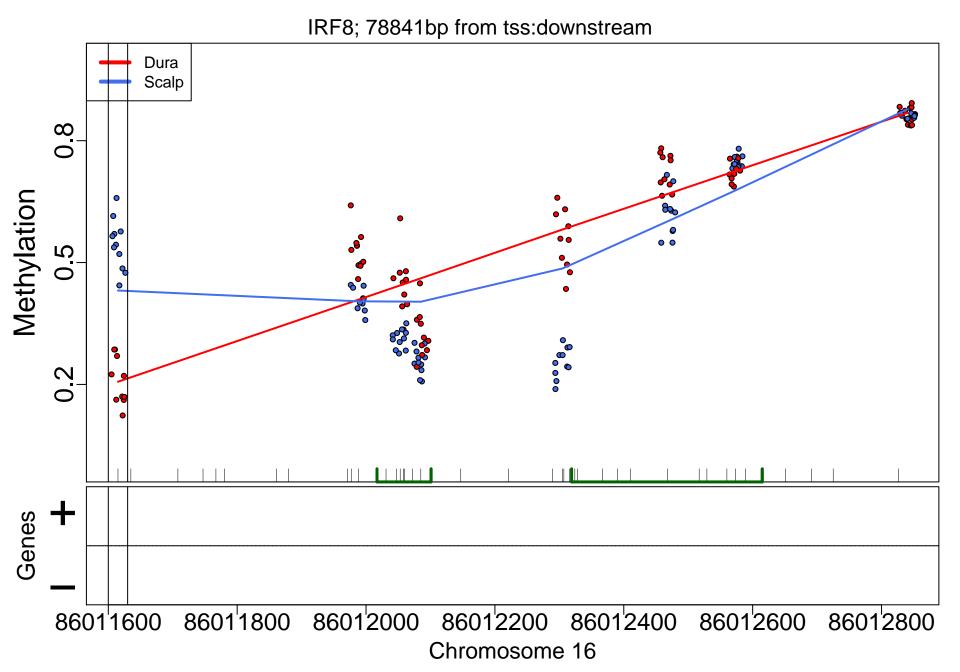


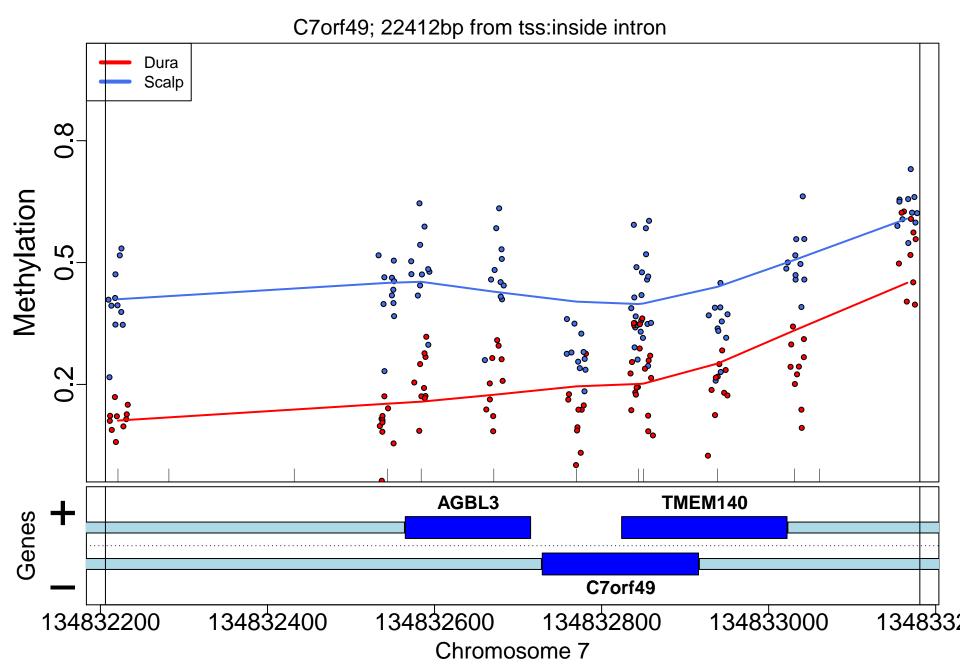


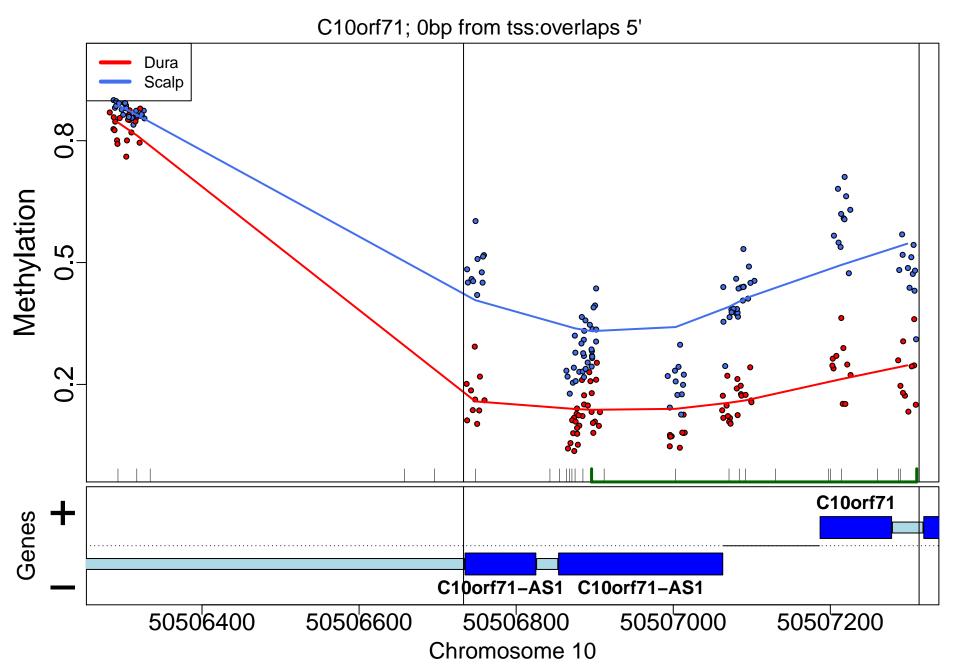




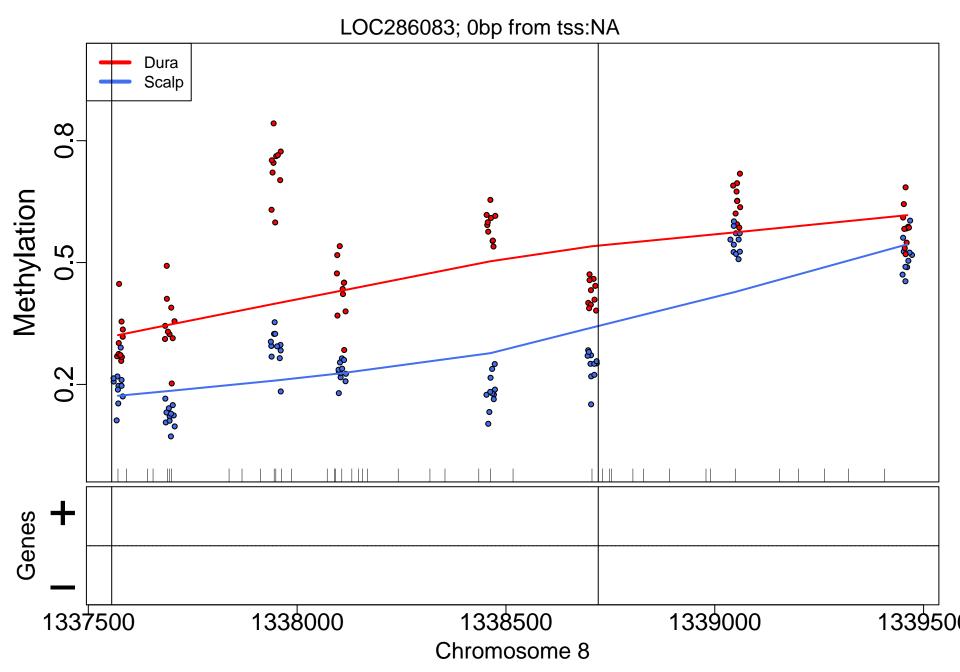


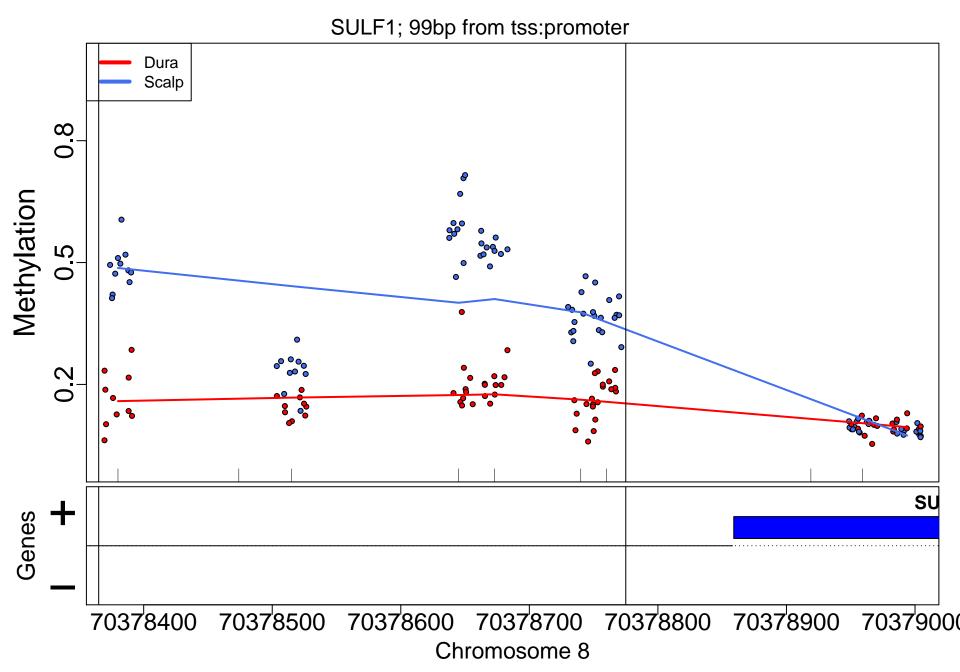


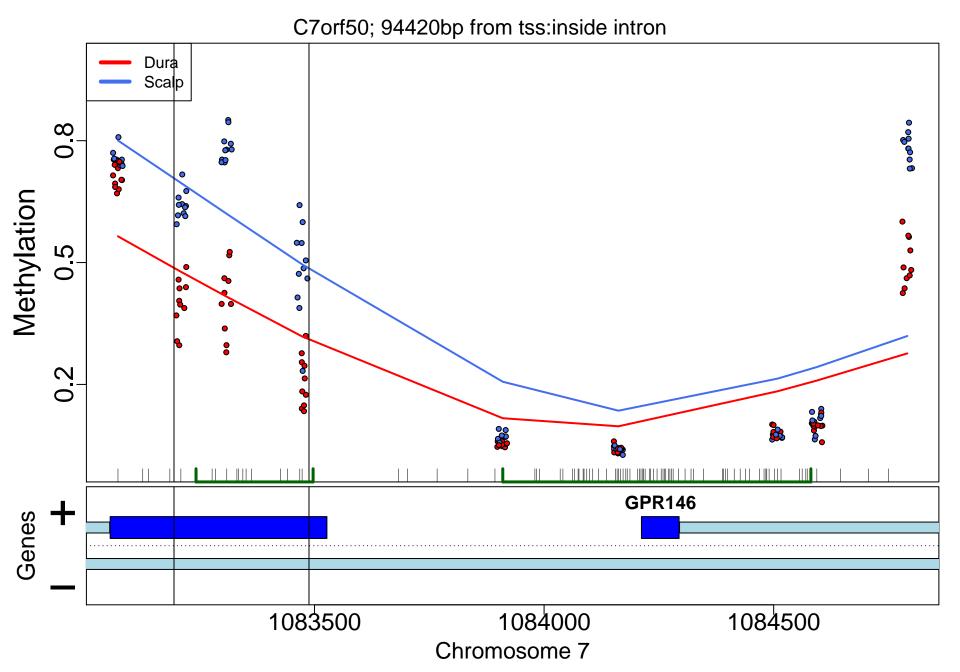




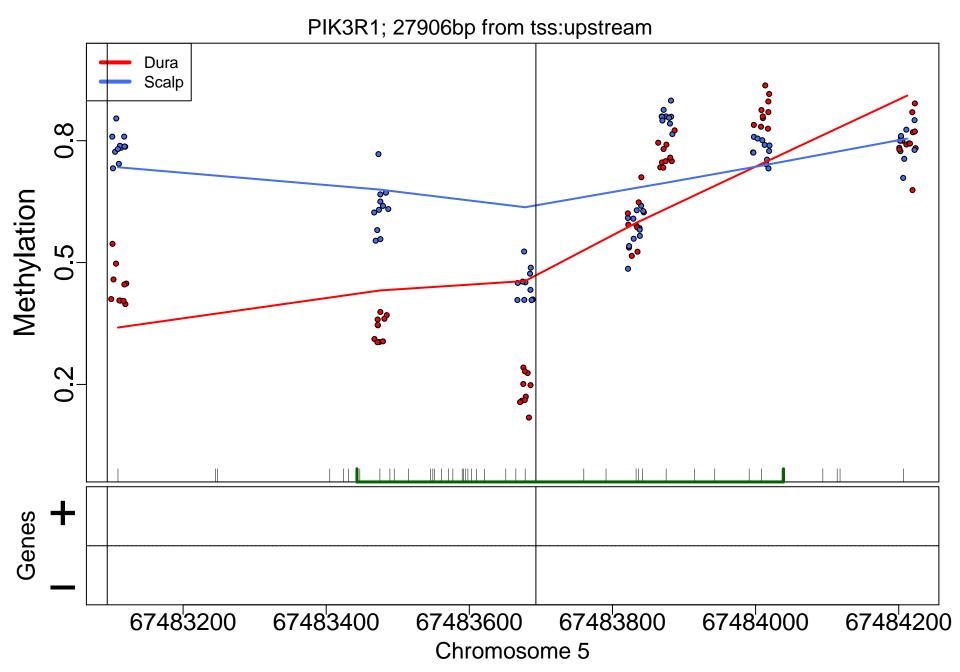
LINC01101; 23076bp from tss:downstream Dura Scalp <u>ω</u>_ Methylation 0.7 Genes 121200000 121200800 121199600 121200400 Chromosome 2

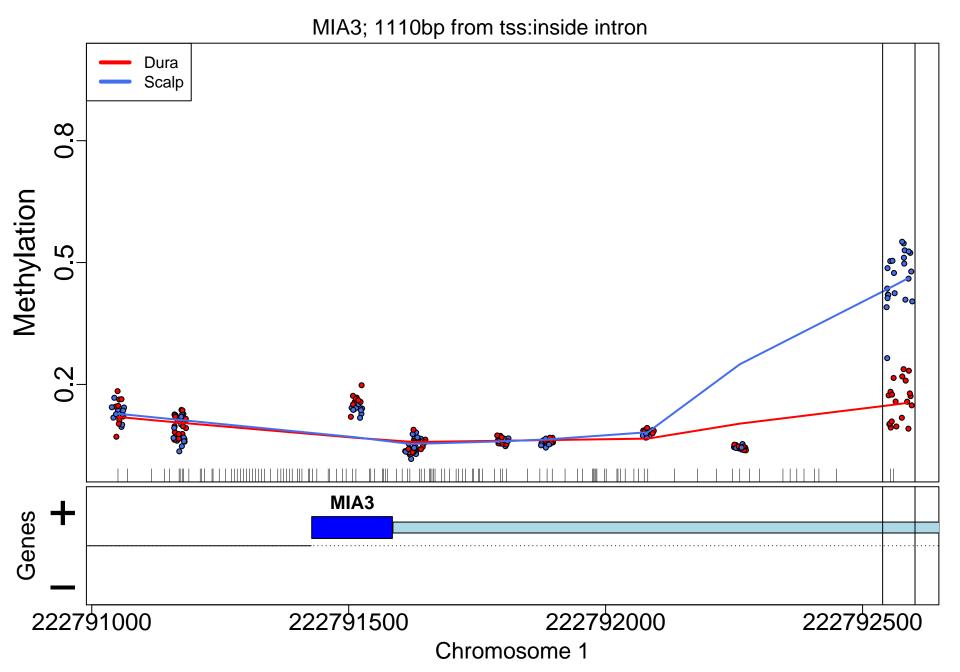


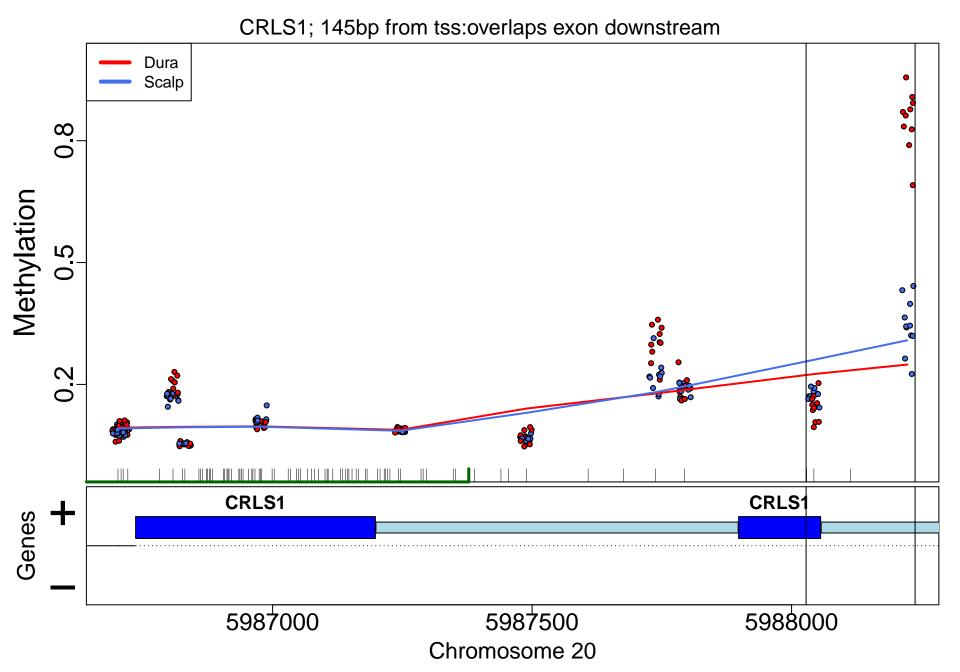


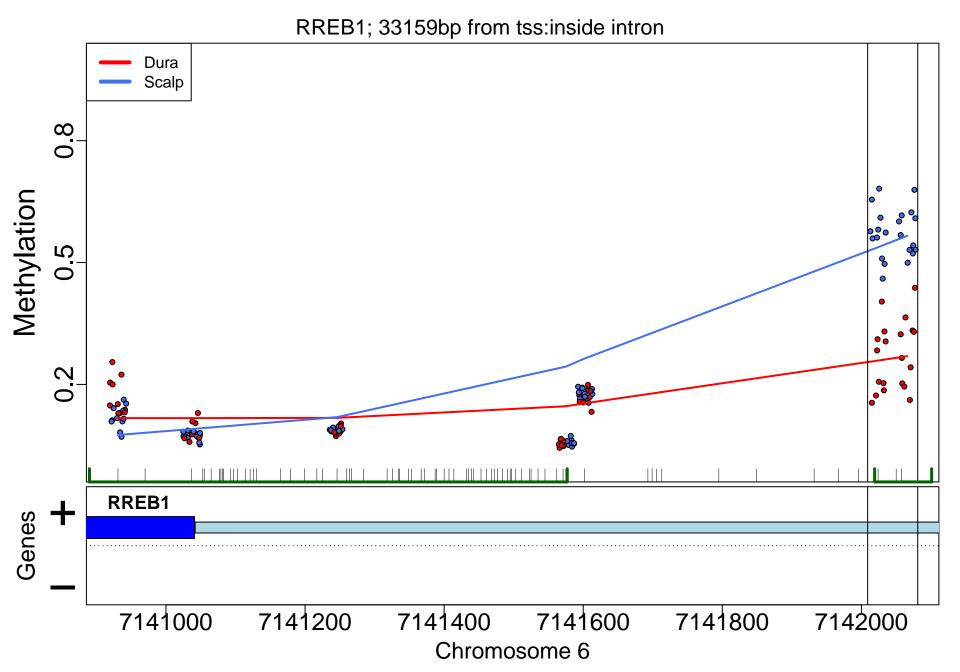


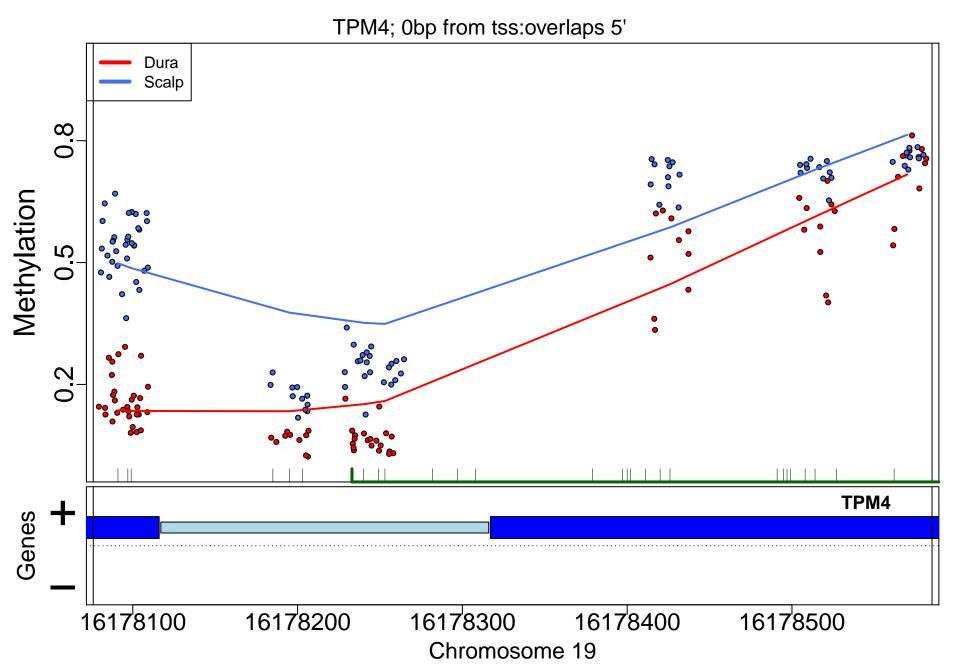
SLC18A2; 348bp from tss:promoter Dura Scalp <u>∞</u>. Methylation SLC18A2 SLC18A2 Genes 119000500 119001000 119001500 Chromosome 10

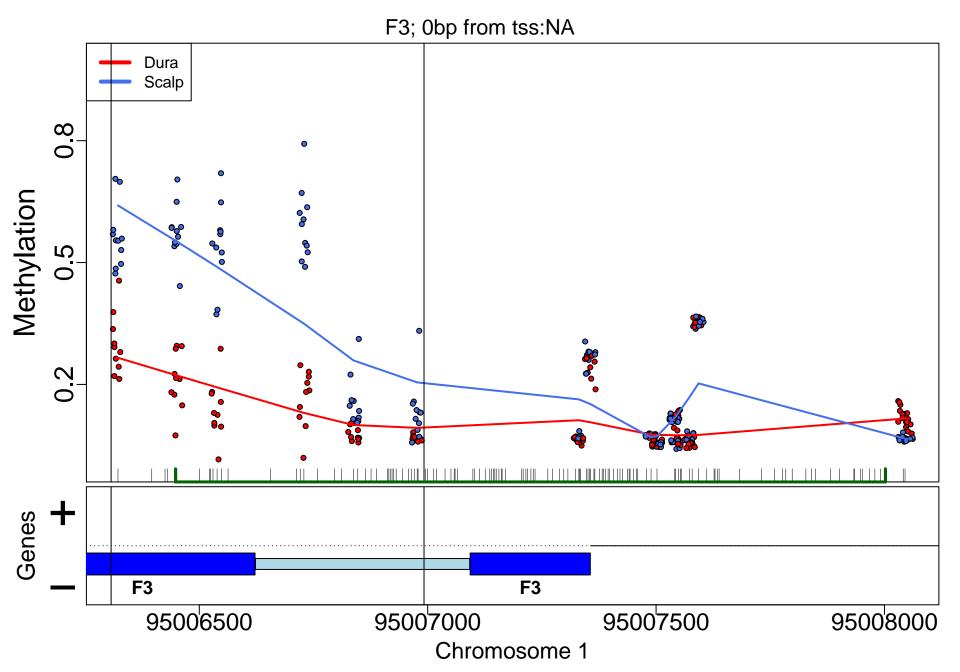


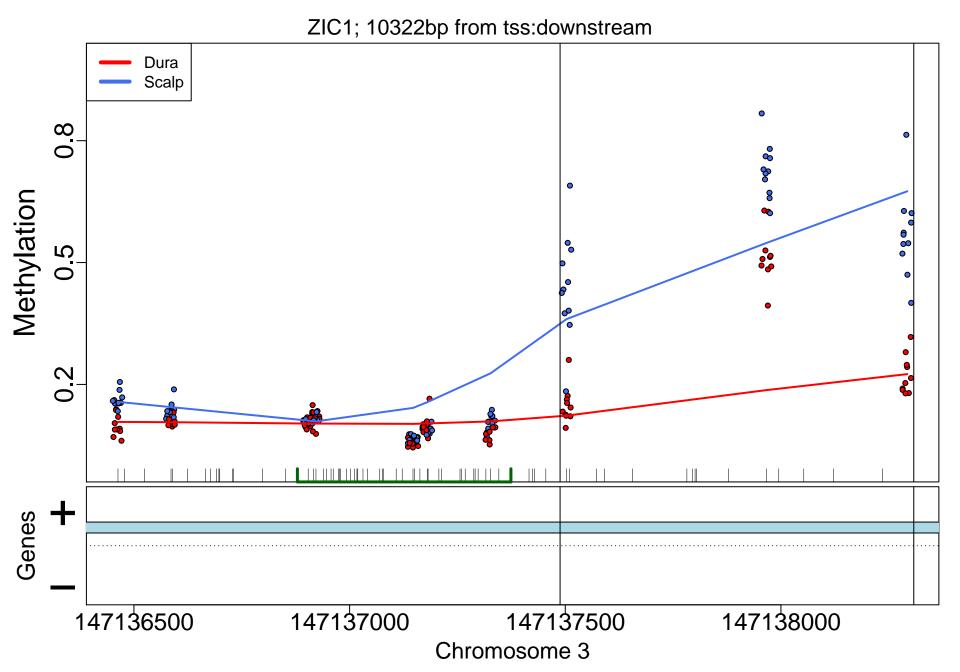


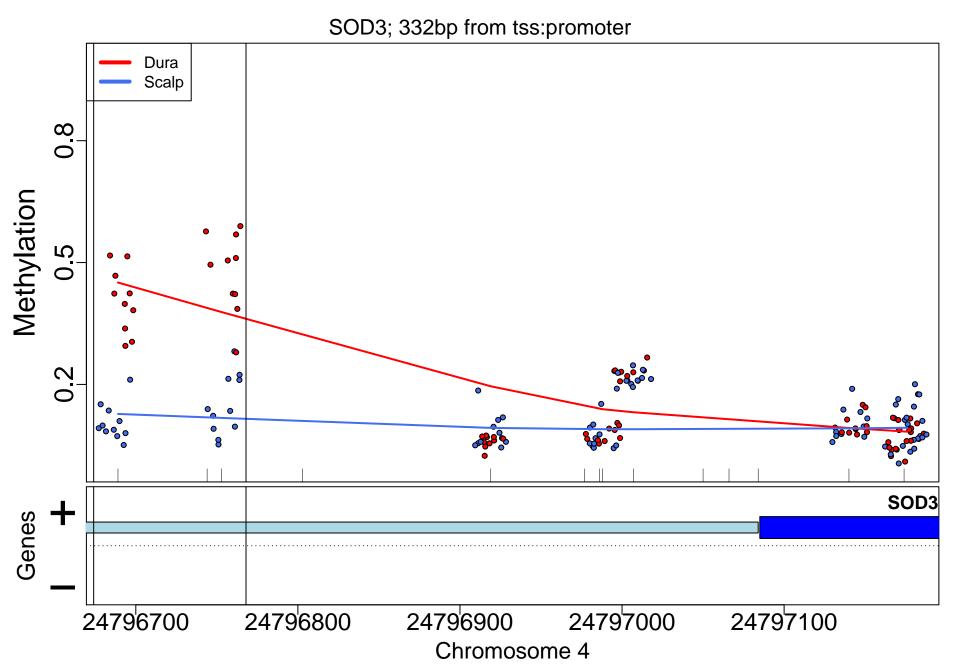




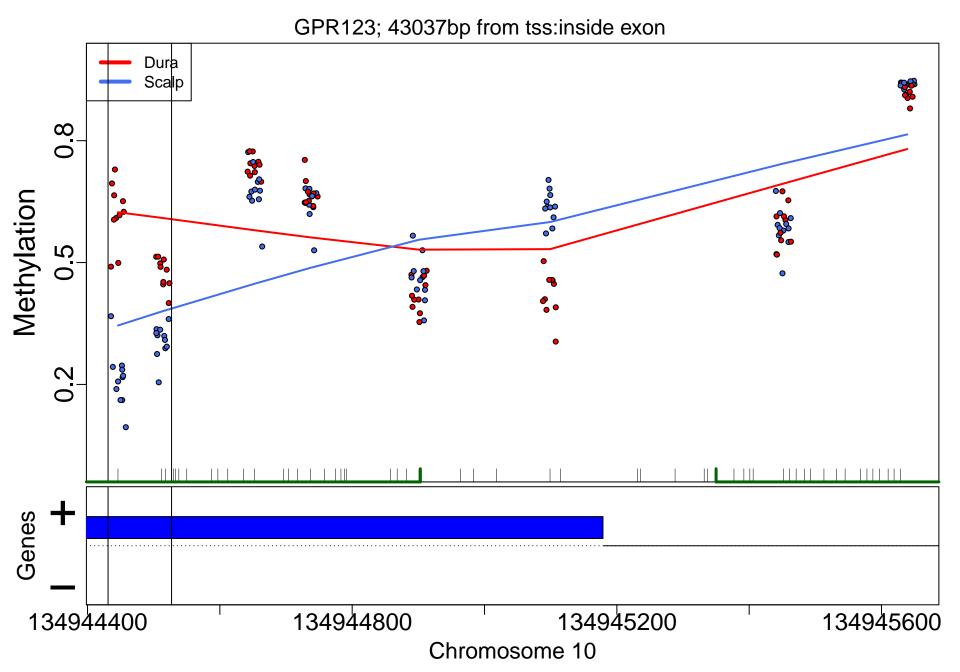




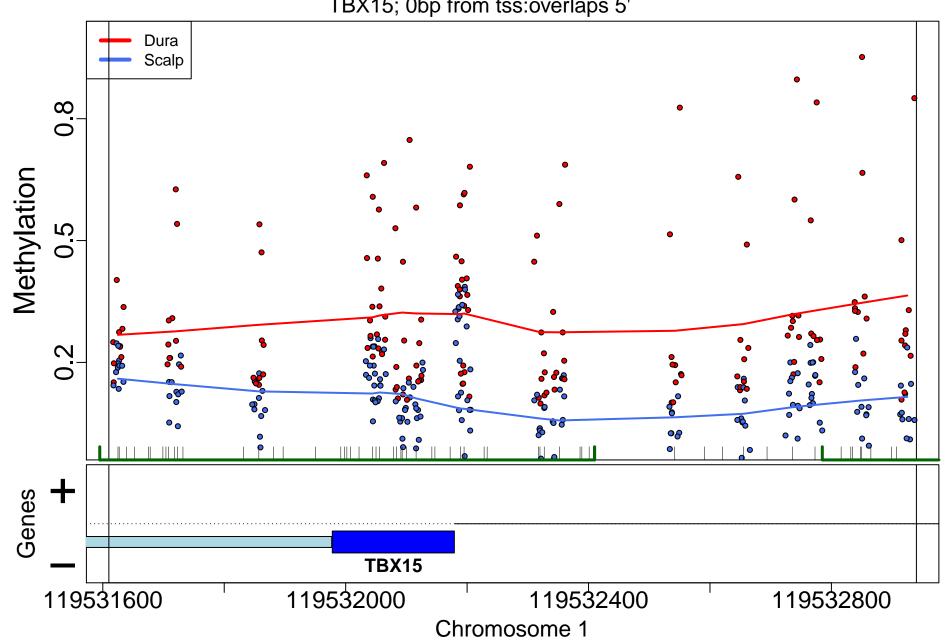


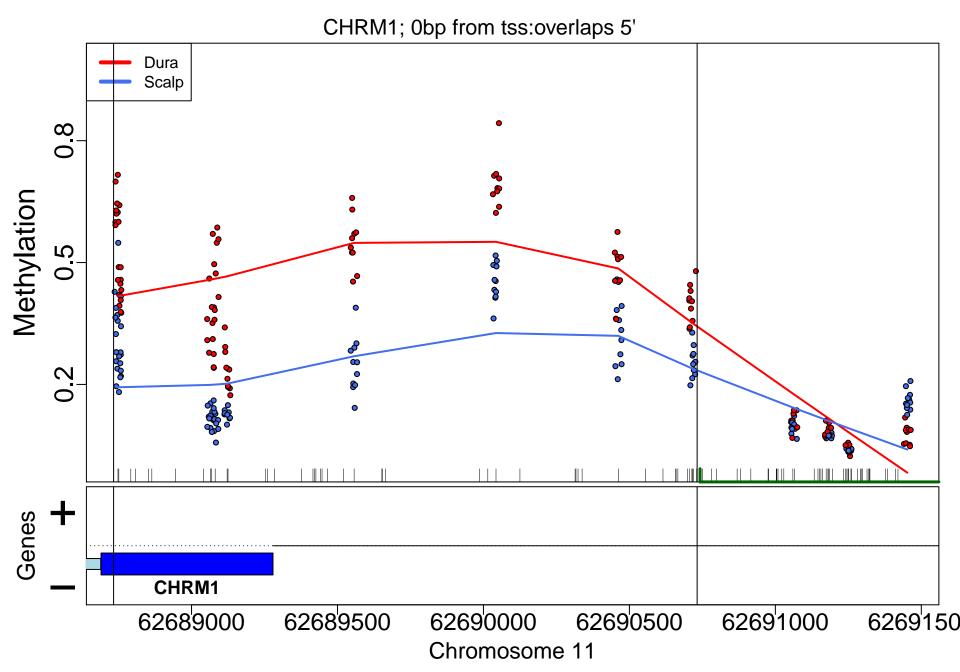


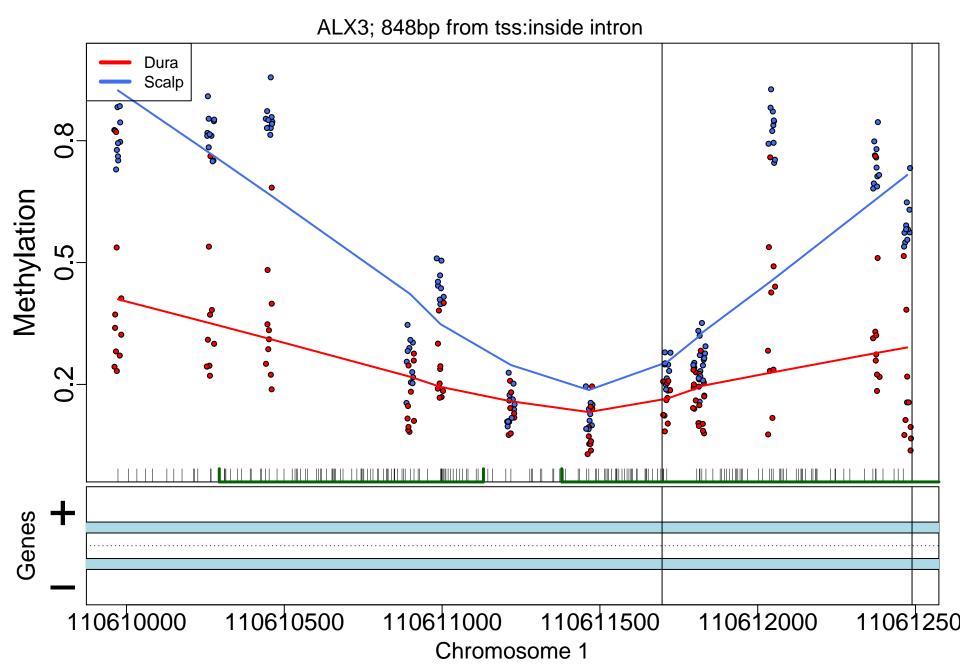
MKI67; 414102bp from tss:upstream Dura Scalp <u>∞</u>. Methylation 88°°° 0.2 Genes 130338600 130338800 130339400 130339000 130339200 130339600 Chromosome 10

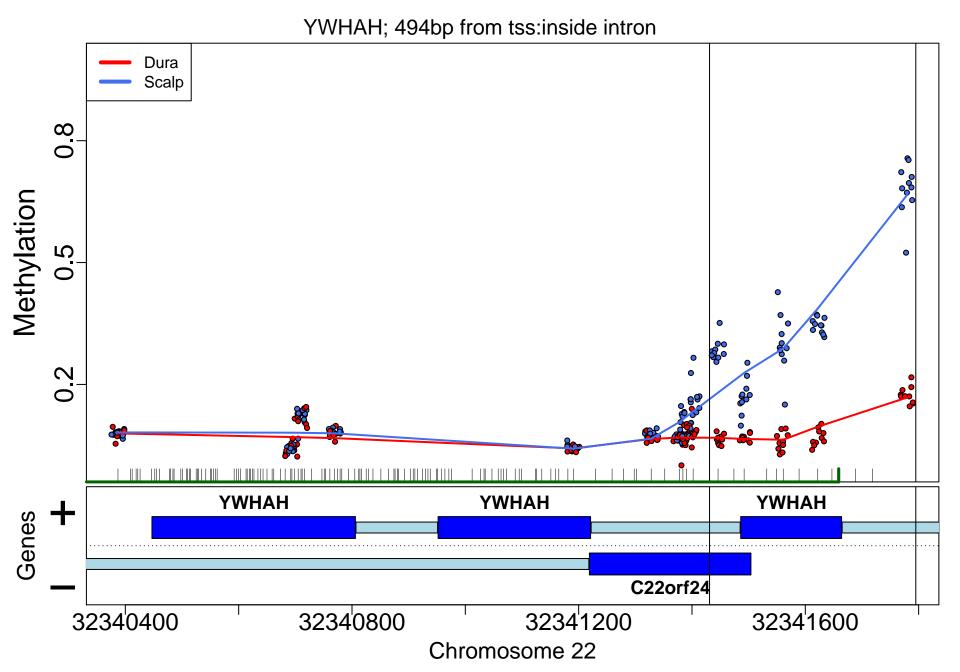


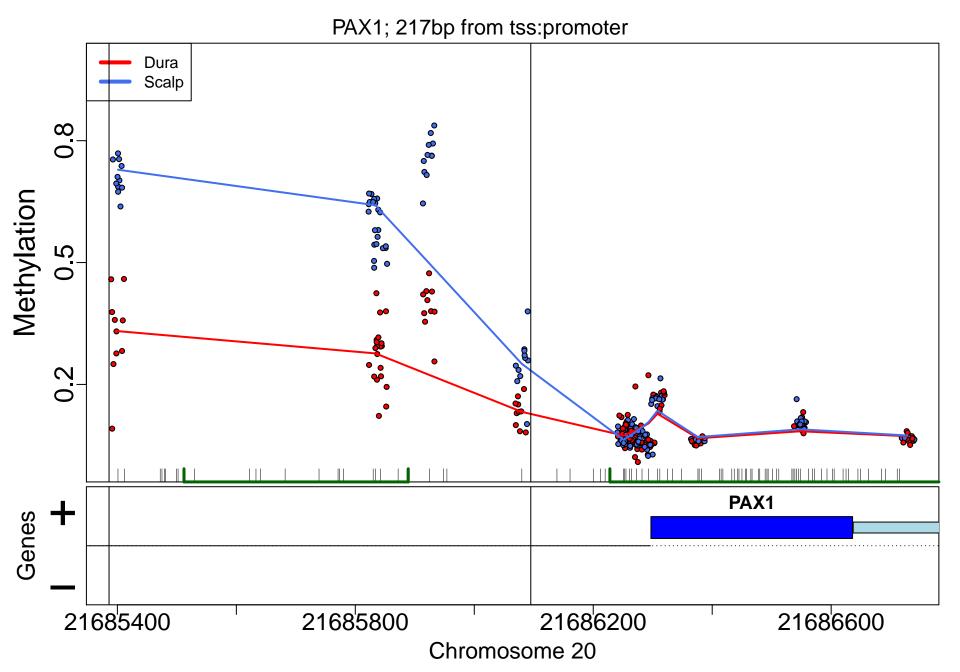
TBX15; 0bp from tss:overlaps 5'

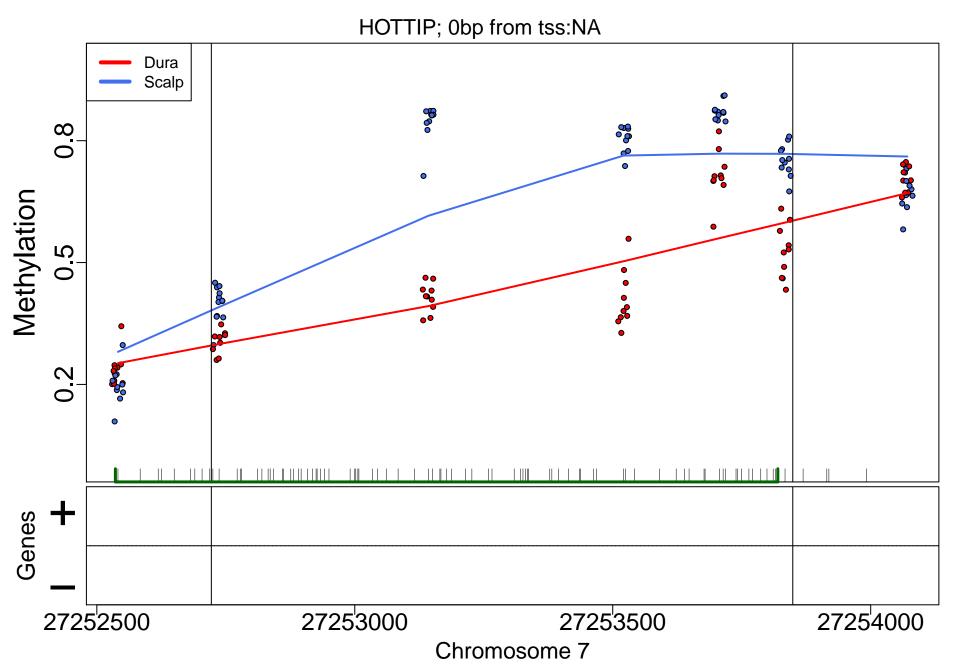


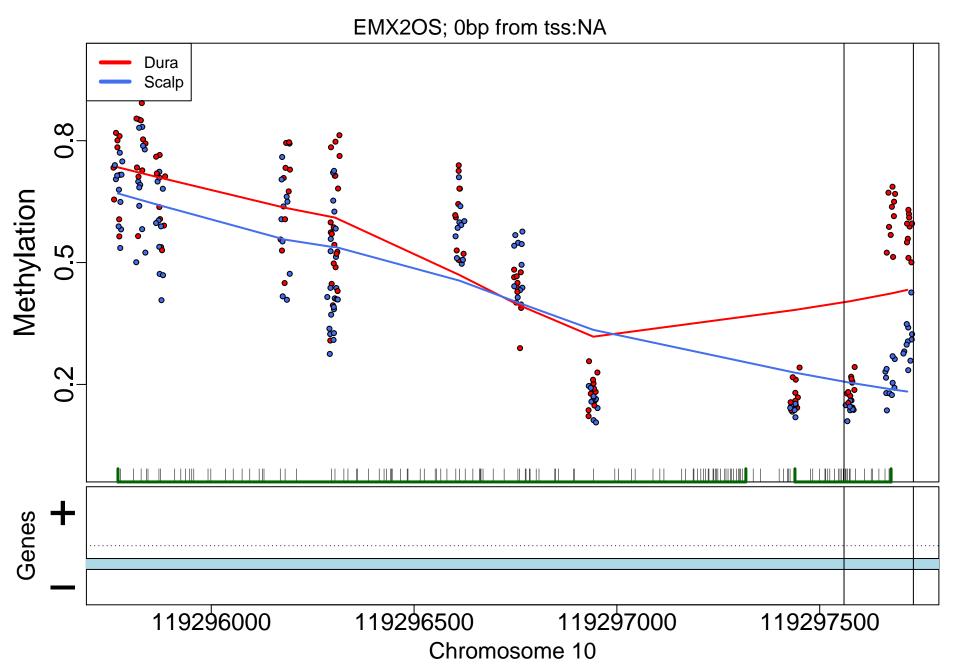


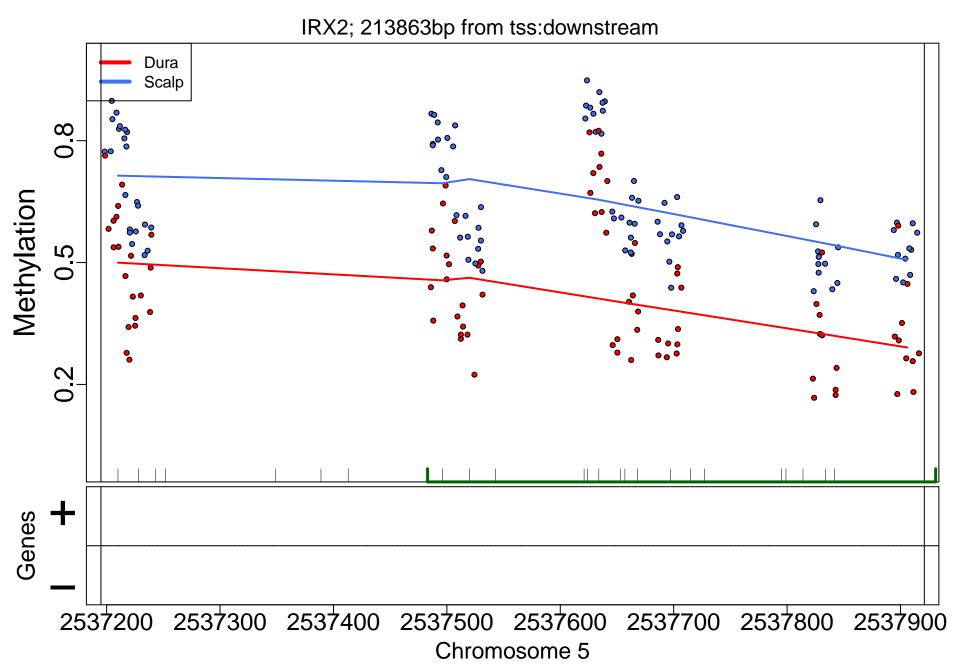


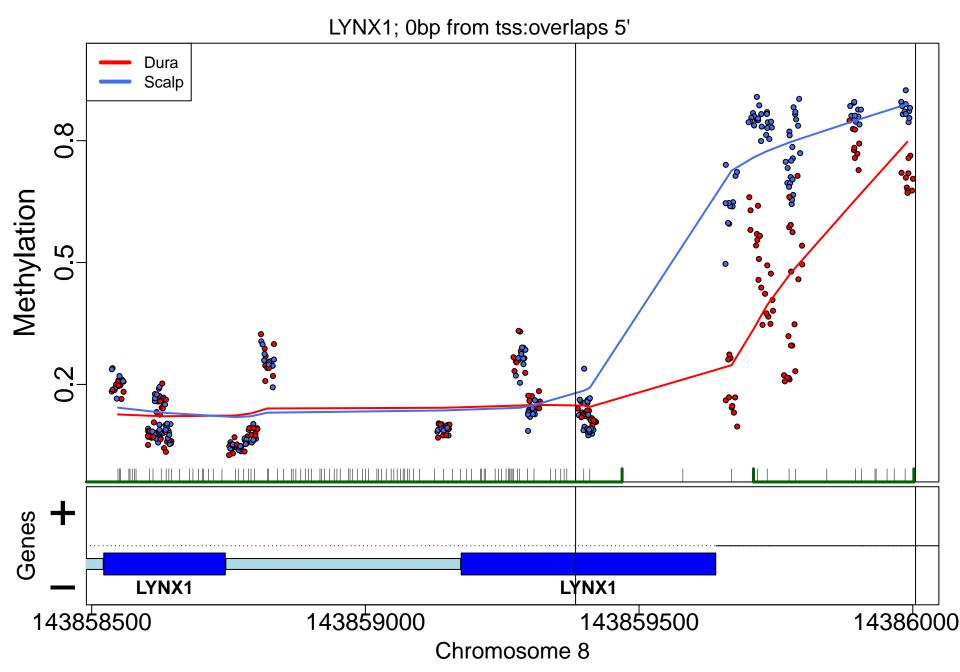


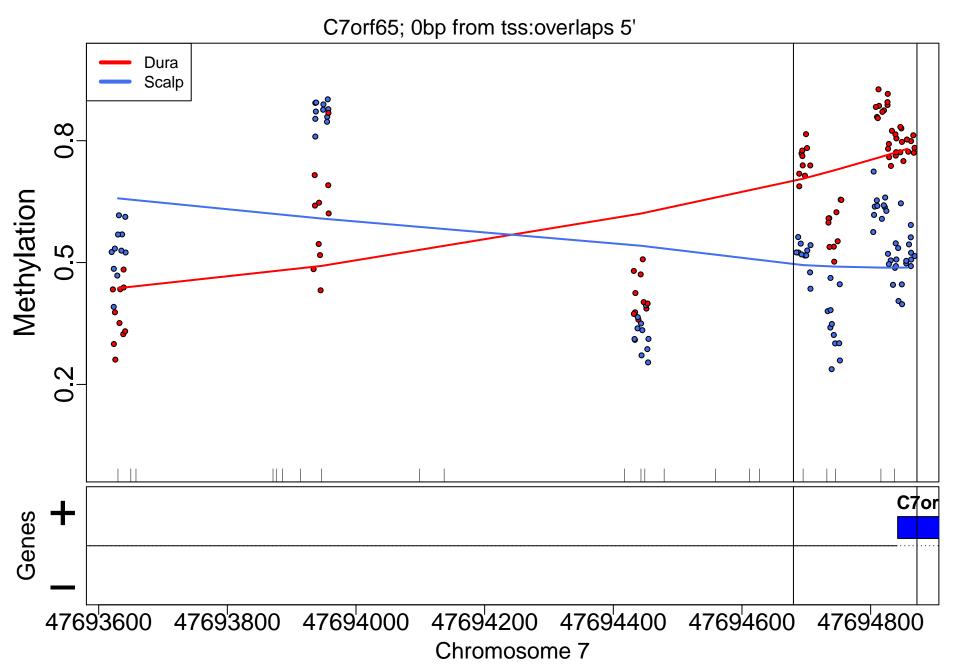


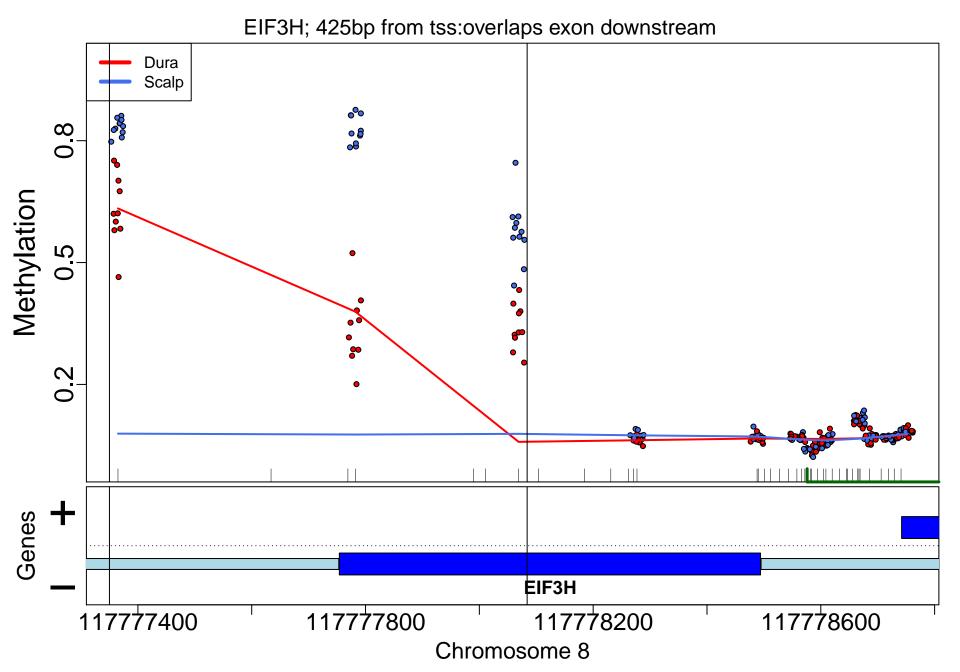


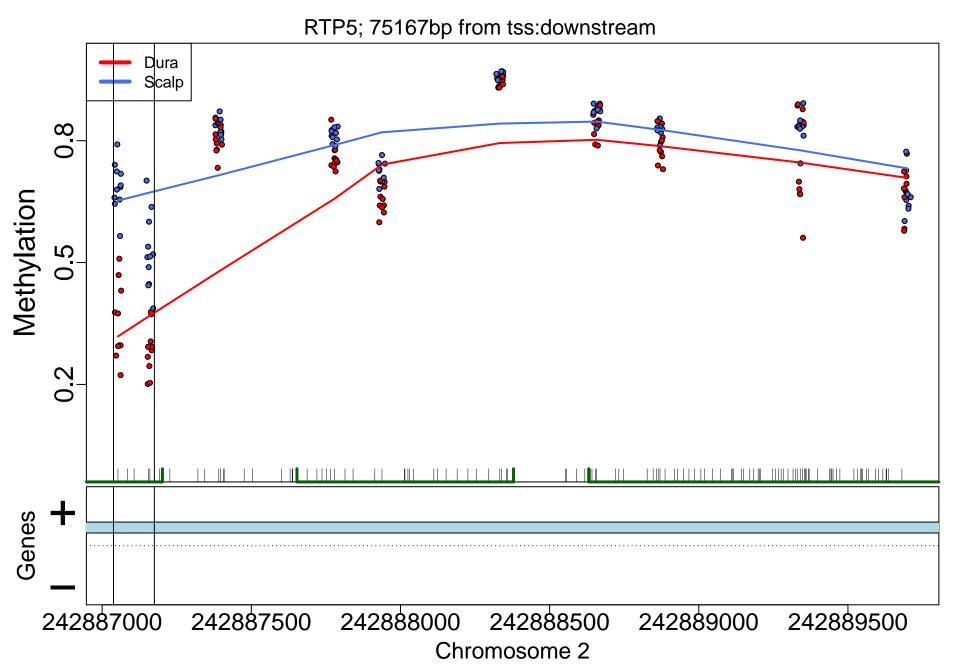


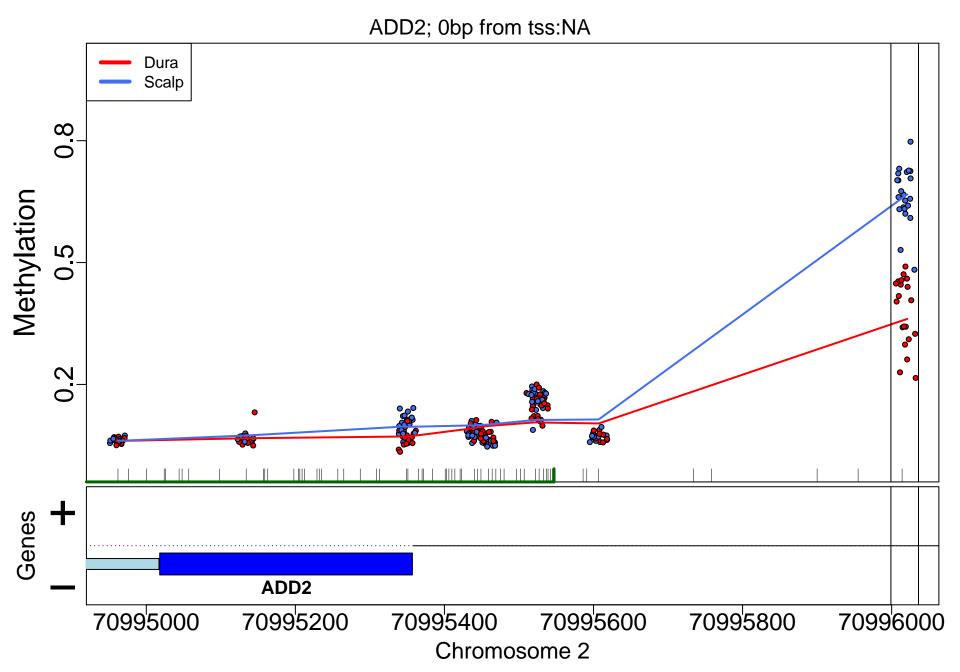


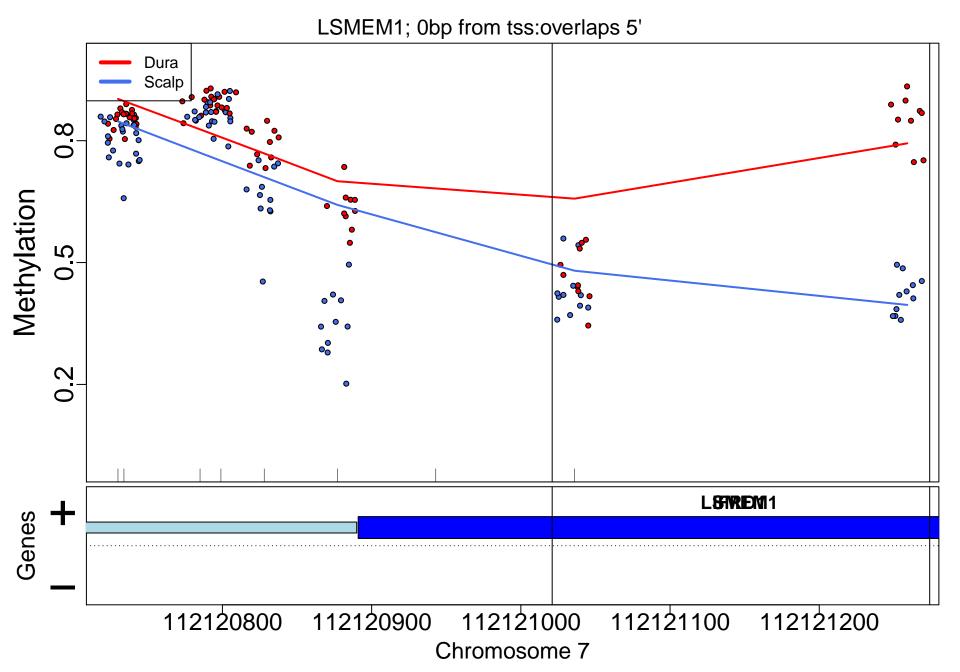


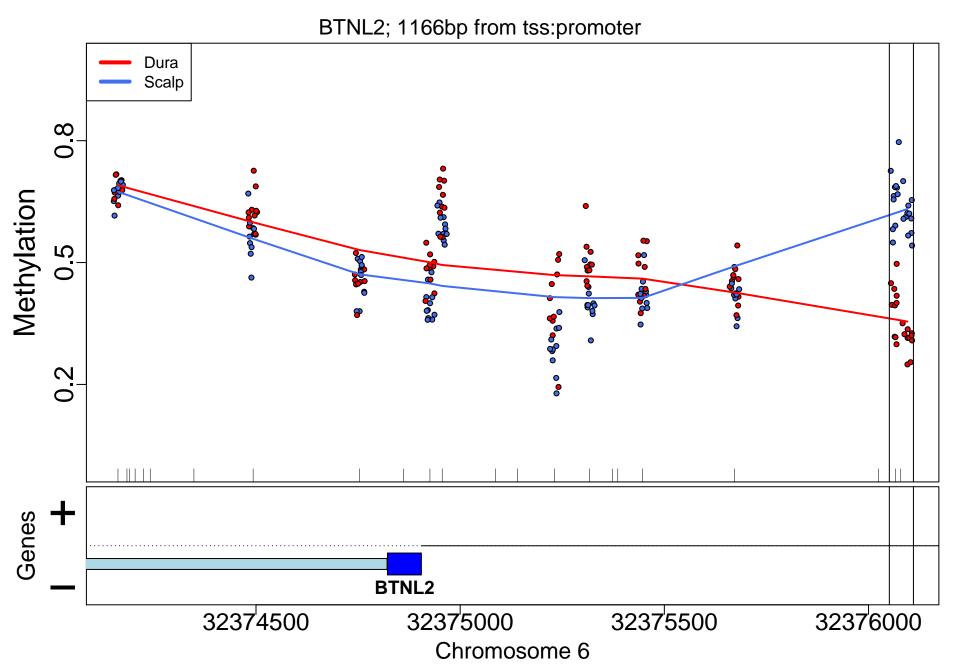


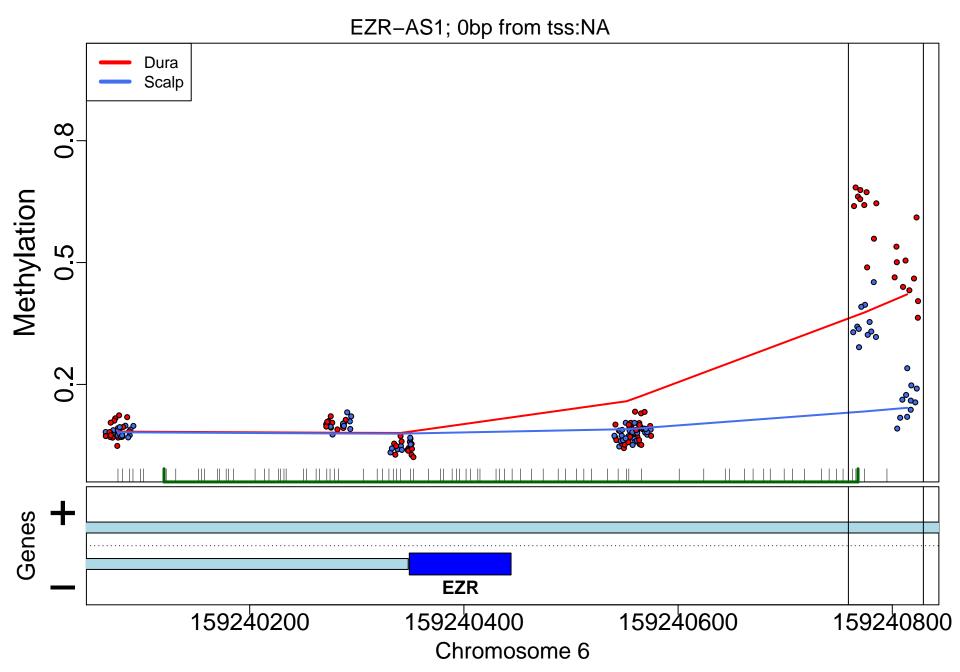


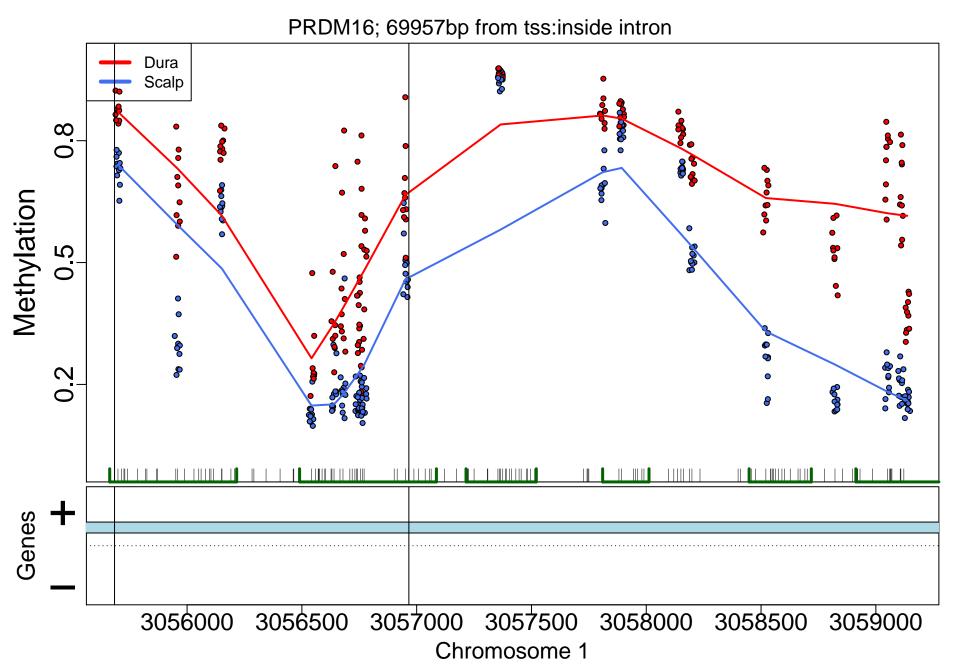


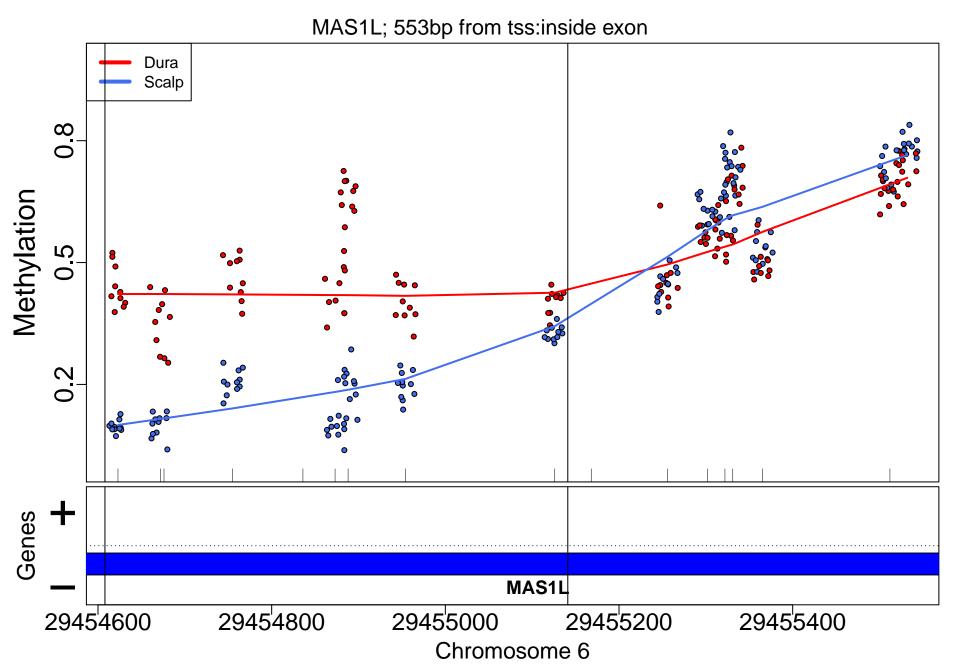


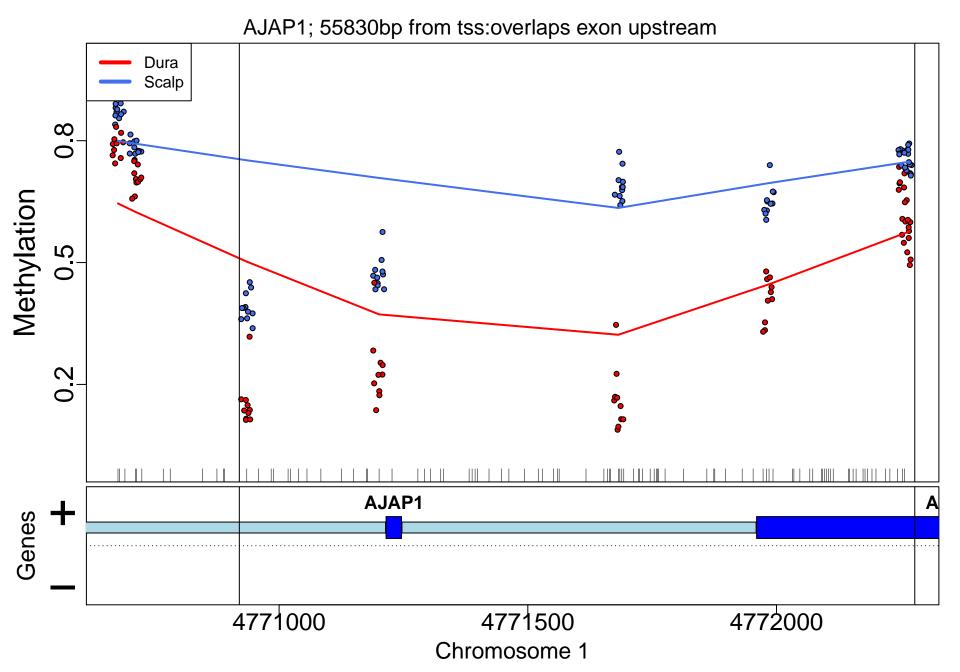


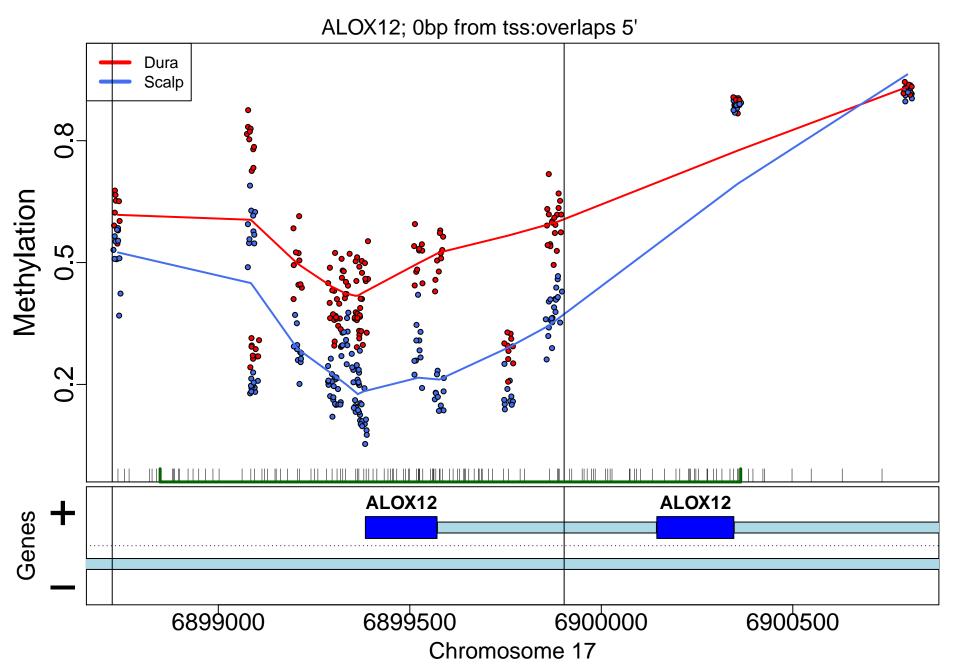


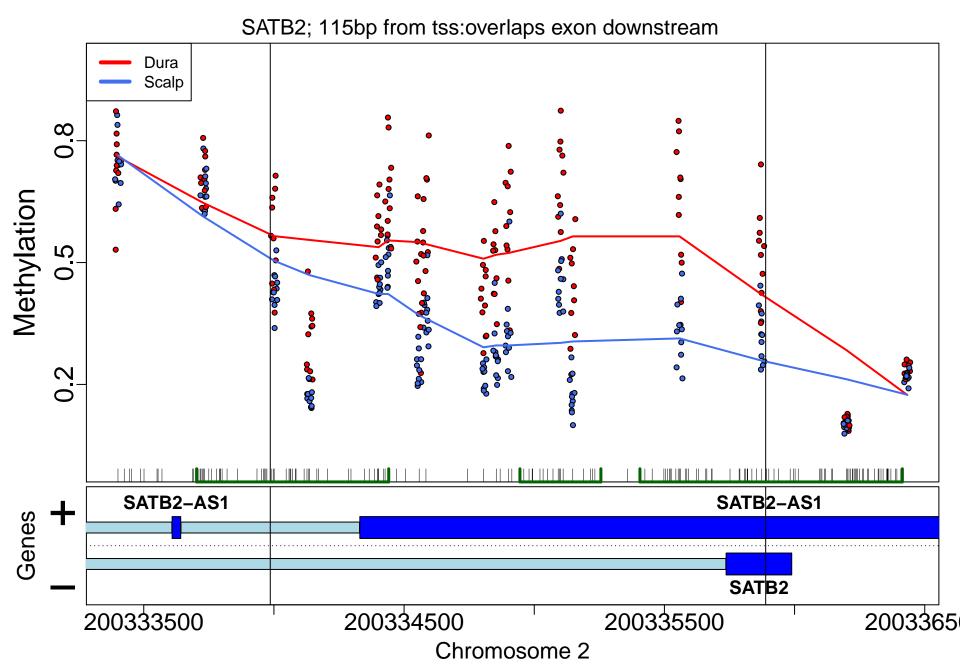


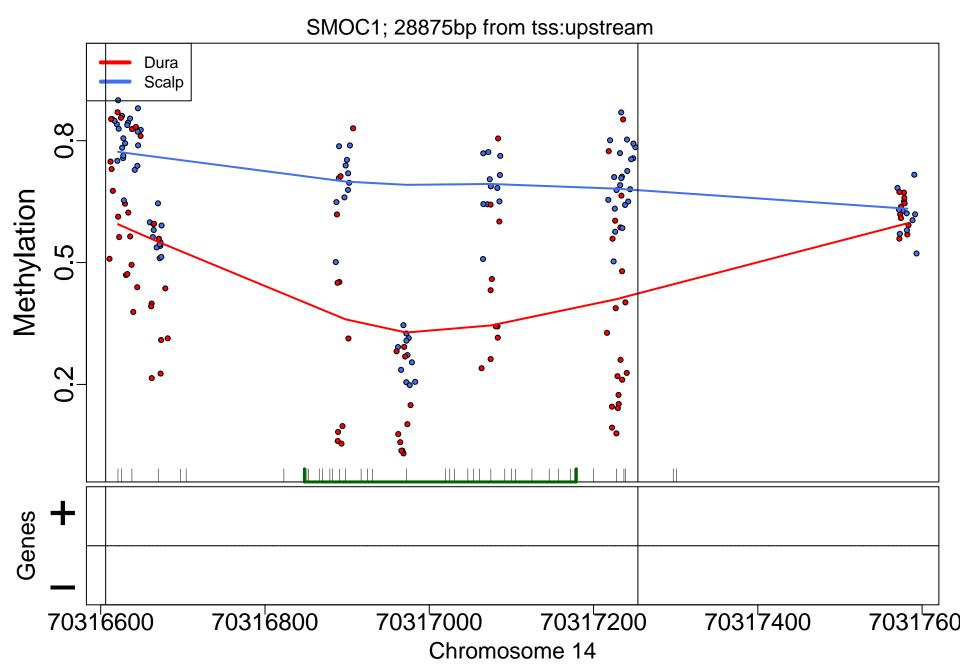


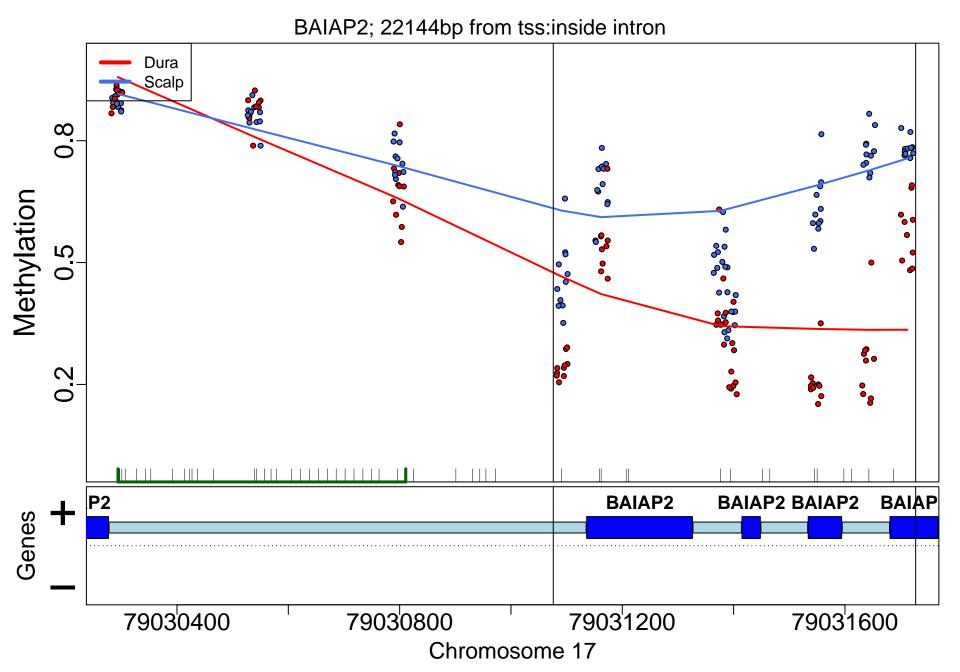


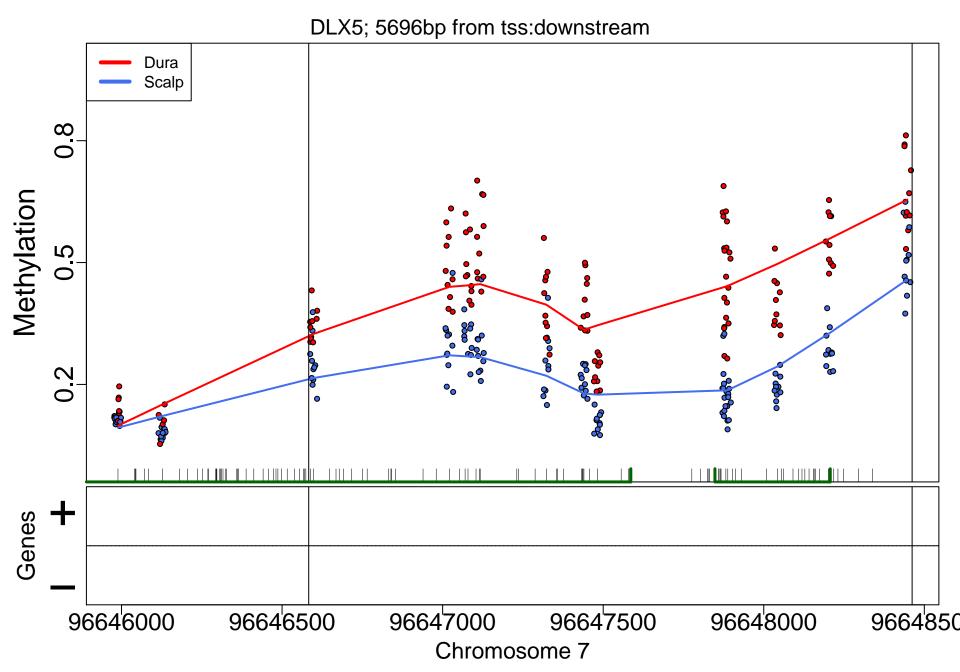


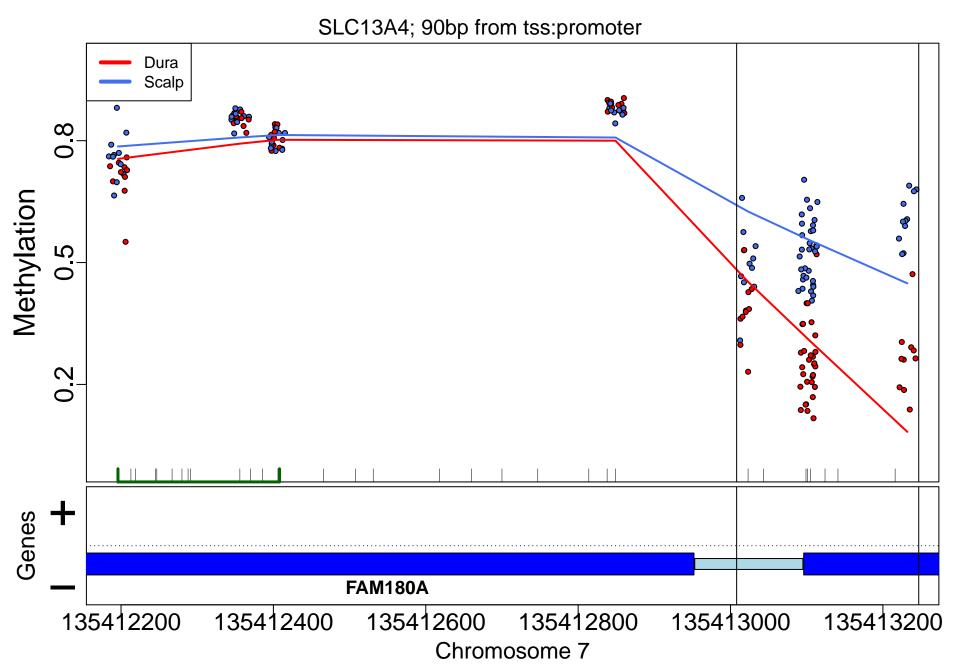


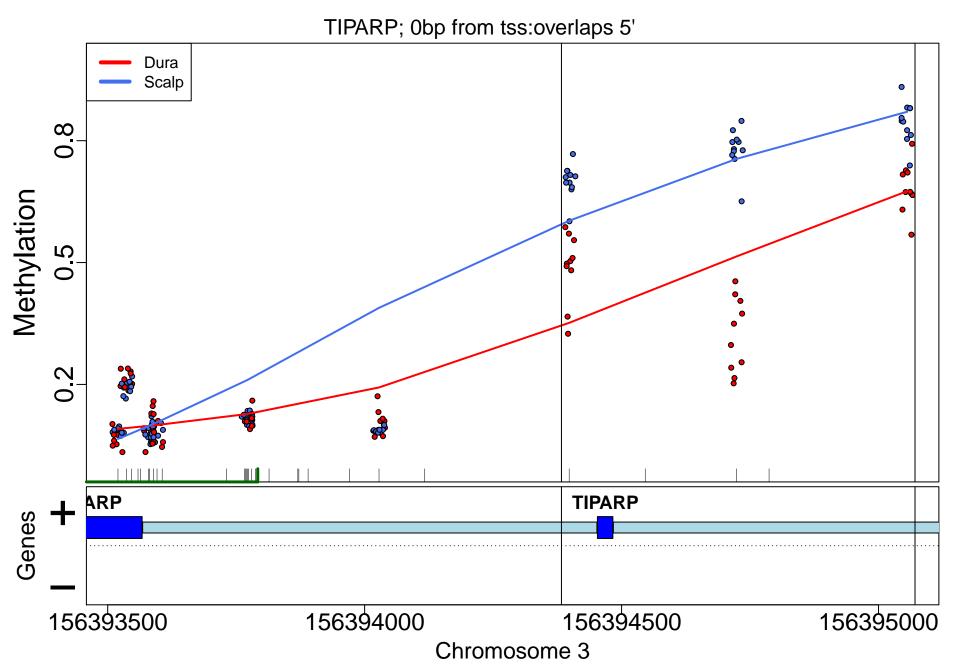


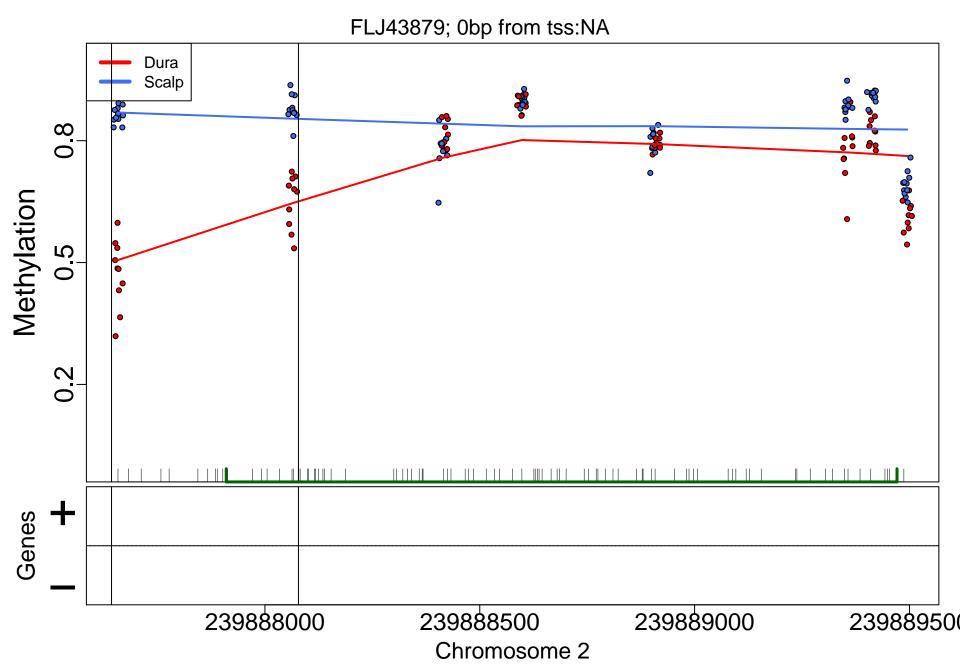




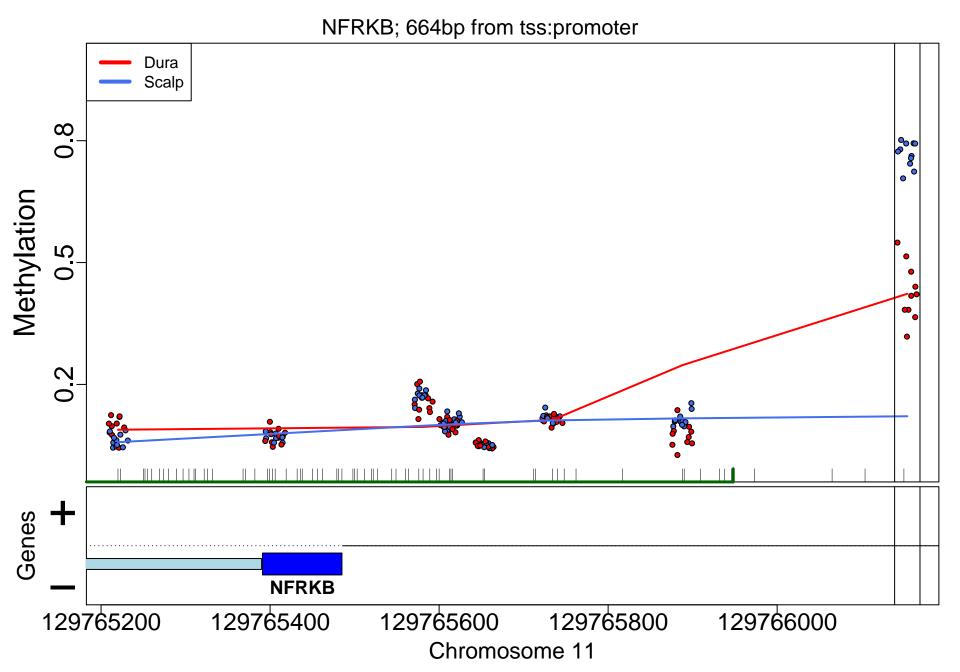




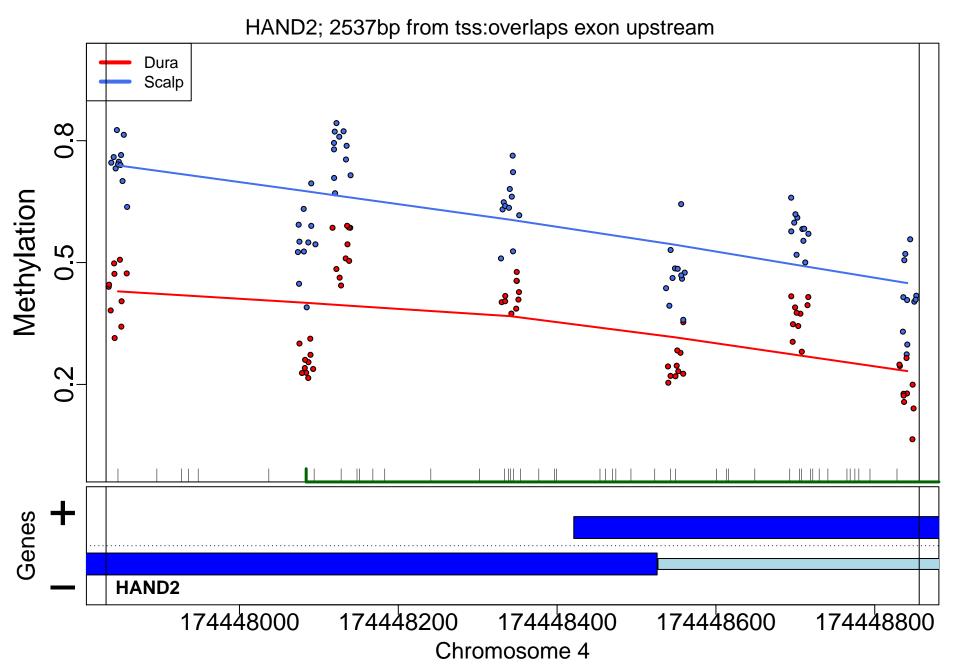


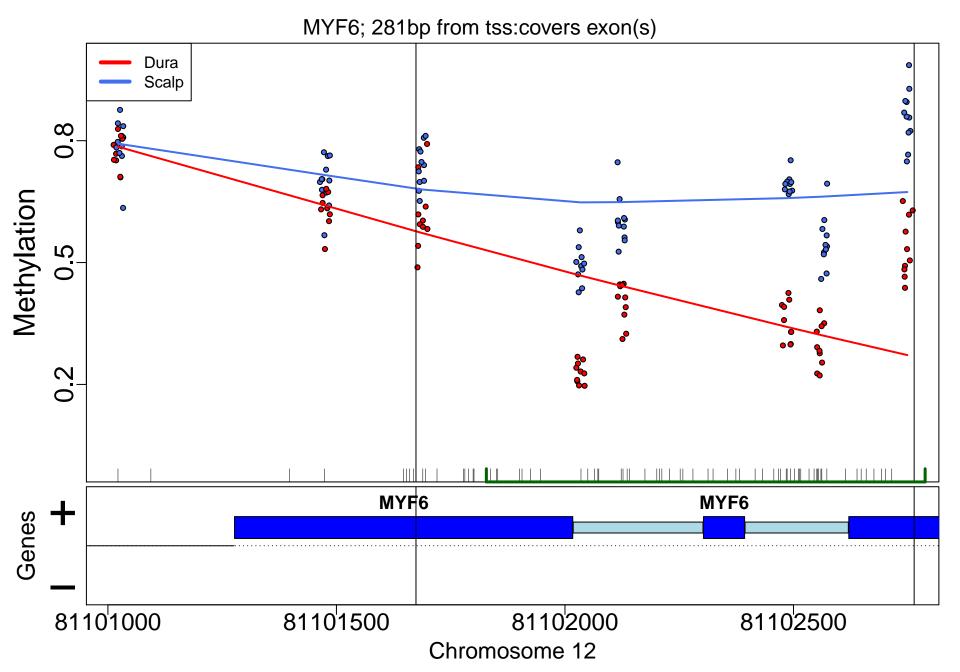


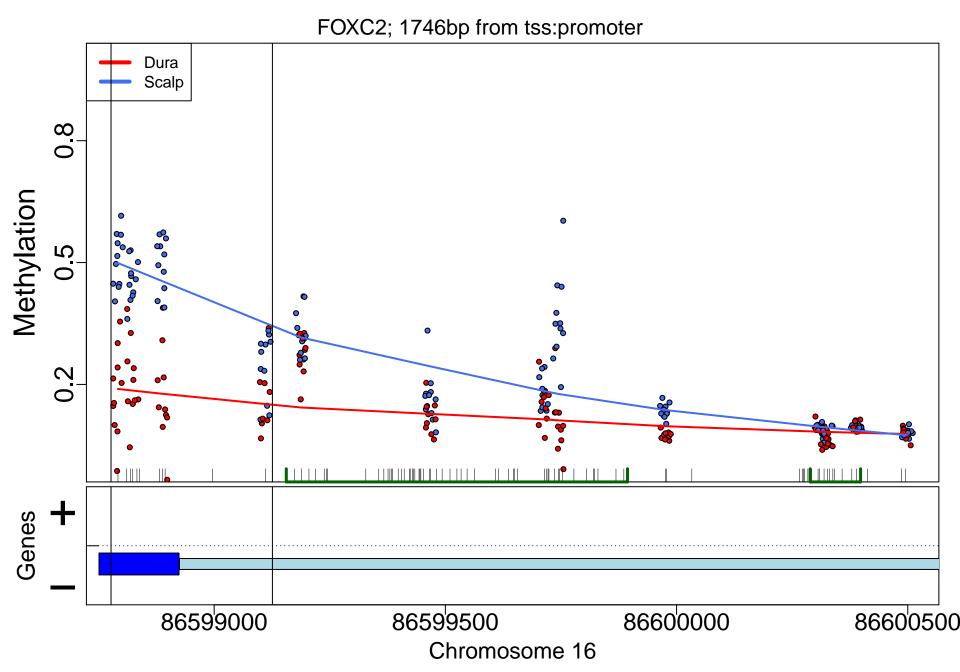
MYCBPAP; 0bp from tss:NA Dura Scalp <u>ω</u>_ Methylation 0.2 **MYCBPAP** Genes 48585200 48585600 48586000 48586400 Chromosome 17

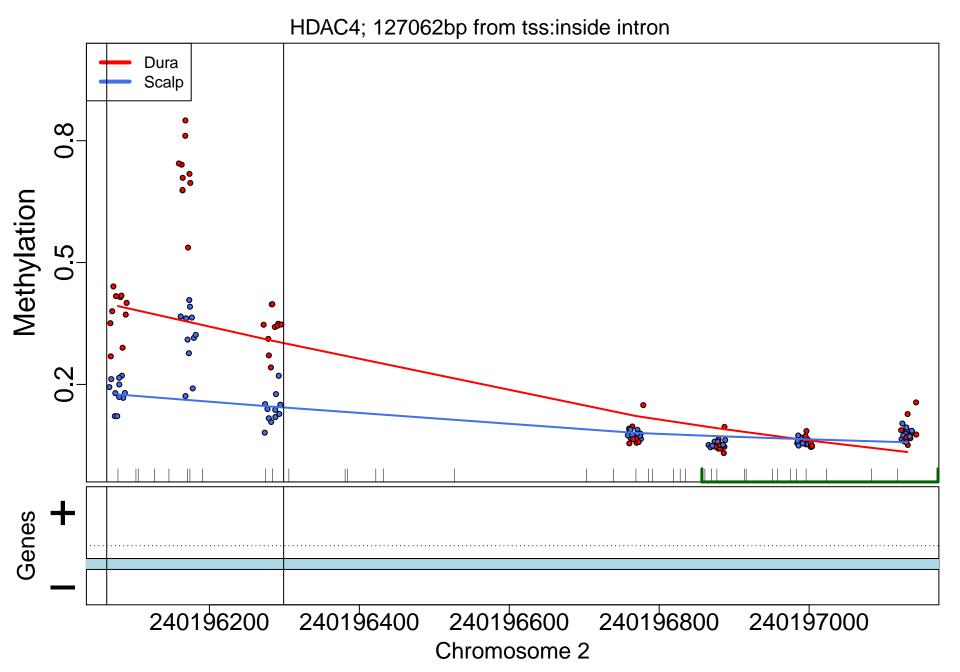


BHLHE22; 5645bp from tss:downstream Dura Scalp <u>∞</u>. Methylation %% % % 0.2 Genes 65498500 65499000 65499500 65500000 Chromosome 8

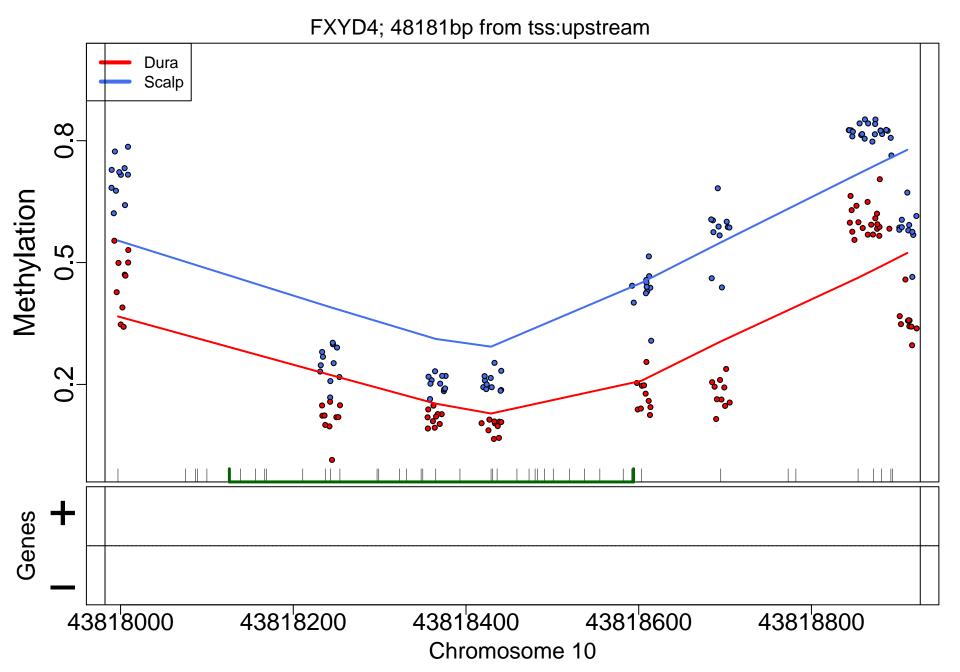


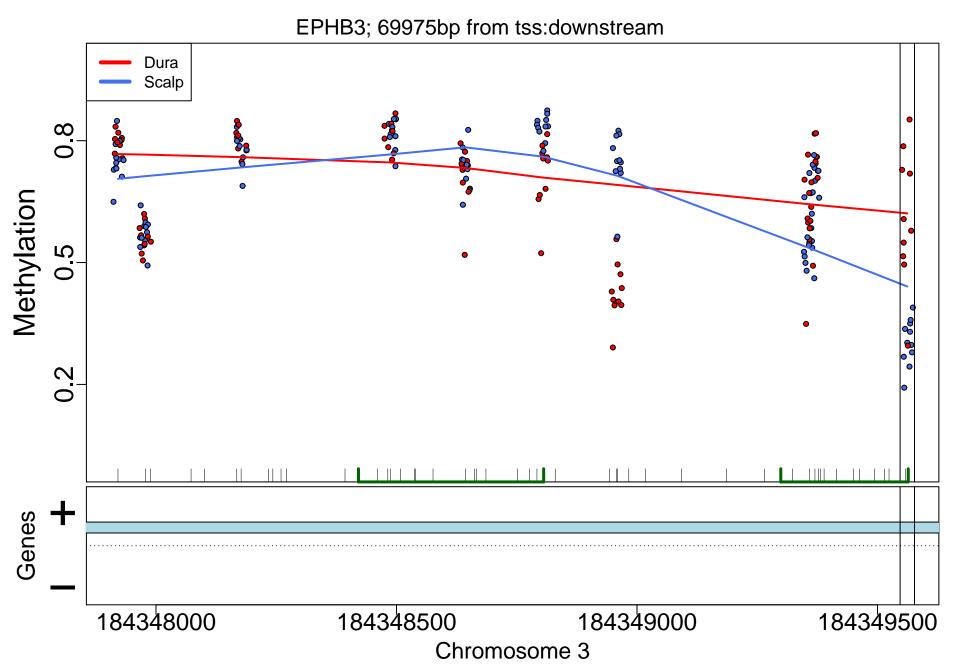


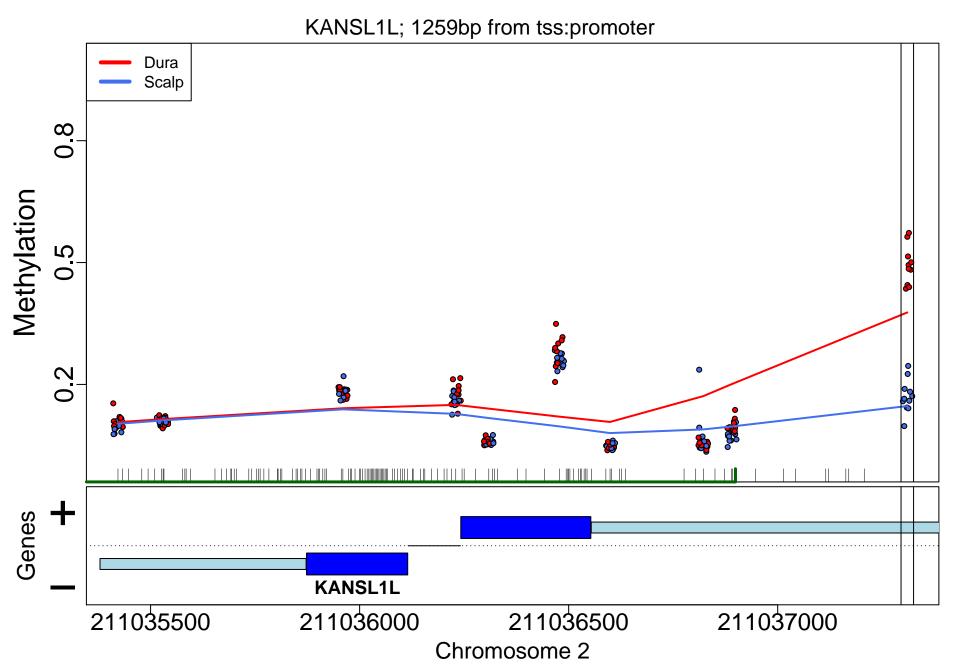


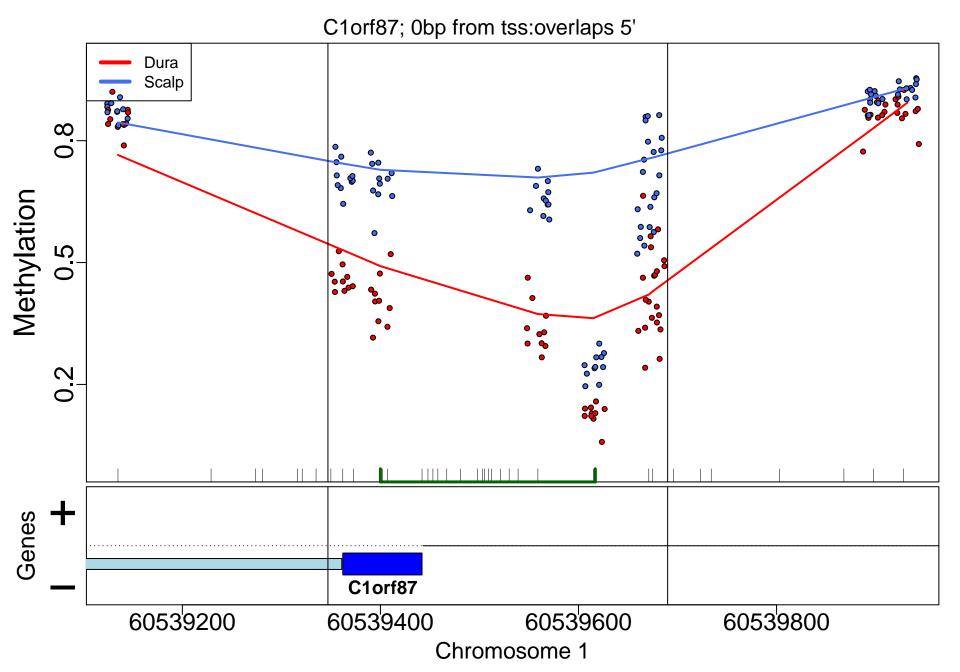


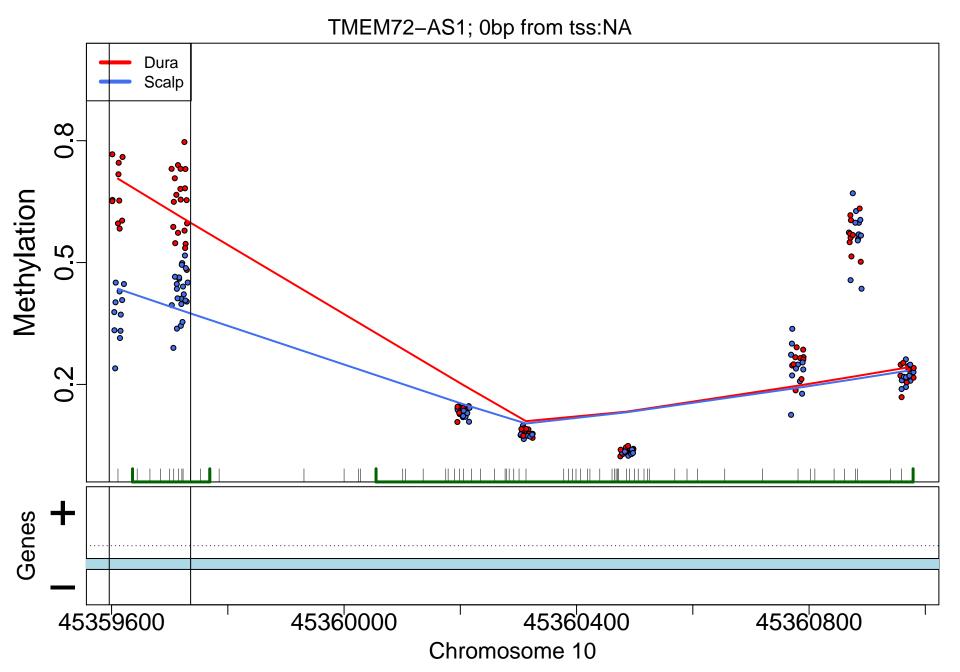
VGLL3; 1107bp from tss:inside intron Dura Scalp <u>ω</u>_ Methylation Genes VGLL3 87039000 87039500 87040000 Chromosome 3

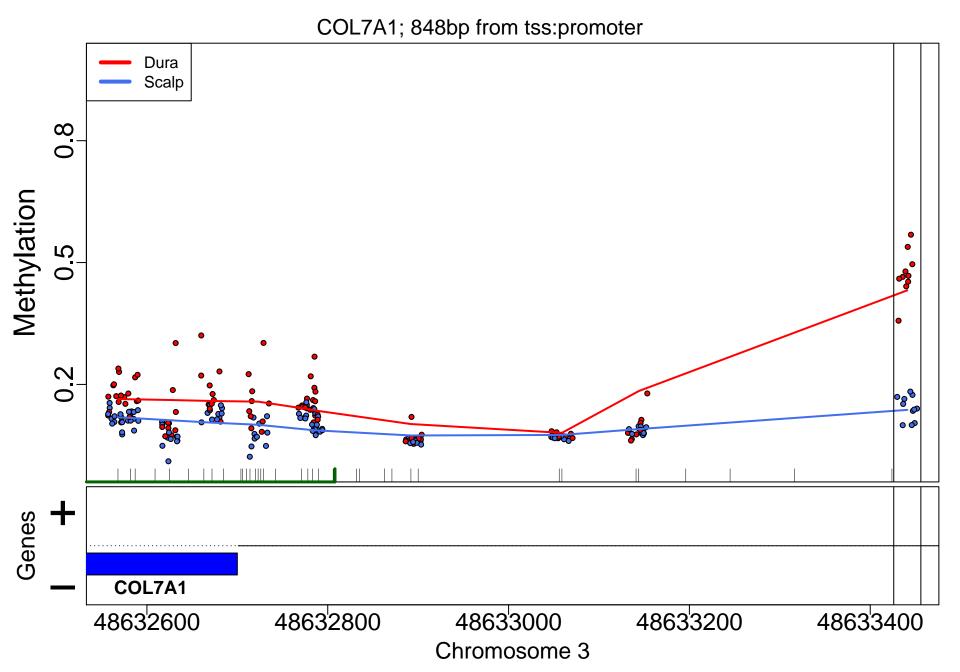


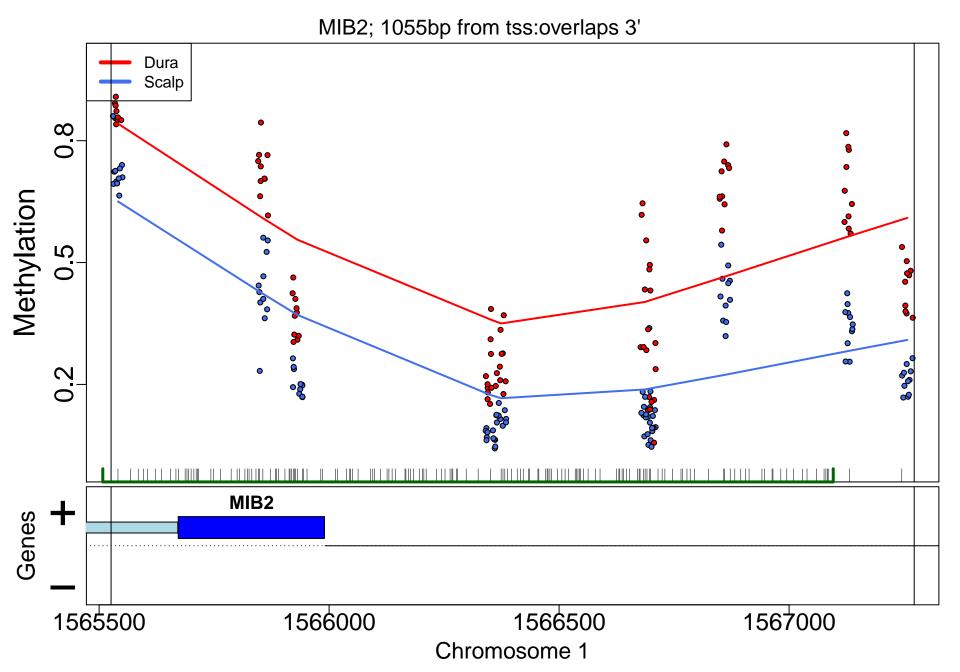


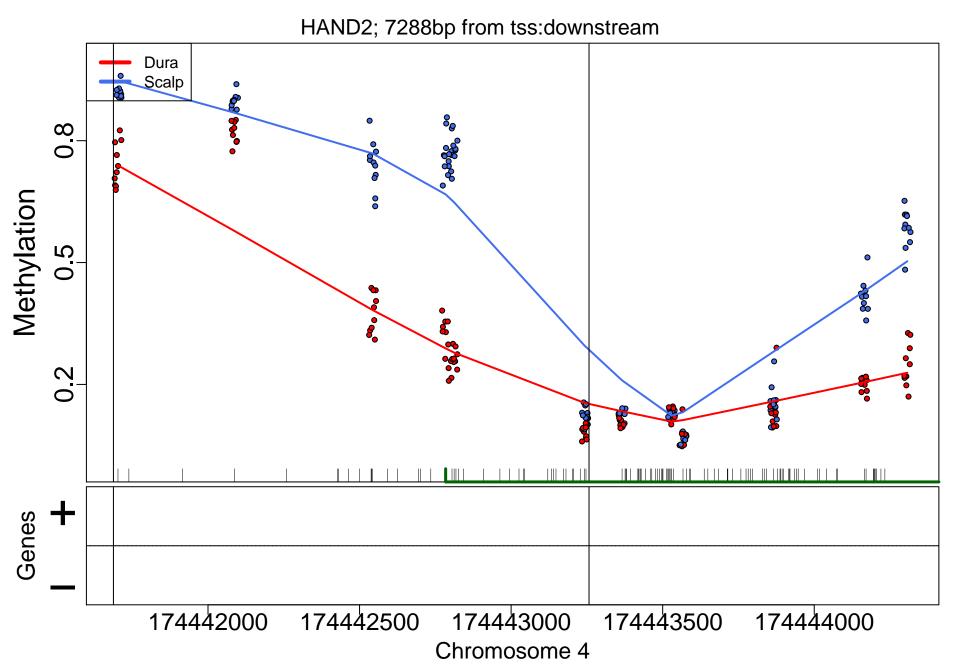


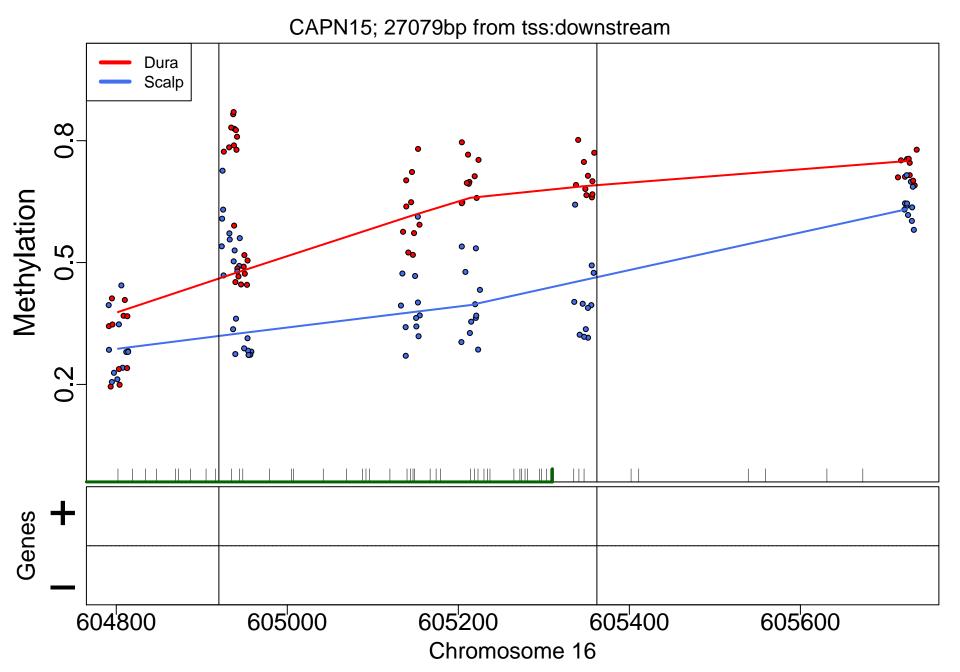


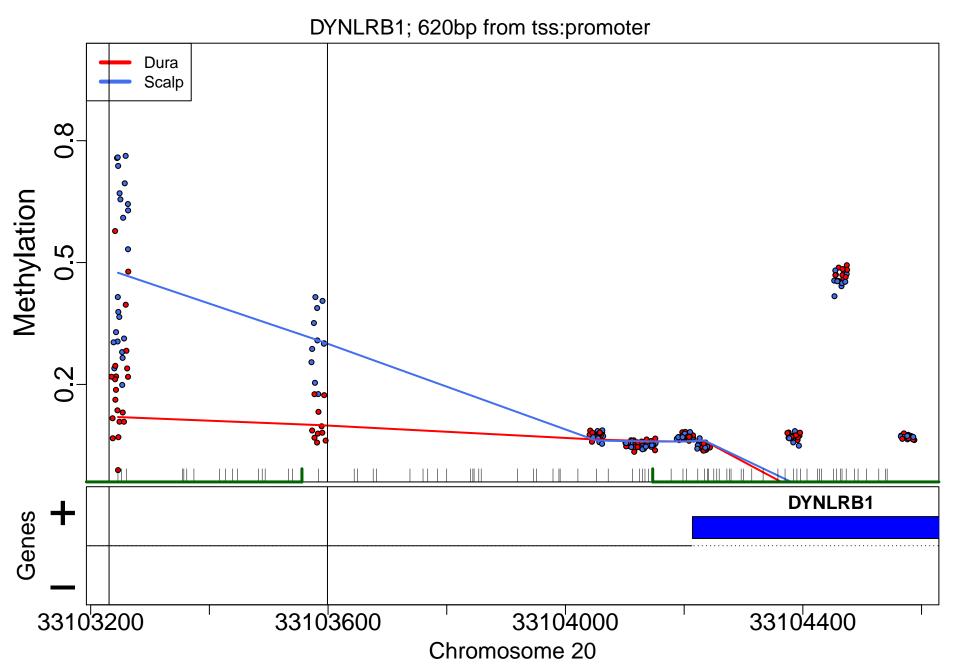


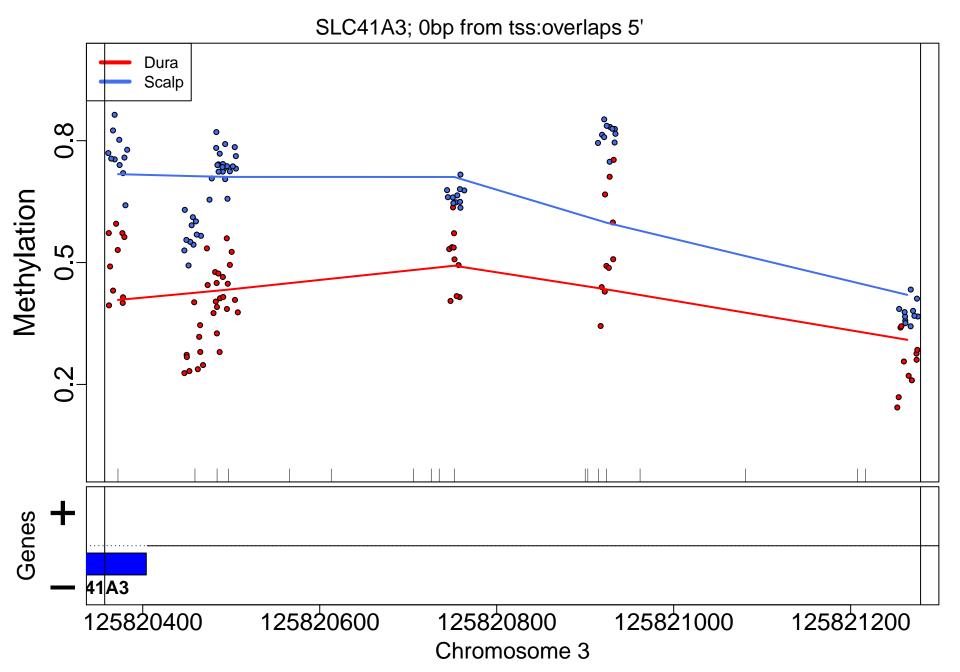


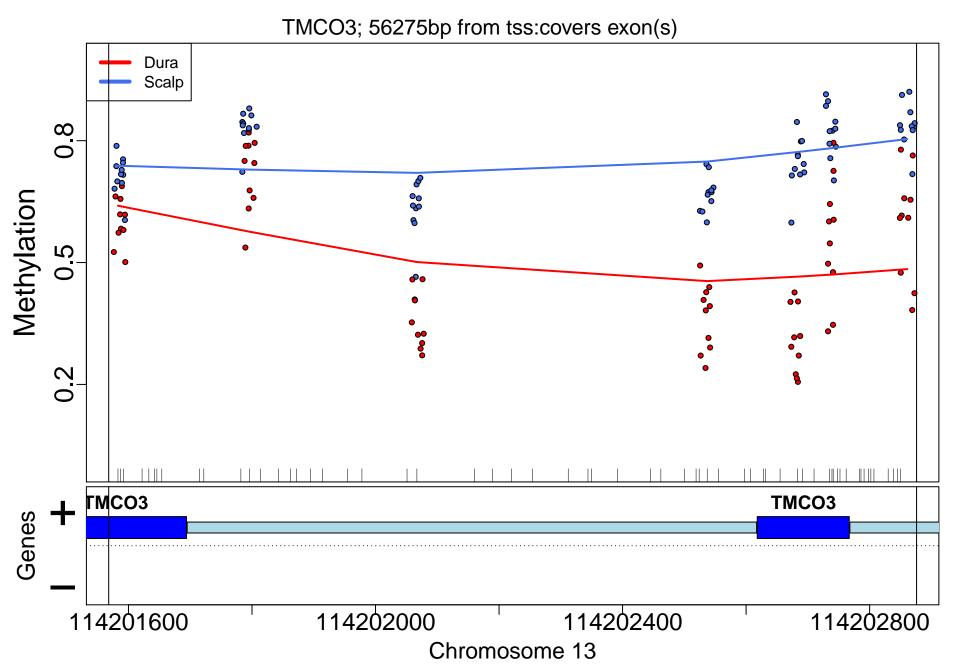


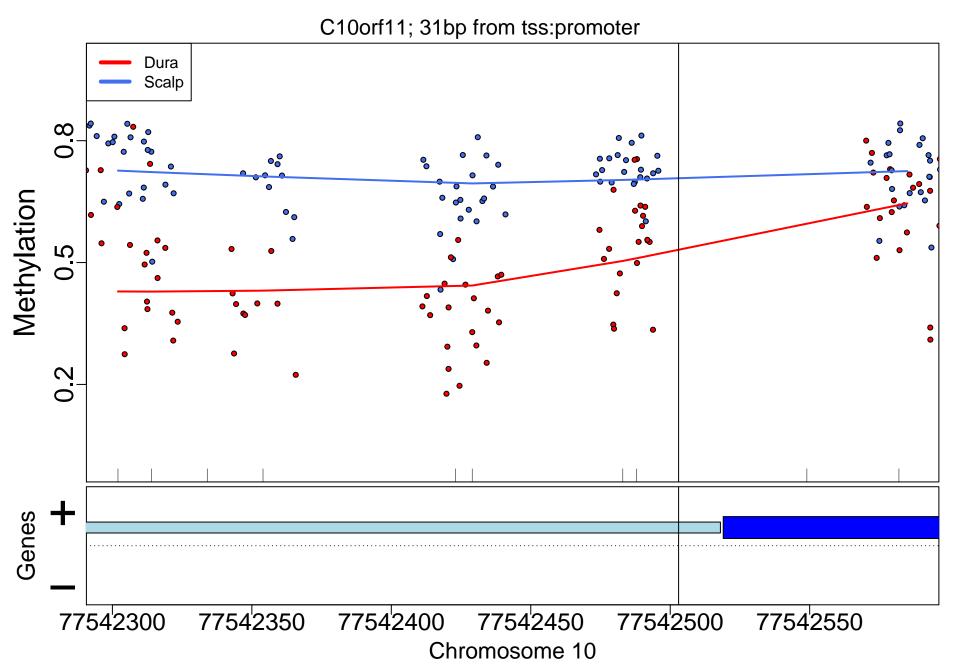


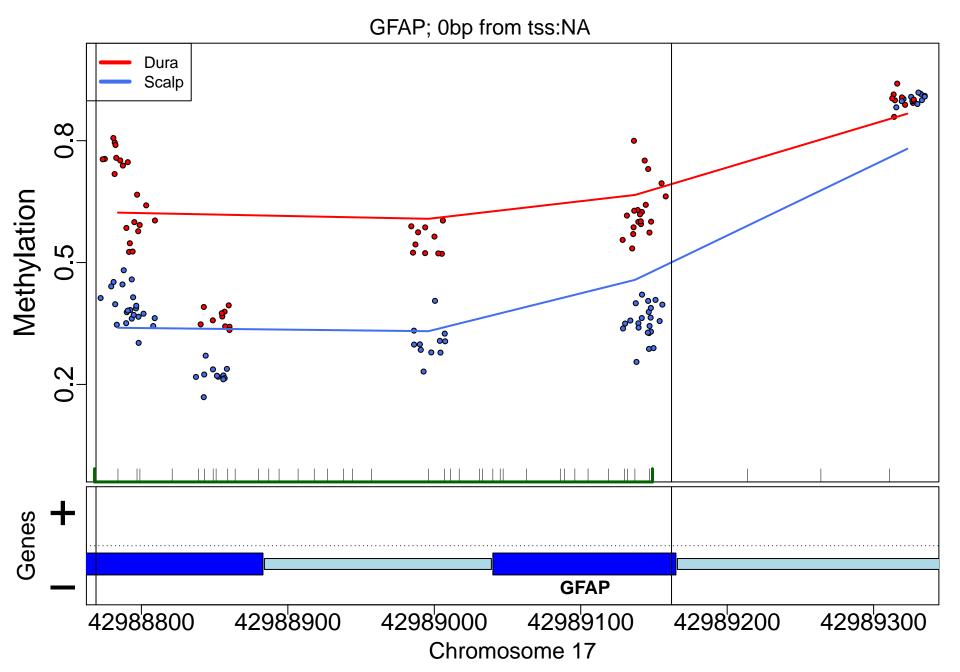


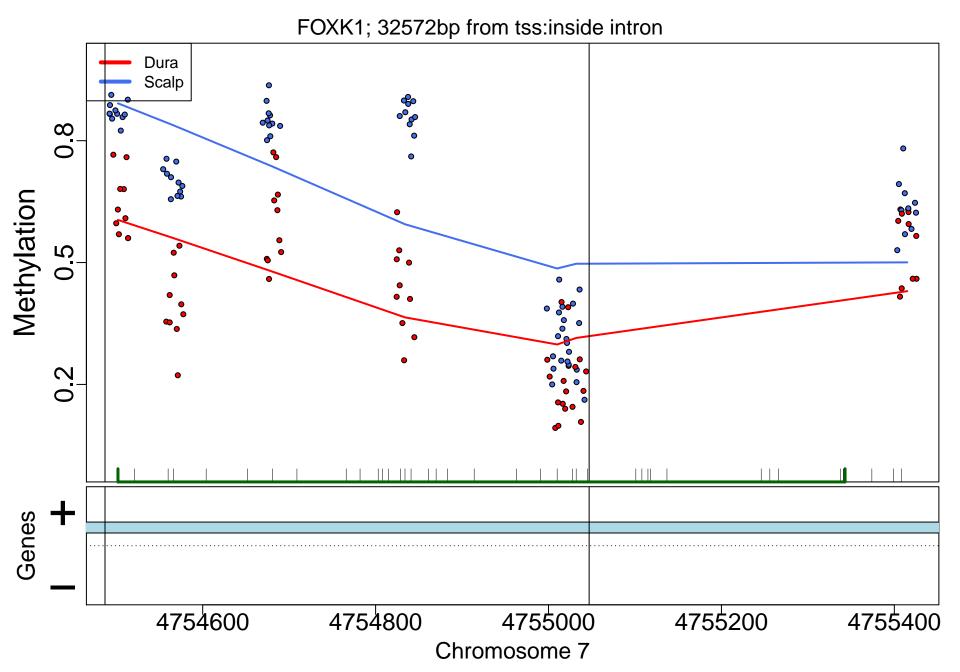


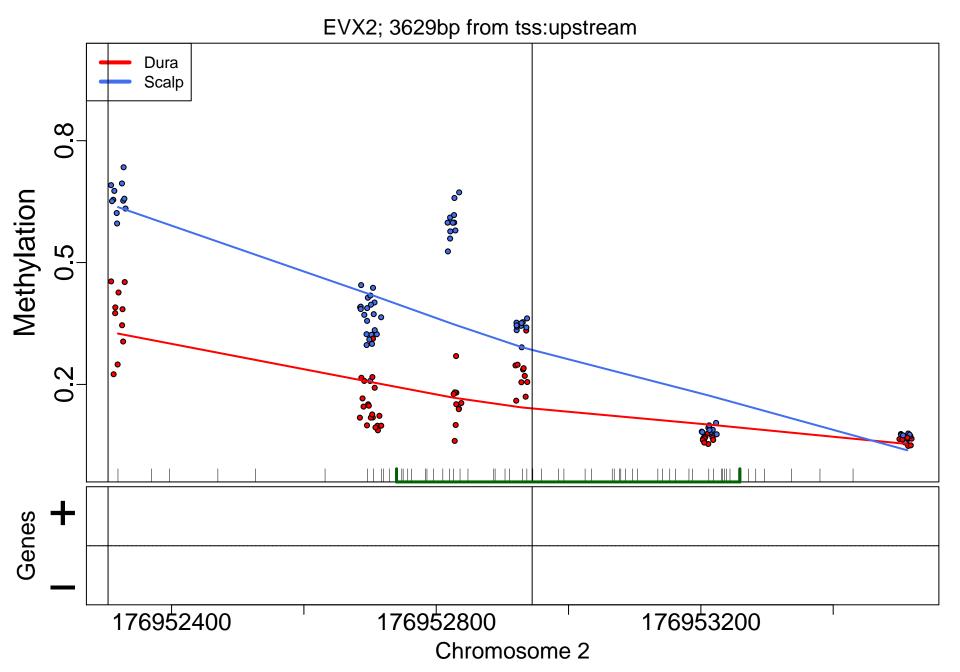


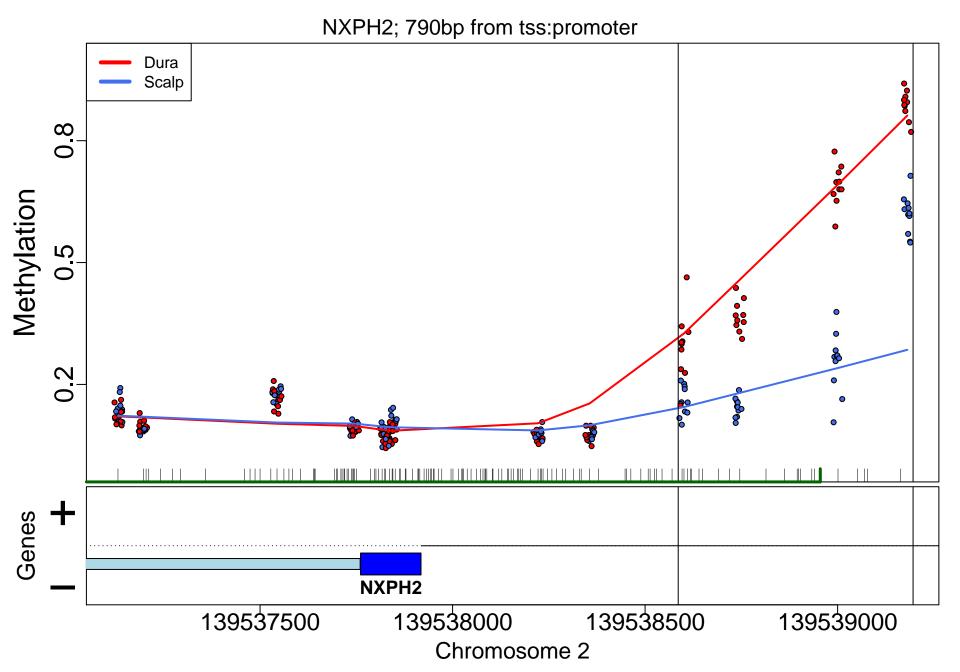


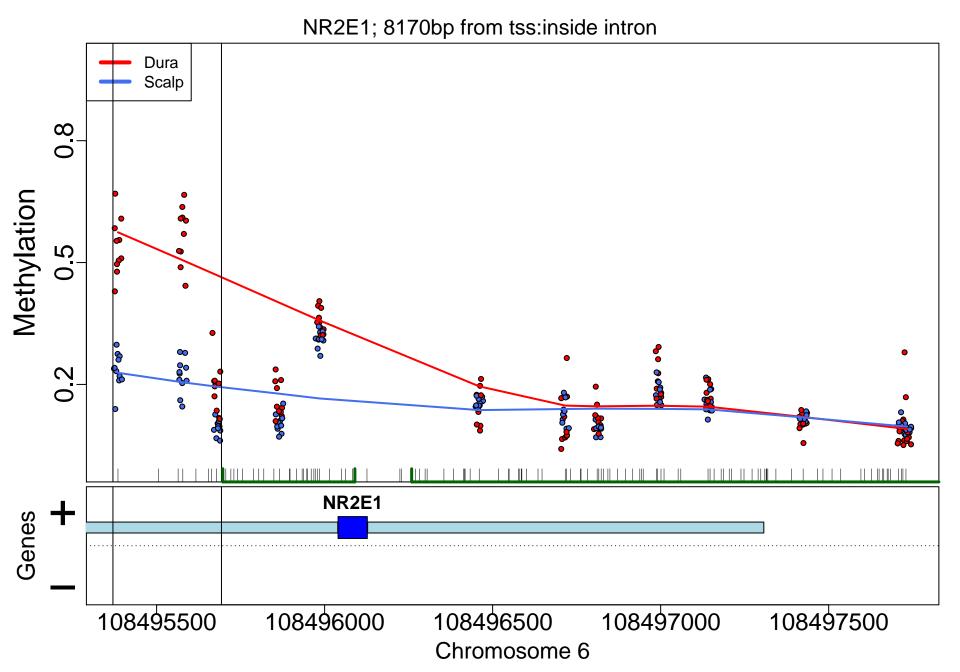


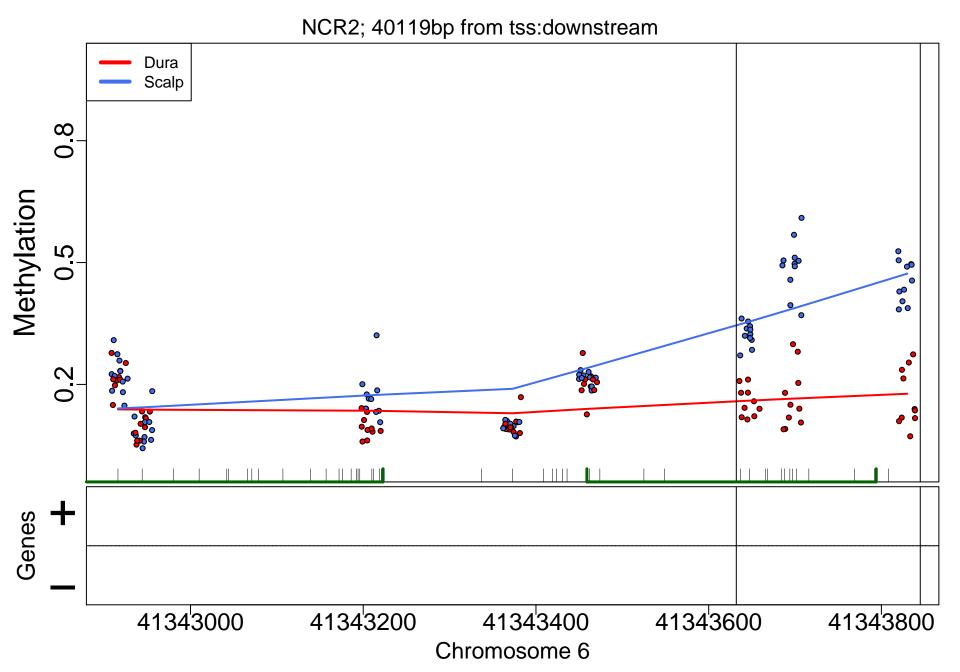




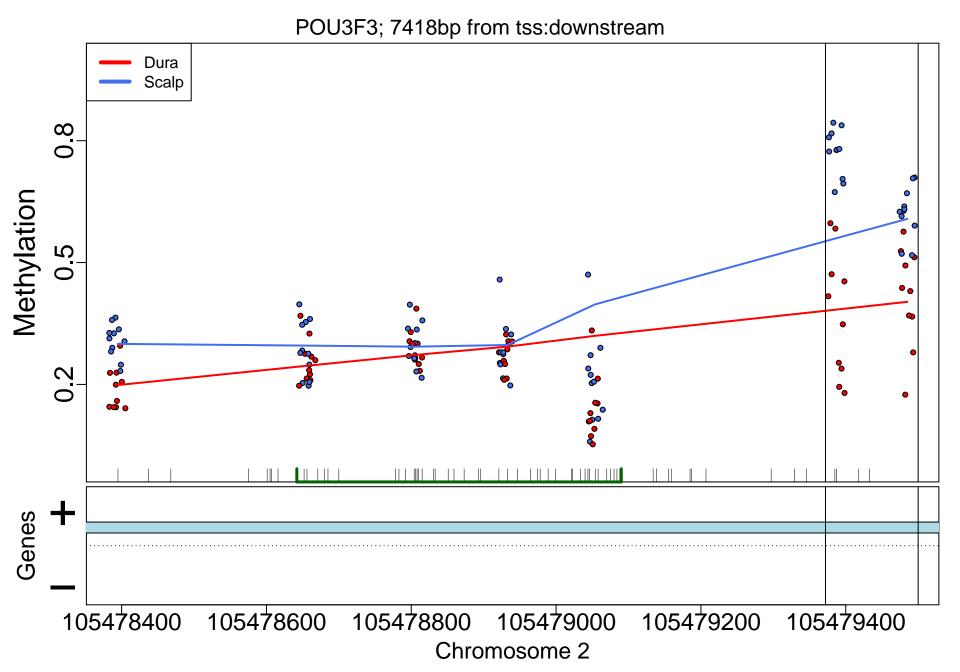


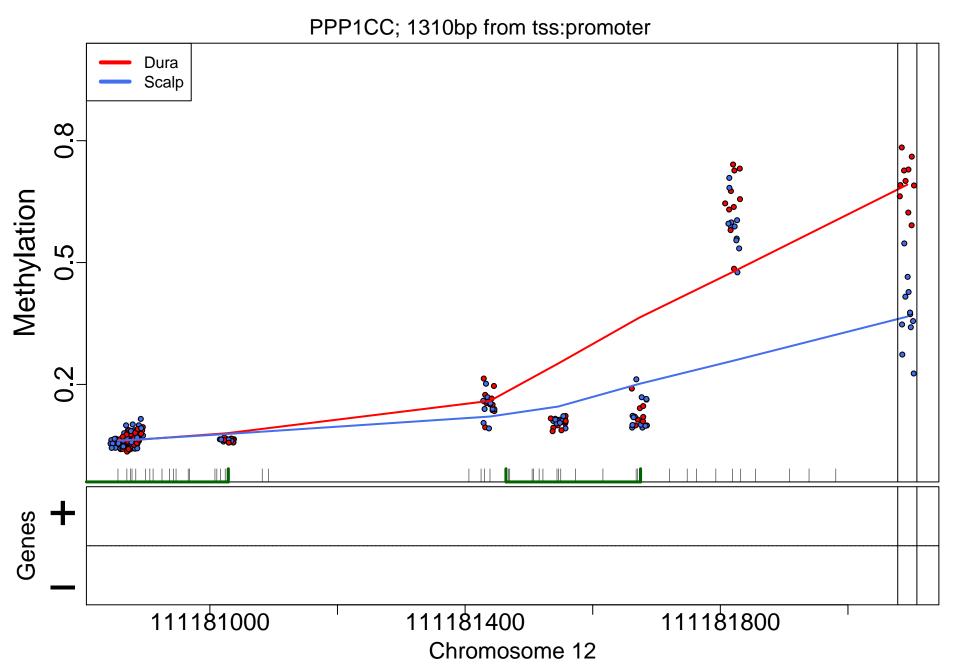


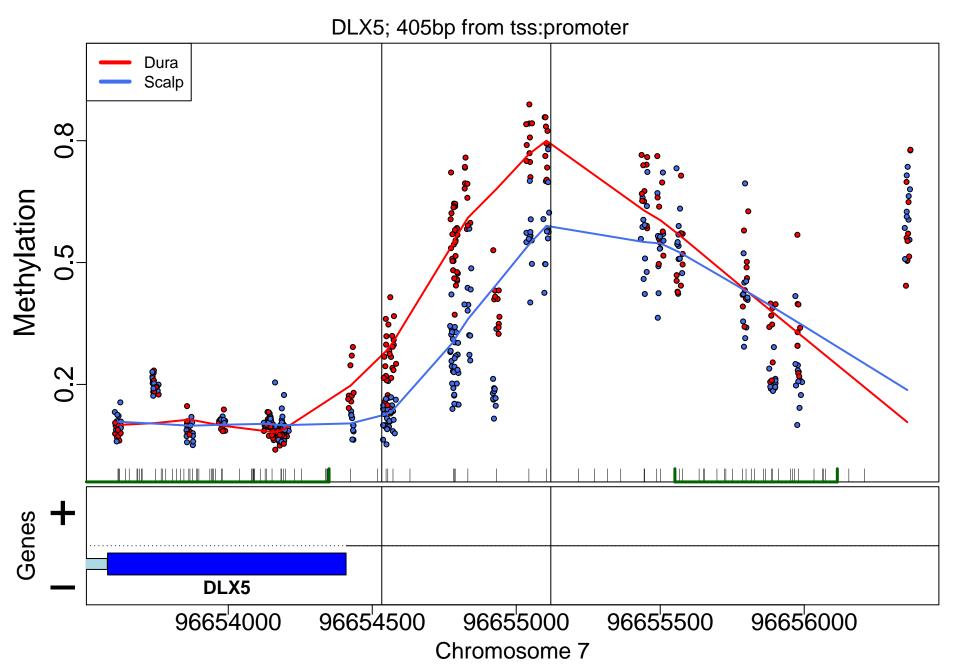


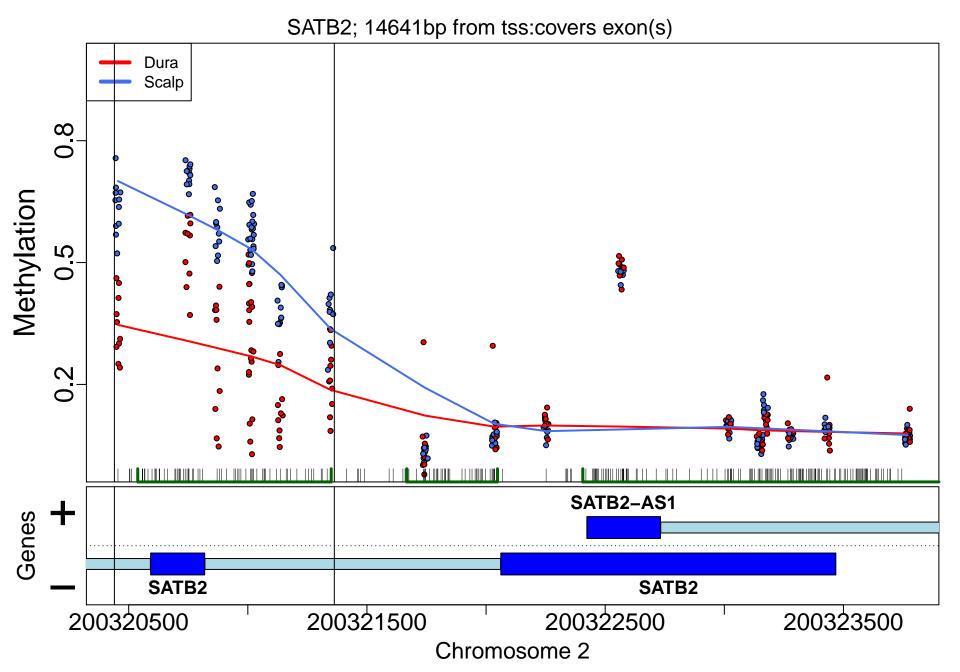


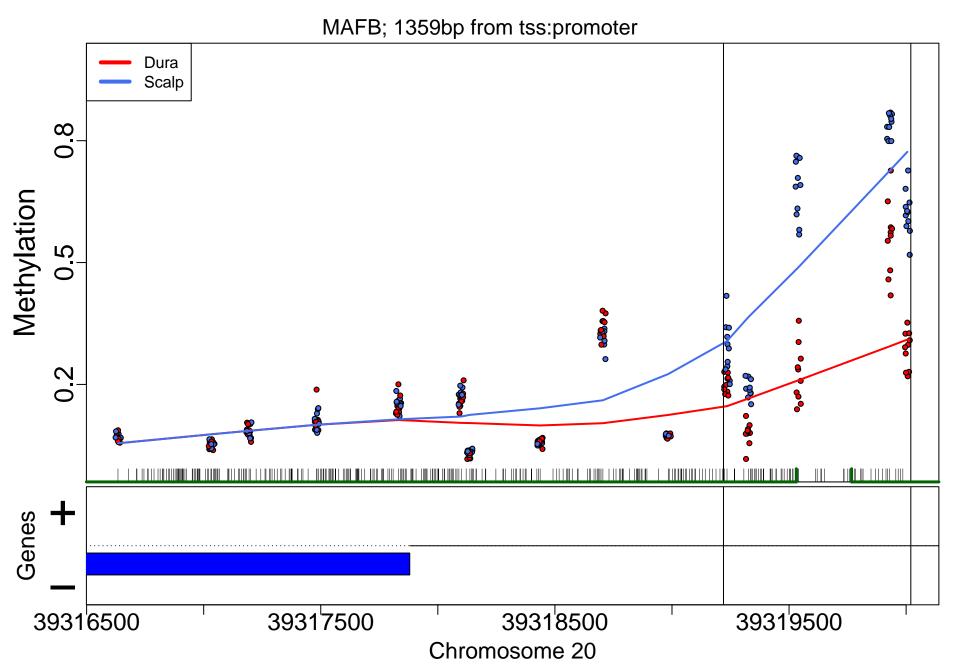
DCAF4; 614bp from tss:promoter Dura Scalp <u>ω</u>_ Methylation 0.7 DCAF4 DCAF4 Genes 73392200 73392600 73393000 73393400 Chromosome 14

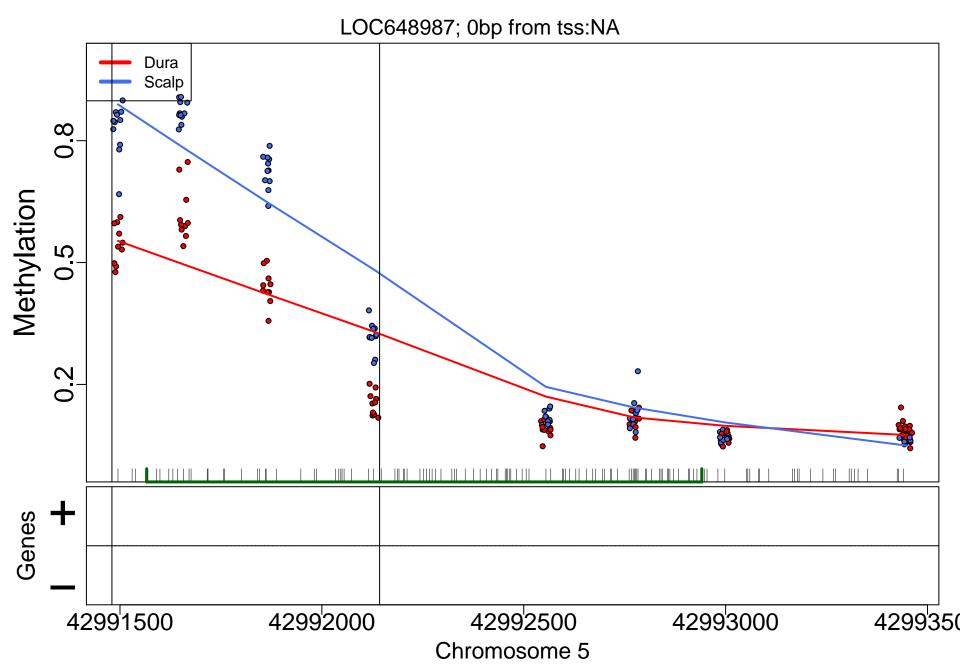


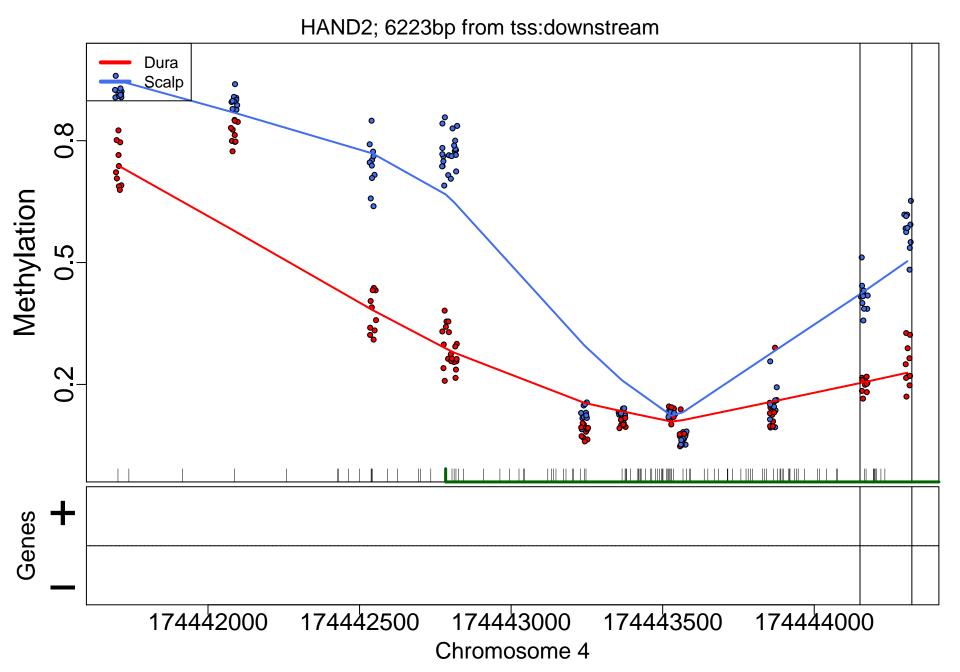


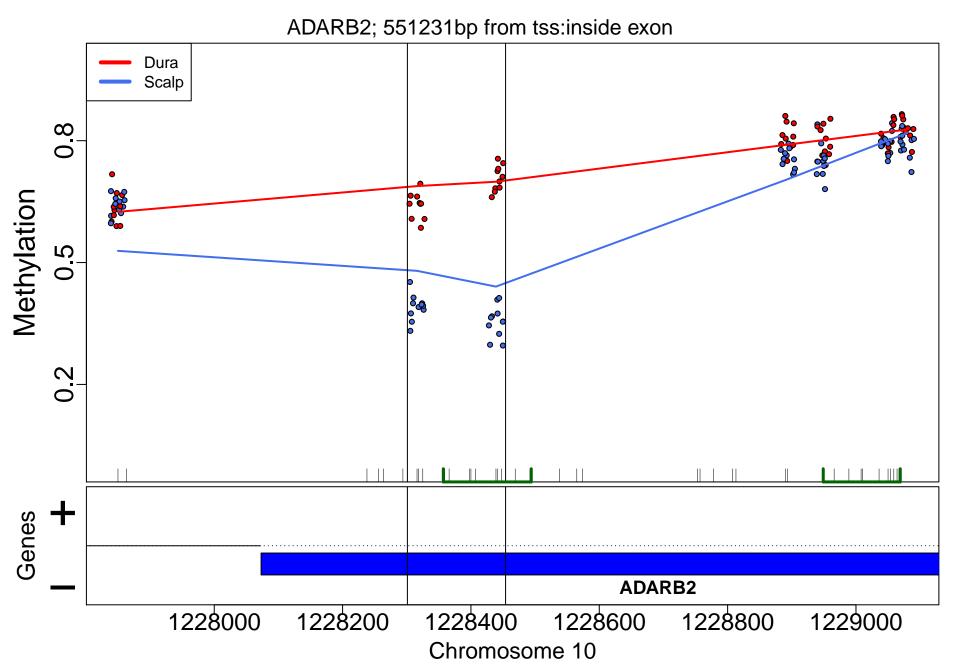


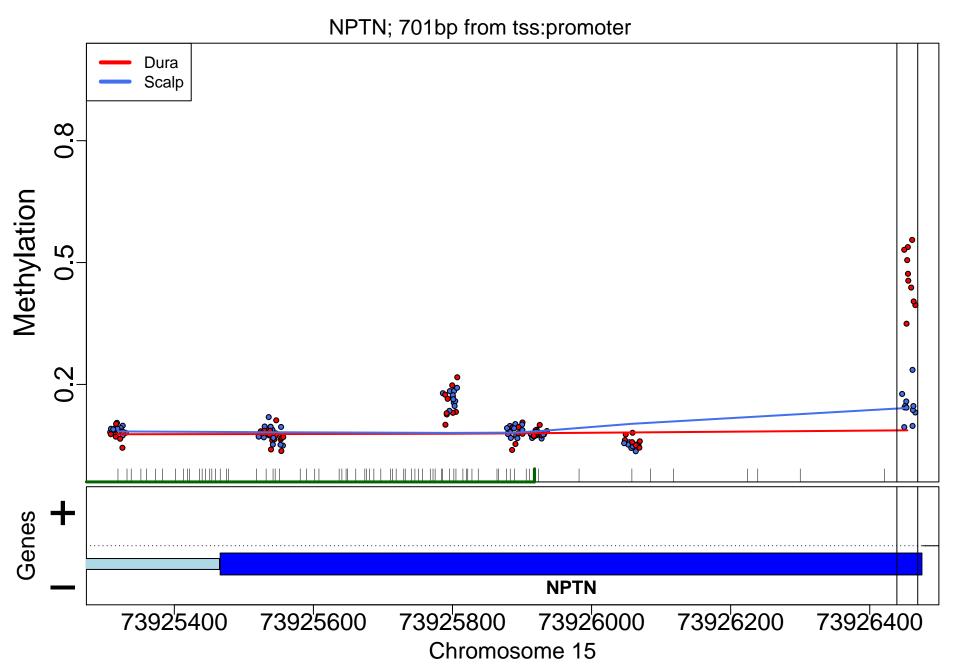


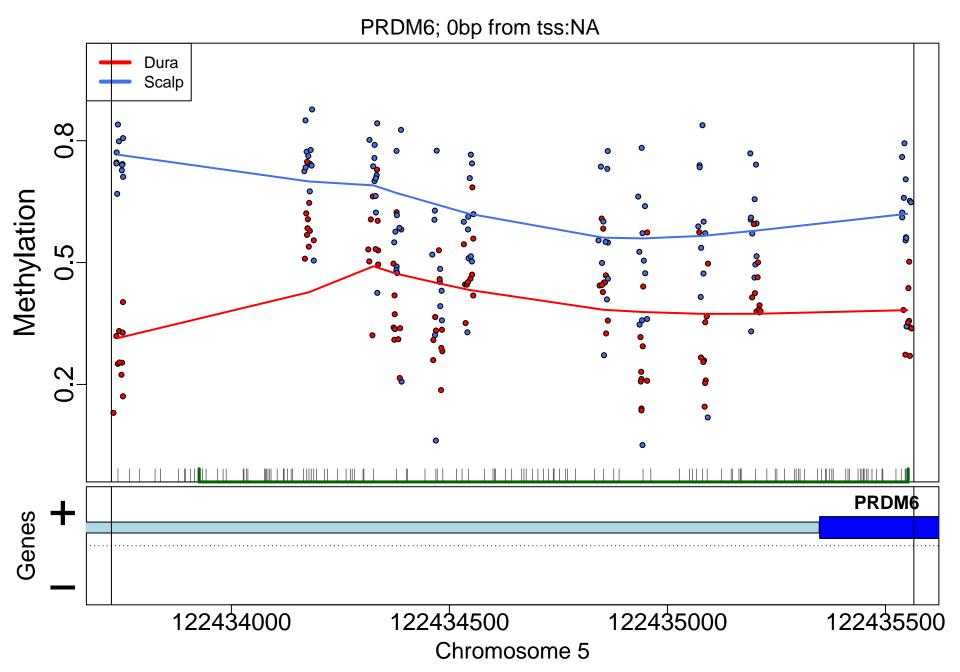


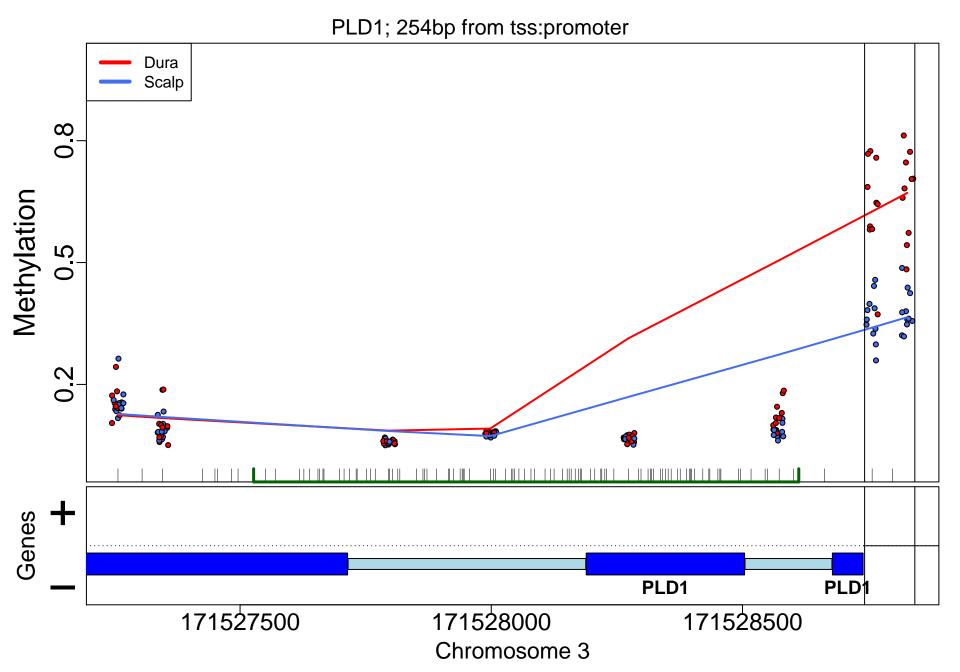


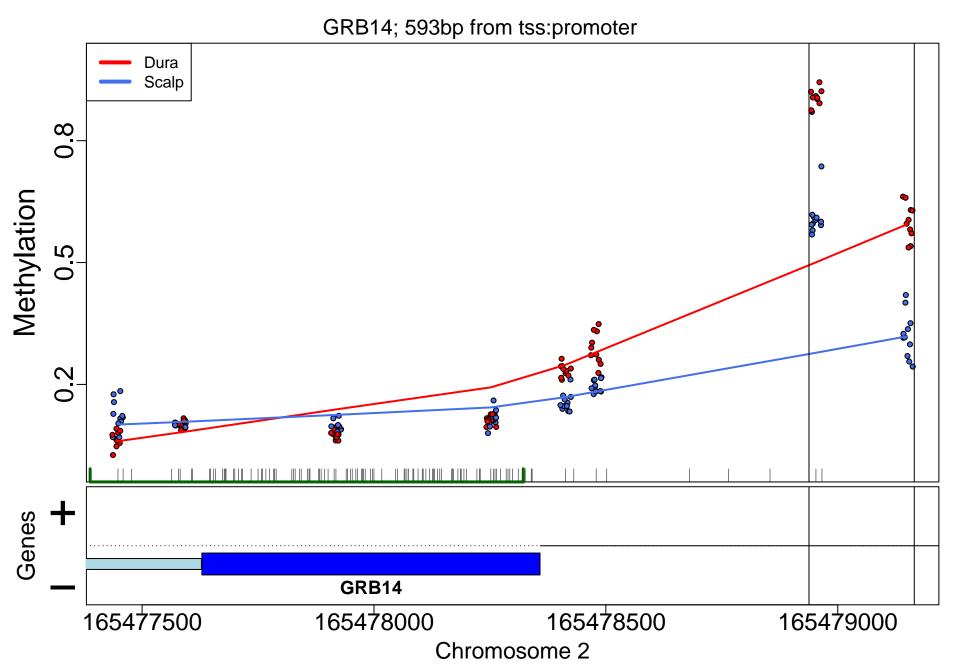


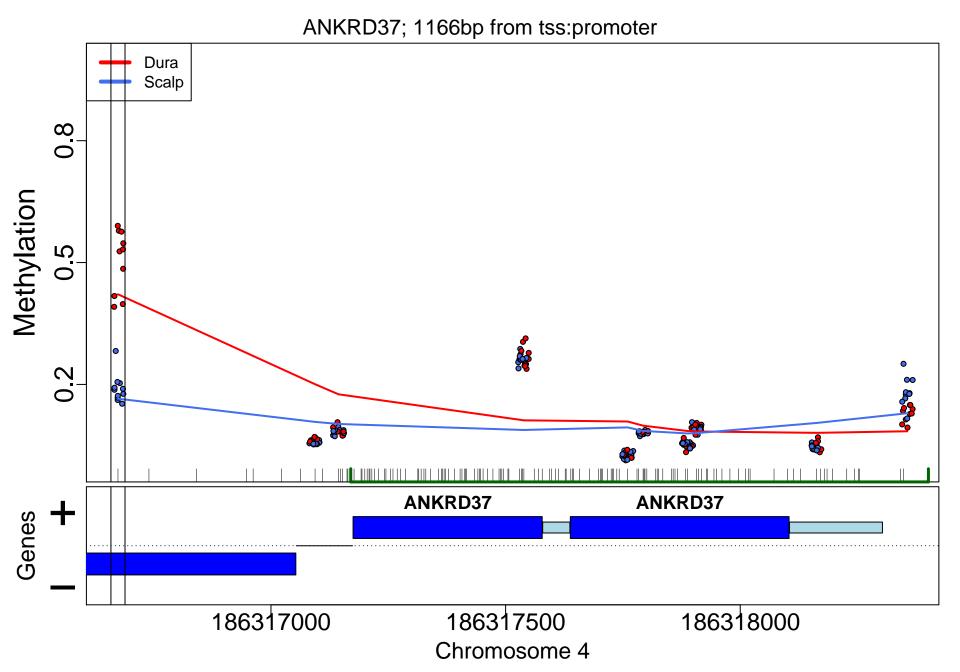


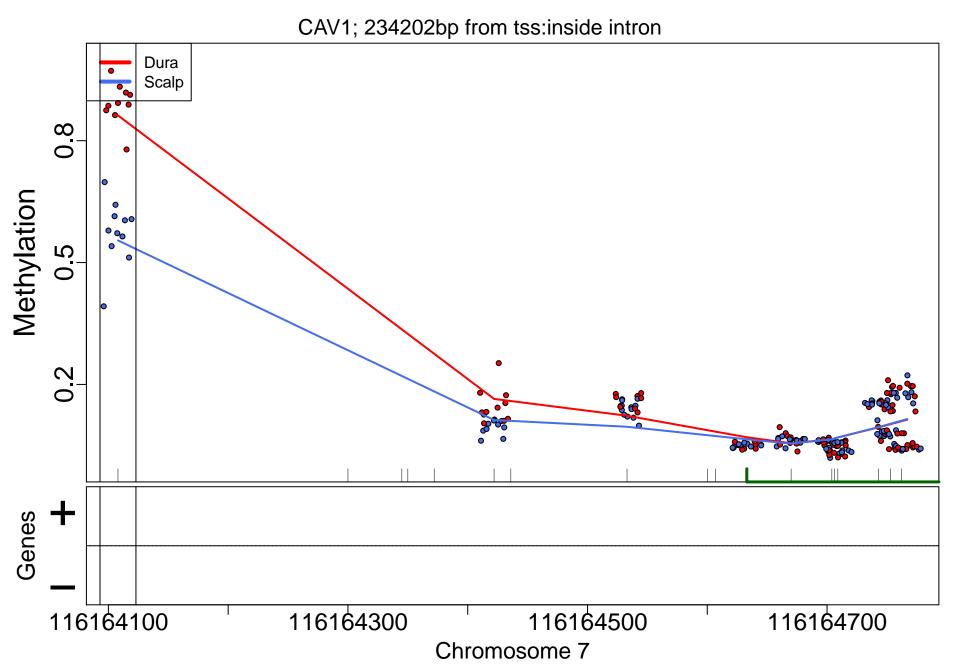


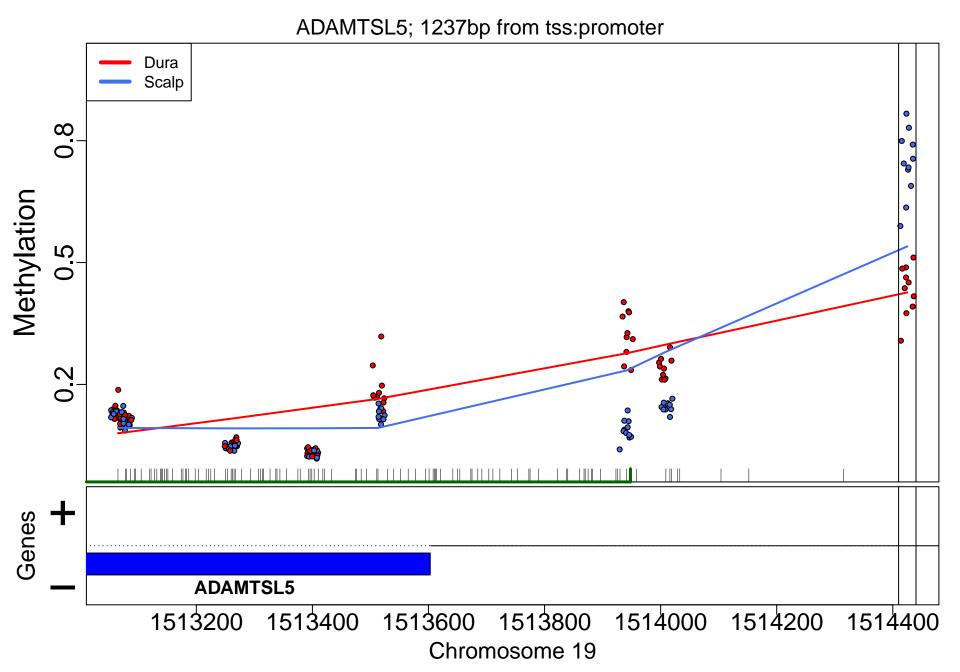


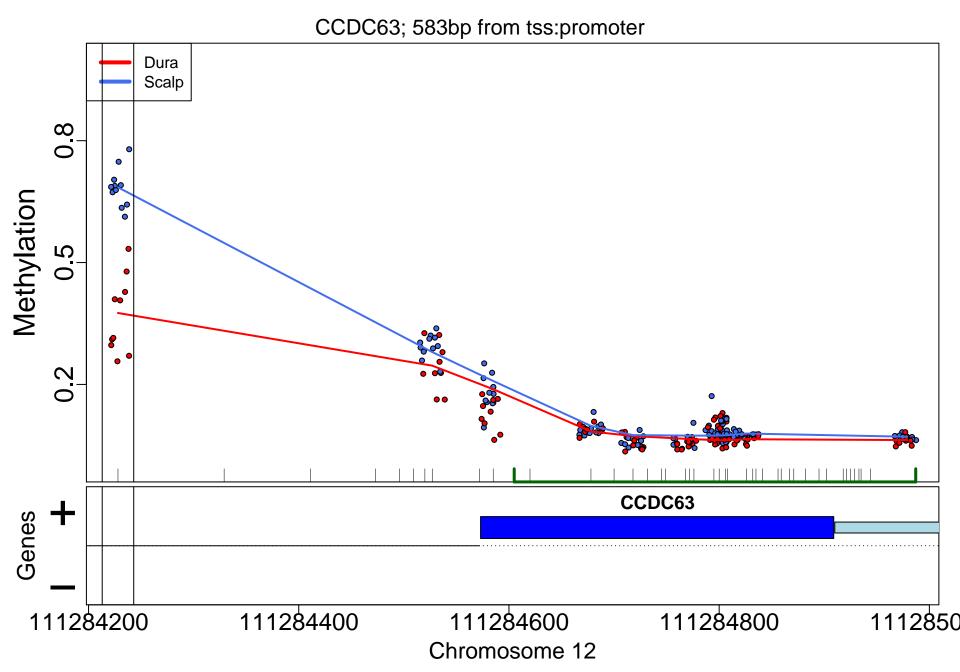




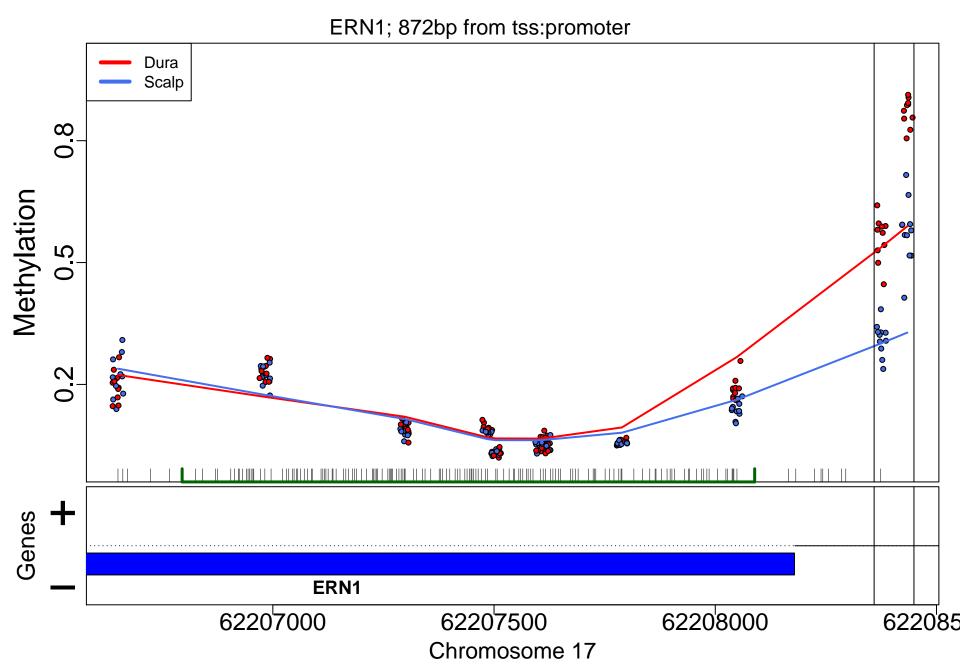


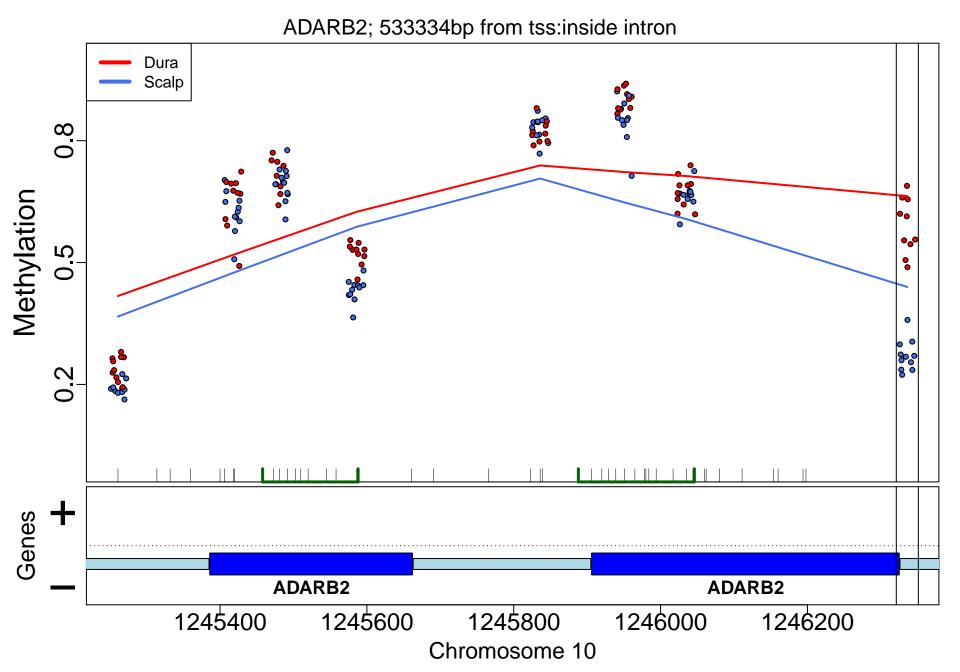


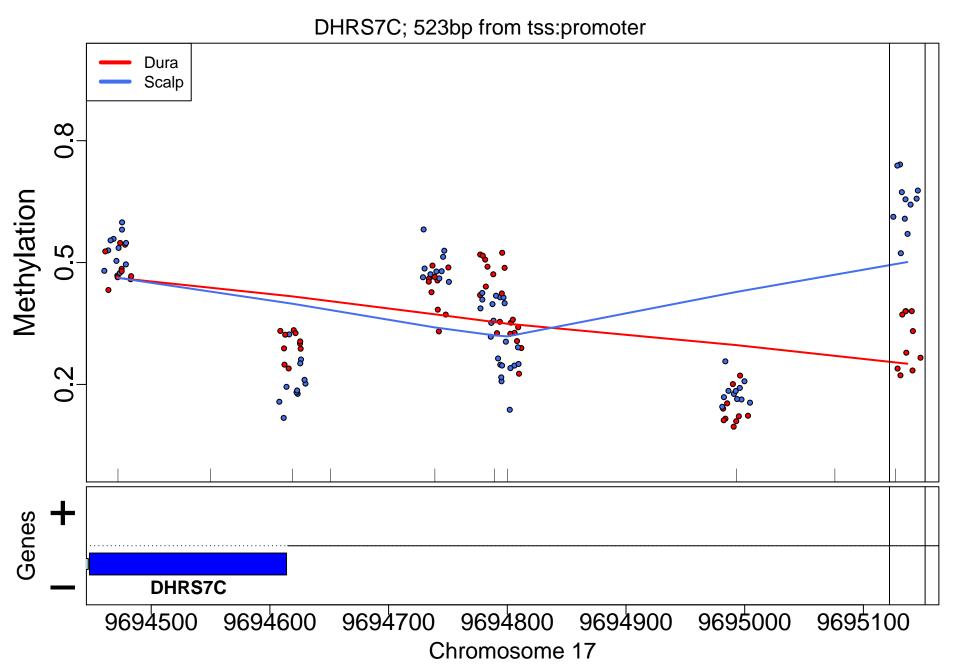




ADPRHL1; 129bp from tss:inside exon Dura Scalp <u>ω</u>_ Methylation 0.5 0.2 Genes ADPRHL1 114107600 114107800 114108000 114108200 114108400 Chromosome 13



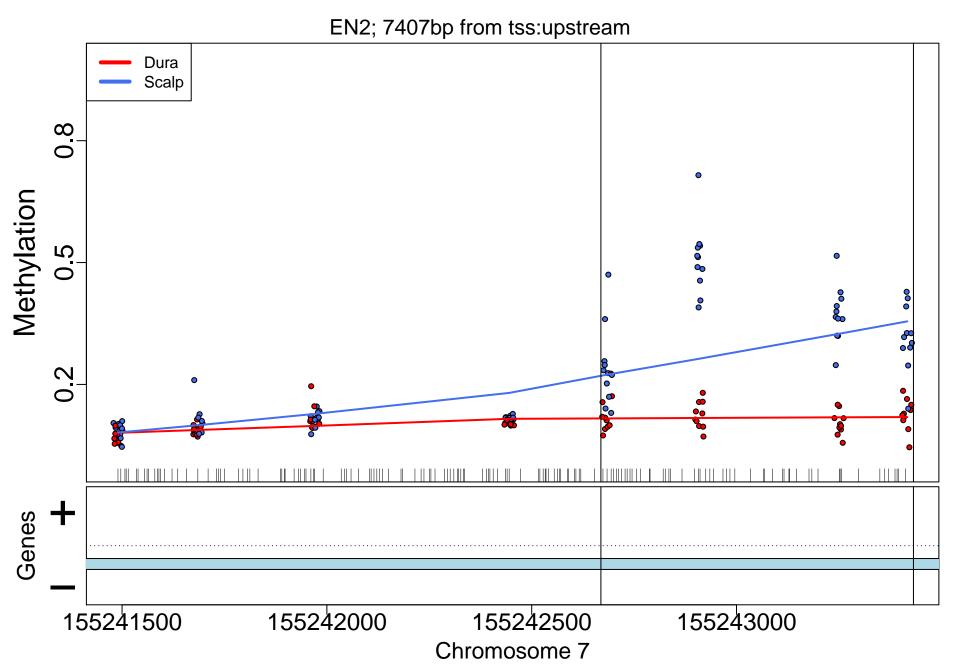


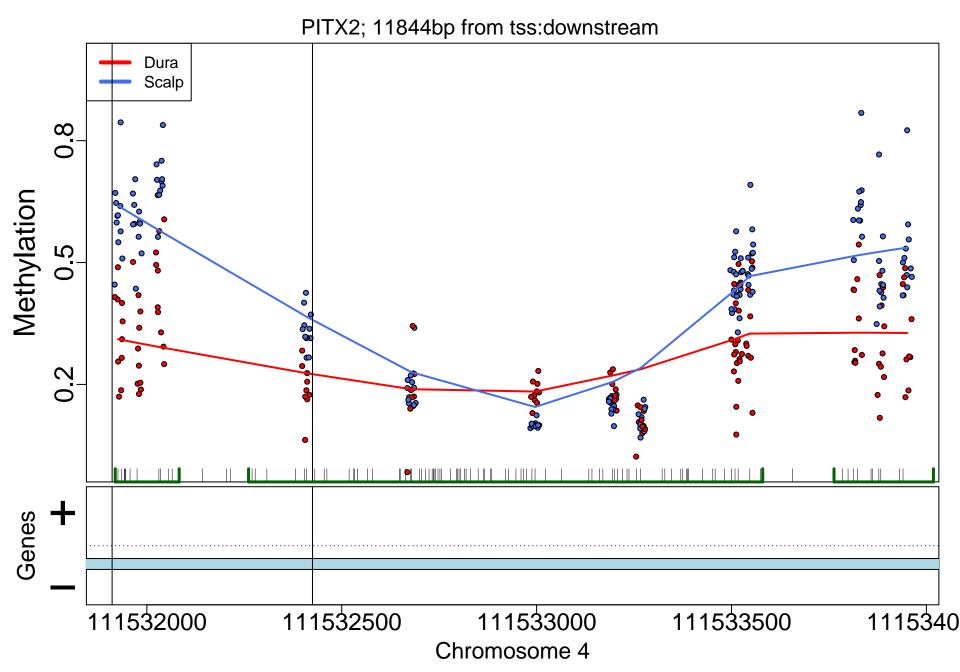


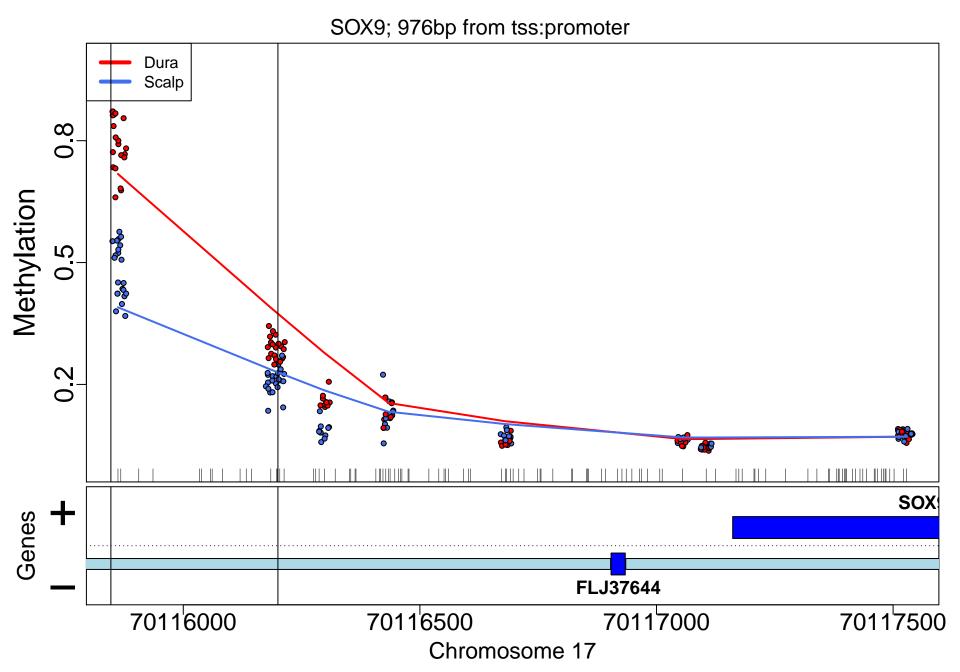
DLX6-AS1; 0bp from tss:NA Dura Scalp 80. Methylation 0.2 Genes 96626000 96625500 96626500 96627000 96627500

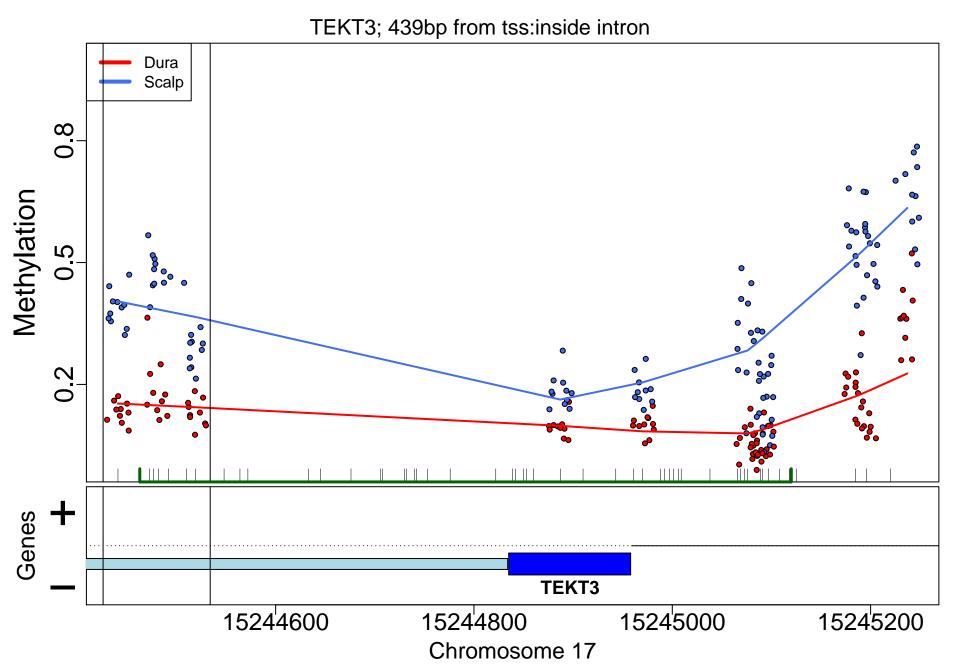
Chromosome 7

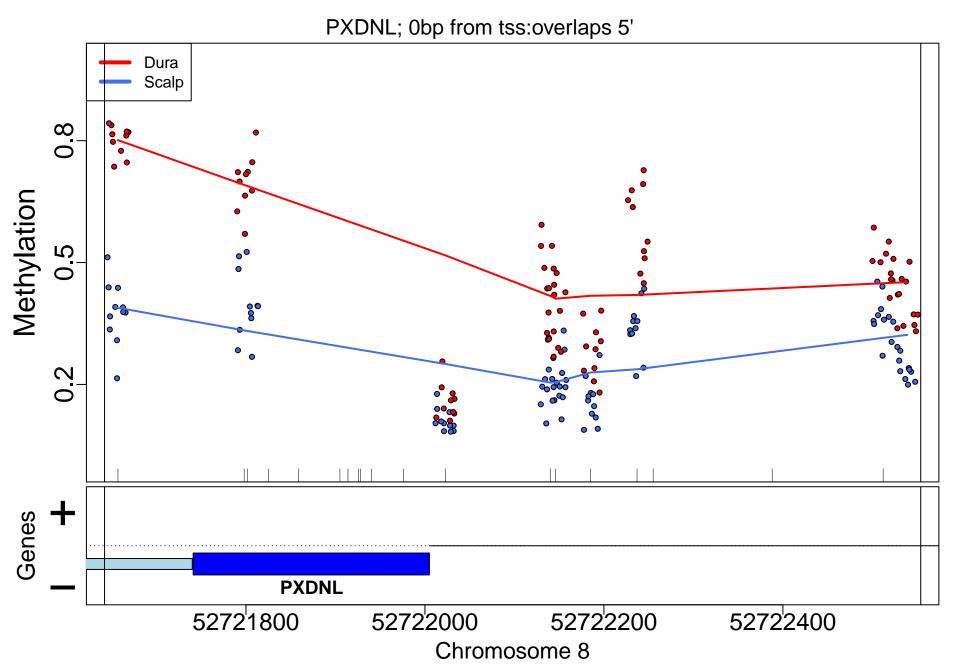
STX18-AS1; 0bp from tss:NA Dura Scalp <u>∞</u>_ Methylation 0.5 0.2 Genes 4763300 4763700 4763400 4763500 4763600 Chromosome 4

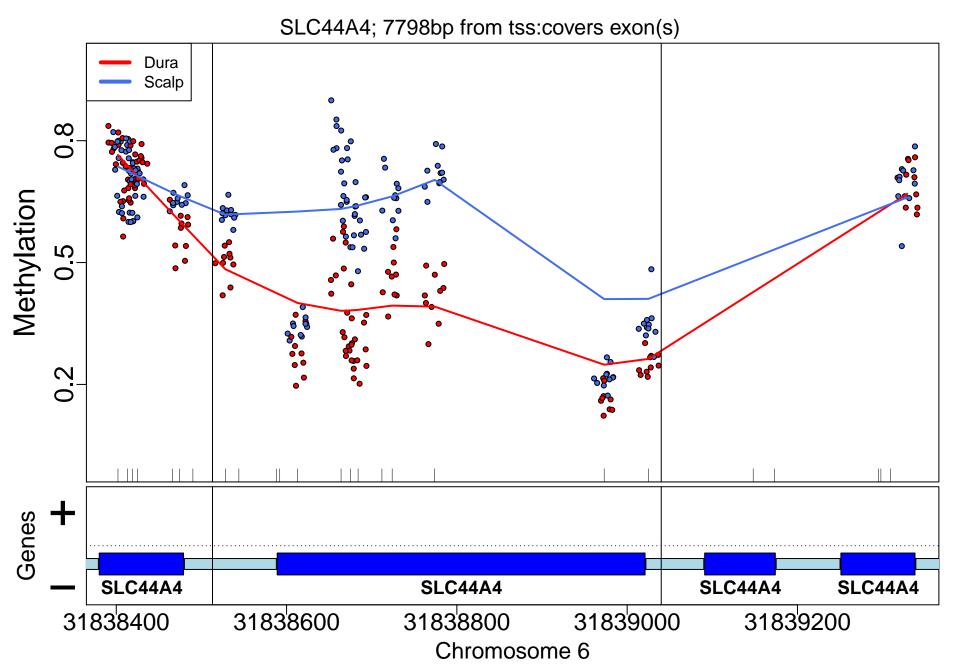


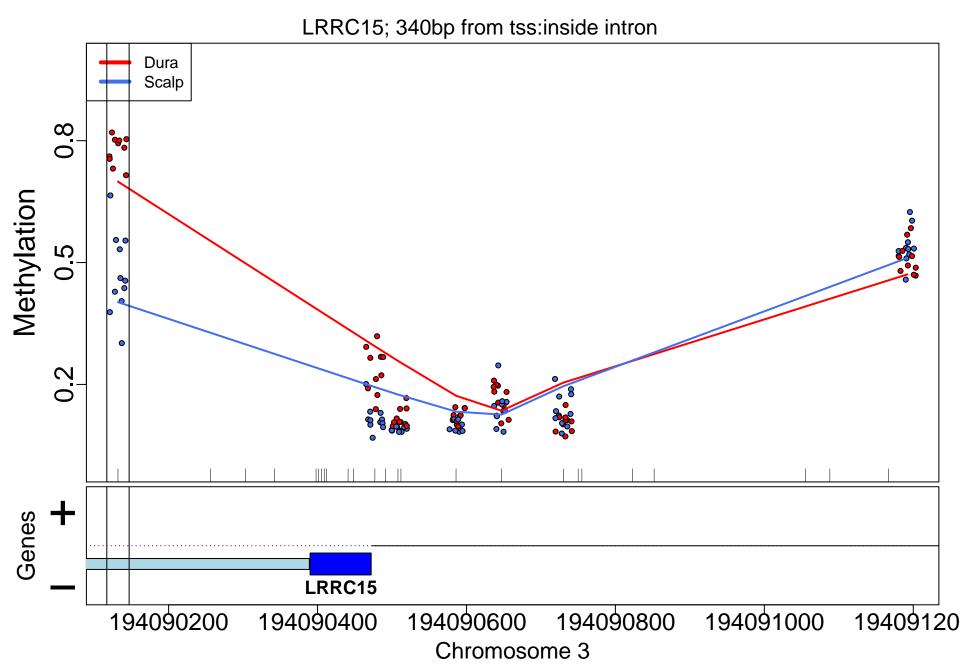


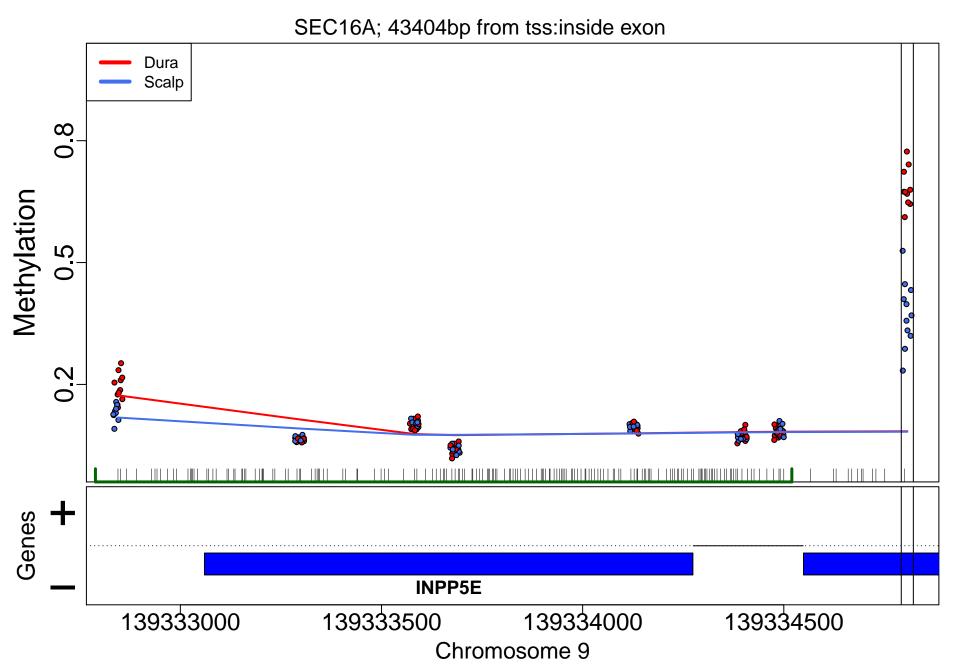




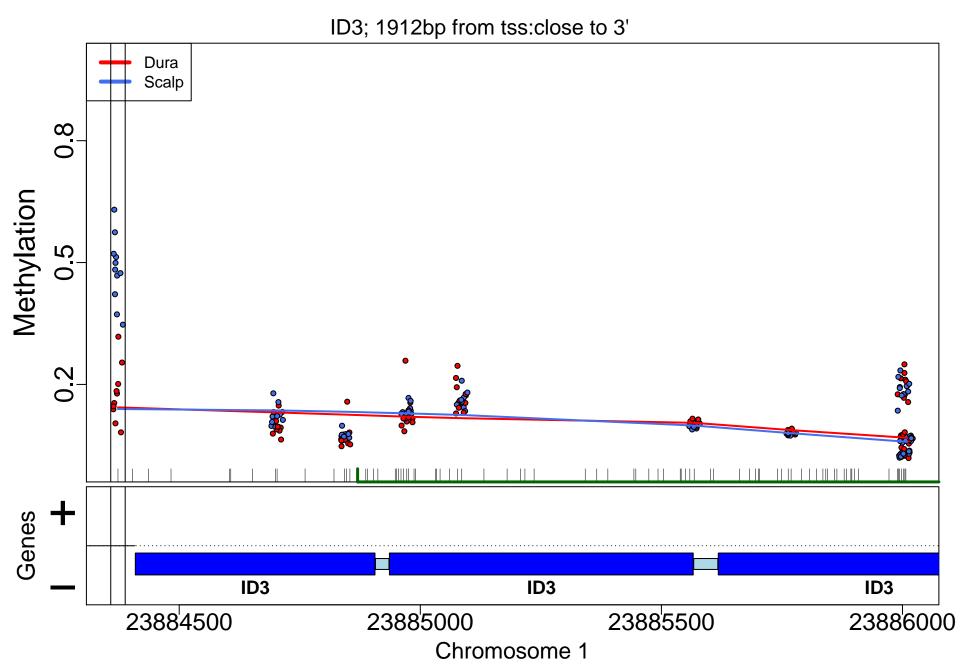


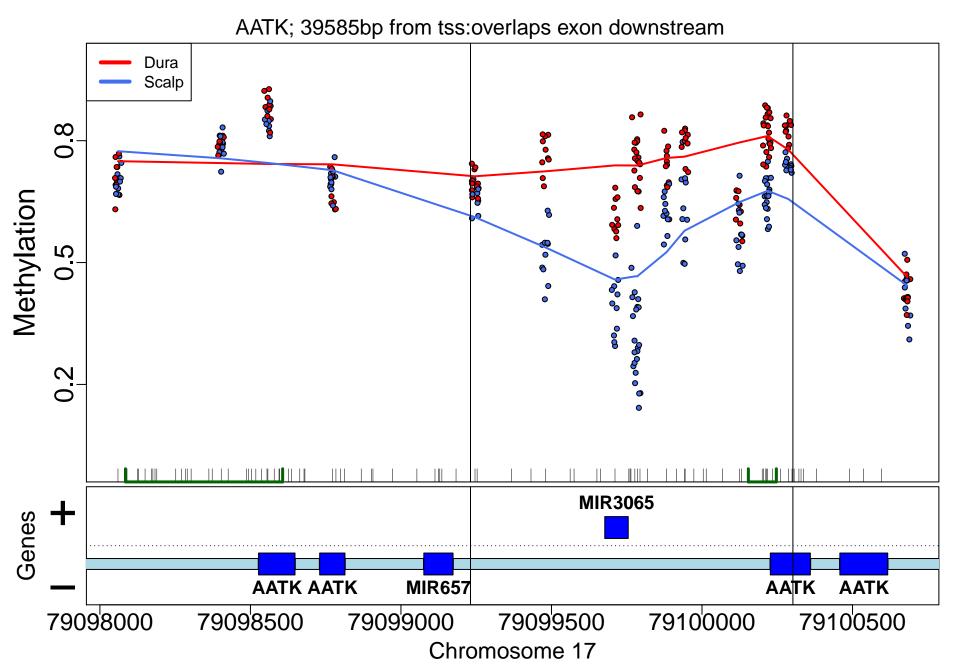


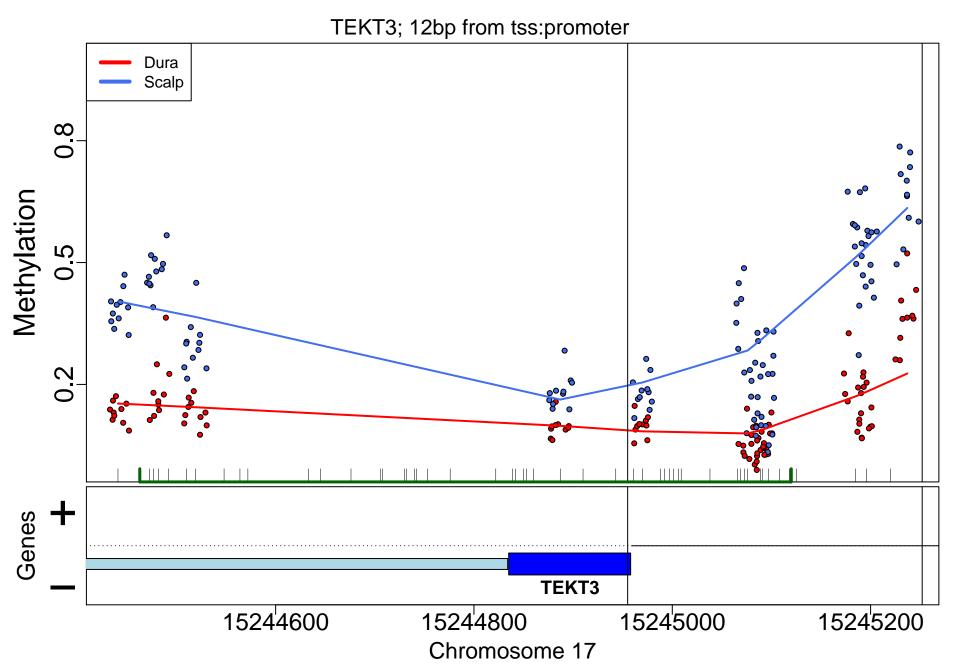


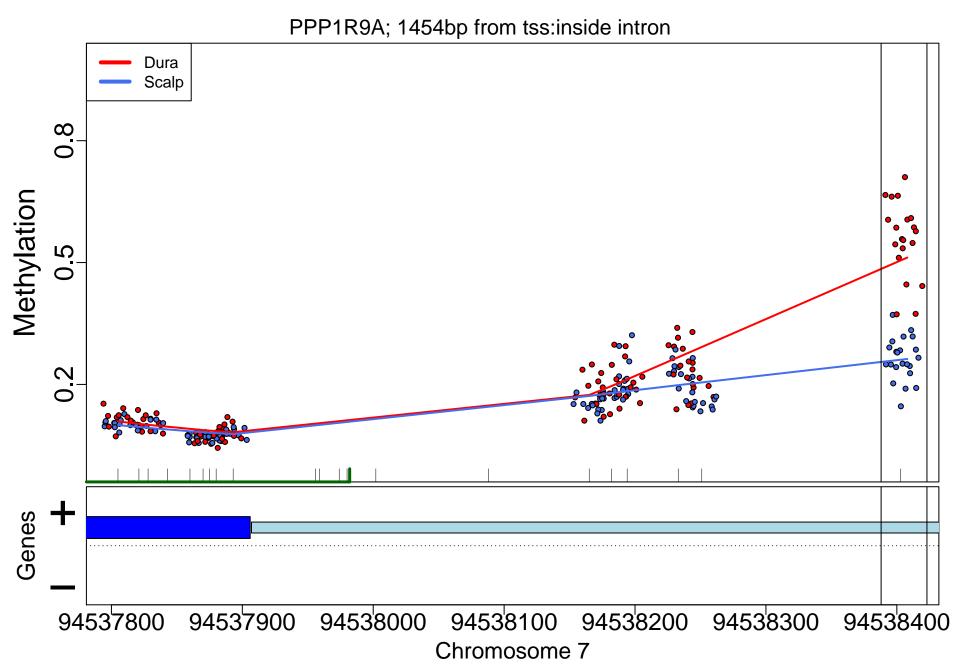


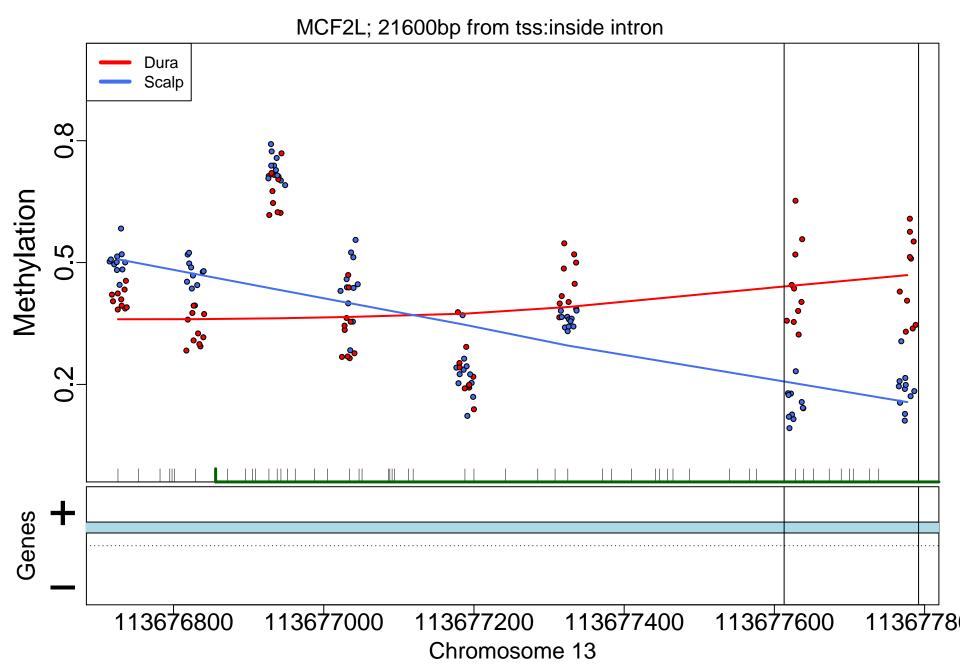
SH3PXD2A; 735bp from tss:inside intron Dura Scalp <u>∞</u>. Methylation Genes SH3PXD2A 105614400 105614800 105615200 105615600 Chromosome 10

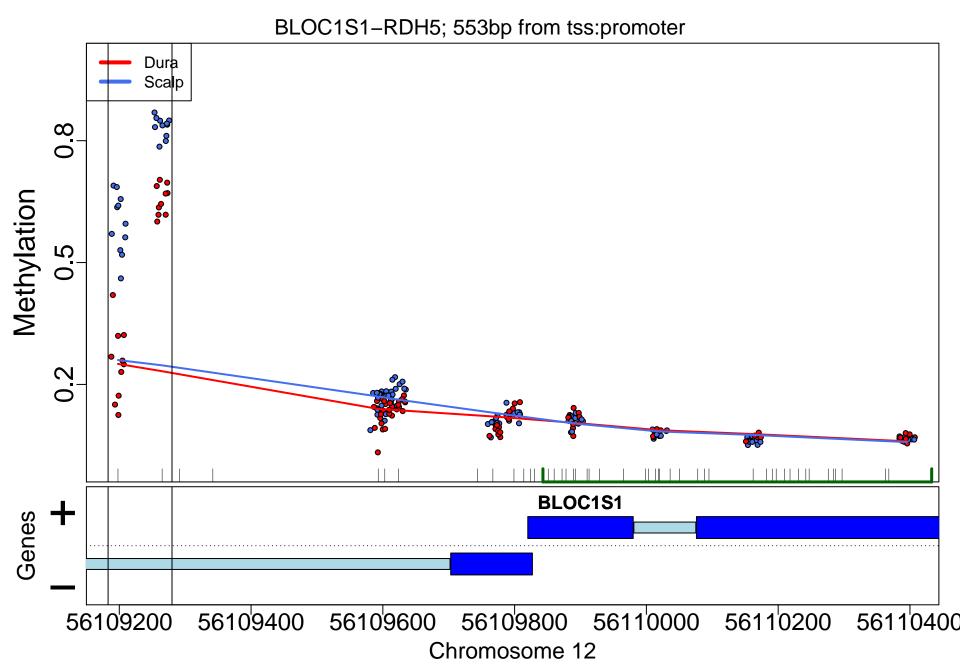


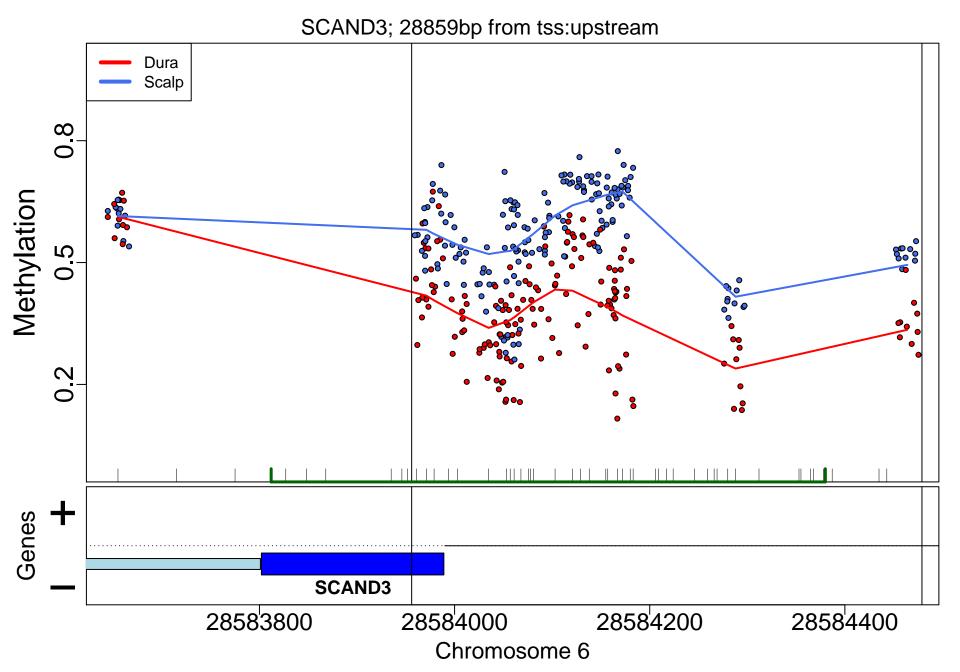




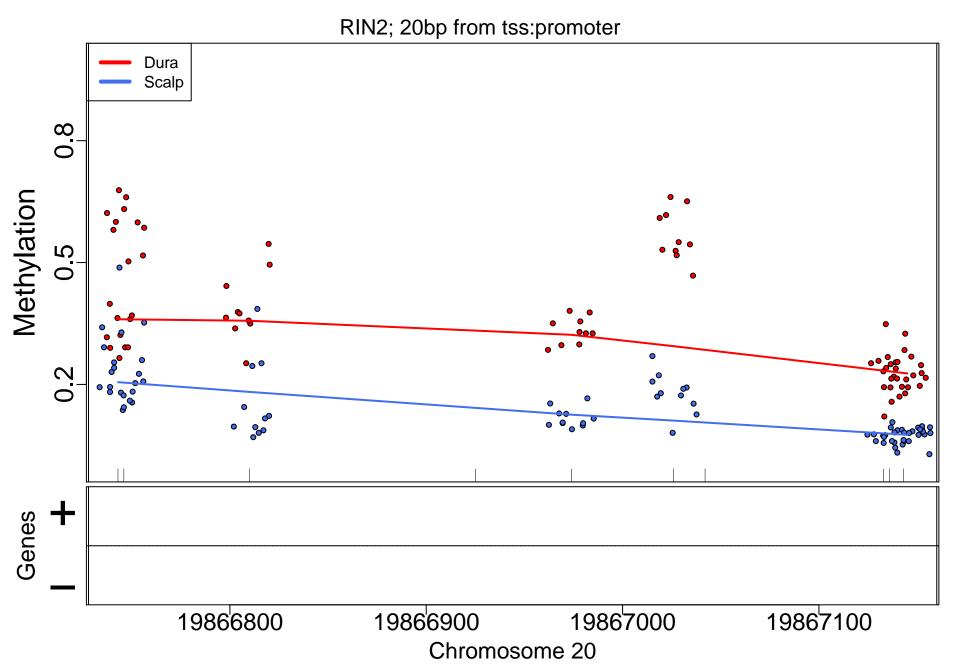






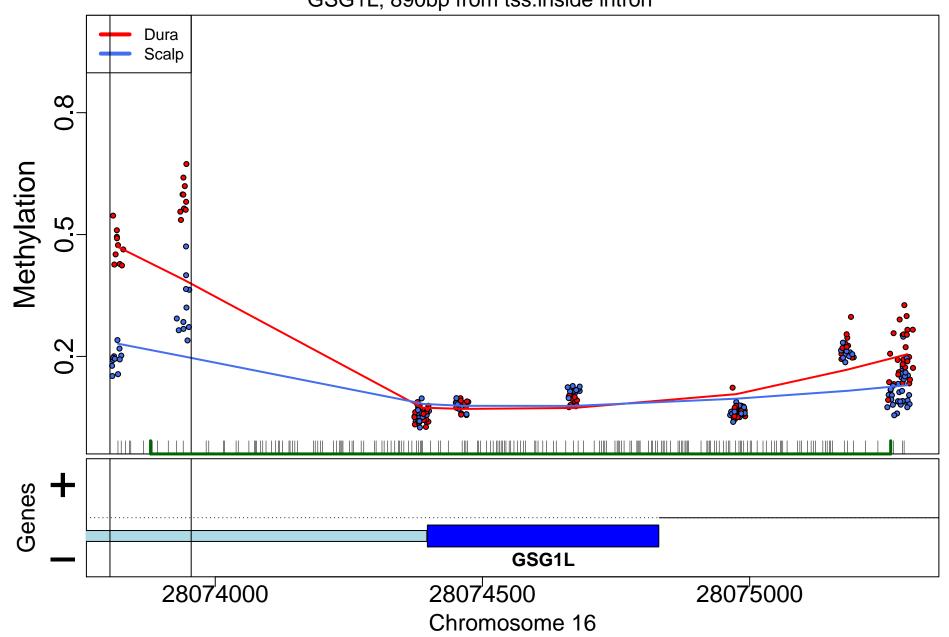


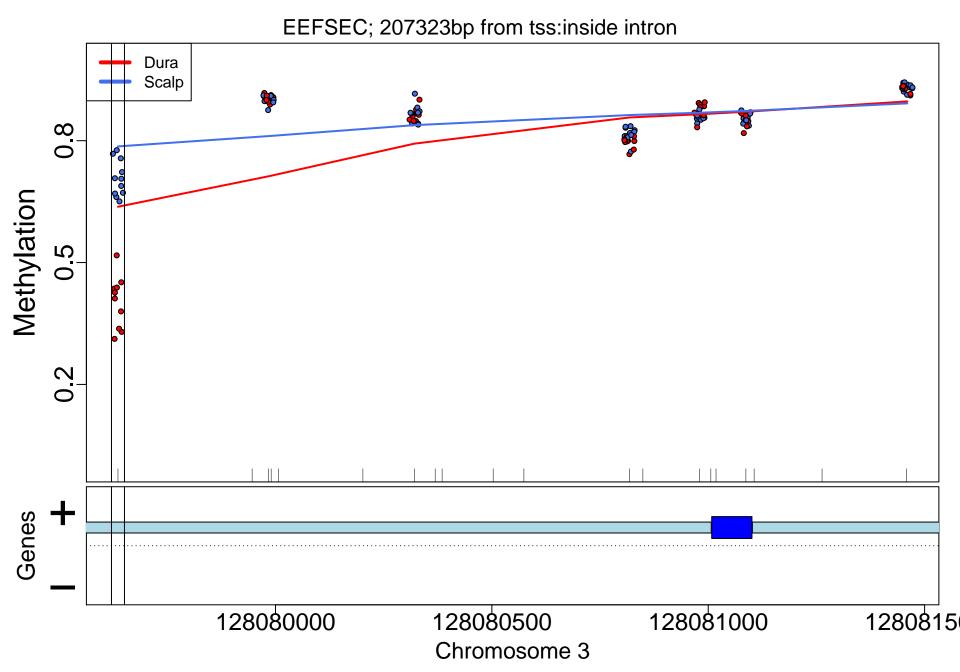
HOXA3; 14055bp from tss:downstream Dura Scalp 80. Methylation 0.2 Genes 27142500 27143000 27143500 27144500 27145000 27144000 Chromosome 7

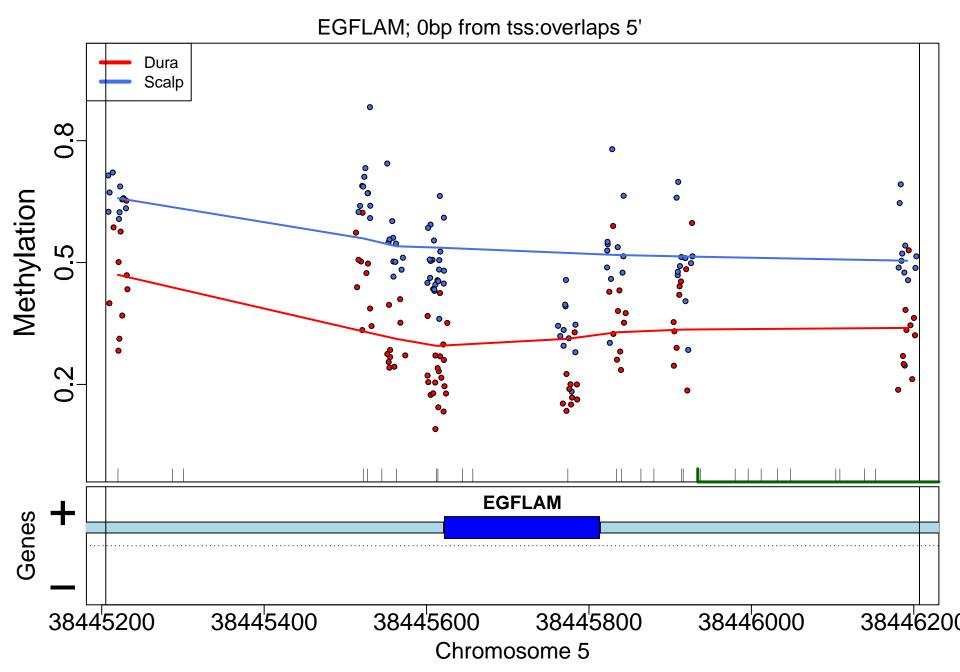


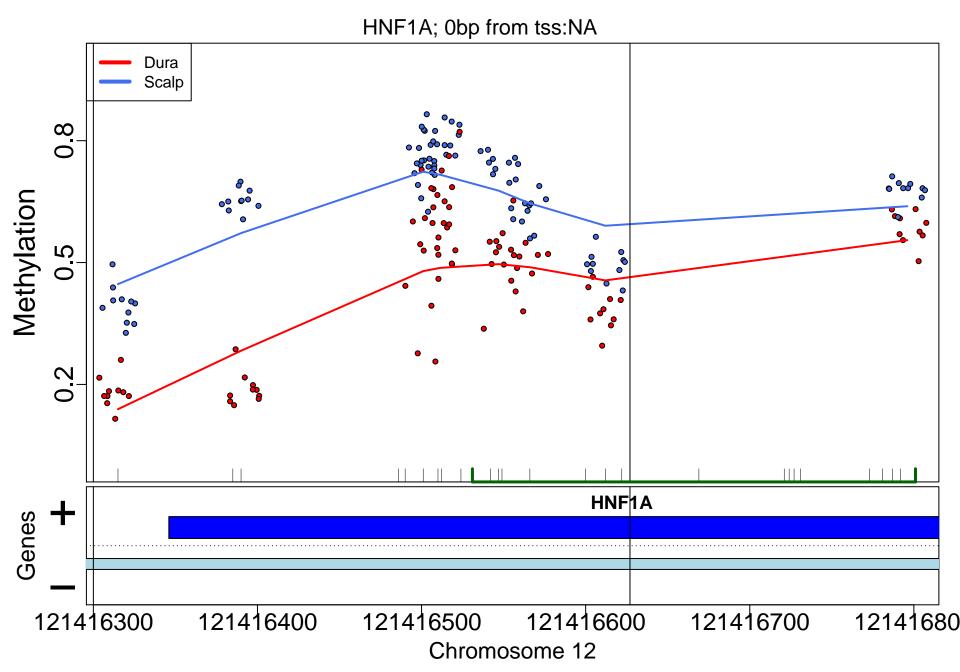
STX2; 19446bp from tss:inside intron Dura Scalp 80. Methylation 0.2 Genes 131303000 131304200 131303400 131303800 Chromosome 12

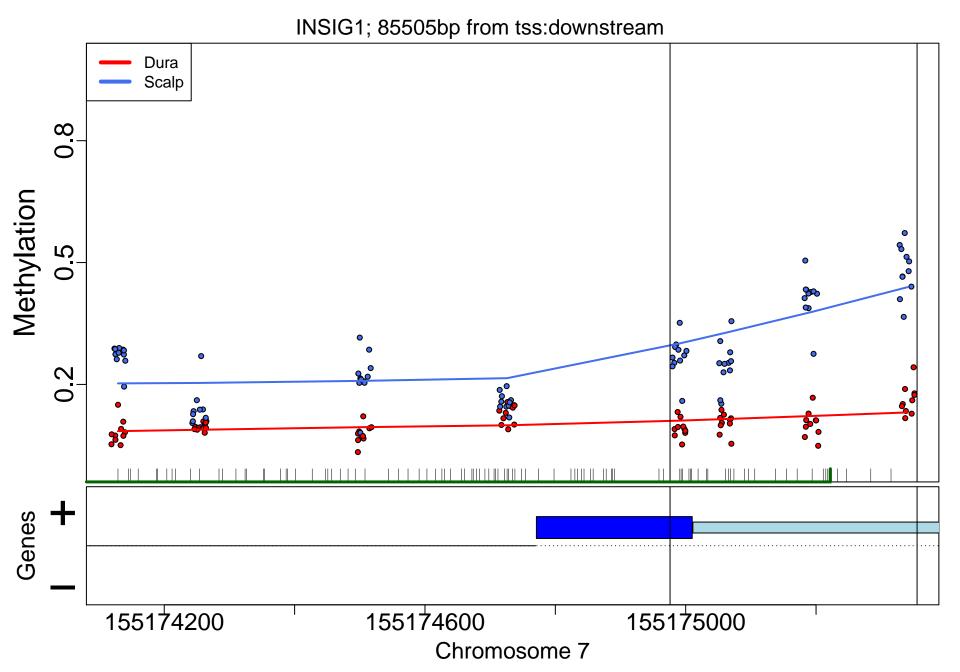
GSG1L; 890bp from tss:inside intron

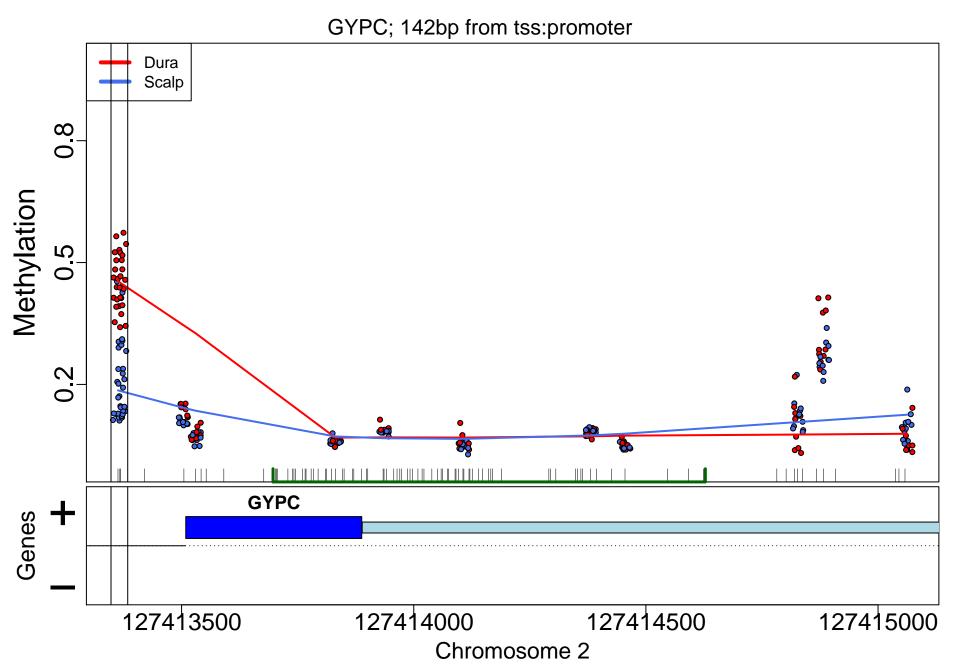


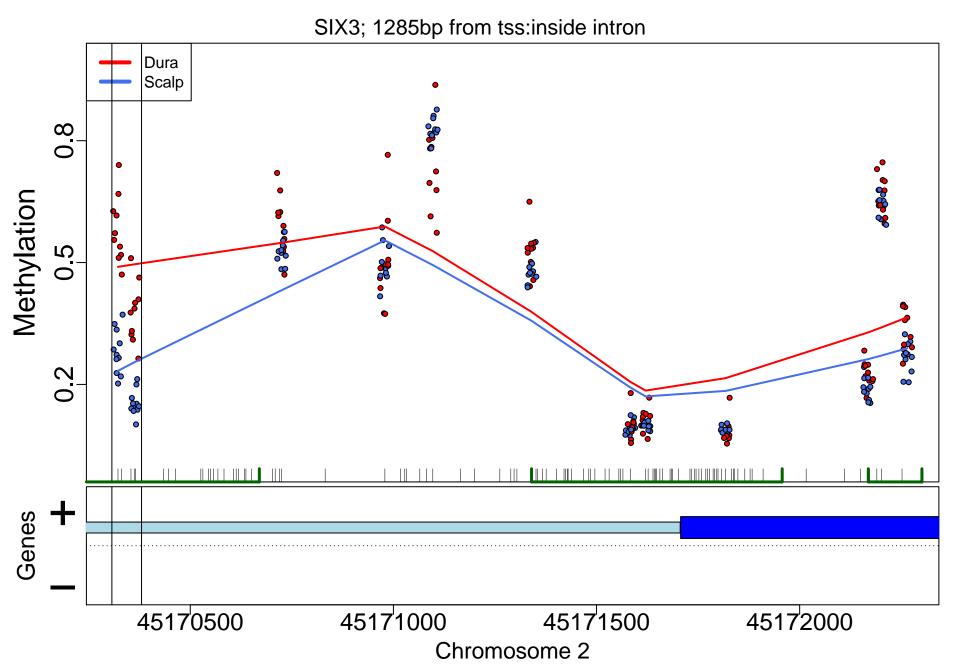




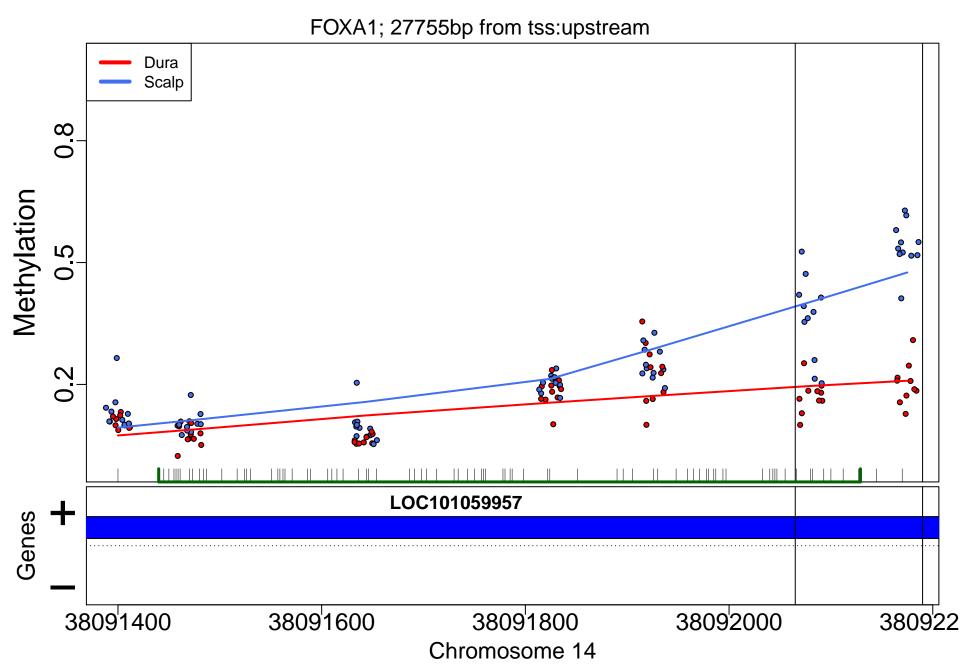




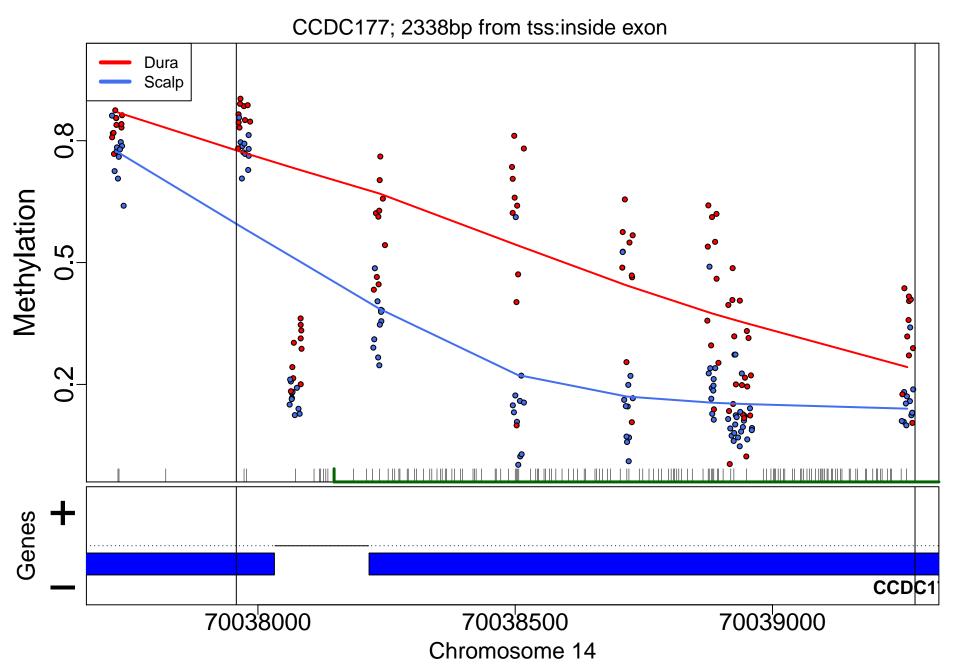


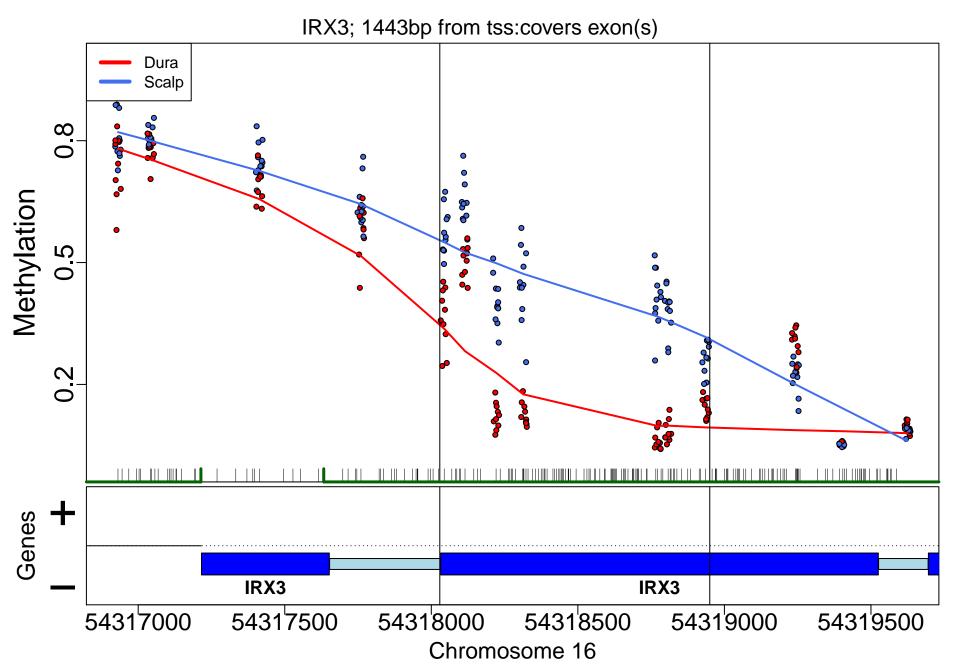


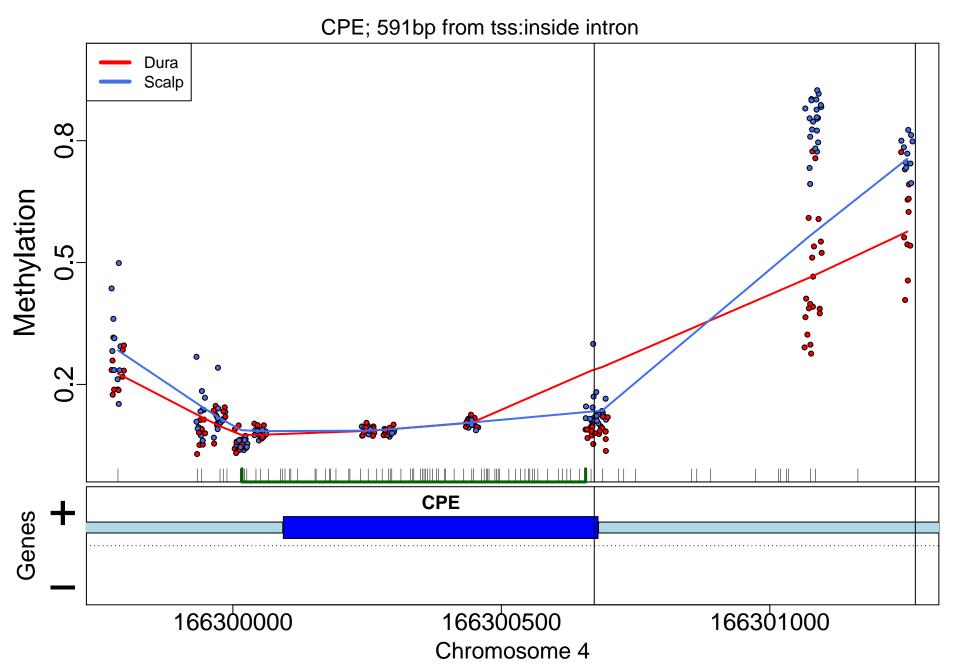
FOXP1; 295082bp from tss:inside intron Dura Scalp <u>∞</u>. Methylation Genes FOXP1 FO 71113500 71114500 71114000 Chromosome 3

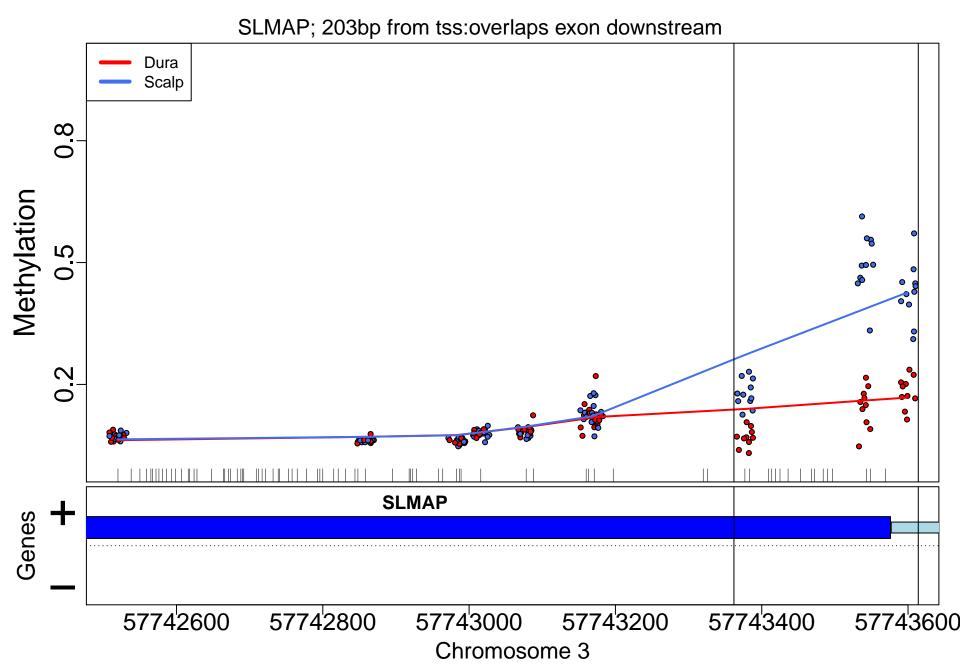


SOX11; 3054bp from tss:inside exon Dura Scalp Methylation SOX11 Genes 5836000 5837000 5836500 Chromosome 2

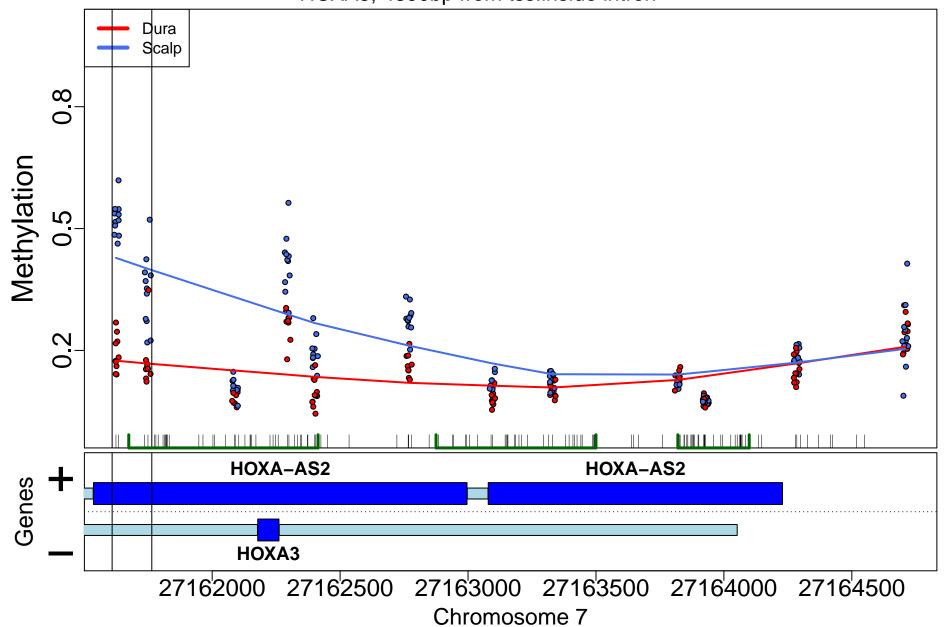


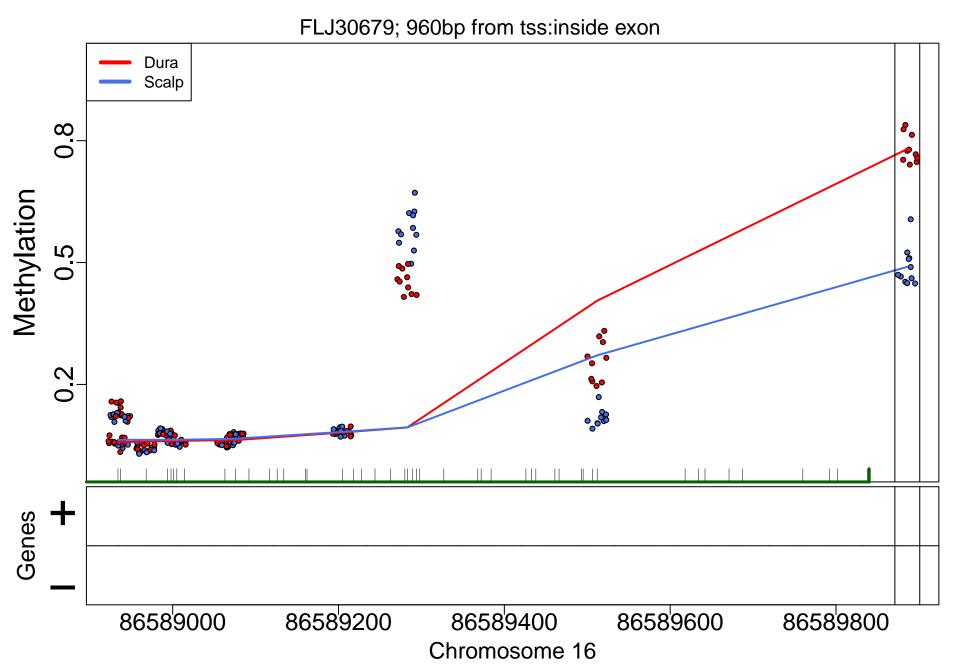


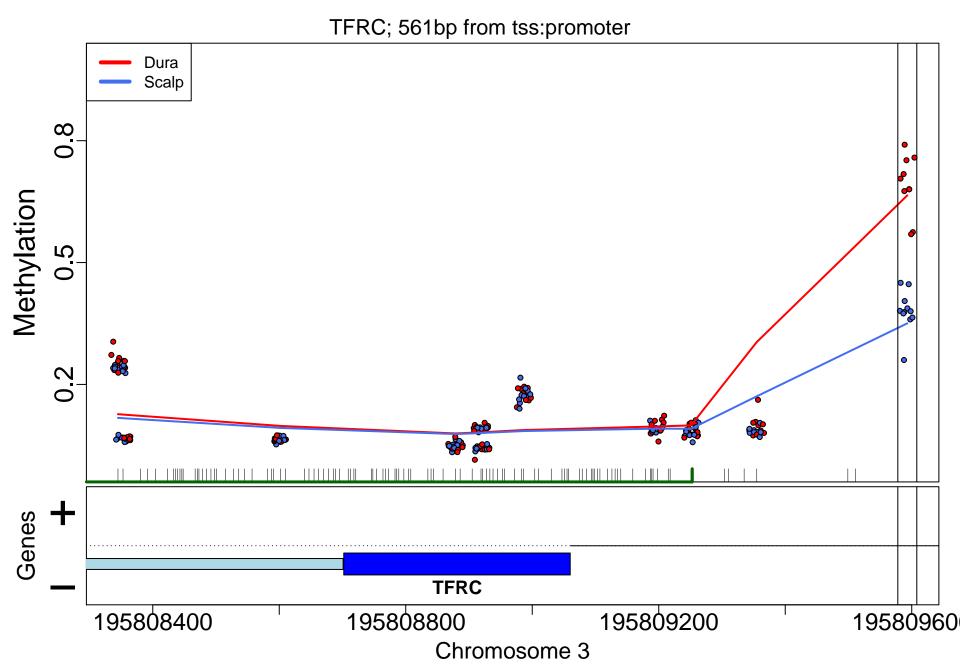


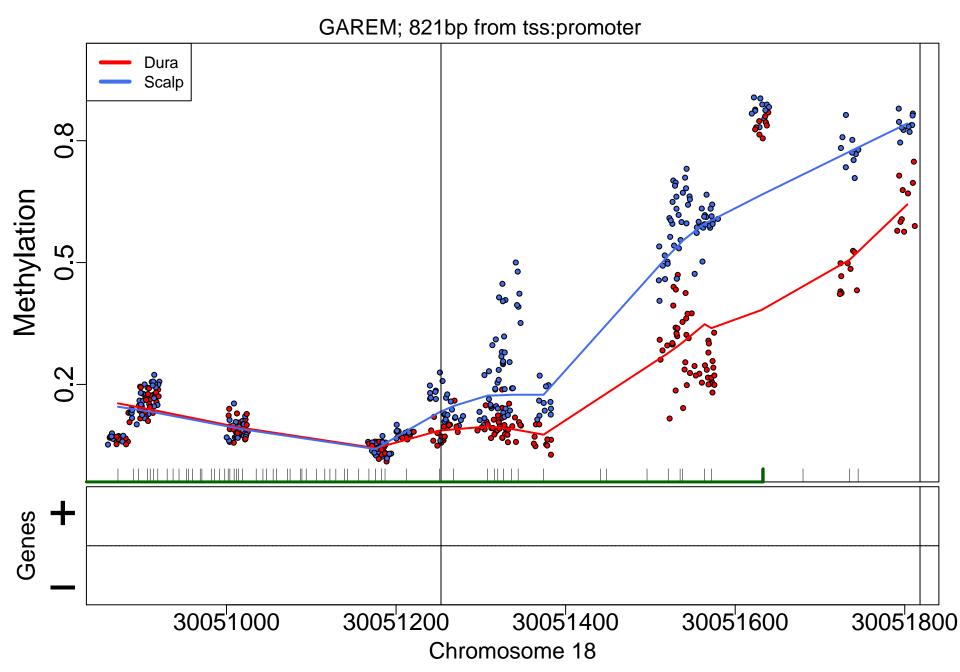


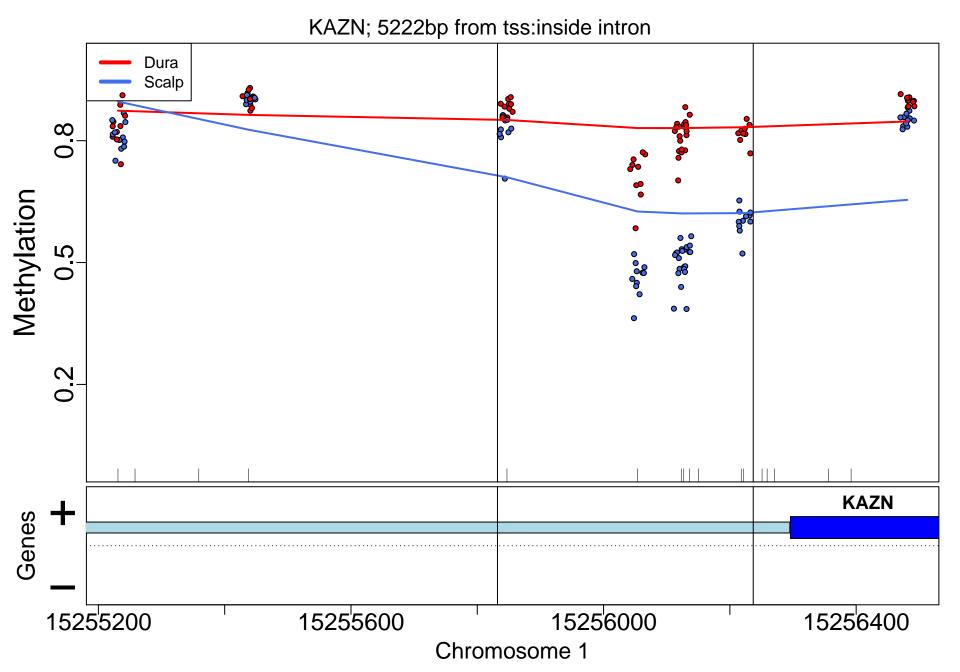
HOXA3; 4890bp from tss:inside intron

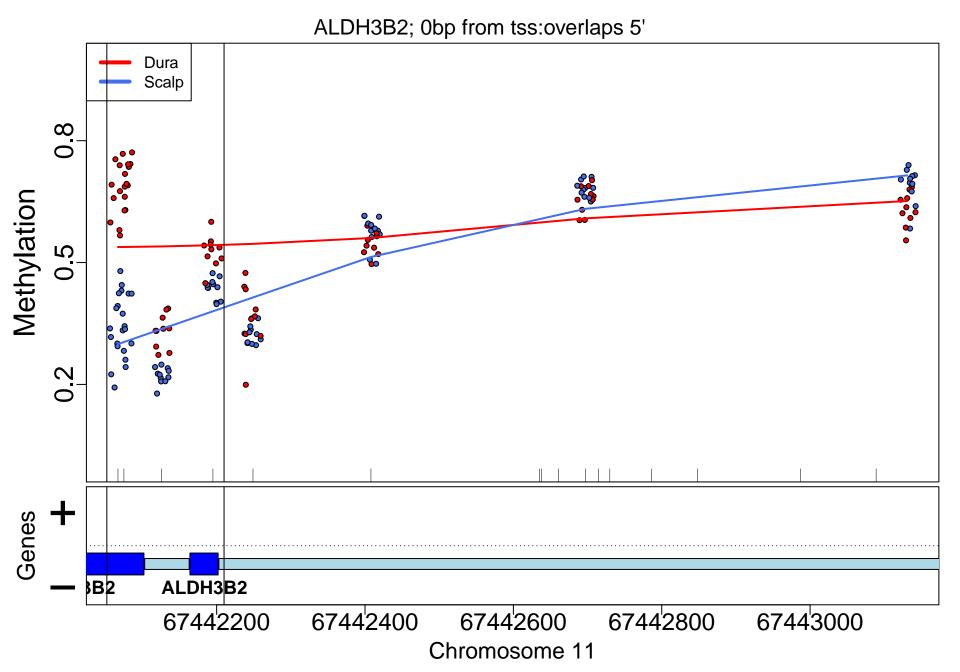


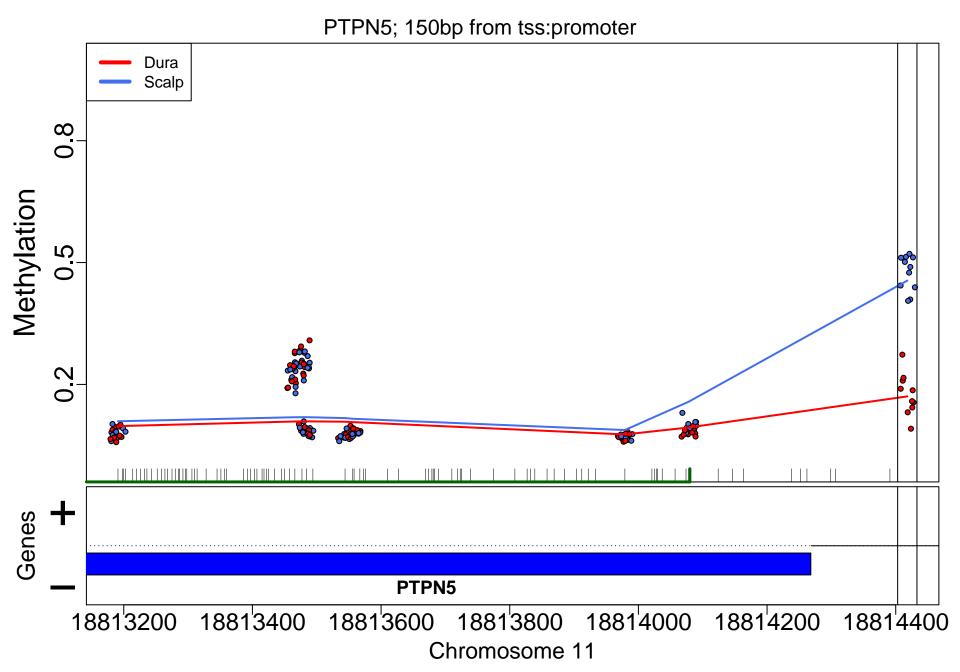


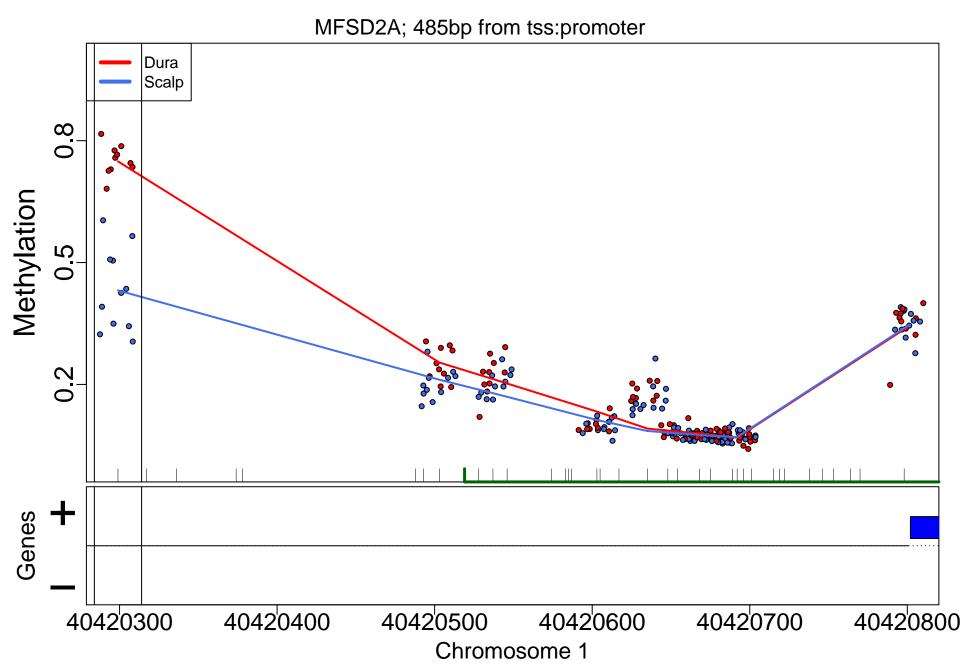


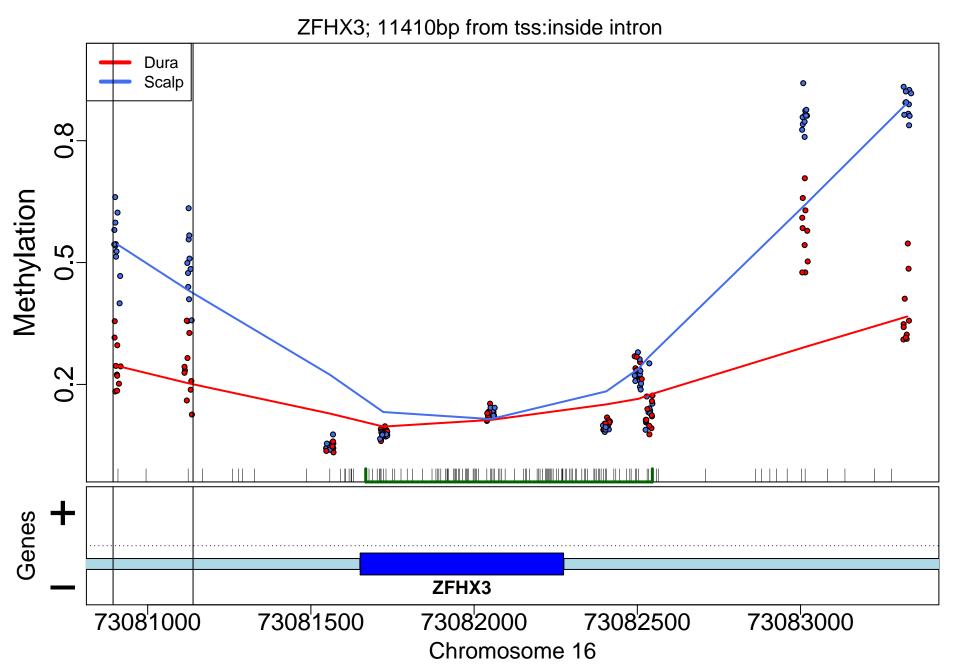


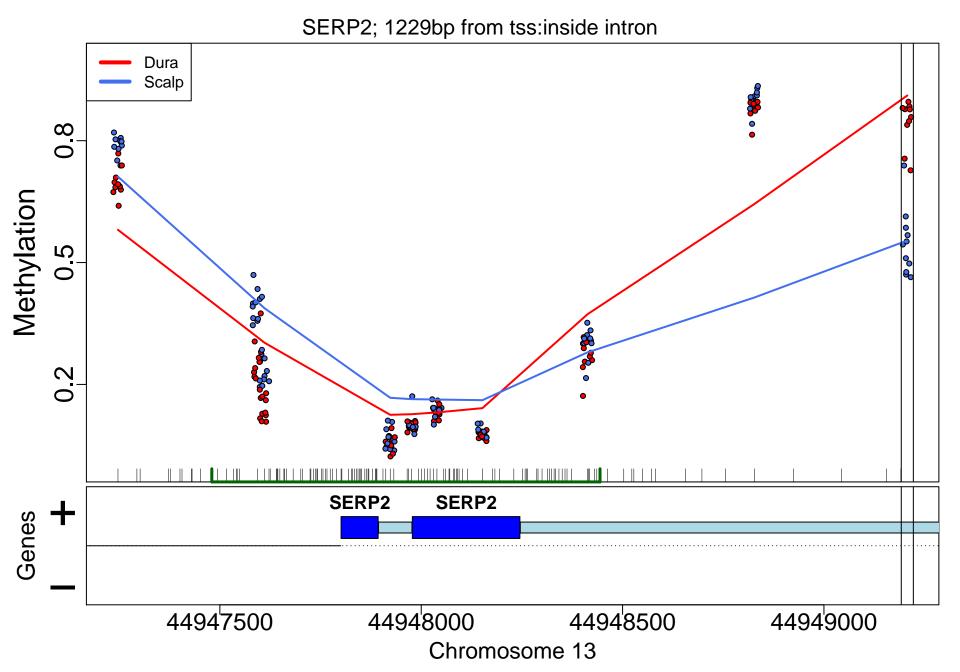


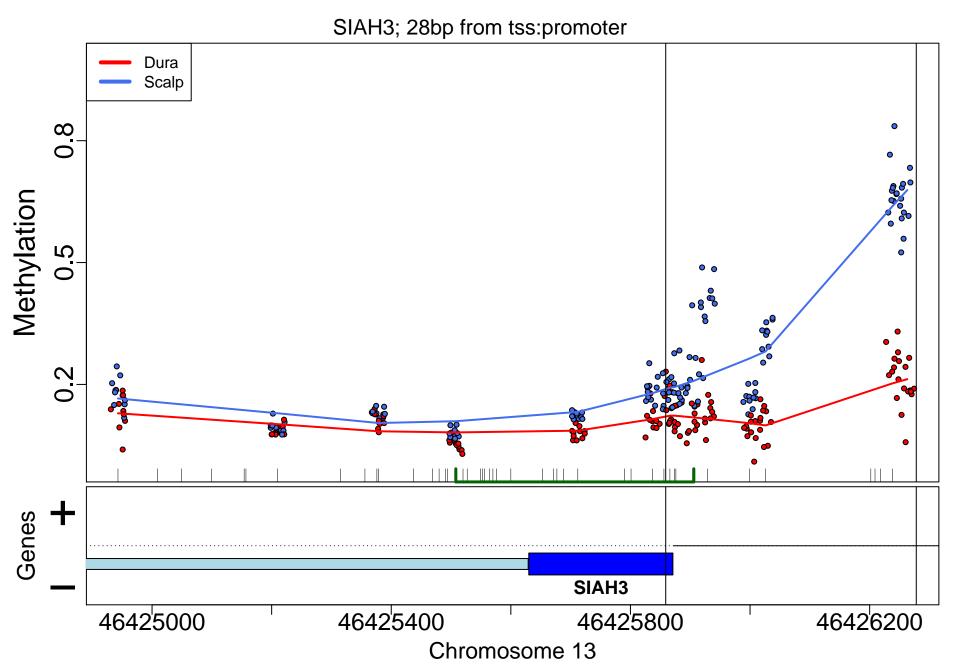


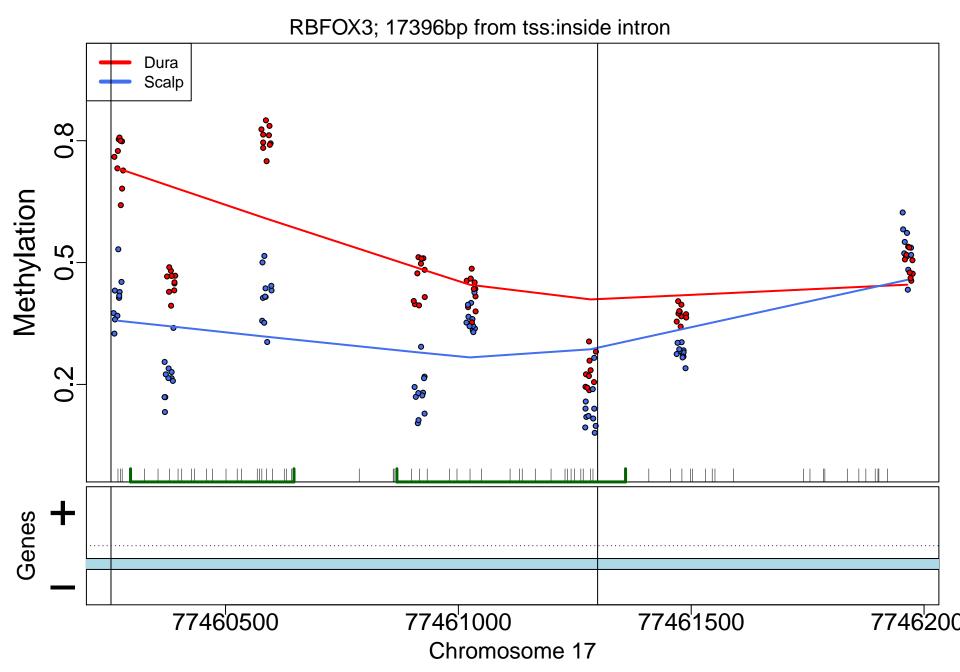


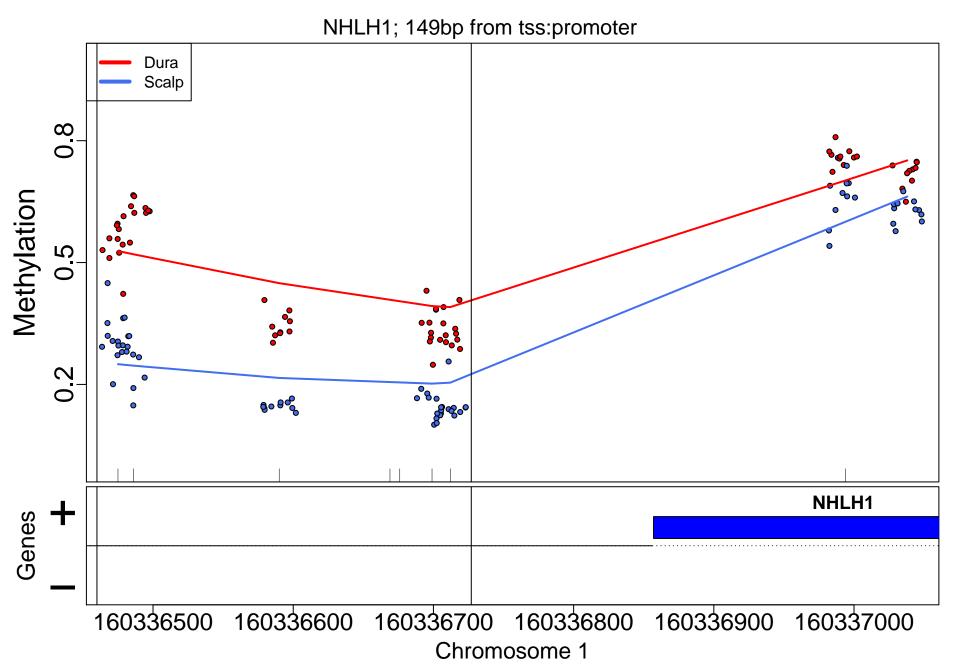


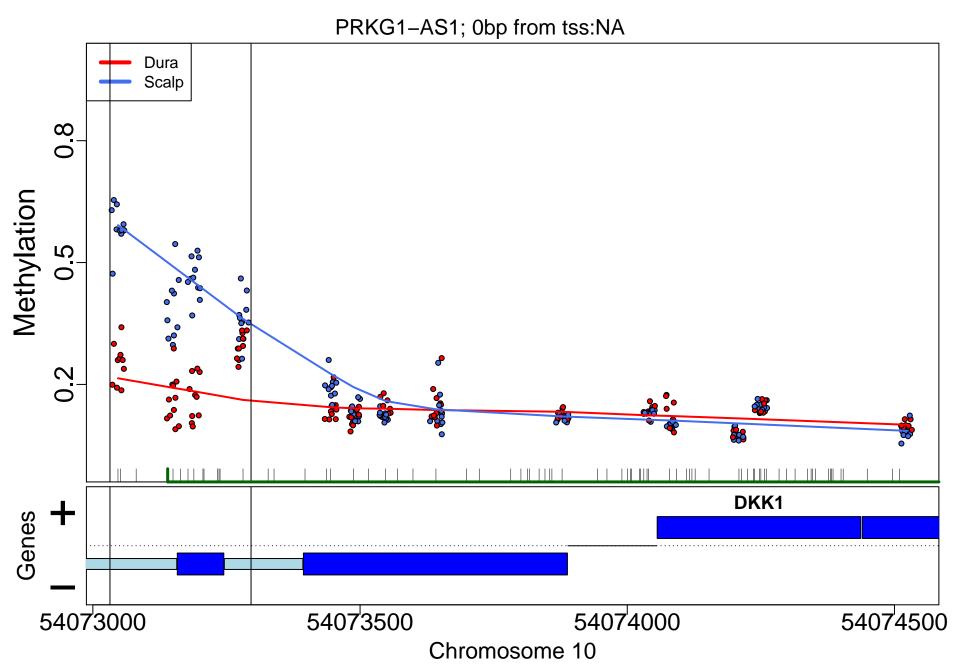


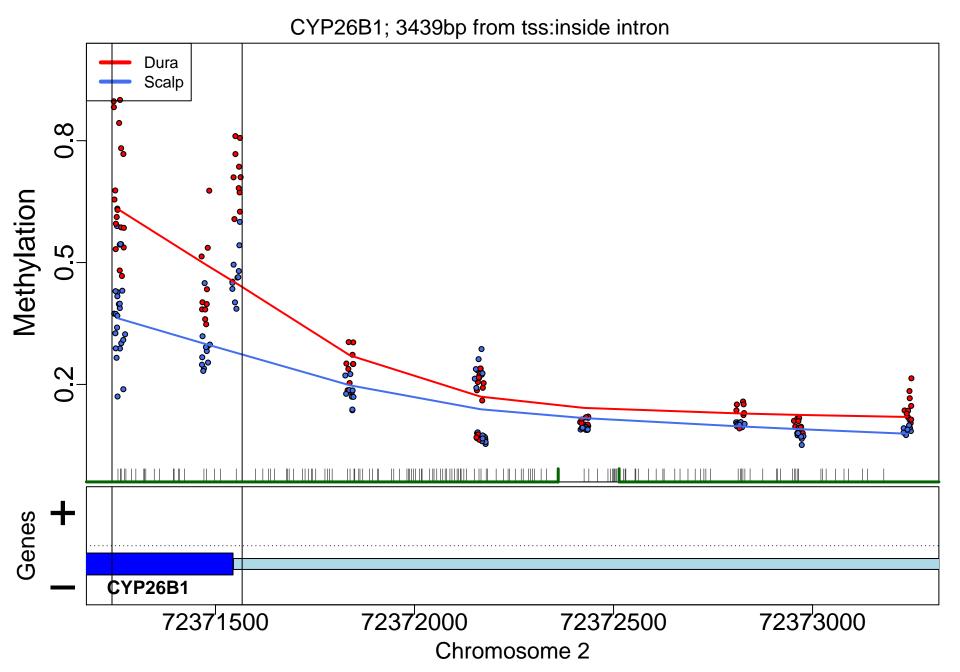


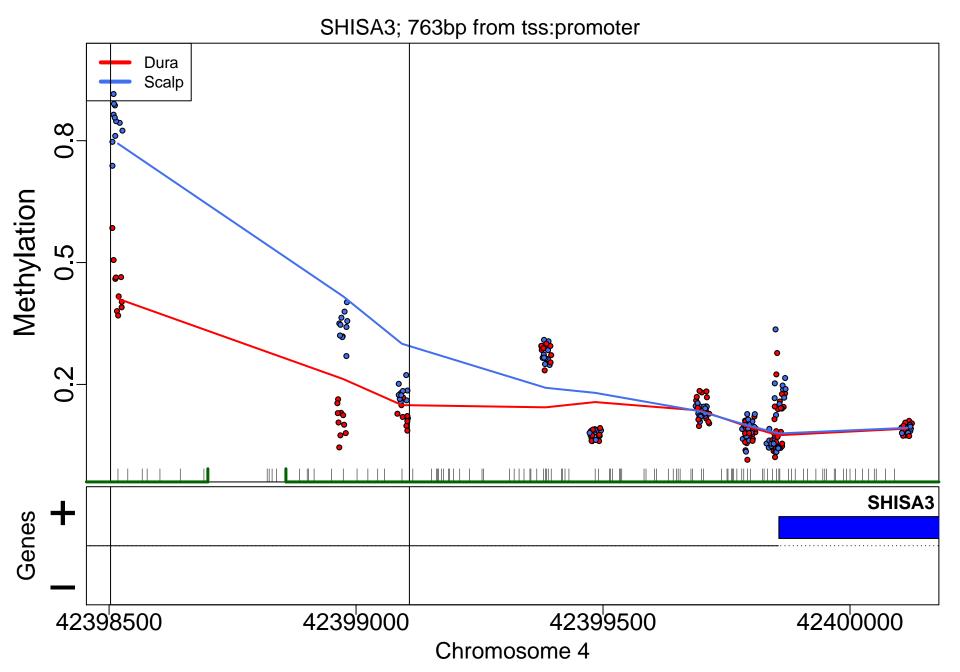


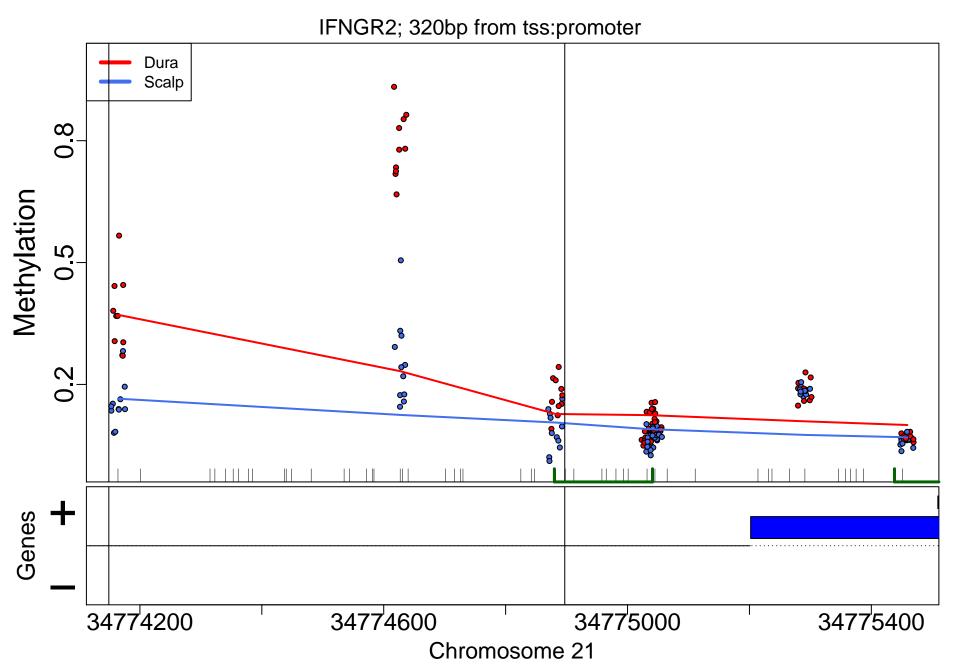


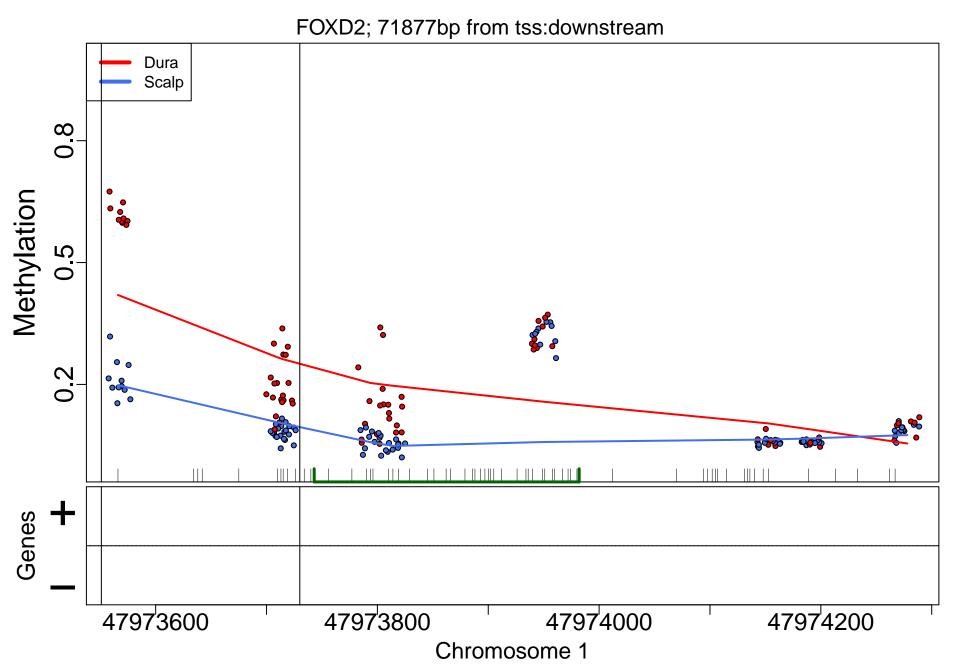


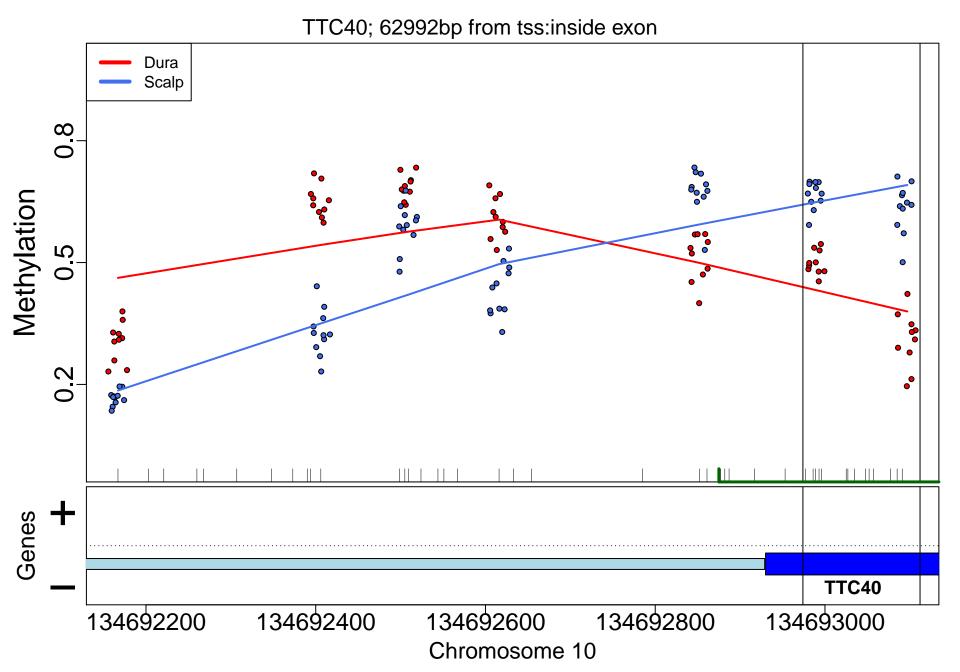


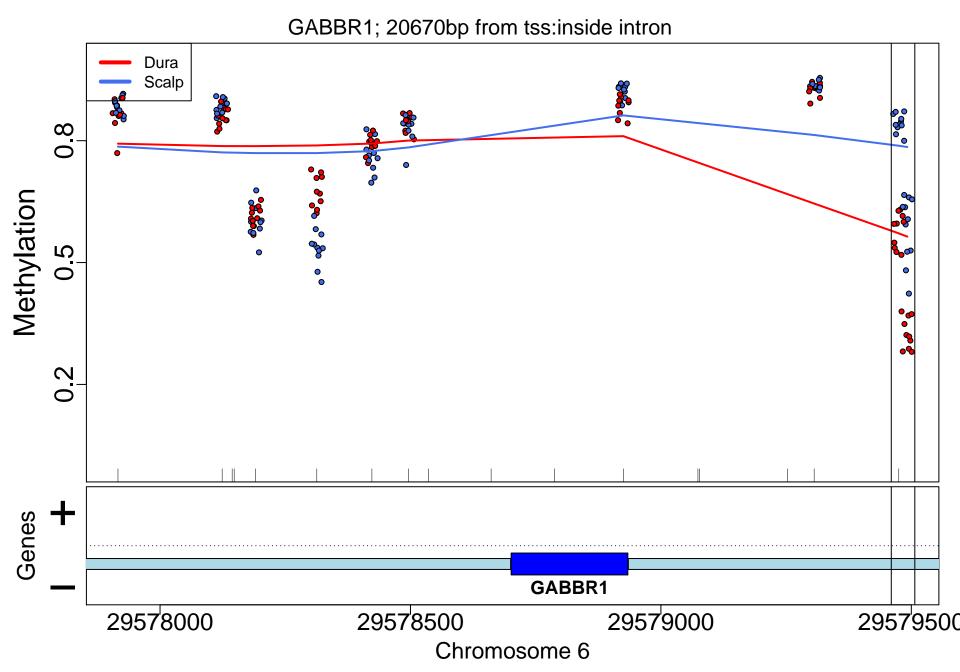


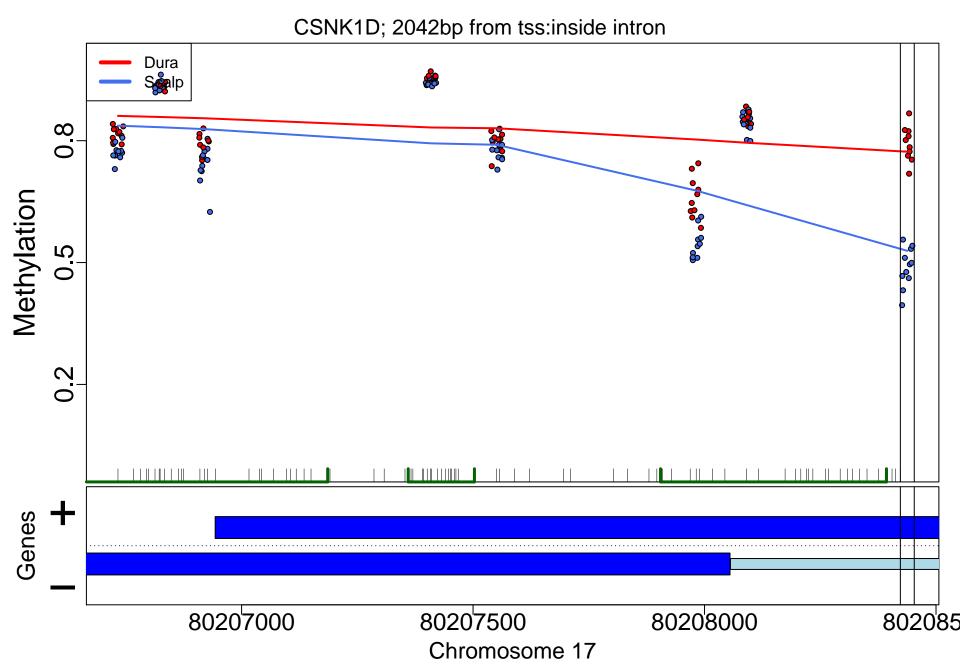


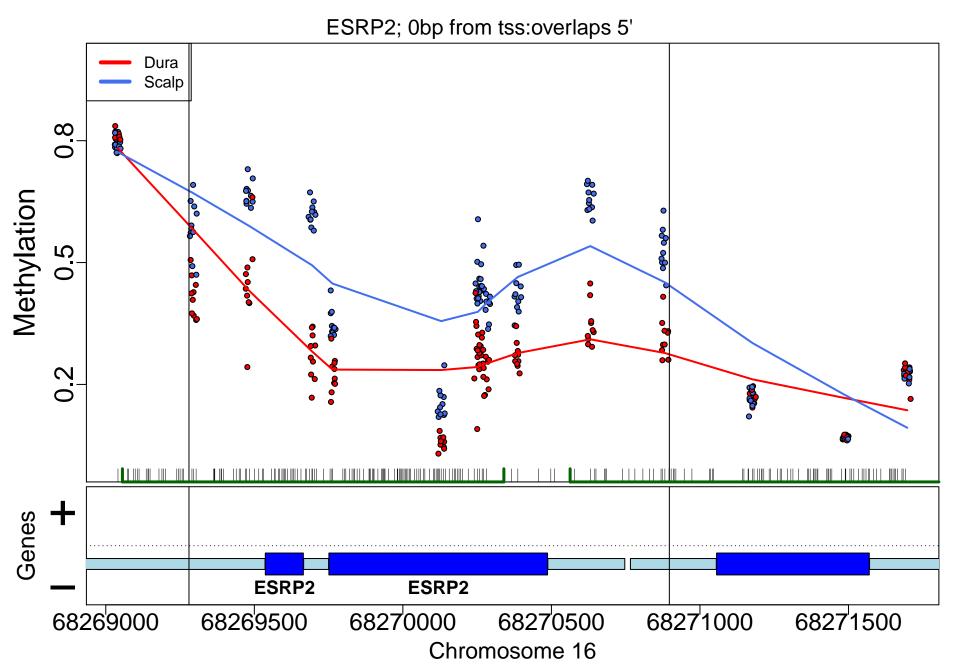


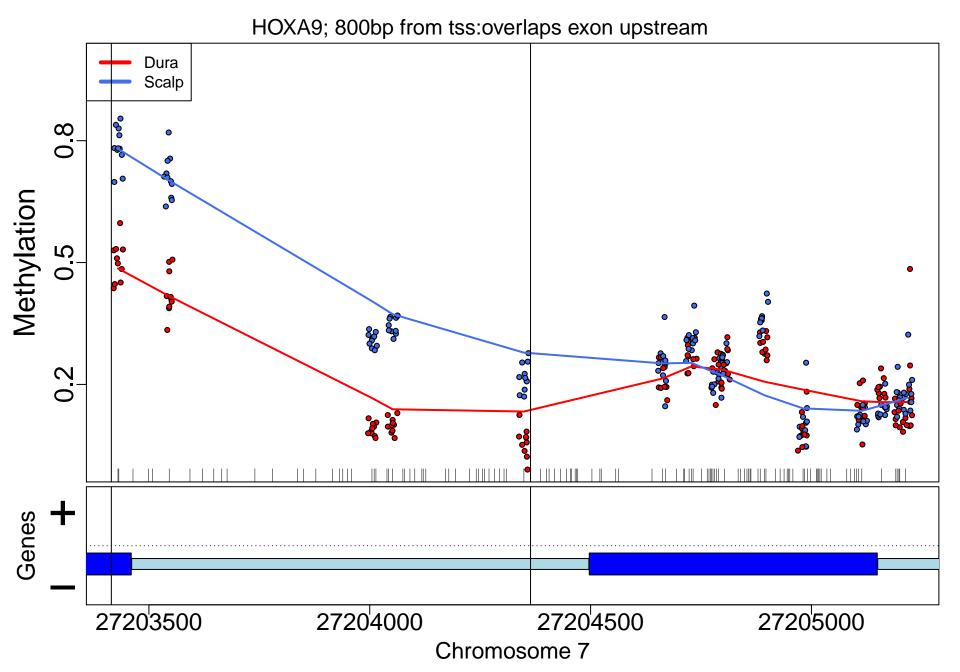


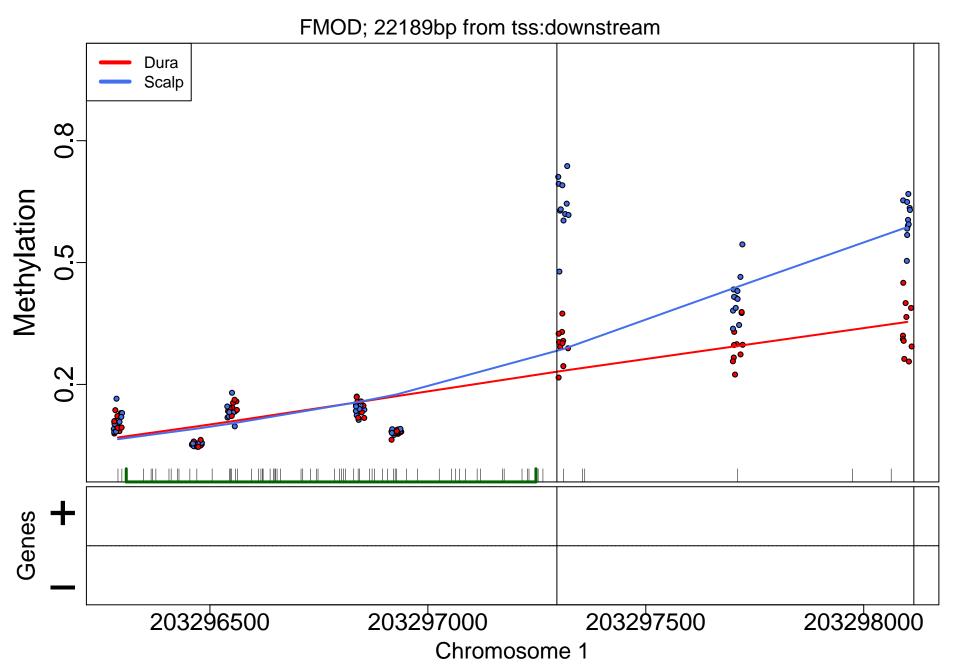


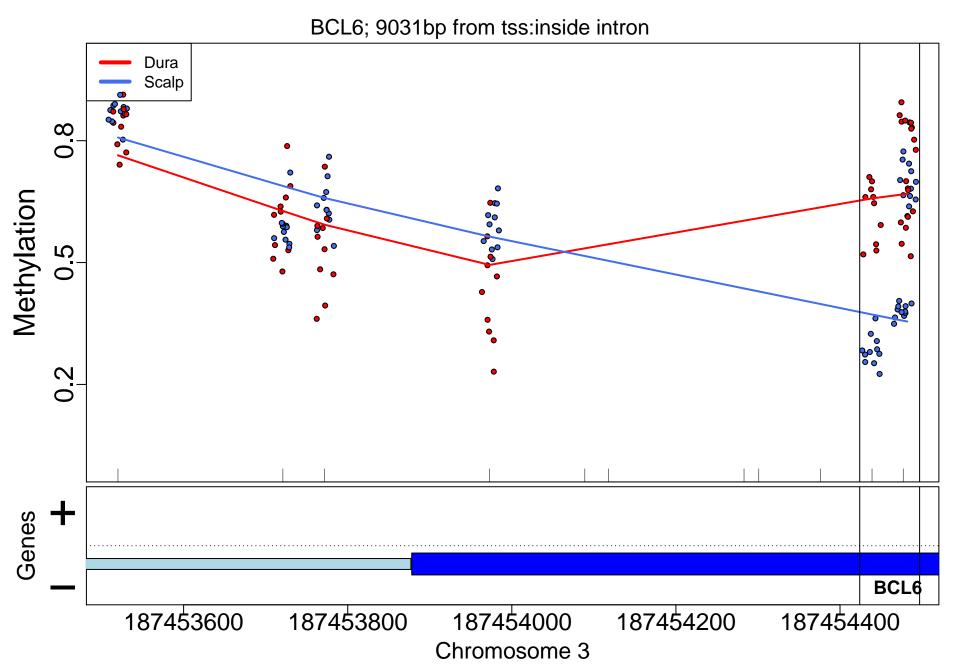


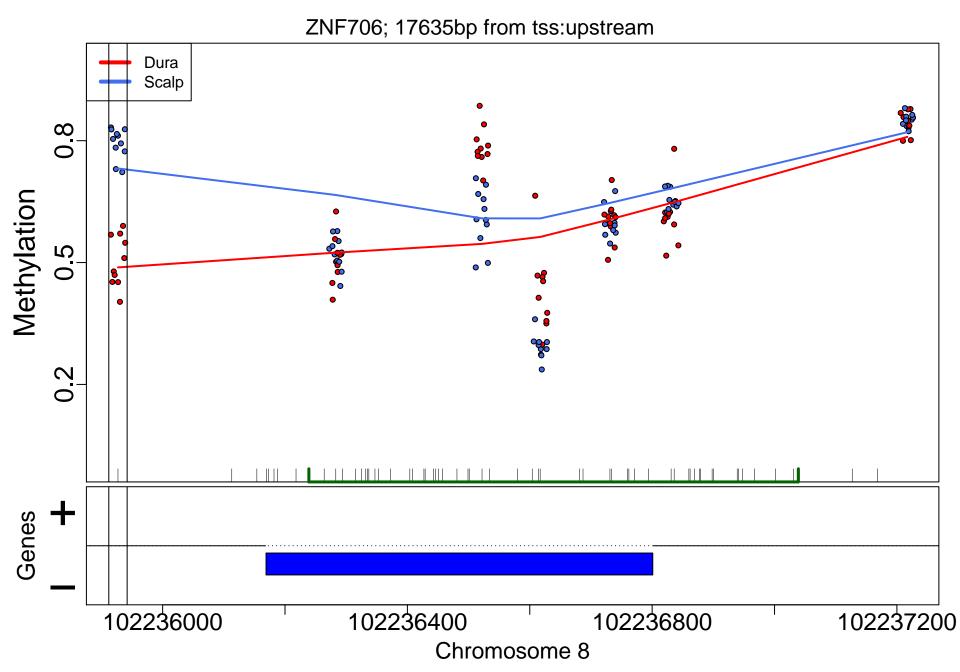


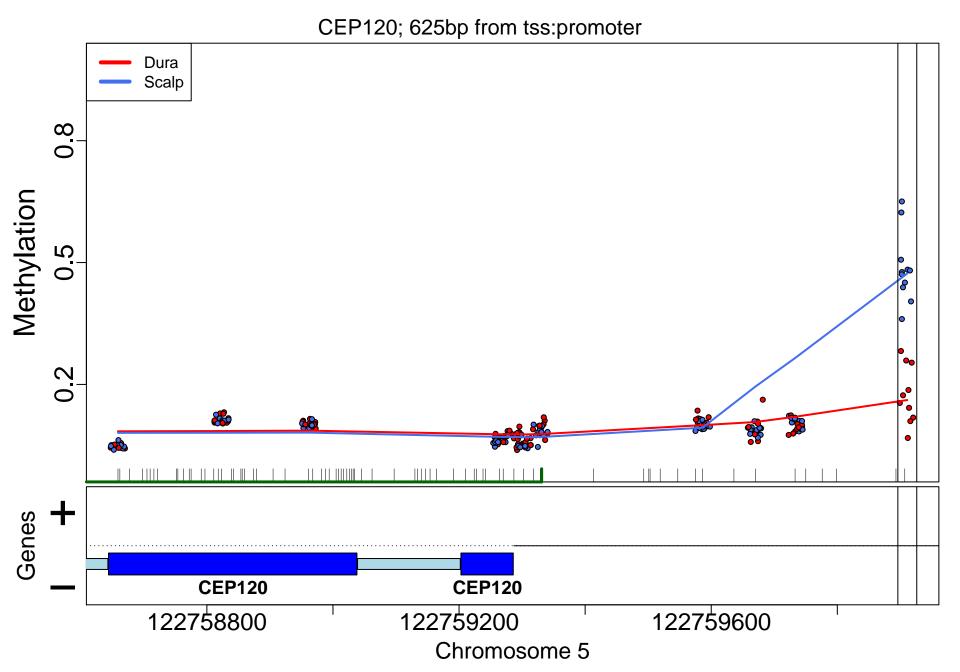


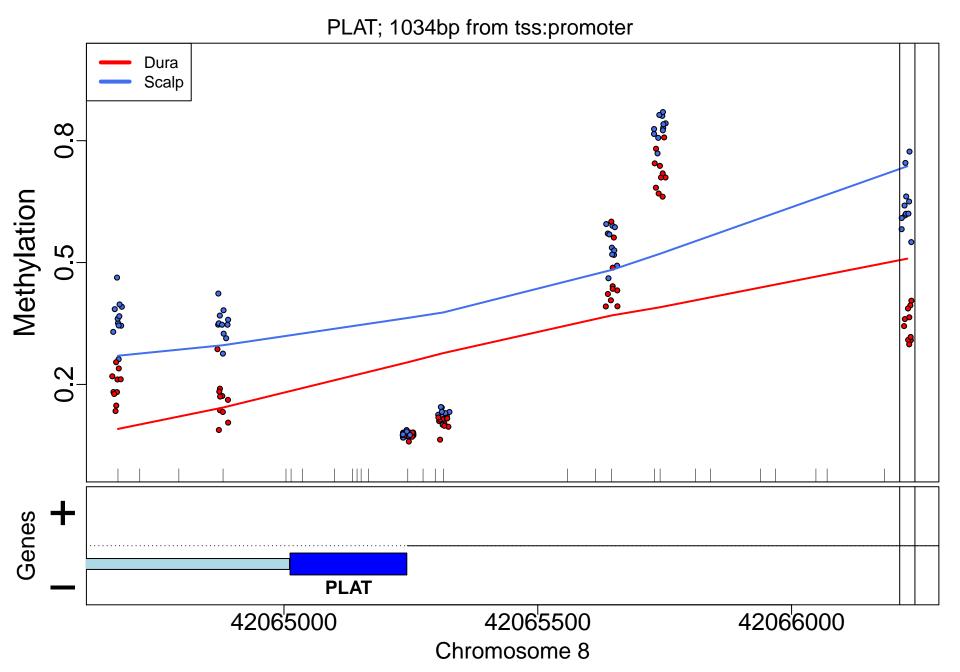


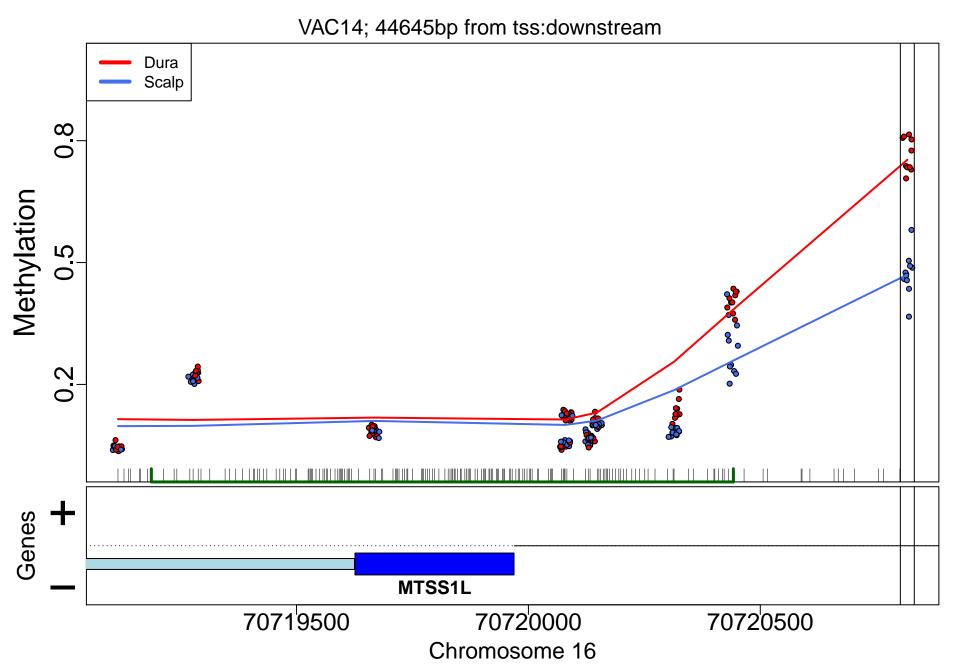


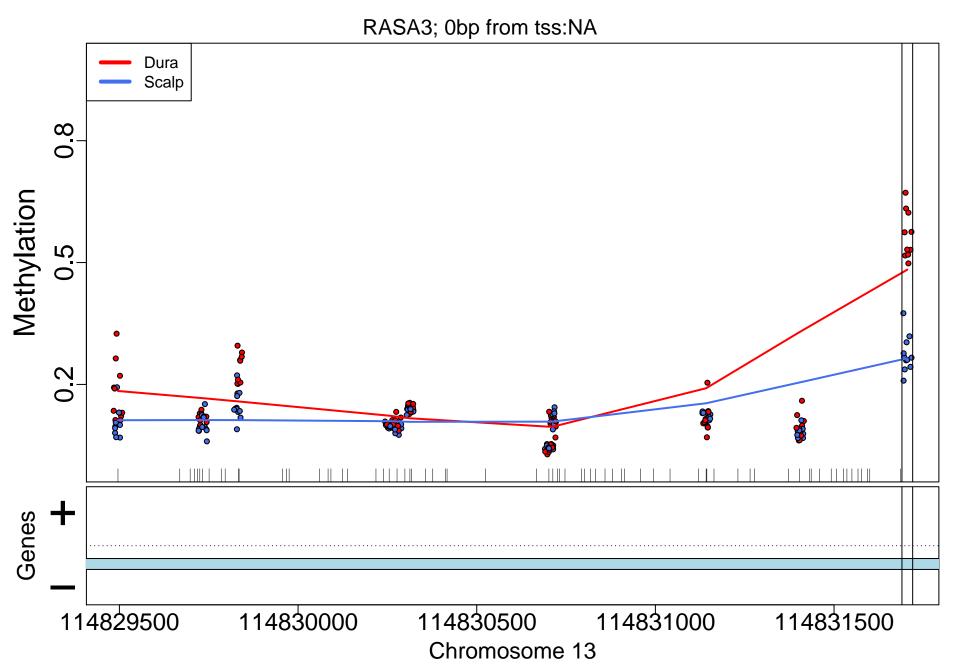


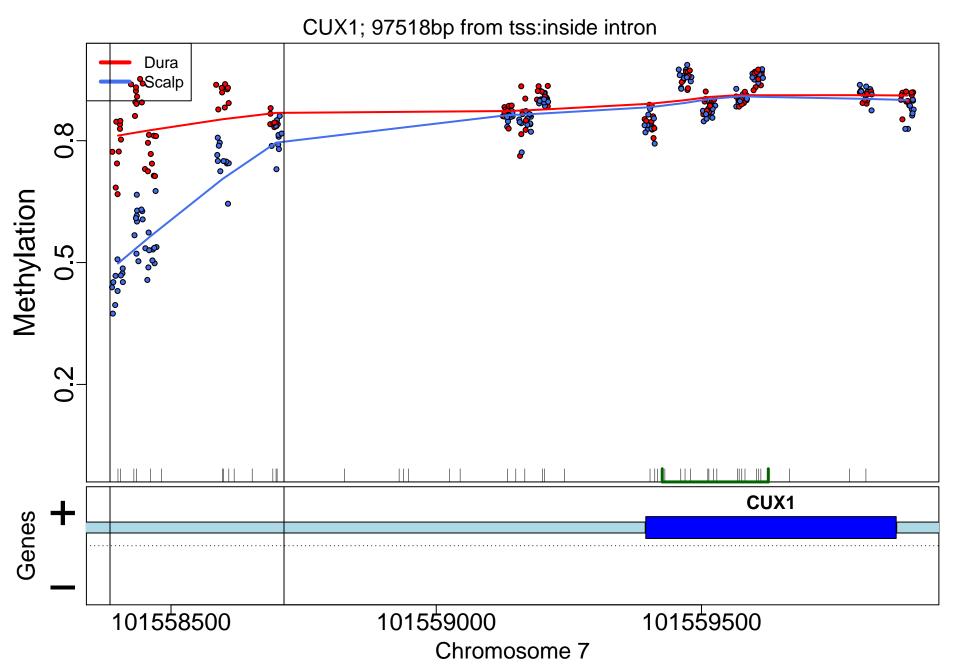


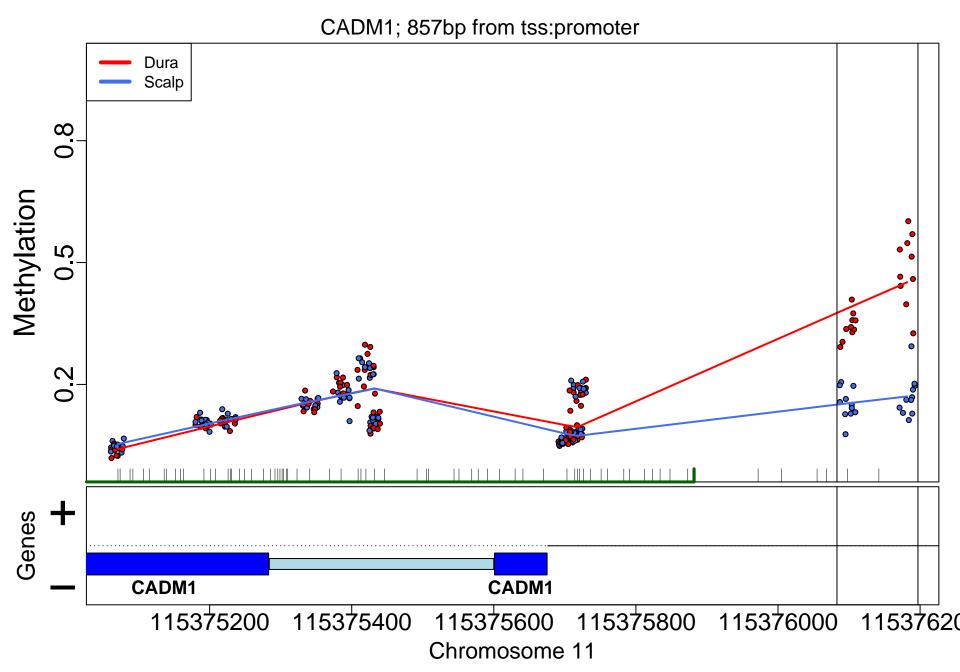


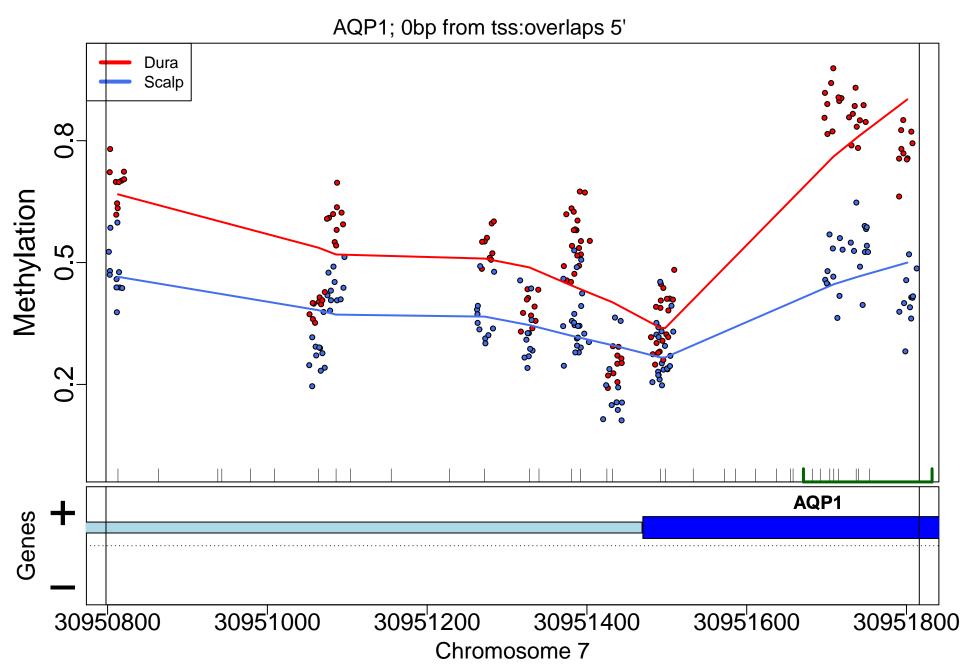




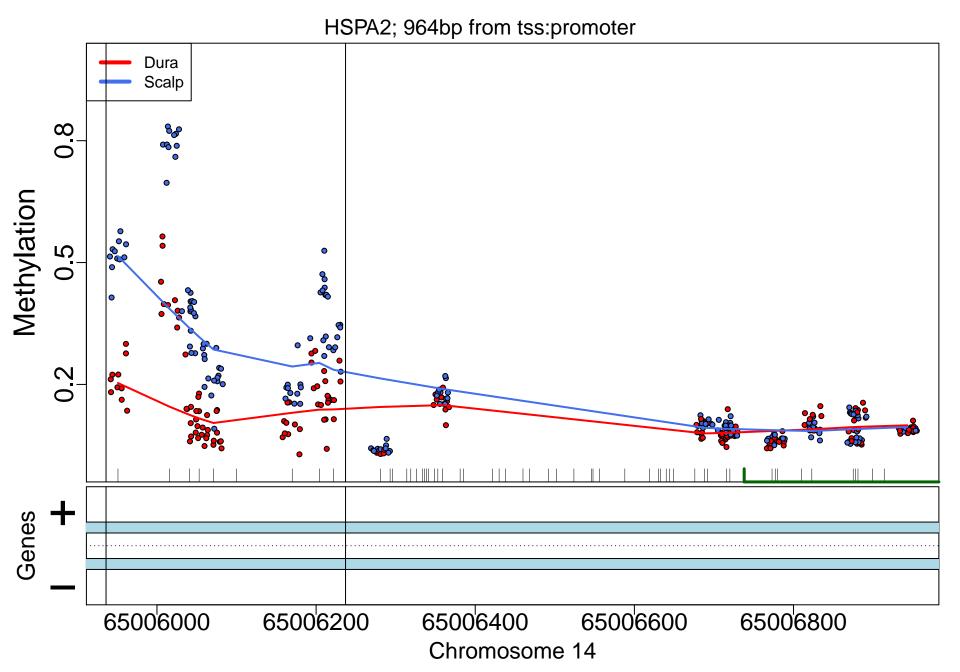


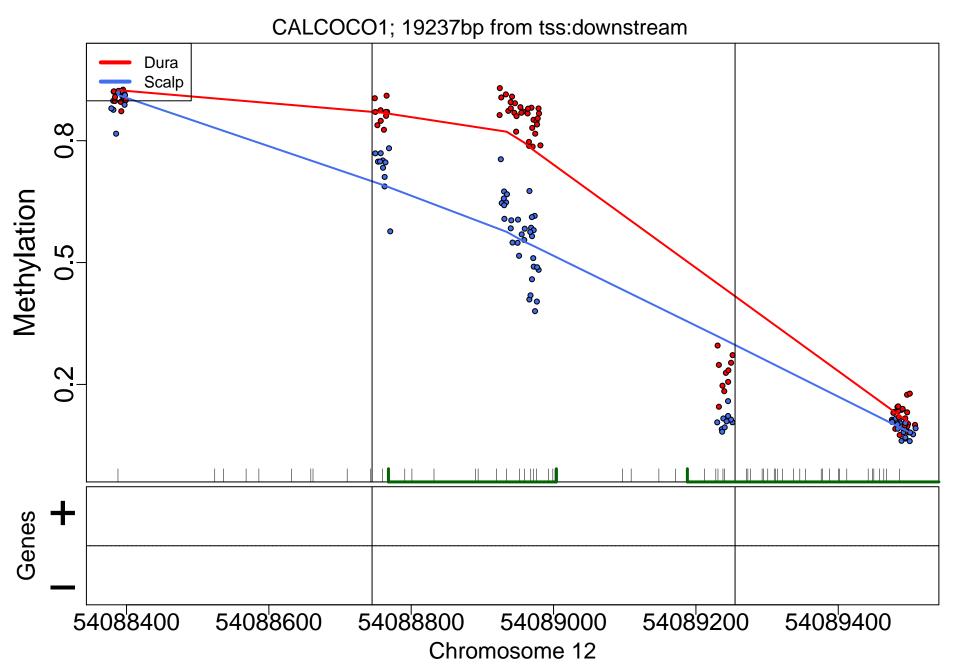


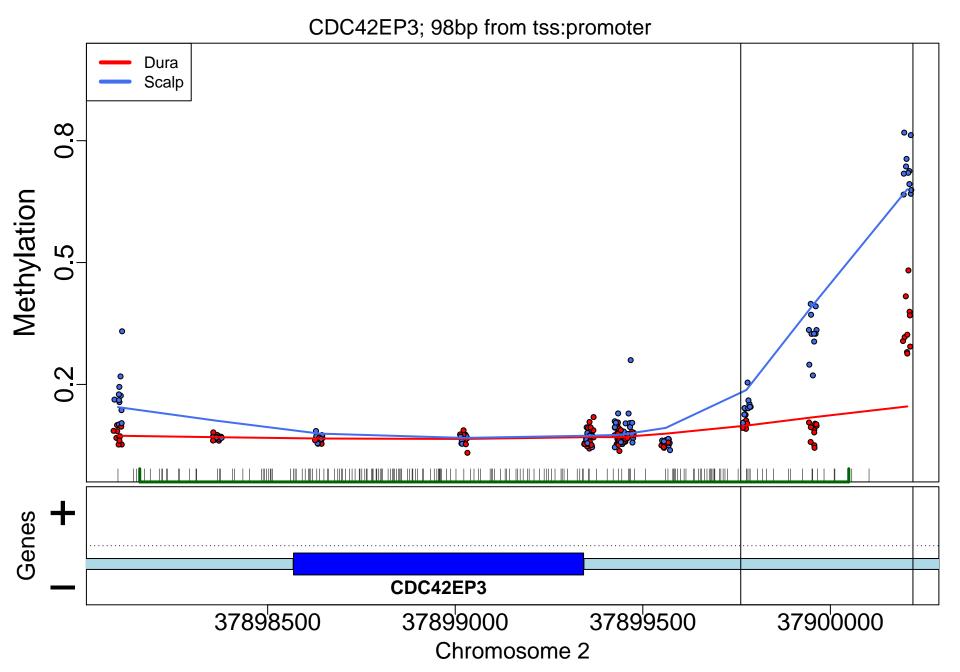


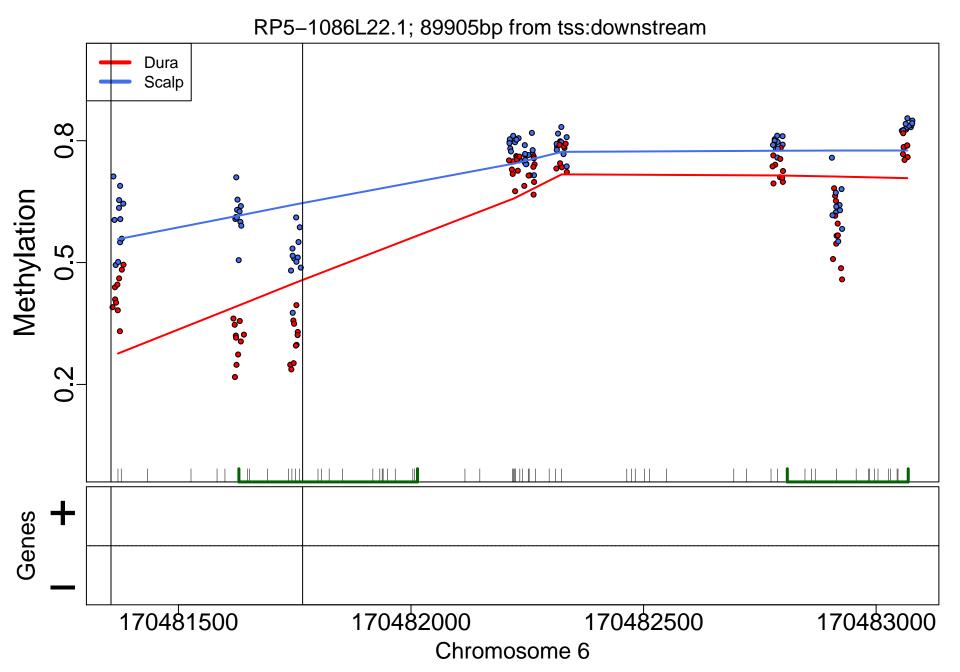


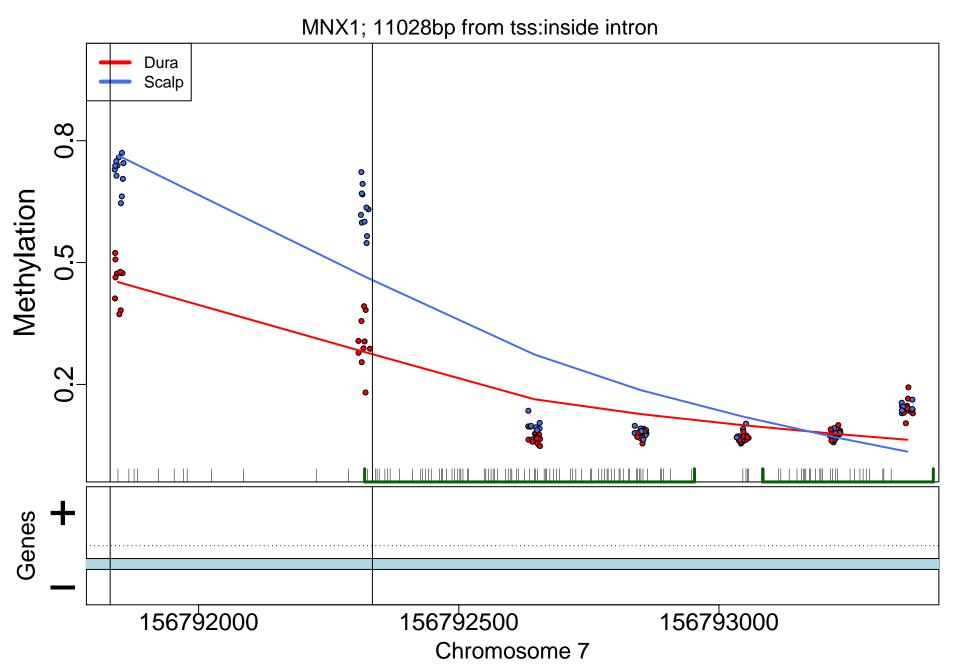
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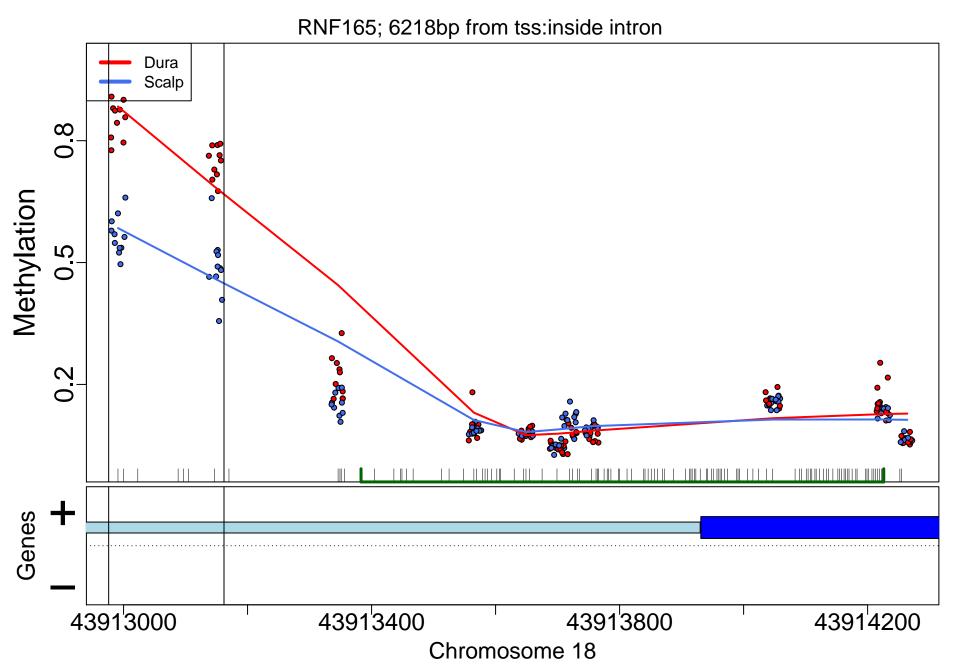


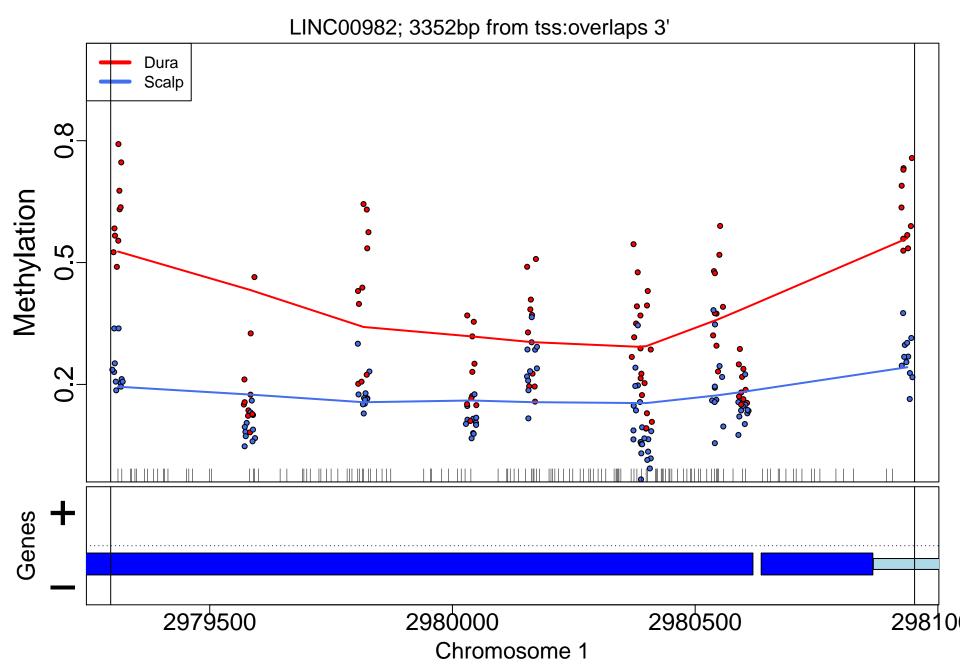


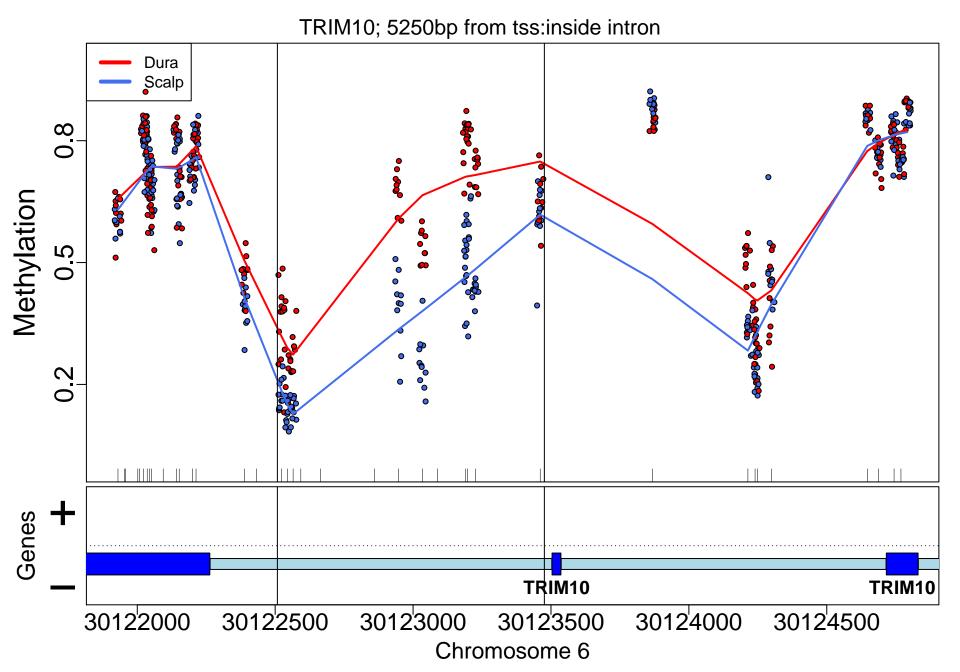


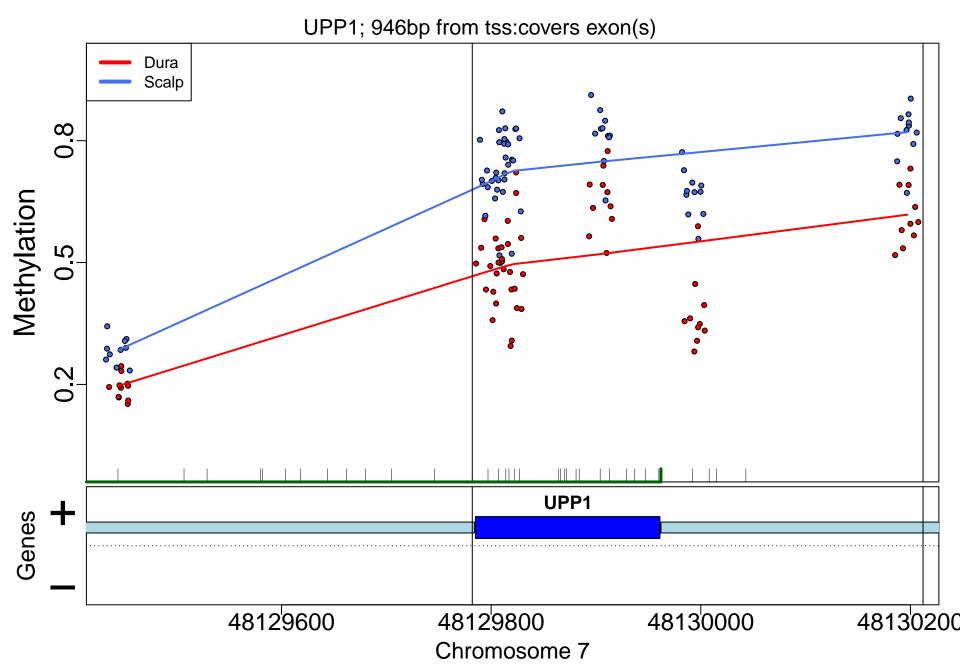


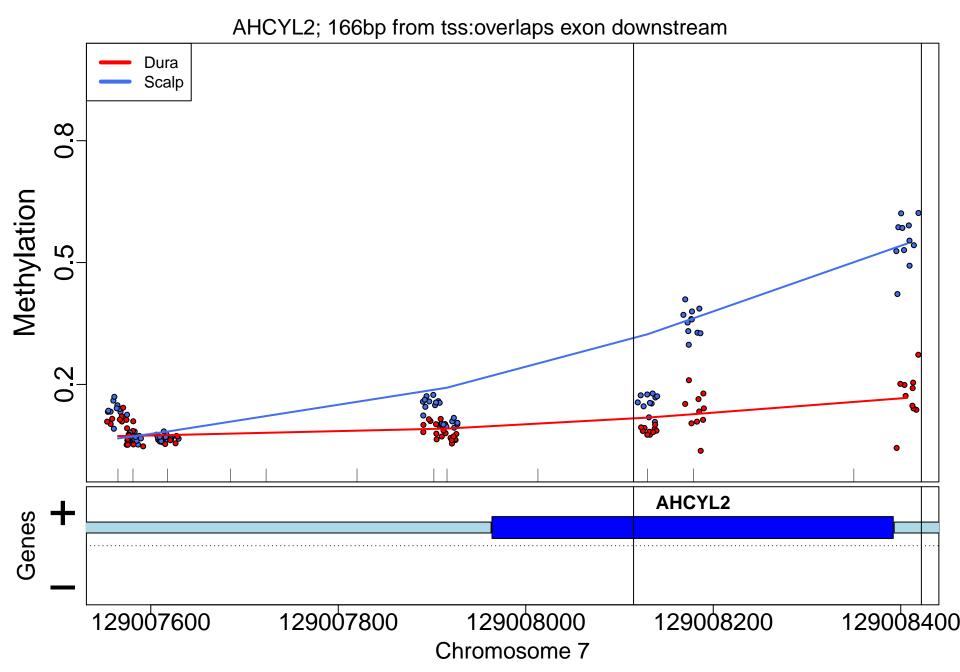


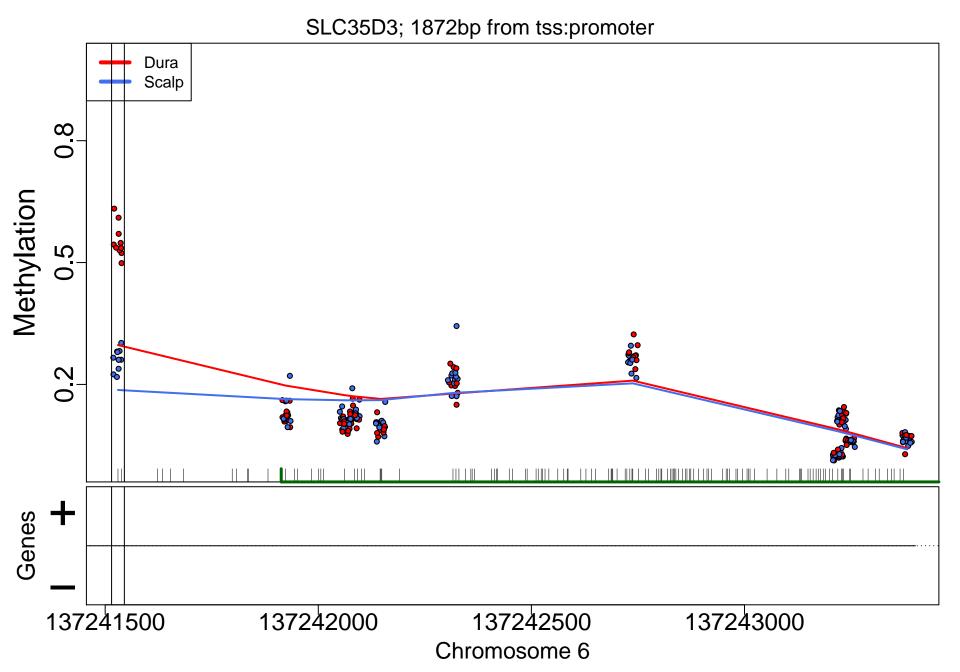


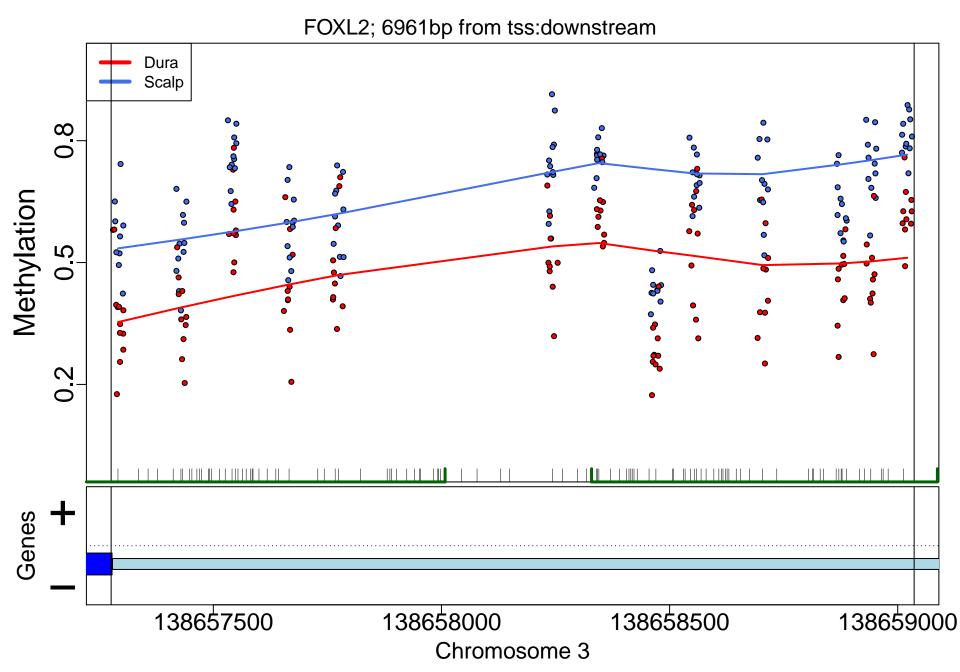


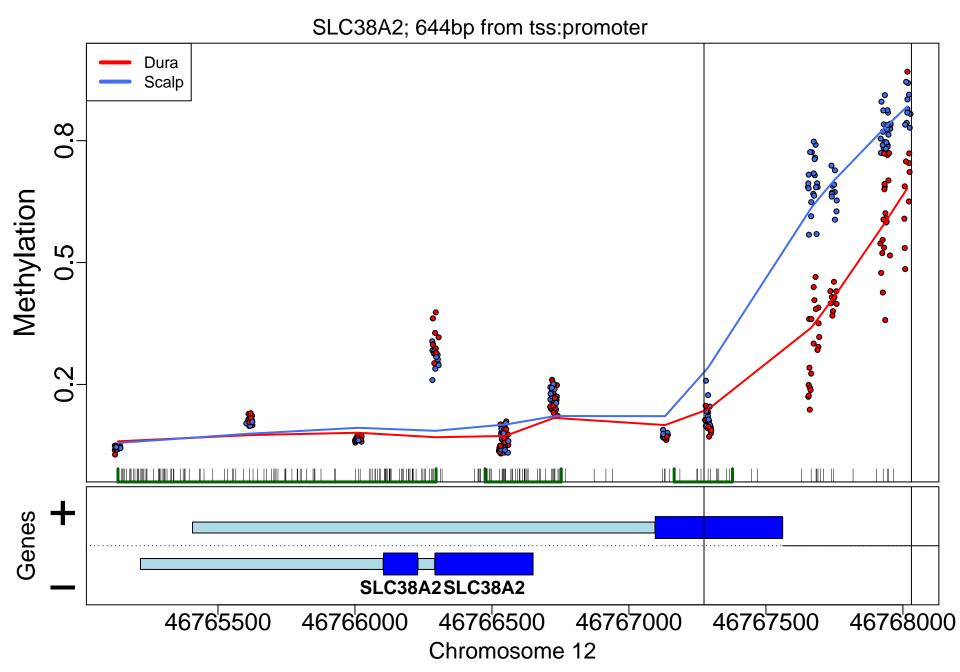


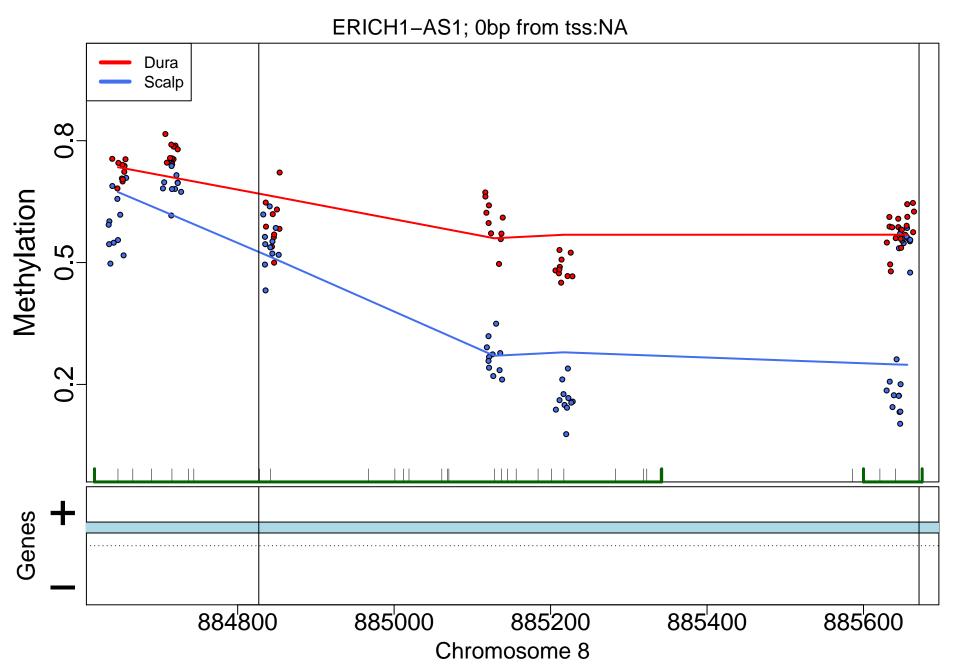


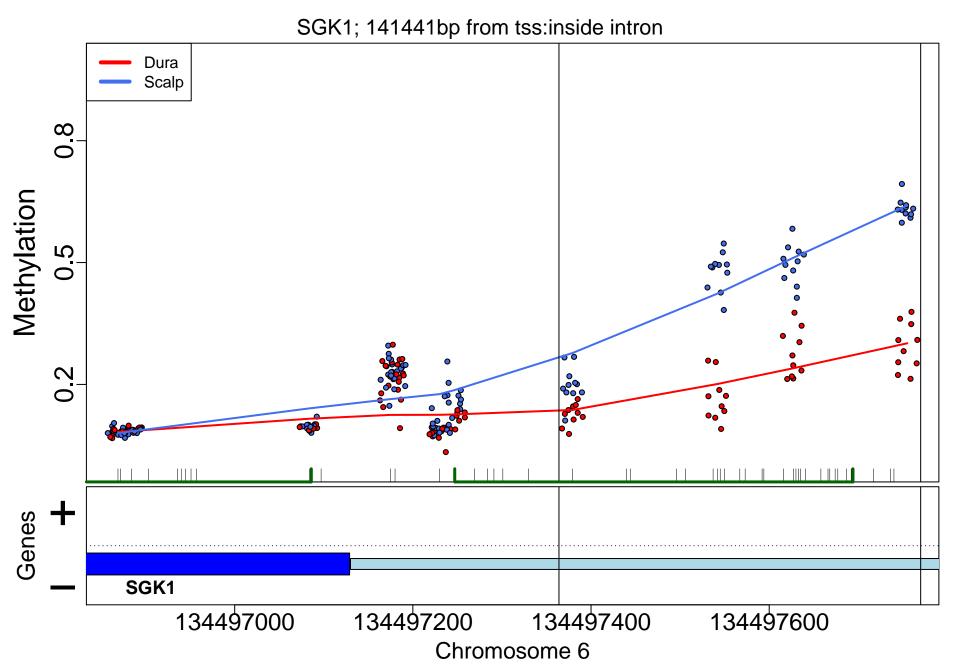


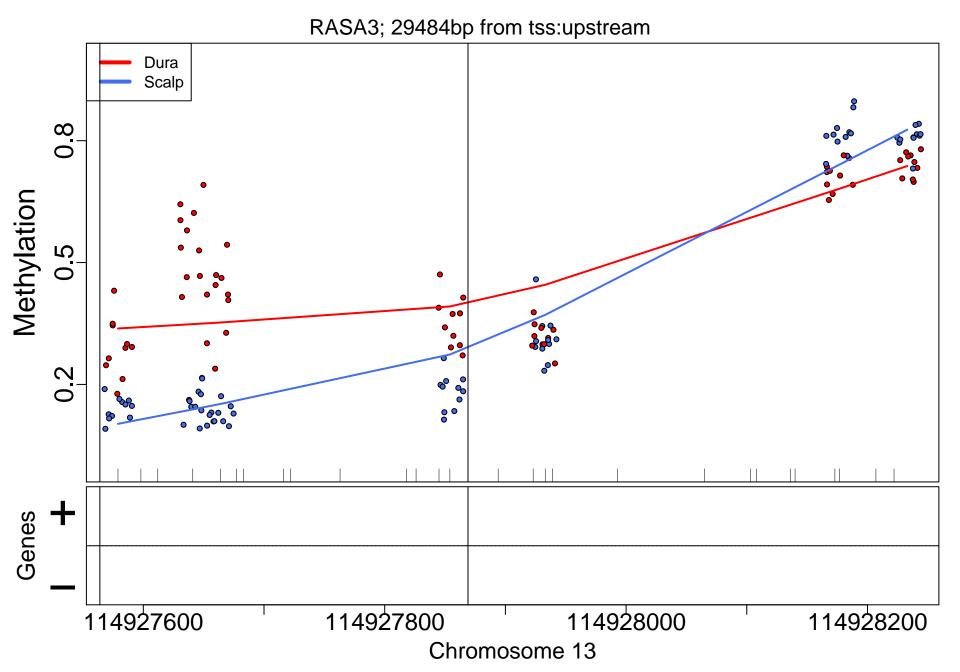


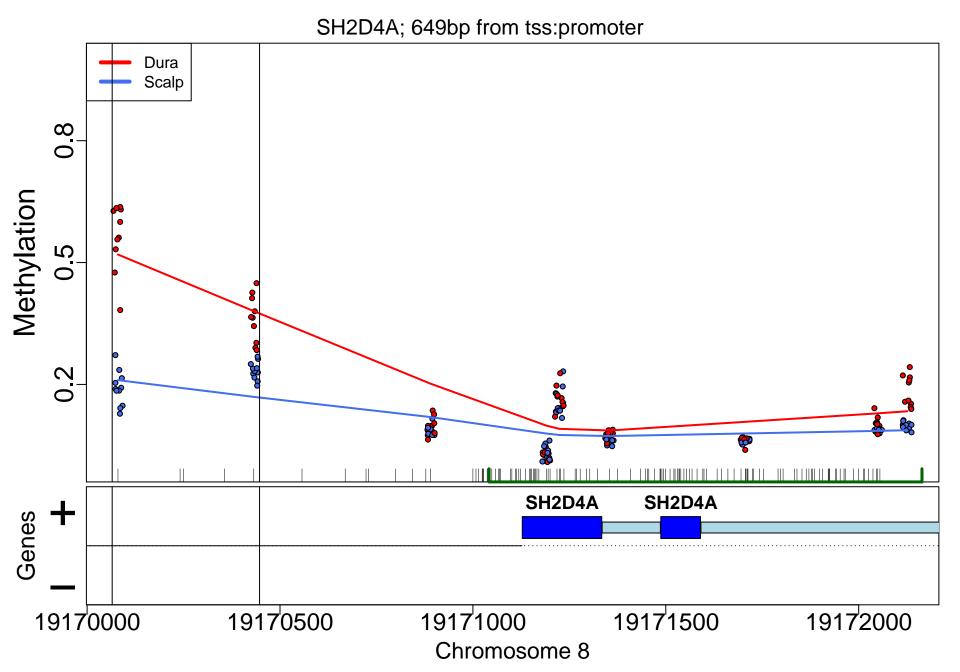


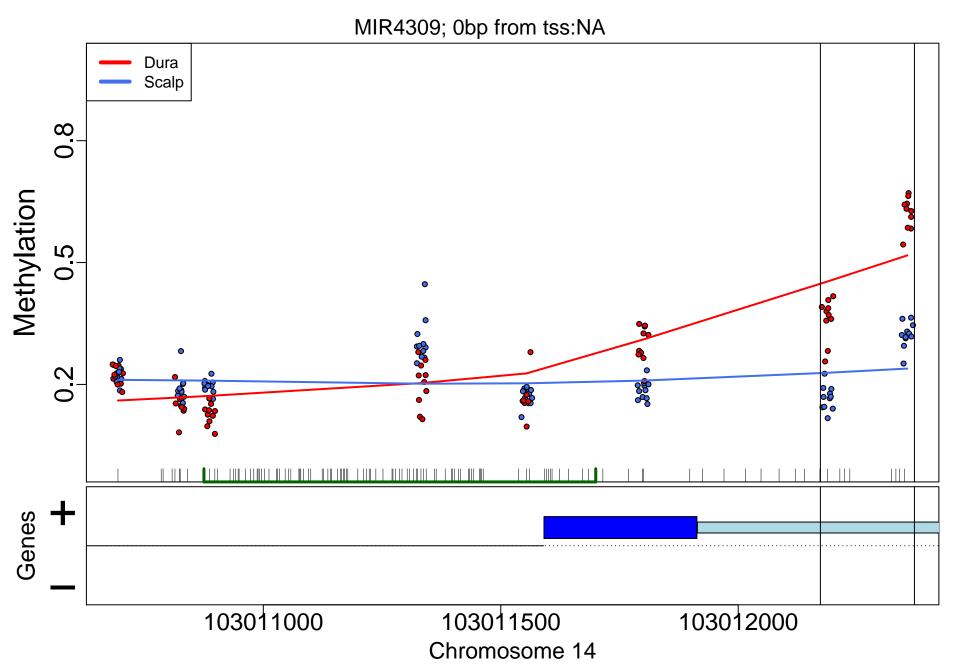


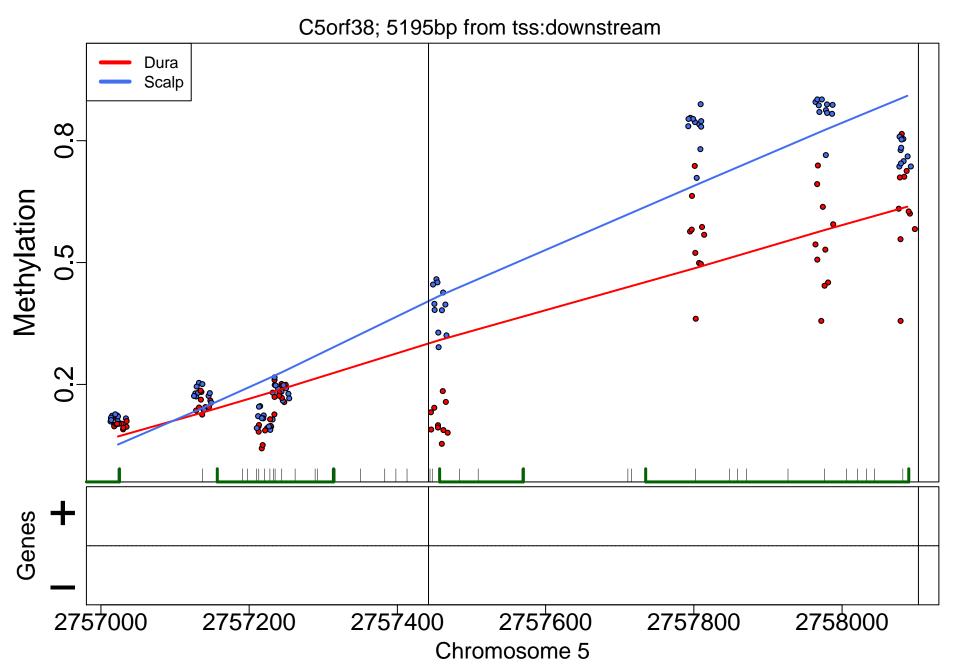


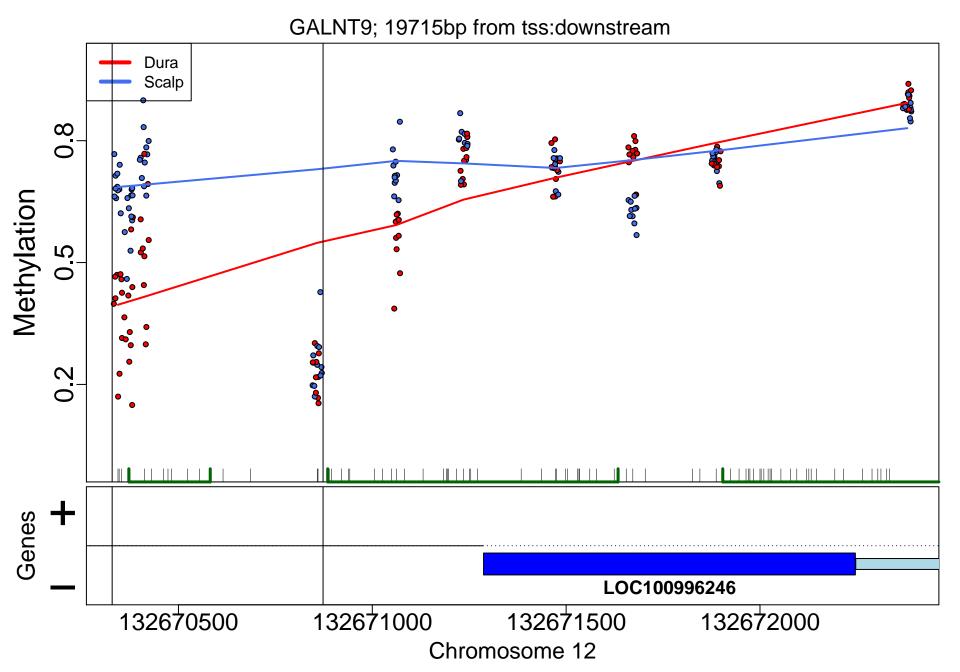


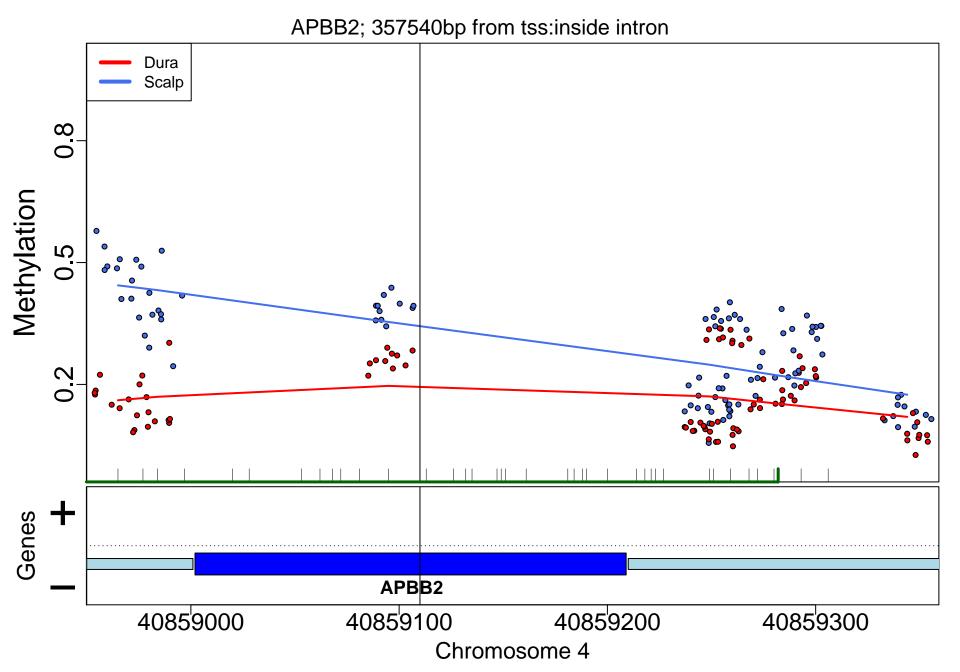


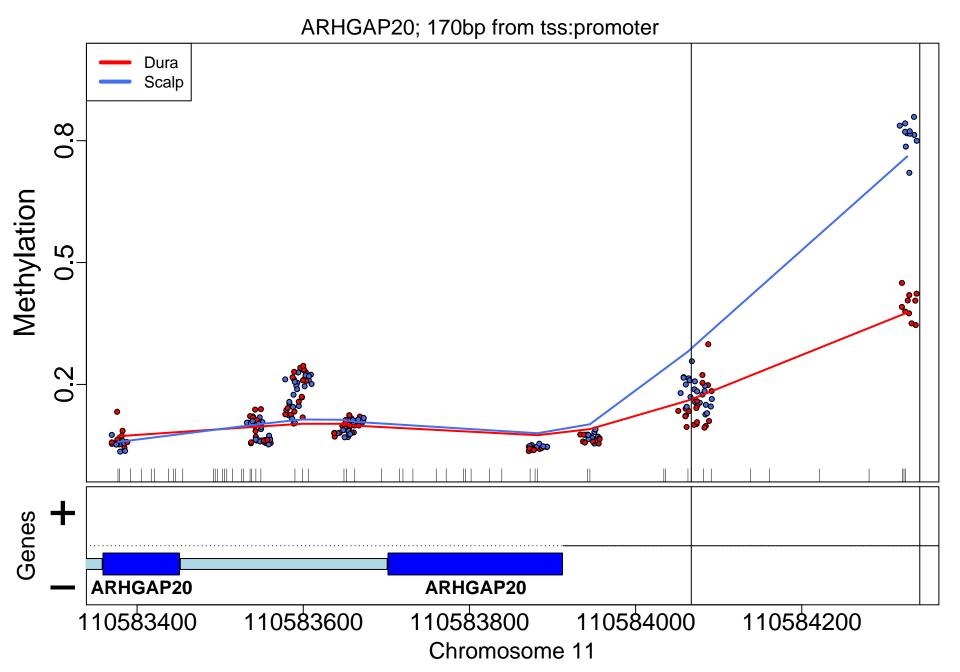


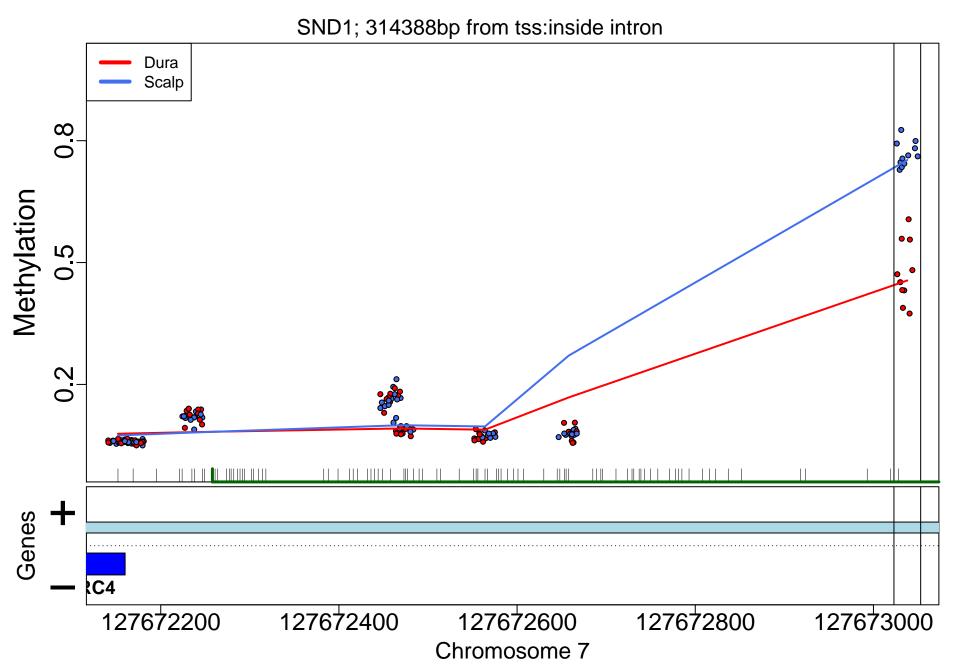


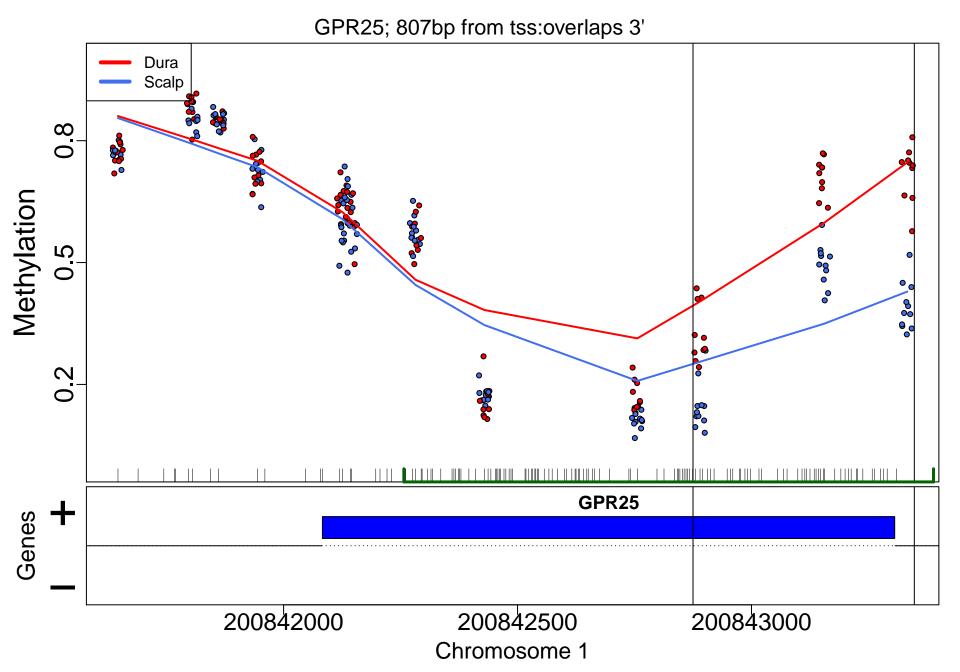


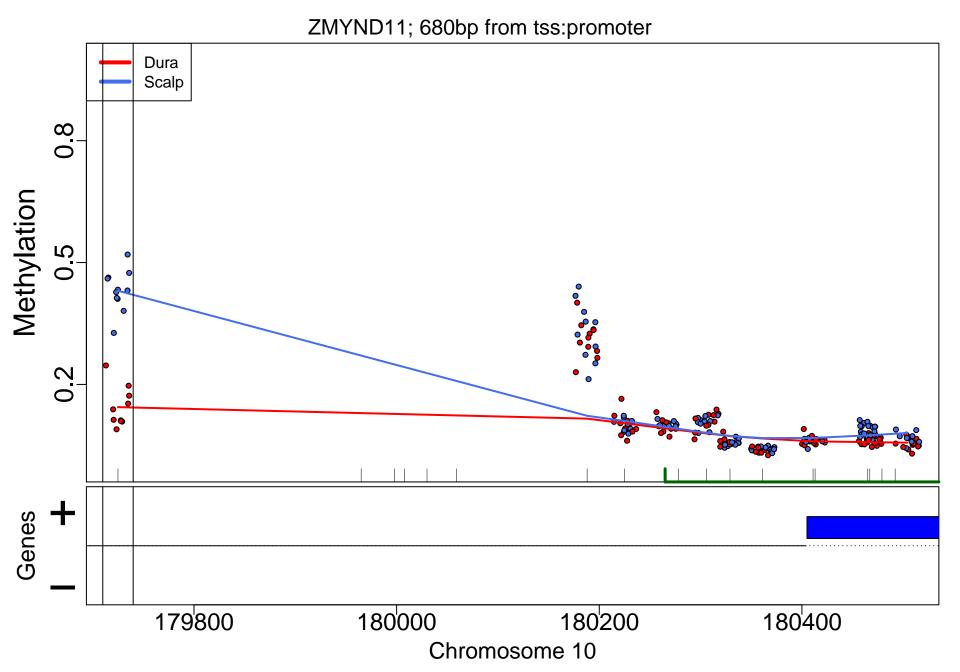


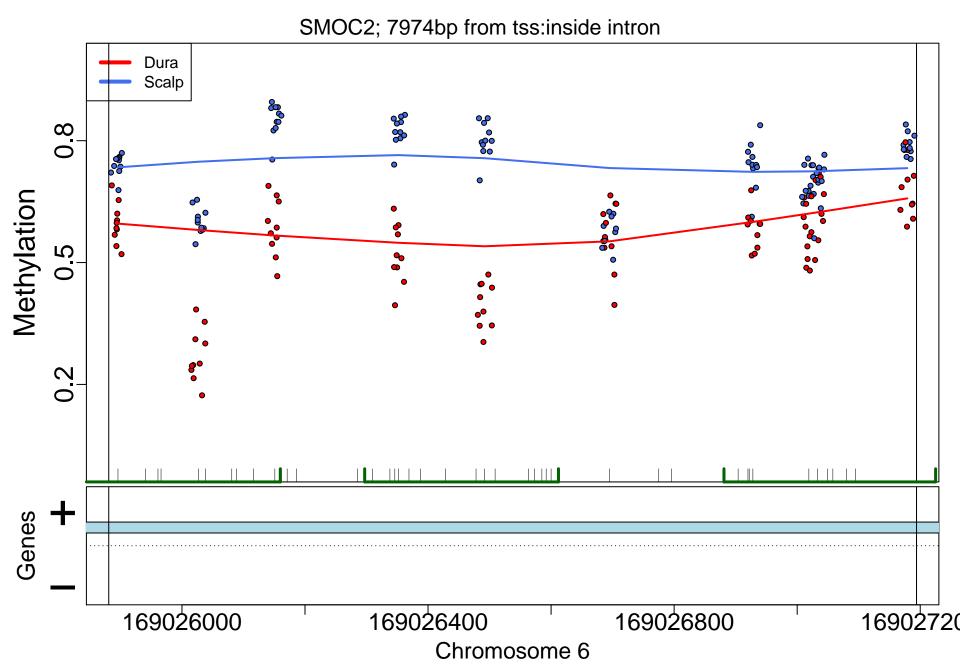


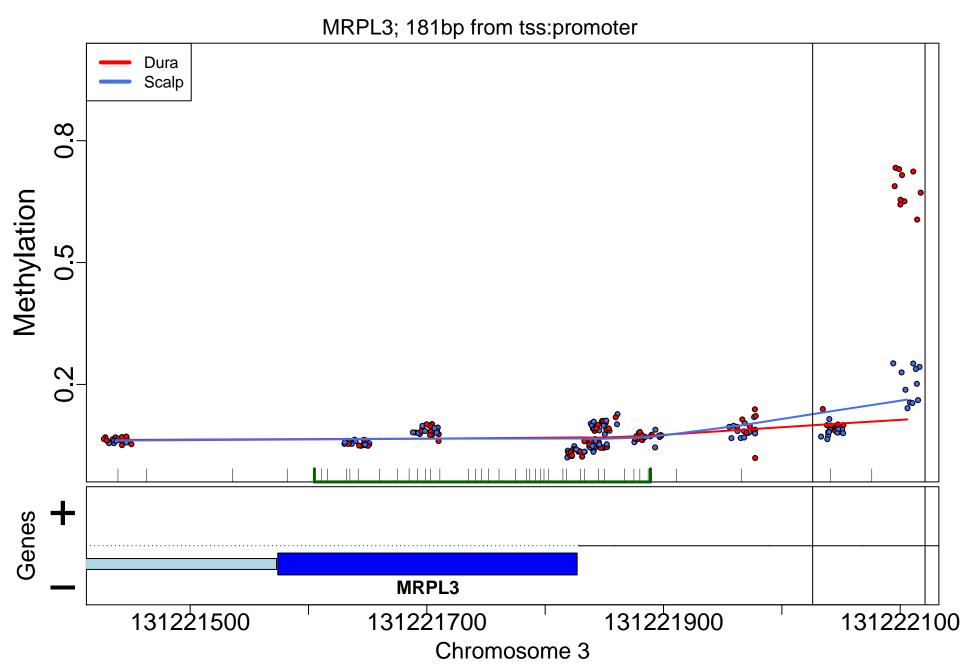


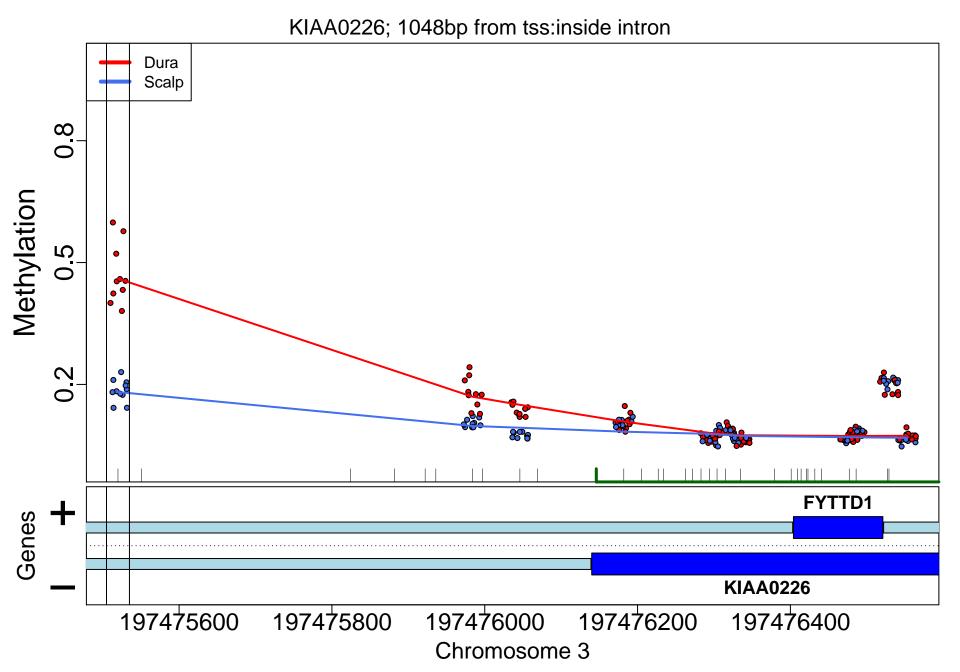


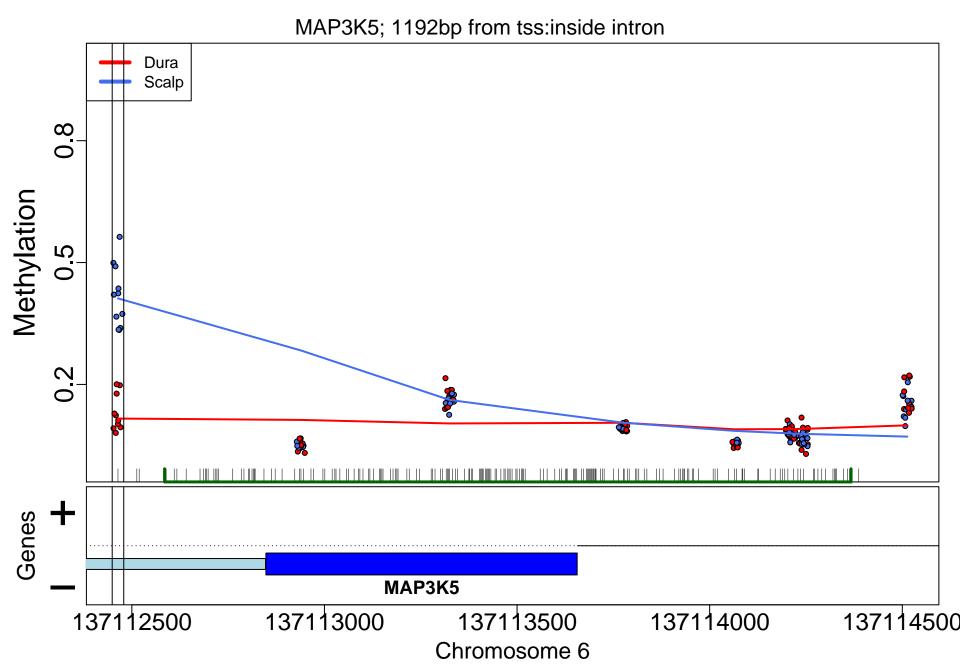


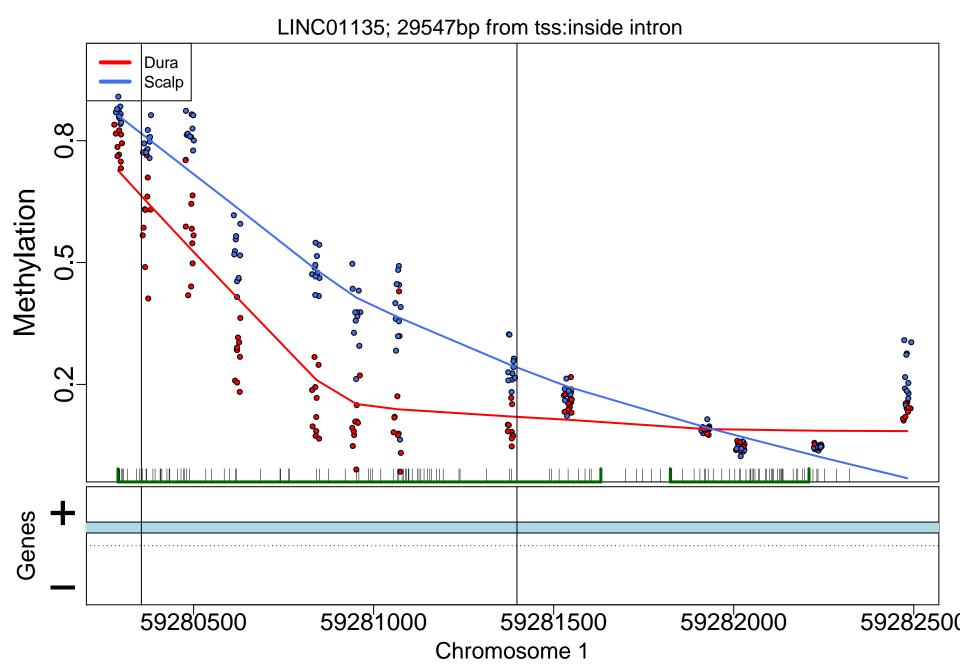


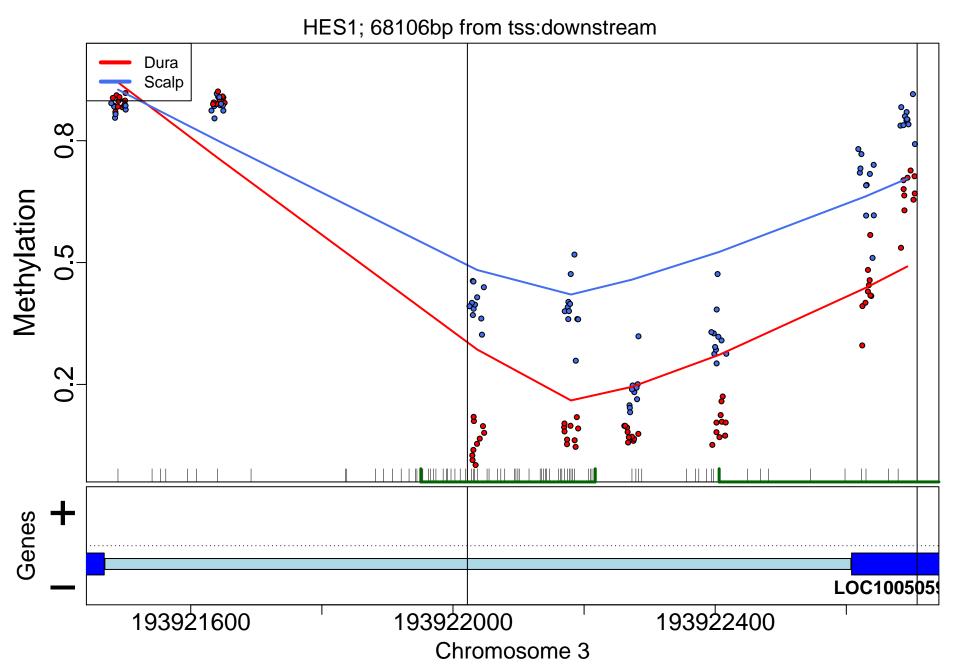


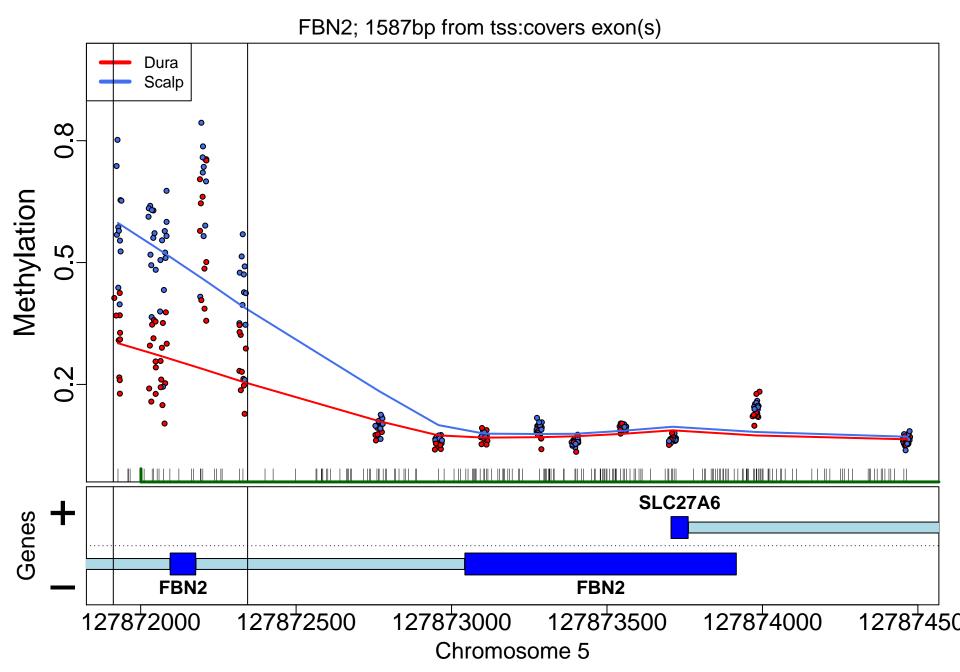


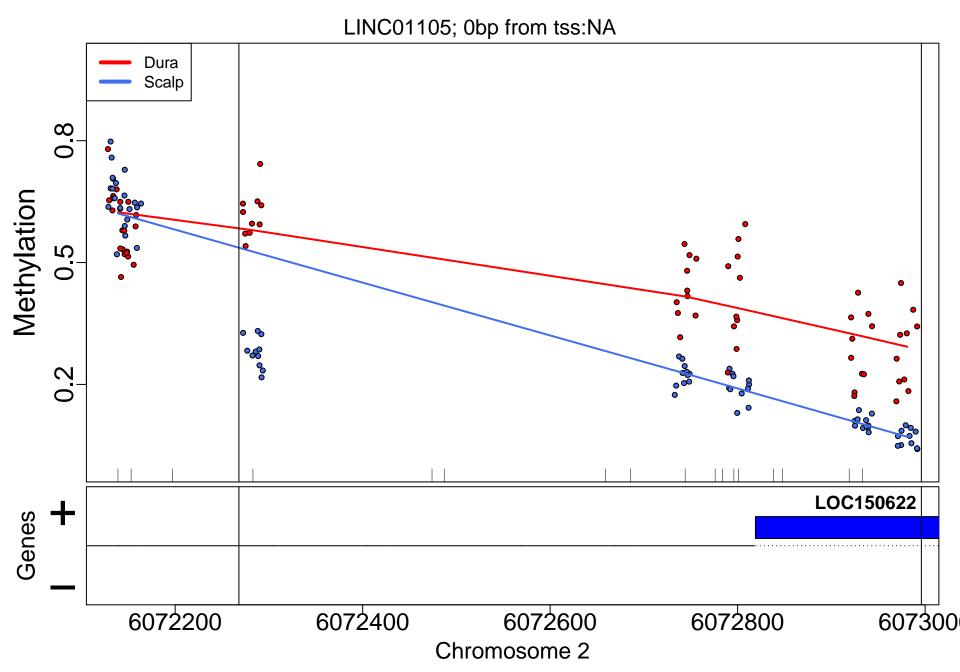


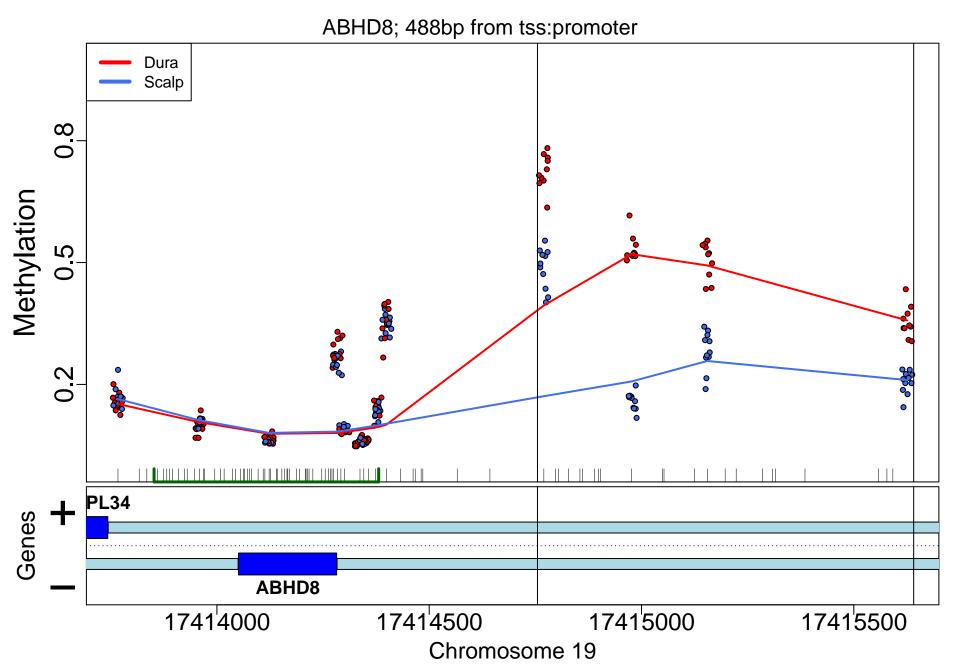


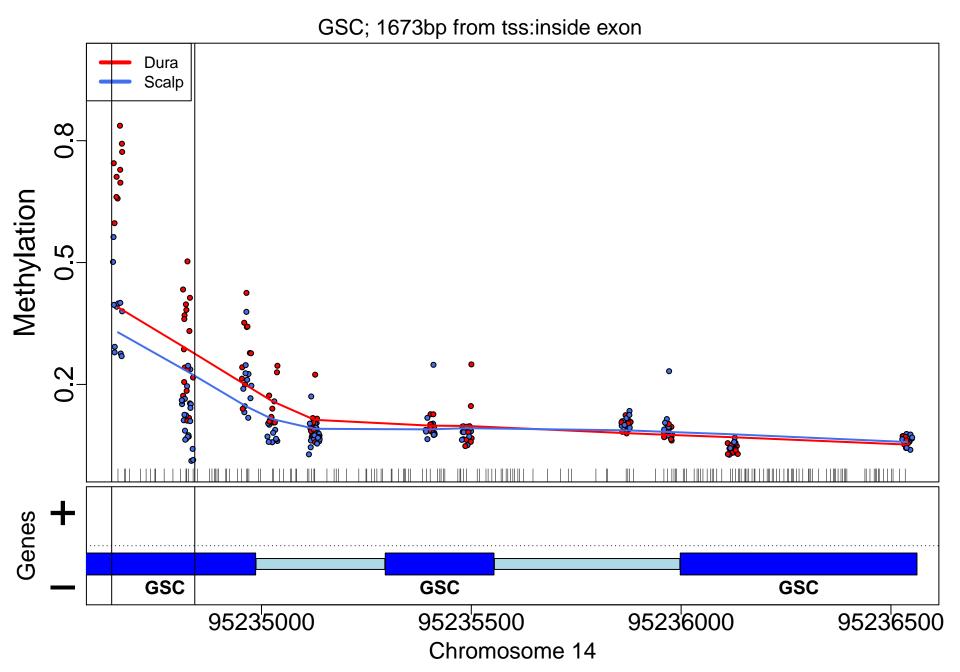


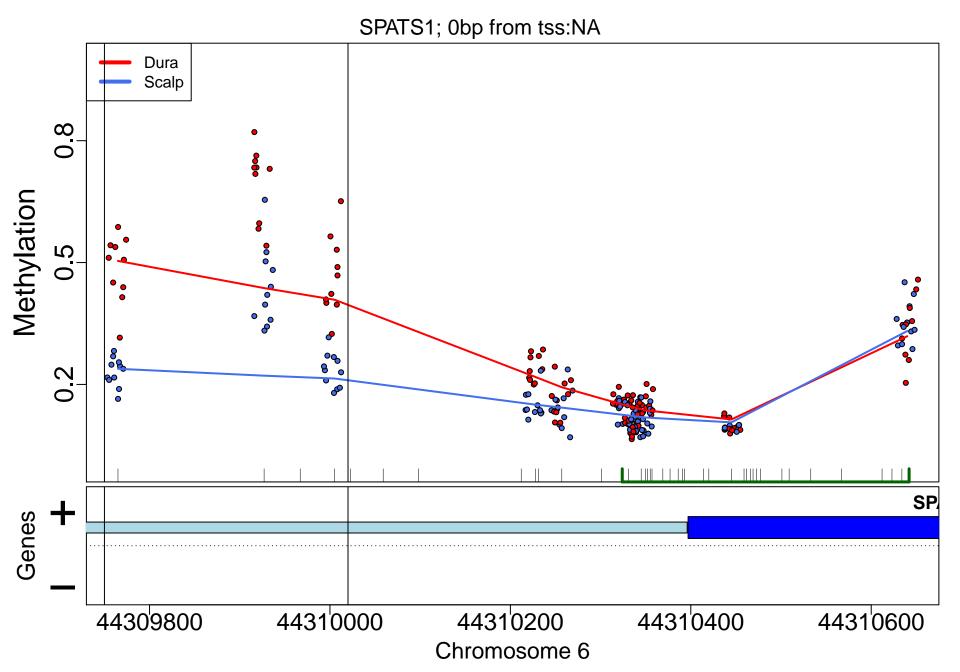


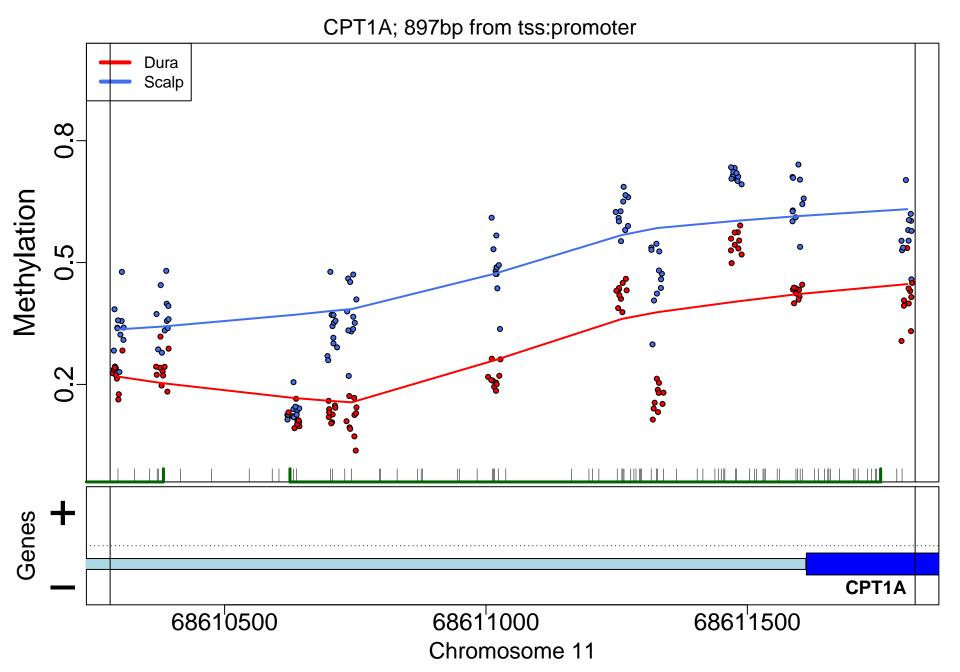


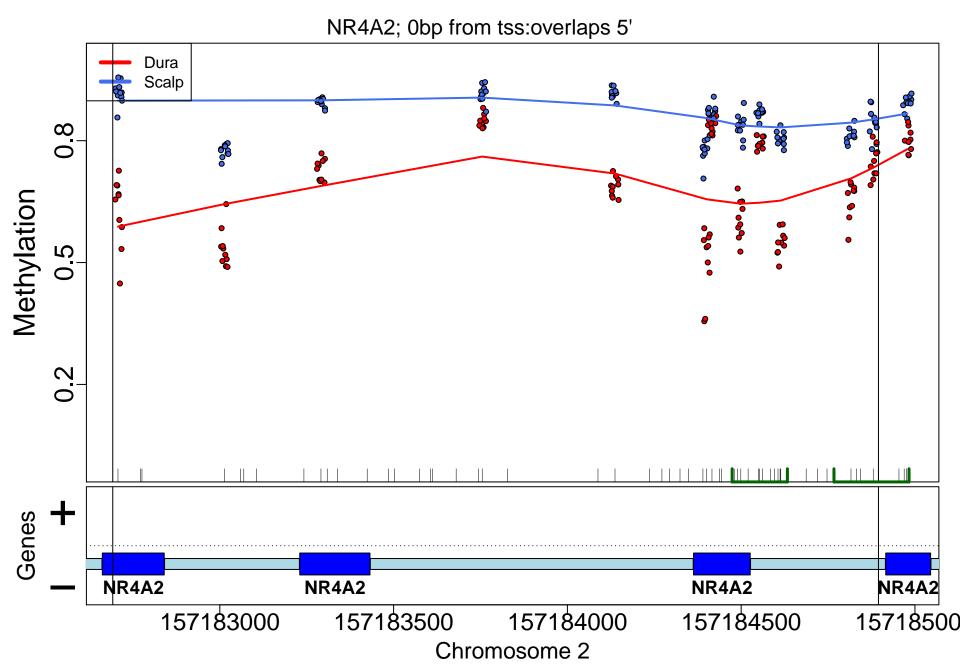


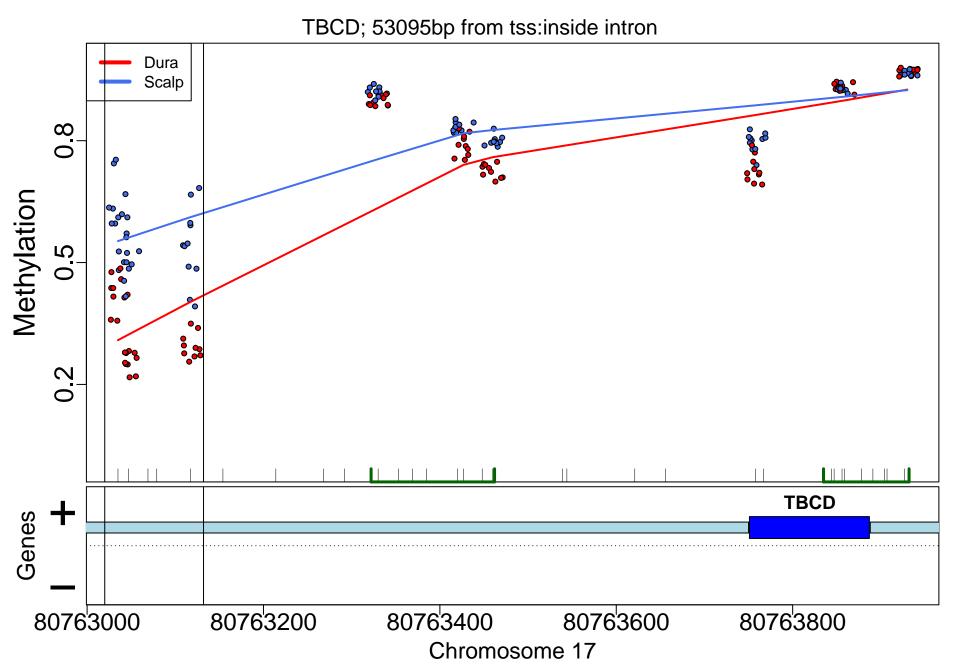


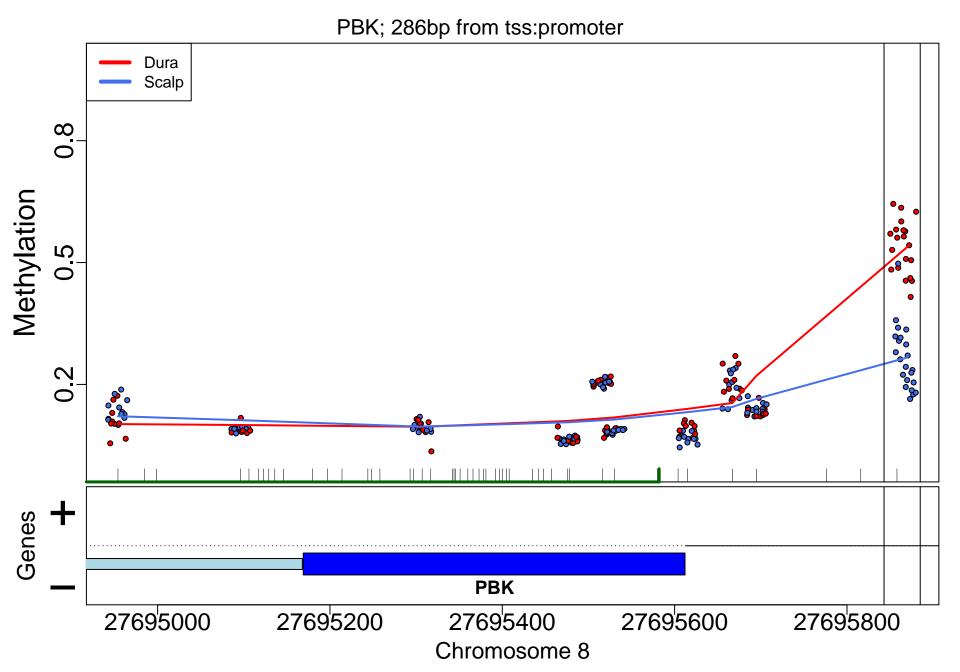


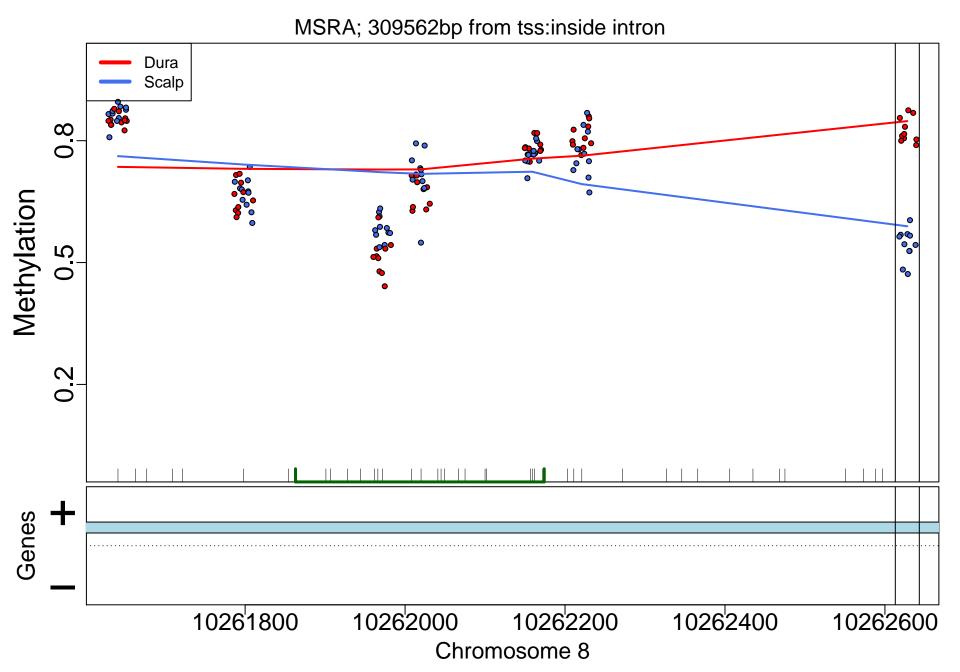




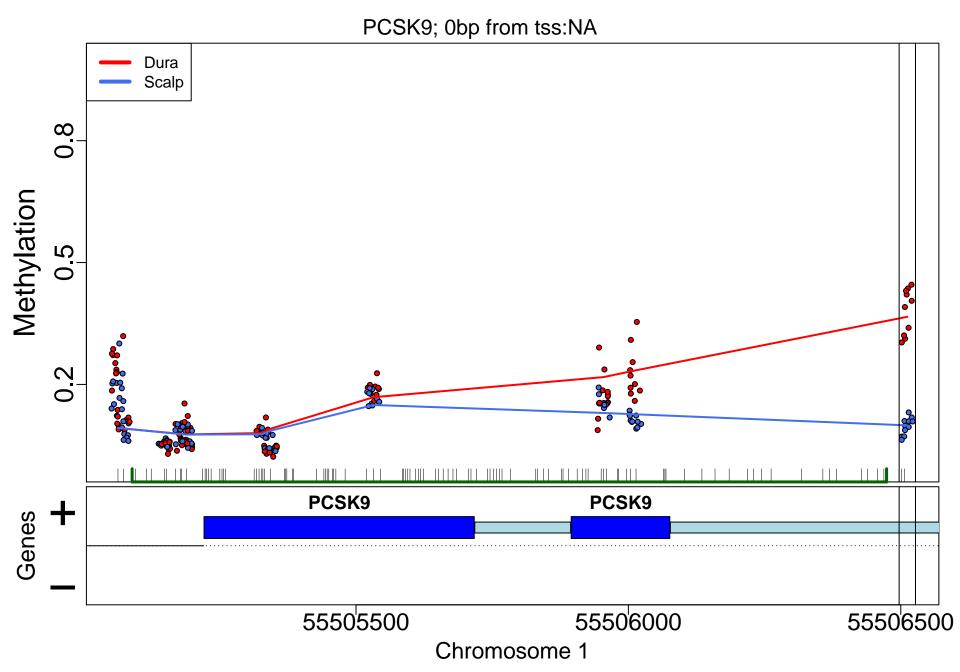


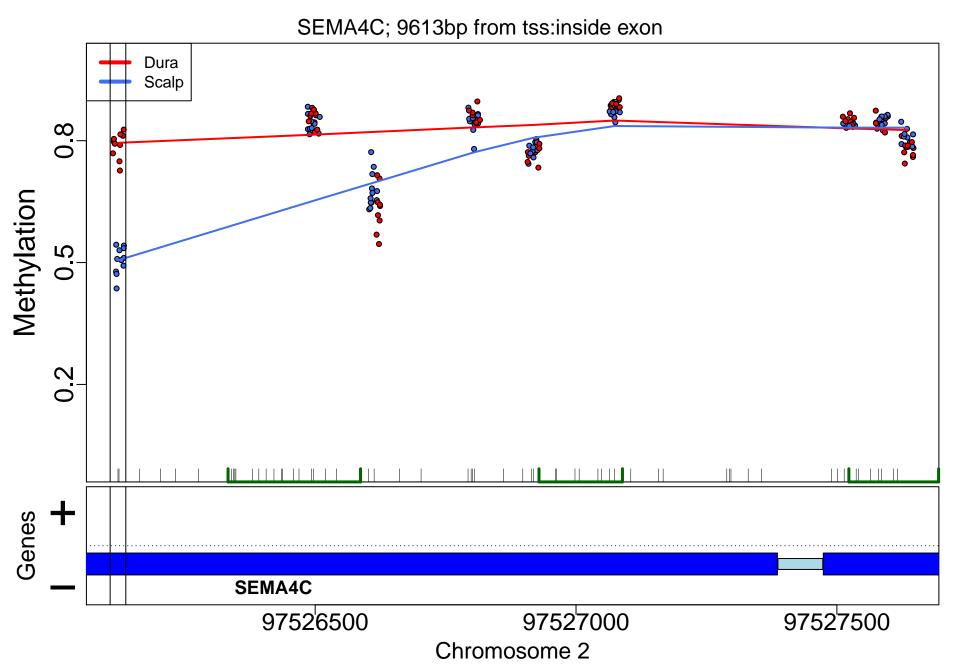


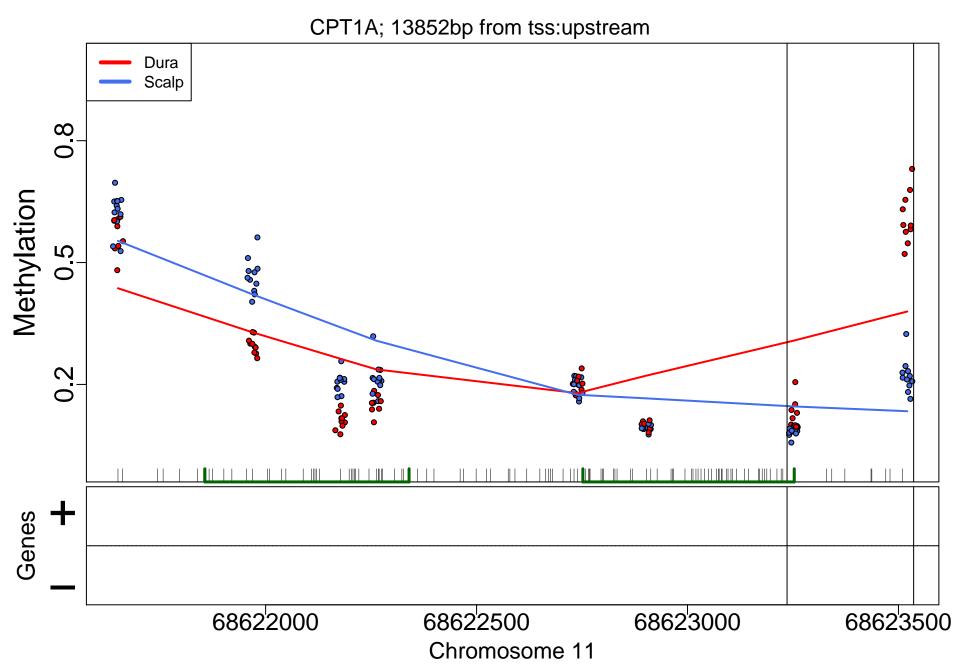


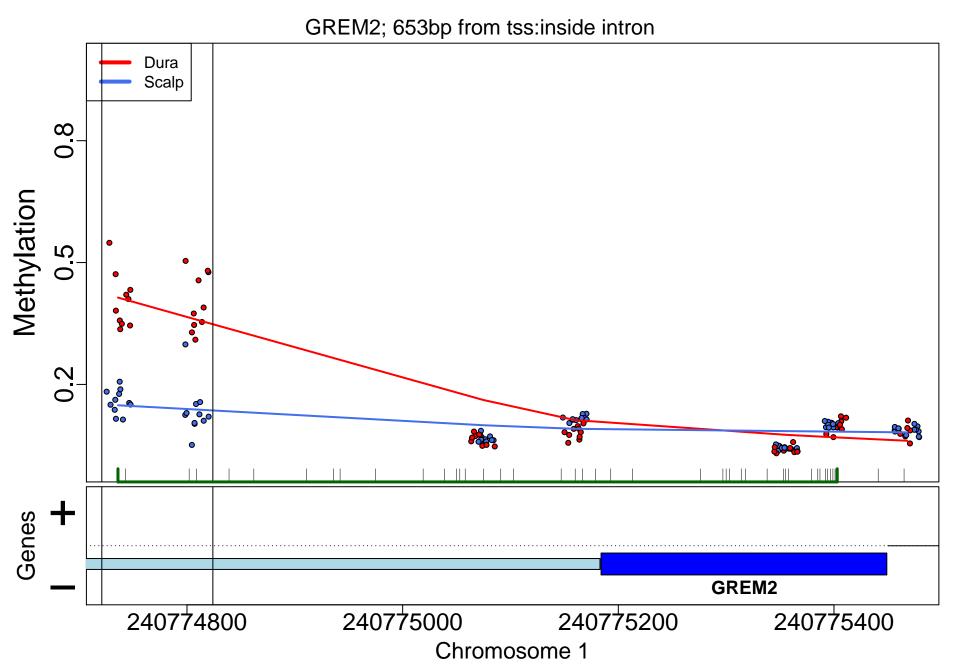


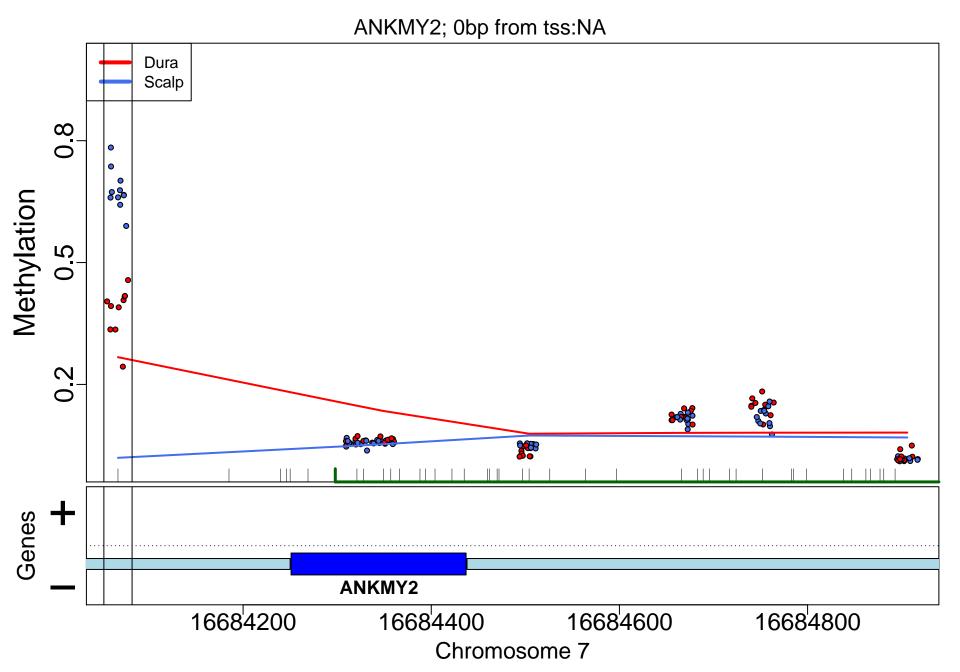
RASA3; 0bp from tss:NA Dura Scalp <u>∞</u>_ Methylation **%** • Genes 114861600 114862000 114861800 114862200 Chromosome 13

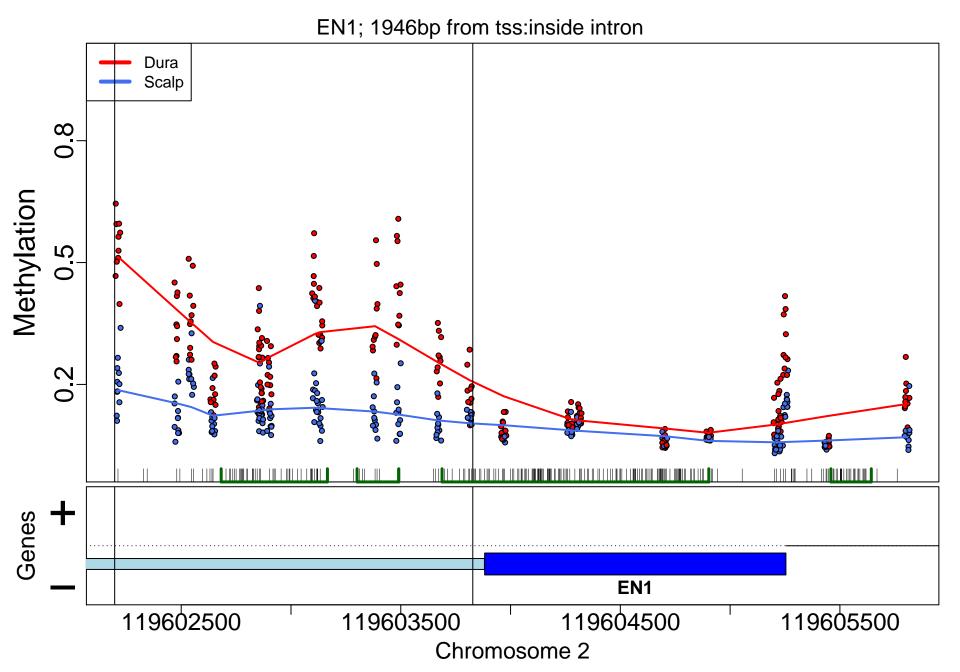


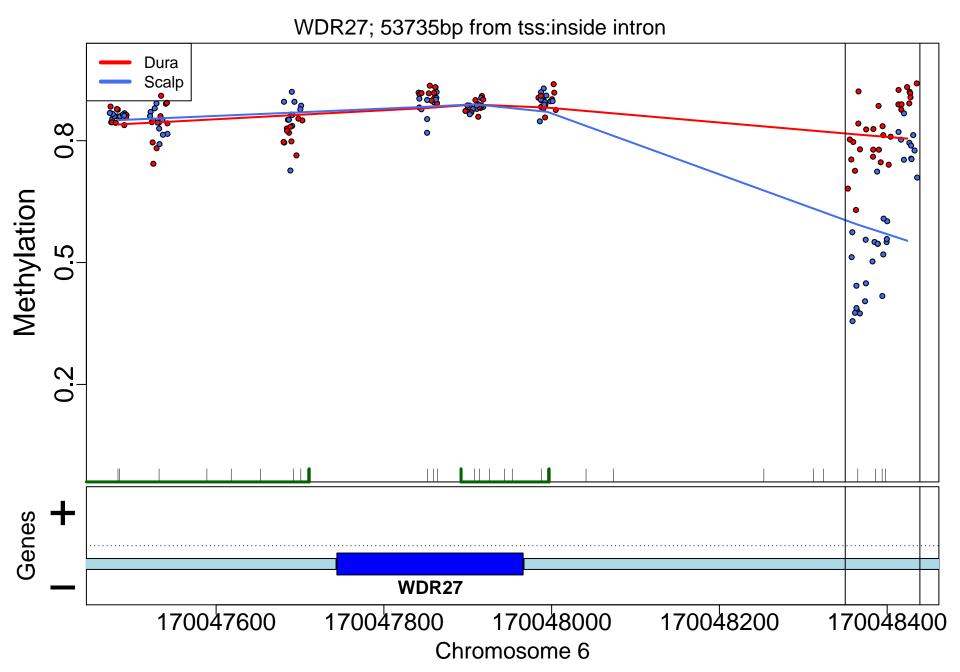


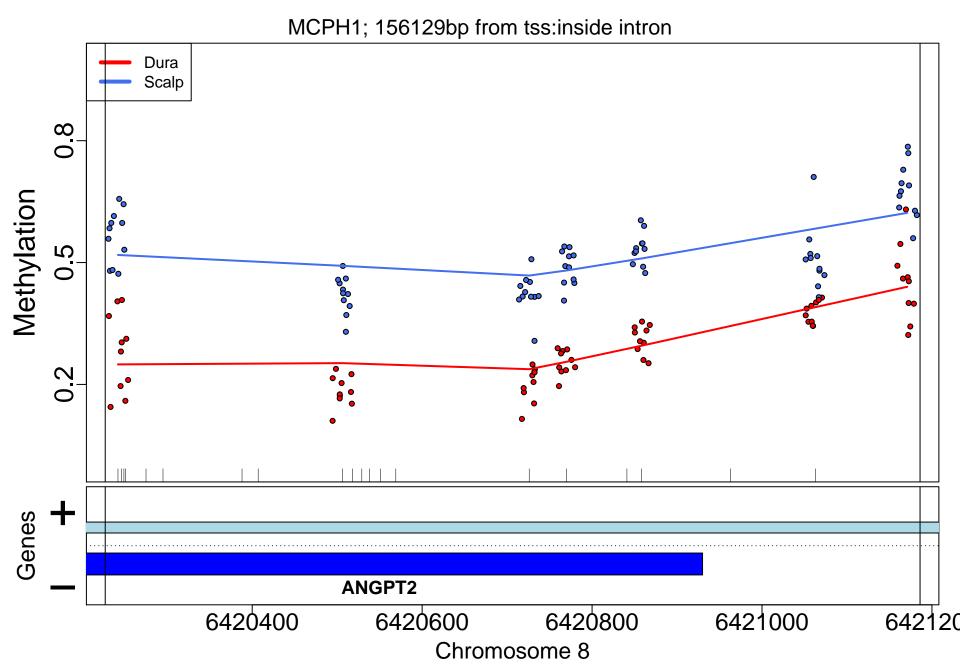


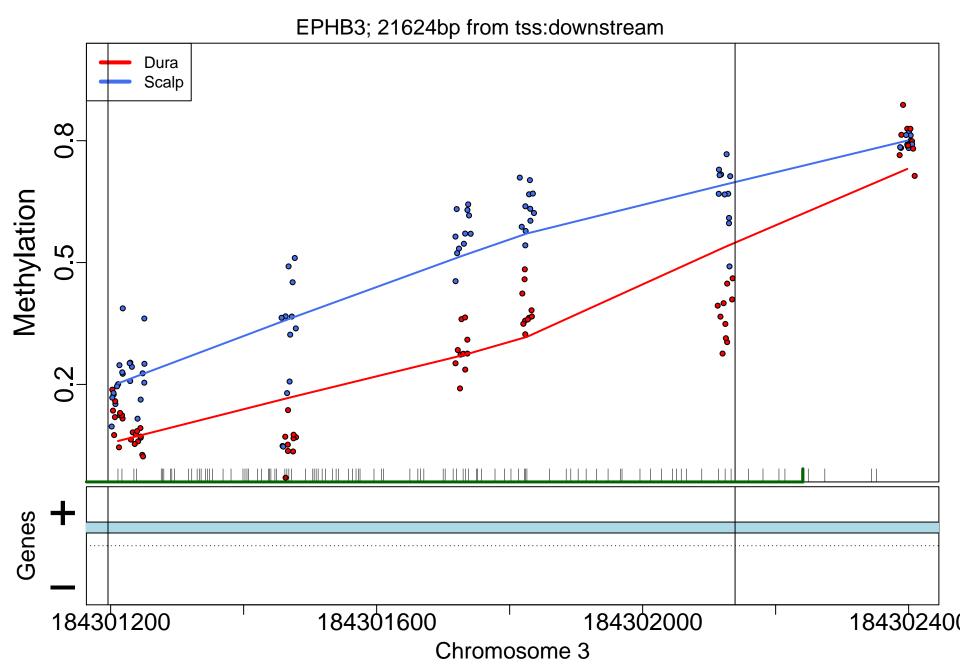


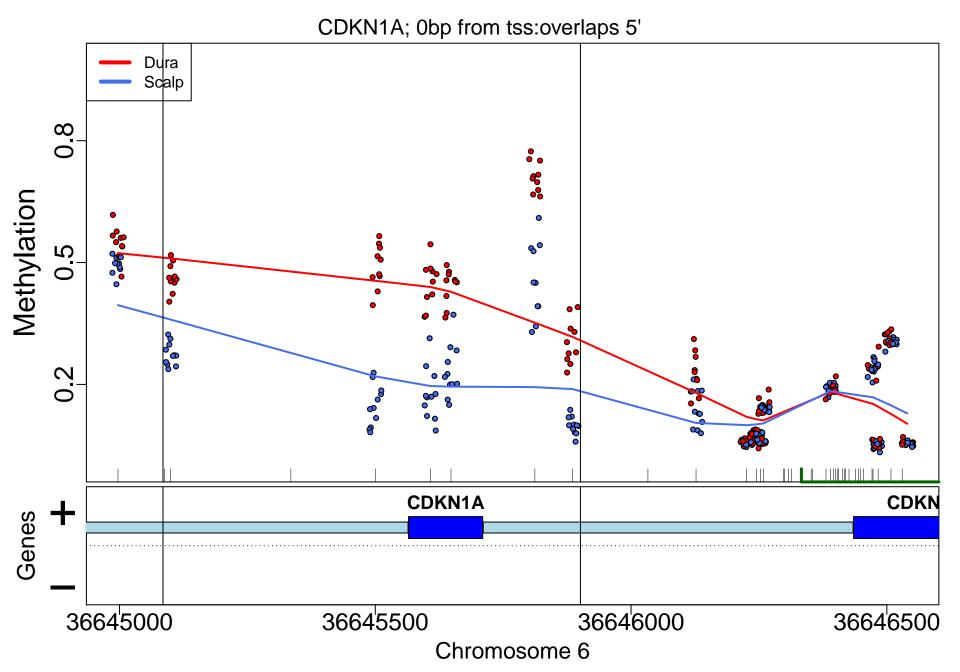


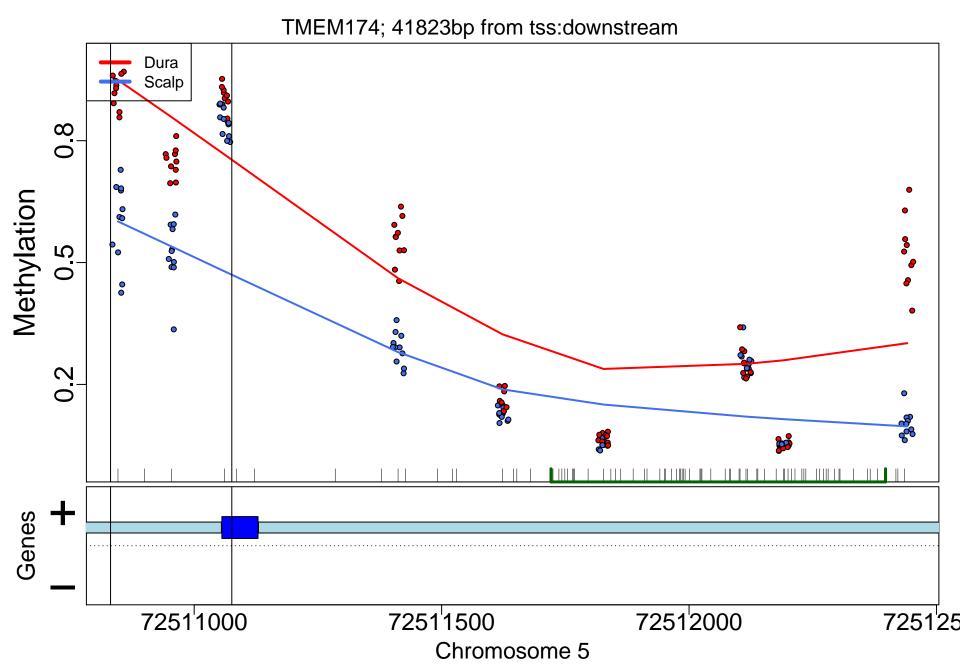


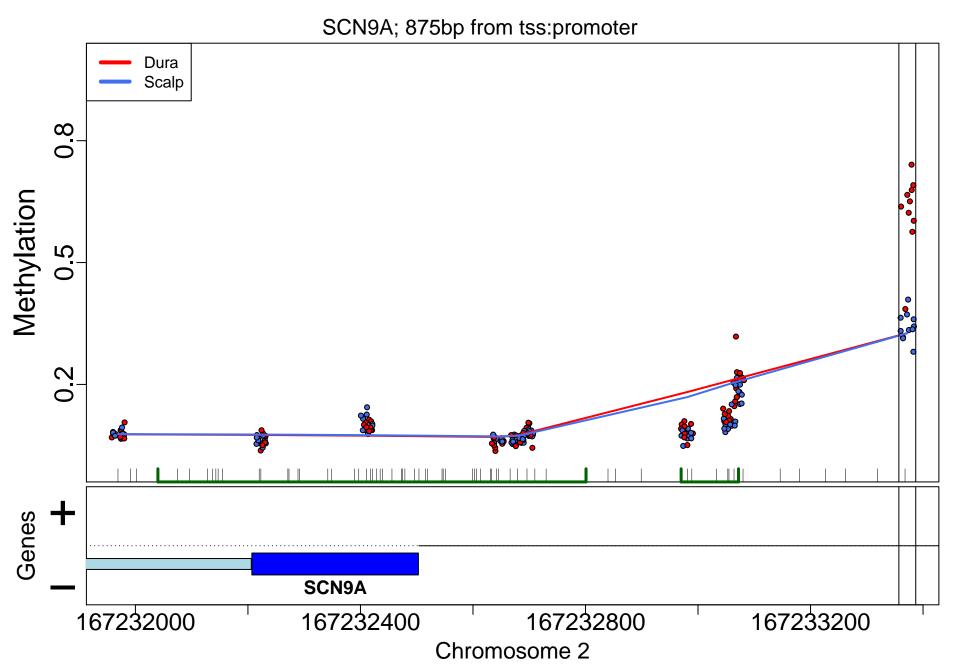


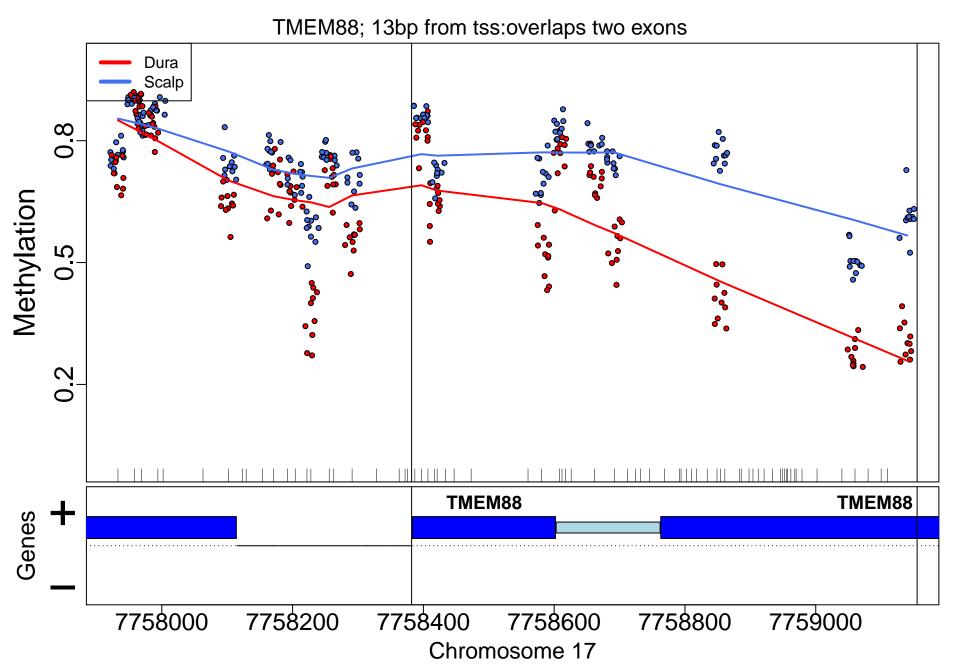


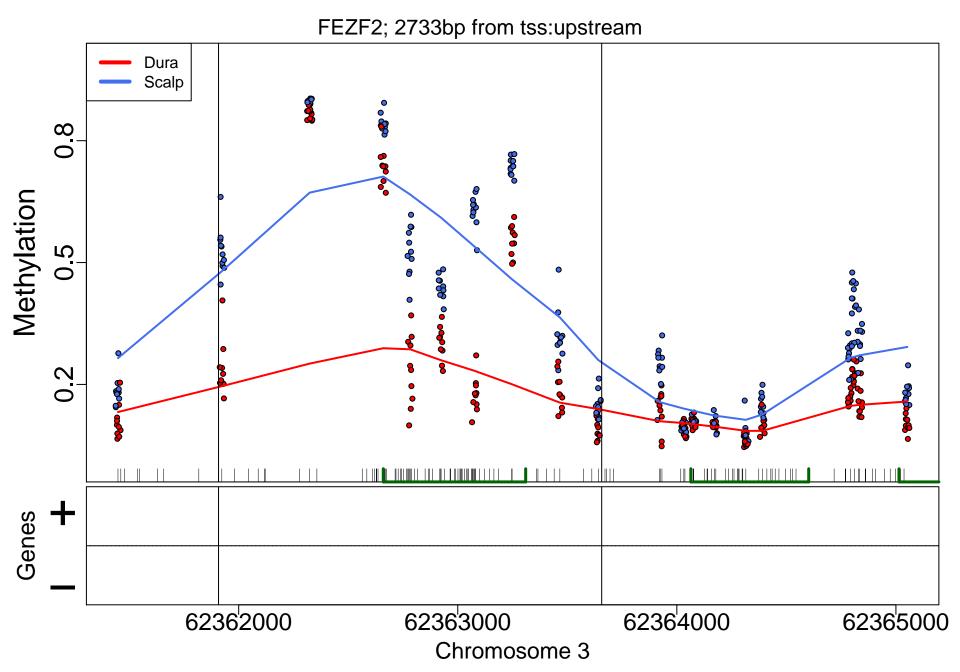


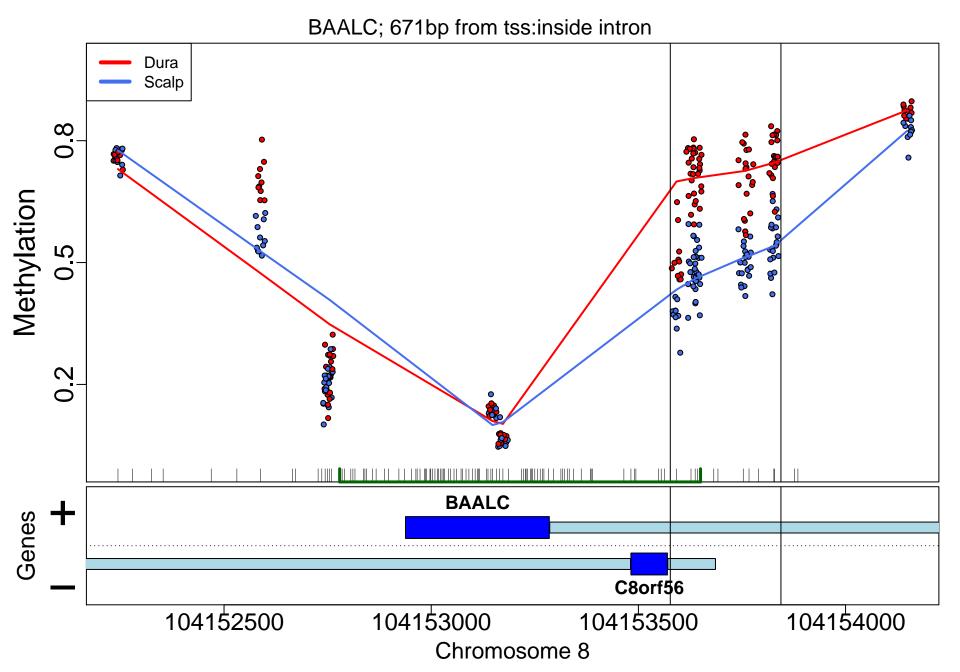




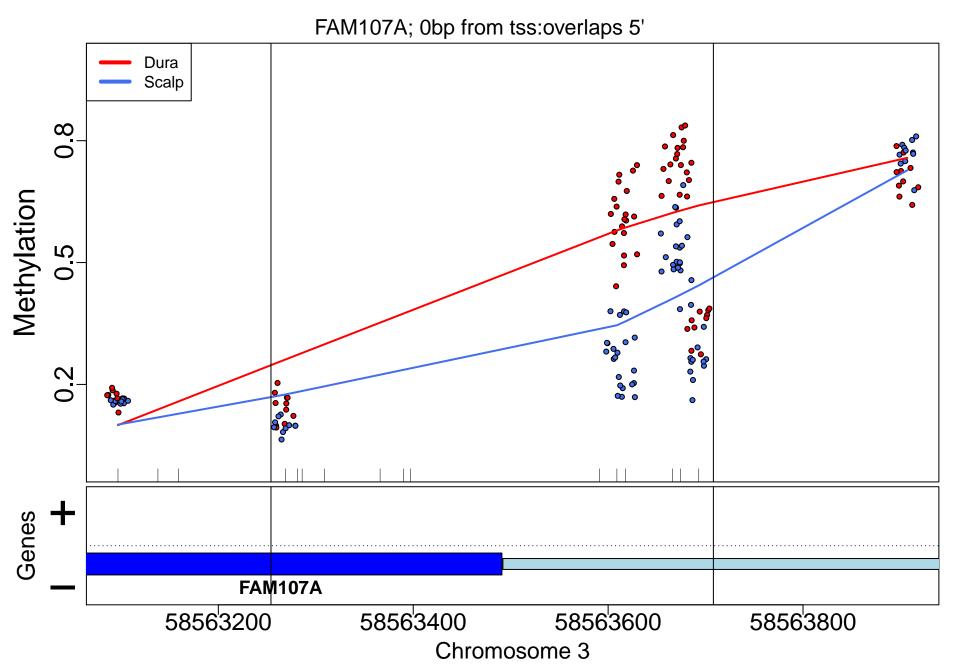


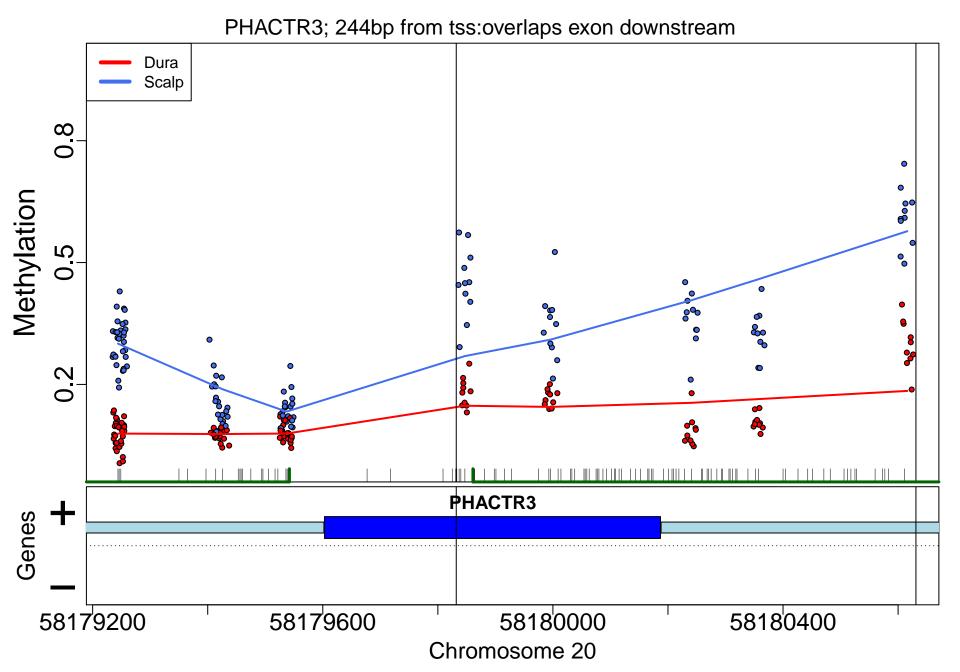


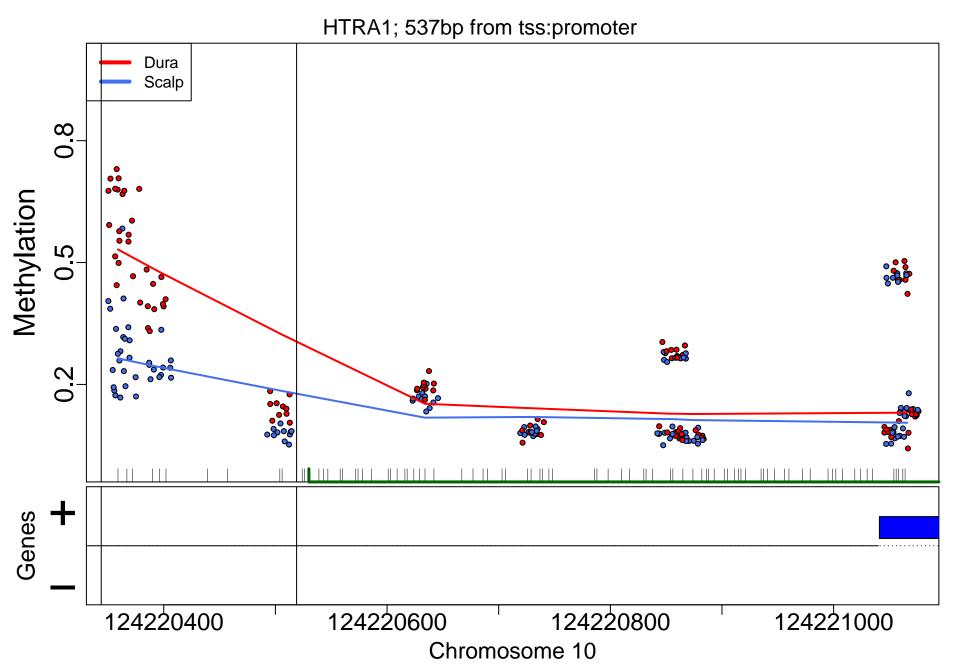




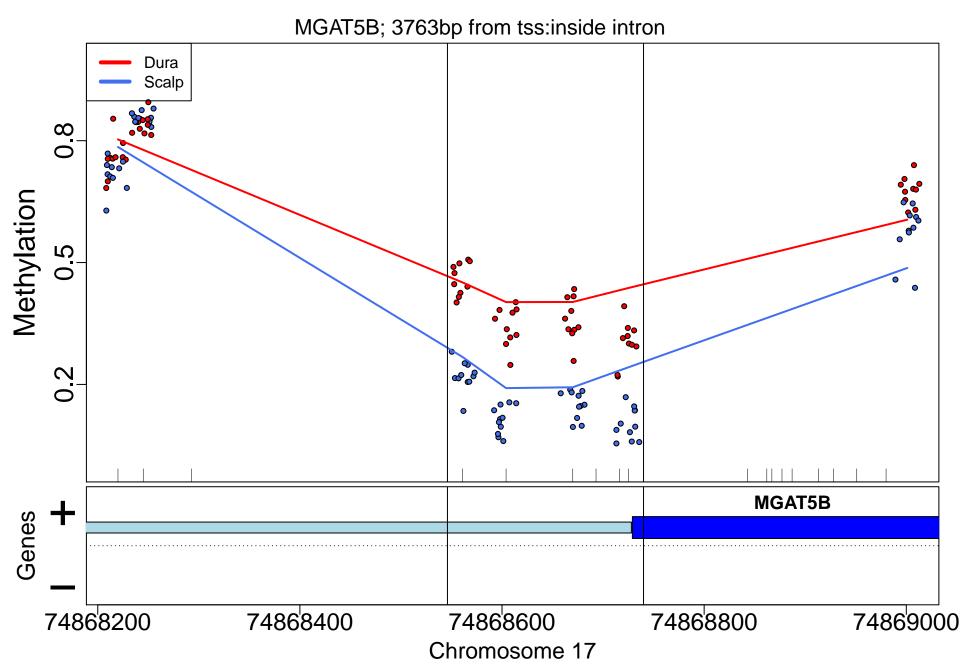
GABBR1; 18723bp from tss:overlaps exon upstream Dura Scalp 80. Methylation 0.2 Genes **GABBR1** 2958 1000 29580900 2958 1300 2958¹1400 2958 1100 2958 1200 Chromosome 6

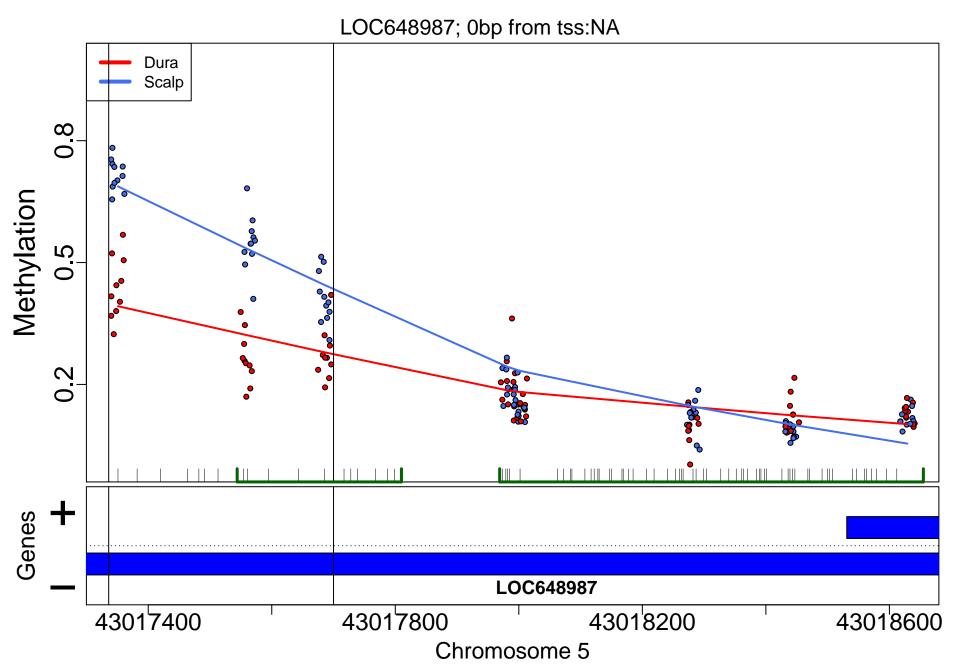


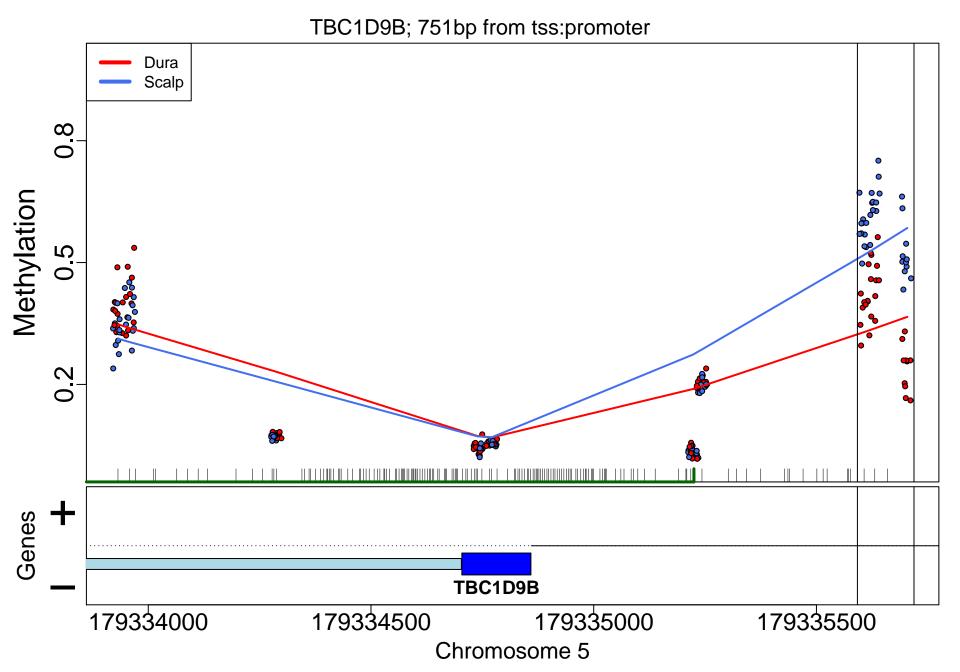


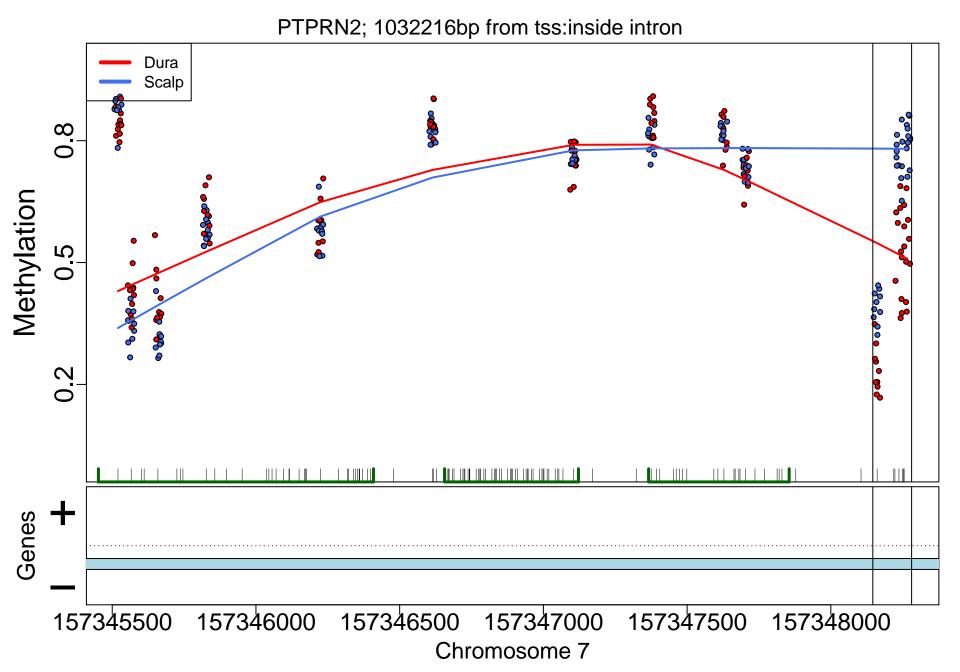


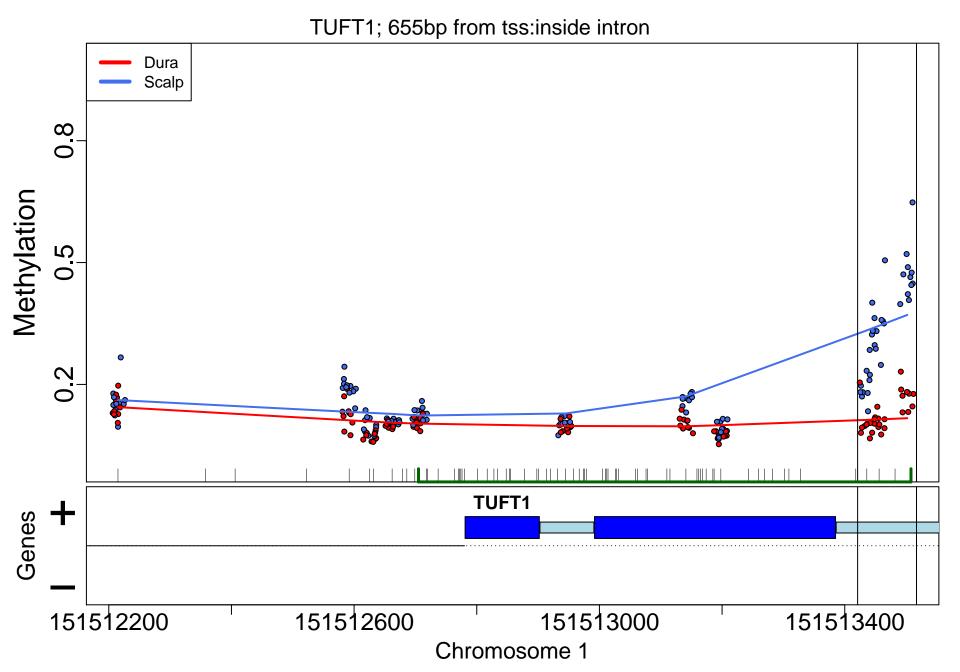
MSX1; 11871bp from tss:downstream Dura Scalp <u>∞</u>. Methylation 0.2 800 Genes 4873600 4873400 4873800 4874000 4874200 4874400 Chromosome 4

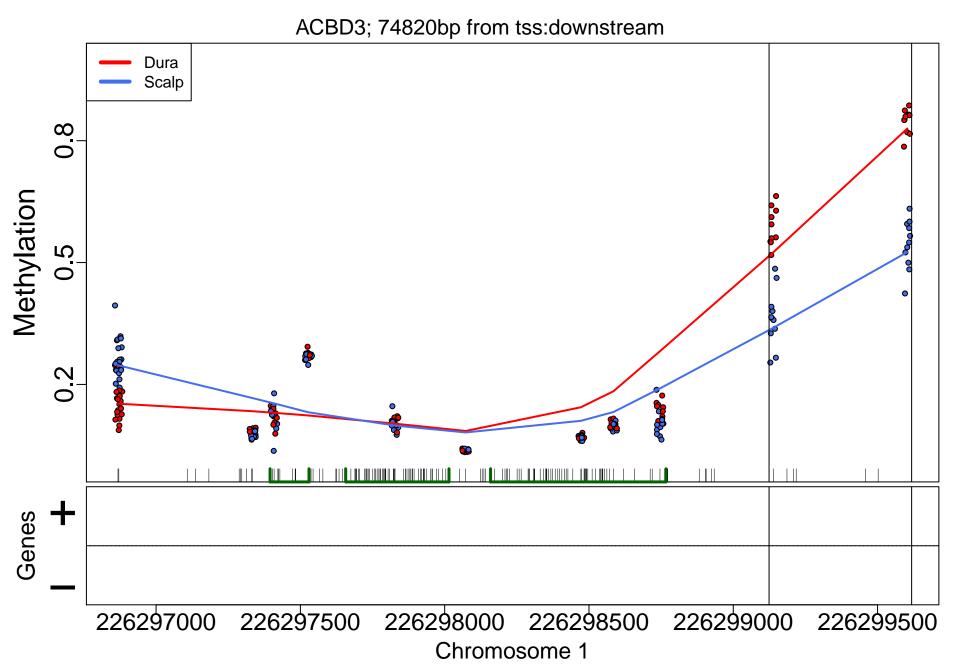


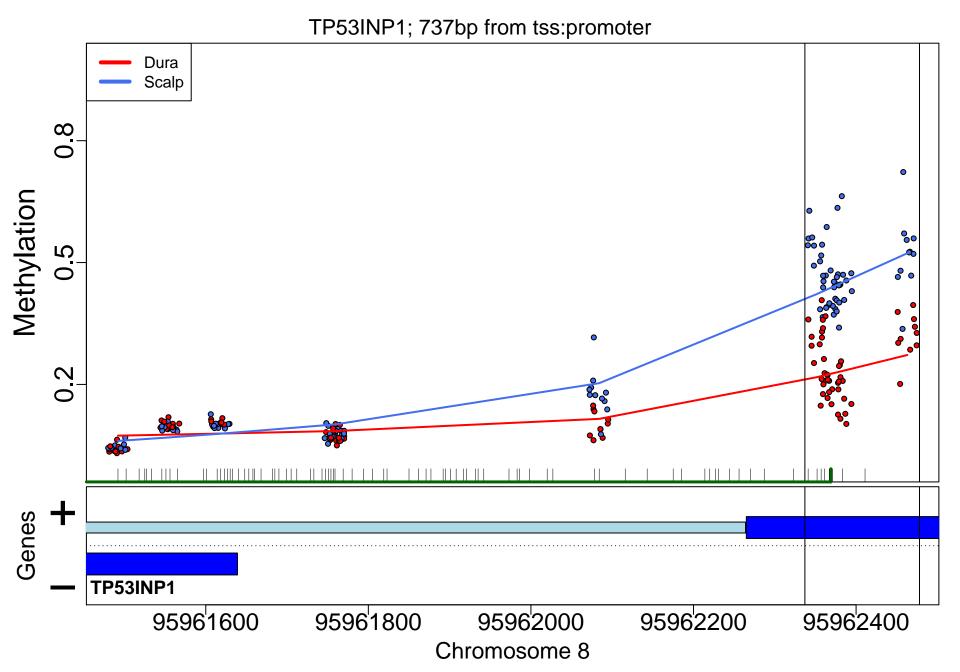


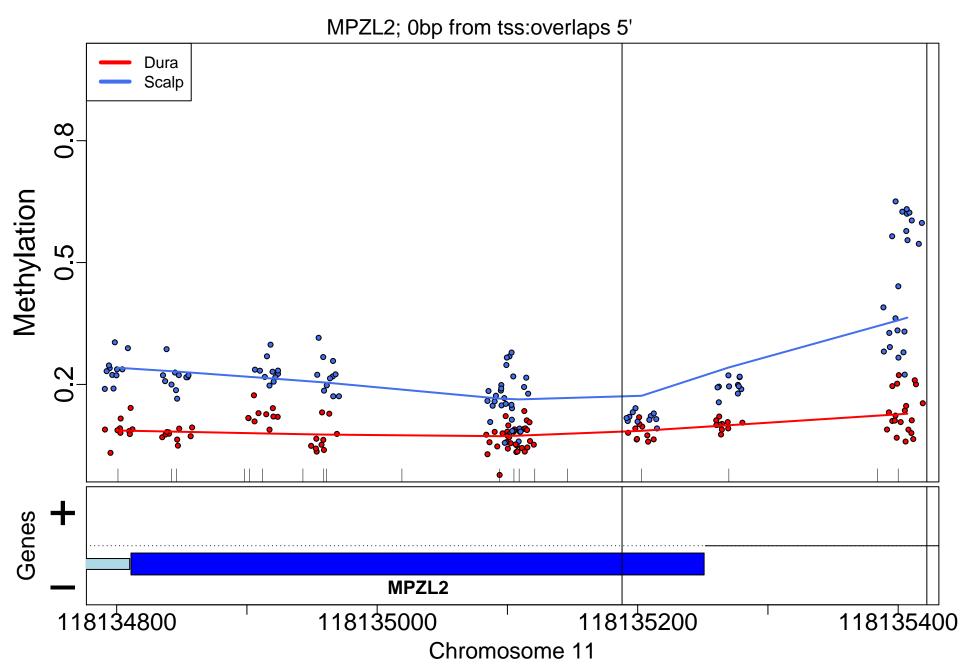


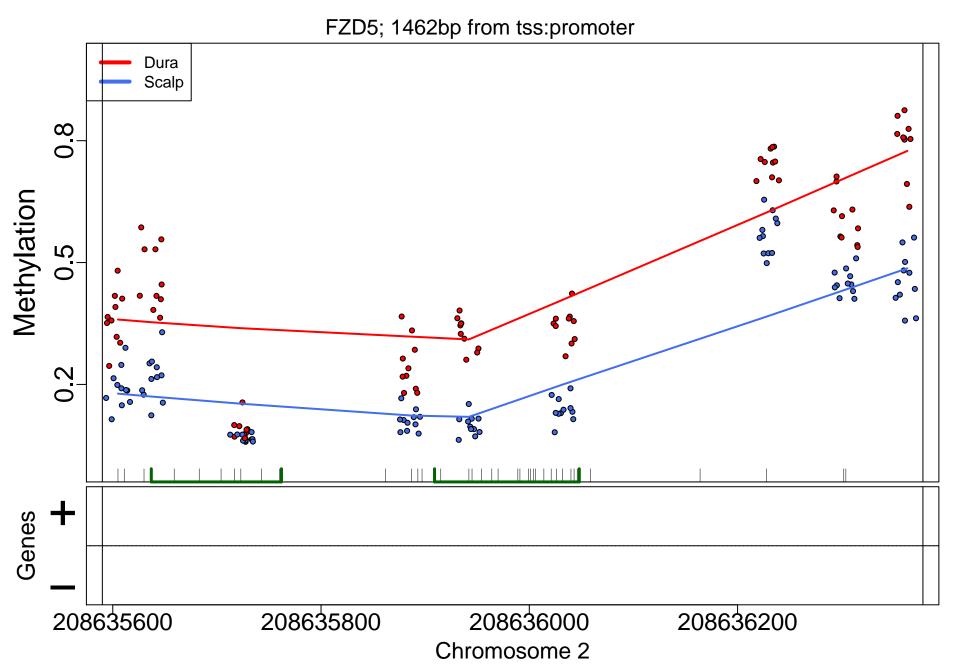


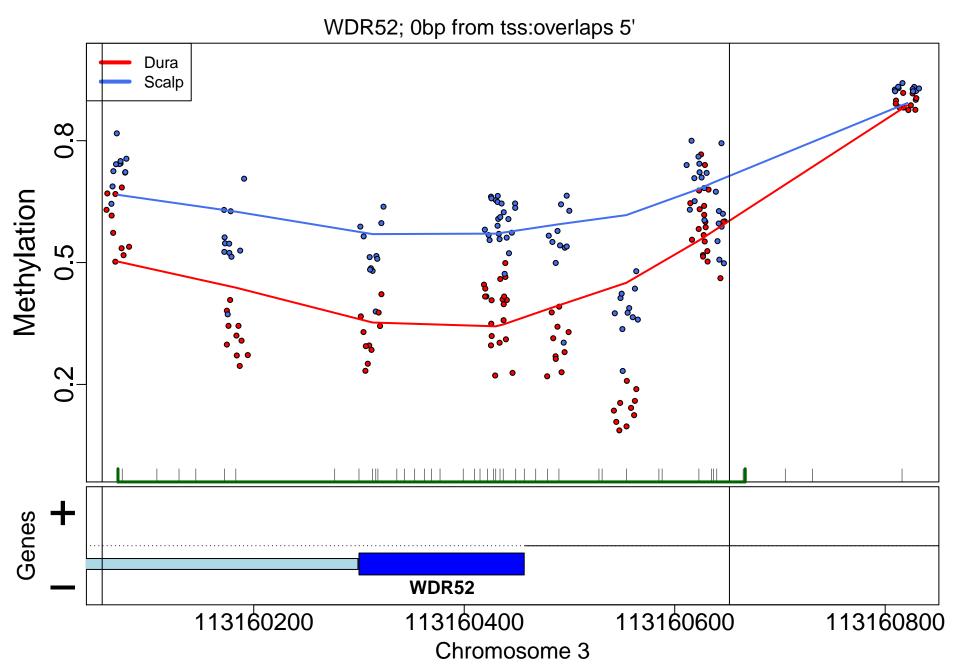


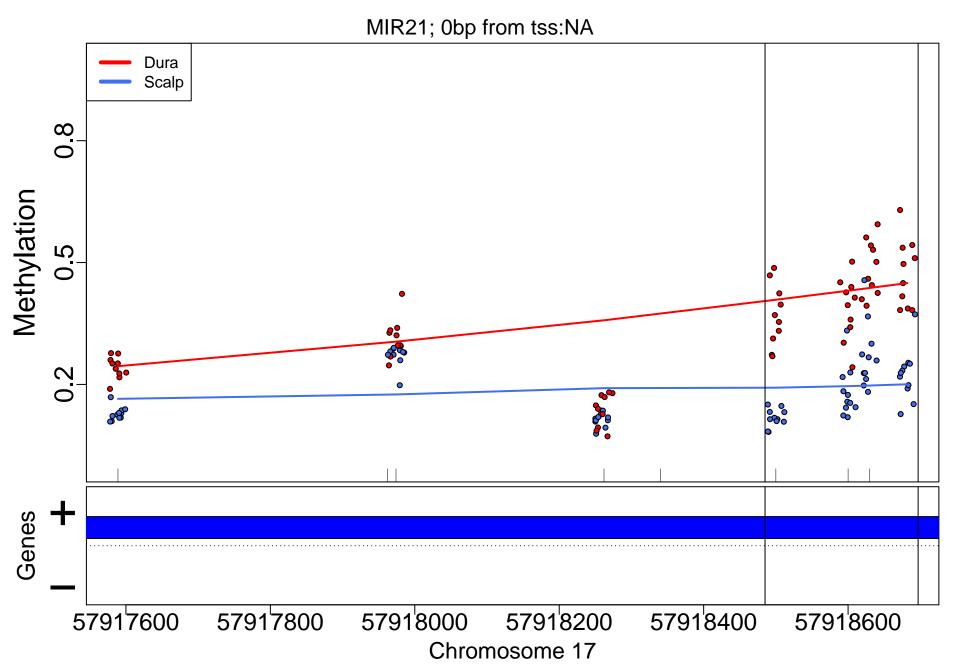


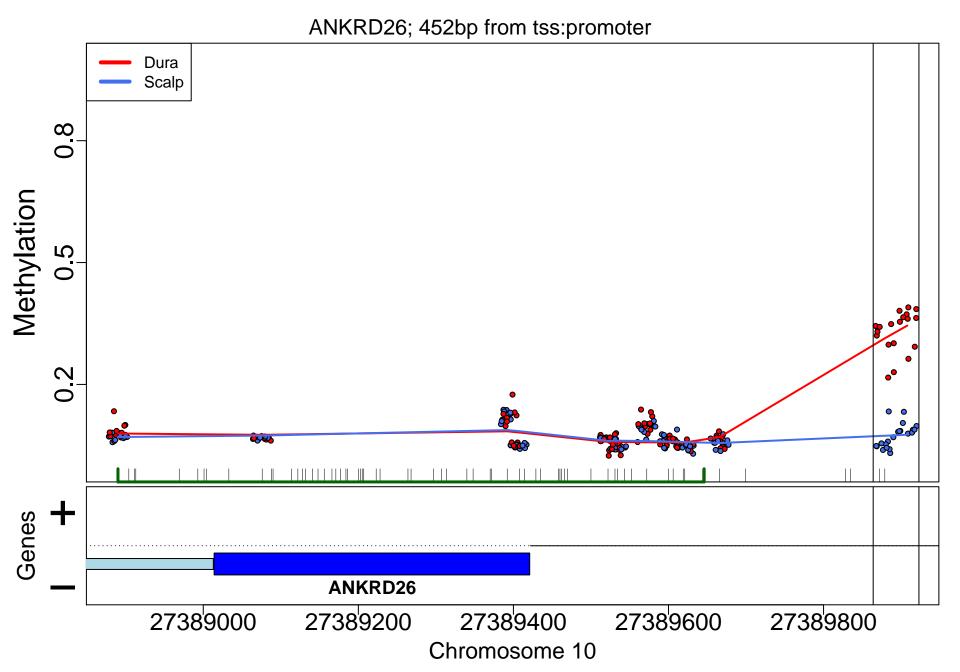


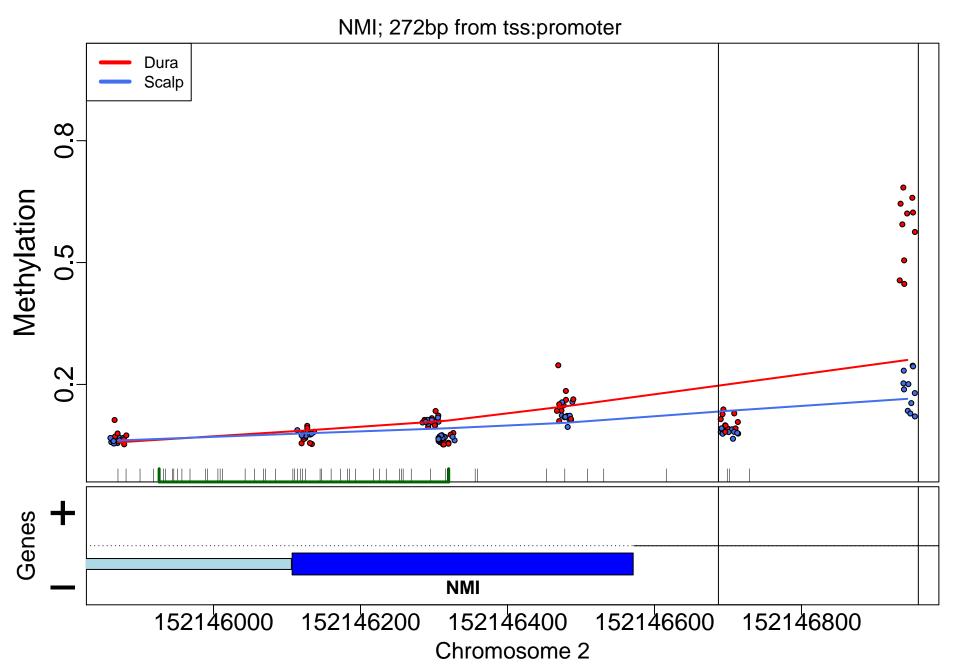


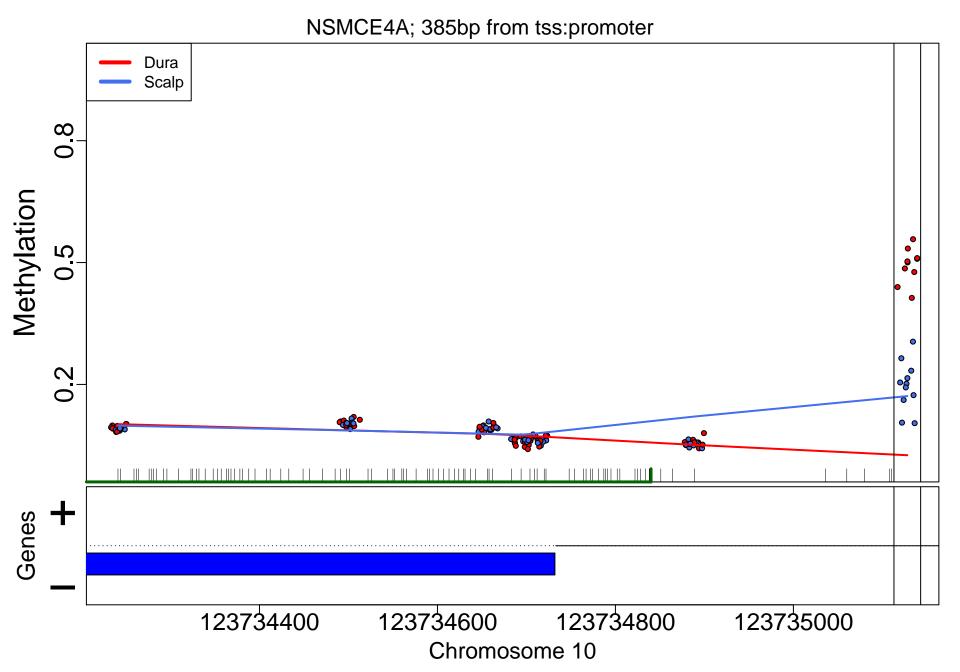


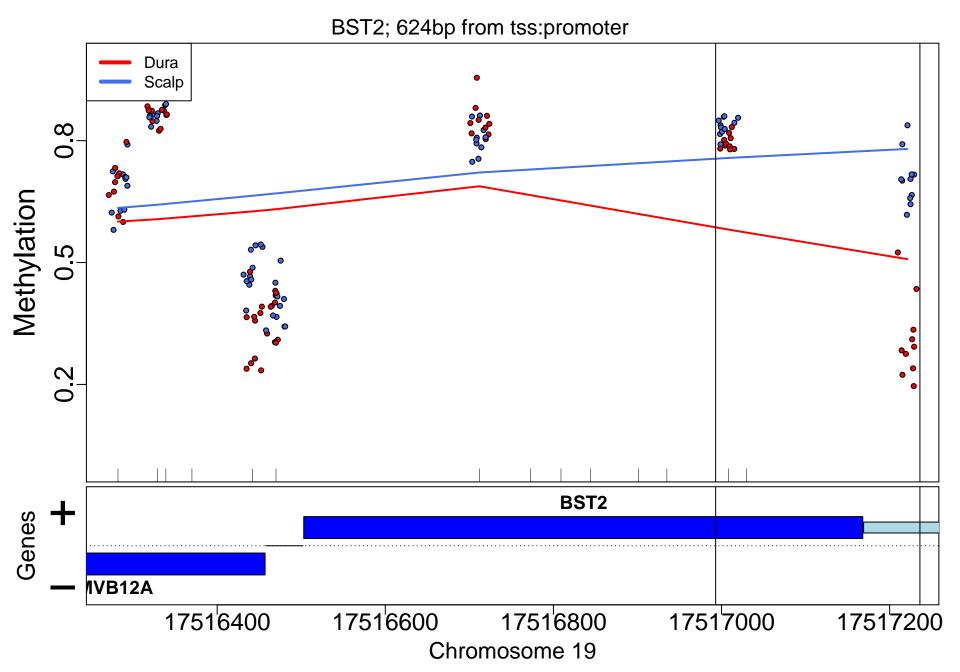


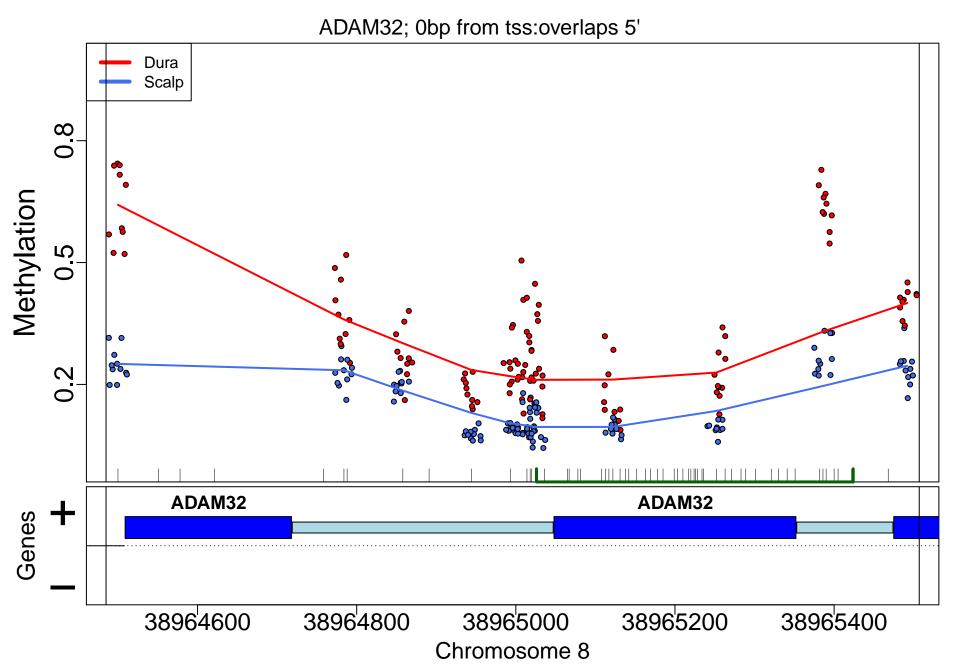


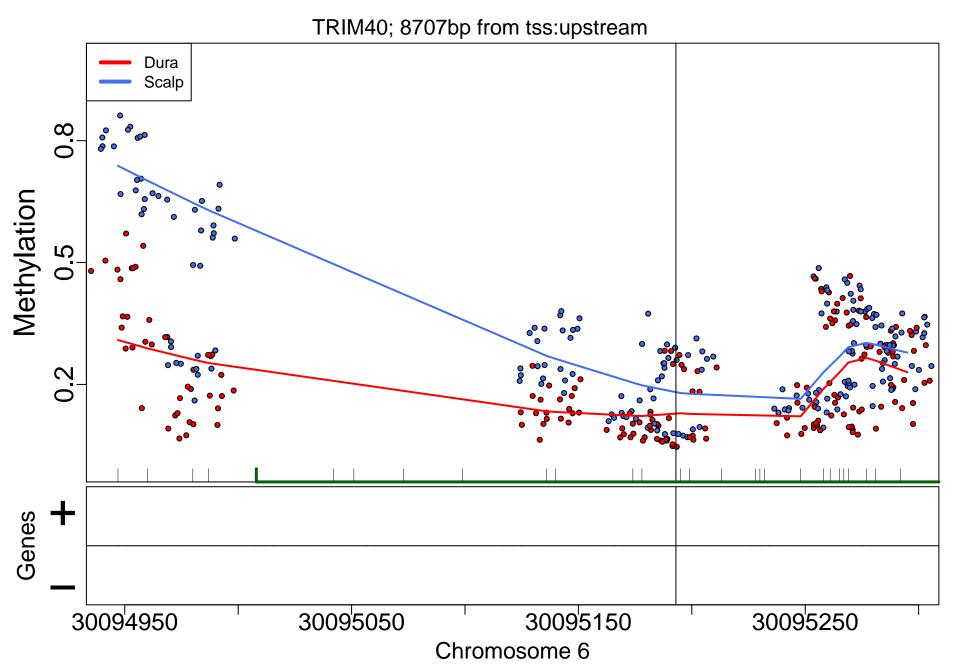


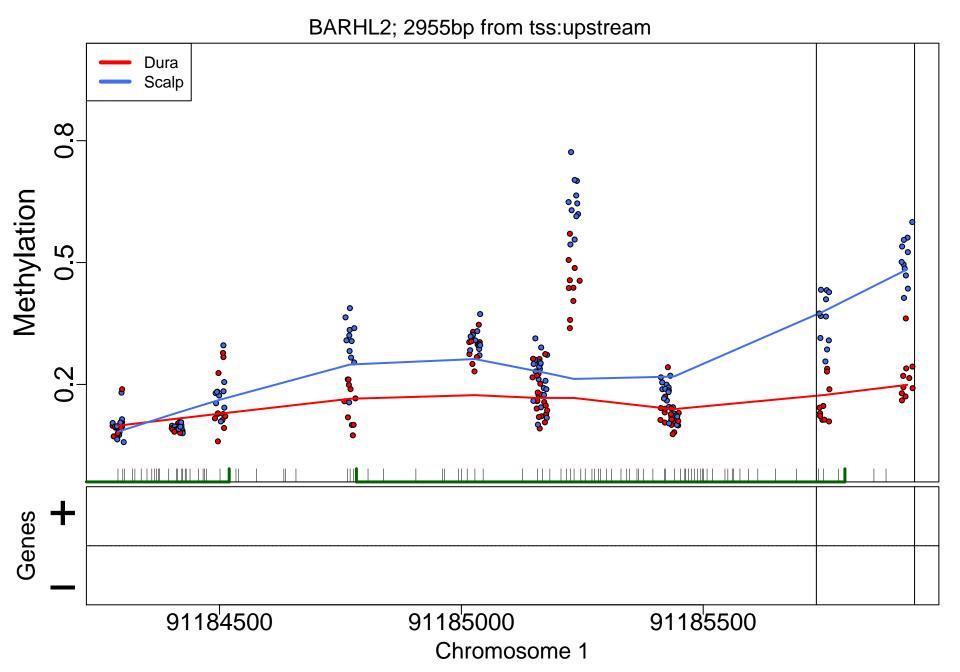


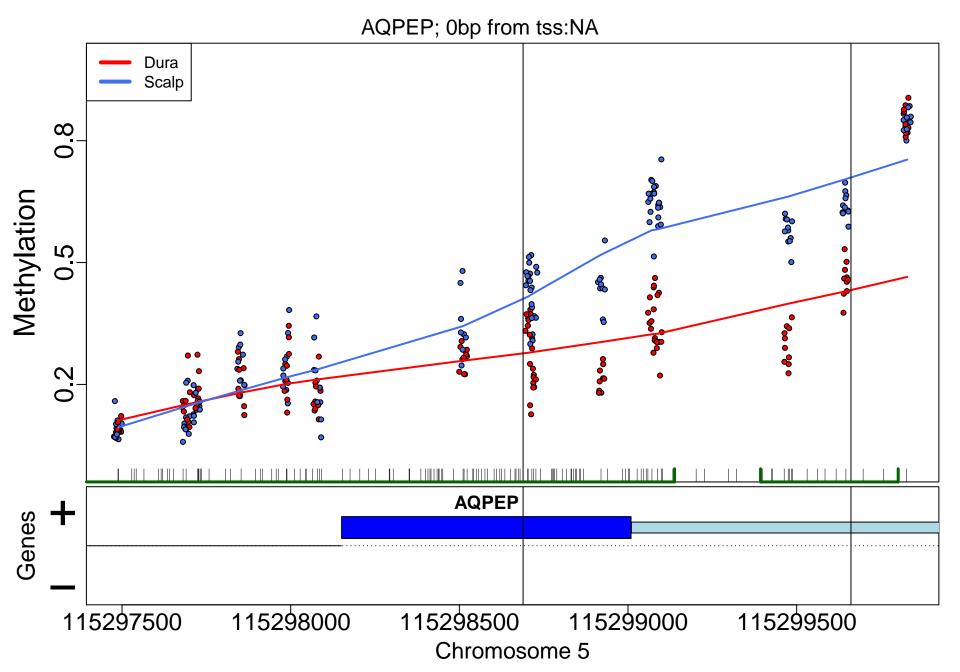


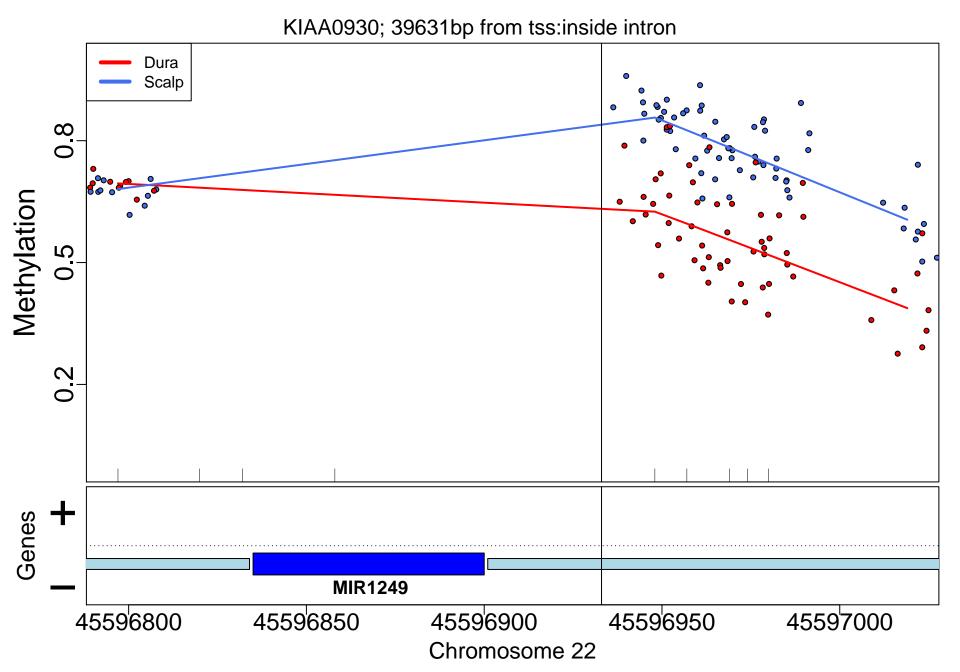


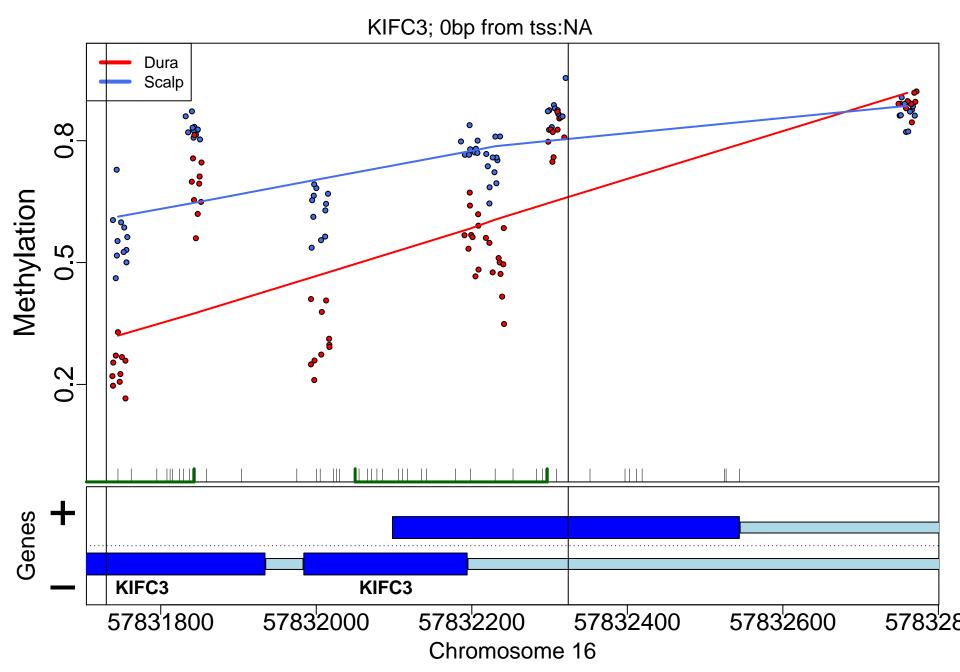


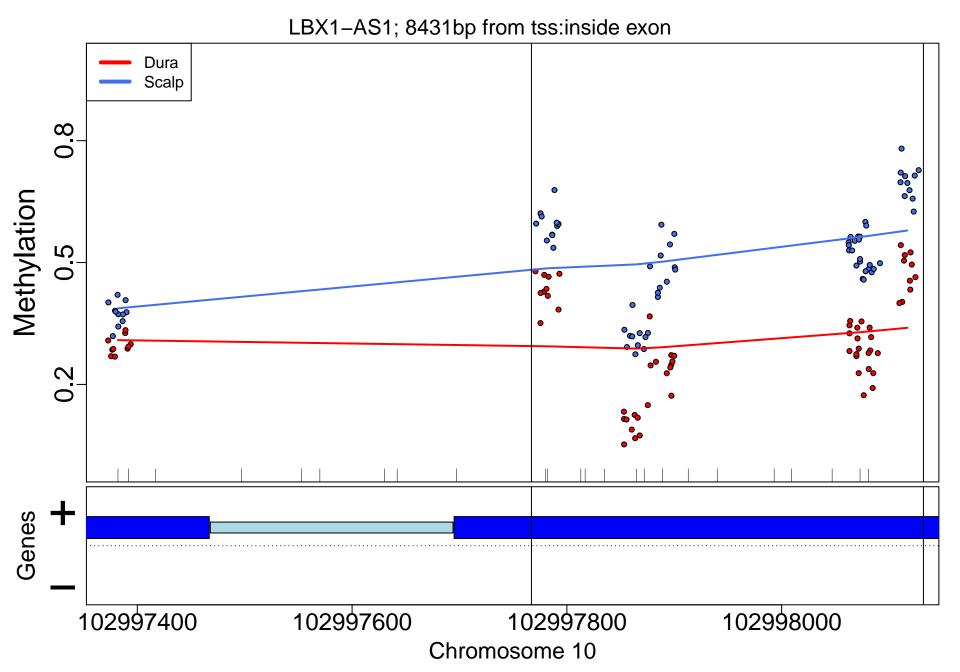


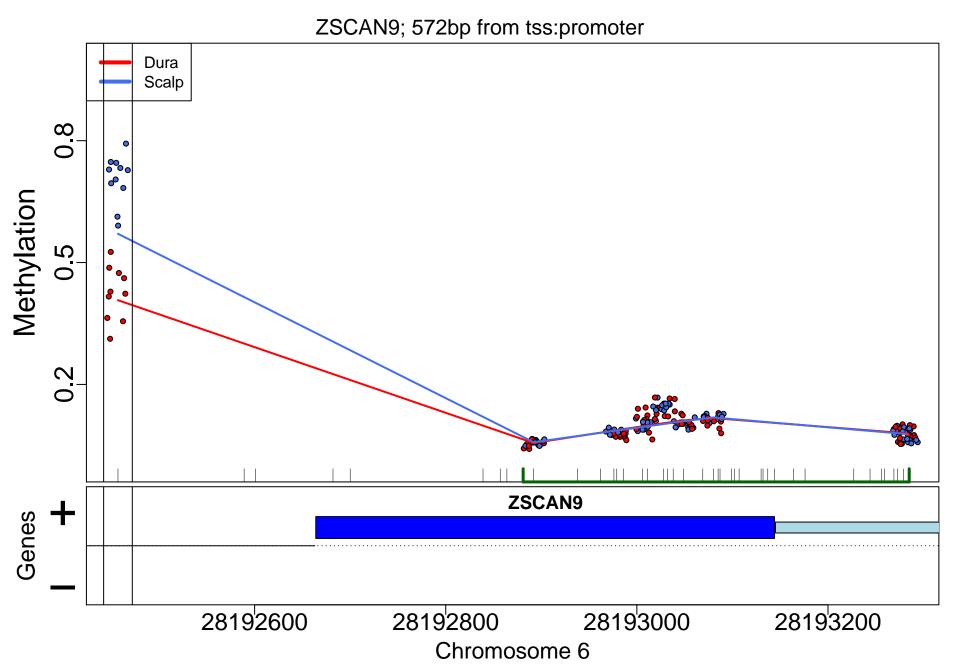


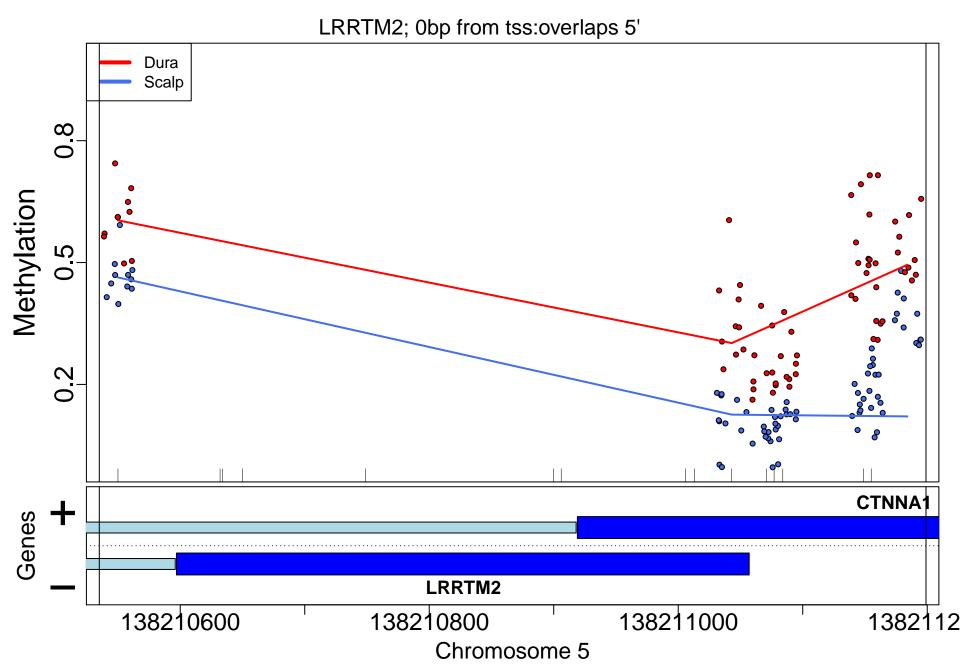


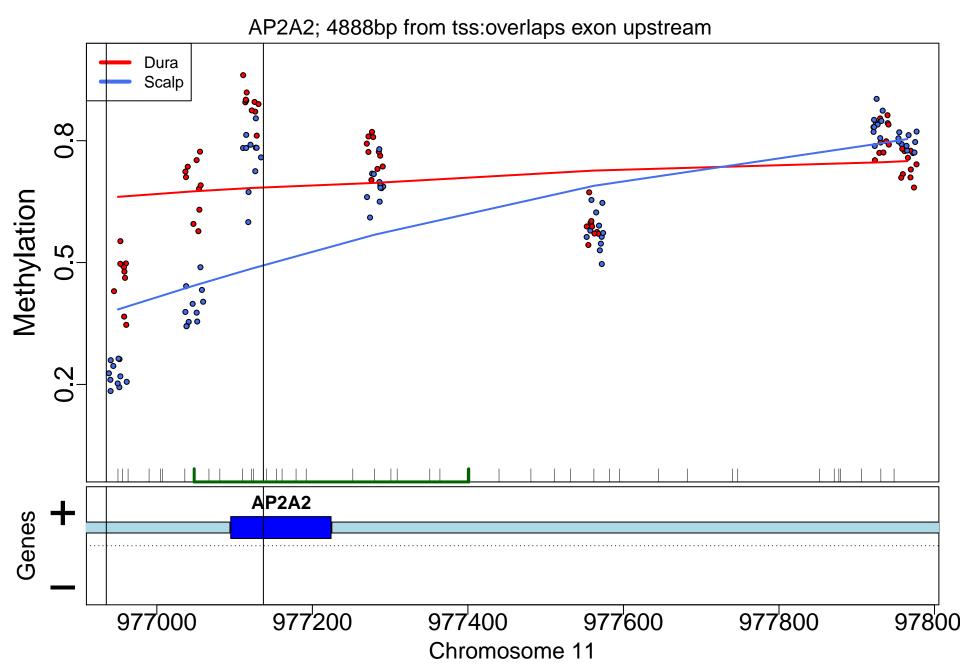


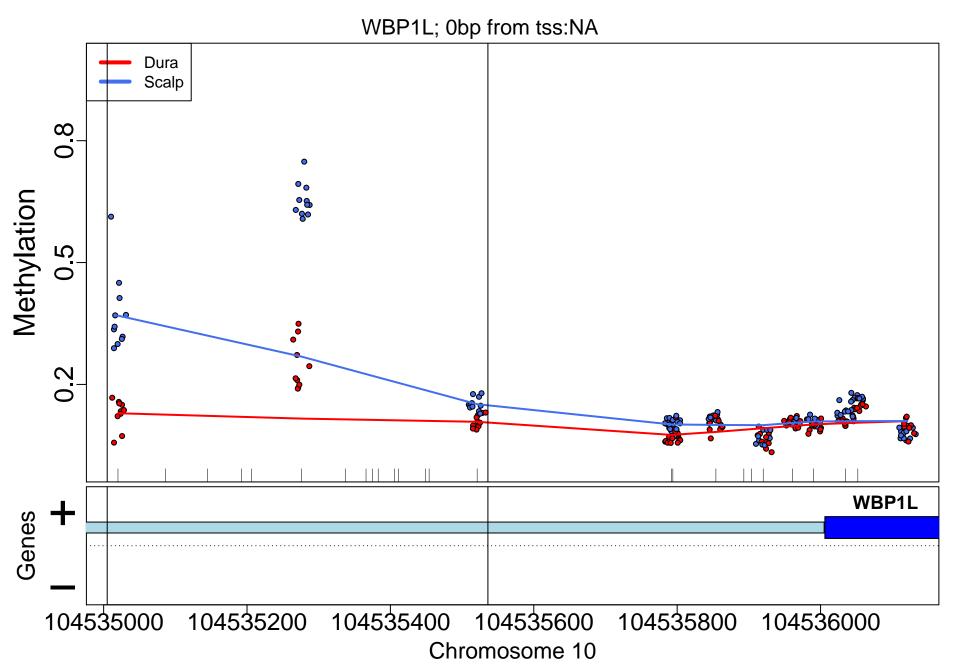


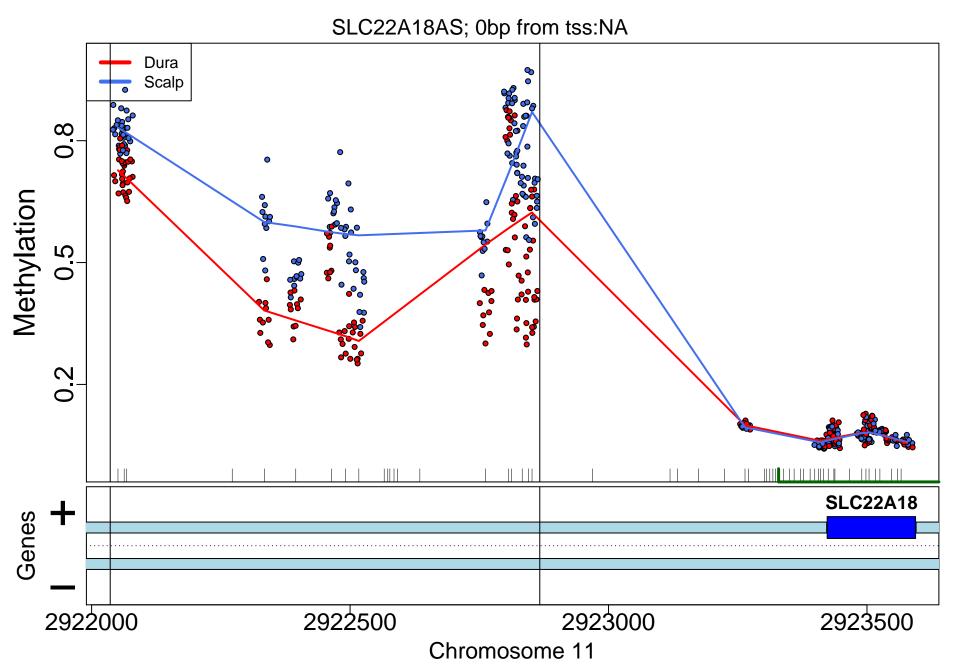


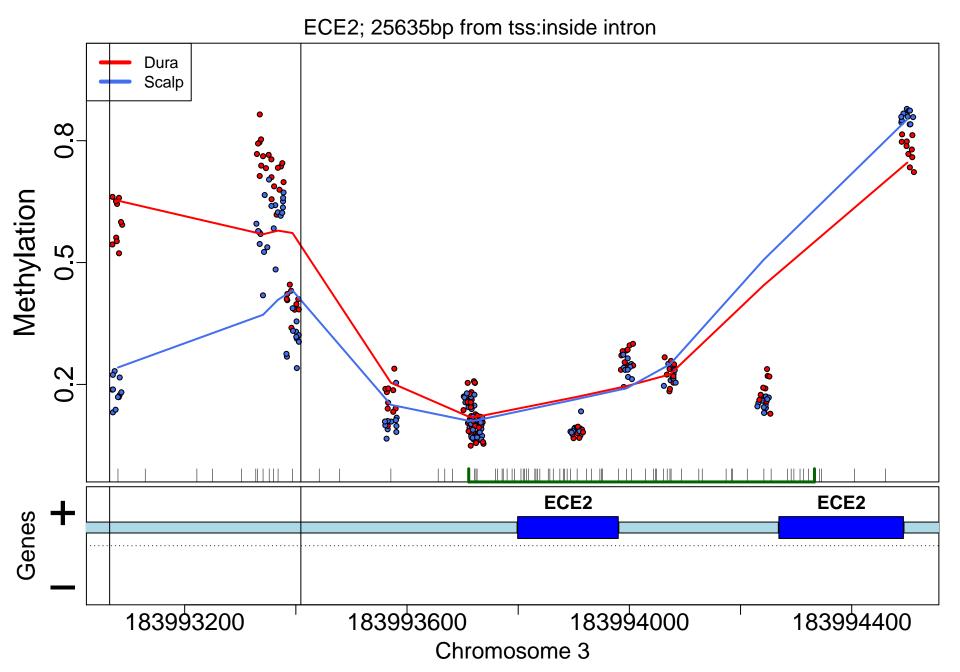


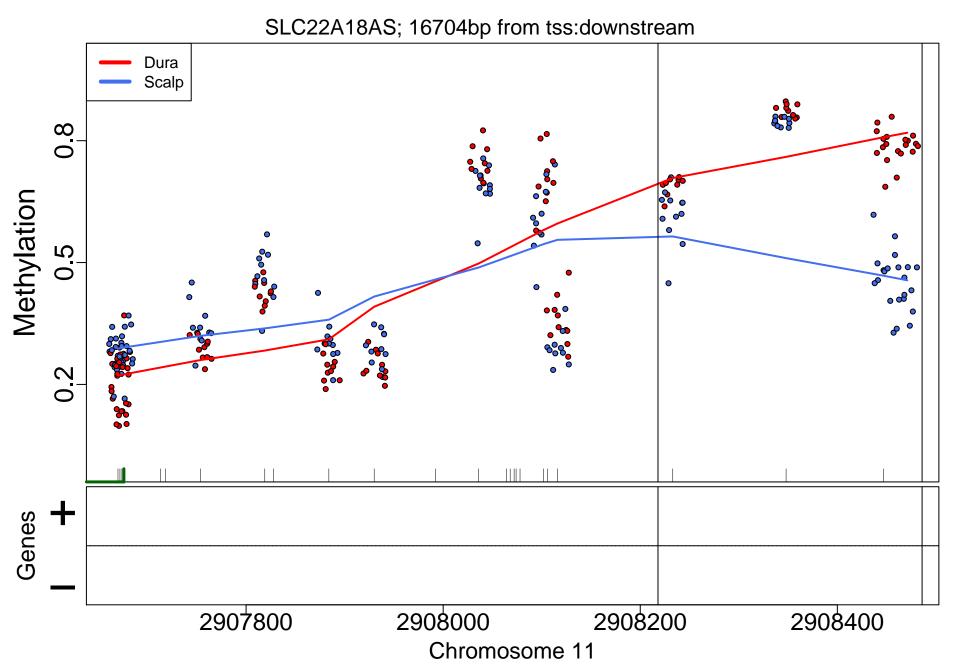


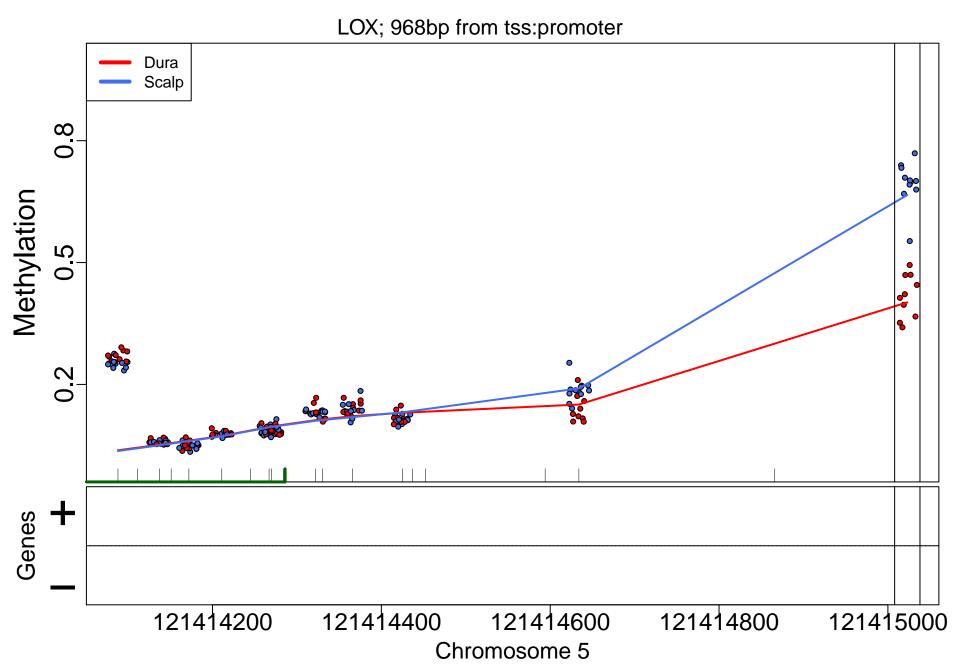


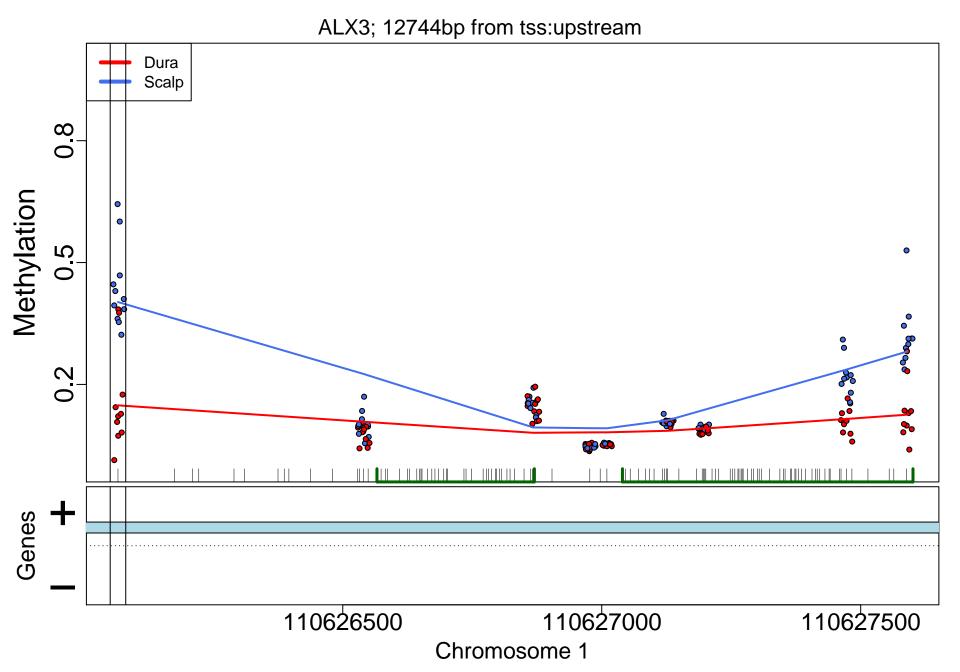






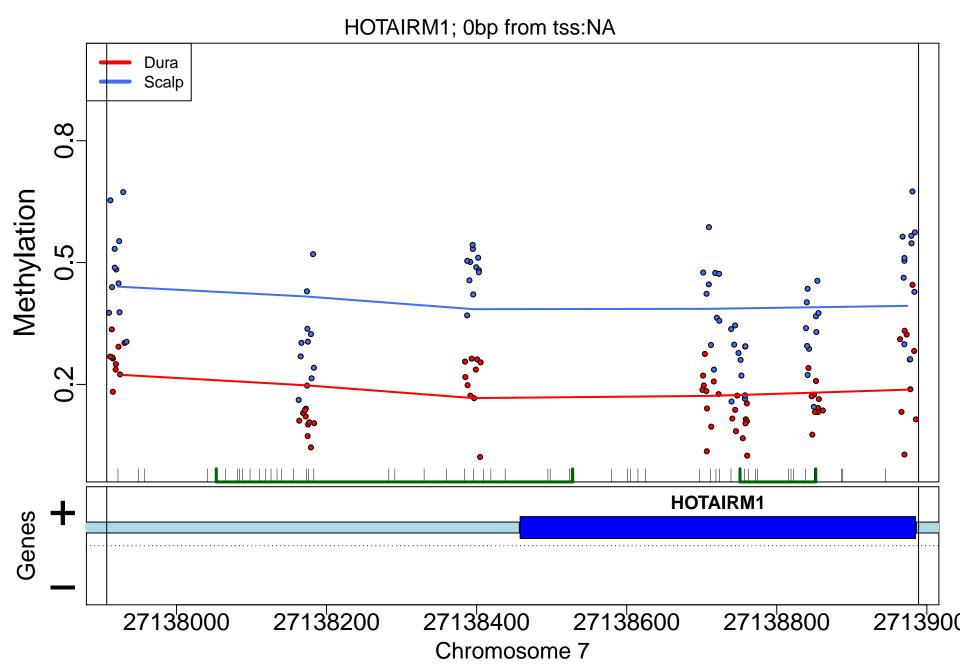


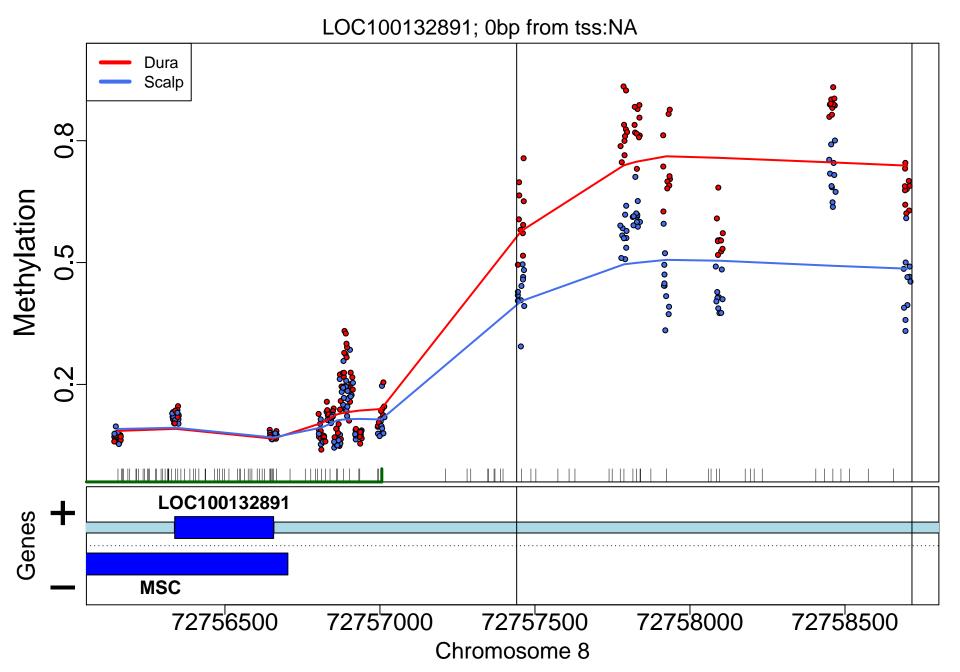


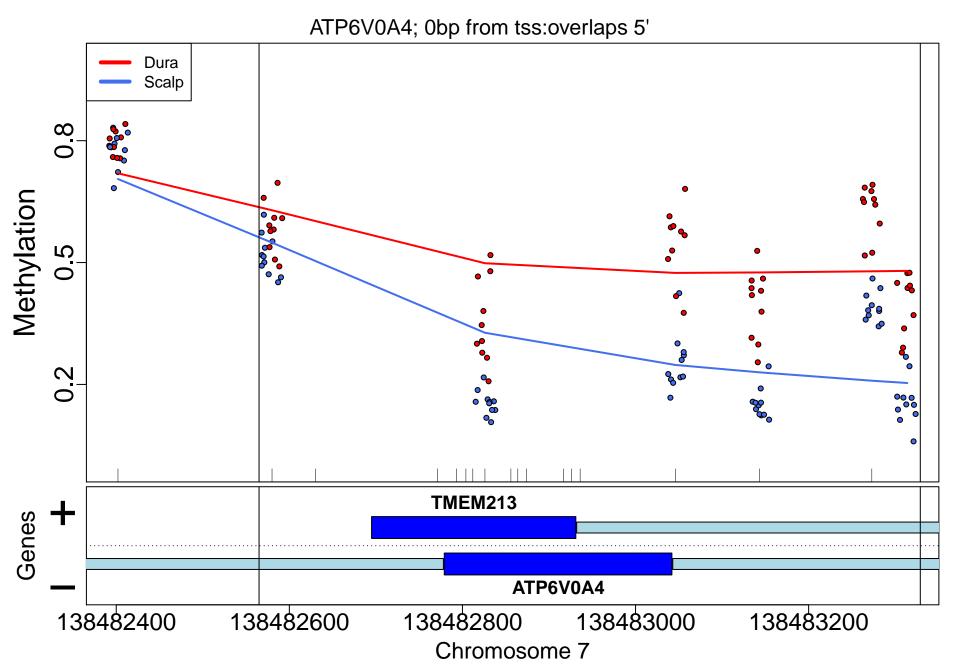


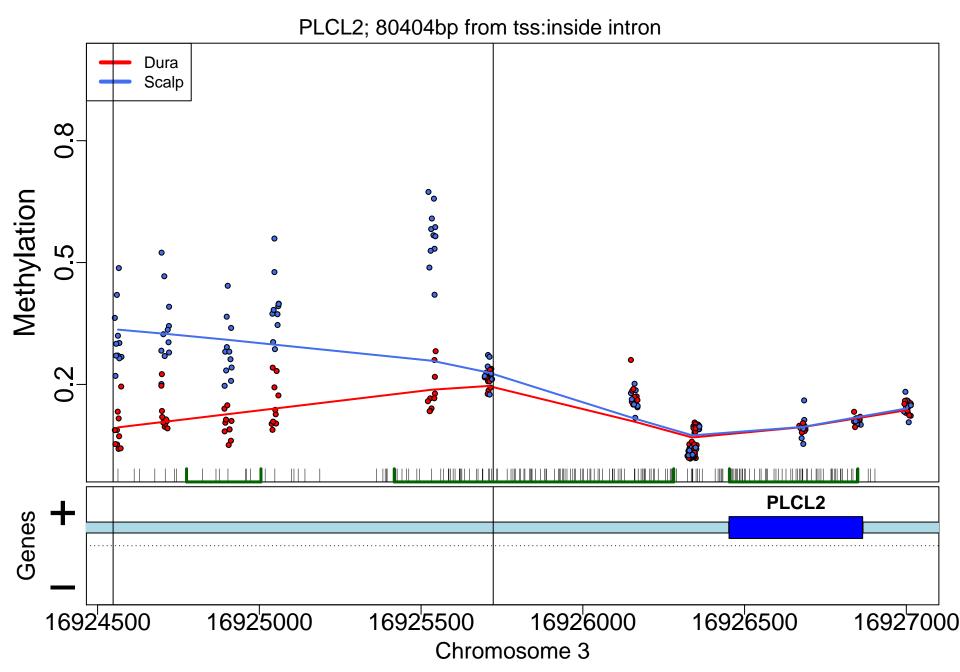
FOXF1; 2495bp from tss:inside exon Dura Scalp 80. Methylation 0.2 FOXF1 Genes 86546400 86546600 86547200 86546800 86547000 86547400

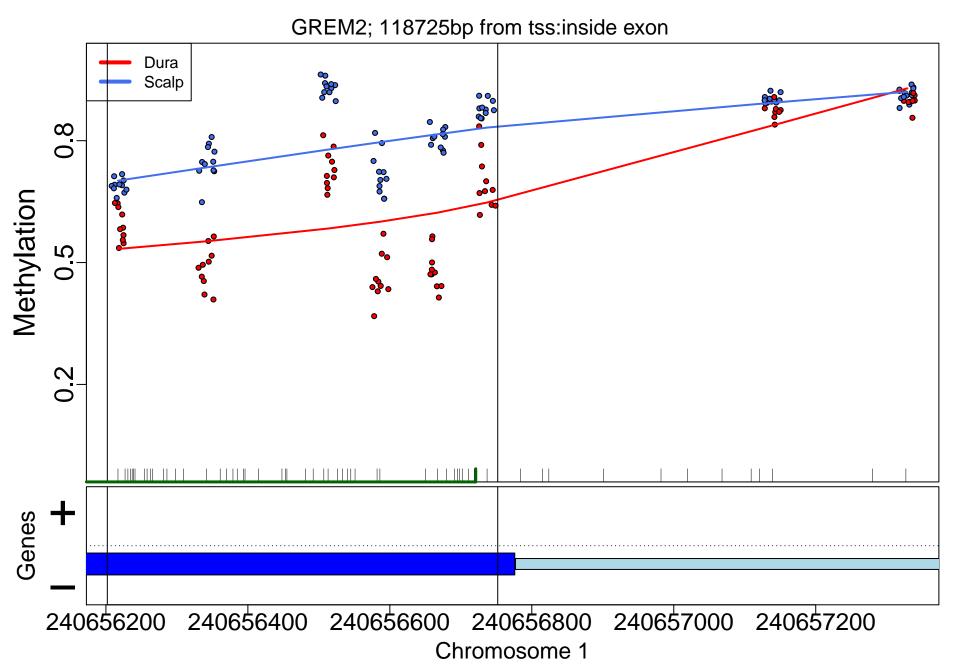
Chromosome 16

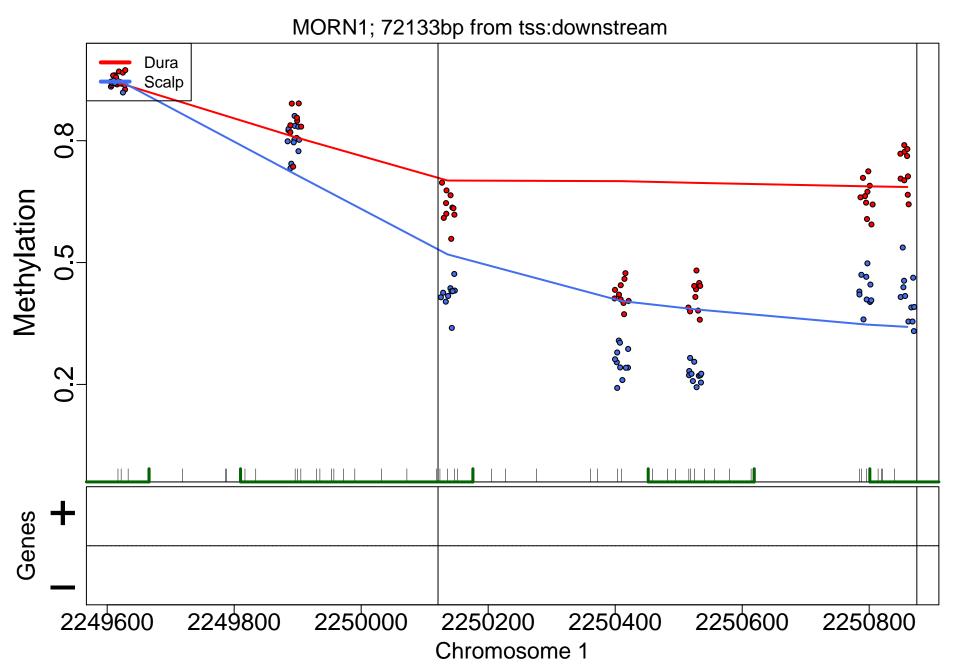


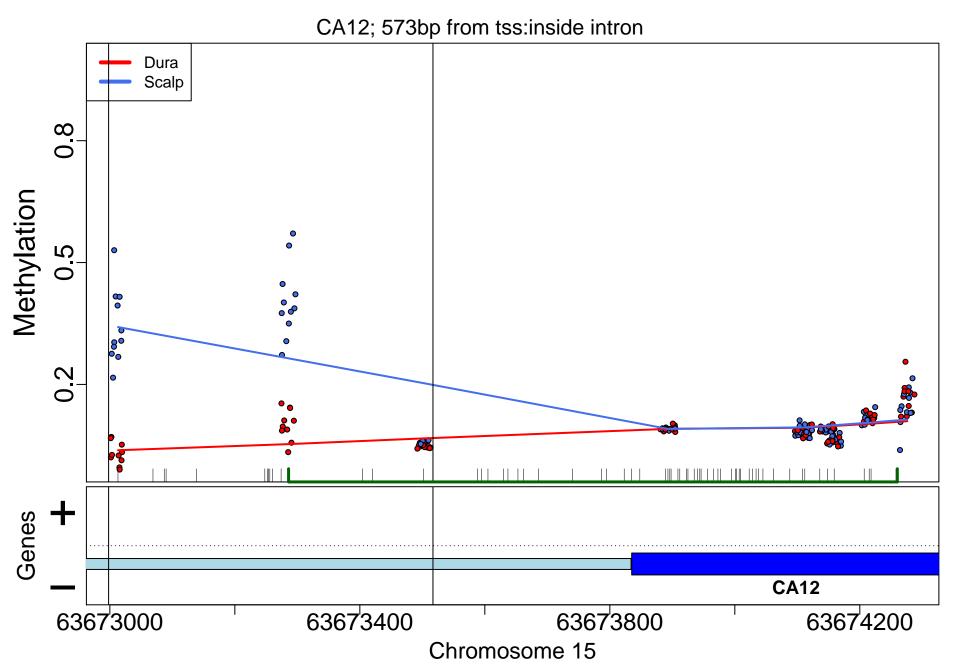


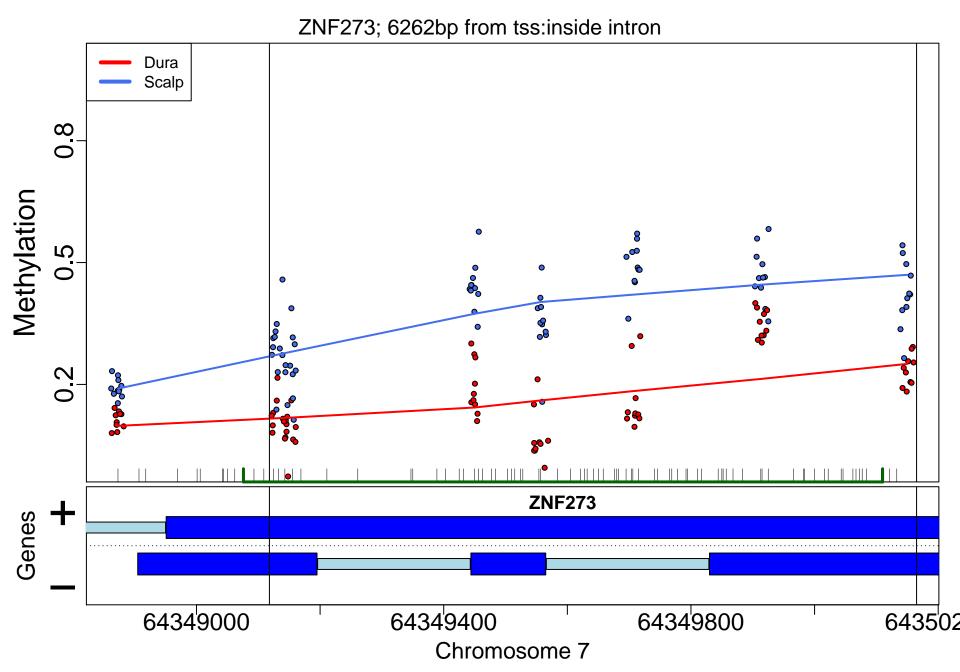


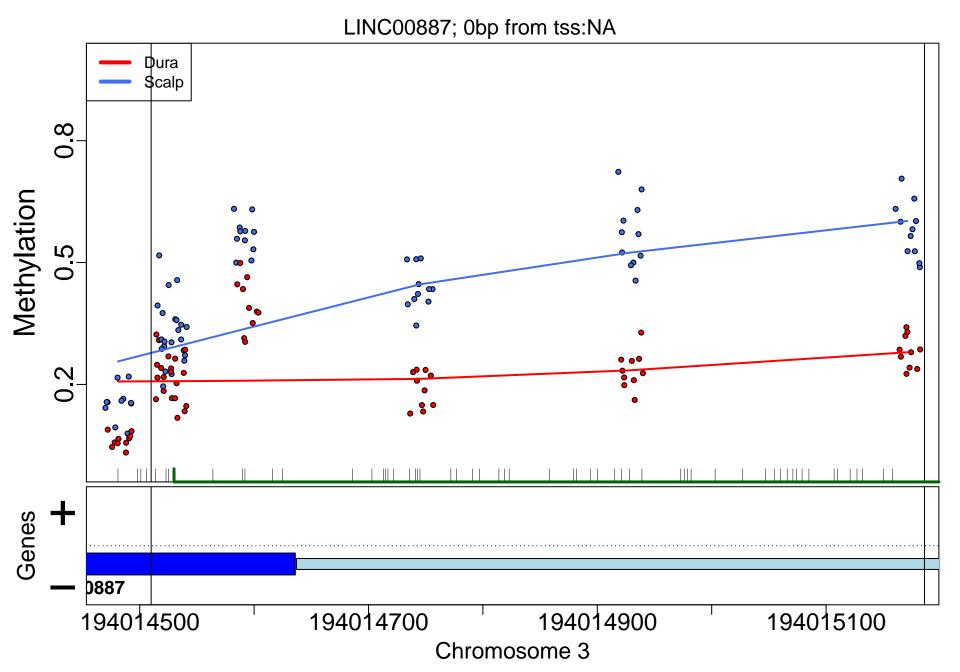


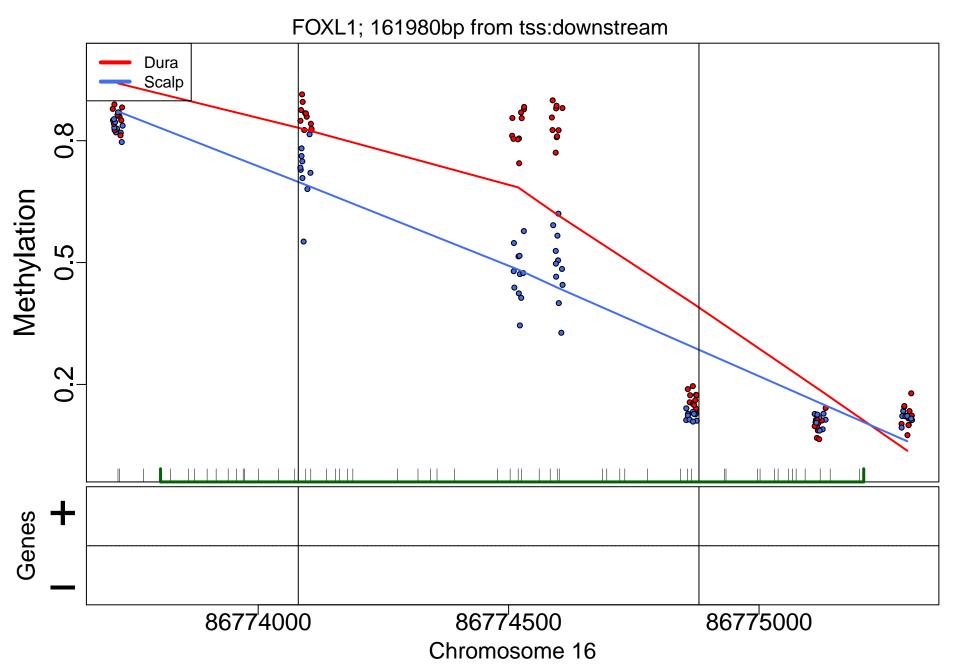


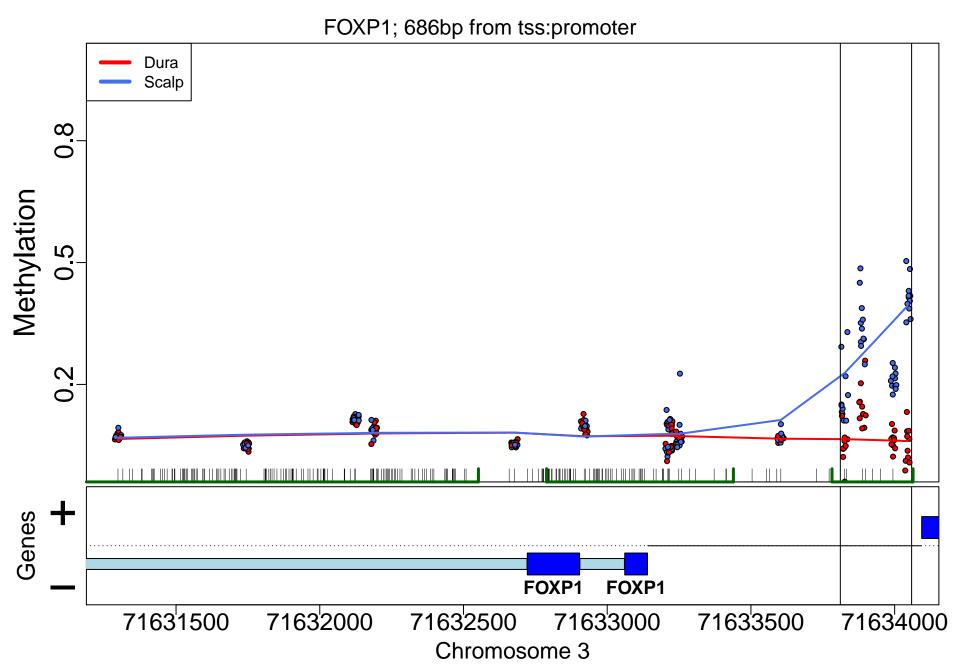


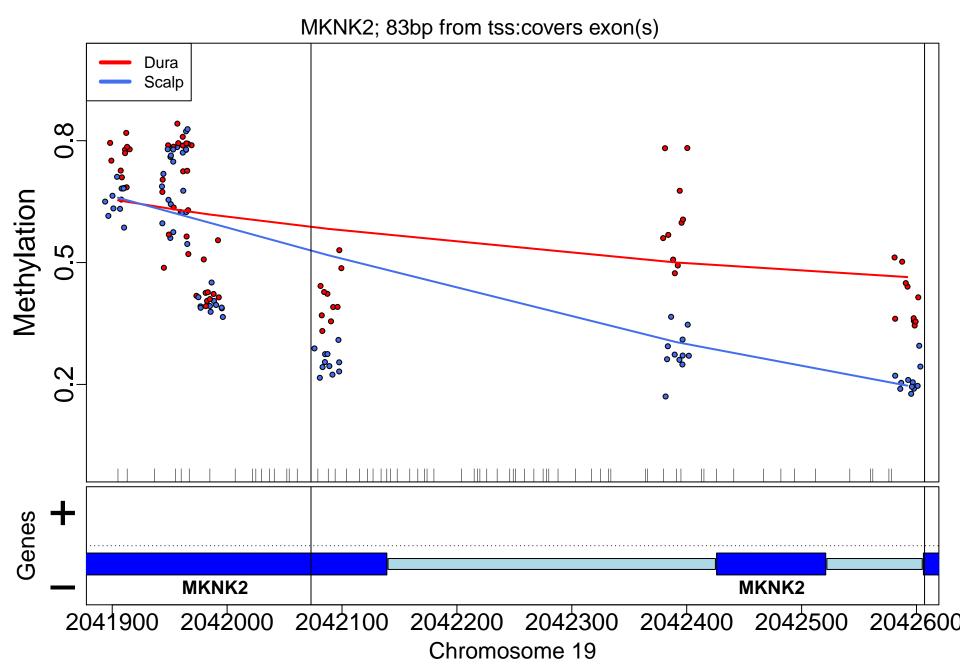


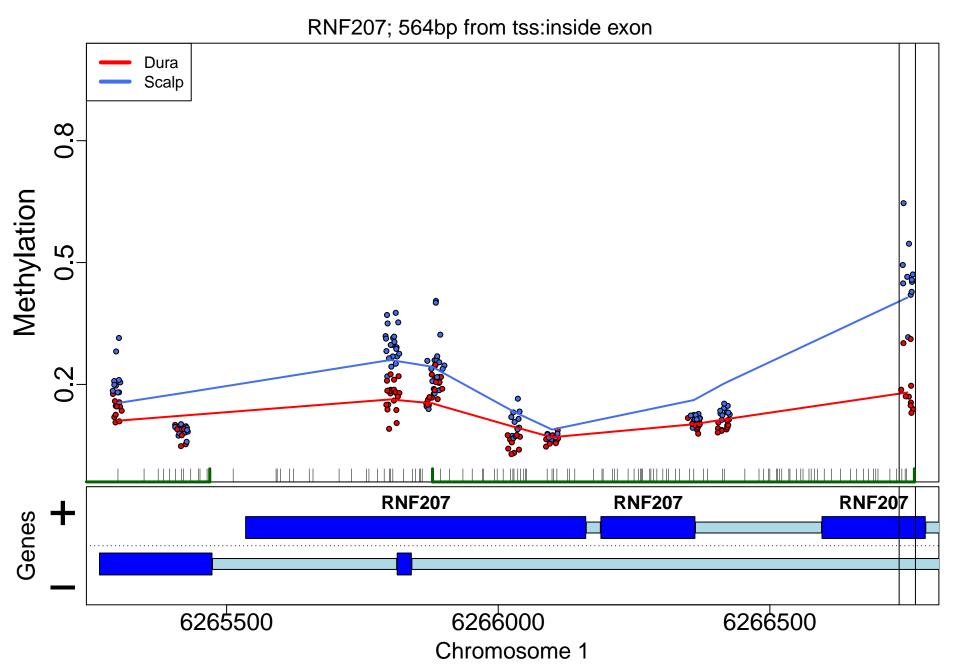


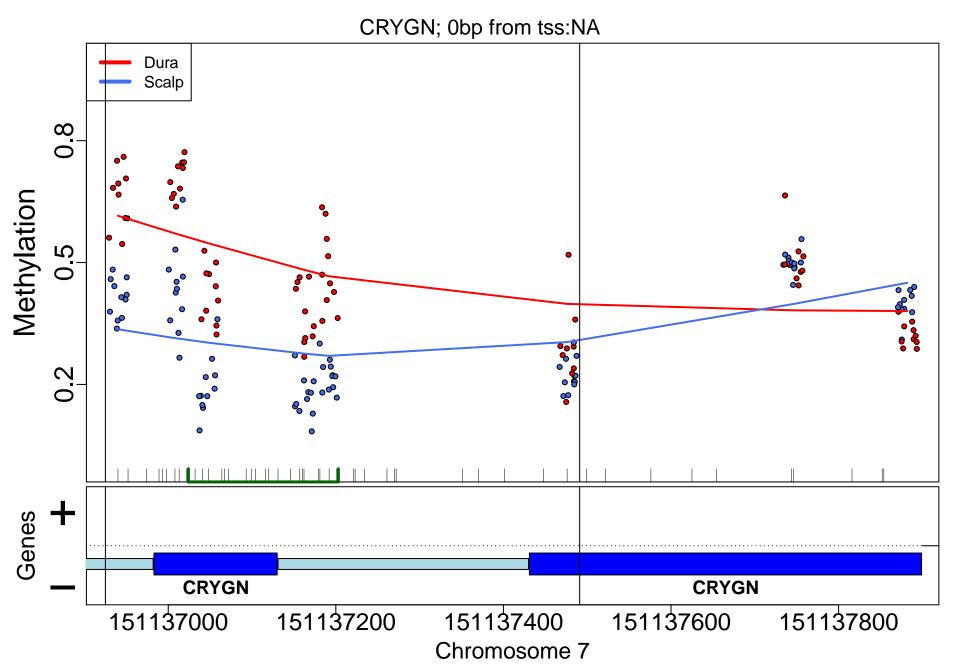




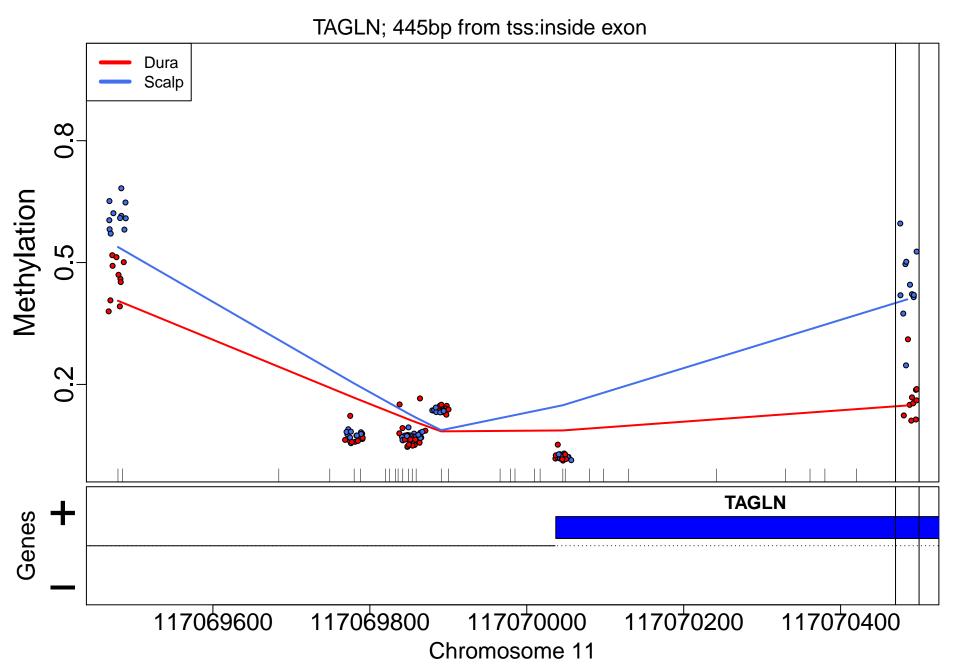


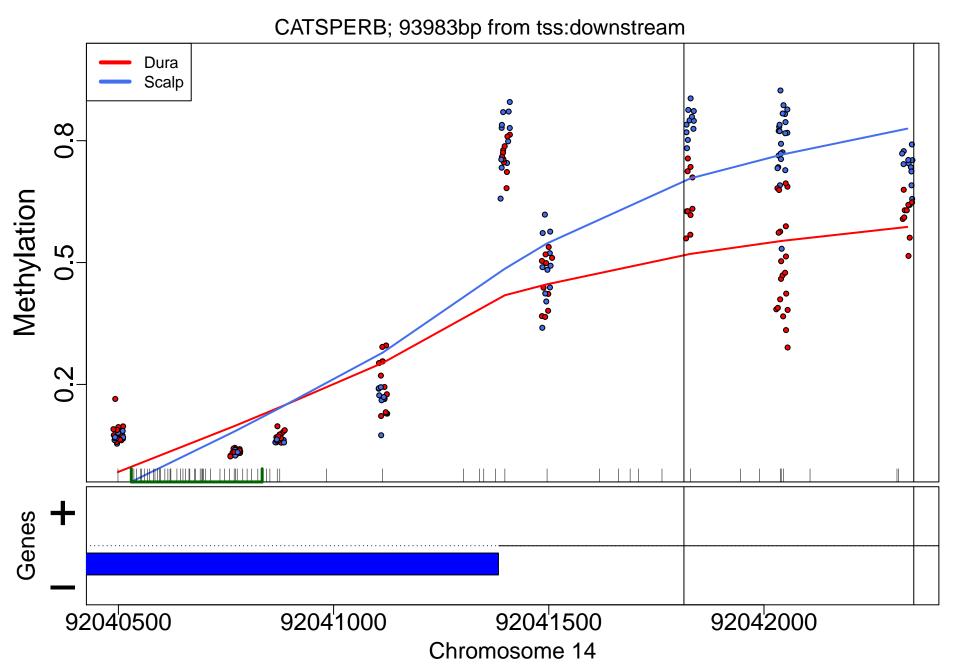


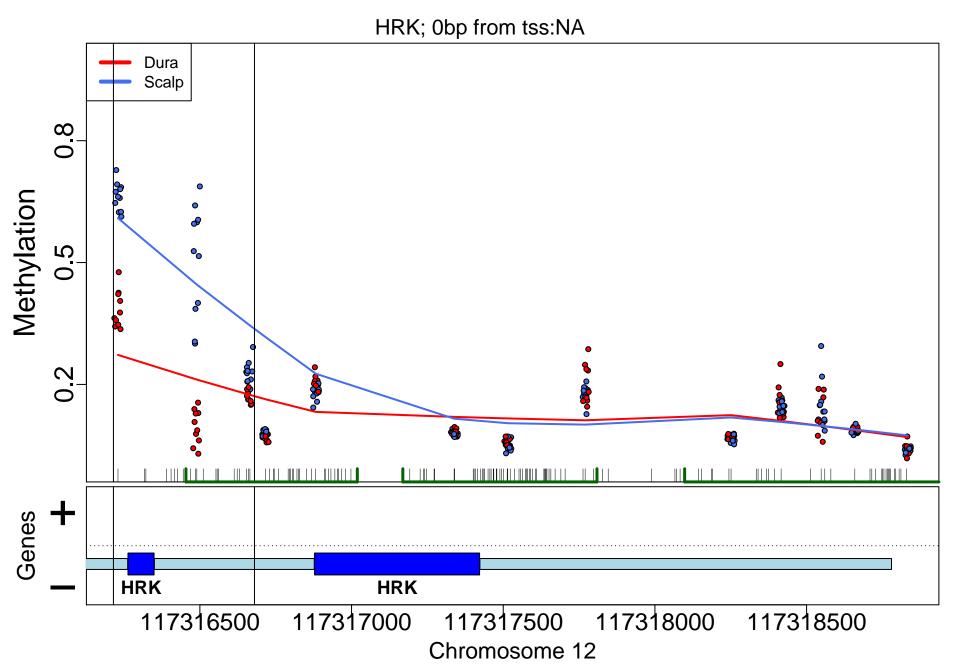


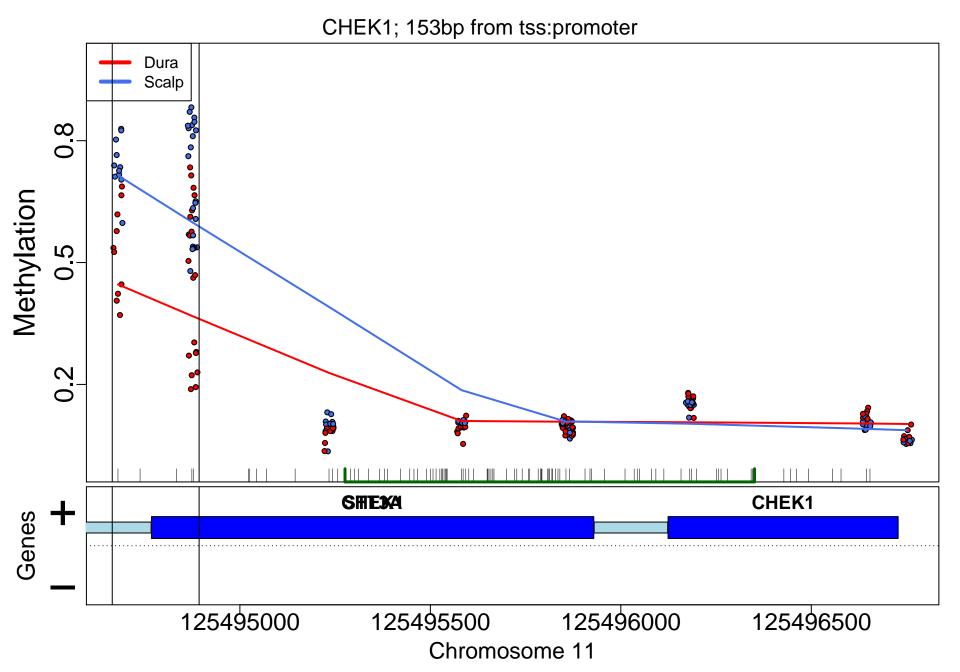


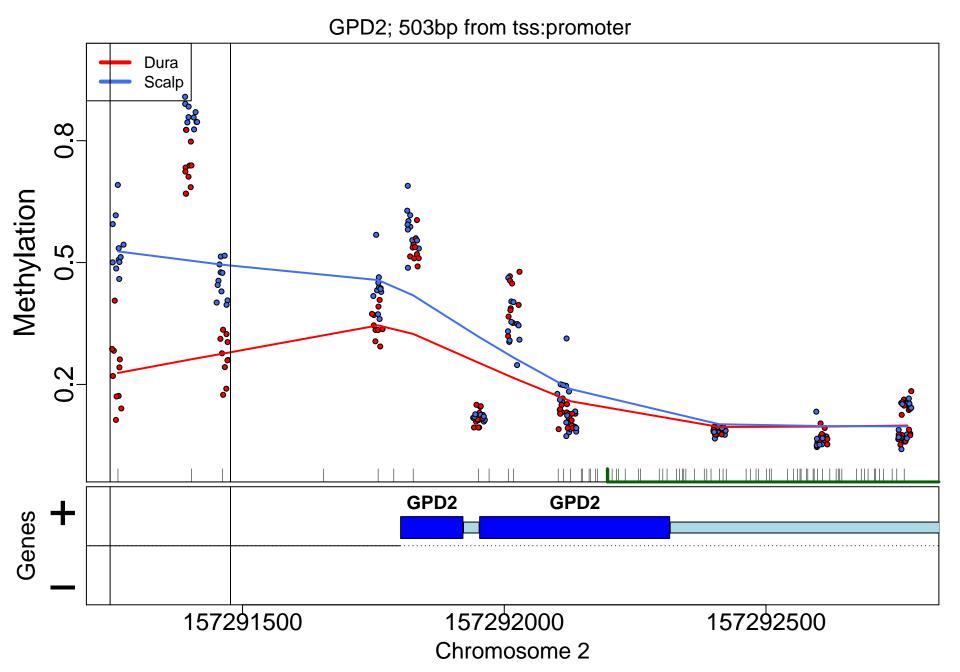
GAD1; 1405bp from tss:promoter Dura Scalp <u>∞</u>. Methylation **8** GAD1 GAD1 Genes 171671500 171672000 171672500 171673000 171673500 Chromosome 2

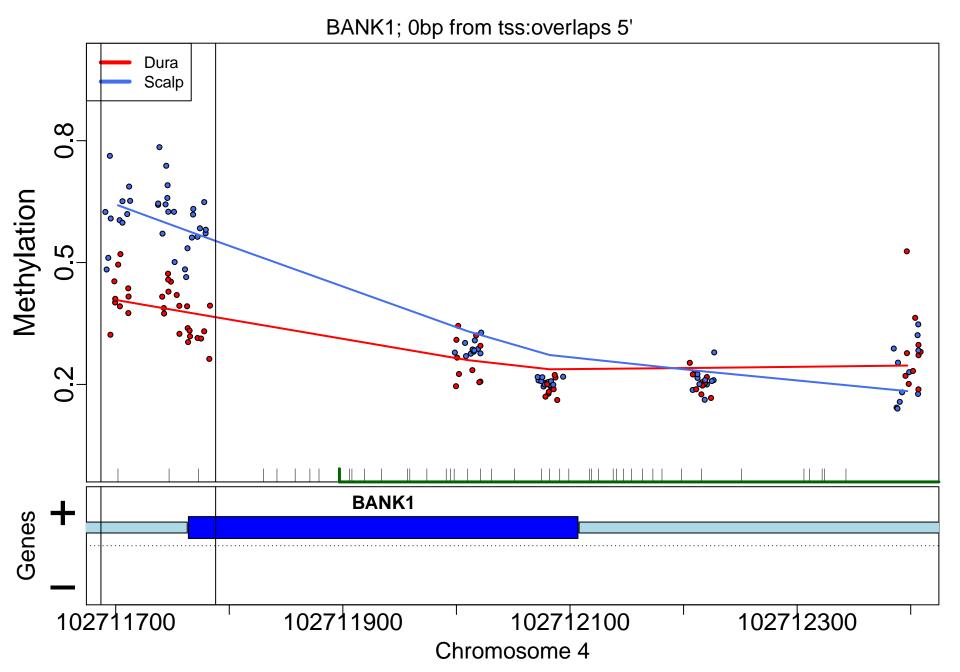


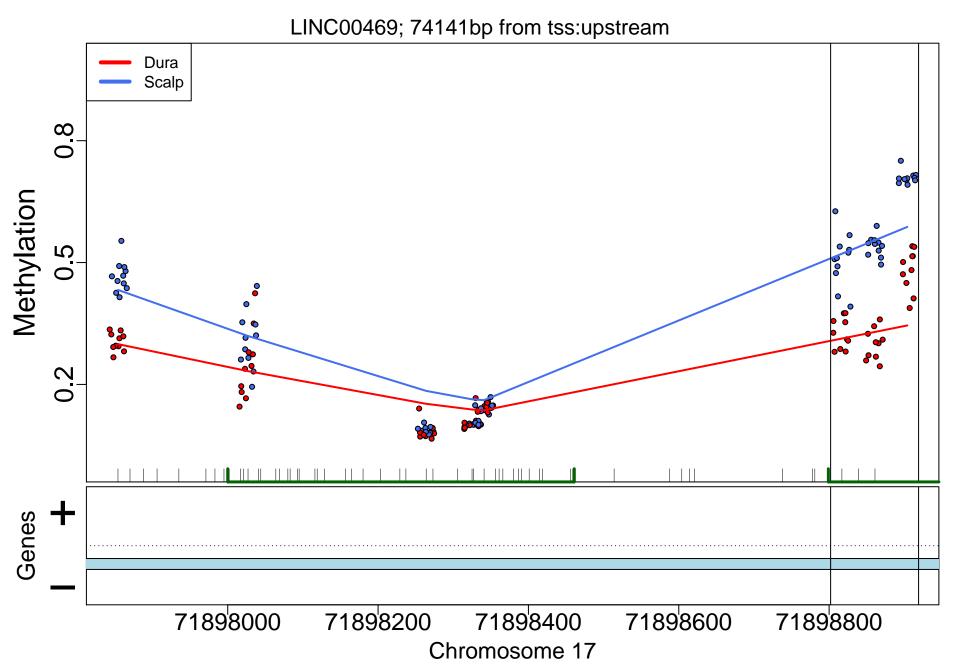


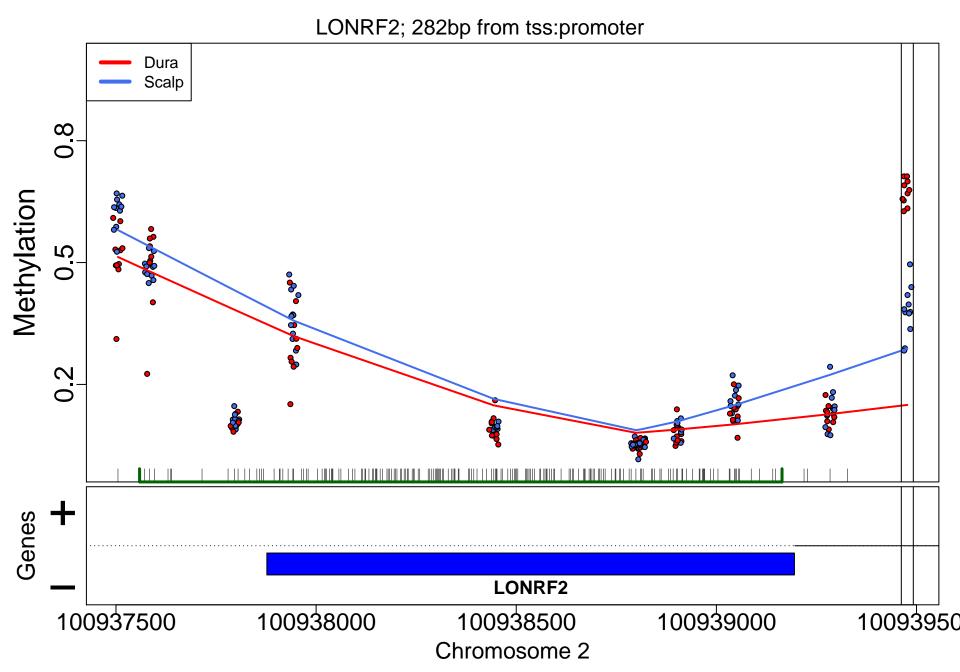


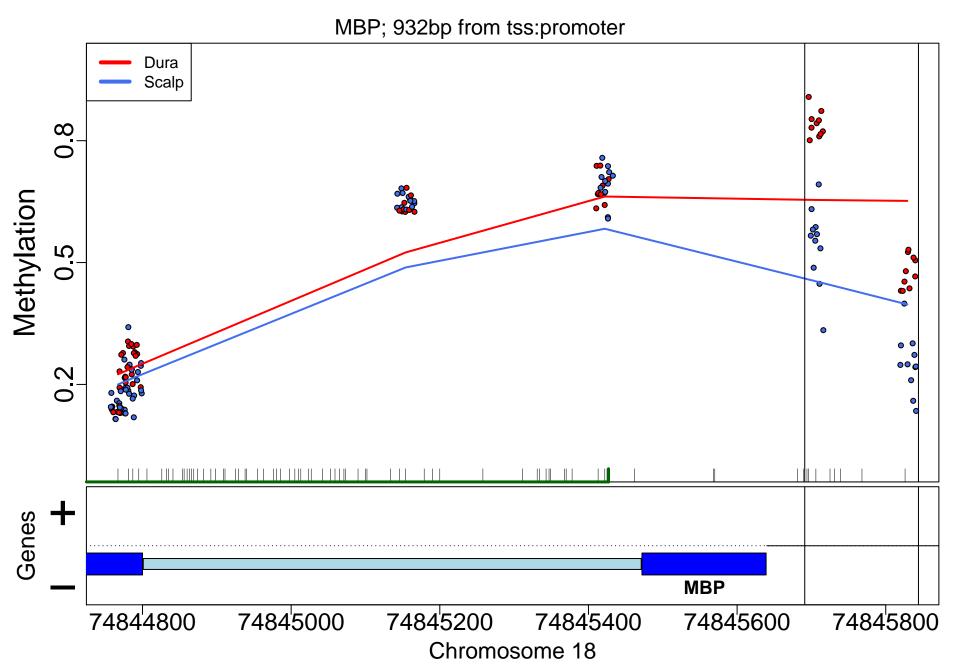


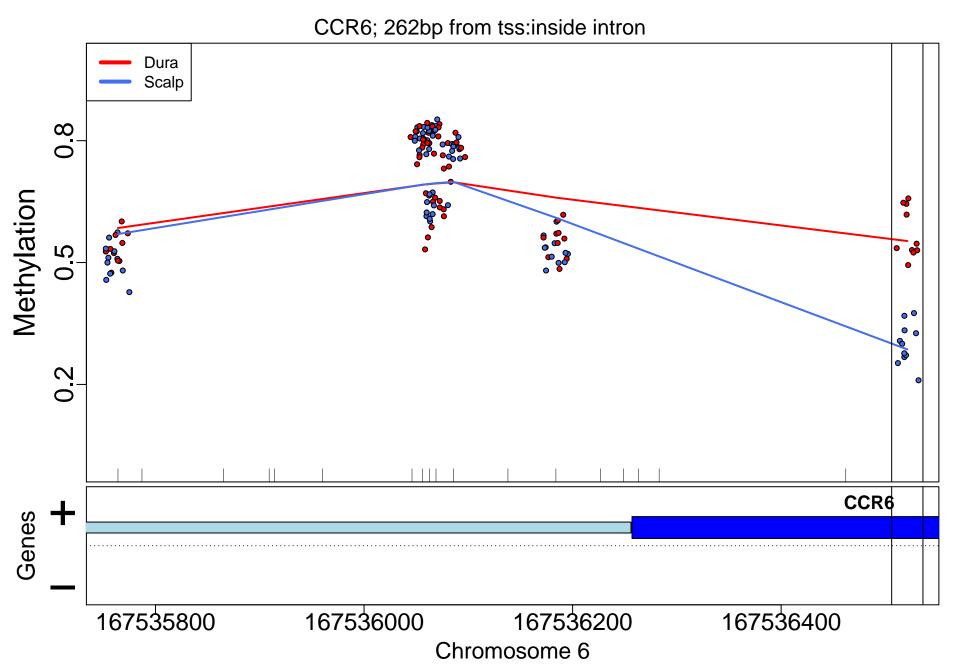


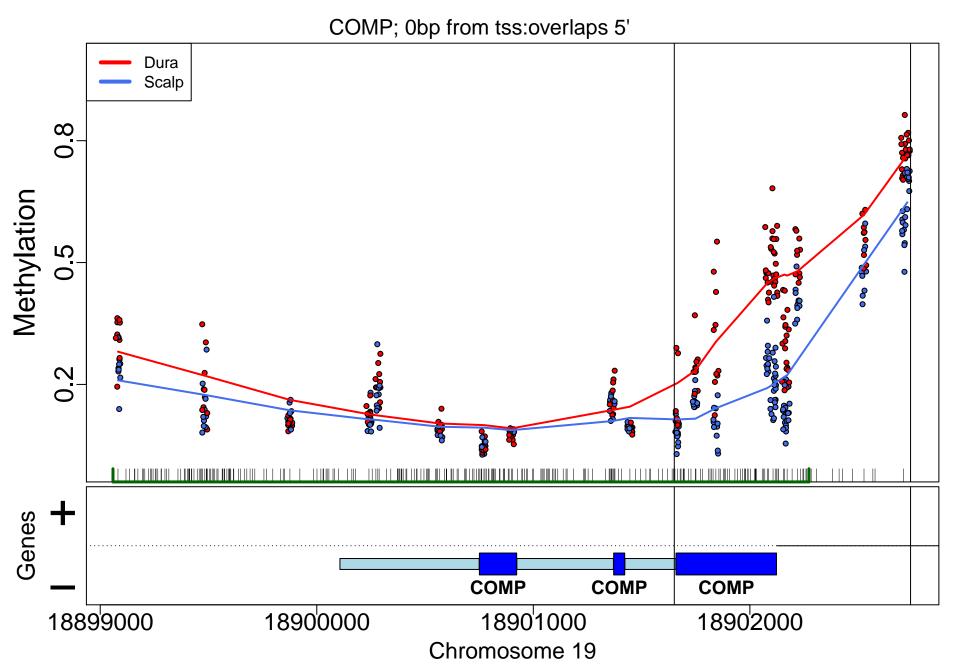


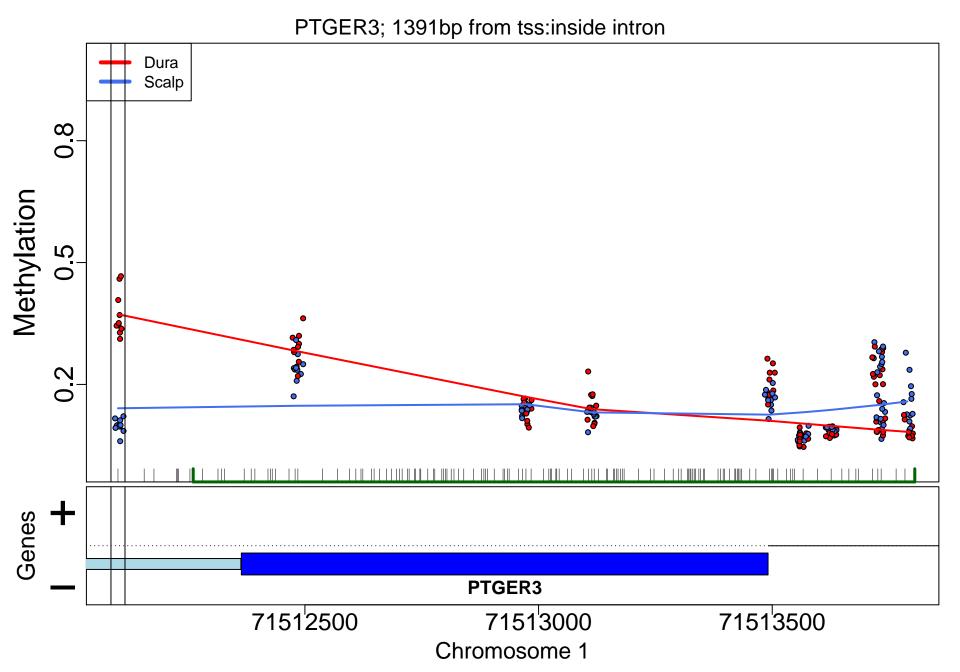


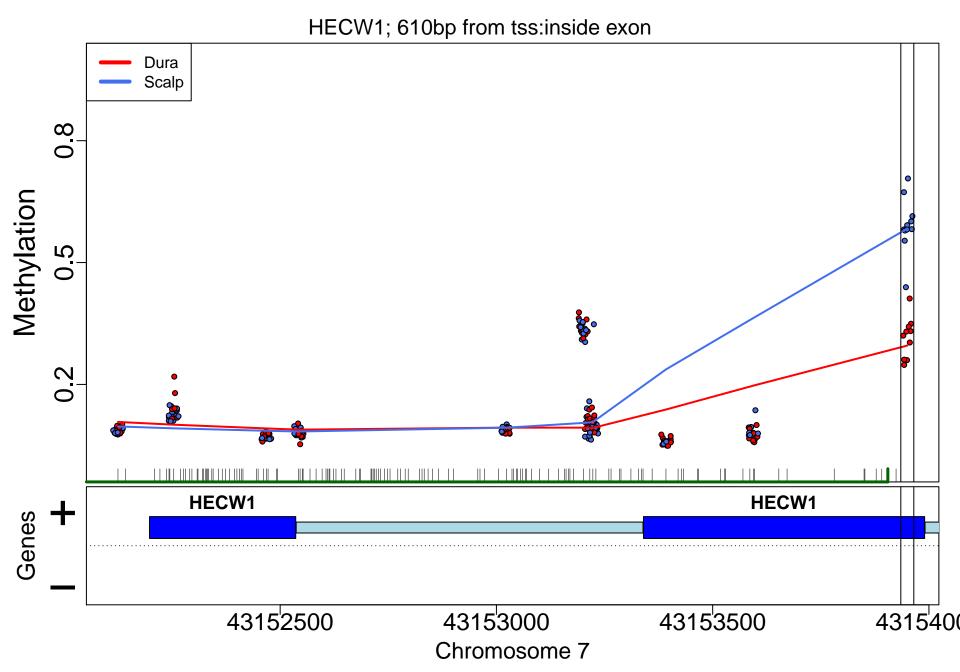


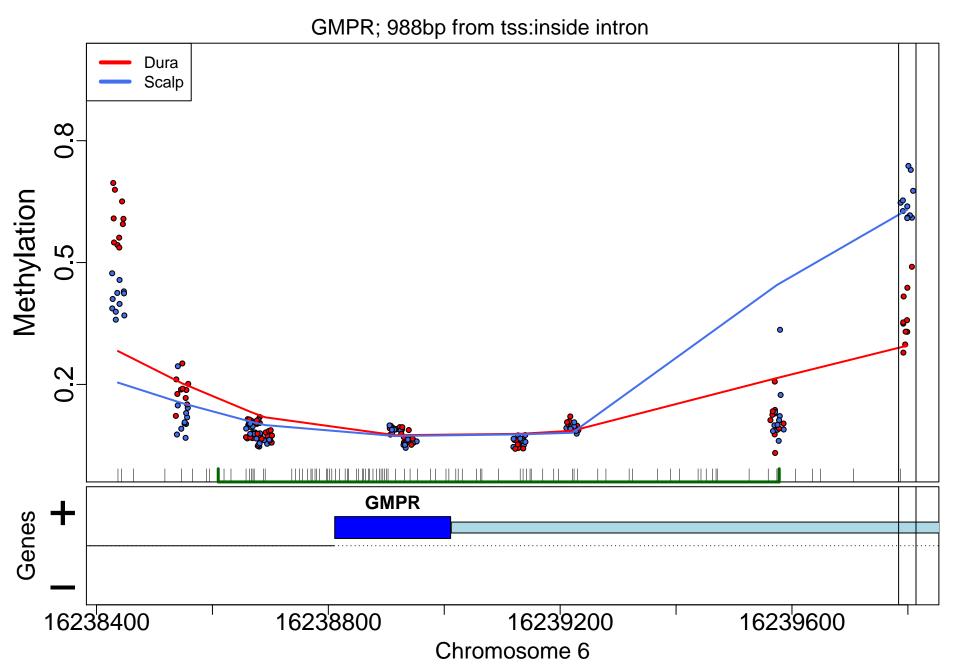


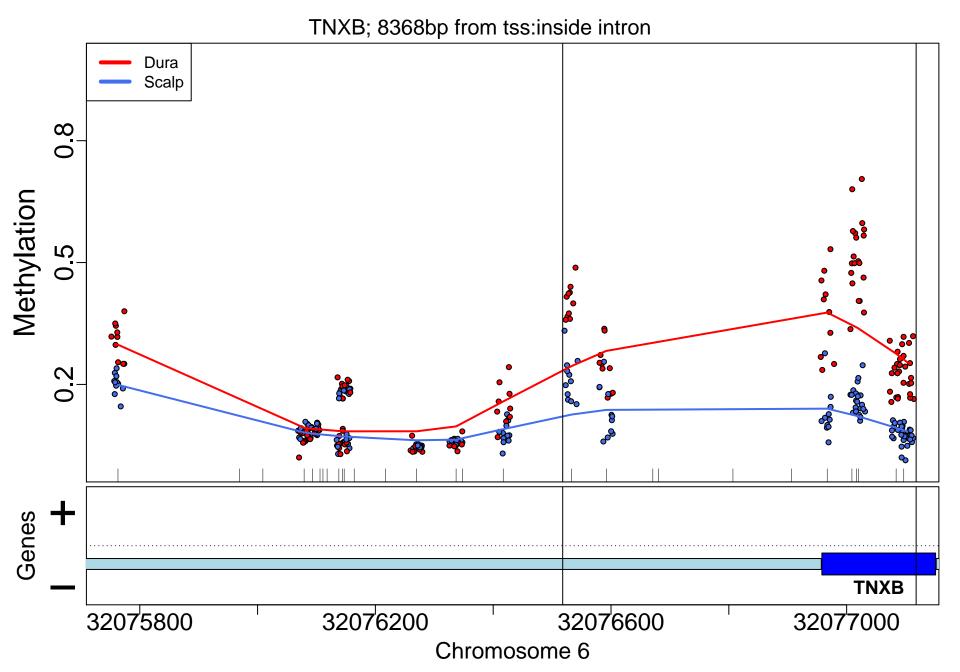


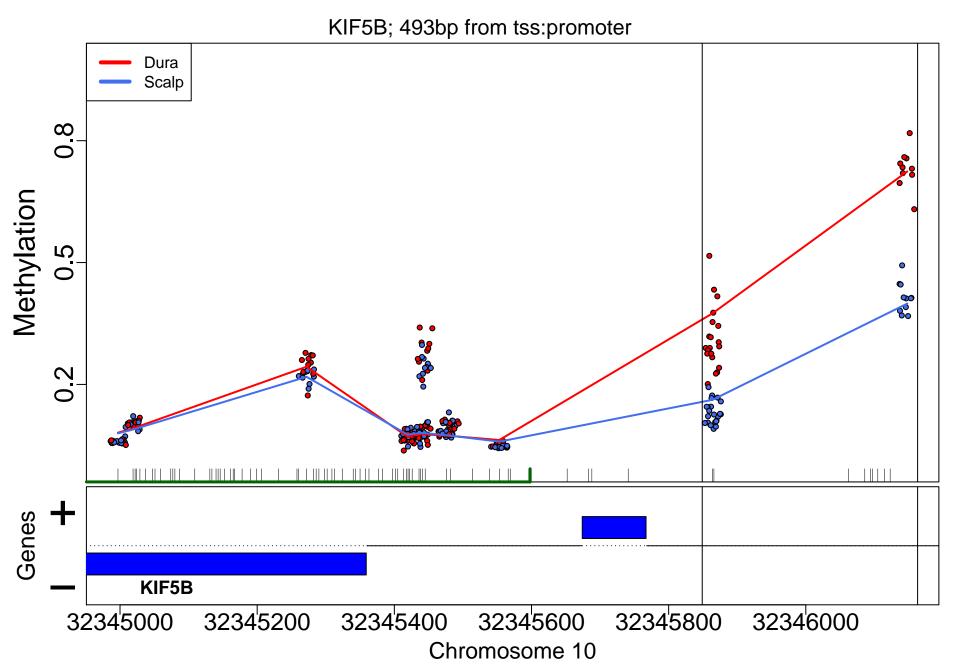


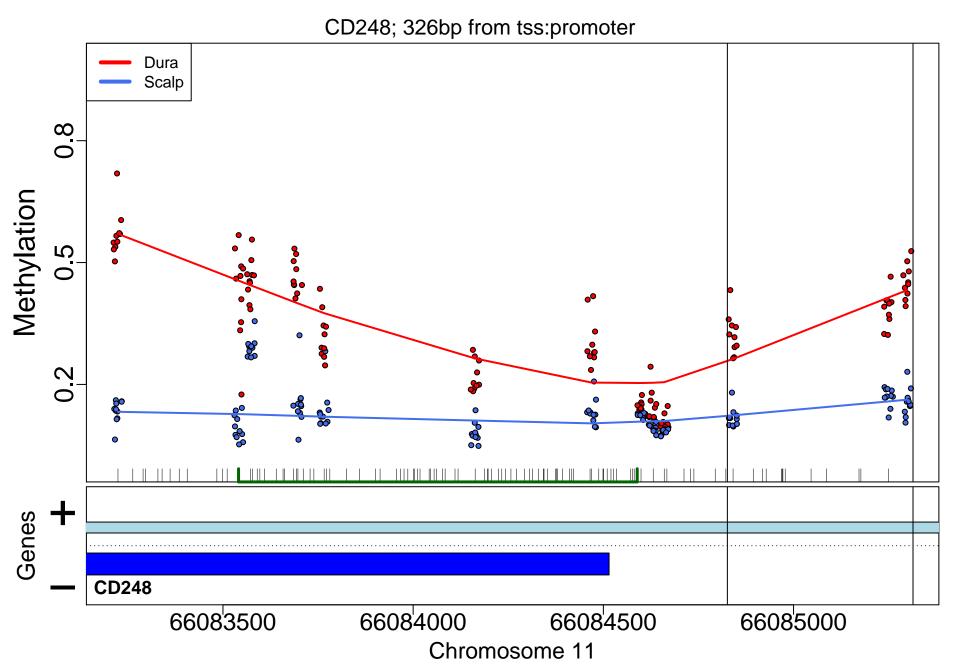


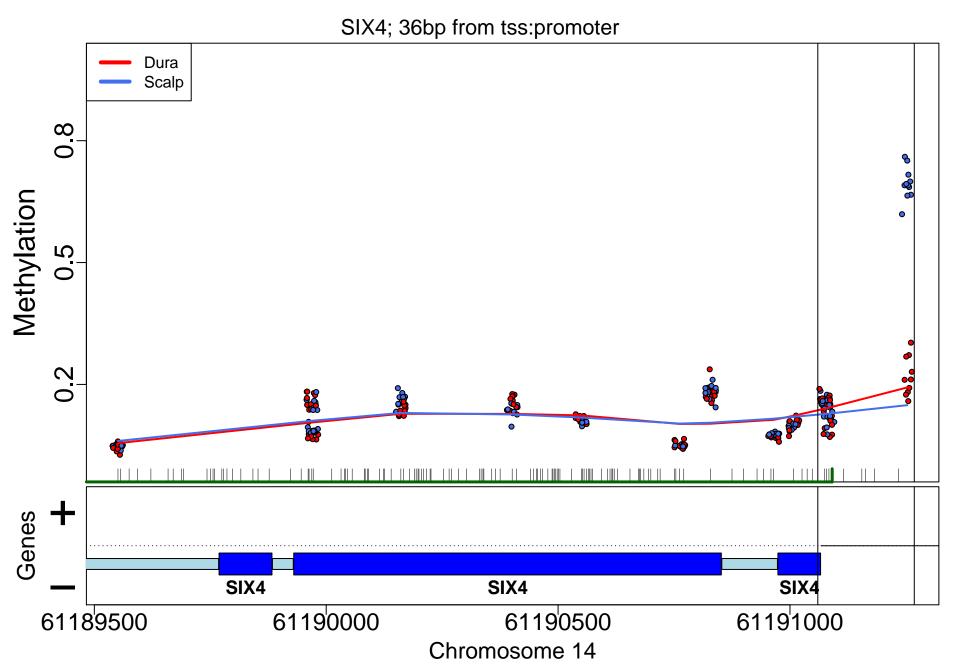


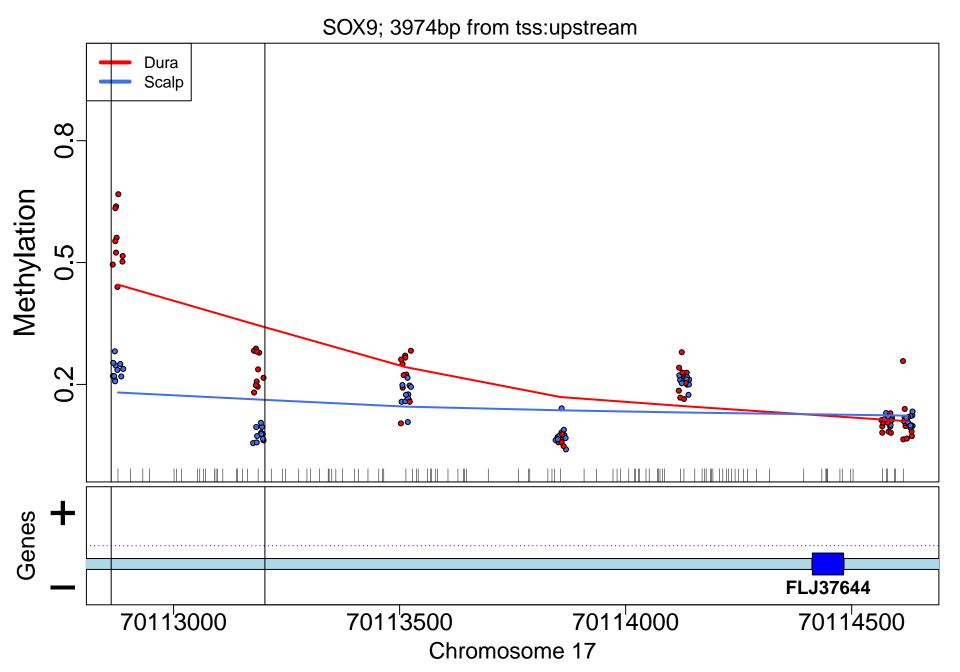


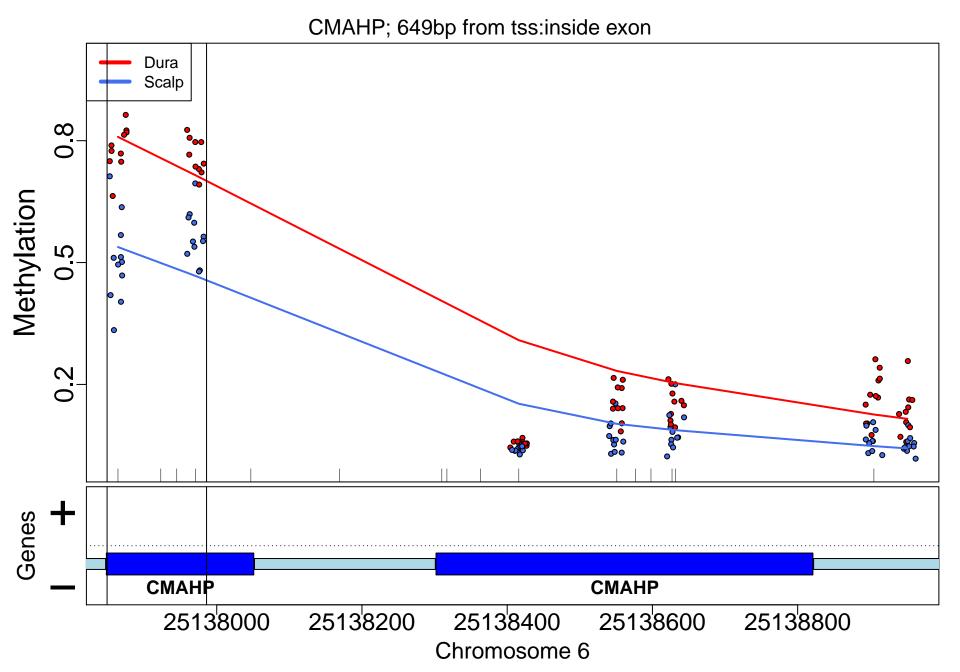


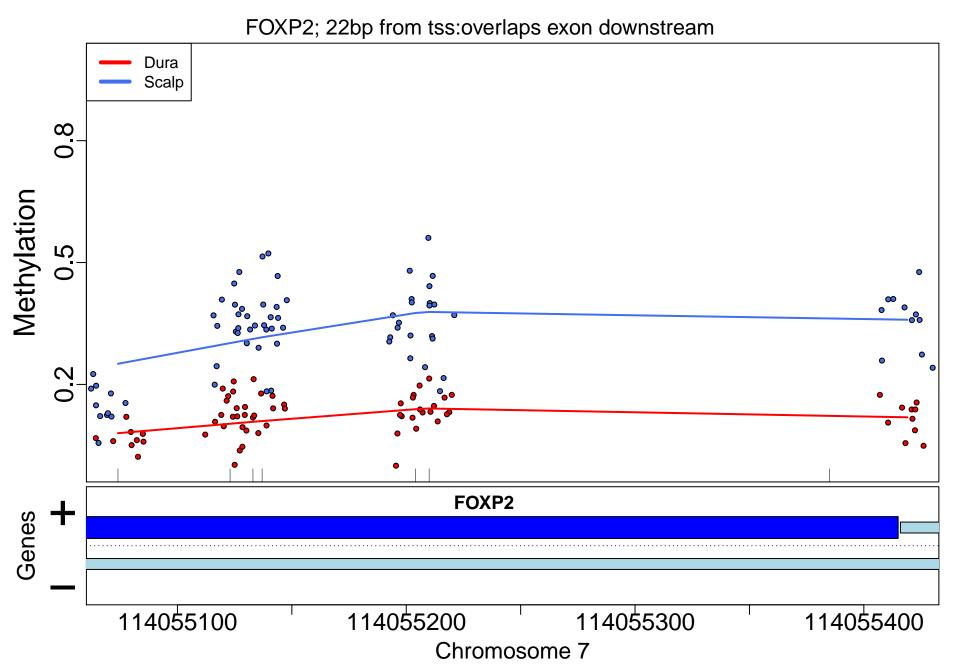


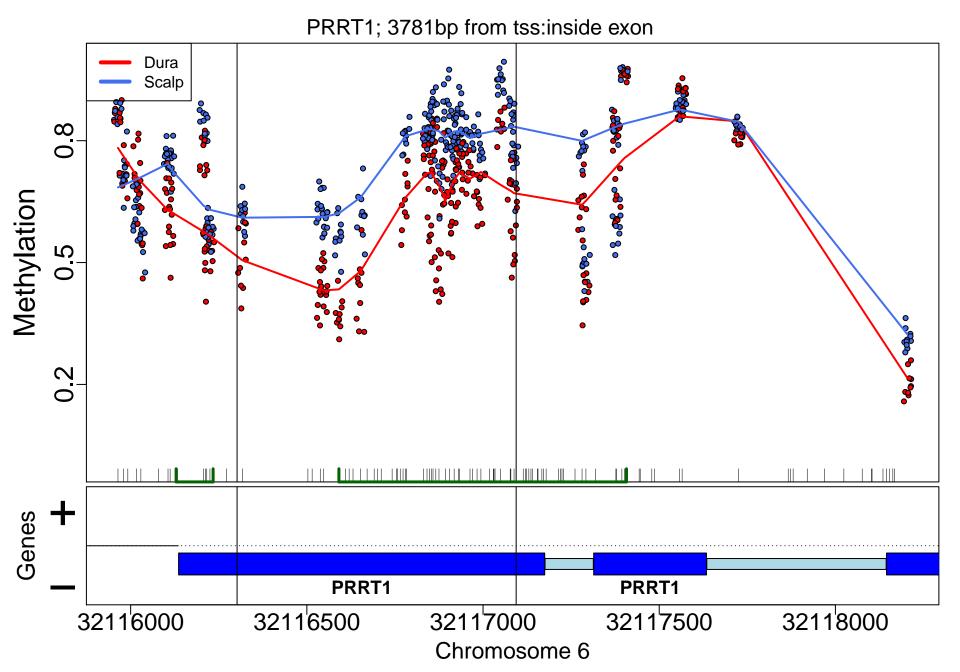


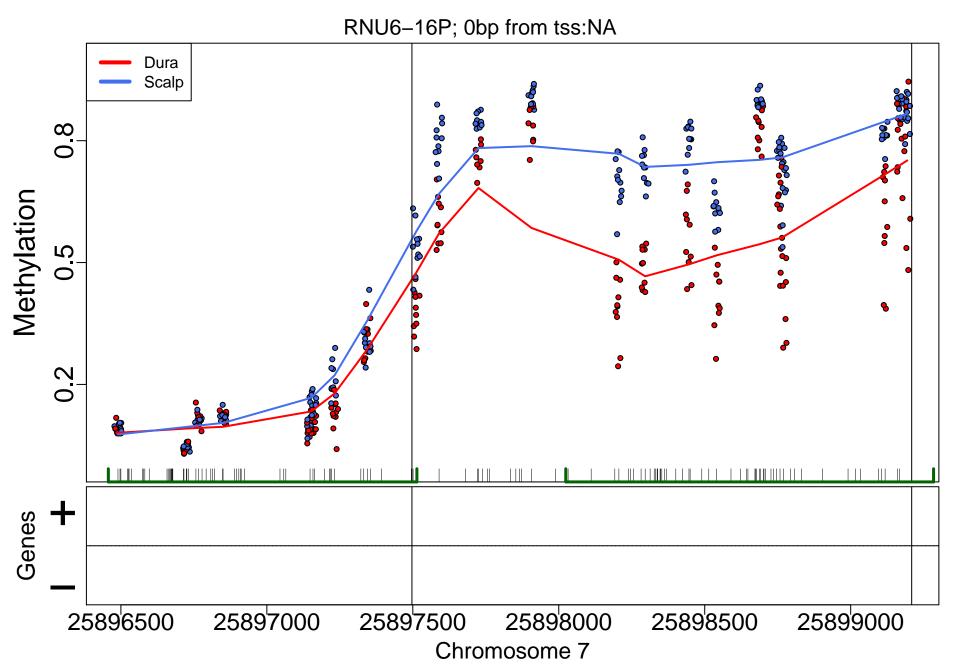


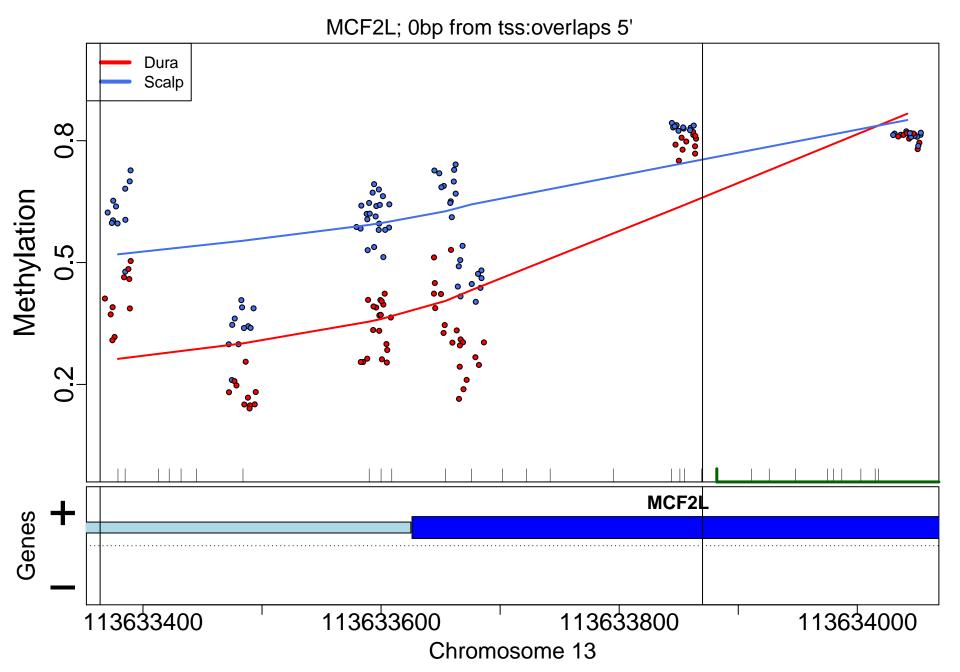


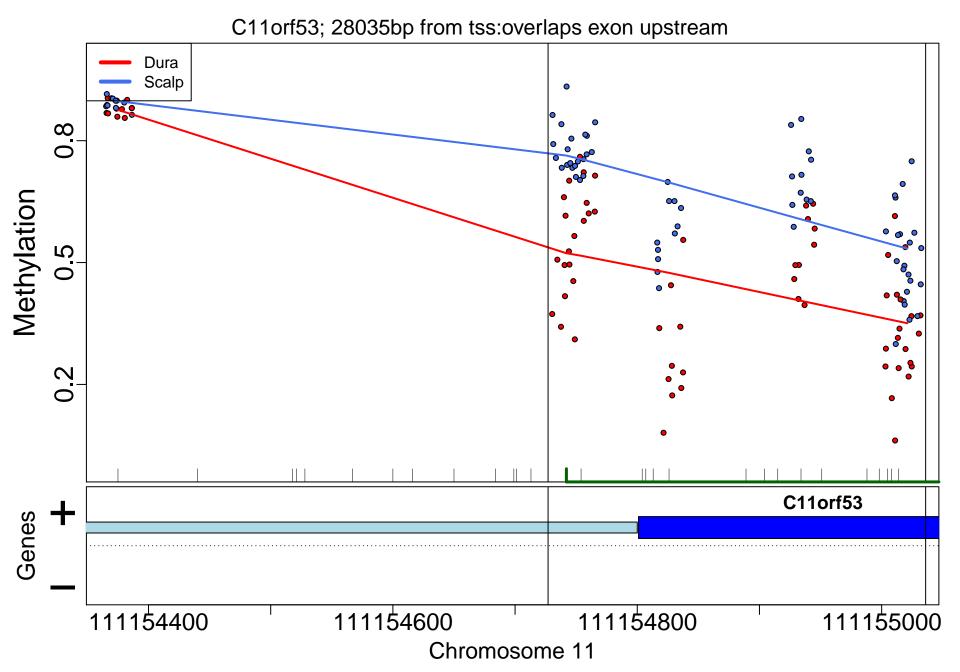


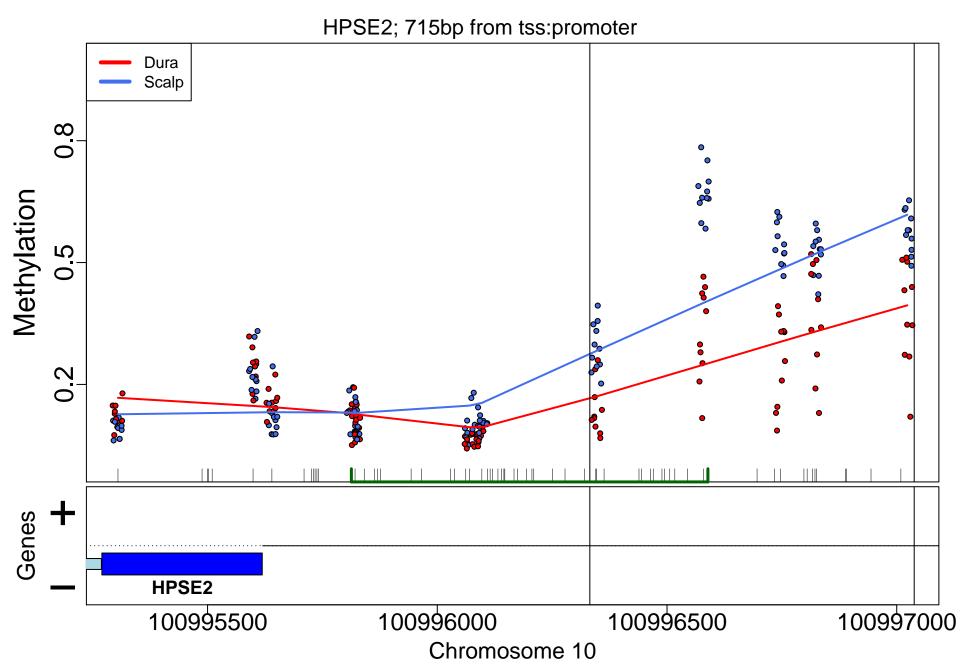


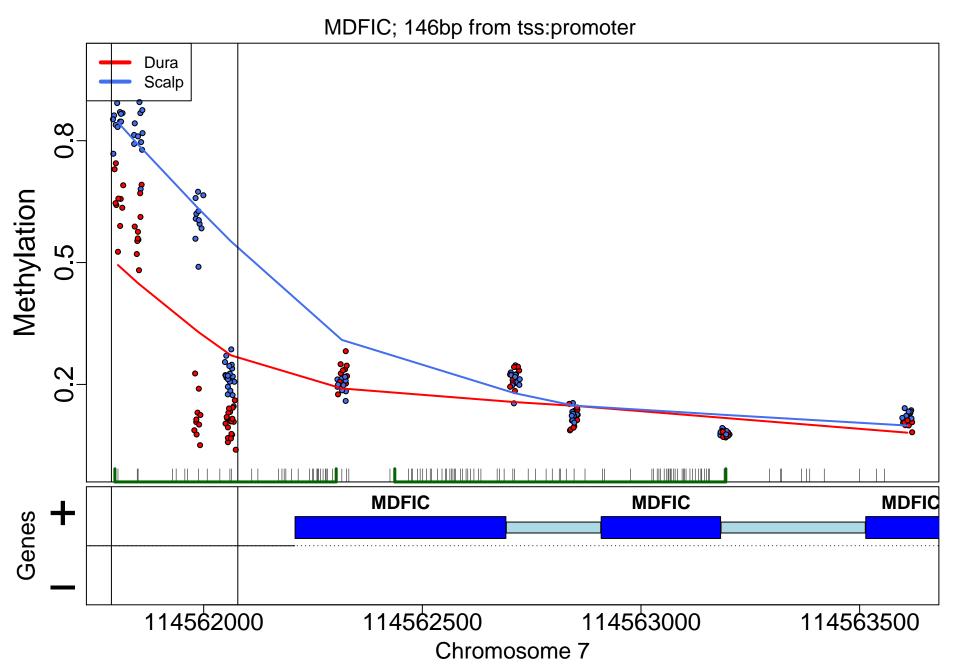


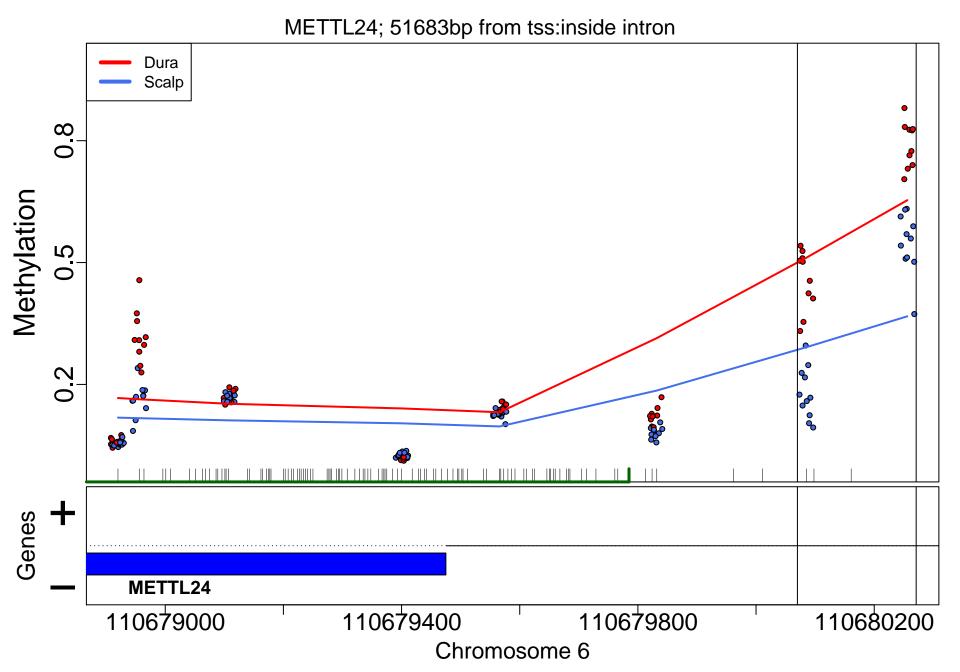


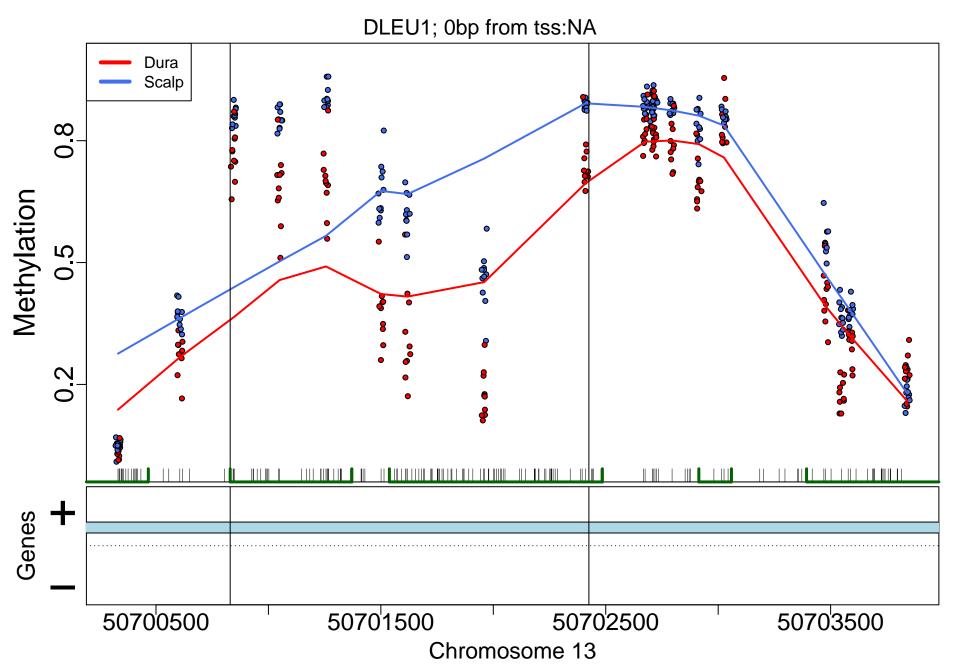


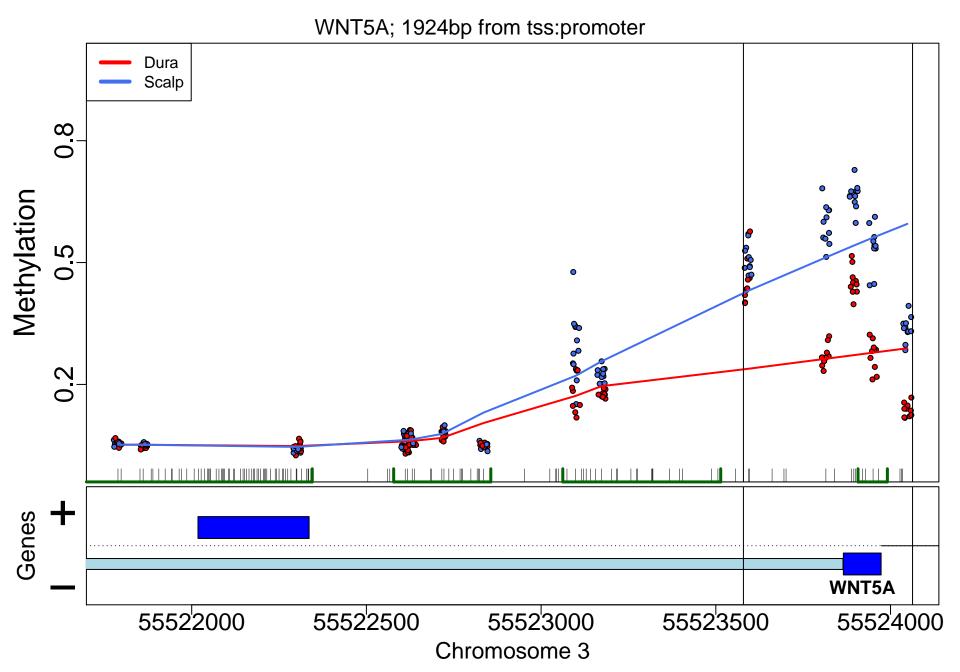


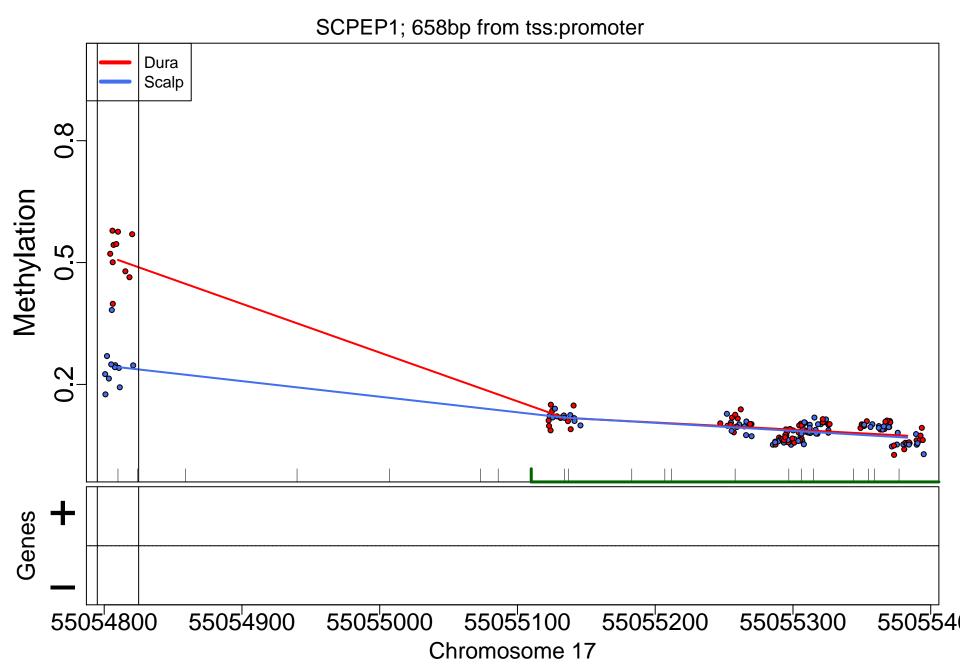


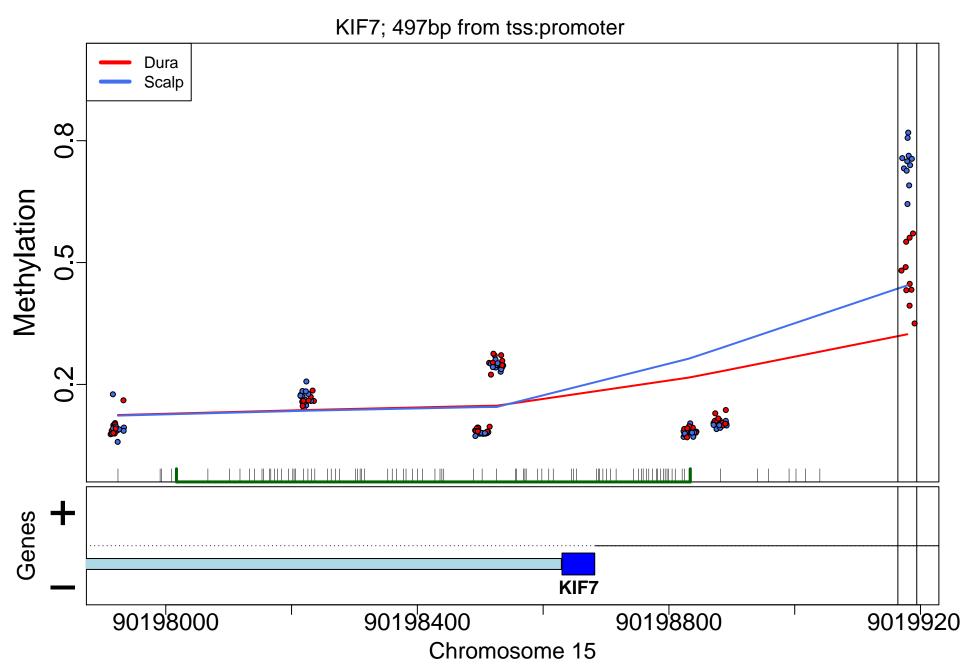












B3GALNT1; 155bp from tss:promoter Dura Scalp 80. Methylation 0.2 NMD3 Genes B3GALNT1 **B3GALNT1** 160823200 160822400 160822800 Chromosome 3

