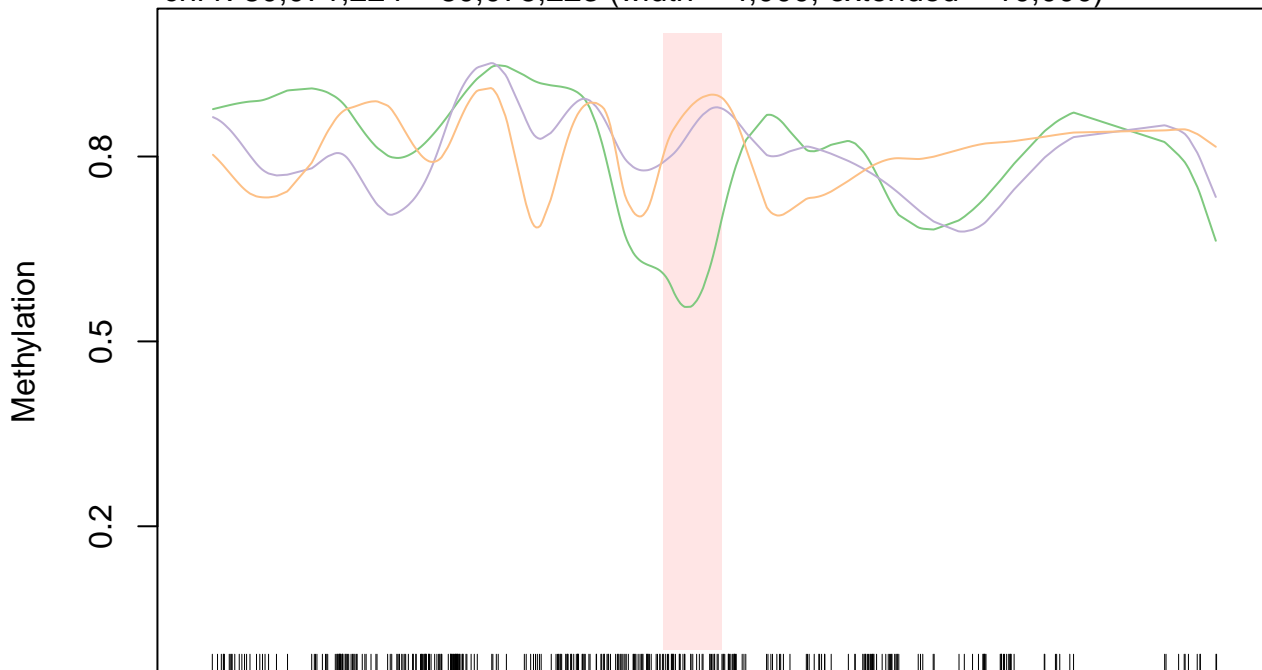


Transcript ENSRNOT00000021456.2 for gene Gipr
chr1: 80,071,224 – 80,075,223 (width = 4,000, extended = 10,000)



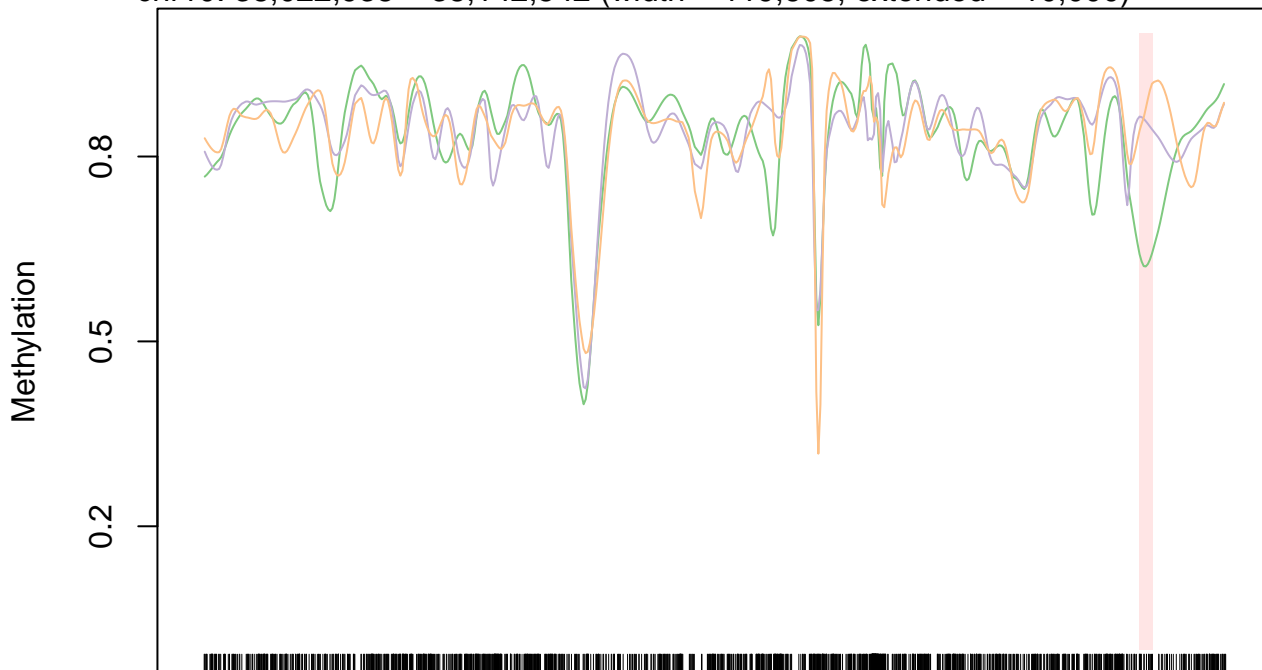
de



genes



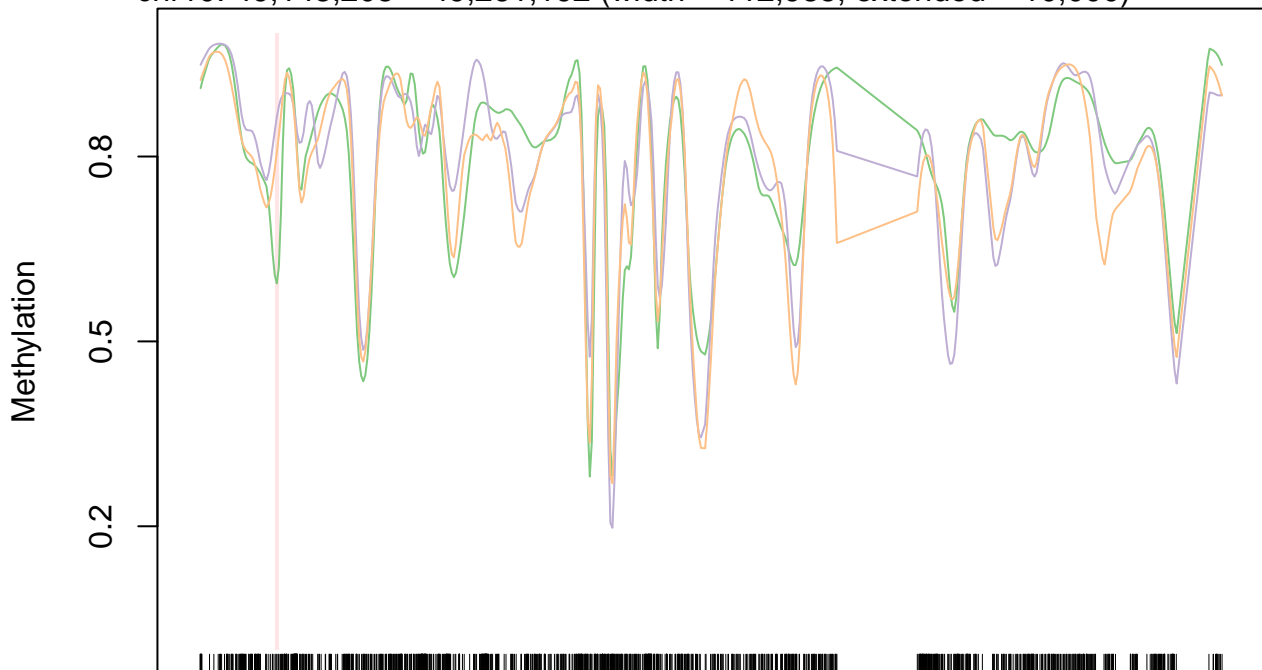
Transcript ENSRNOT00000042688.2 for gene LOC100362027
chr10: 38,022,938 – 38,142,542 (width = 119,605, extended = 10,000)



de

genes

Transcript ENSRNOT00000091100.1 for gene Fbxw10
chr10: 49,148,208 – 49,261,192 (width = 112,985, extended = 10,000)



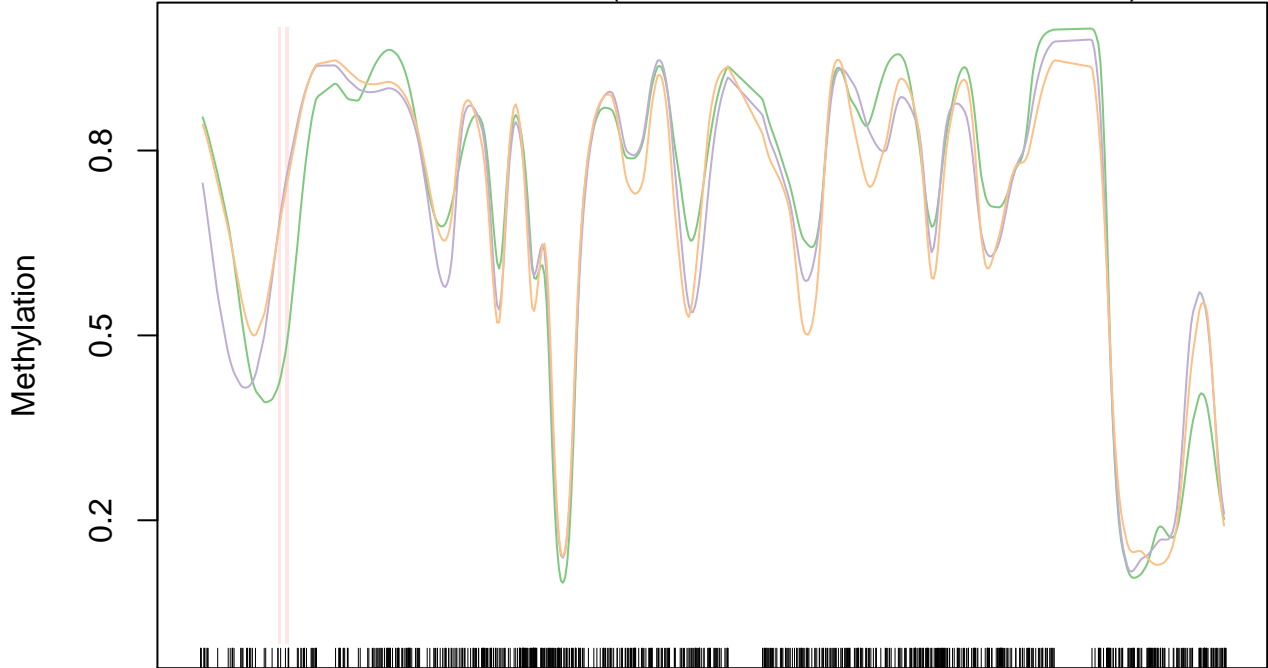
de



genes



Transcript ENSRNOT00000057724.4 for gene Plch1
chr2: 154,226,746 – 154,336,356 (width = 109,611, extended = 10,000)



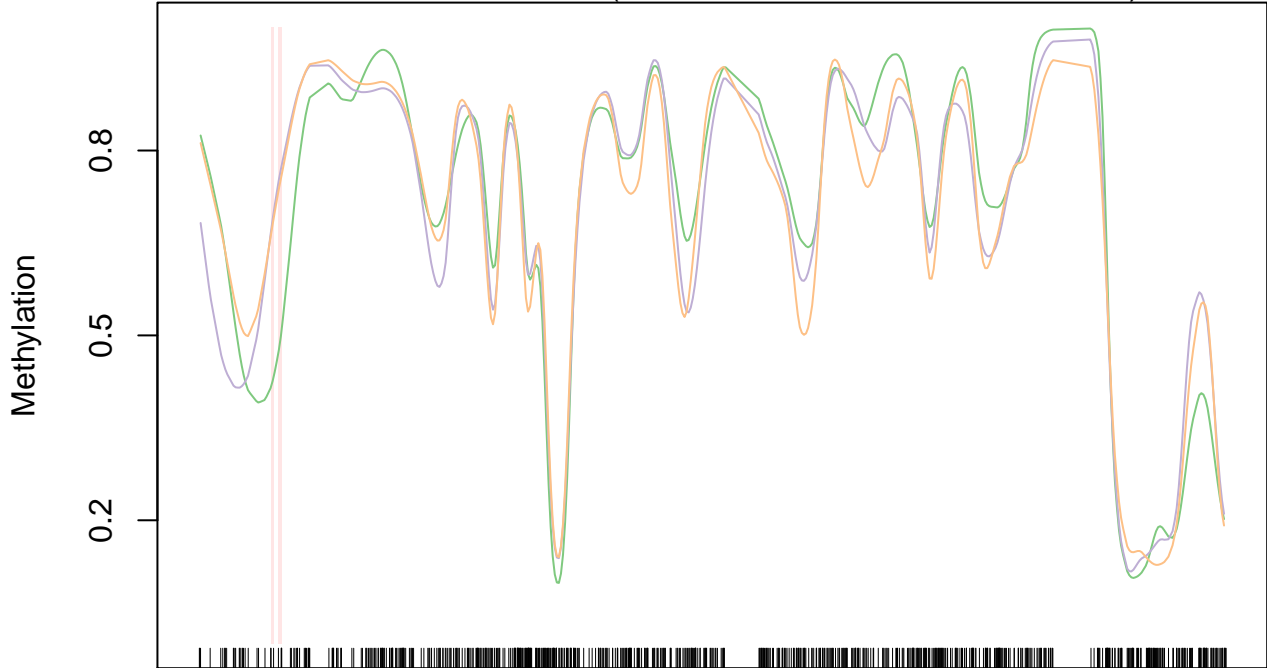
de



genes



Transcript ENSRNOT00000057724.4 for gene Plch1
chr2: 154,227,707 – 154,336,356 (width = 108,650, extended = 10,000)



de

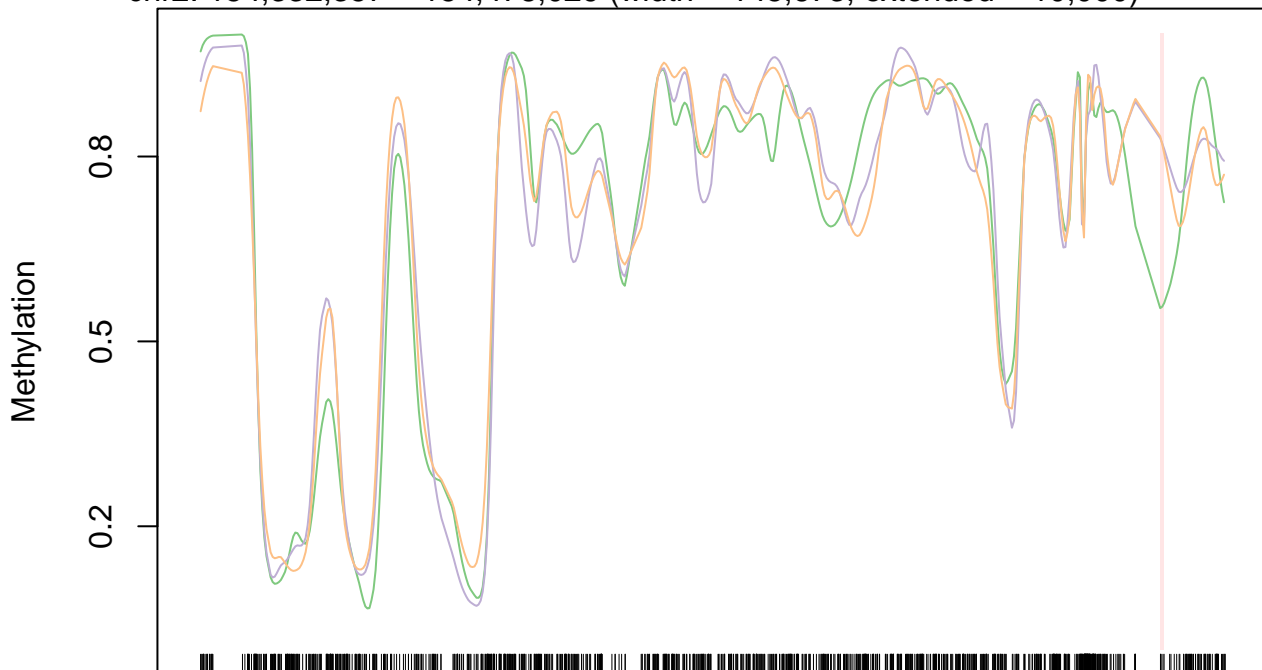


genes



Transcript ENSRNOT00000057724.4 for gene Plch1

chr2: 154,332,357 – 154,478,029 (width = 145,673, extended = 10,000)



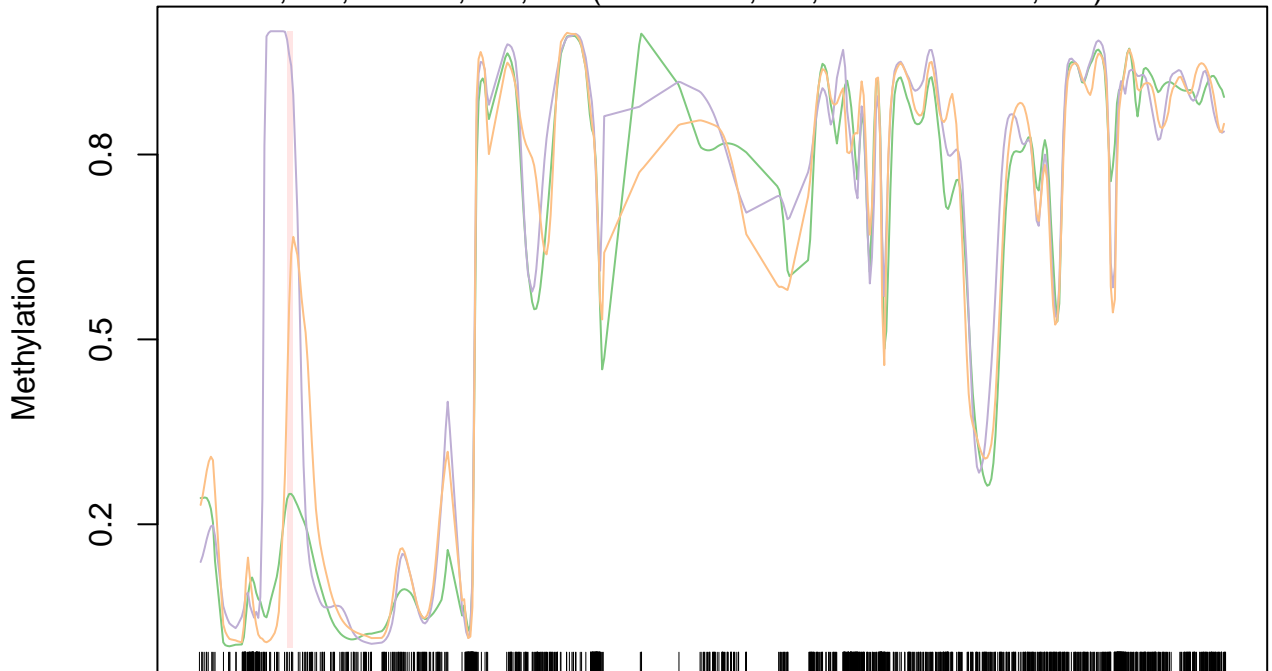
de



genes



Transcript ENSRNOT00000087861.1 for gene Vars
chr20: 4,900,369 – 4,995,688 (width = 95,320, extended = 10,000)



de



genes



Transcript ENSRNOT00000087883.1 for gene Foxj2

chr4: 155,600,141 – 155,656,909 (width = 56,769, extended = 10,000)

Methylation

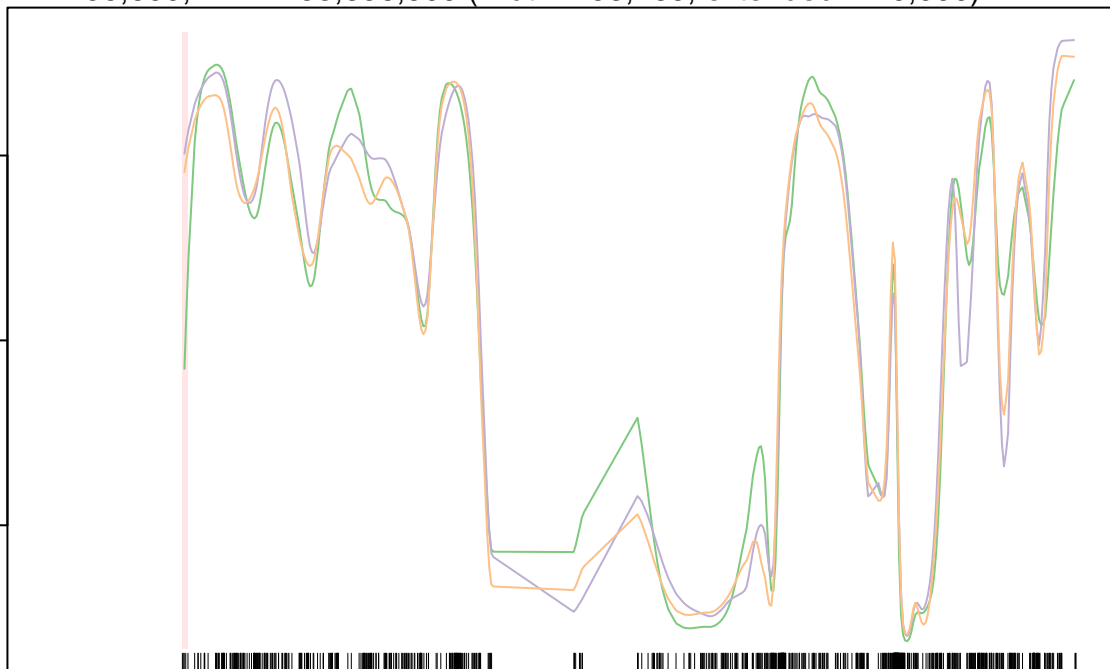
0.8

0.5

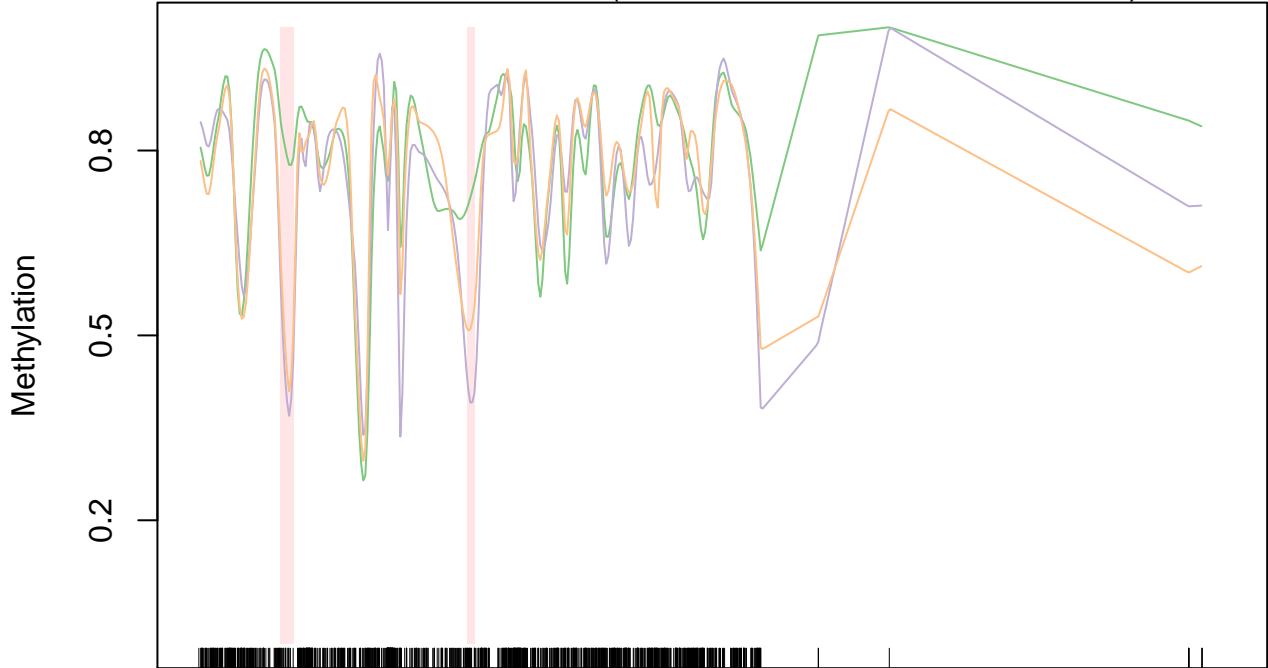
0.2

de

genes



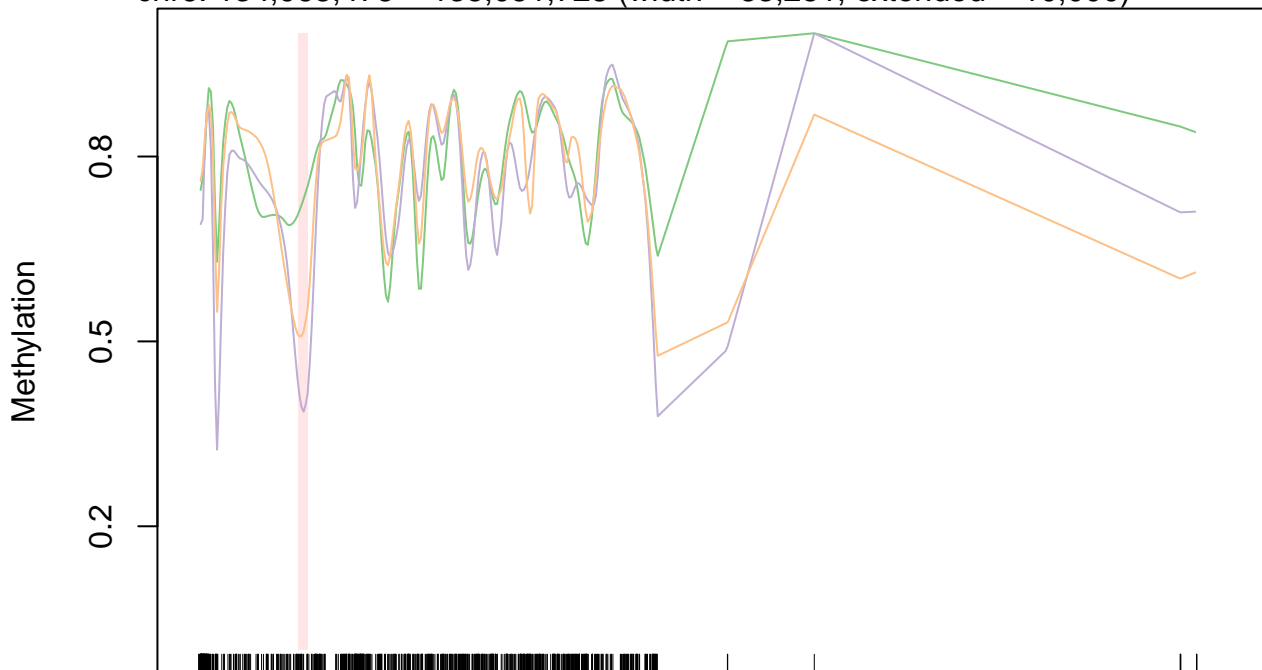
Transcript ENSRNOT00000009556.7 for gene Hsp90aa1
chr6: 134,945,484 – 135,051,728 (width = 106,245, extended = 10,000)



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genes

Transcript ENSRNOT00000009556.7 for gene Hsp90aa1
chr6: 134,968,478 – 135,051,728 (width = 83,251, extended = 10,000)



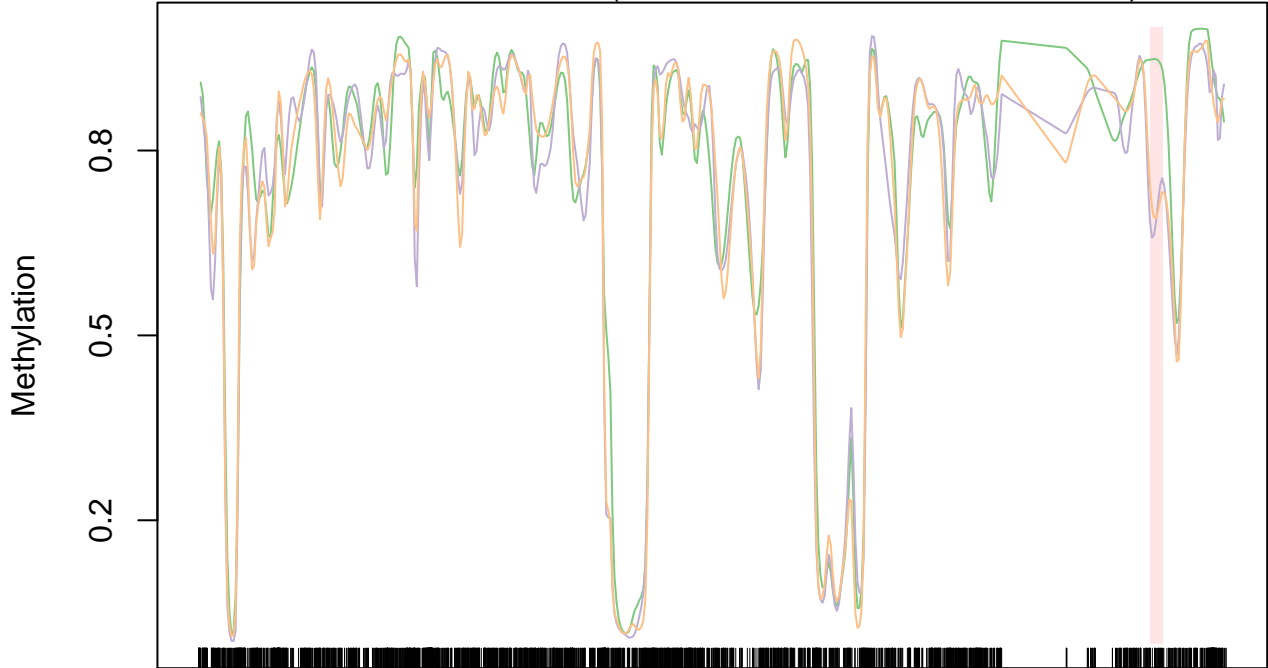
de



genes



Transcript ENSRNOT00000082699.1 for gene Adcy6
chr7: 140,288,797 – 140,430,348 (width = 141,552, extended = 10,000)



de 

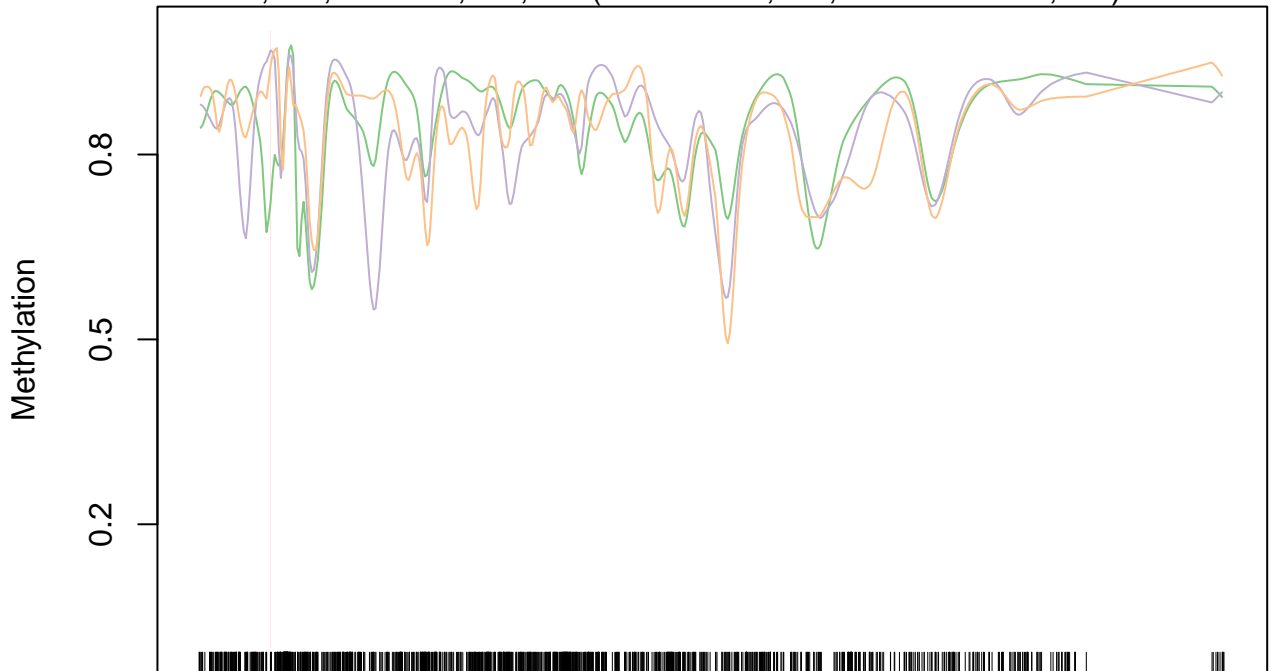
genes 







Transcript ENSRNOT00000087270.1 for gene Tcf12
chr8: 78,276,169 – 78,399,123 (width = 122,955, extended = 10,000)



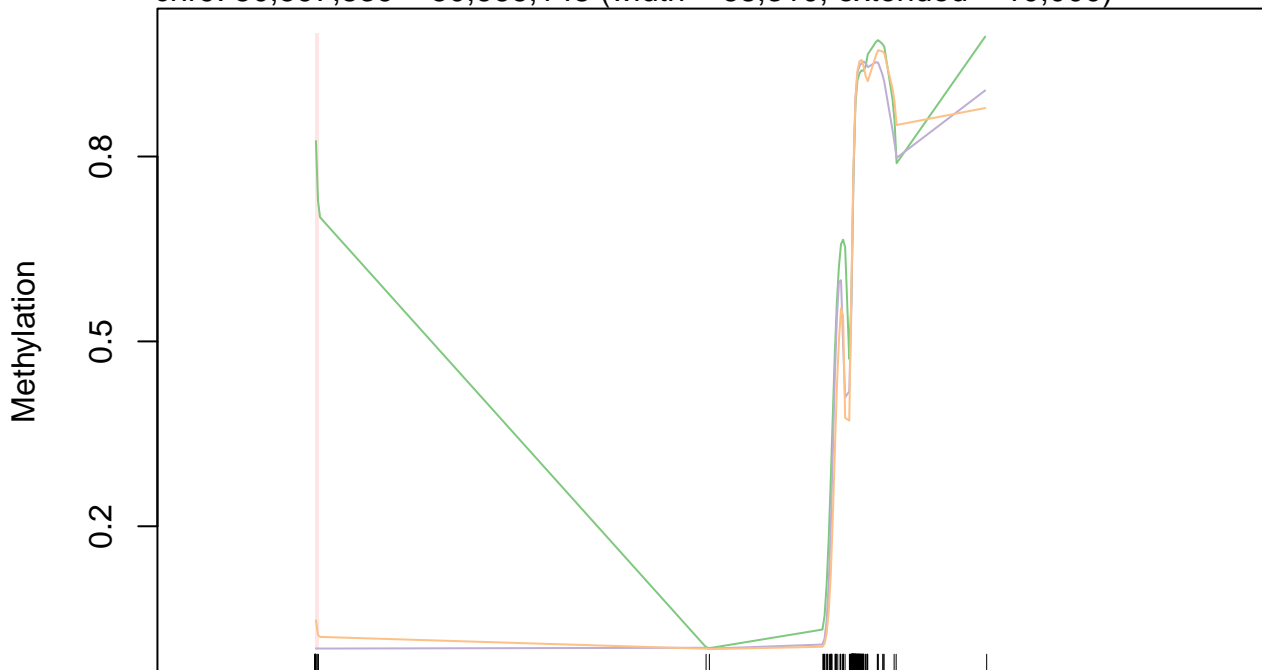
de



genes



Transcript ENSRNOT00000015393.5 for gene Tpp2
chr9: 50,597,839 – 50,666,148 (width = 68,310, extended = 10,000)



de

genes