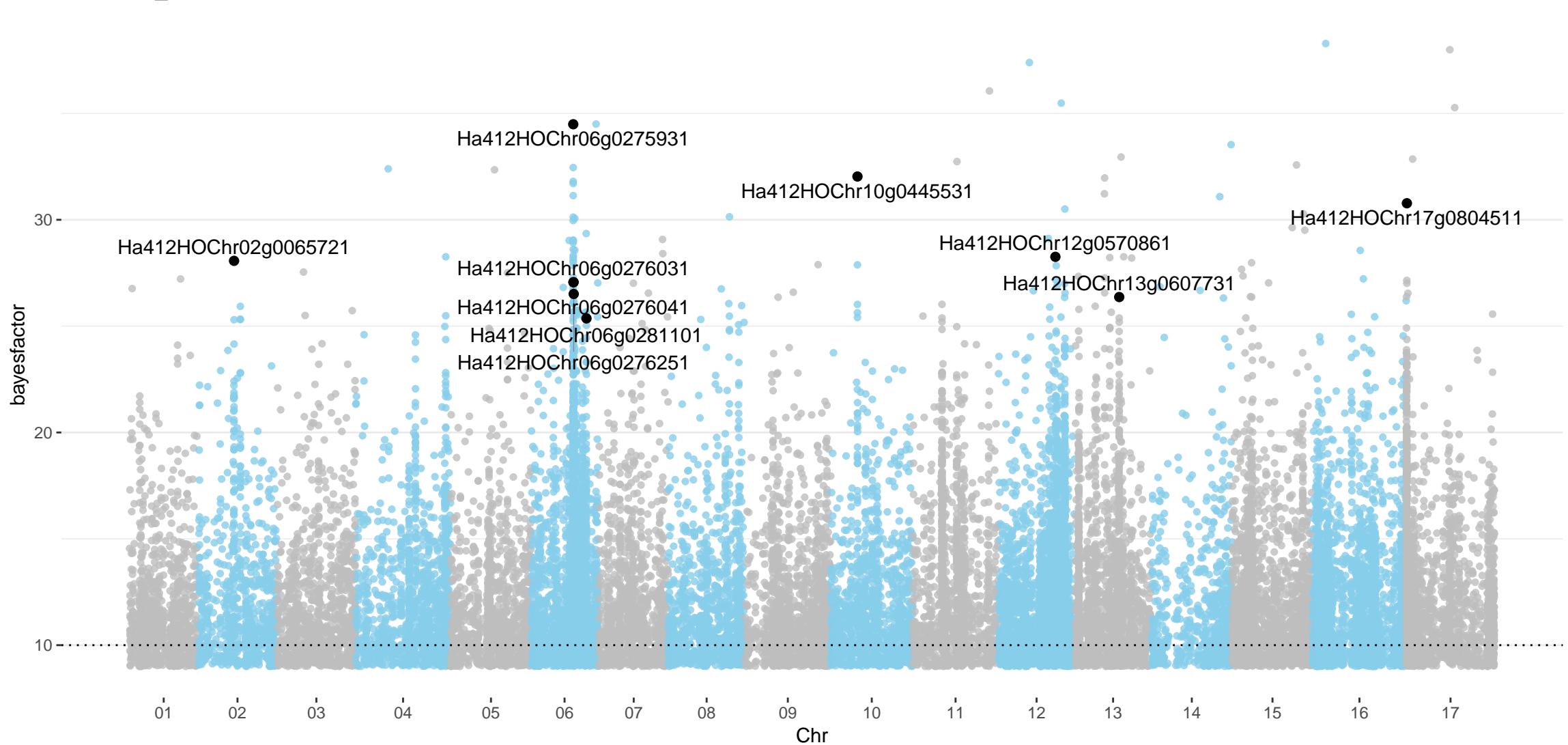
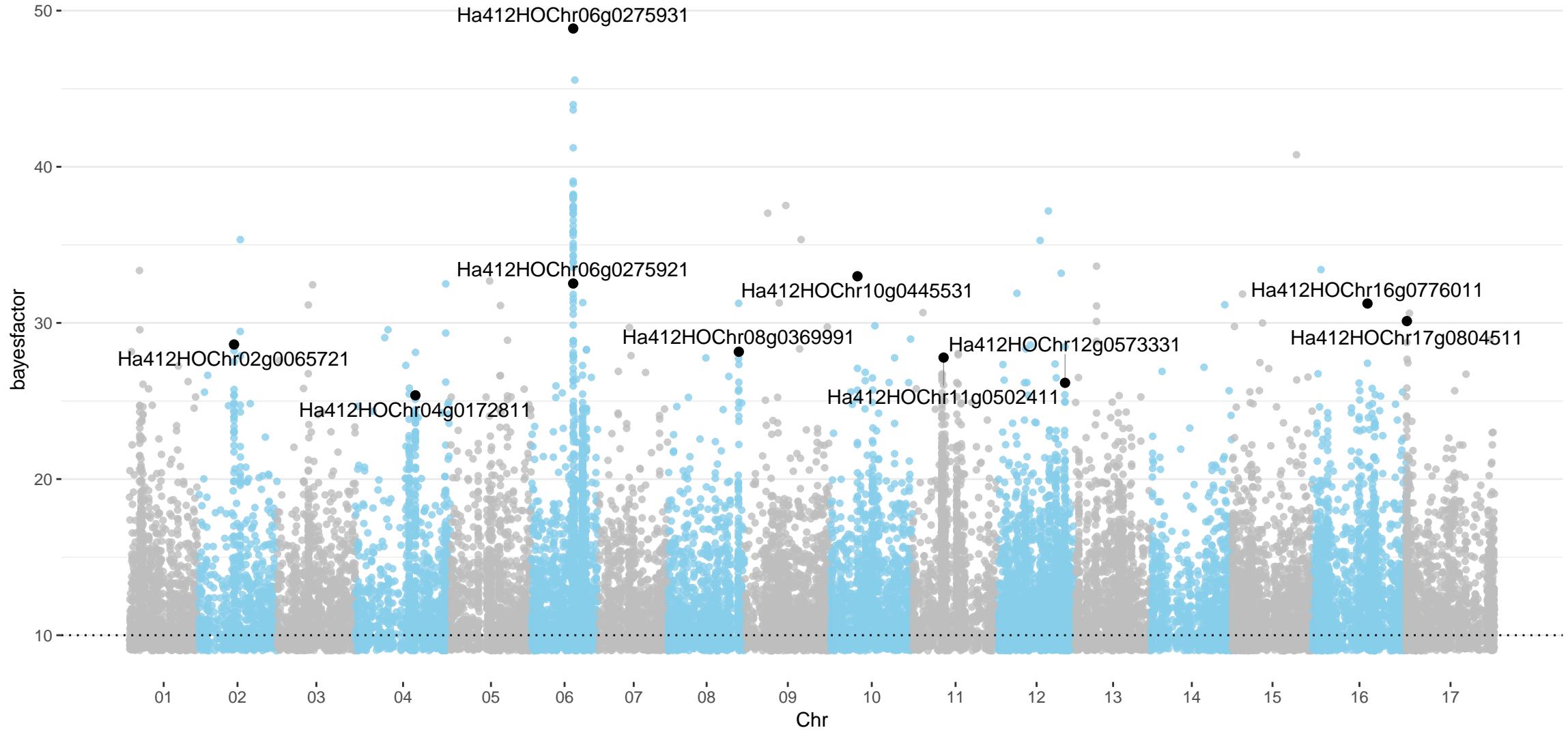


elevation\_e

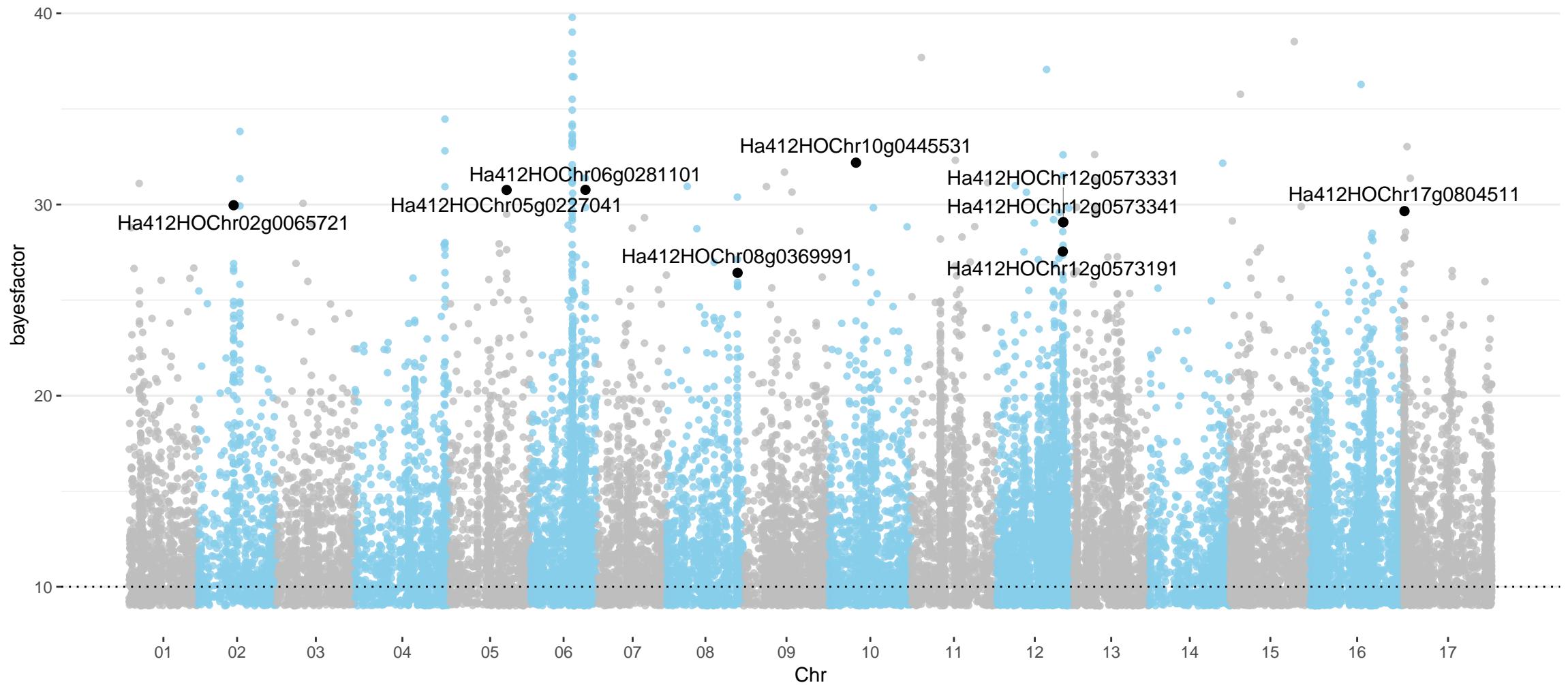


MAT\_e



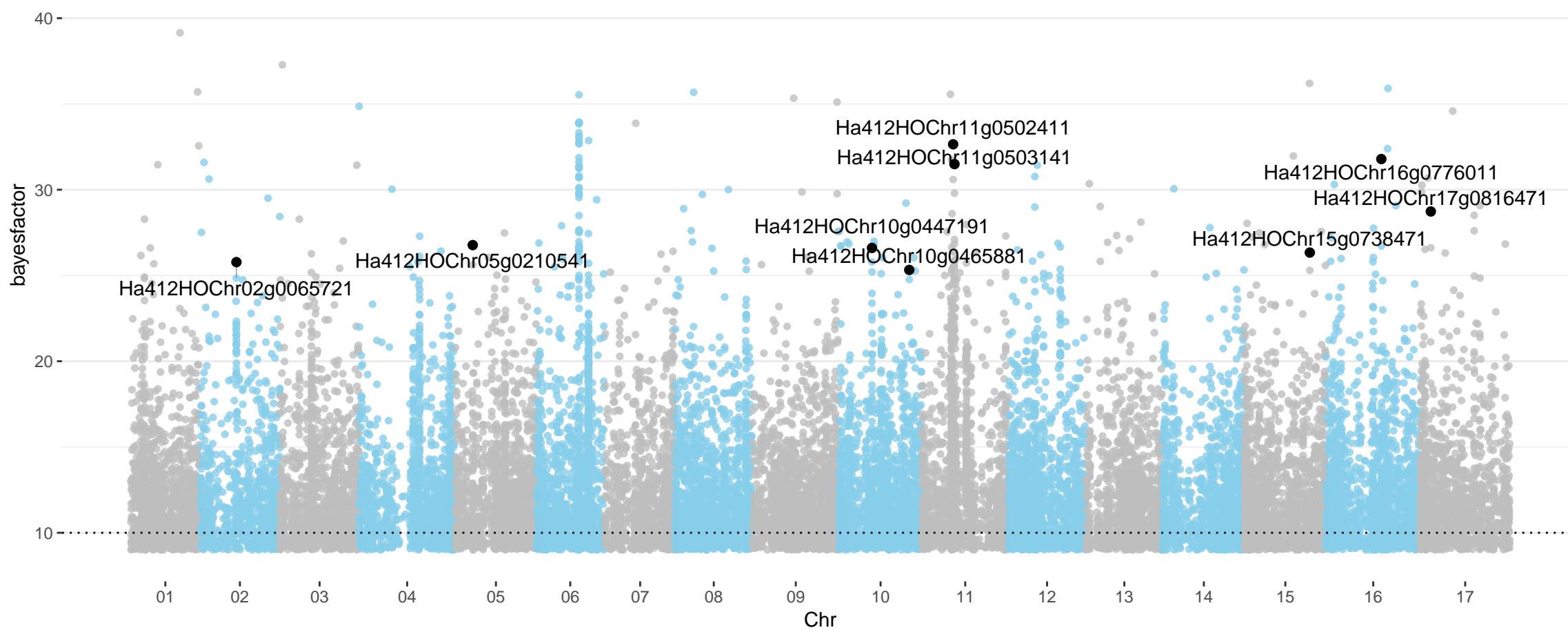
MWMT\_e

Ha412HOChr06g0275931

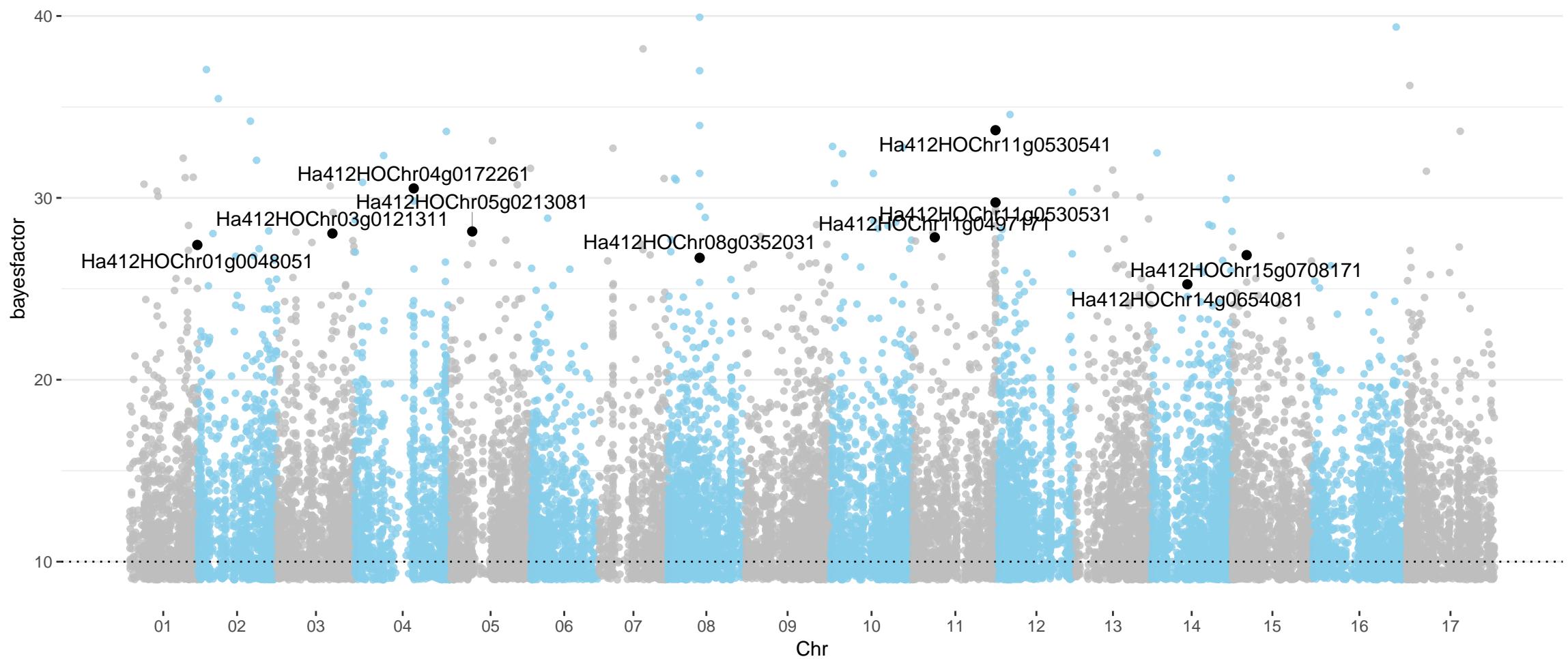


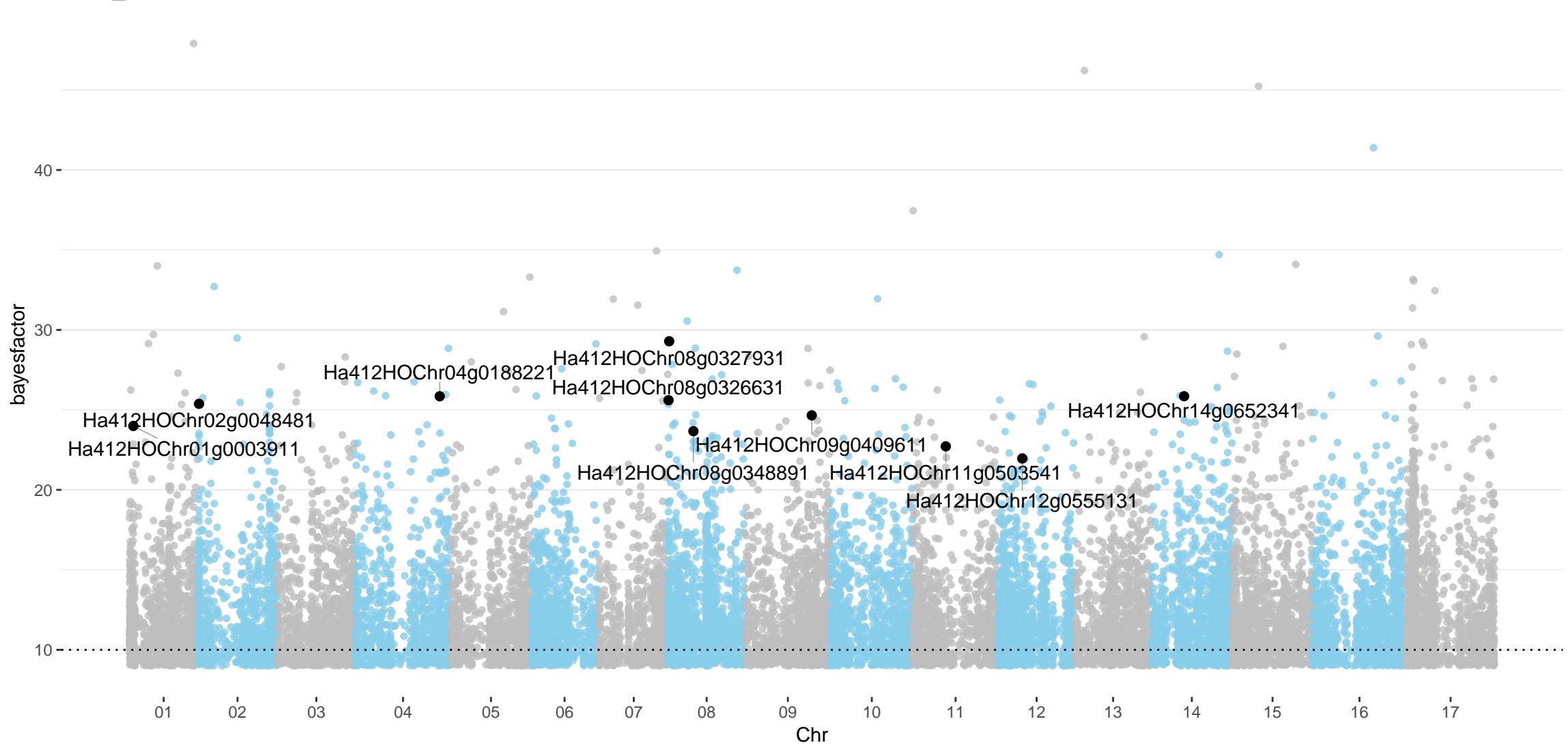
MCMT\_e

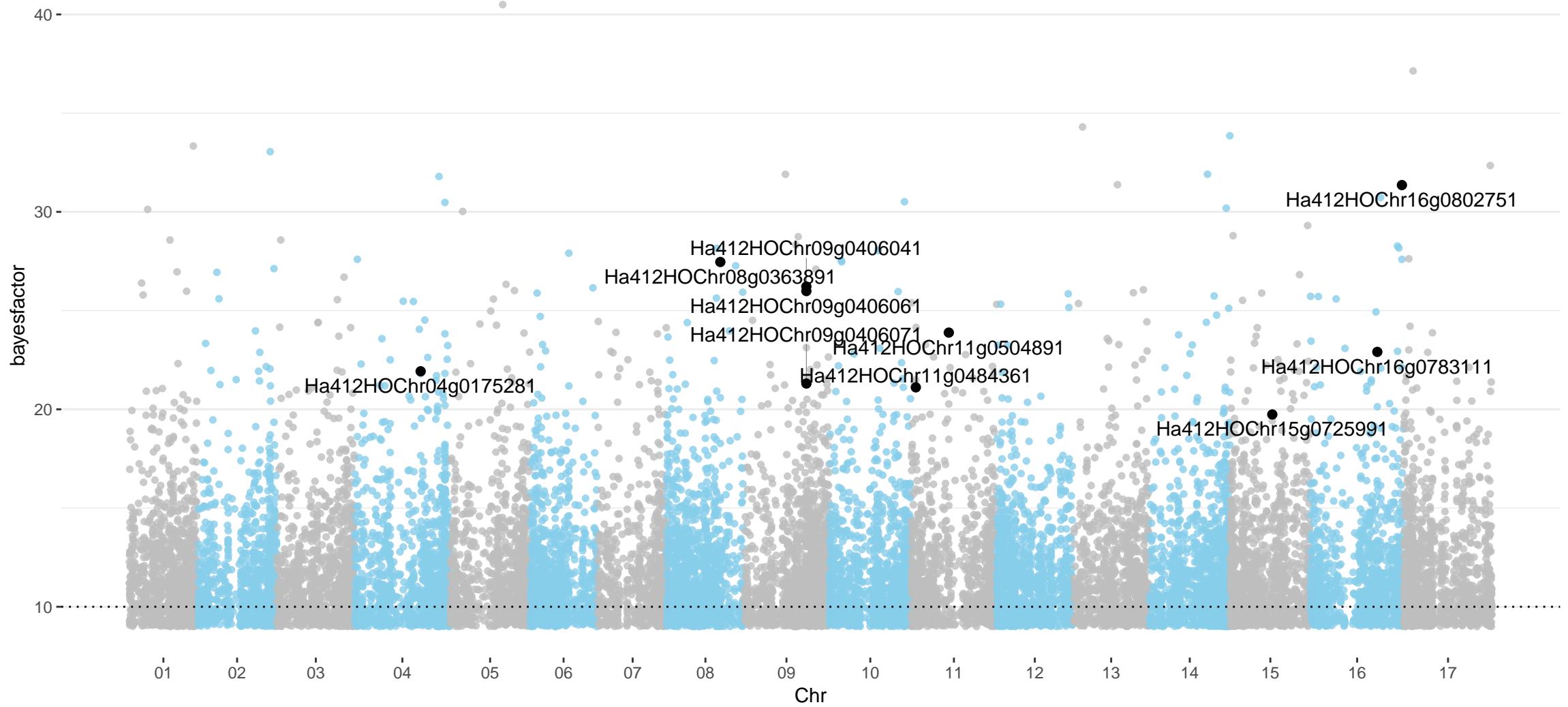
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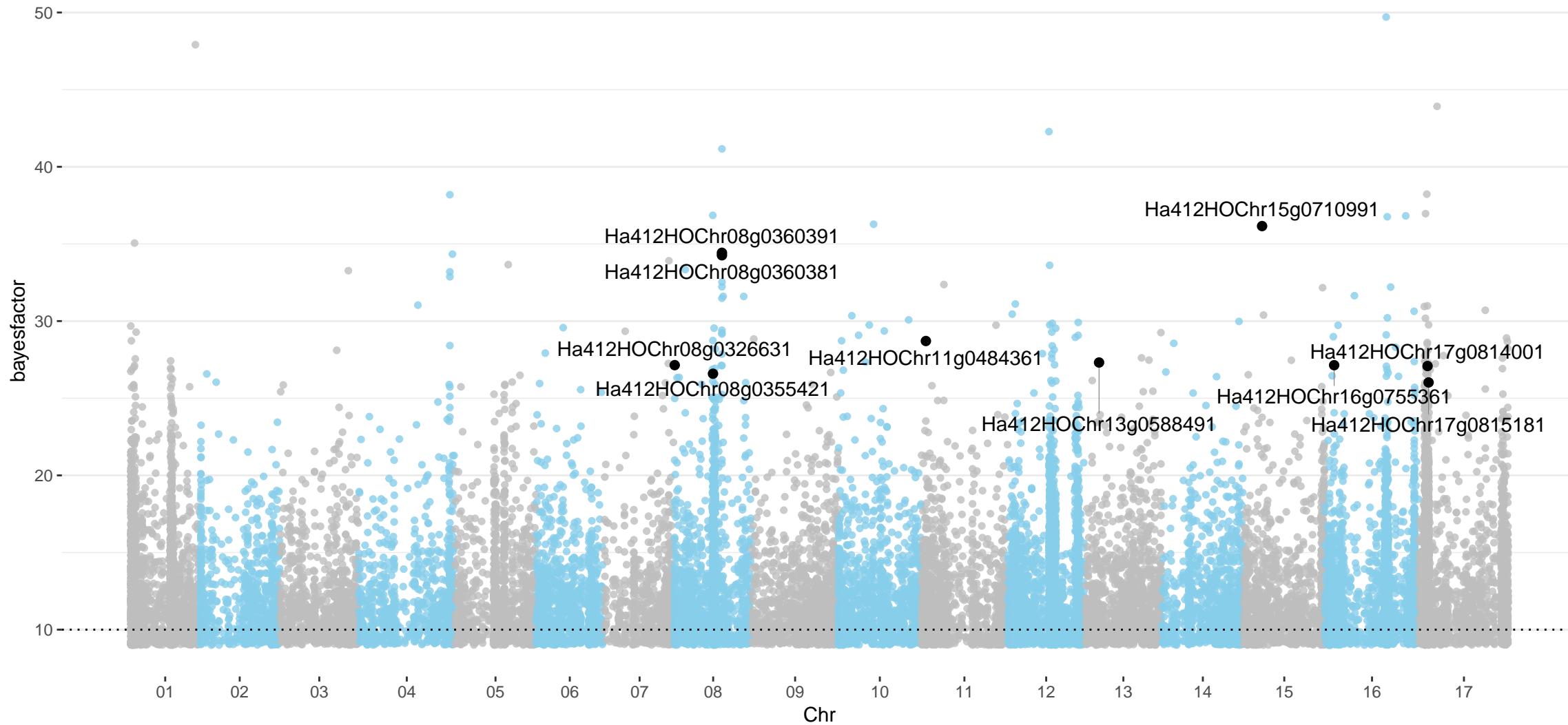
TD\_e



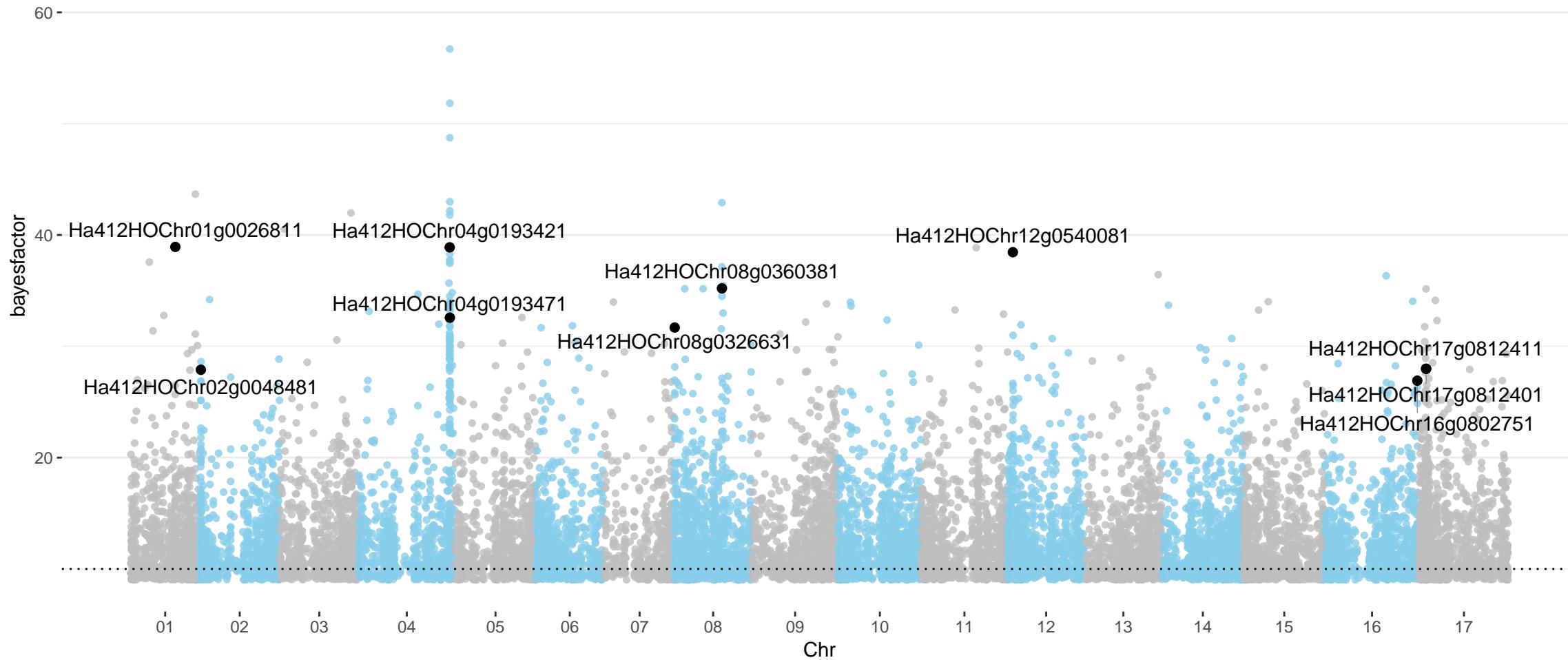




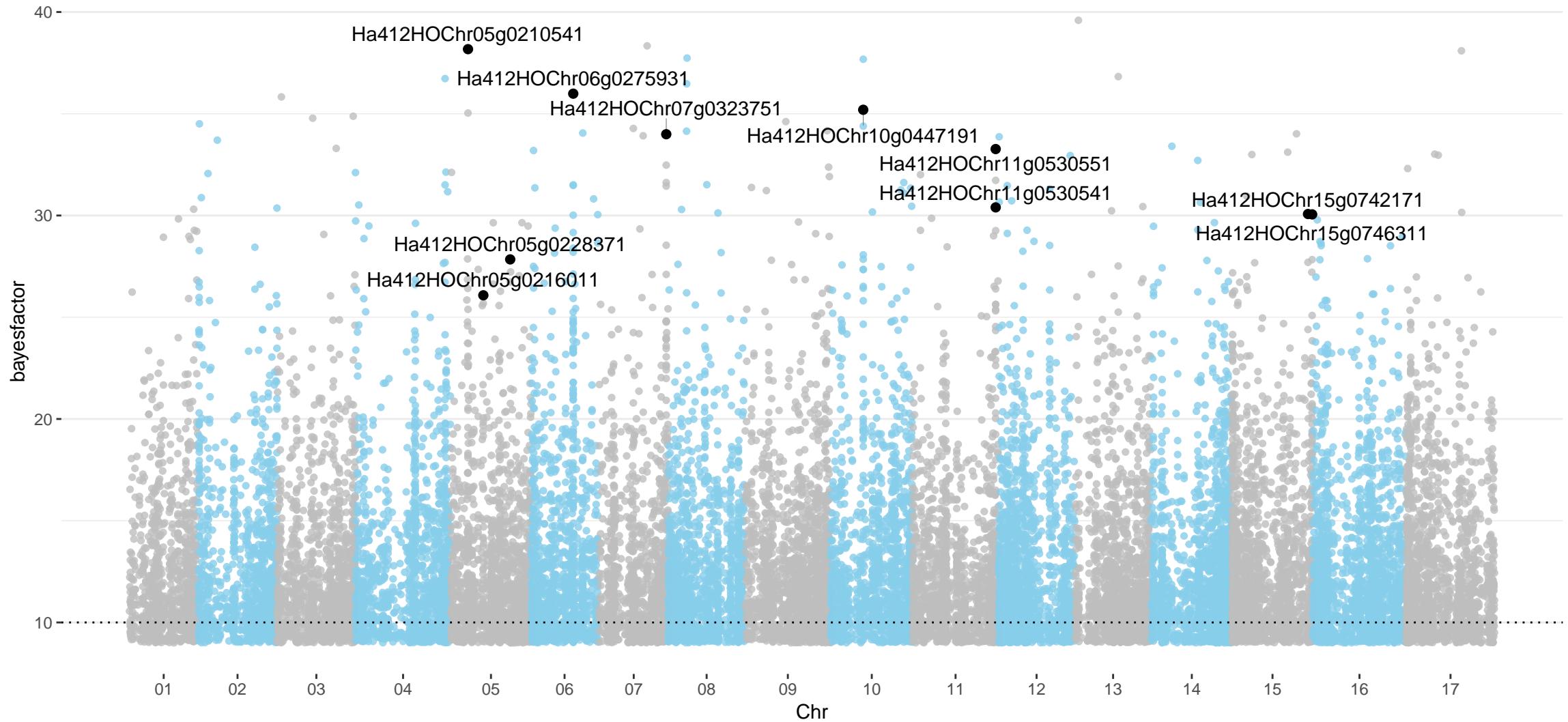
## AHM\_e



SHM\_e



DD\_0\_e



DD5\_e

Ha412HOChr06g0275931

Ha412HOChr10g0445531

Ha412HOChr01g0006621

Ha412HOChr01g0006601  
Ha412HOChr02g0065721

Ha412HOChr01g0006831

Ha412HOChr06g0275921

Ha412HOChr05g0227041  
Ha412HOChr06g0279821

Ha412HOChr08g0369991

bayesfactor

40

20

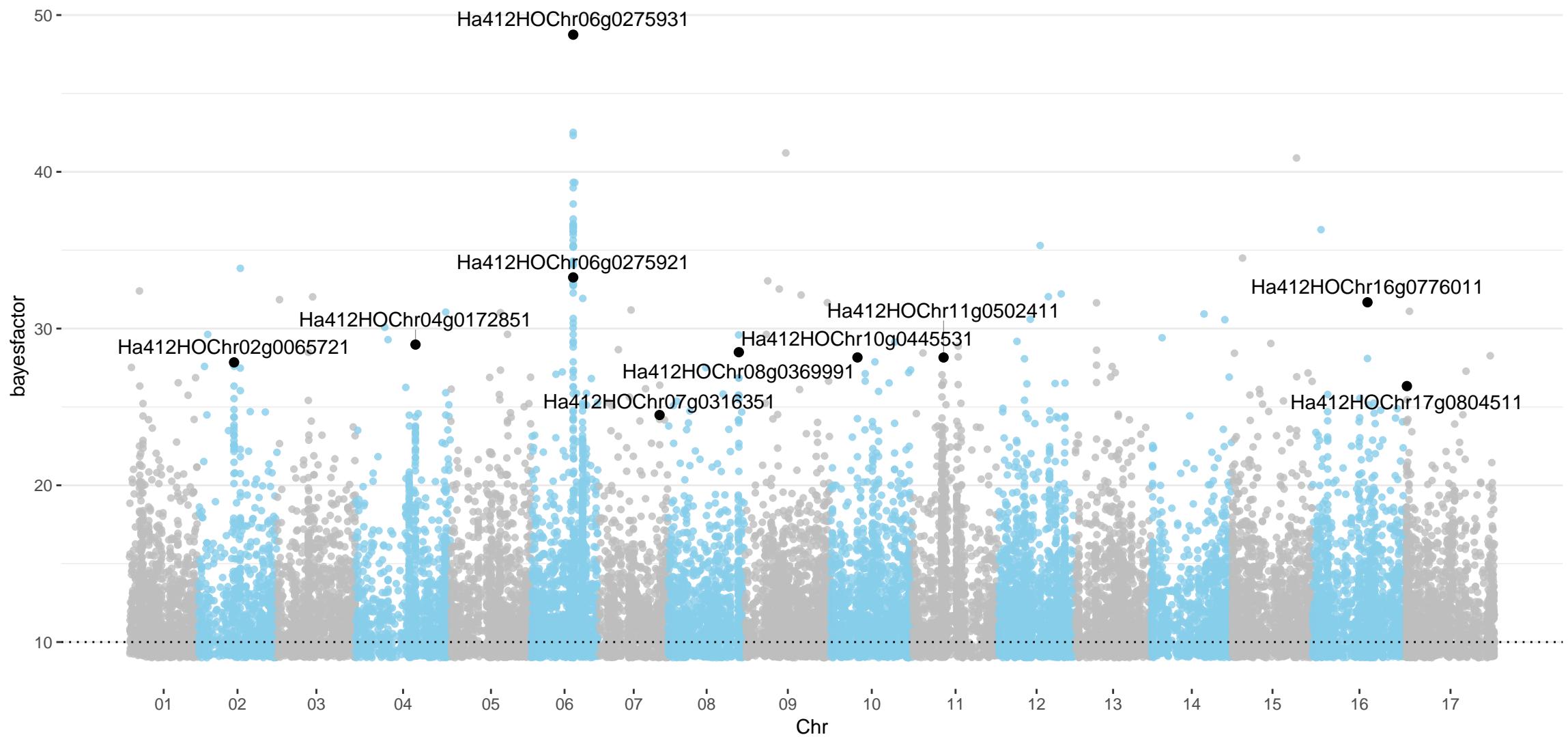
10

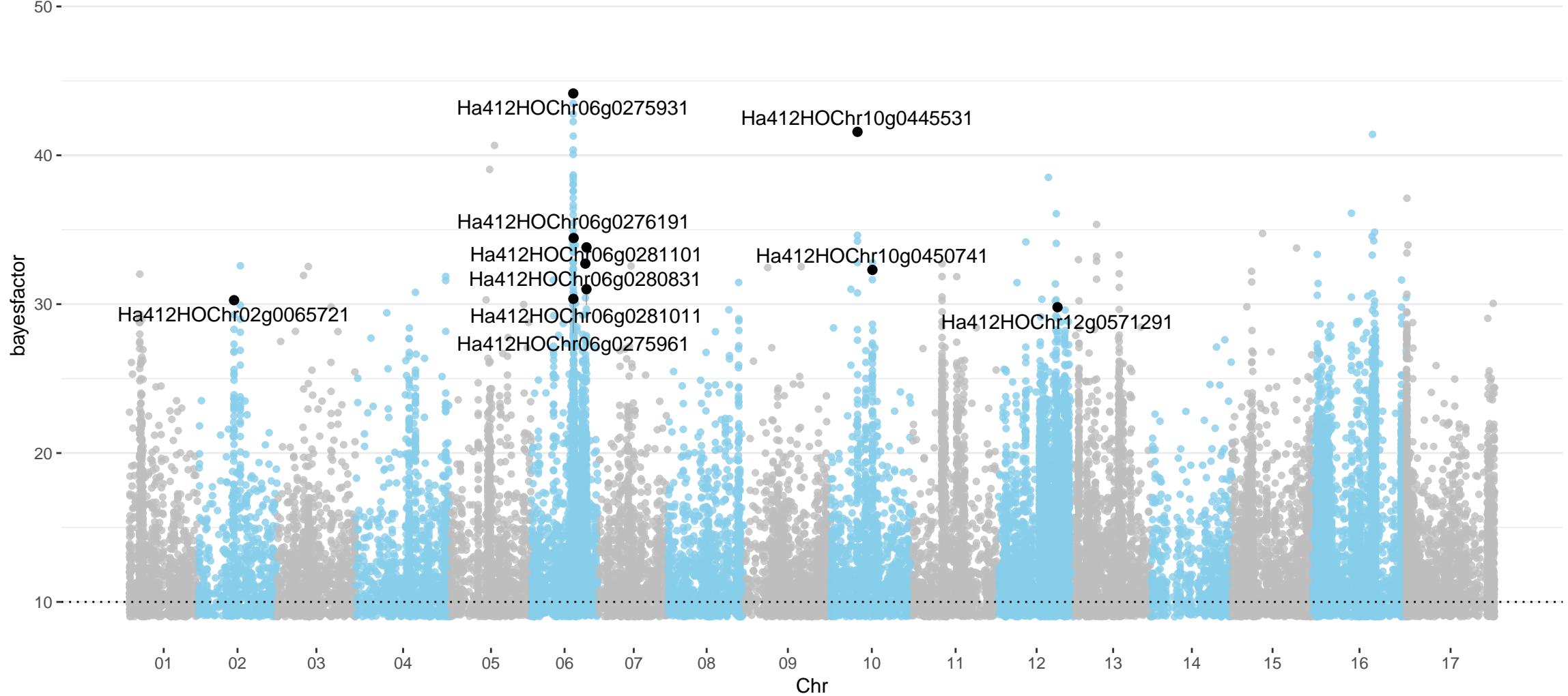
01 02 03 04 05 06 07 08 09 10 11 12 13 14 15 16 17

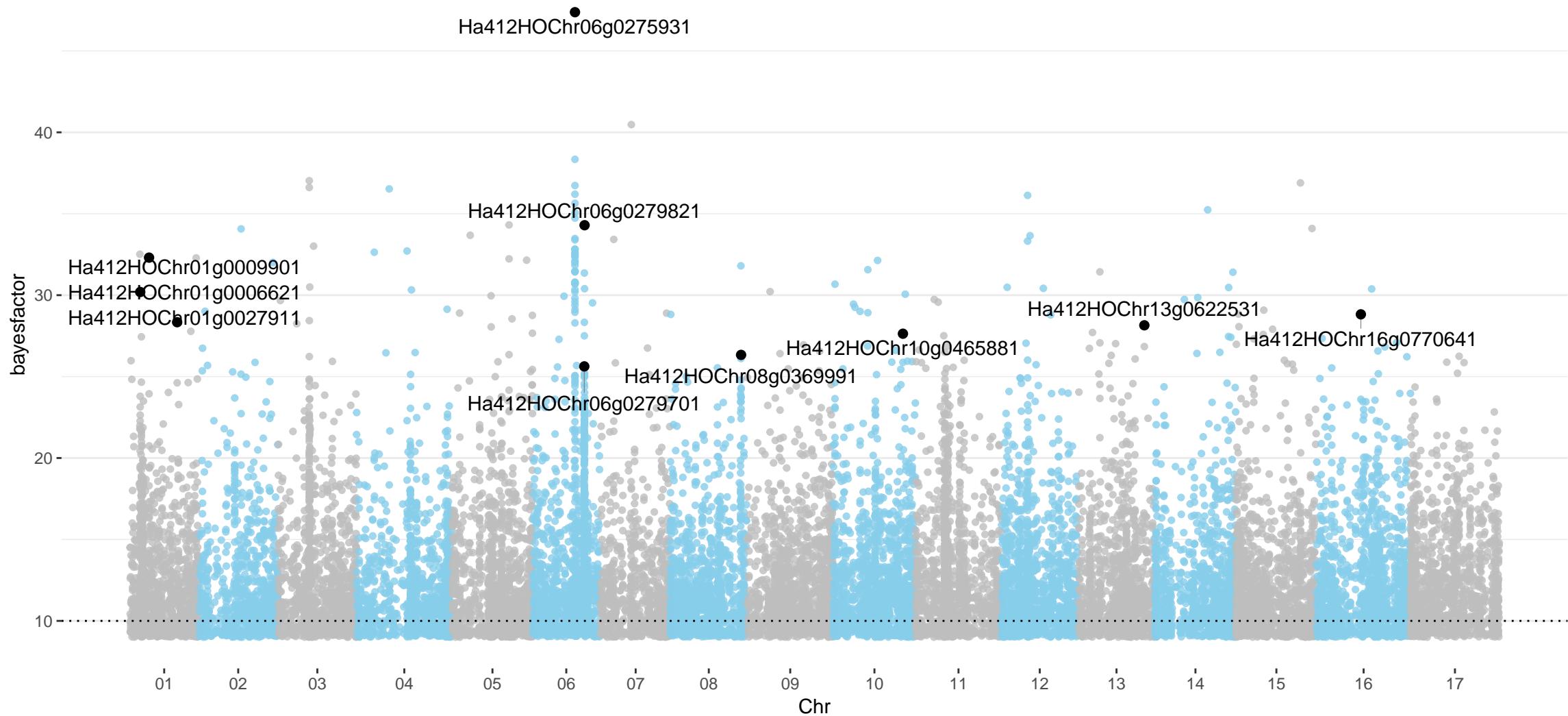
Chr

A genomic plot showing bayesfactor values across chromosomes 1 to 17. The y-axis ranges from 10 to 40. A vertical red line is at Chr 6. Black dots represent specific peaks labeled in the text. Blue dots represent other data points. Grey dots form the background. Horizontal grey lines are at y=10, 20, 30, 40.

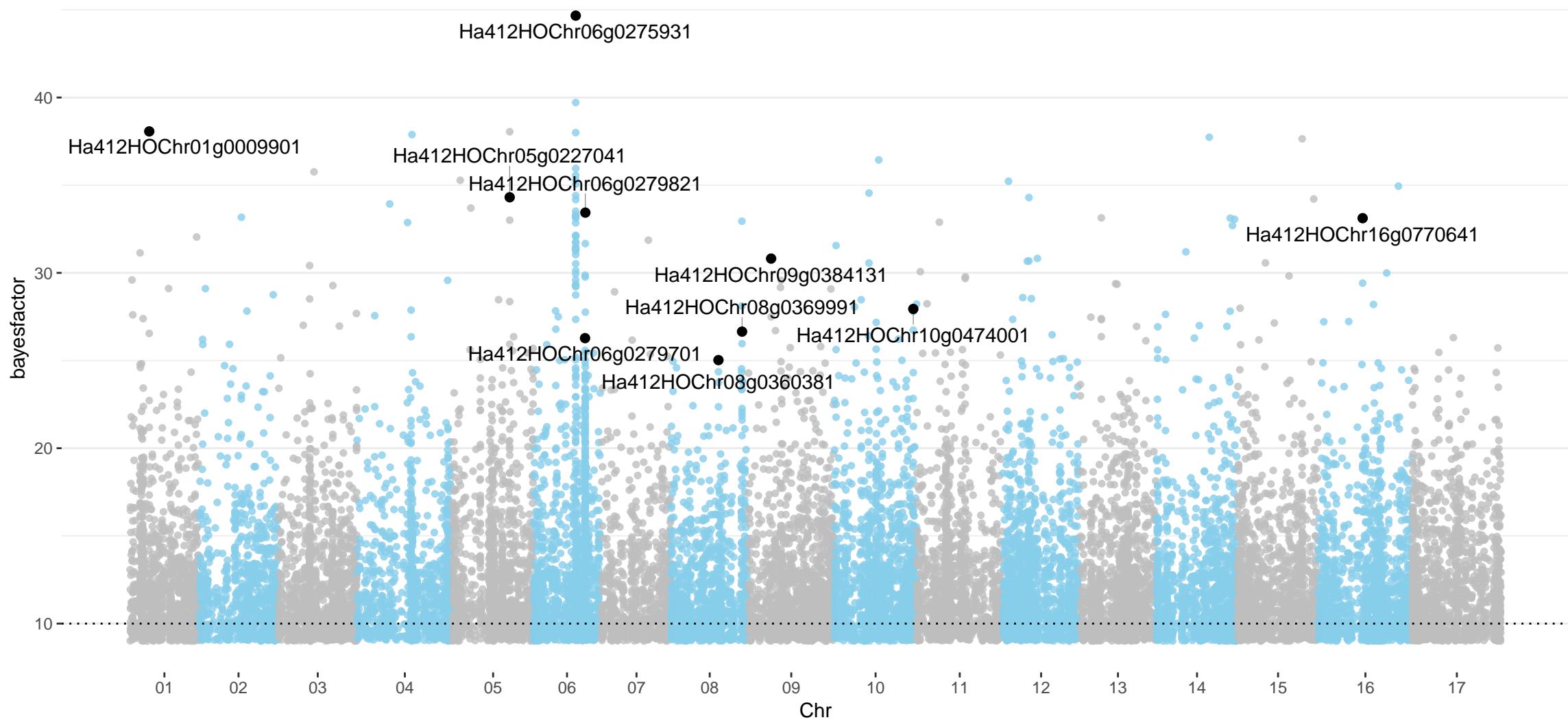
DD\_18\_e







bFFP\_e



eFFP\_e

Ha412HOChr06g0275931

Ha412HOChr07g0293701

Ha412HOChr06g0279821

Ha412HOChr06g0279751  
Ha412HOChr07g0293711  
Ha412HOChr06g0279701

Ha412HOChr11g0501111

Ha412HOChr11g0502411

Ha412HOChr10g0465881

Ha412HOChr13g0622531

bayesfactor

40

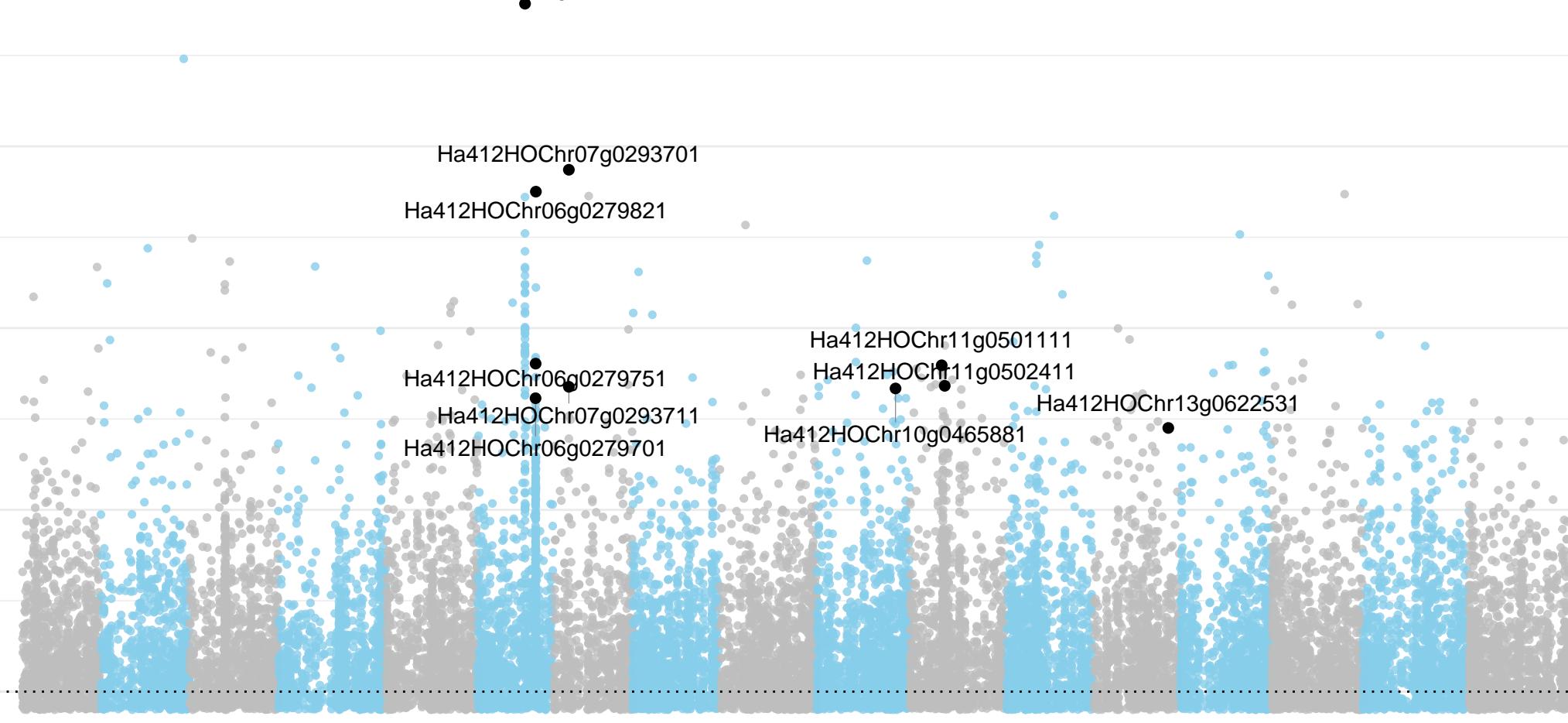
30

20

10

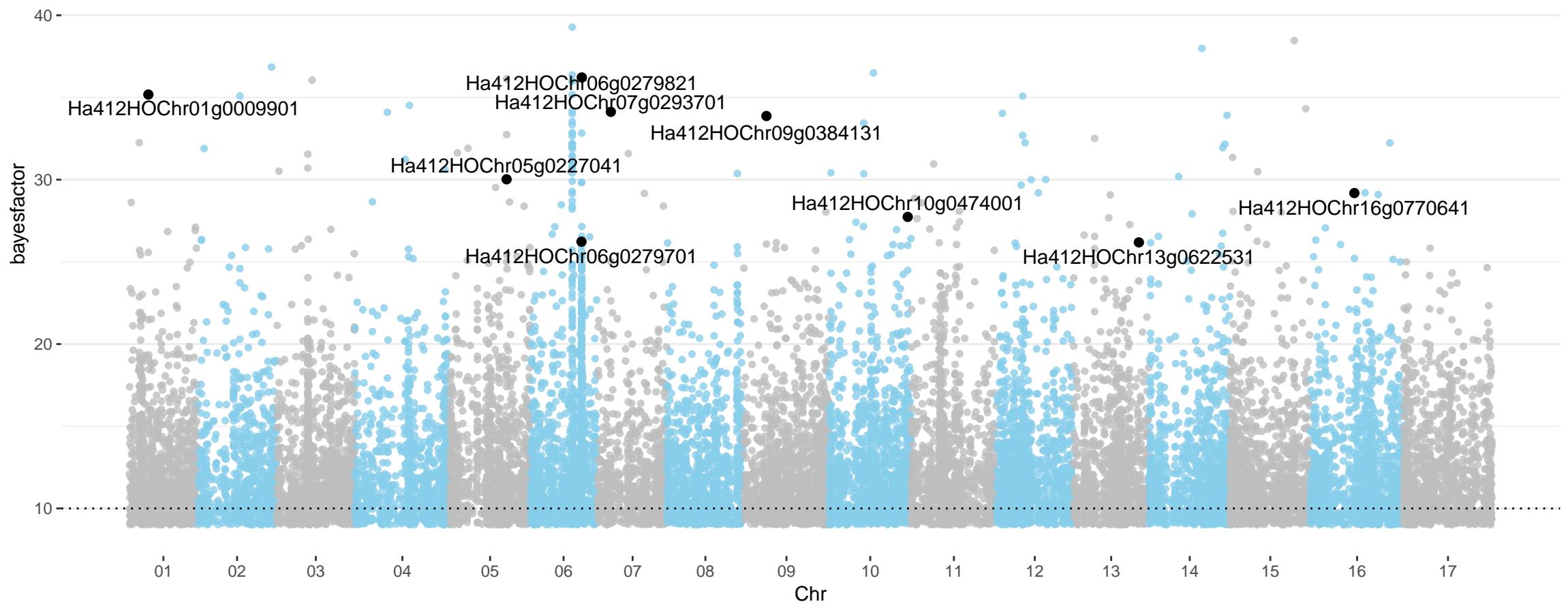
01 02 03 04 05 06 07 08 09 10 11 12 13 14 15 16 17

Chr



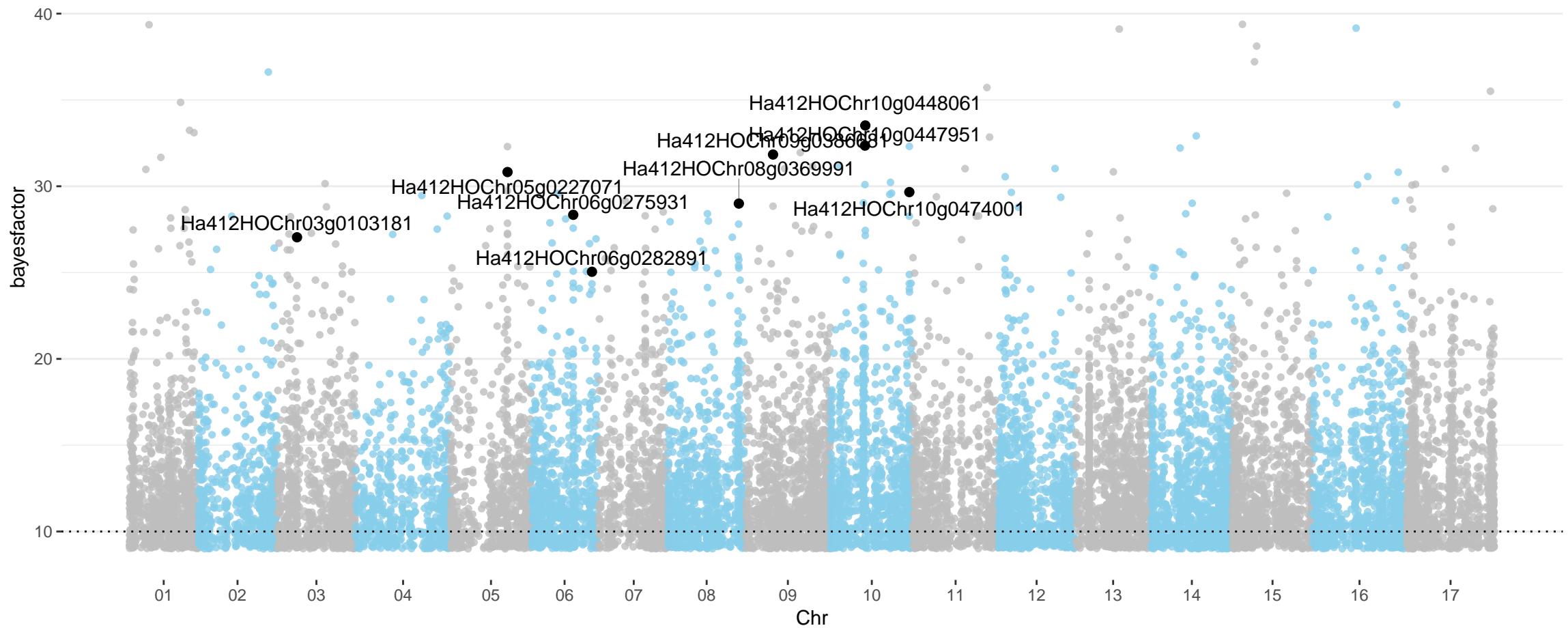
FFP\_e

Ha412HOChr06g0275931



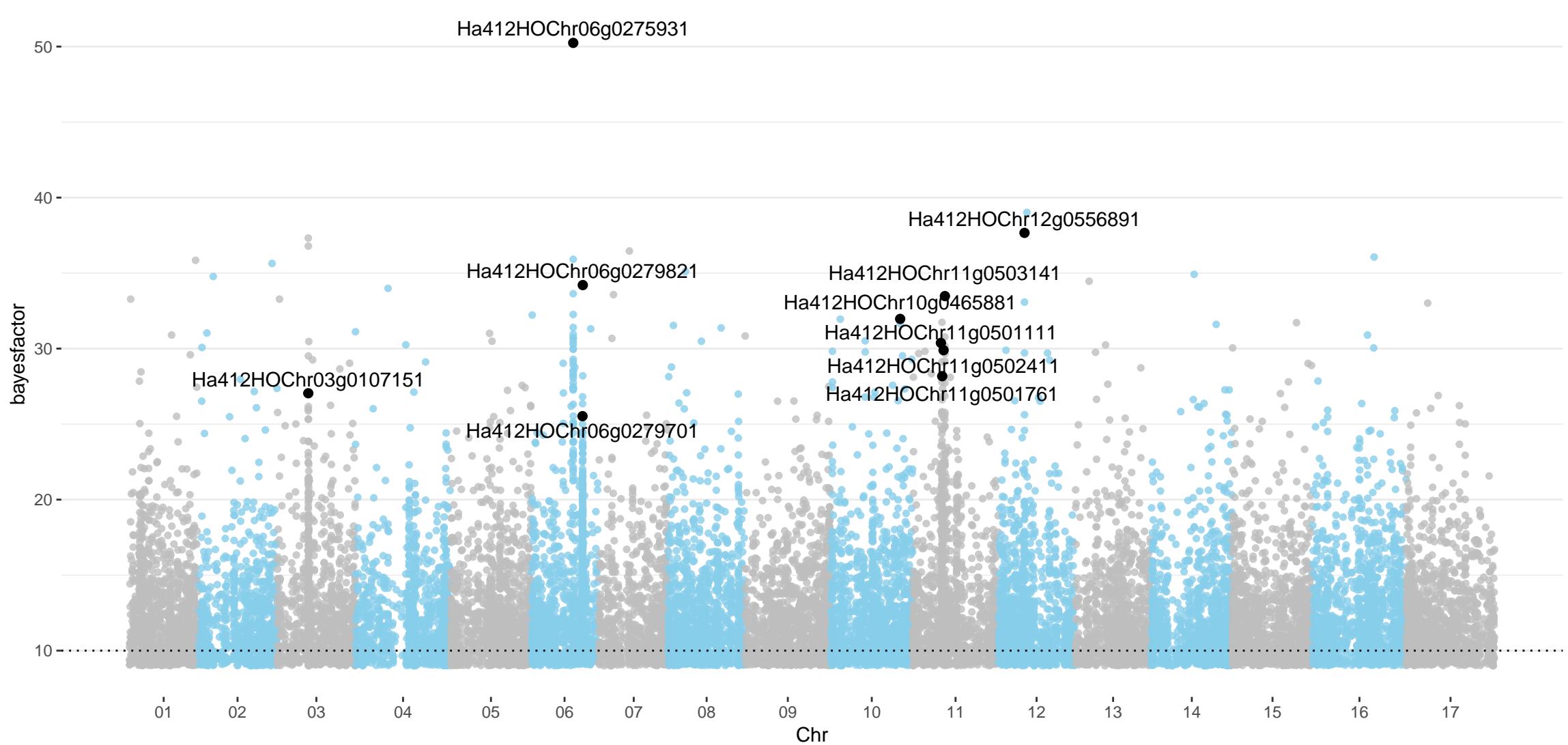
PAS\_e

Ha412HOChr05g0227041

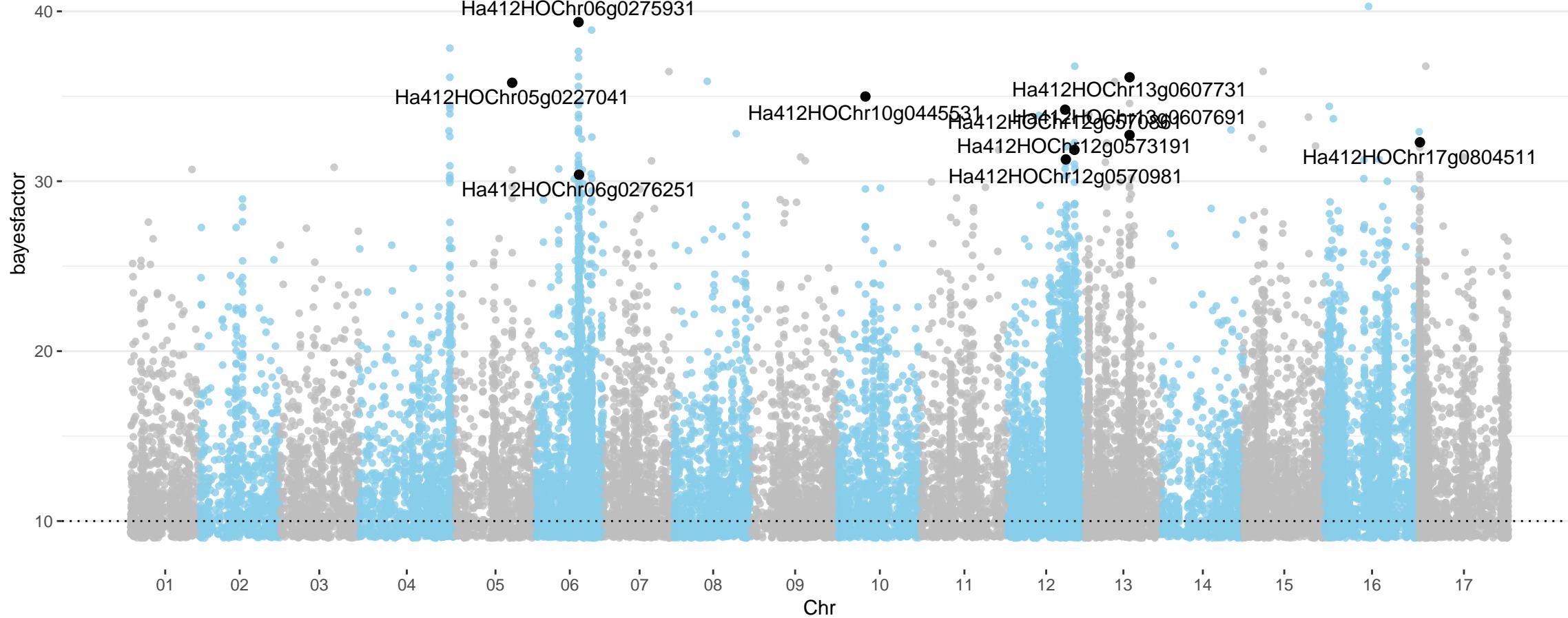


EMT\_e

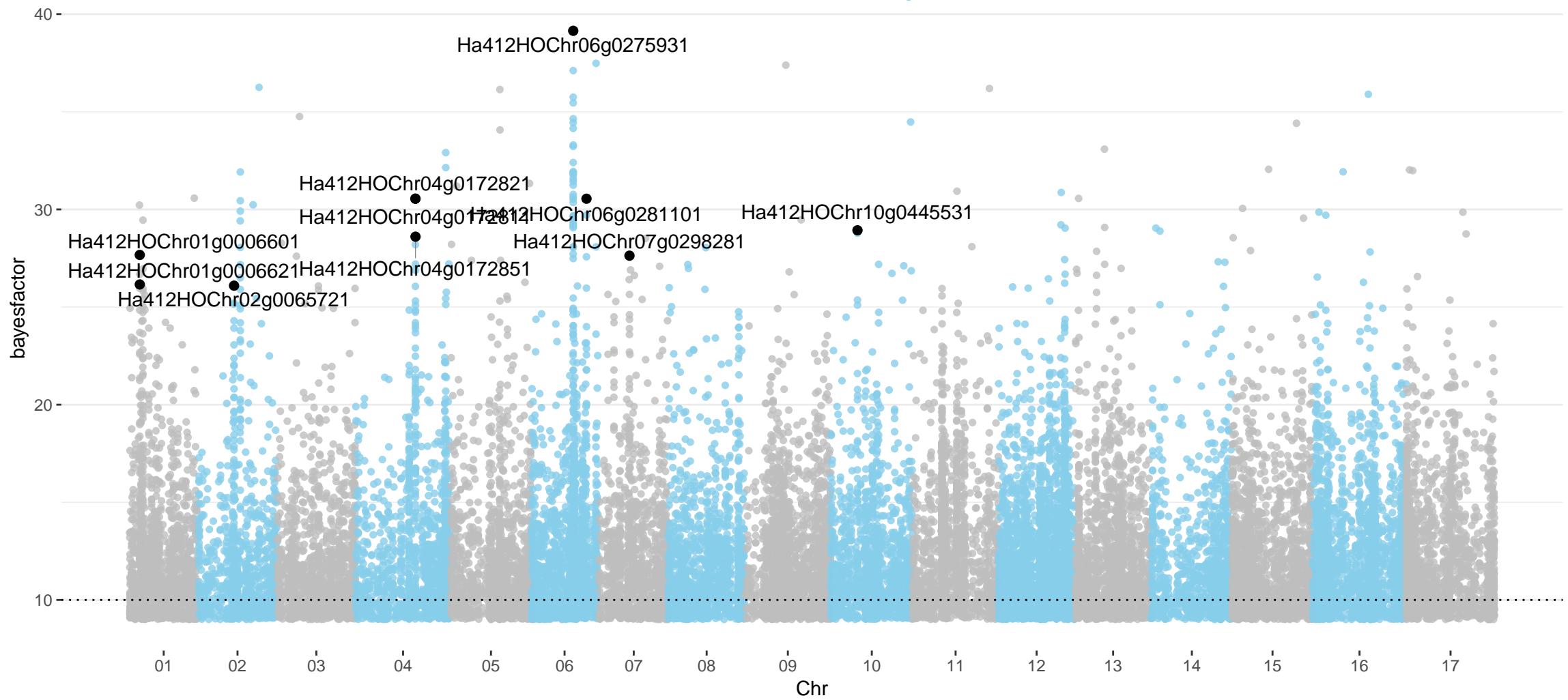
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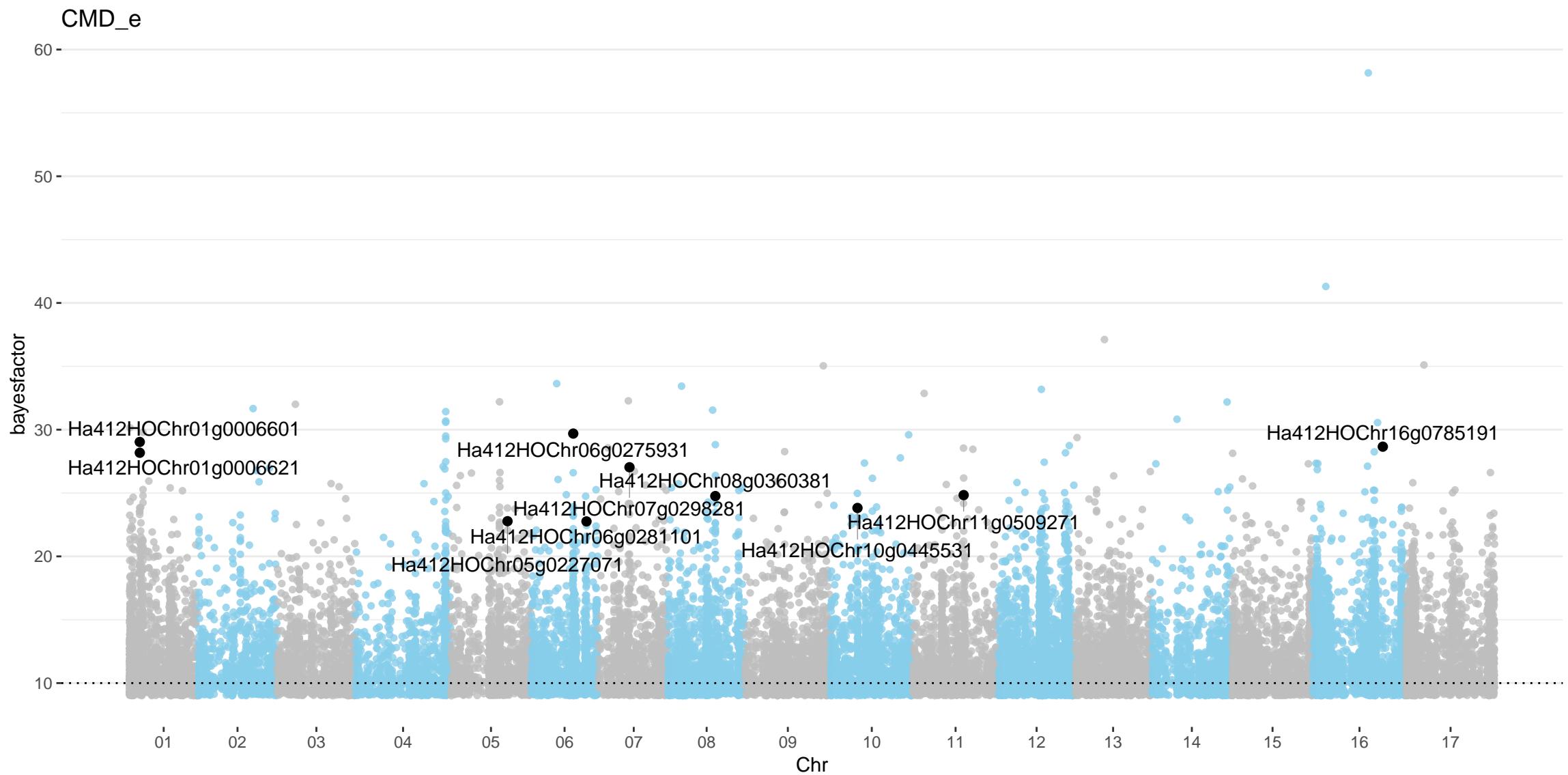


EXT\_e

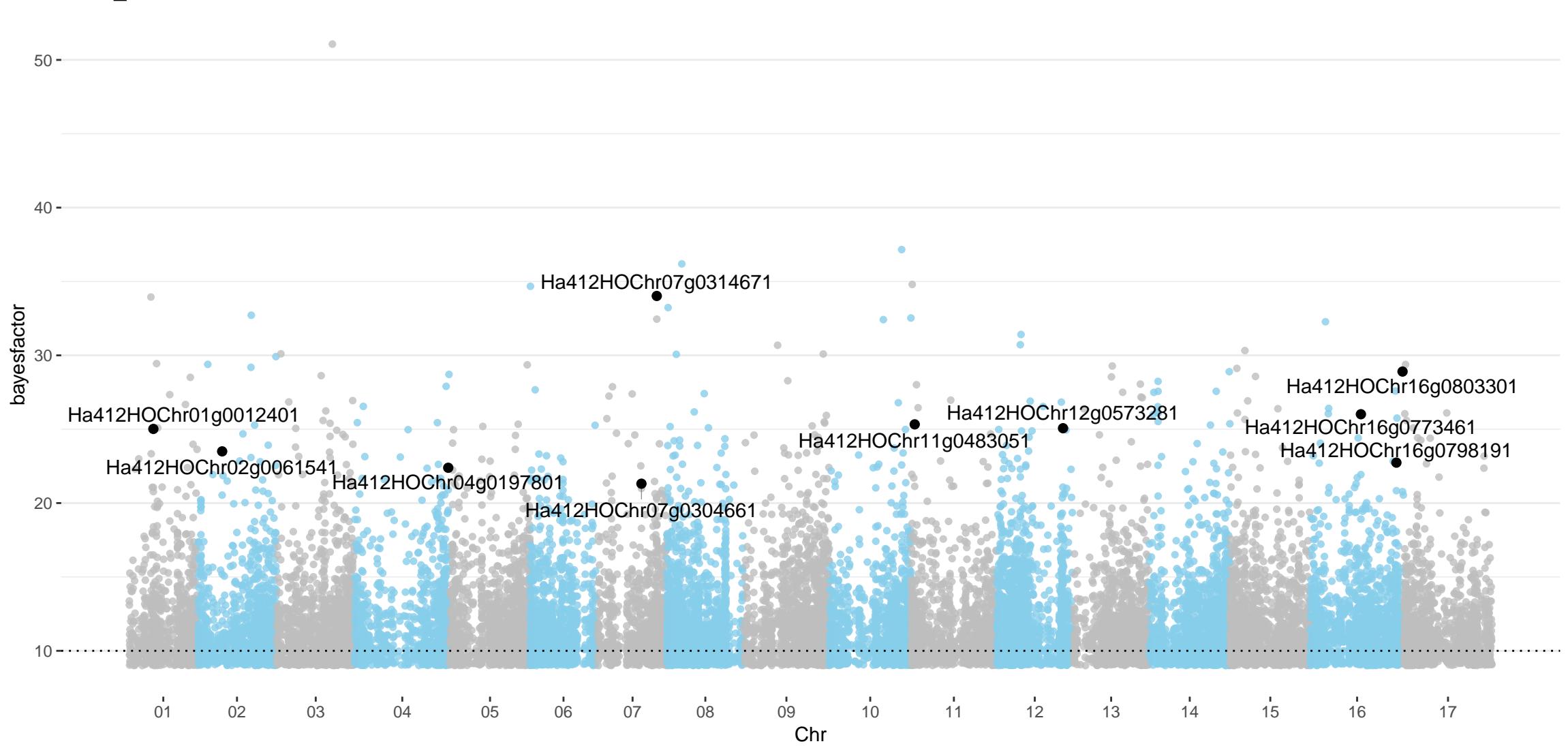


Eref\_e





MAR\_e



RH\_e

Ha412HOChr07g0301541

bayesfactor

40

30

20

10

01 02 03 04 05 06 07 08 09 10 11 12 13 14 15 16 17

Chr

Ha412HOChr07g0299581

Ha412HOChr07g0299571

Ha412HOChr06g0279821

Ha412HOChr05g0227731

Ha412HOChr09g0430281

Ha412HOChr10g0465881

Ha412HOChr13g0632591

Ha412HOChr16g0793421

Ha412HOChr13g0606911