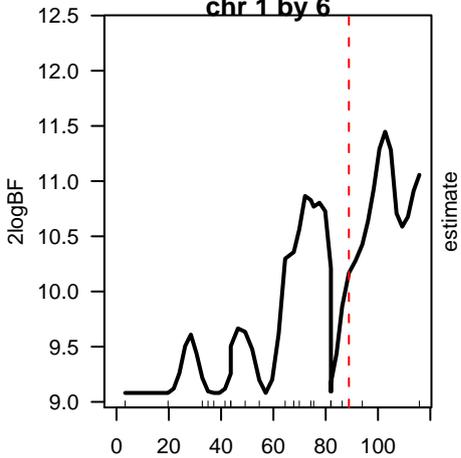
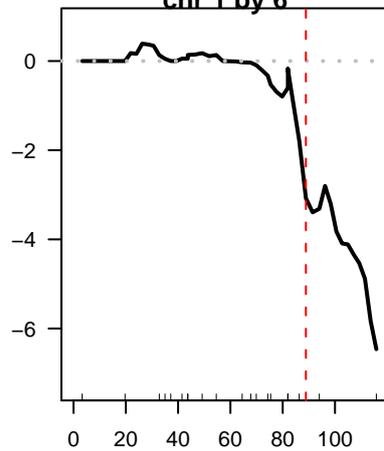
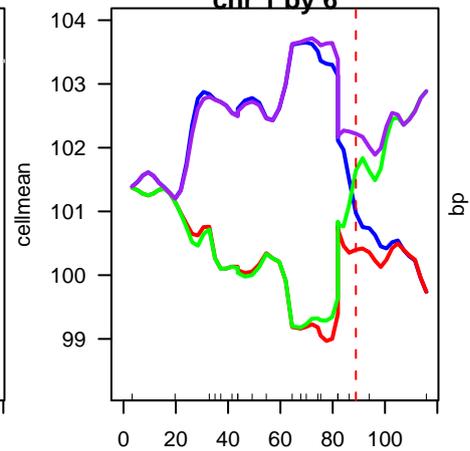


2logBF**chr 1 by 6**

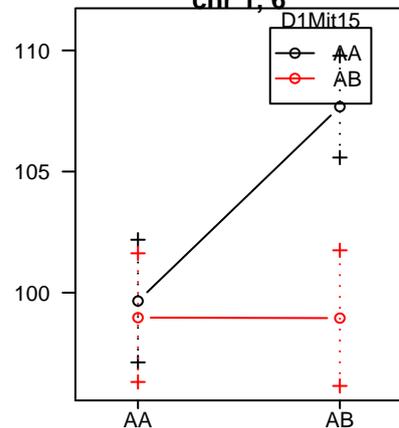
Map position (cM)
2logBF

estimate**chr 1 by 6**

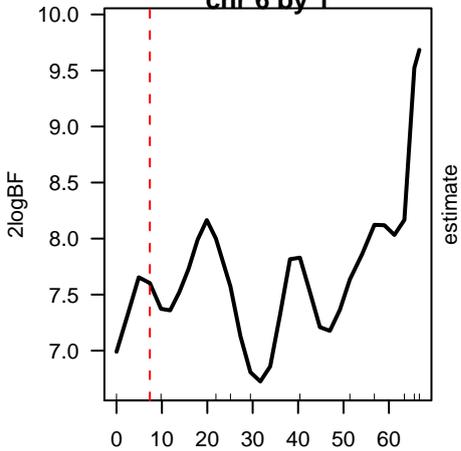
Map position (cM)
estimate

cellmean**chr 1 by 6**

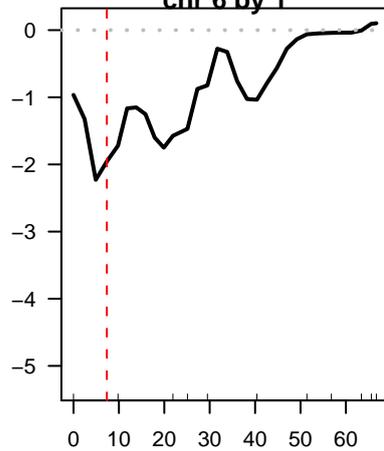
AA=blue, AH=purple, HA=green, HH=red
Map position (cM)
cellmean

interaction**chr 1, 6**

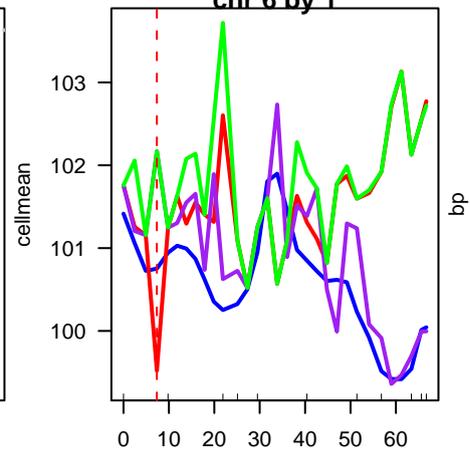
D1Mit15
AA AB
bp
interaction

chr 6 by 1

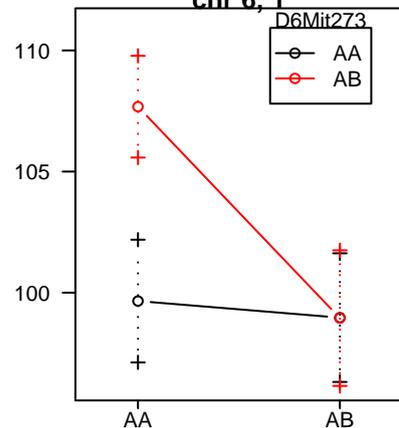
Map position (cM)
2logBF

chr 6 by 1

Map position (cM)
estimate

chr 6 by 1

AA=blue, HA=green, AH=purple, HH=red
Map position (cM)
cellmean

chr 6, 1

D6Mit273
AA AB
bp
interaction

D1Mit15