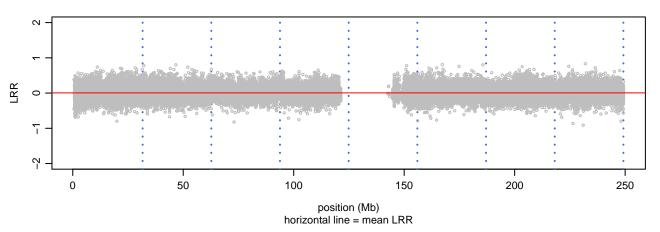
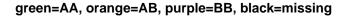
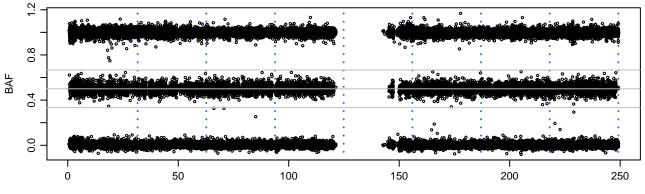
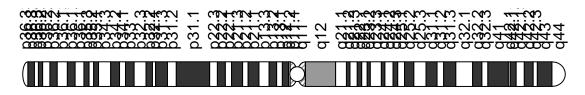
Scan 280 - Chromosome 1 - CLN5_W26X_G04



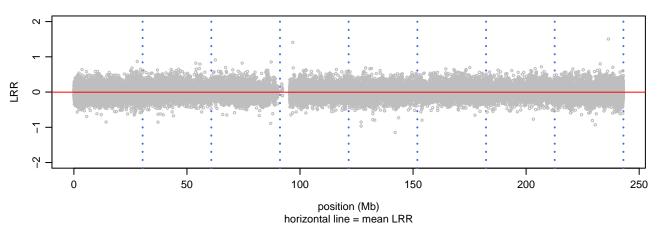




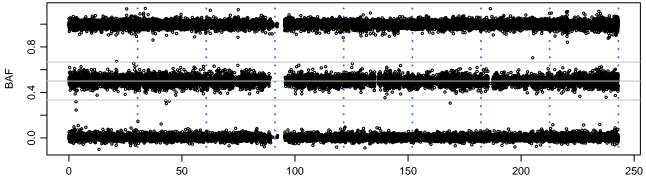
position (Mb) horizontal line = 0.5000 0.3333 0.6667



Scan 280 – Chromosome 2 – CLN5_W26X_G04

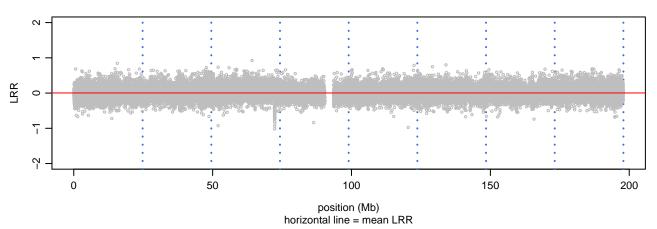


green=AA, orange=AB, purple=BB, black=missing

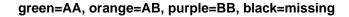


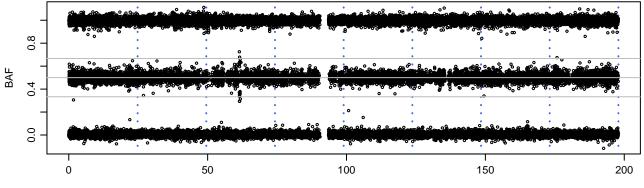
position (Mb) horizontal line = 0.5000 0.3333 0.6667



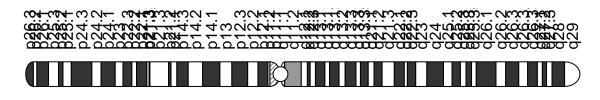


Scan 280 - Chromosome 3 - CLN5_W26X_G04

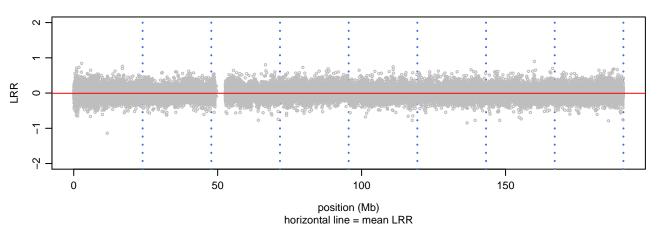


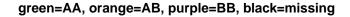


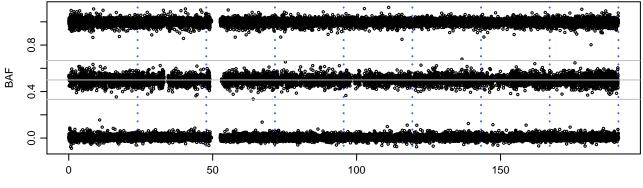
position (Mb) horizontal line = 0.5000 0.3333 0.6667



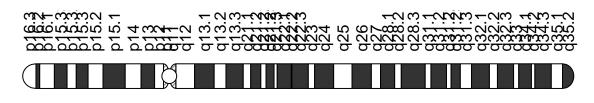
Scan 280 - Chromosome 4 - CLN5_W26X_G04



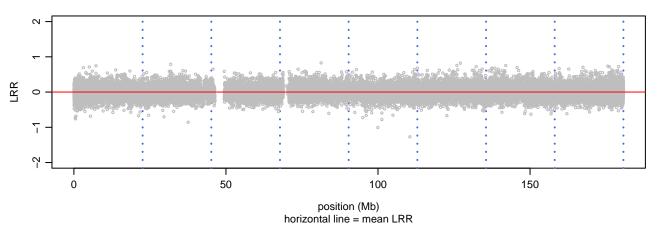




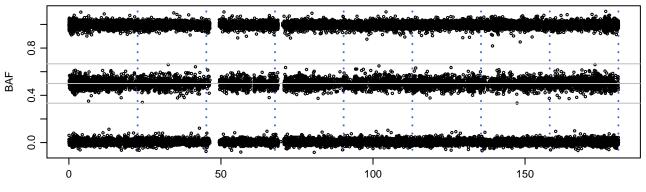
position (Mb) horizontal line = 0.5000 0.3333 0.6667



Scan 280 - Chromosome 5 - CLN5_W26X_G04



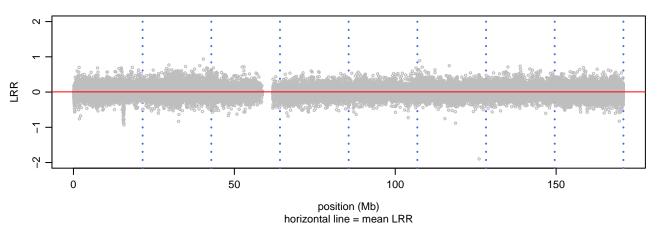
green=AA, orange=AB, purple=BB, black=missing



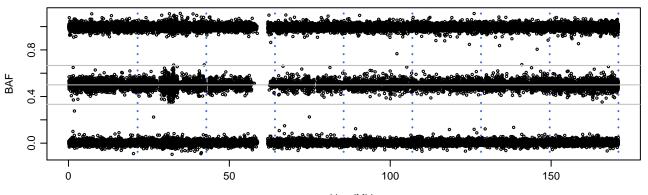
position (Mb) horizontal line = 0.5000 0.3333 0.6667



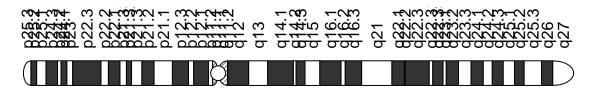
Scan 280 - Chromosome 6 - CLN5_W26X_G04



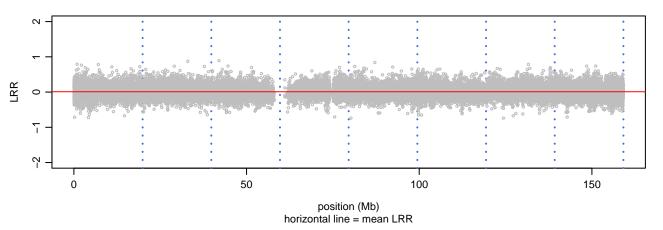
green=AA, orange=AB, purple=BB, black=missing



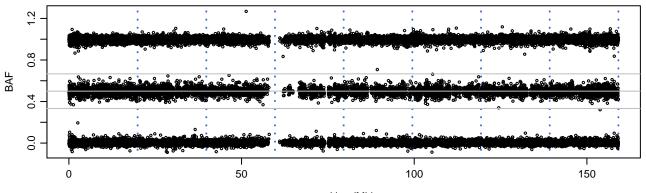
position (Mb) horizontal line = 0.5000 0.3333 0.6667



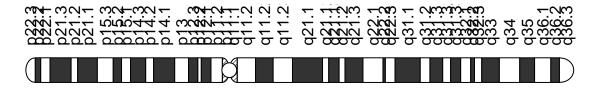
Scan 280 - Chromosome 7 - CLN5_W26X_G04



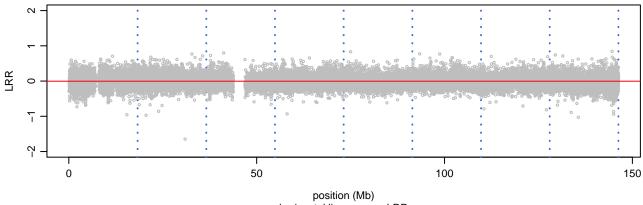
green=AA, orange=AB, purple=BB, black=missing



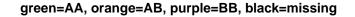
position (Mb) horizontal line = 0.5000 0.3333 0.6667

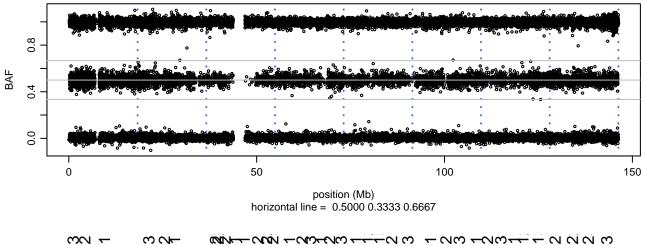


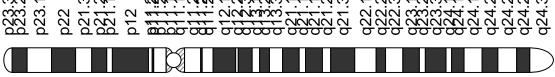
Scan 280 - Chromosome 8 - CLN5_W26X_G04



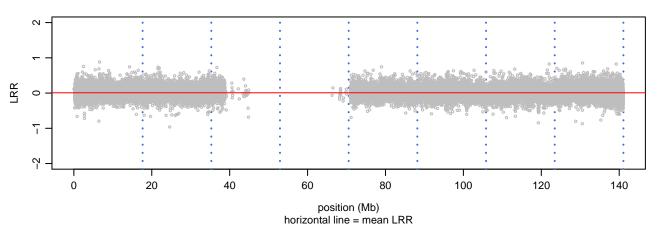
horizontal line = mean LRR



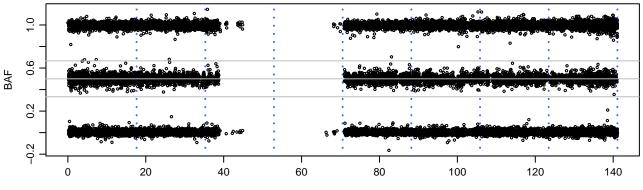




Scan 280 - Chromosome 9 - CLN5_W26X_G04



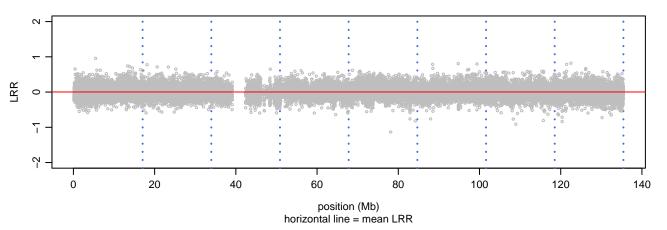
green=AA, orange=AB, purple=BB, black=missing



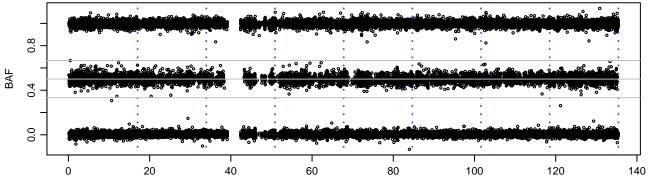
position (Mb) horizontal line = 0.5000 0.3333 0.6667



Scan 280 - Chromosome 10 - CLN5_W26X_G04

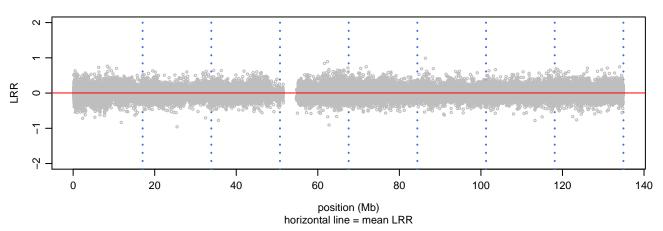


green=AA, orange=AB, purple=BB, black=missing

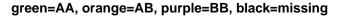


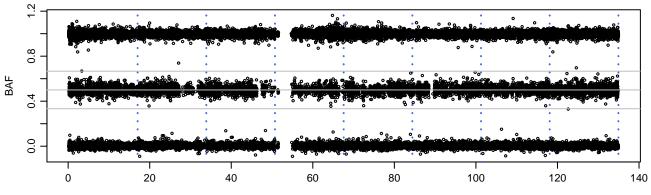
position (Mb) horizontal line = 0.5000 0.3333 0.6667



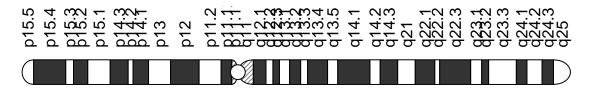


Scan 280 - Chromosome 11 - CLN5_W26X_G04

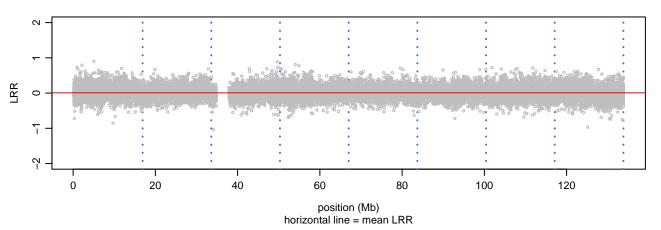




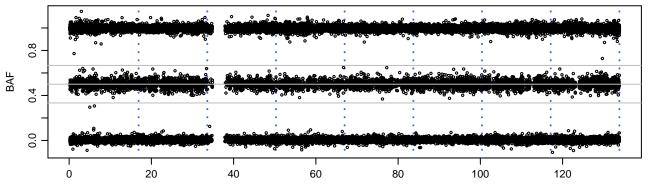
position (Mb) horizontal line = 0.5000 0.3333 0.6667



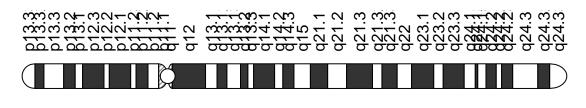
Scan 280 - Chromosome 12 - CLN5_W26X_G04

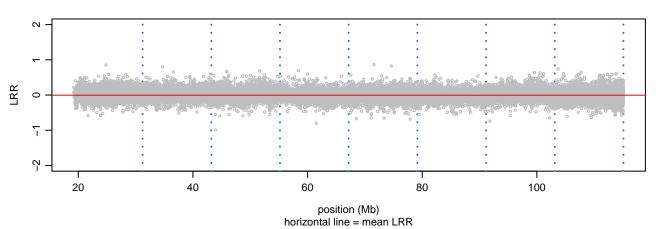


green=AA, orange=AB, purple=BB, black=missing

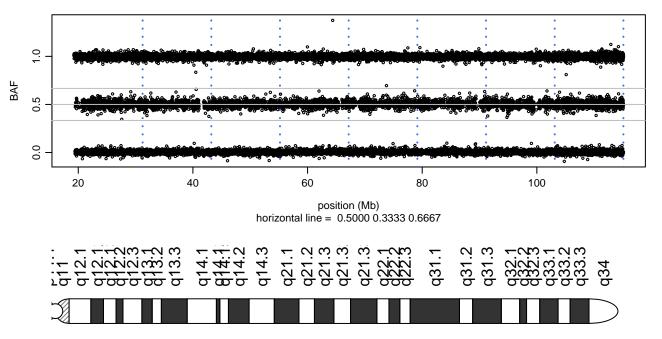


position (Mb) horizontal line = 0.5000 0.3333 0.6667

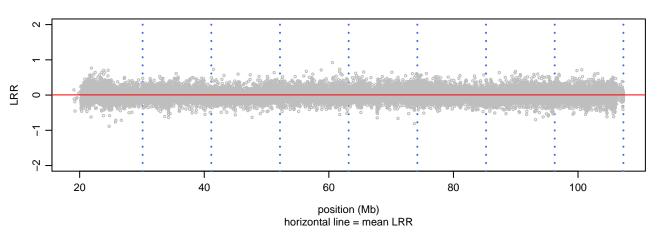




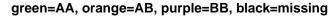


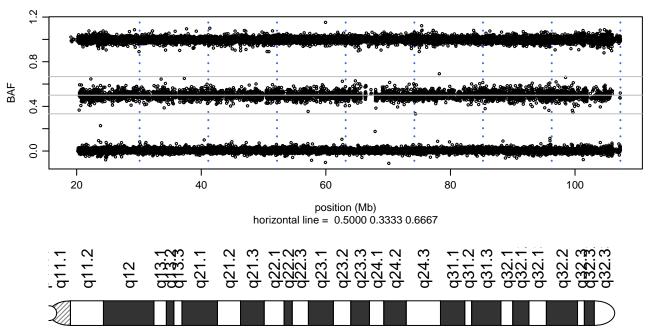


Scan 280 - Chromosome 13 - CLN5_W26X_G04

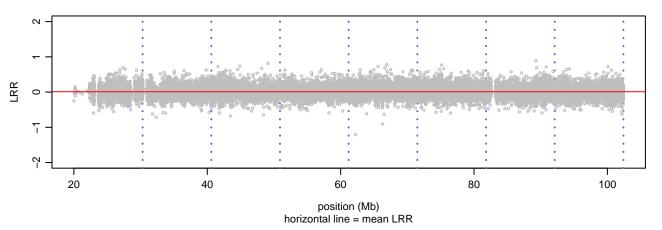




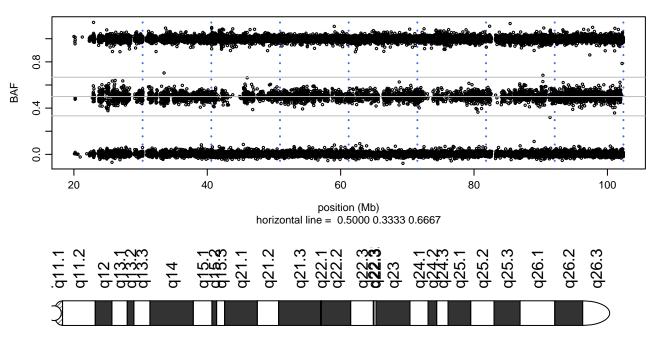




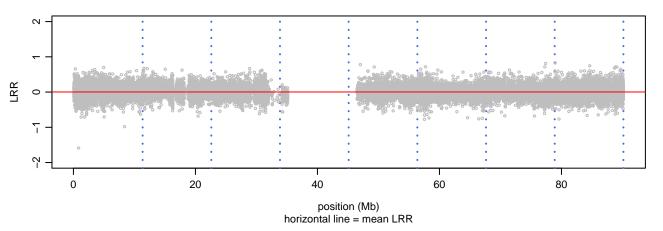
Scan 280 – Chromosome 15 – CLN5_W26X_G04



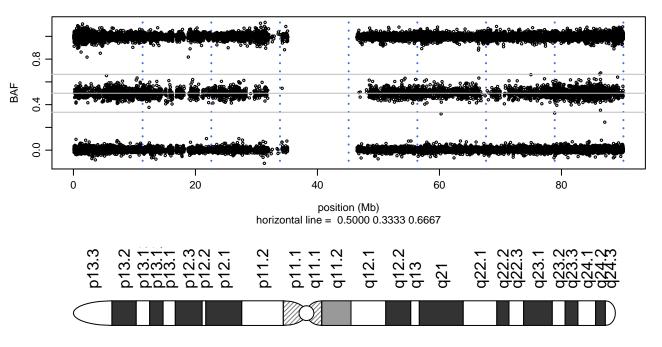
green=AA, orange=AB, purple=BB, black=missing

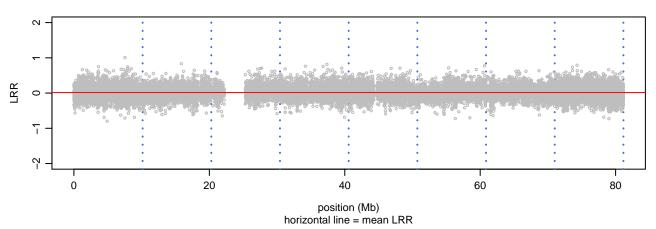


Scan 280 - Chromosome 16 - CLN5_W26X_G04

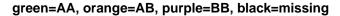


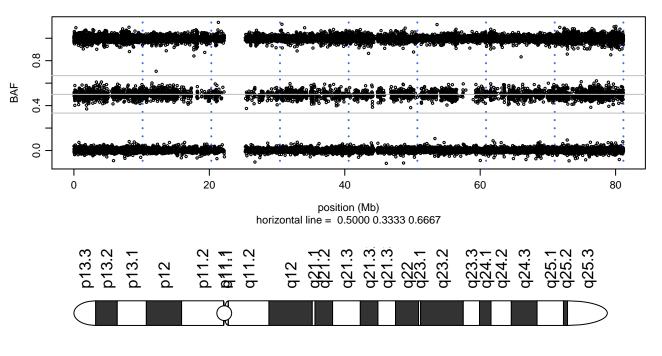
green=AA, orange=AB, purple=BB, black=missing

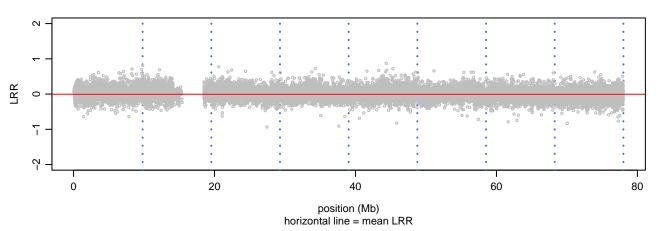




Scan 280 - Chromosome 17 - CLN5_W26X_G04

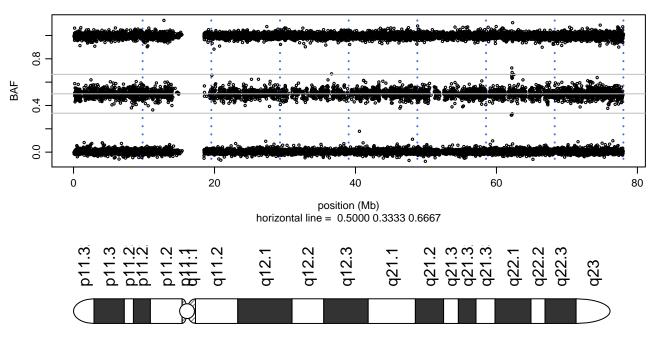




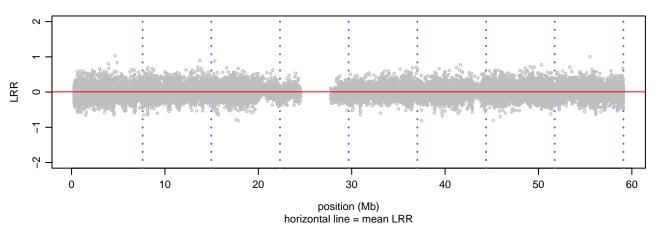


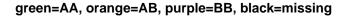
Scan 280 - Chromosome 18 - CLN5_W26X_G04

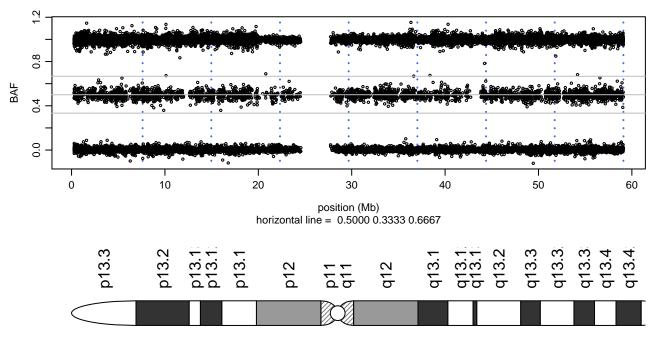
green=AA, orange=AB, purple=BB, black=missing



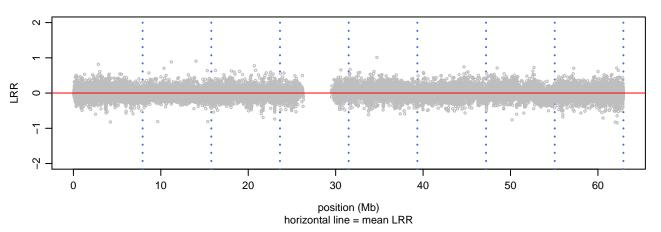
Scan 280 - Chromosome 19 - CLN5_W26X_G04

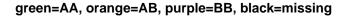


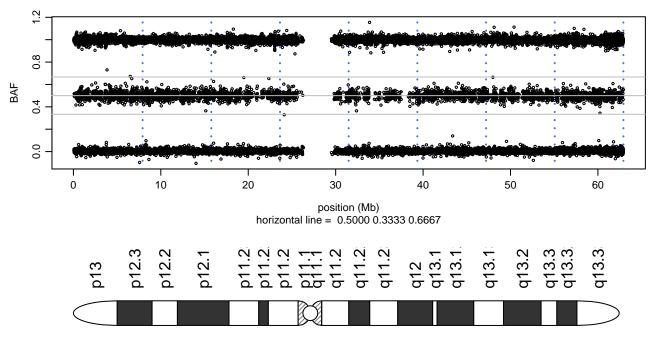




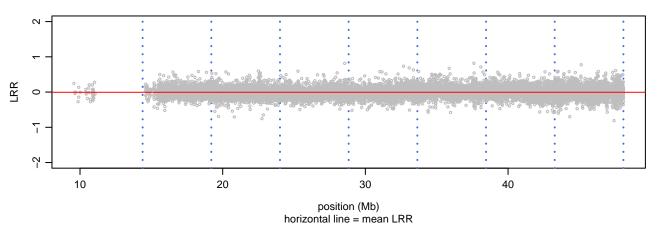




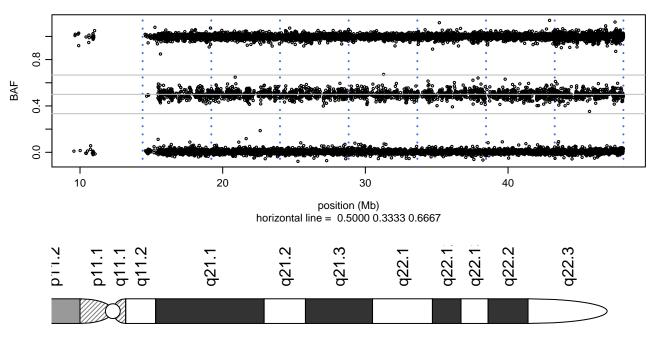




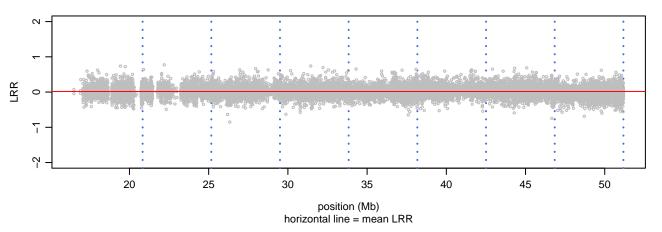




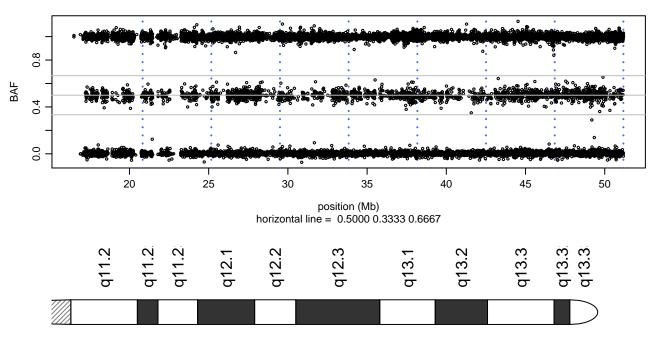
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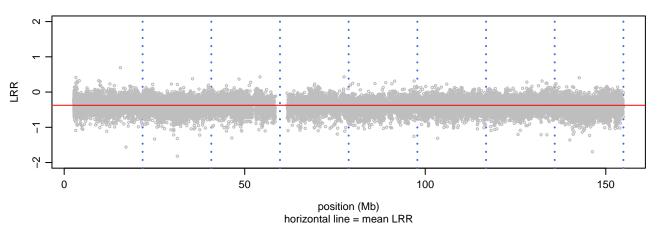


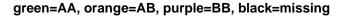


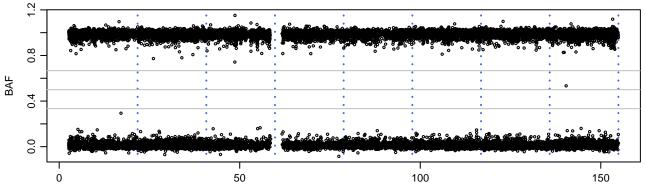
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Scan 280 – Chromosome X – CLN5_W26X_G04

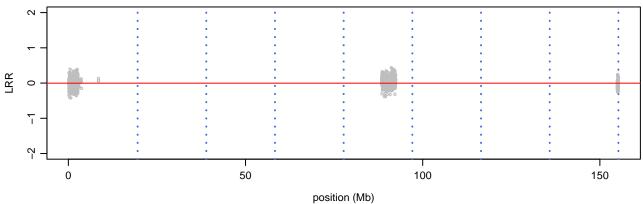






position (Mb) horizontal line = 0.5000 0.3333 0.6667

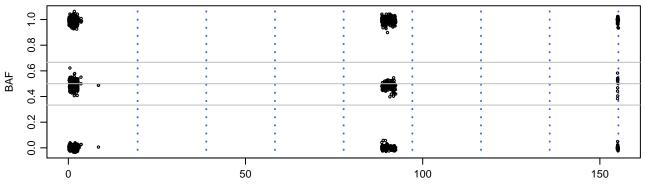




Scan 280 – Chromosome Y – CLN5_W26X_G04

horizontal line = mean LRR

green=AA, orange=AB, purple=BB, black=missing



position (Mb) horizontal line = 0.5000 0.3333 0.6667

