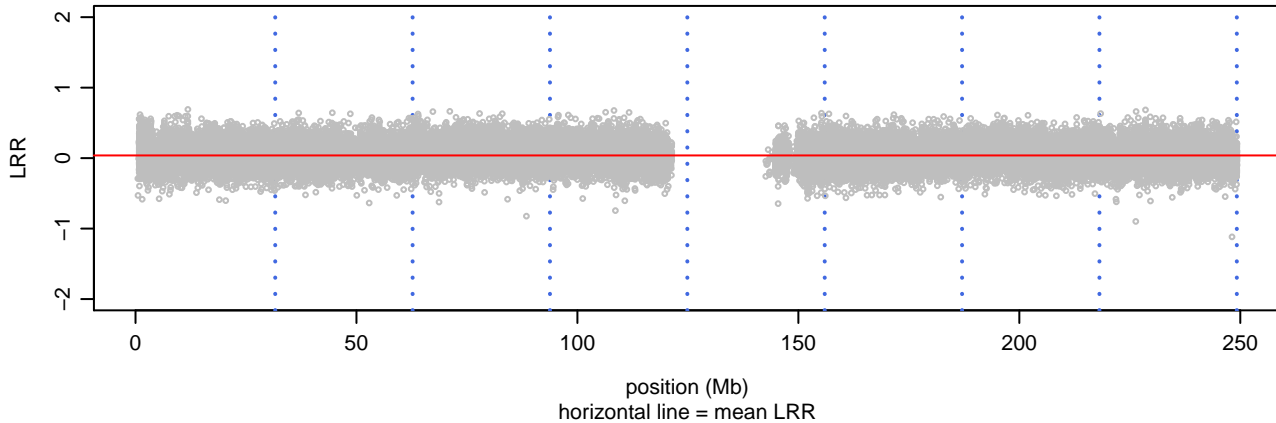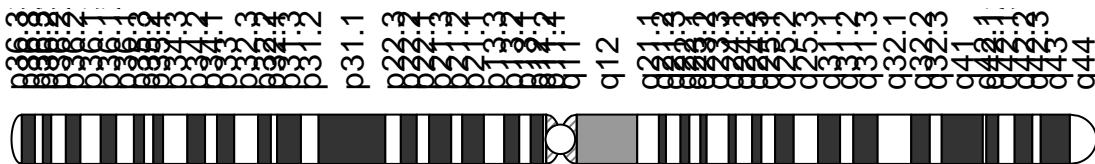# Scan 280 – Chromosome 1 – SNCB_P123H_D11

position (Mb)
horizontal line = mean LRR

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

position (Mb)
horizontal line = 0.5000 0.3333 0.6667
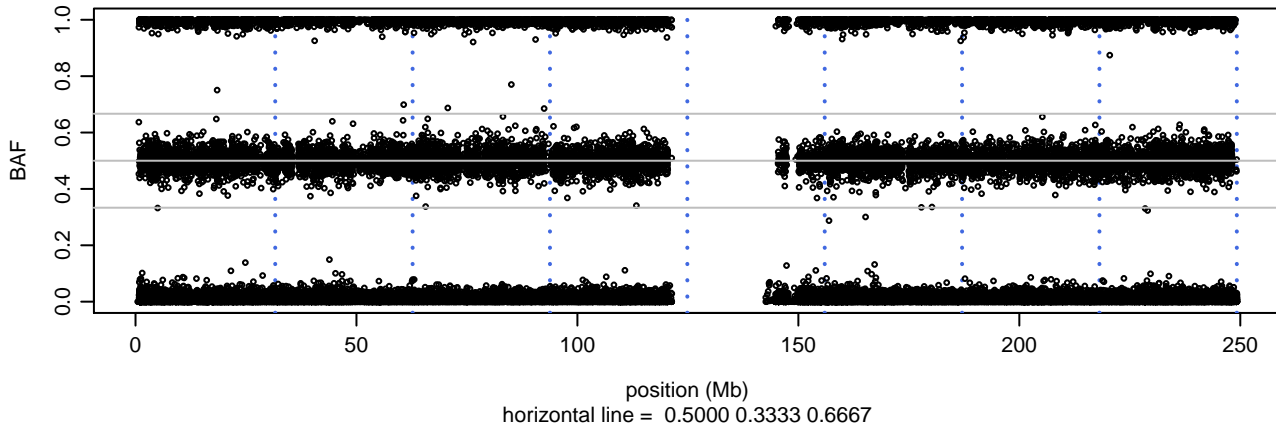
Scan 280 – Chromosome 2 – SNCB_P123H_D11

LRR

position (Mb)
horizontal line = mean LRR

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

BAF

position (Mb)
horizontal line = 0.5000 0.3333 0.6667
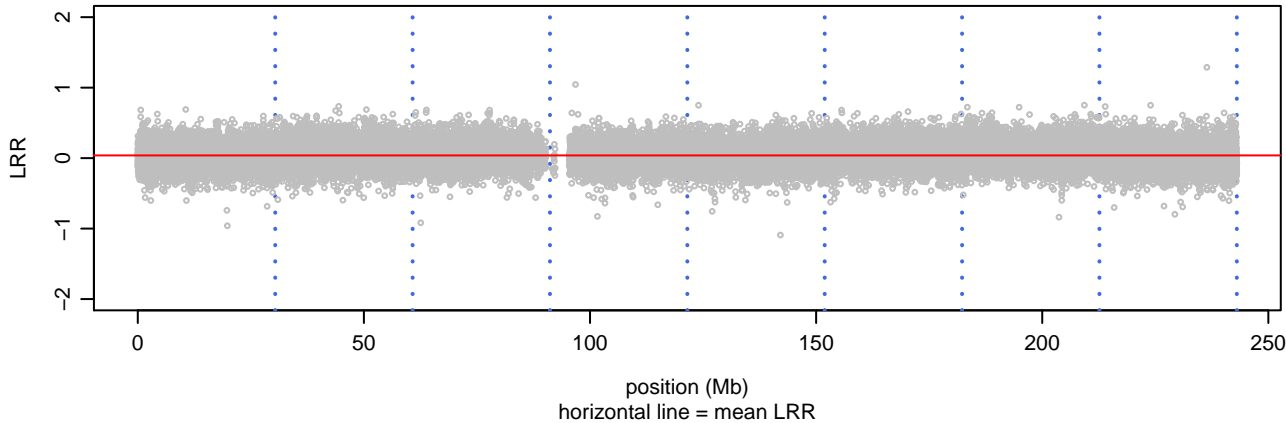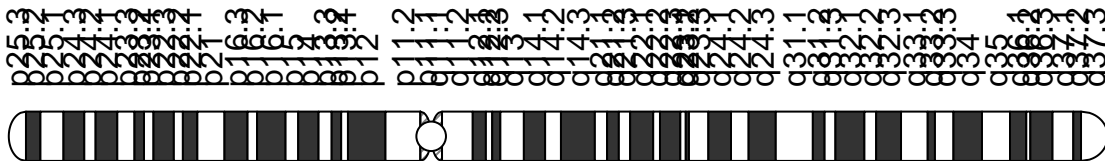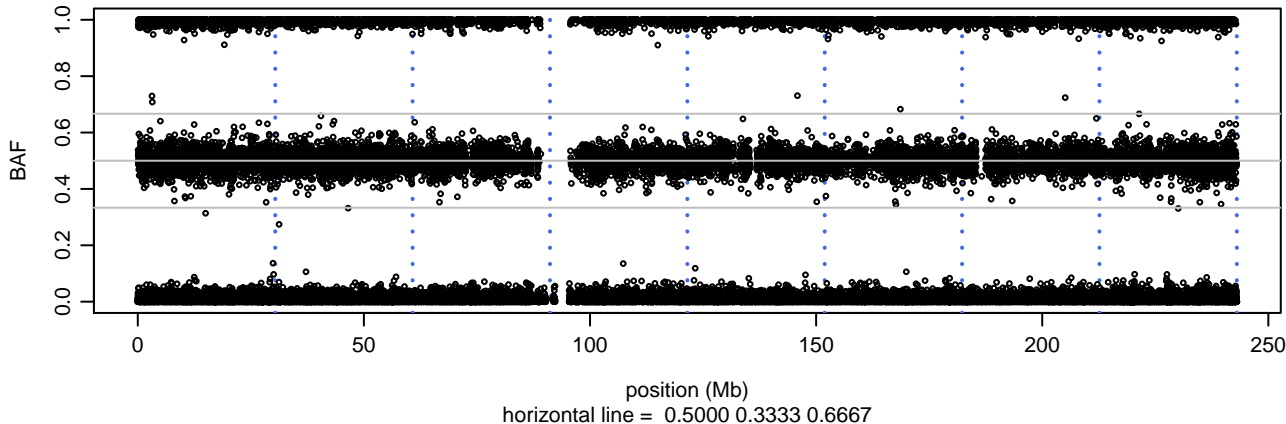
Scan 280 – Chromosome 3 – SNCB_P123H_D11

position (Mb)
horizontal line = mean LRR

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

position (Mb)
horizontal line = 0.5000 0.3333 0.6667
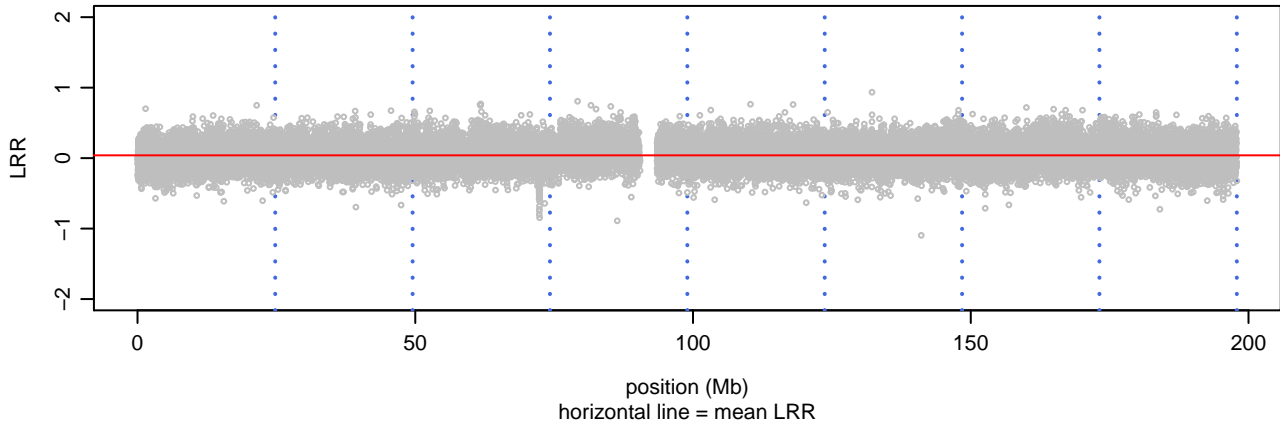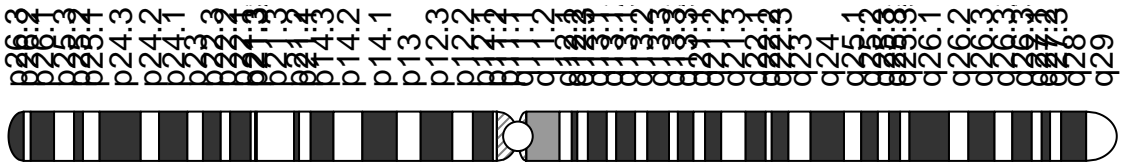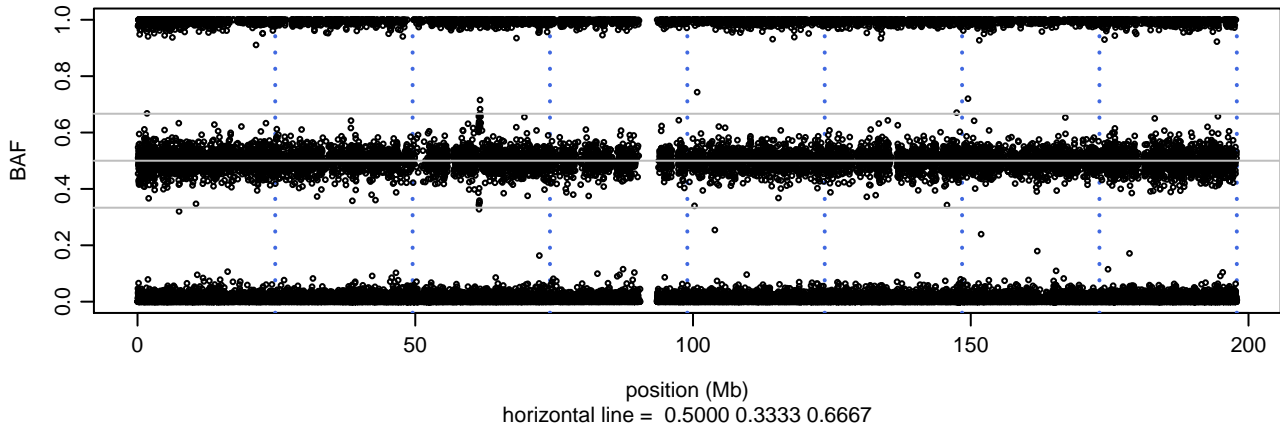
Scan 280 – Chromosome 4 – SNCB_P123H_D11

LRR

position (Mb)
horizontal line = mean LRR

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

BAF

position (Mb)
horizontal line = 0.5000 0.3333 0.6667
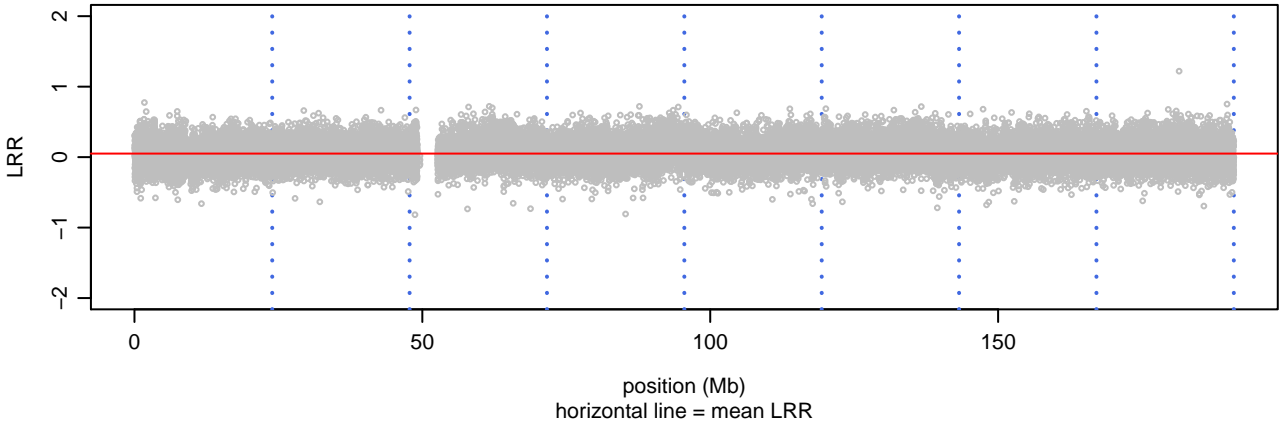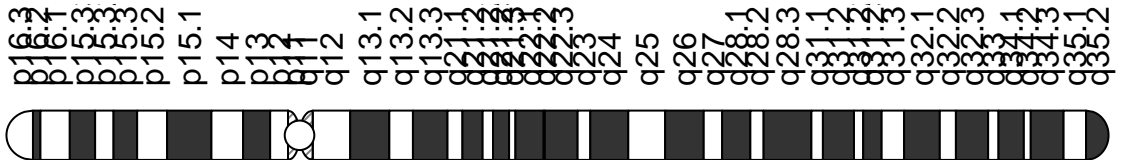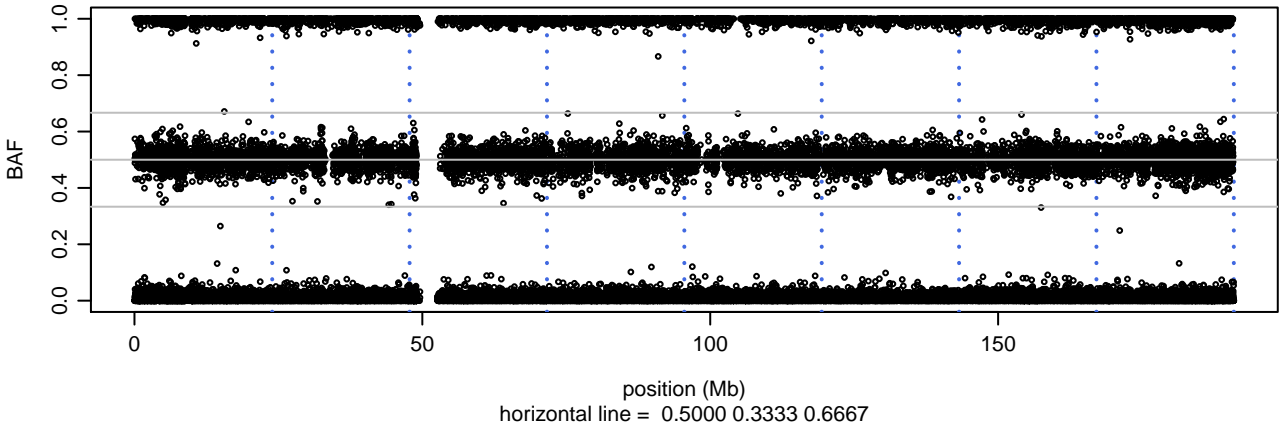
Scan 280 – Chromosome 5 – SNCB_P123H_D11

LRR

position (Mb)
horizontal line = mean LRR

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

BAF

position (Mb)
horizontal line = 0.5000 0.3333 0.6667
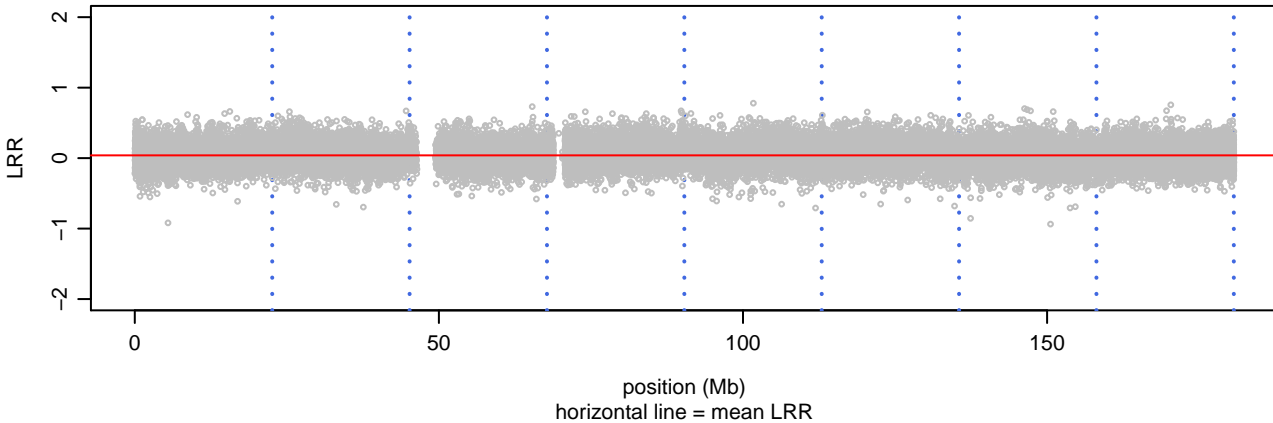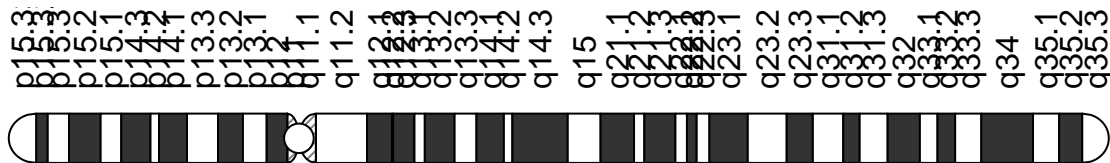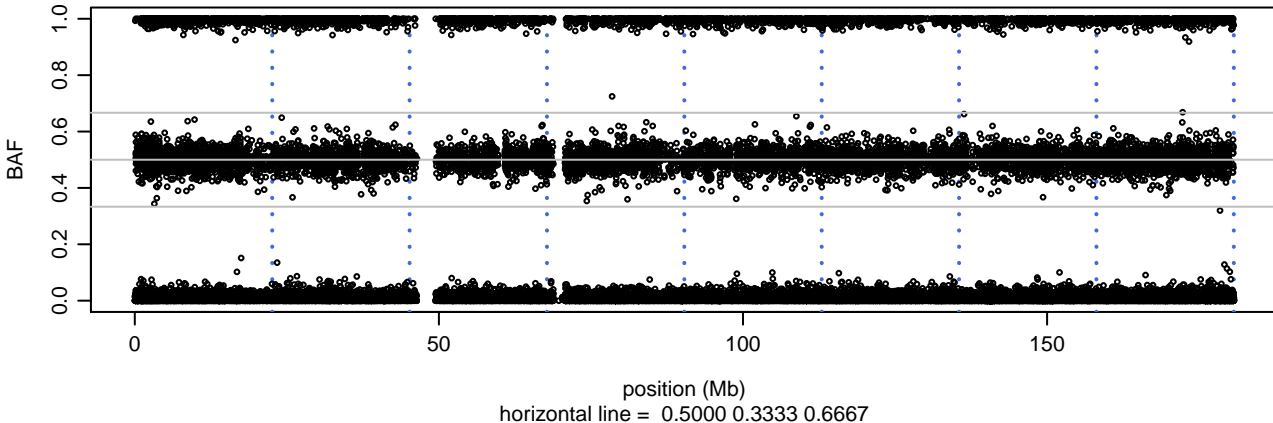
Scan 280 – Chromosome 6 – SNCB_P123H_D11

position (Mb)
horizontal line = mean LRR

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

position (Mb)
horizontal line = 0.5000 0.3333 0.6667

Scan 280 – Chromosome 7 – SNCB_P123H_D11

position (Mb)
horizontal line = mean LRR

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

position (Mb)
horizontal line = 0.5000 0.3333 0.6667
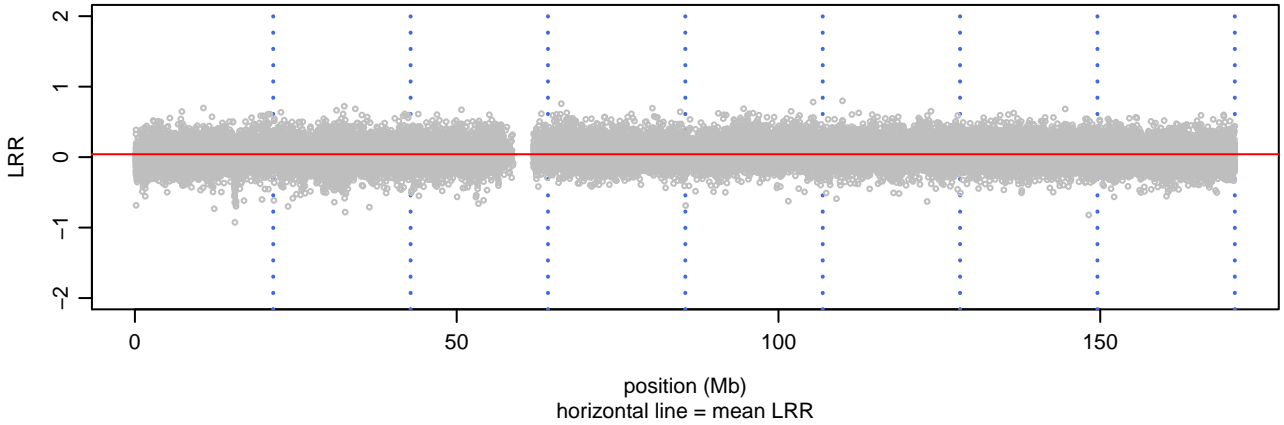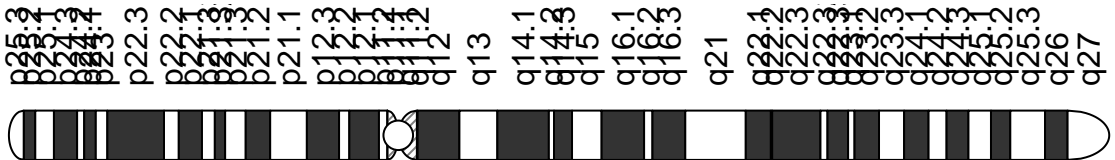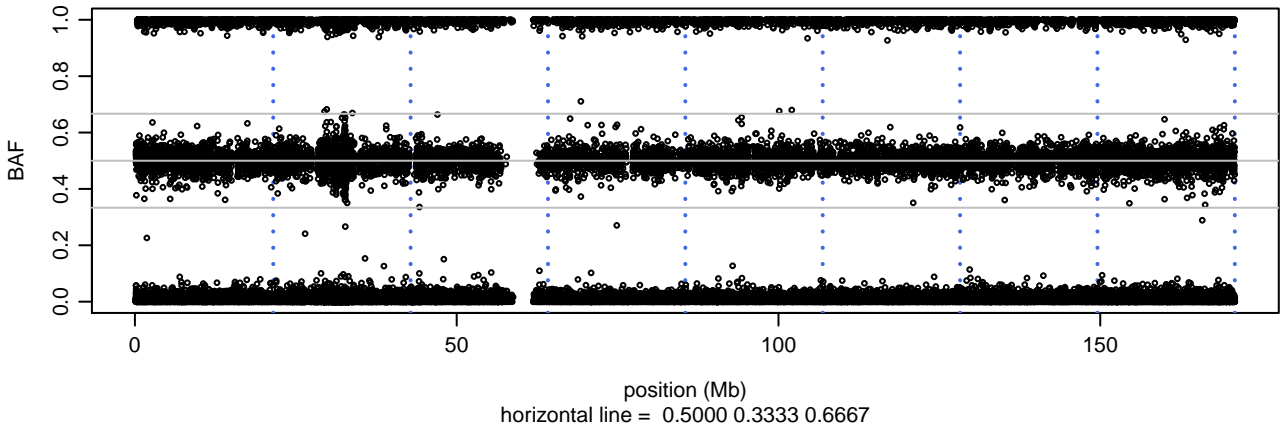
Scan 280 – Chromosome 8 – SNCB_P123H_D11

position (Mb)
horizontal line = mean LRR

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

position (Mb)
horizontal line = 0.5000 0.3333 0.6667
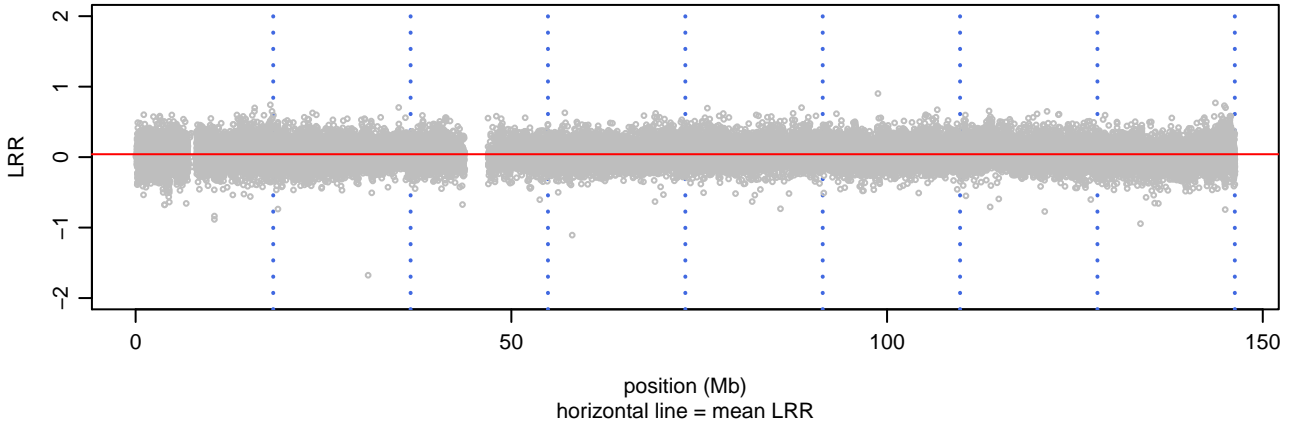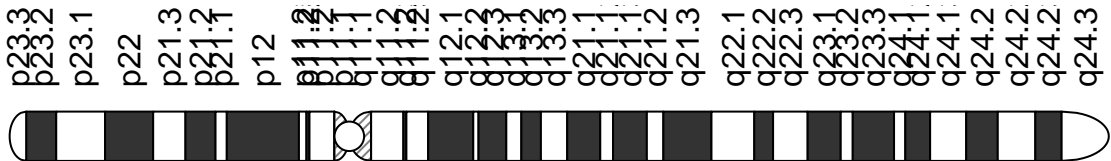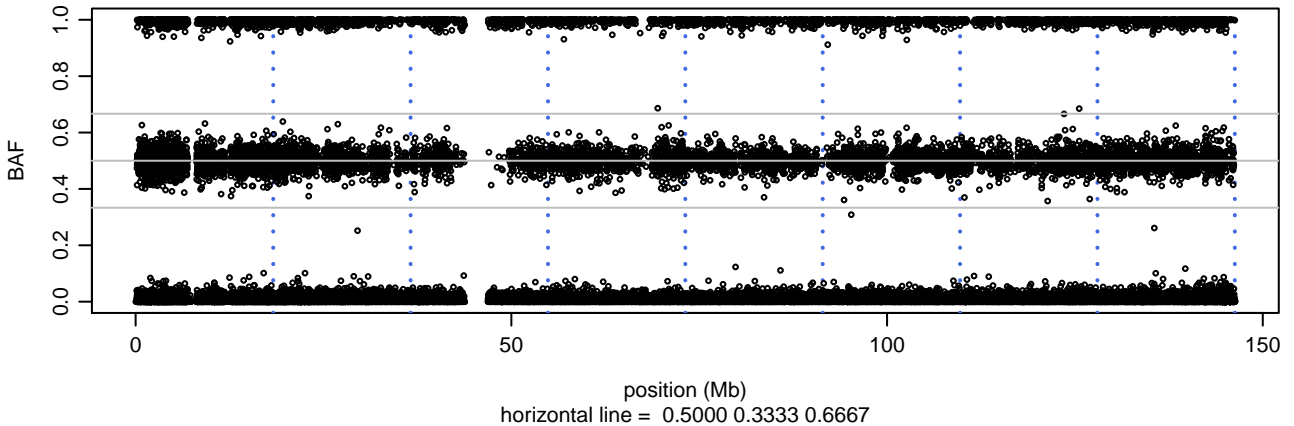
Scan 280 – Chromosome 9 – SNCB_P123H_D11

position (Mb)
horizontal line = mean LRR

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

position (Mb)
horizontal line = 0.5000 0.3333 0.6667

Scan 280 – Chromosome 10 – SNCB_P123H_D11

Scan 280 – Chromosome 11 – SNCB_P123H_D11

LRR

position (Mb)
horizontal line = mean LRR

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

BAF

position (Mb)
horizontal line = 0.5000 0.3333 0.6667
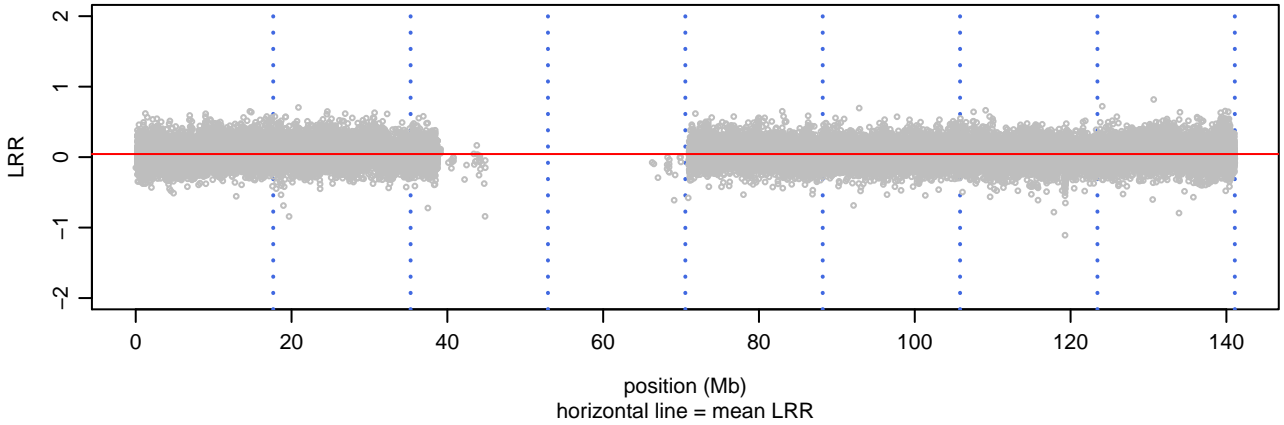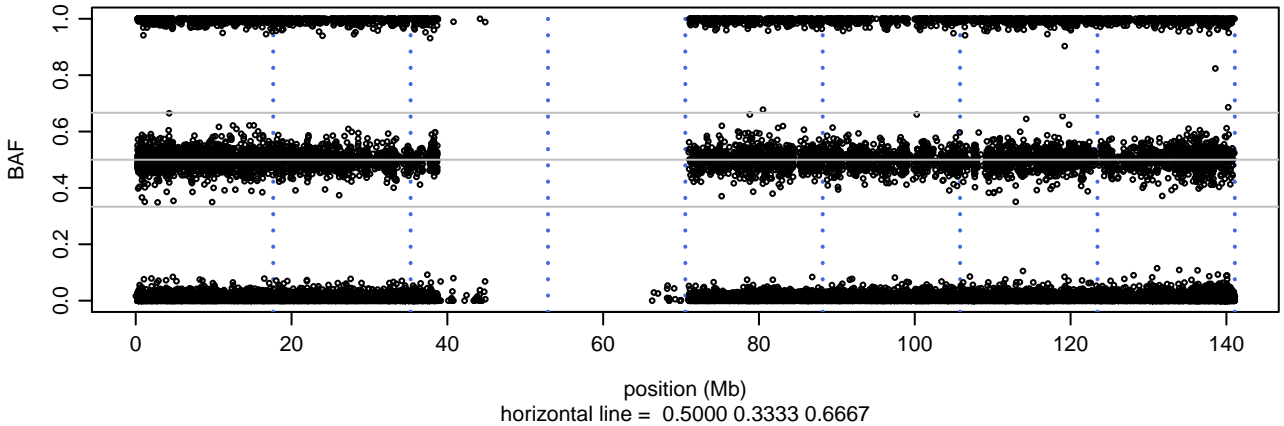
Scan 280 − Chromosome 12 − SNCB_P123H_D11

position (Mb)
horizontal line = mean LRR

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

position (Mb)
horizontal line = 0.5000 0.3333 0.6667
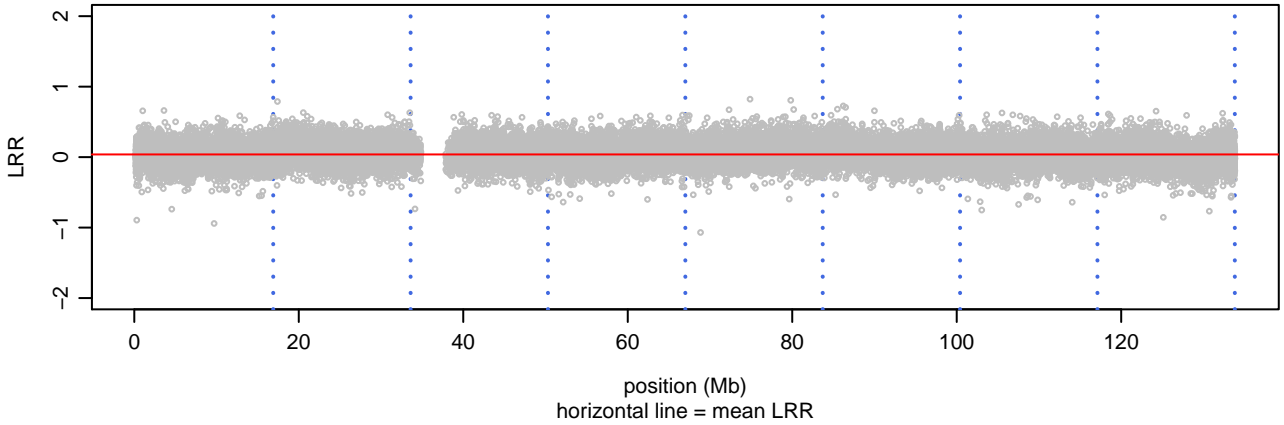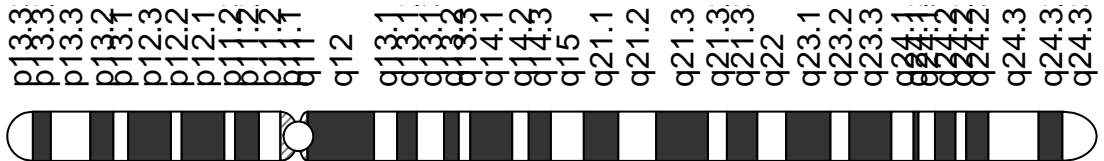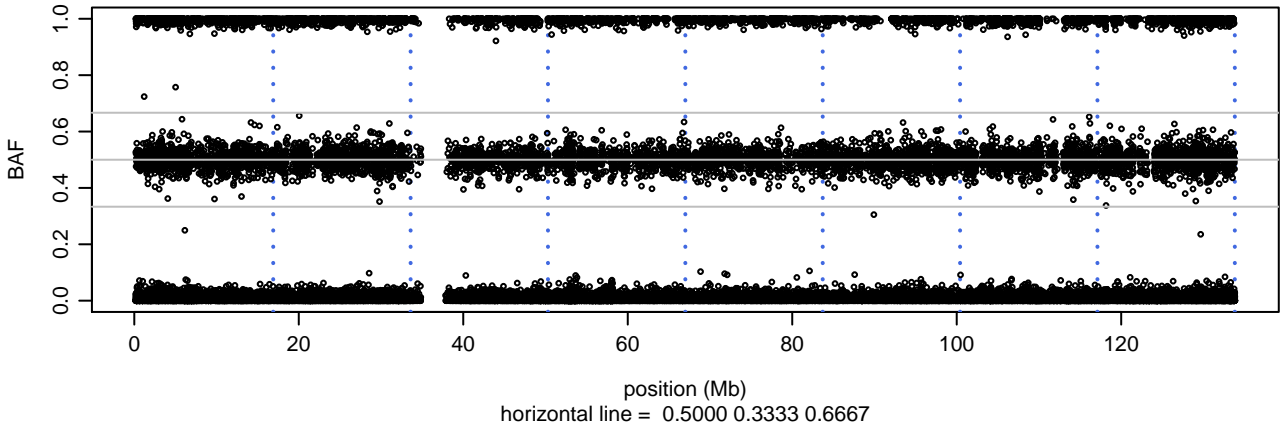
Scan 280 – Chromosome 13 – SNCB_P123H_D11

position (Mb)
horizontal line = mean LRR

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

position (Mb)
horizontal line = 0.5000 0.3333 0.6667
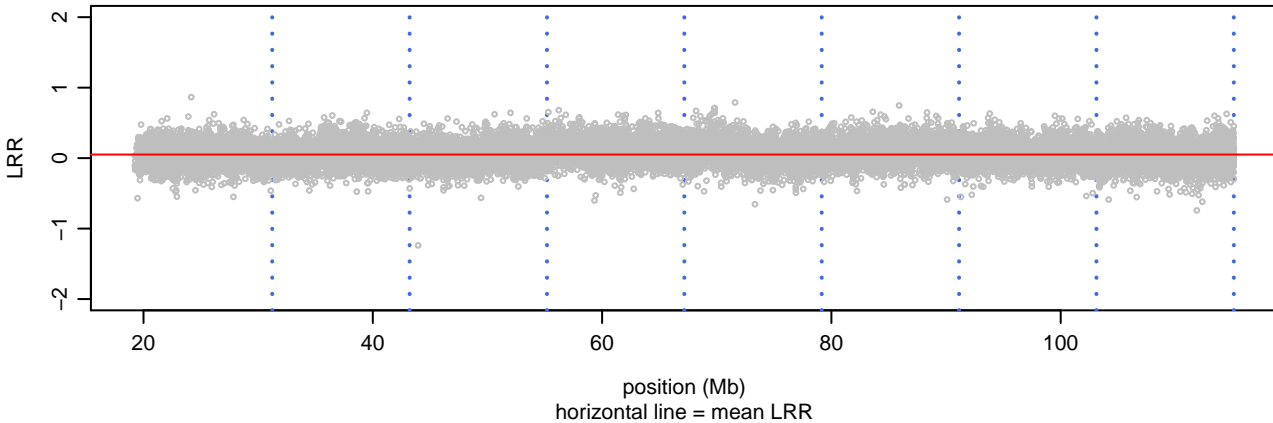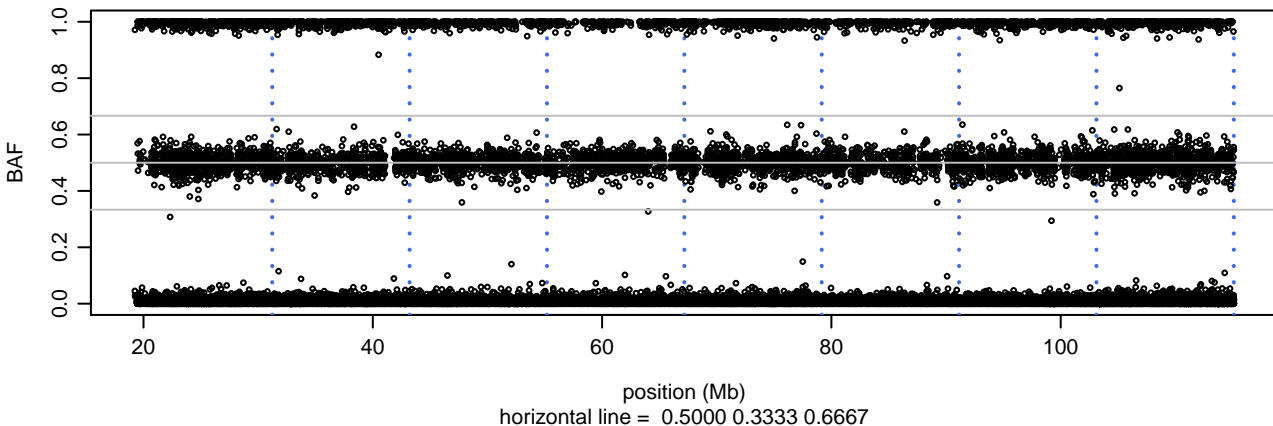
Scan 280 − Chromosome 14 − SNCB_P123H_D11

position (Mb)
horizontal line = mean LRR

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

position (Mb)
horizontal line = 0.5000 0.3333 0.6667
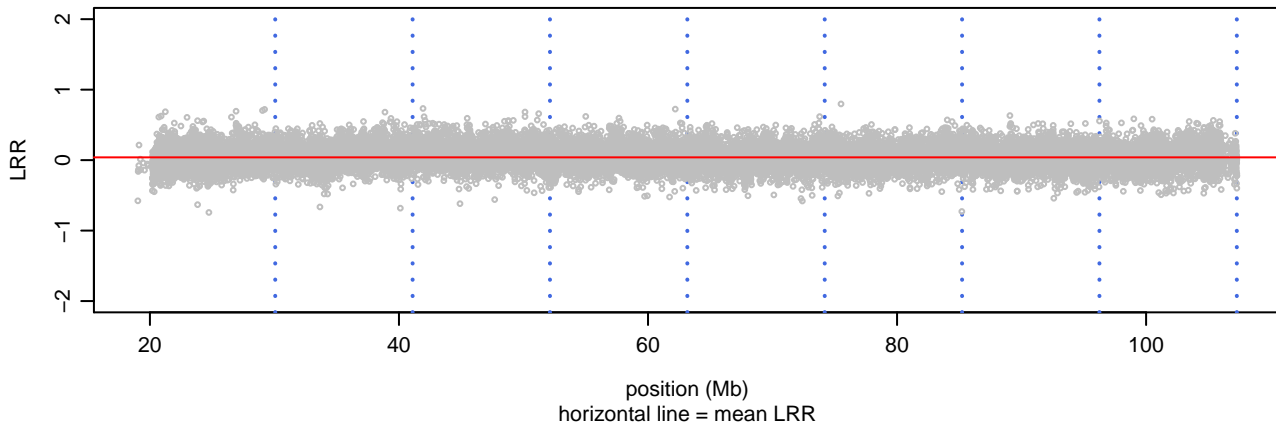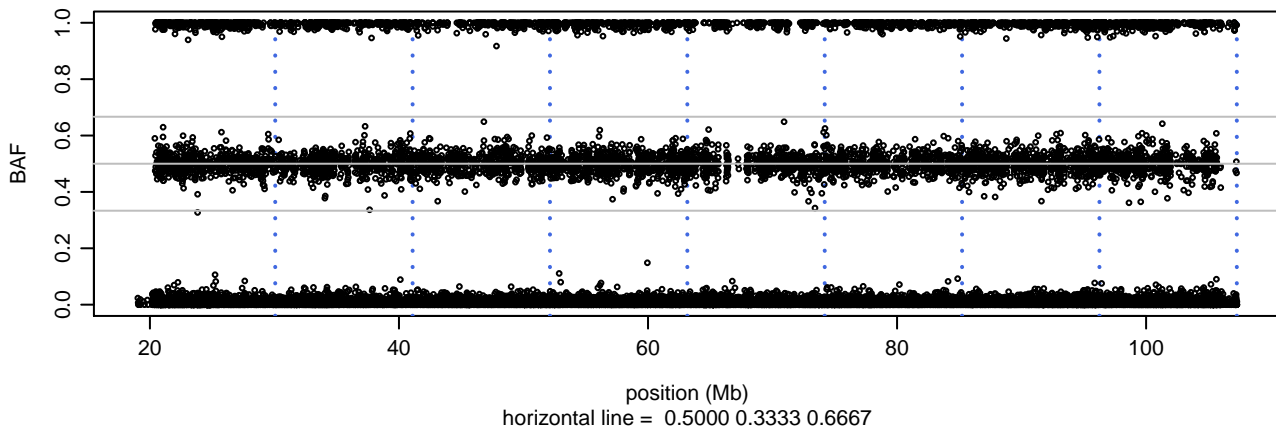
Scan 280 – Chromosome 15 – SNCB_P123H_D11
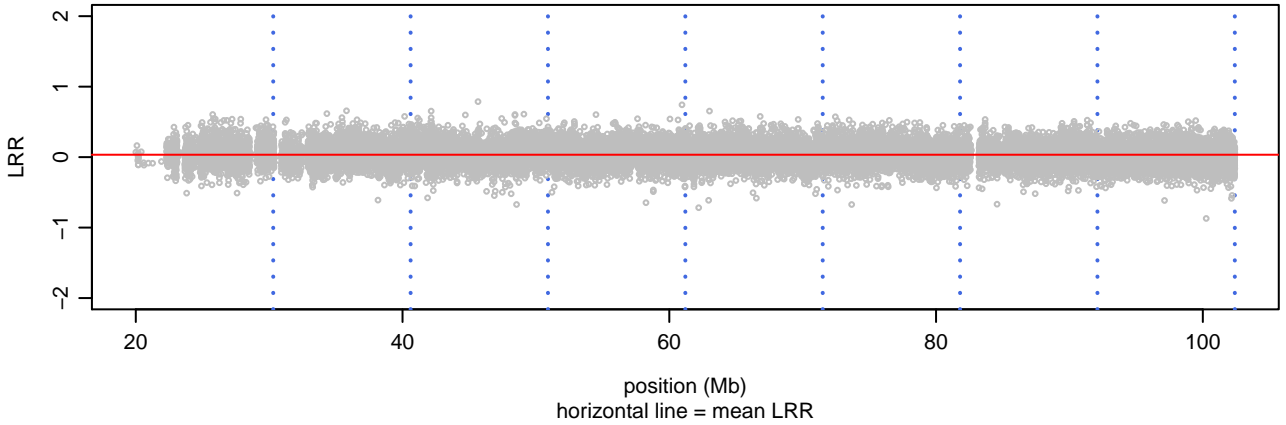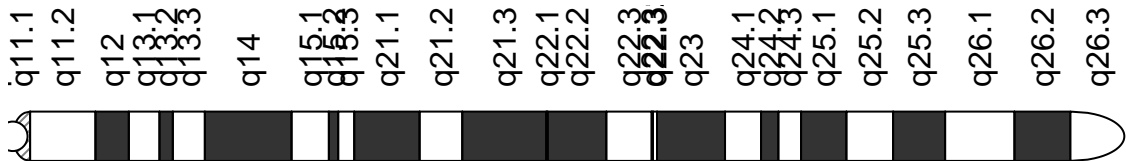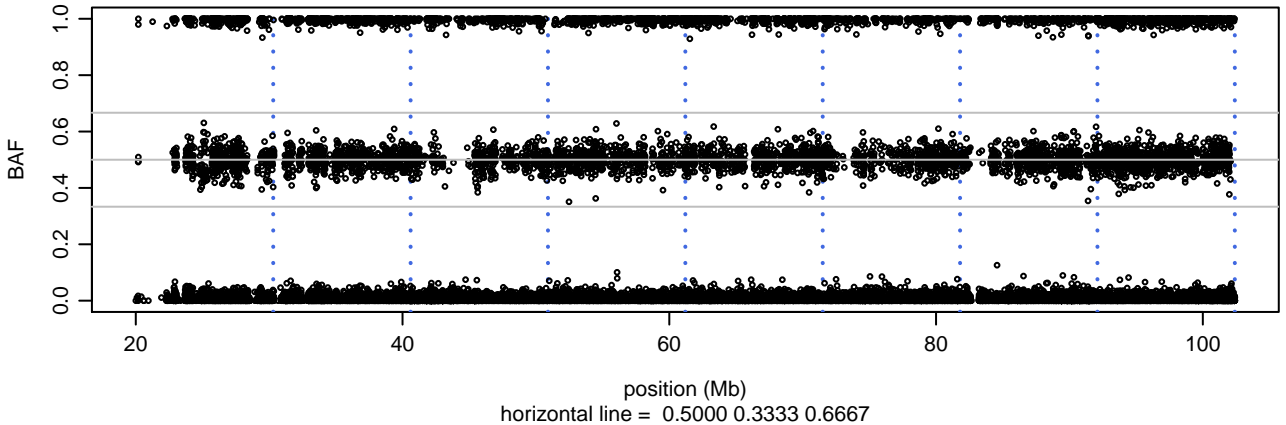
Scan 280 – Chromosome 16 – SNCB_P123H_D11

position (Mb)
horizontal line = mean LRR

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

position (Mb)
horizontal line = 0.5000 0.3333 0.6667
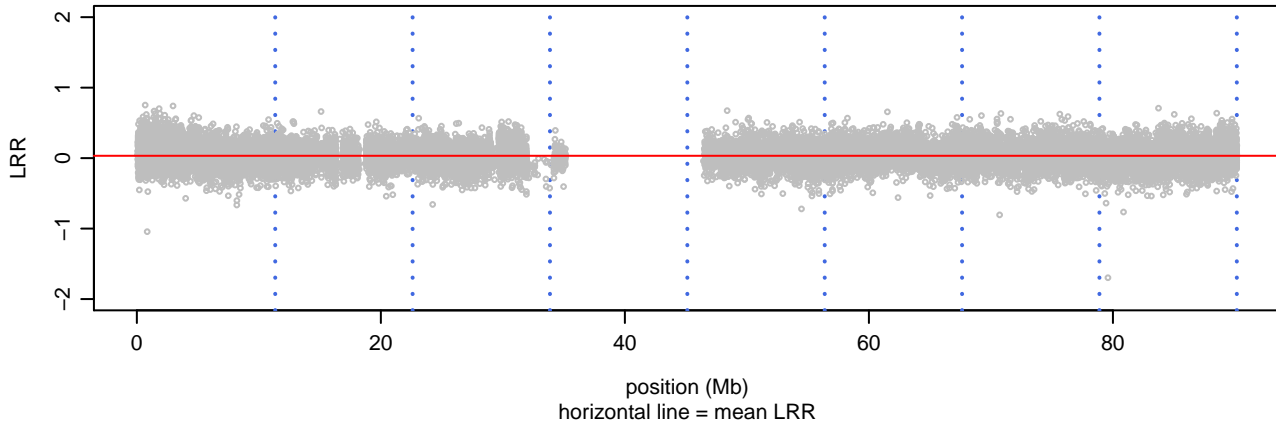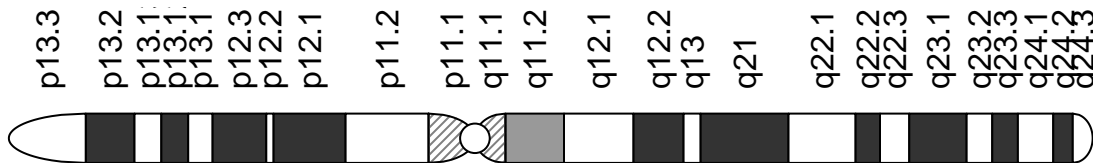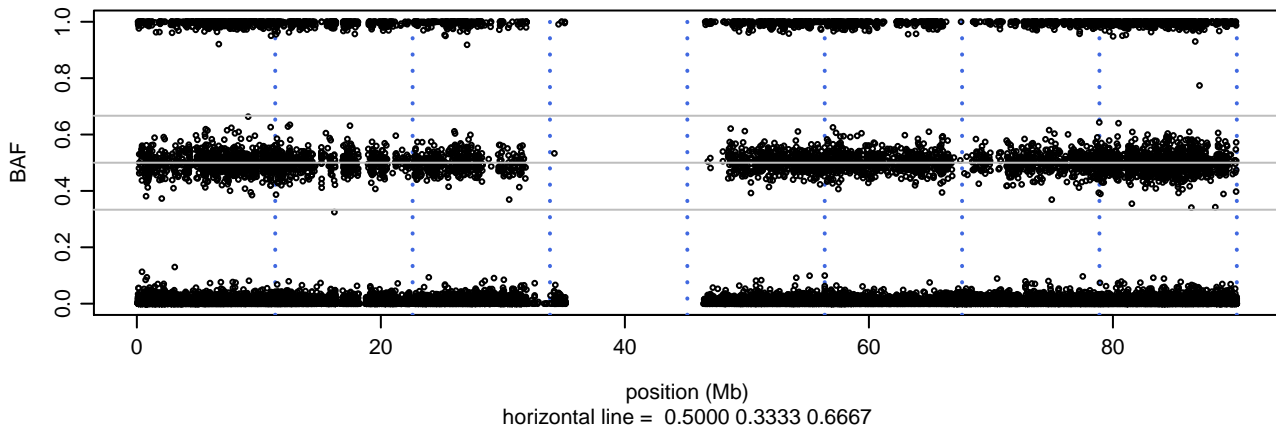
Scan 280 – Chromosome 17 – SNCB_P123H_D11
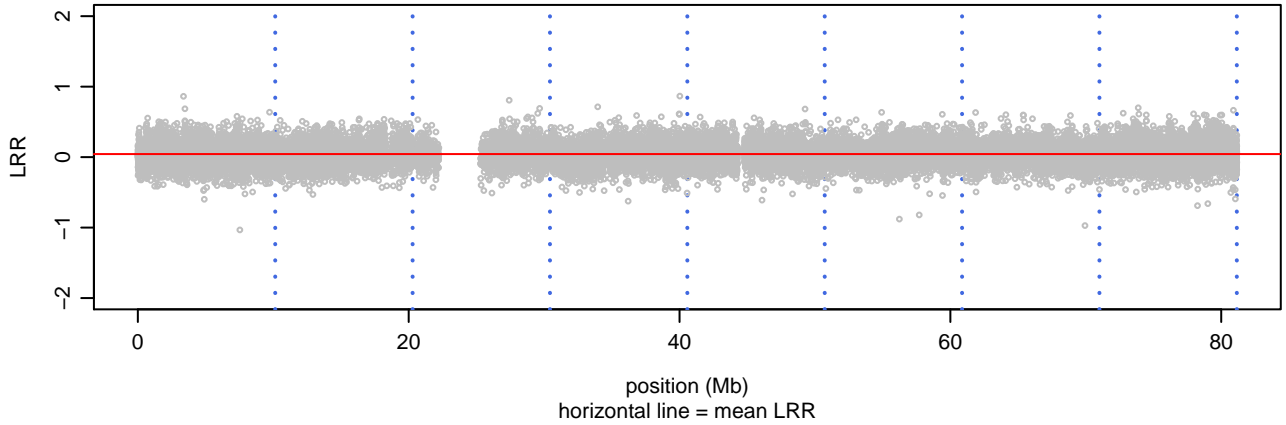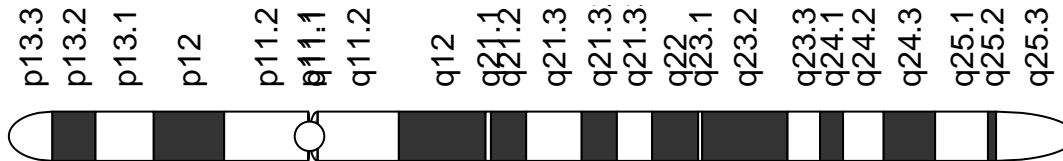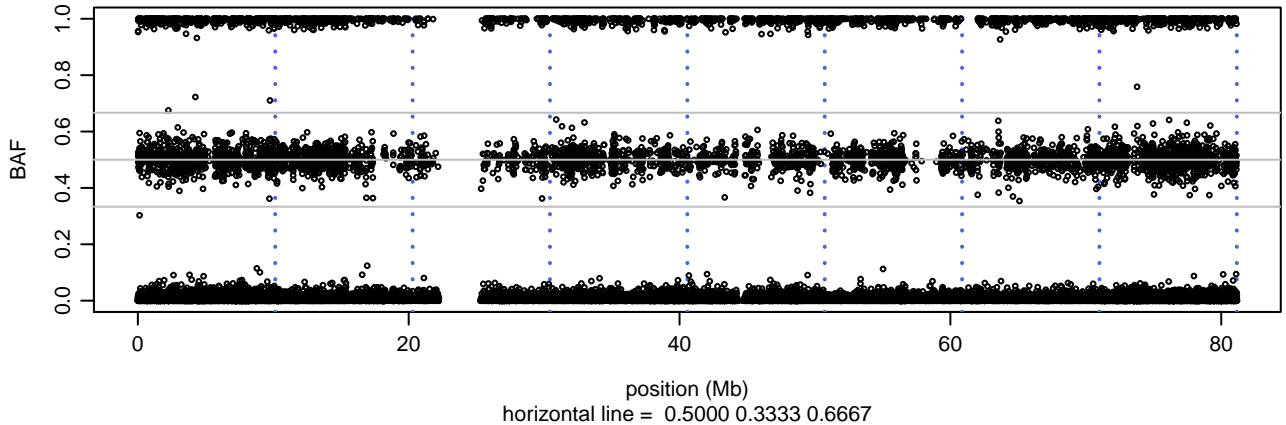
Scan 280 – Chromosome 18 – SNCB_P123H_D11

LRR

position (Mb)
horizontal line = mean LRR

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

BAF

position (Mb)
horizontal line = 0.5000 0.3333 0.6667

p11.3 p11.3 p11.2 p11.2 p11.2 p11.1 q11.2 q12.1 q12.2 q12.3 q21.1 q21.2 q21.3 q21.3 q21.3 q22.1 q22.2 q22.3 q23
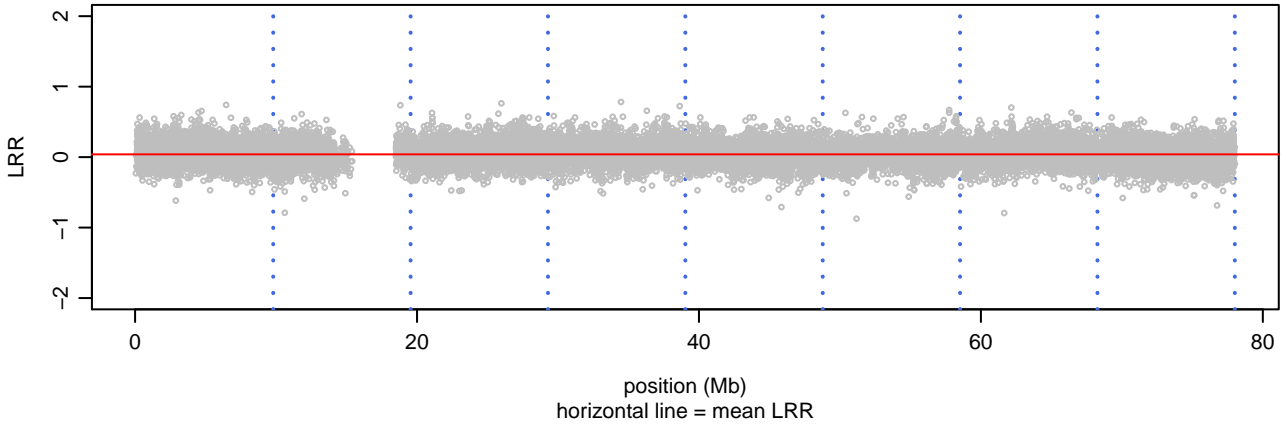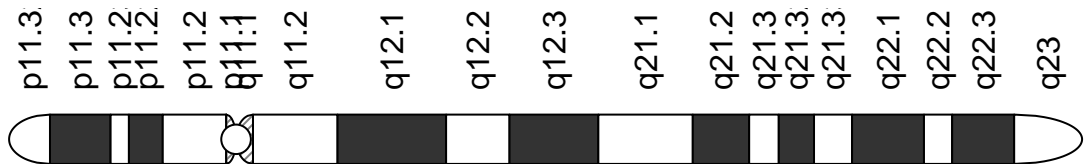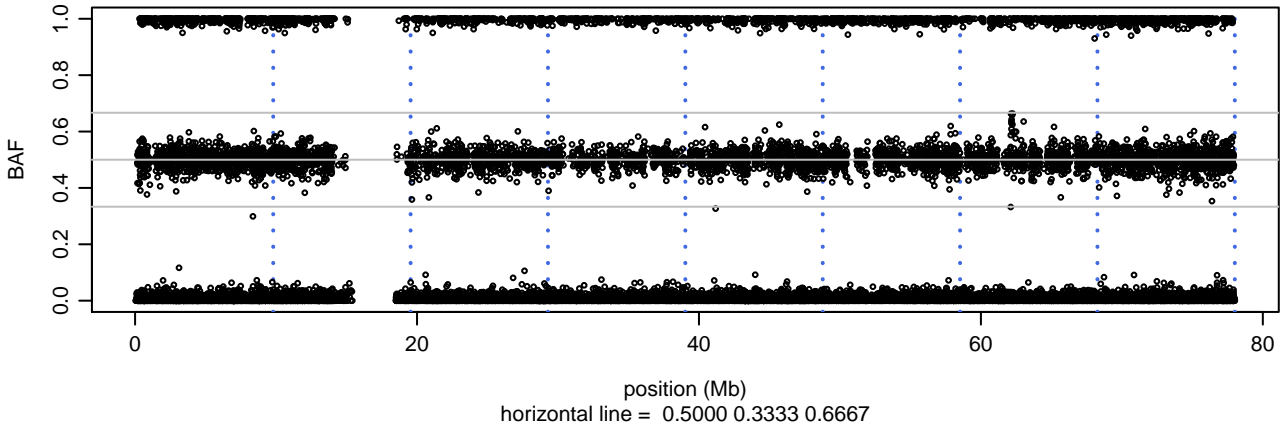
Scan 280 – Chromosome 19 – SNCB_P123H_D11

LRR

position (Mb)
horizontal line = mean LRR

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

BAF

position (Mb)
horizontal line = 0.5000 0.3333 0.6667
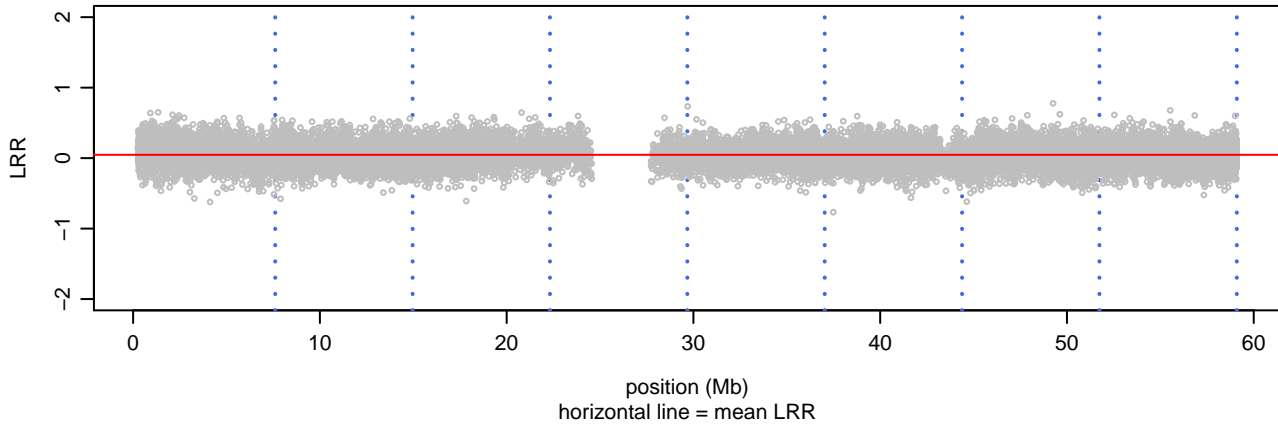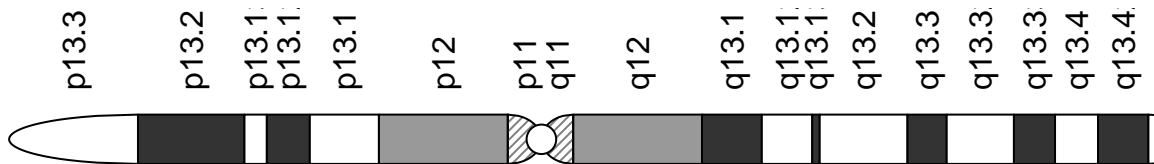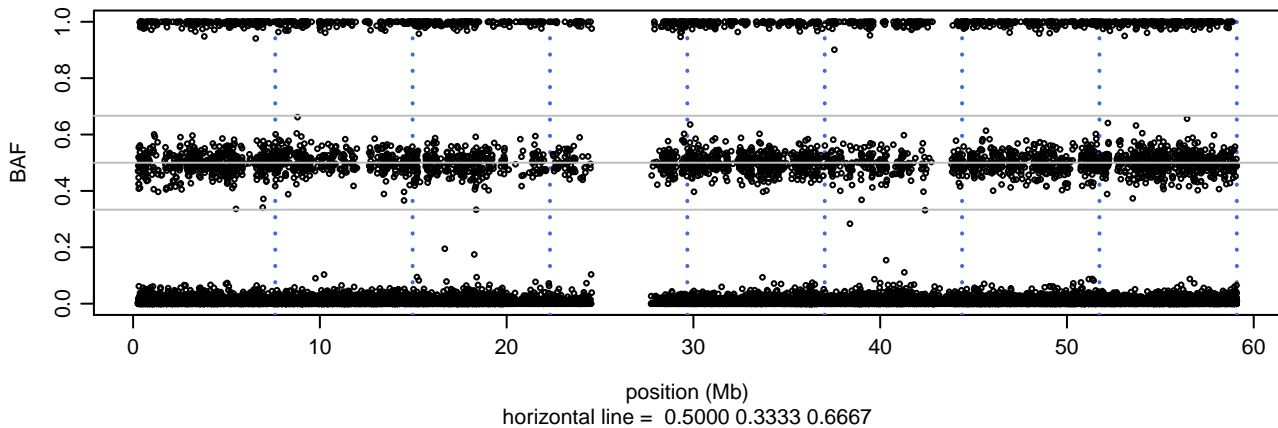
Scan 280 – Chromosome 20 – SNCB_P123H_D11

LRR

position (Mb)
horizontal line = mean LRR

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

BAF

position (Mb)
horizontal line = 0.5000 0.3333 0.6667
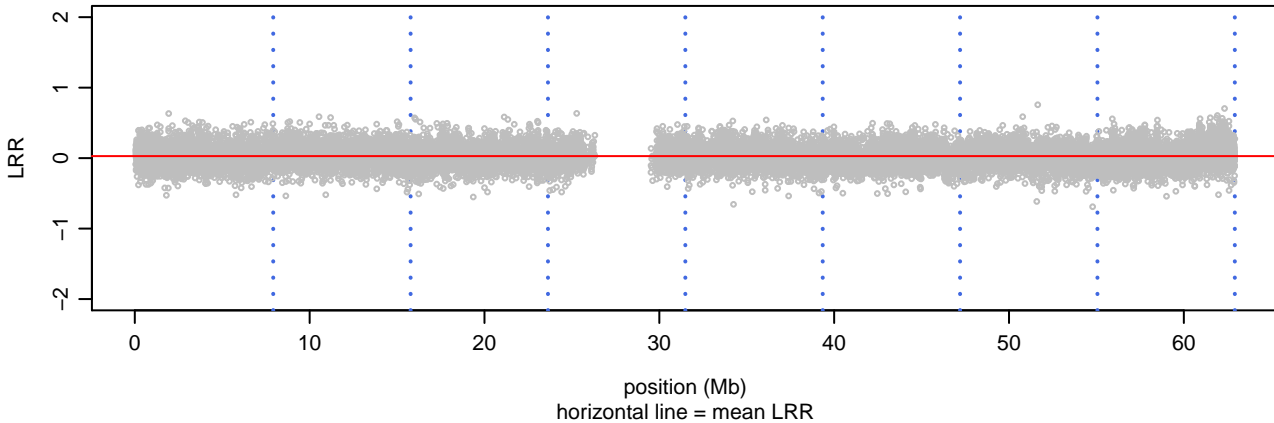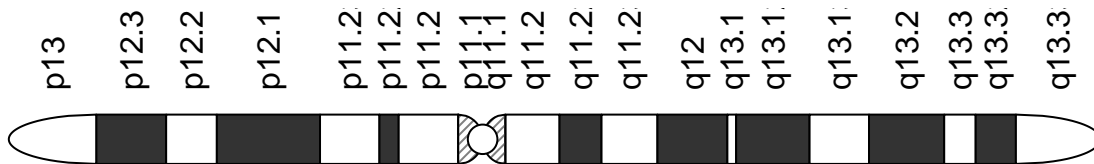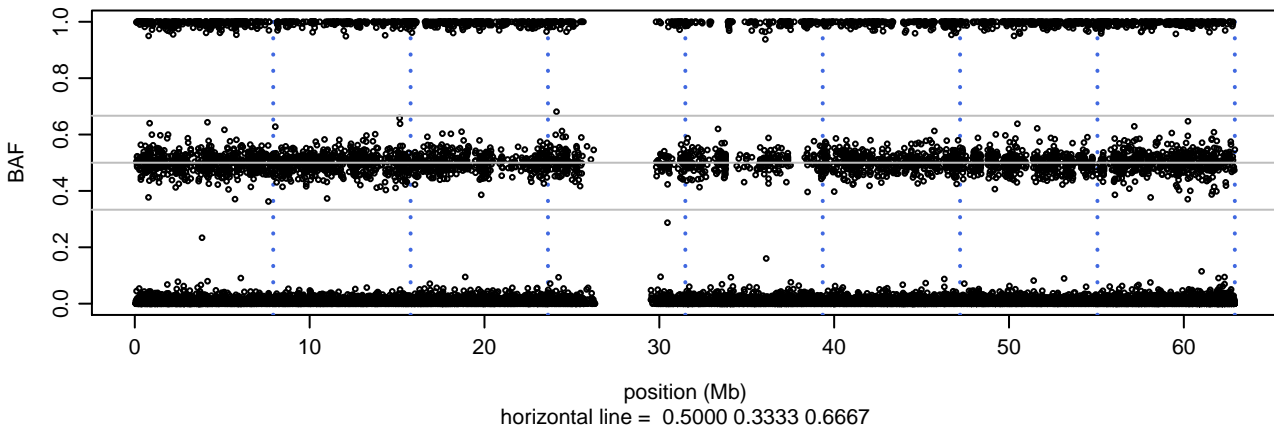
Scan 280 – Chromosome 21 – SNCB_P123H_D11
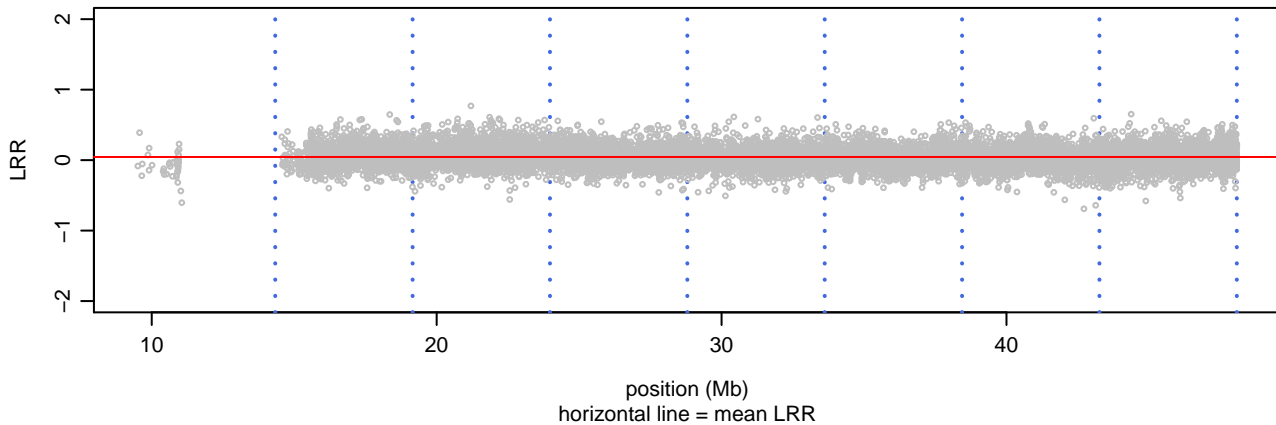
position (Mb)
horizontal line = mean LRR

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

position (Mb)
horizontal line = 0.5000 0.3333 0.6667
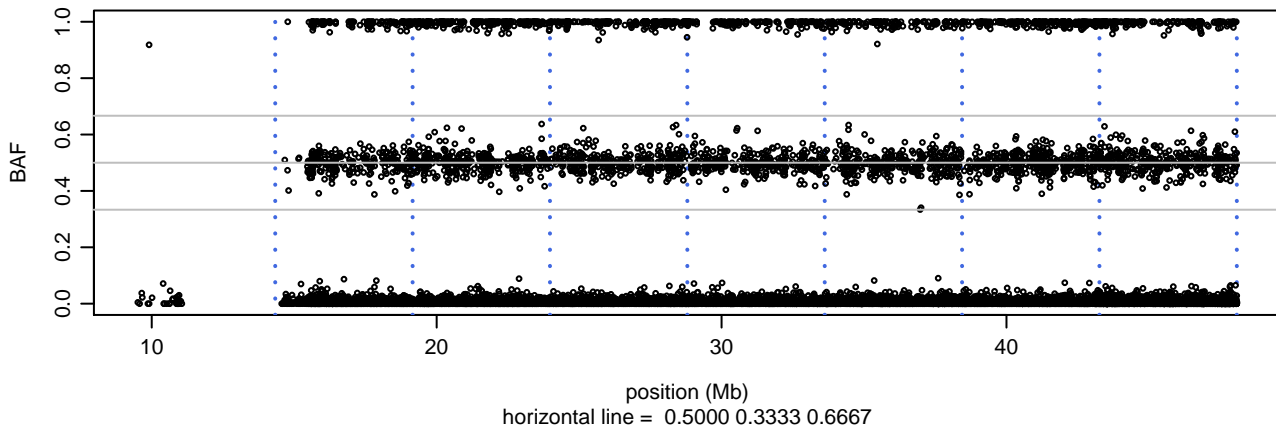
Scan 280 – Chromosome 22 – SNCB_P123H_D11
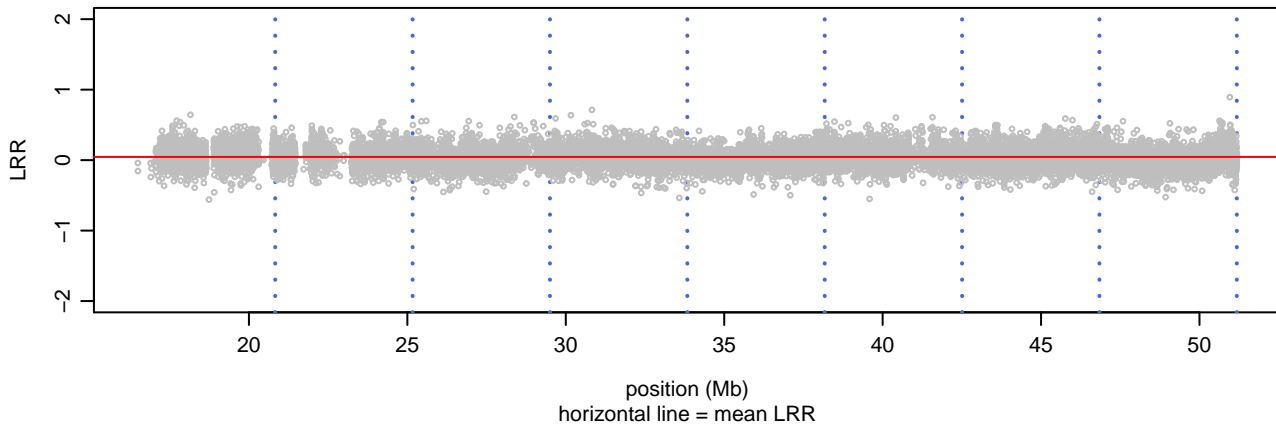
position (Mb)
horizontal line = mean LRR

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

position (Mb)
horizontal line = 0.5000 0.3333 0.6667

Scan 280 – Chromosome X – SNCB_P123H_D11

LRR

position (Mb)
horizontal line = mean LRR

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

BAF

position (Mb)
horizontal line = 0.5000 0.3333 0.6667
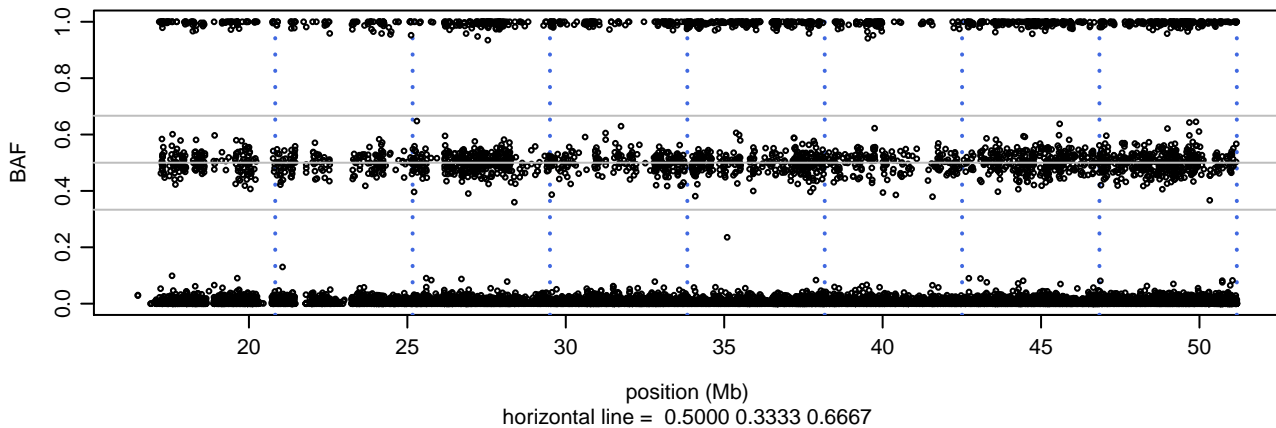
Scan 280 – Chromosome Y – SNCB_P123H_D11

position (Mb)
horizontal line = mean LRR

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

position (Mb)
horizontal line = 0.5000 0.3333 0.6667