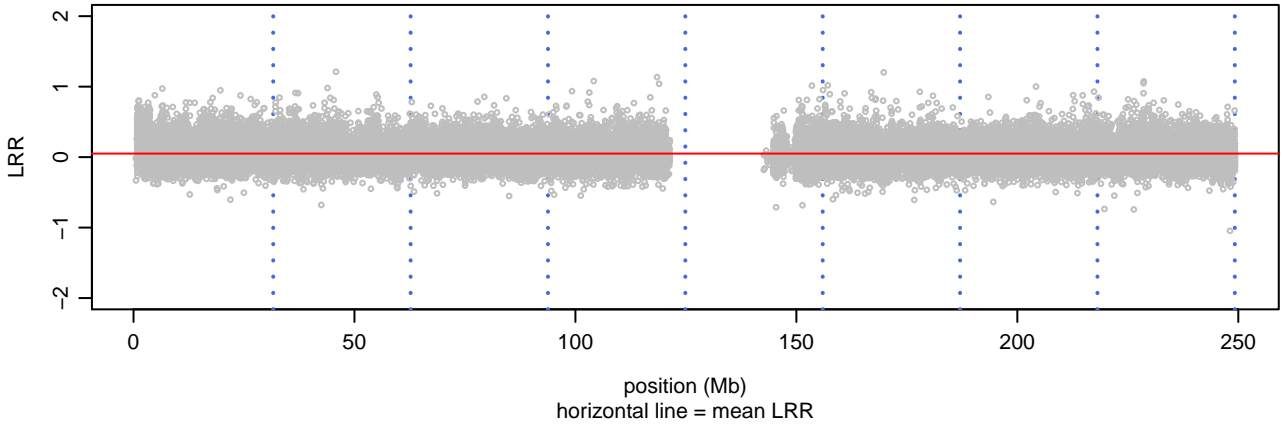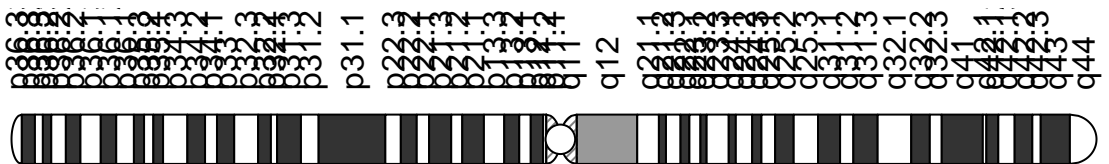**Scan 280 – Chromosome 1 – FUS_R216C_A01**

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
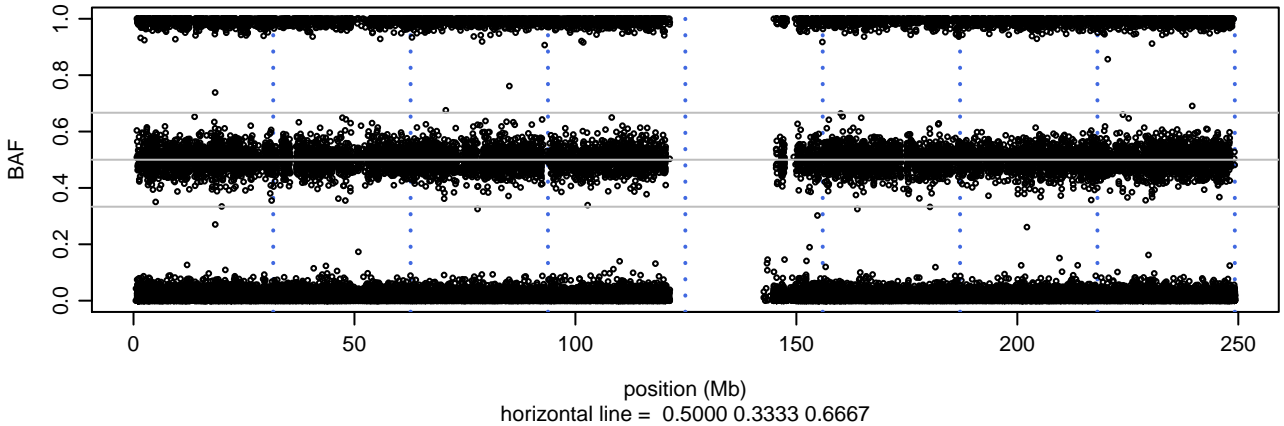
Scan 280 – Chromosome 2 – FUS_R216C_A01

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
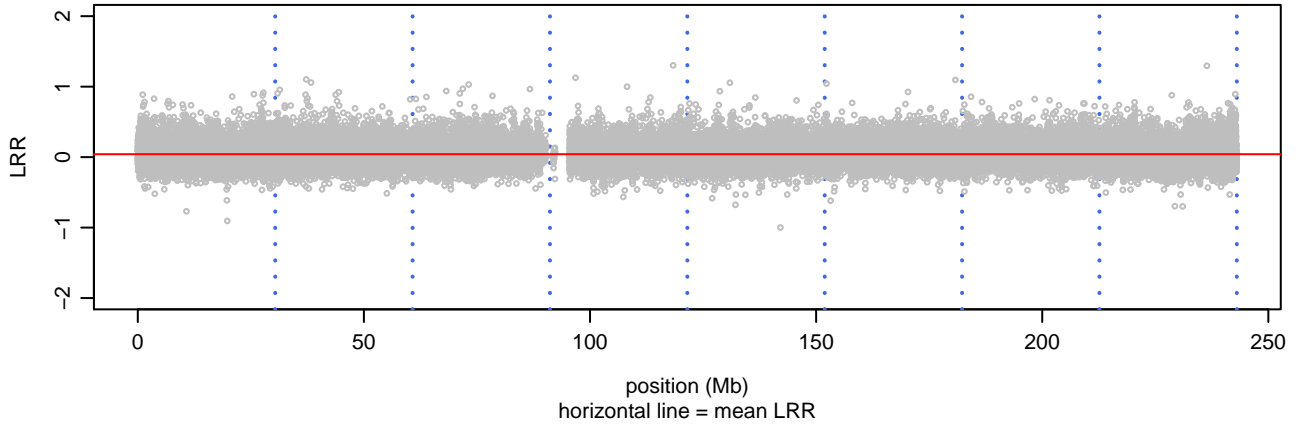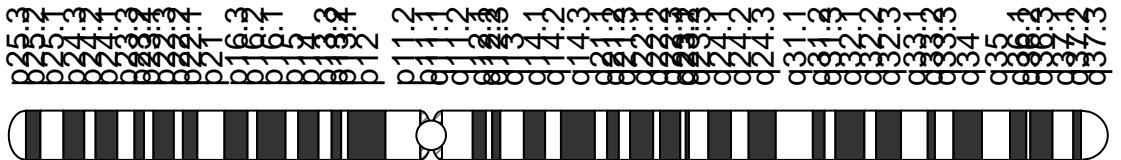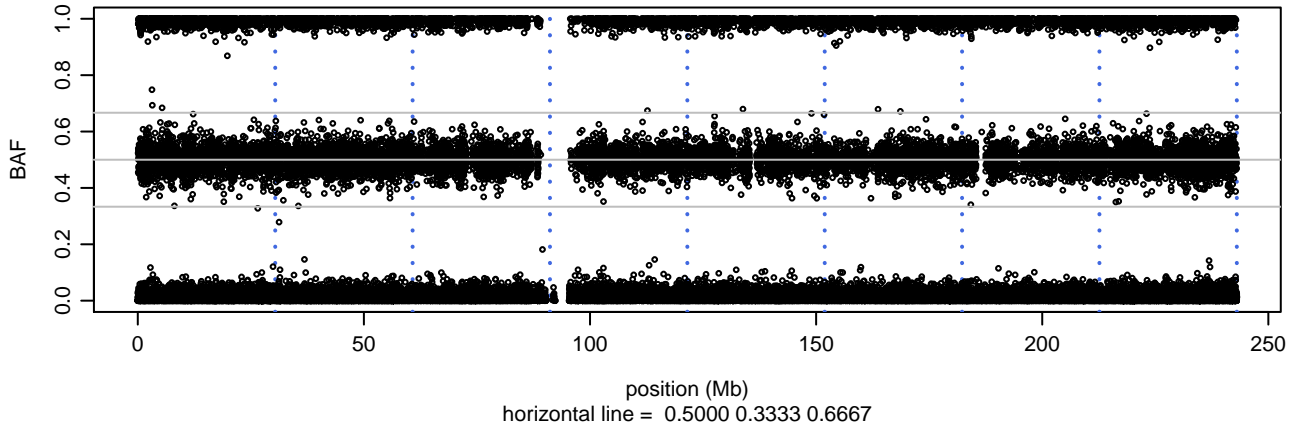
Scan 280 – Chromosome 3 – FUS_R216C_A01

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
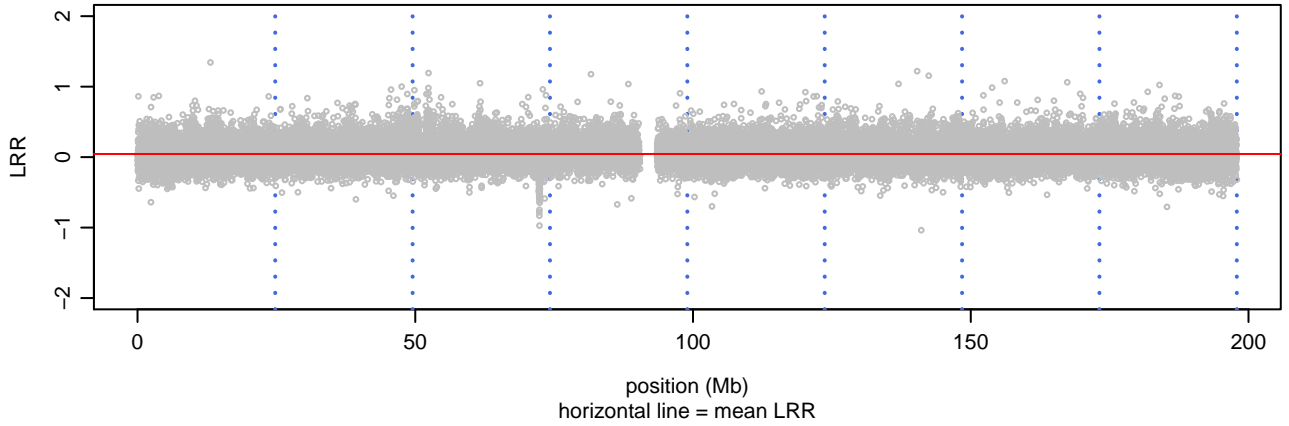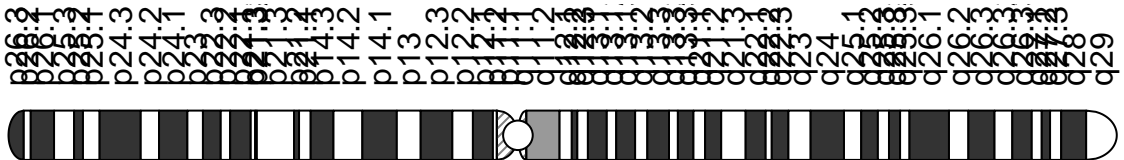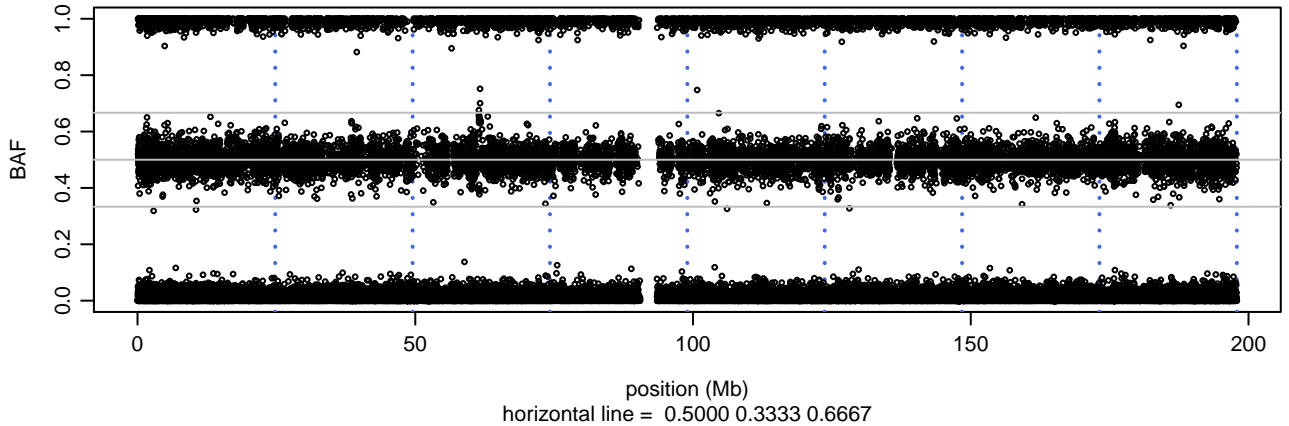
Scan 280 – Chromosome 4 – FUS_R216C_A01

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 – FUS_R216C_A01

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 – FUS_R216C_A01

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 7 – FUS_R216C_A01

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 8 – FUS_R216C_A01

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 – FUS_R216C_A01

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
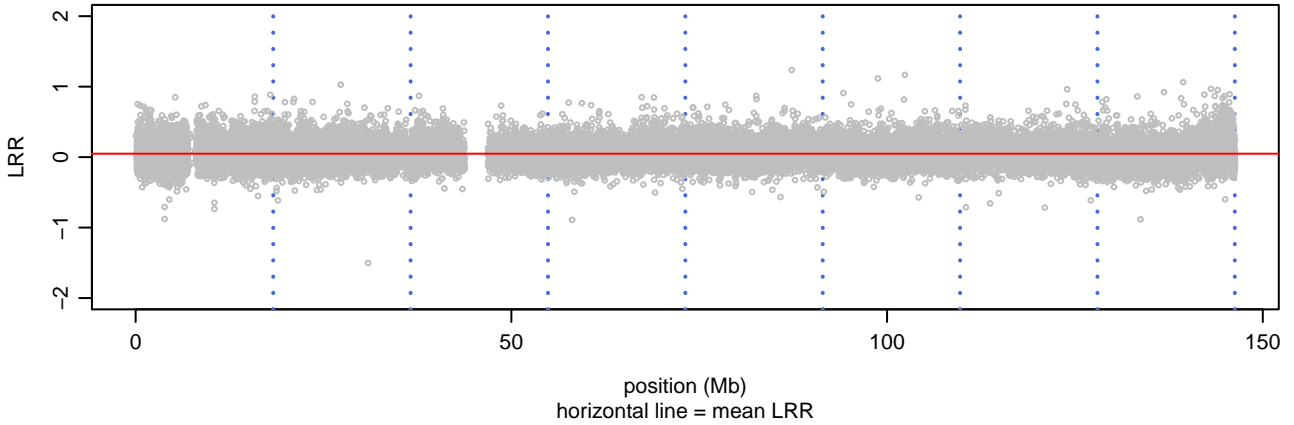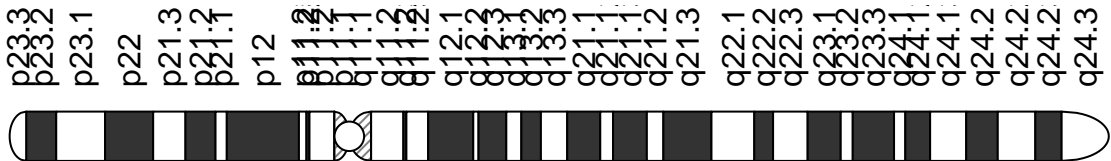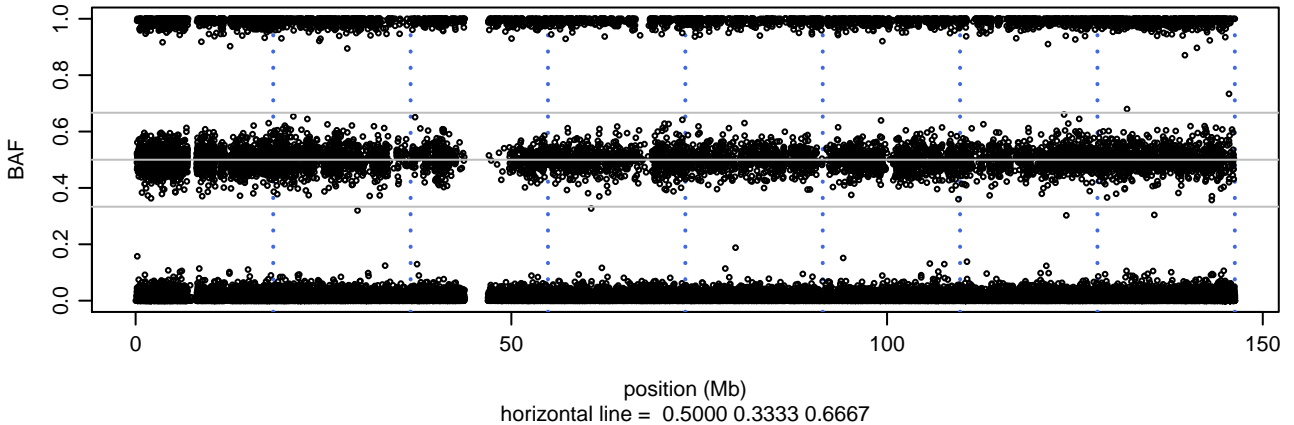
Scan 280 – Chromosome 10 – FUS_R216C_A01

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
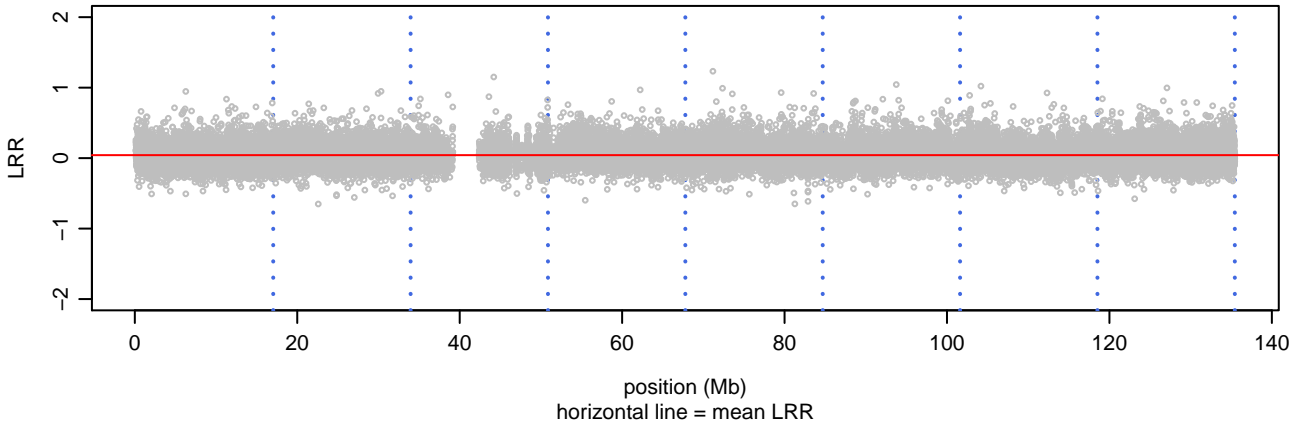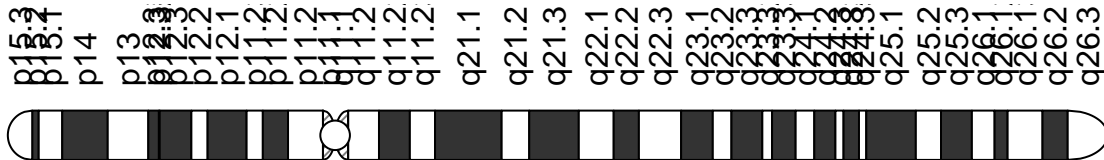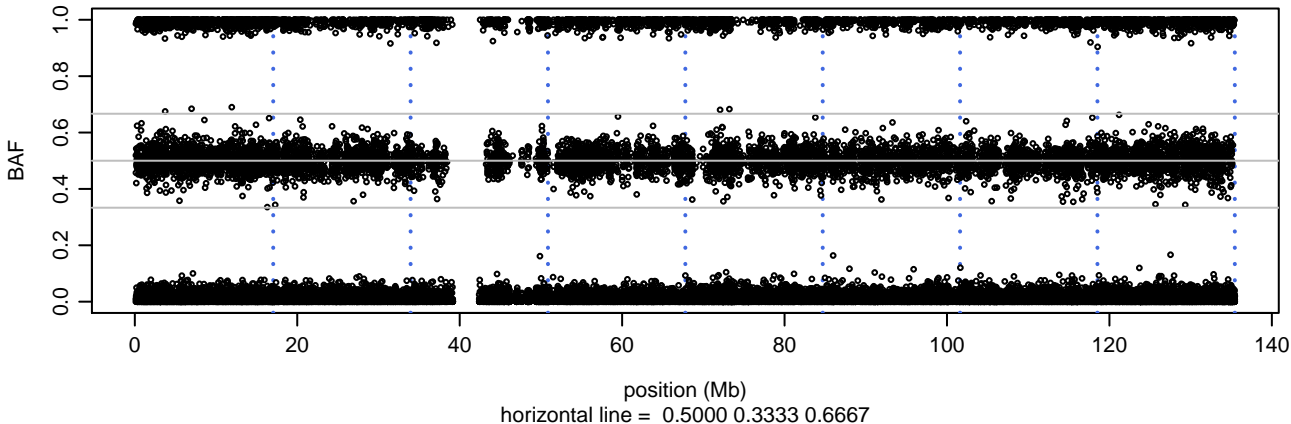
Scan 280 – Chromosome 11 – FUS_R216C_A01

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 12 – FUS_R216C_A01

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
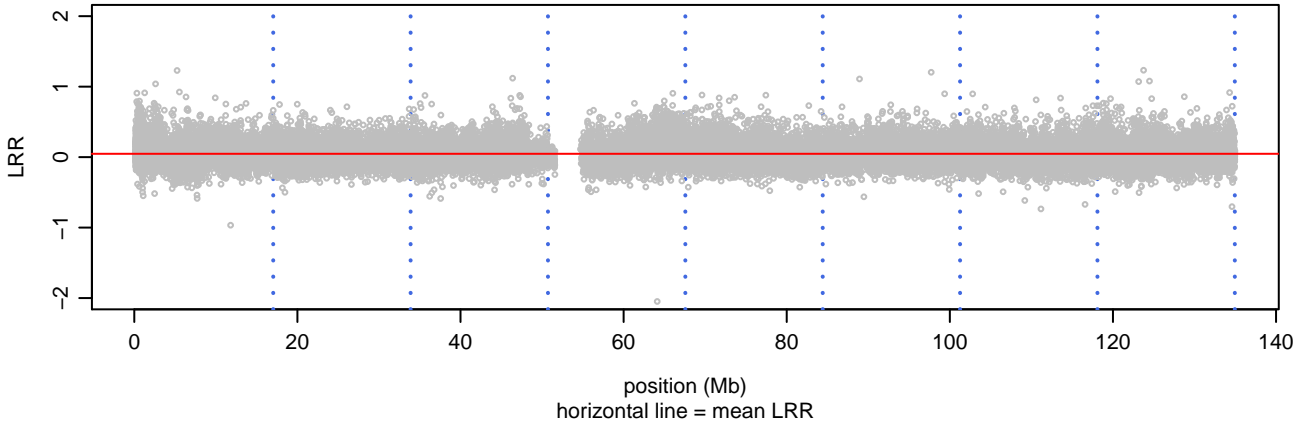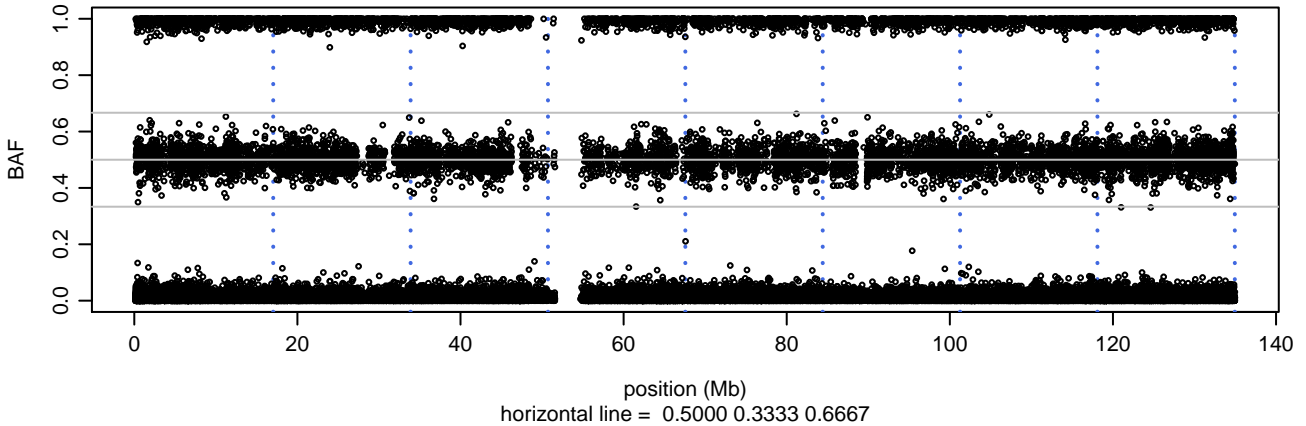
Scan 280 – Chromosome 13 – FUS_R216C_A01

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
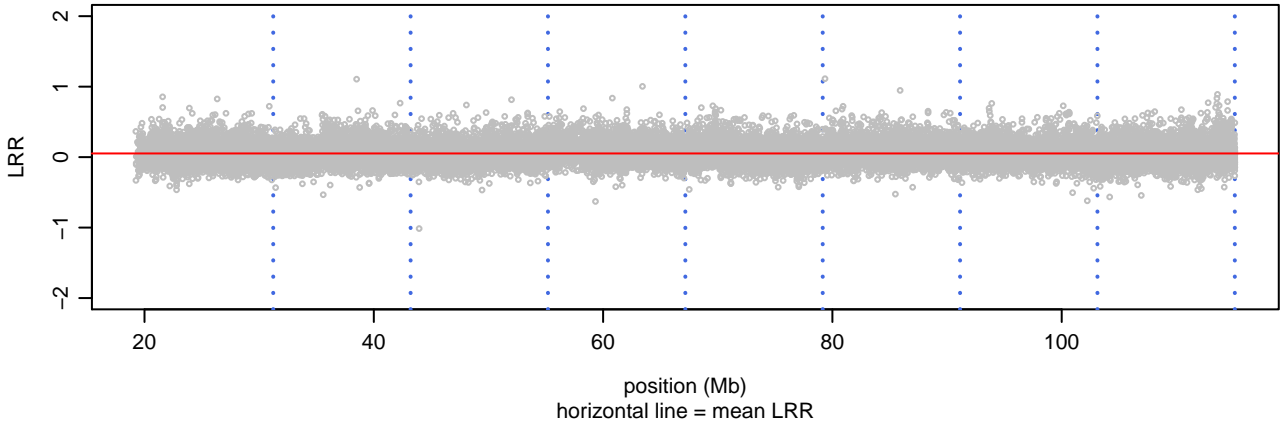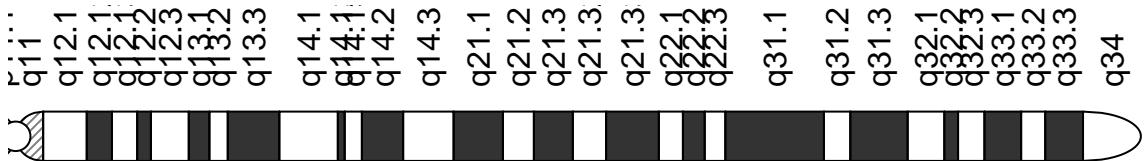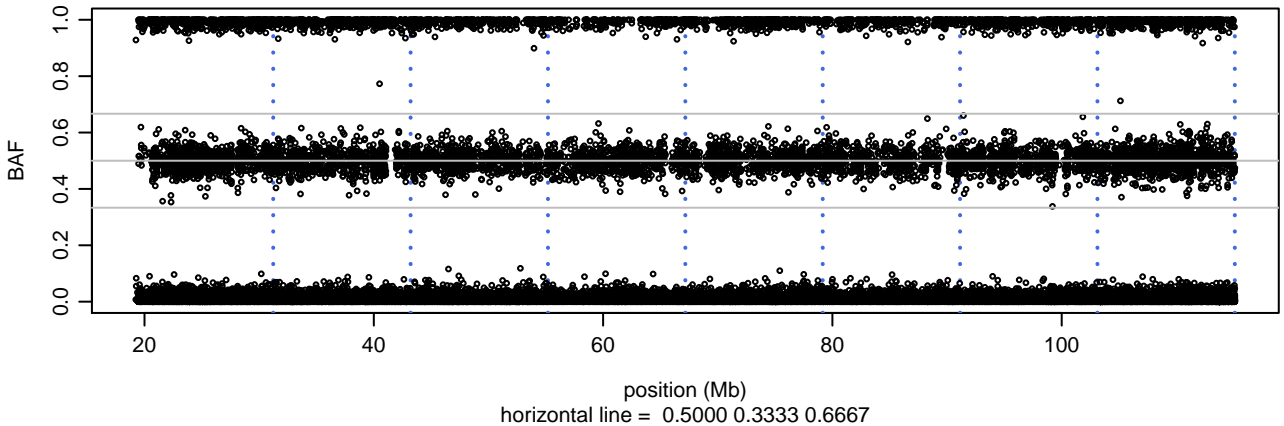
Scan 280 – Chromosome 14 – FUS_R216C_A01

position (Mb)
horizontal line = mean LRR

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

position (Mb)
horizontal line = 0.5000 0.3333 0.6667
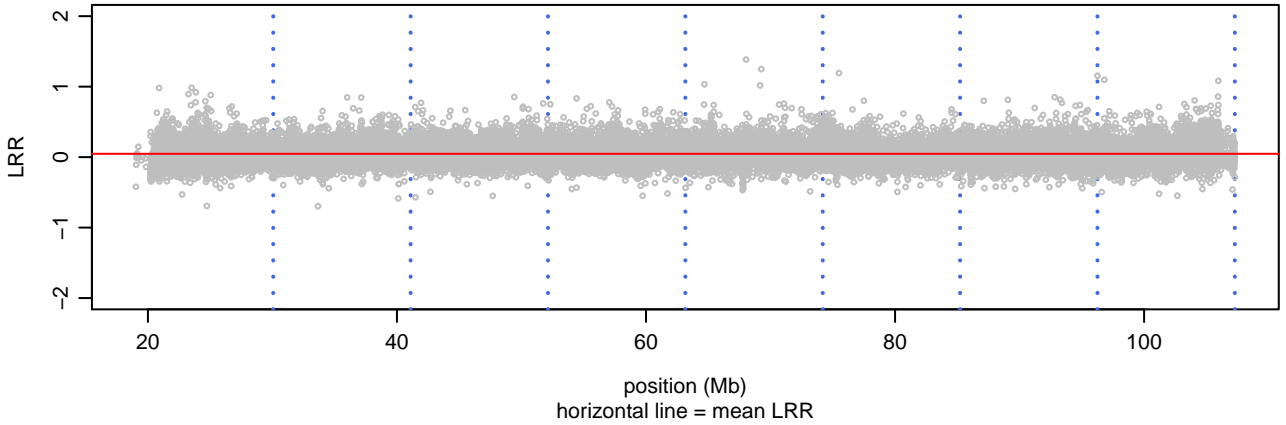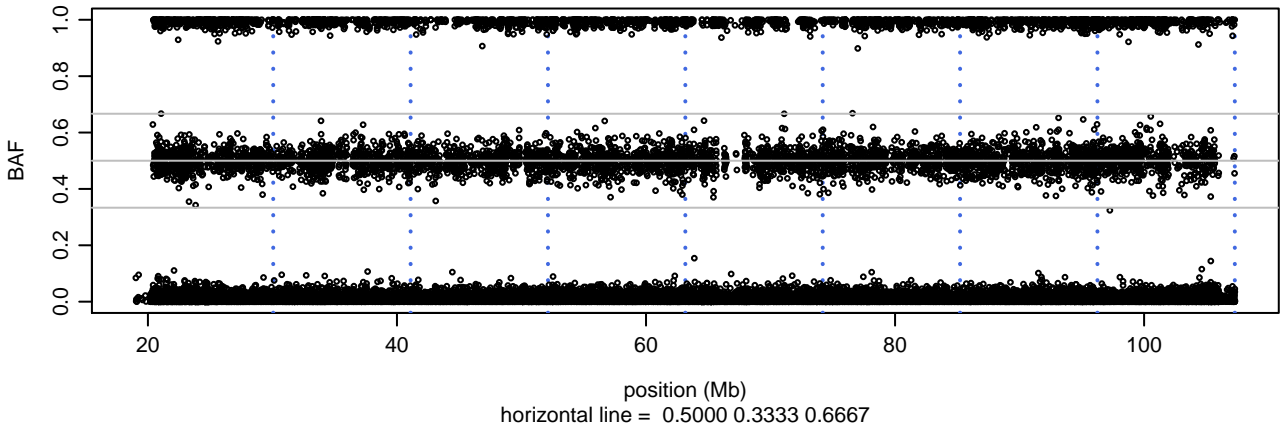
Scan 280 – Chromosome 15 – FUS_R216C_A01

position (Mb)
horizontal line = mean LRR

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

position (Mb)
horizontal line = 0.5000 0.3333 0.6667
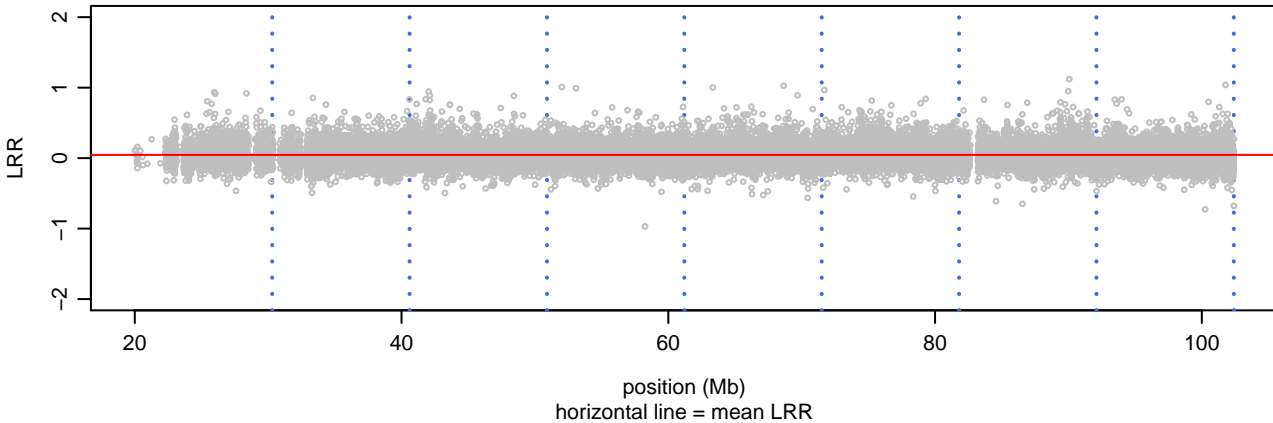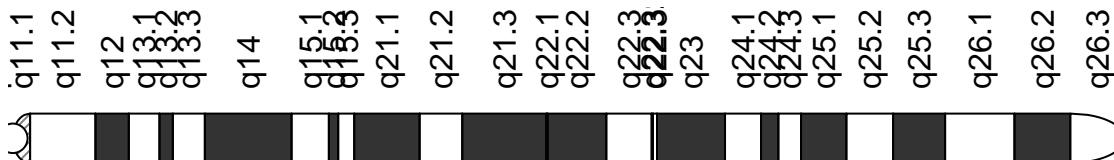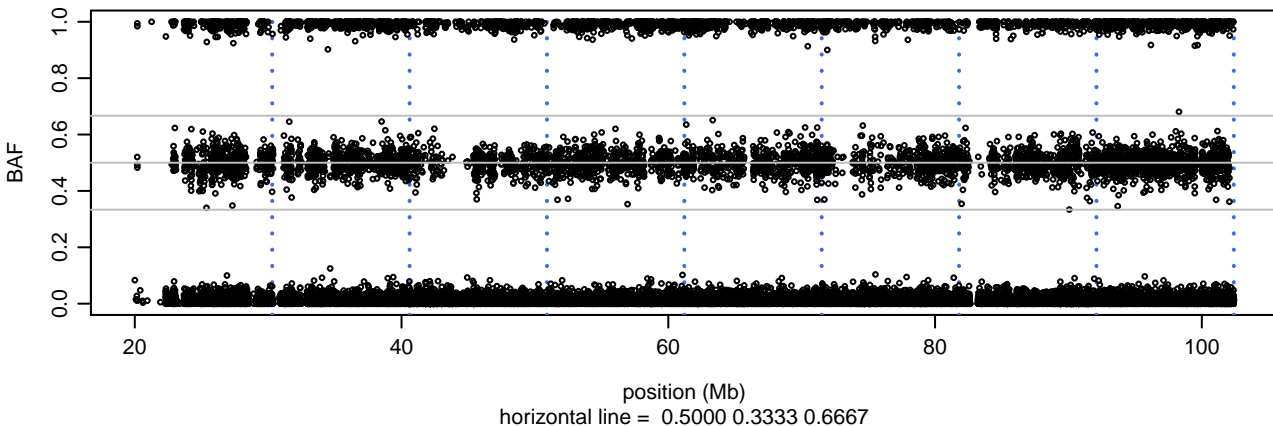
Scan 280 – Chromosome 16 – FUS_R216C_A01

position (Mb)
horizontal line = mean LRR

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

position (Mb)
horizontal line = 0.5000 0.3333 0.6667
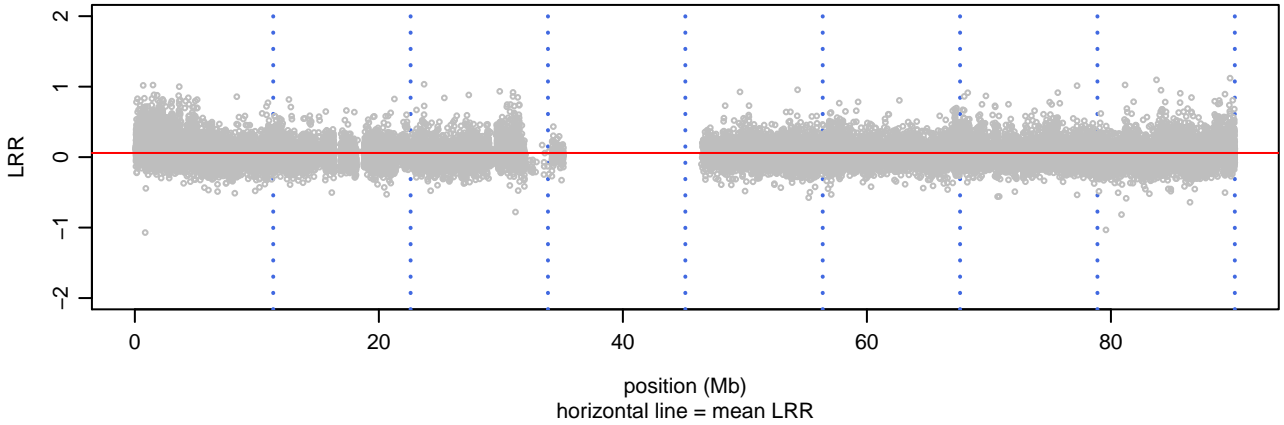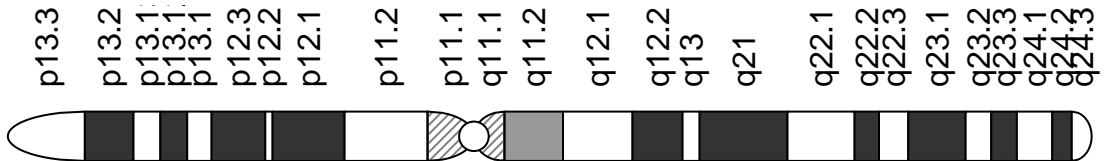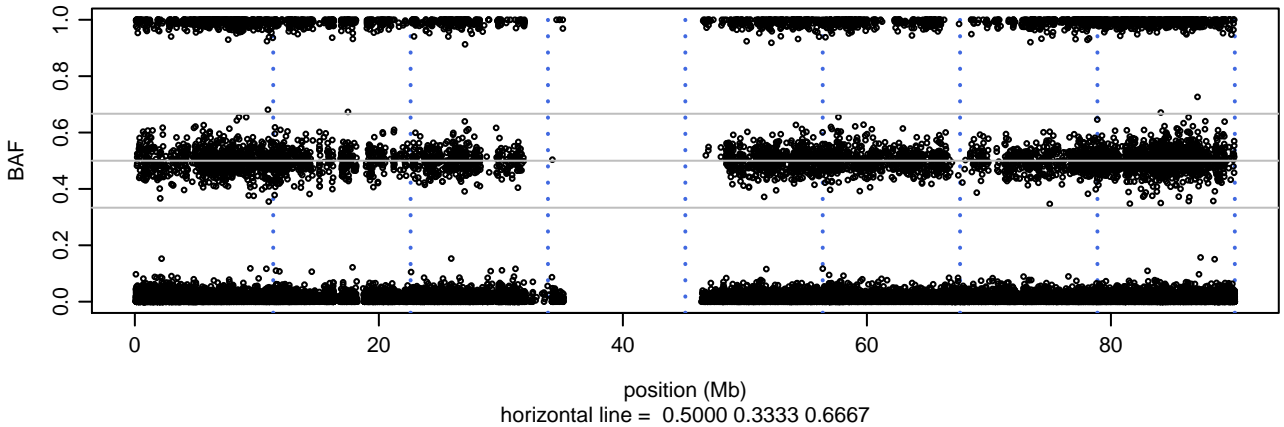
Scan 280 – Chromosome 17 – FUS_R216C_A01

position (Mb)
horizontal line = mean LRR

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

position (Mb)
horizontal line = 0.5000 0.3333 0.6667
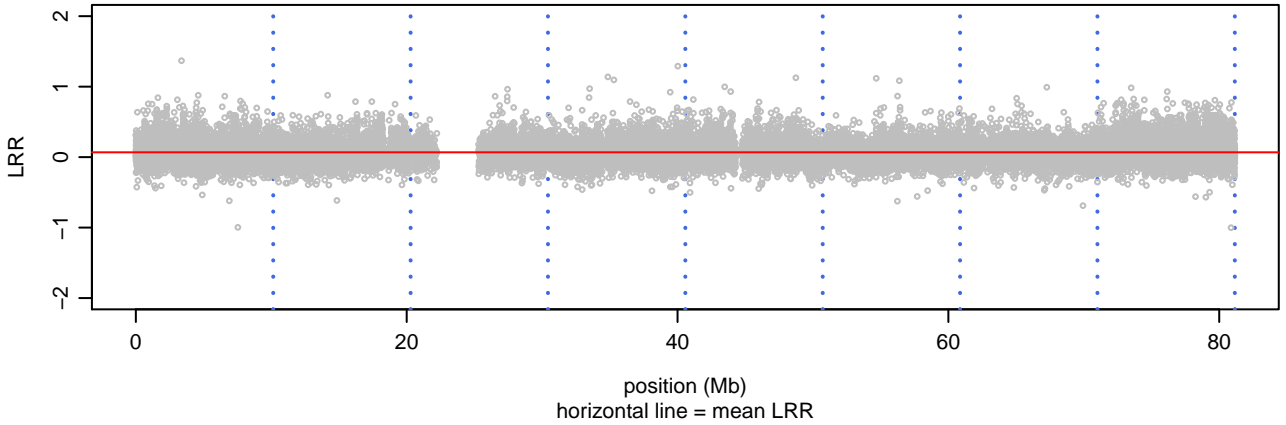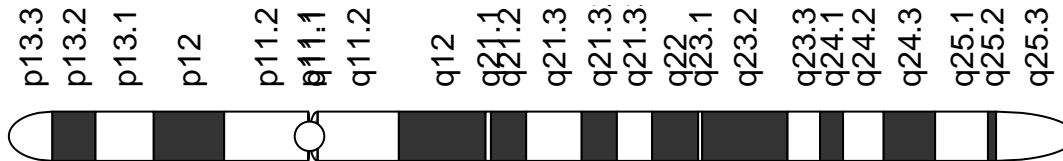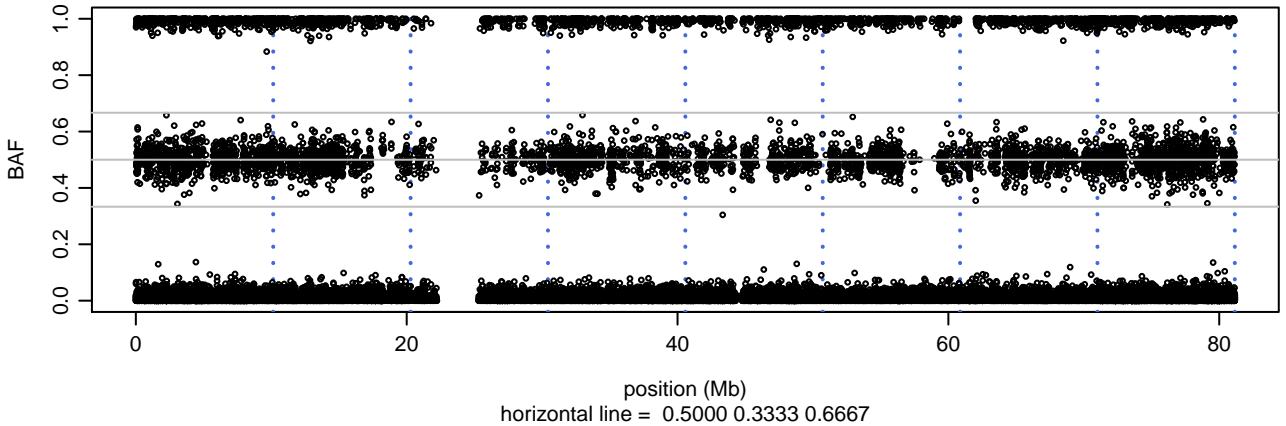
Scan 280 – Chromosome 18 – FUS_R216C_A01

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 q11.1 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
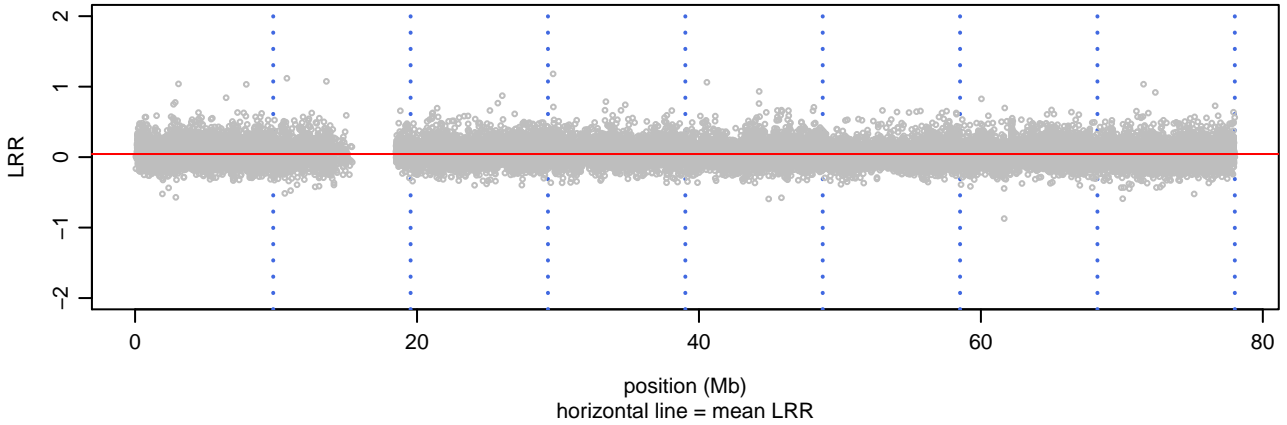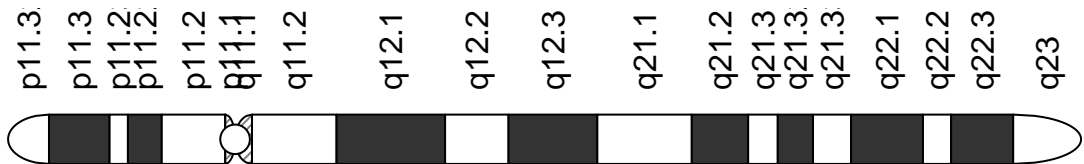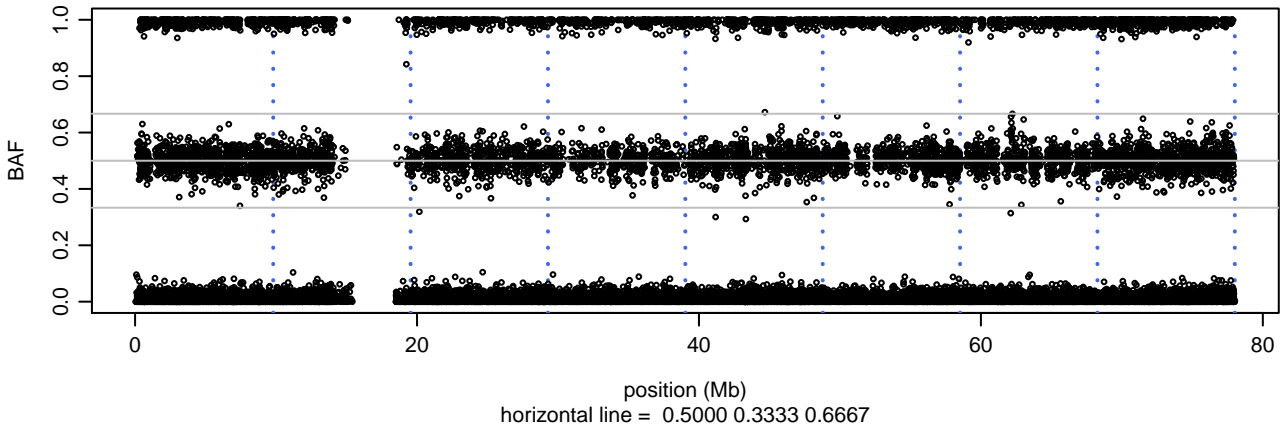
Scan 280 – Chromosome 19 – FUS_R216C_A01

position (Mb)
horizontal line = mean LRR

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

position (Mb)
horizontal line = 0.5000 0.3333 0.6667
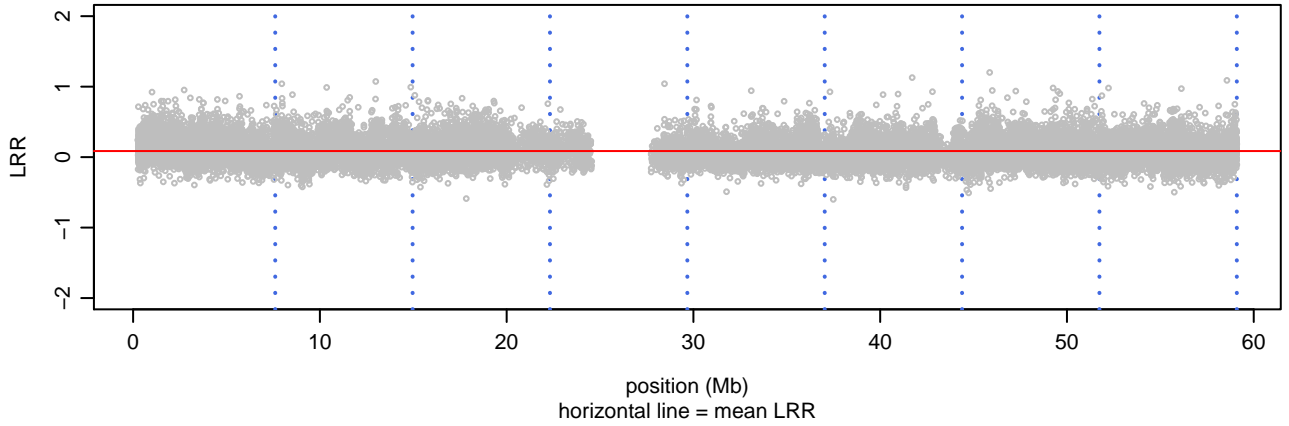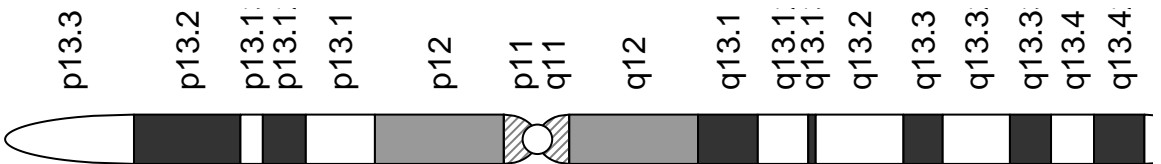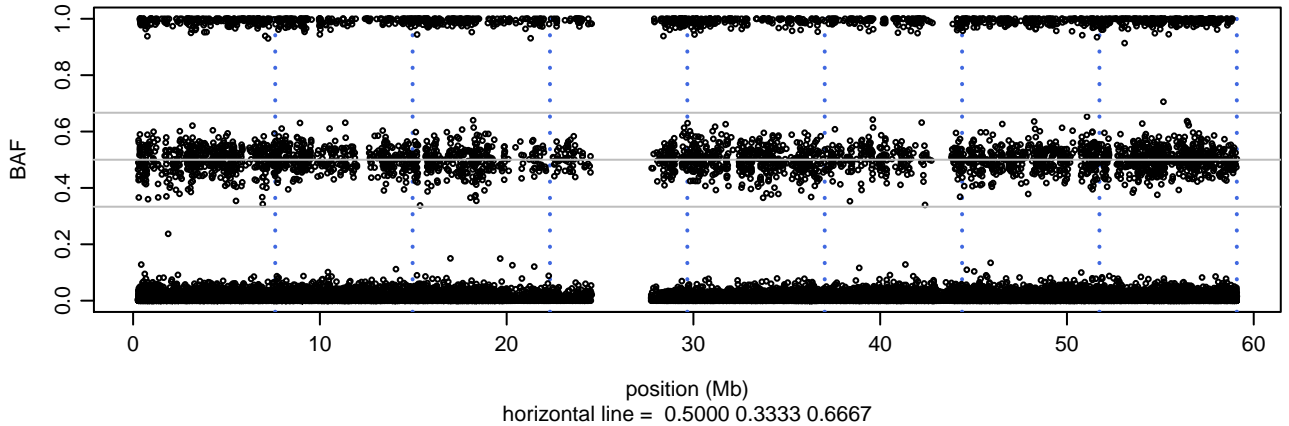
Scan 280 – Chromosome 20 – FUS_R216C_A01

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
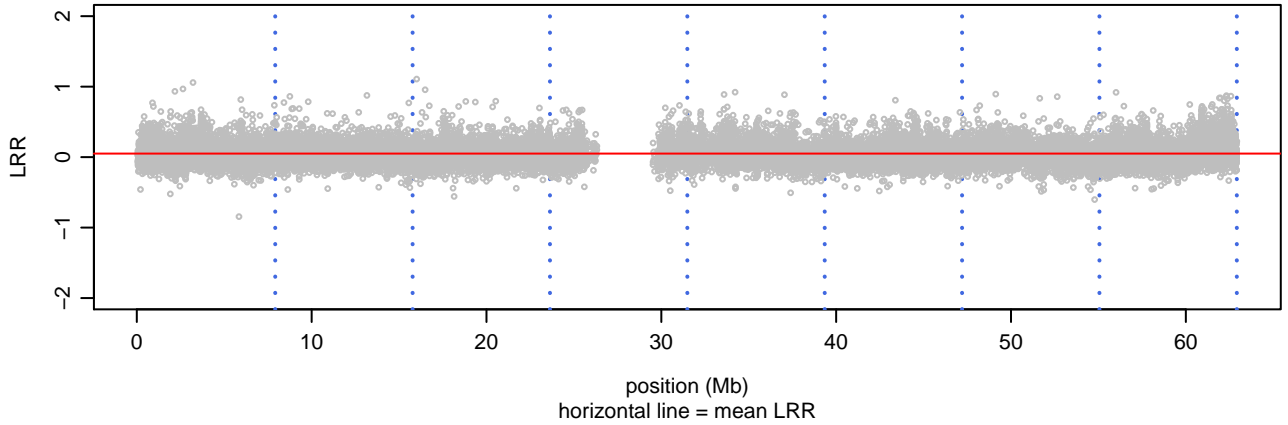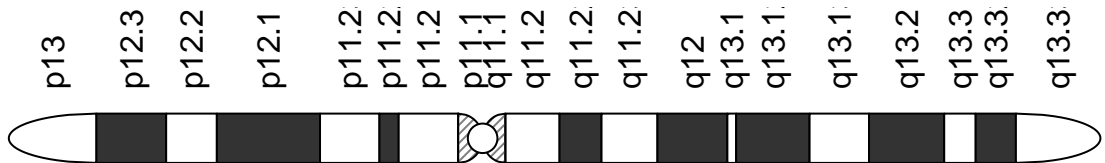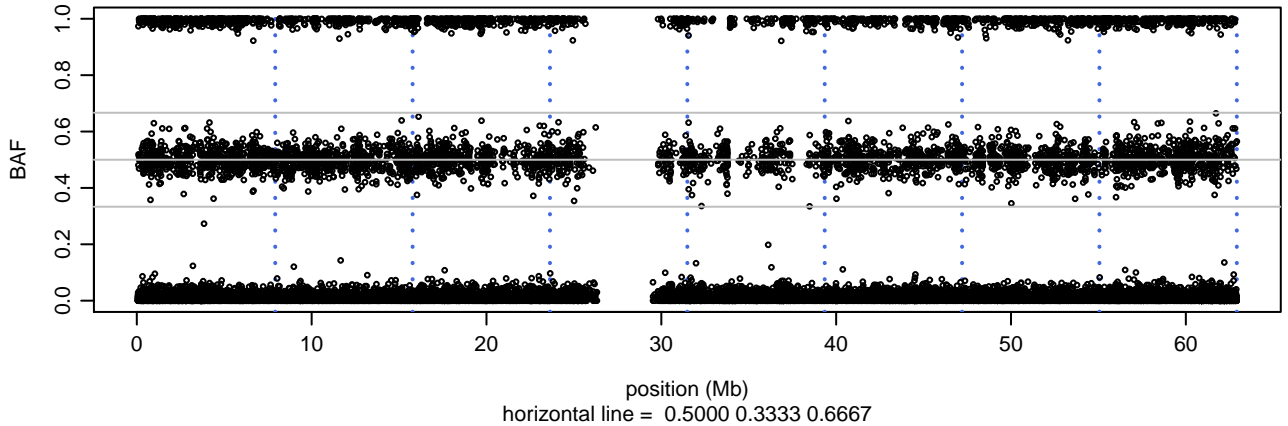
Scan 280 – Chromosome 21 – FUS_R216C_A01
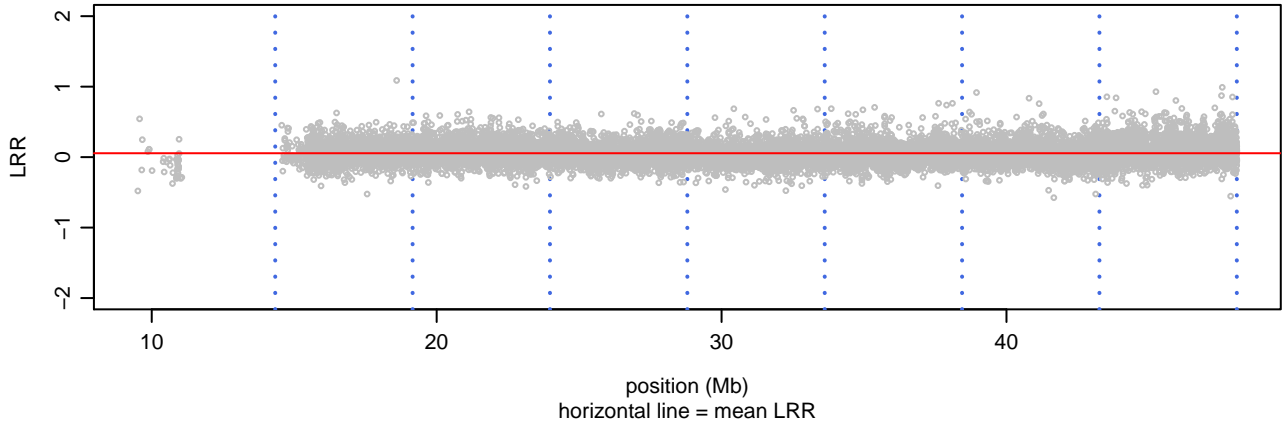
position (Mb)
horizontal line = mean LRR

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

position (Mb)
horizontal line = 0.5000 0.3333 0.6667
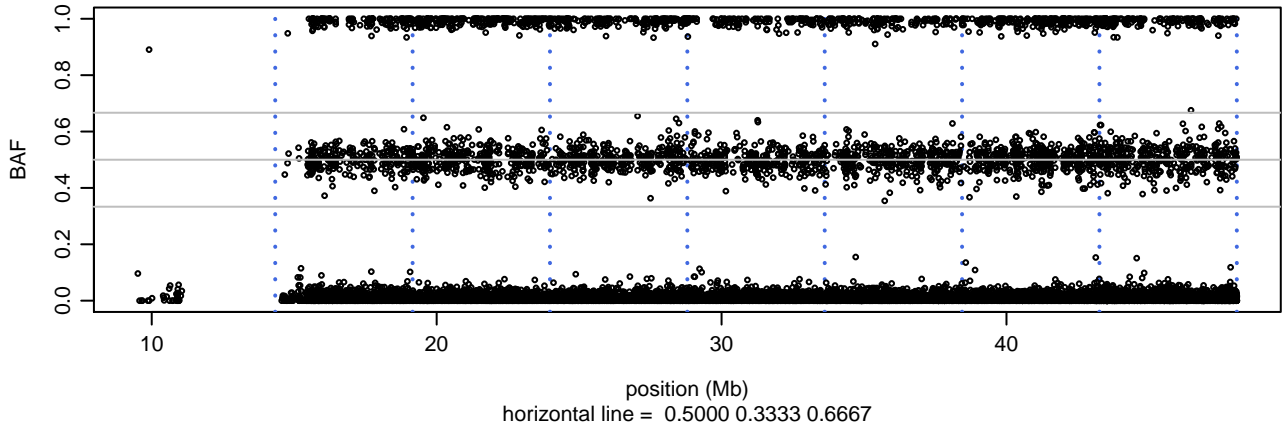
Scan 280 – Chromosome 22 – FUS_R216C_A01
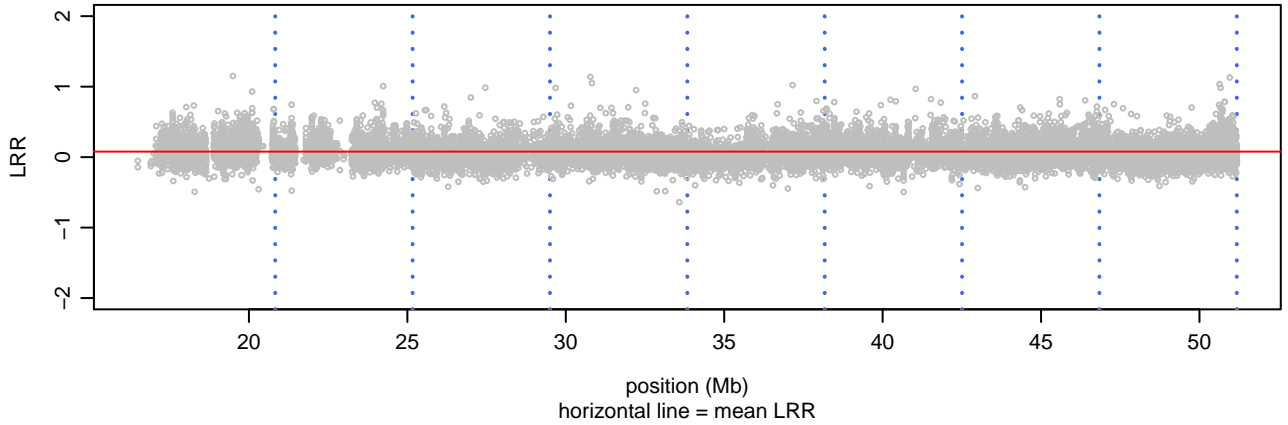
position (Mb)
horizontal line = mean LRR

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

position (Mb)
horizontal line = 0.5000 0.3333 0.6667

Scan 280 – Chromosome X – FUS_R216C_A01

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

p22.3
p22.33
p22.3
p22.2
p22.1
p22.1
p22.1
p21.3
p21.2
p21.1
p11.4
p11.3
p11.22
p11.2
p11.1
p11.1
p11.1
q11.2
q12
q13.1
q13.3
q13.3
q21.1
q21.2
q21.31
q21.31
q21.33
q22.1
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q23
q24
q25
q26.2
q26.3
q27.1
q27.2
q27.3
q28
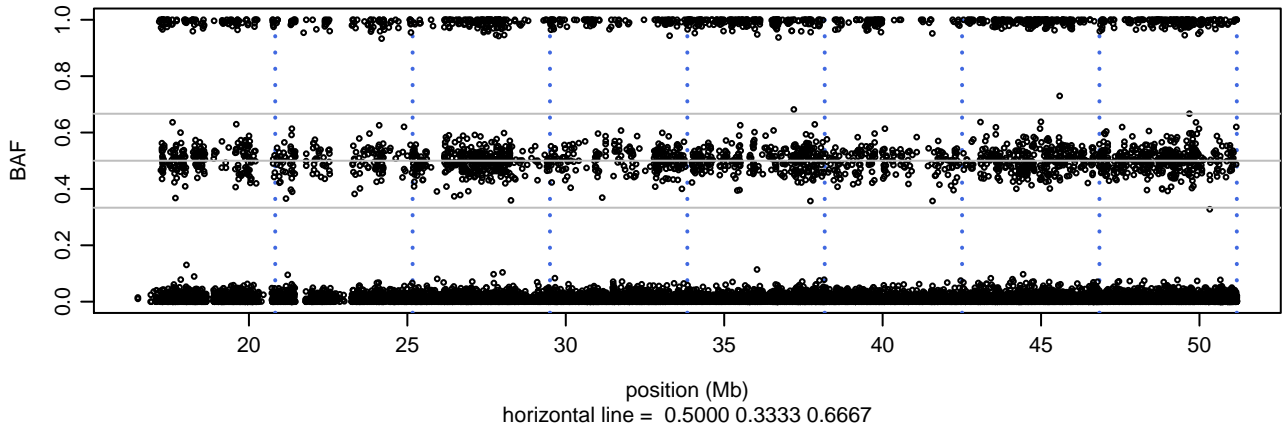
Scan 280 – Chromosome Y – FUS_R216C_A01

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