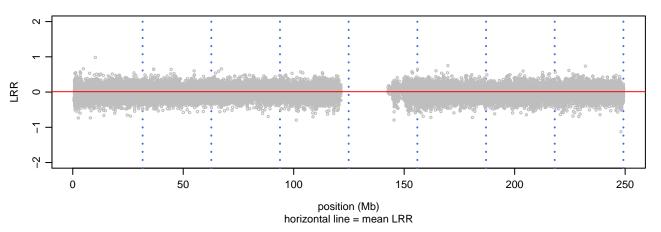
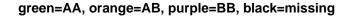
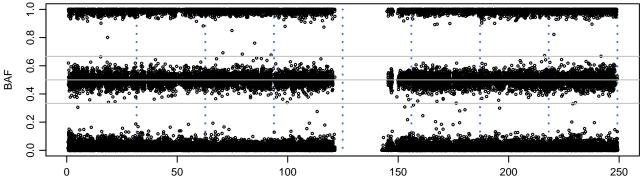
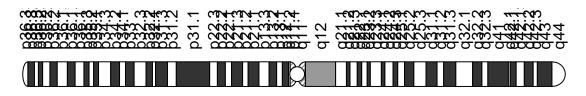
Scan 280 – Chromosome 1 – SNCA_E46K_A02



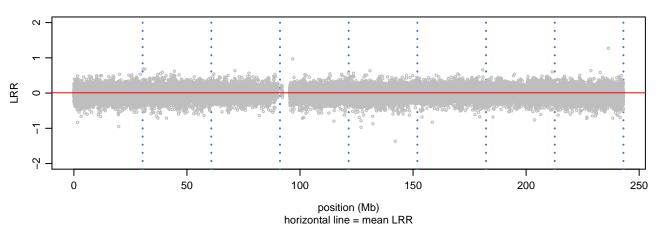




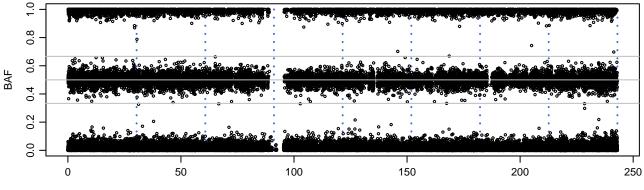
position (Mb) horizontal line = 0.5000 0.3333 0.6667



Scan 280 – Chromosome 2 – SNCA_E46K_A02



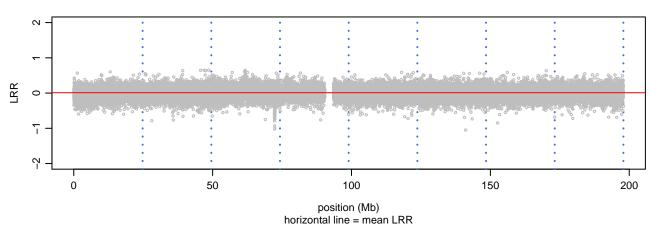
green=AA, orange=AB, purple=BB, black=missing

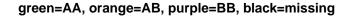


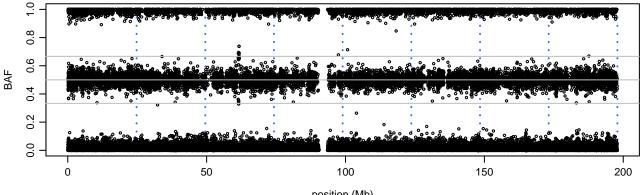
position (Mb) horizontal line = 0.5000 0.3333 0.6667



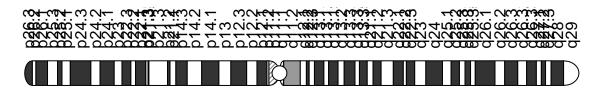
Scan 280 – Chromosome 3 – SNCA_E46K_A02



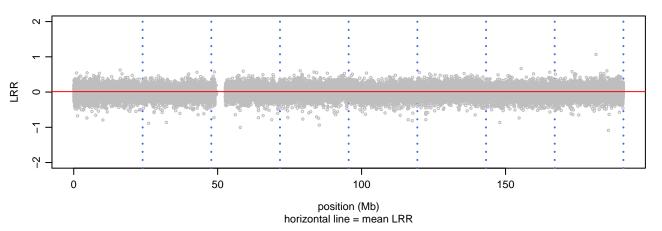


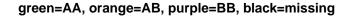


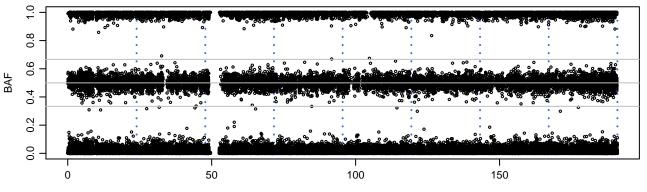
position (Mb) horizontal line = 0.5000 0.3333 0.6667



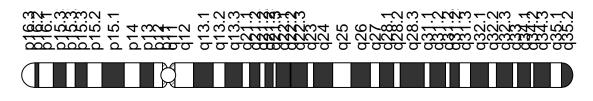
Scan 280 – Chromosome 4 – SNCA_E46K_A02



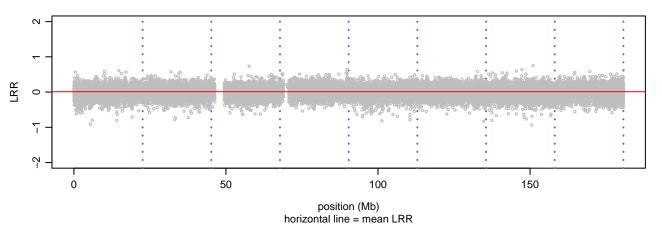


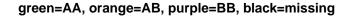


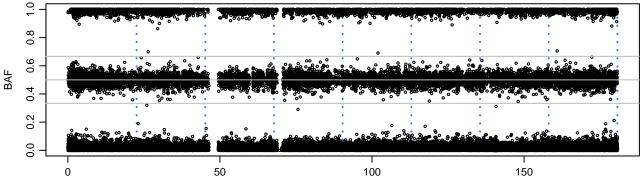
position (Mb) horizontal line = 0.5000 0.3333 0.6667



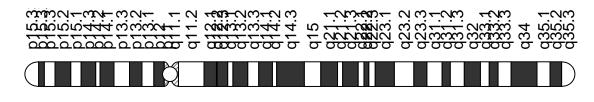
Scan 280 – Chromosome 5 – SNCA_E46K_A02



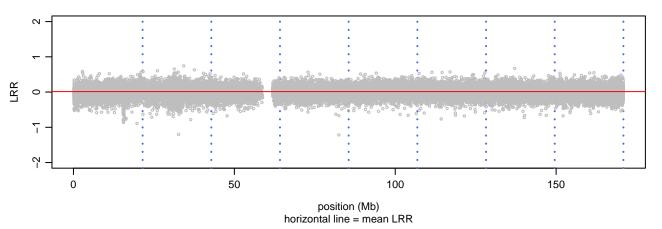




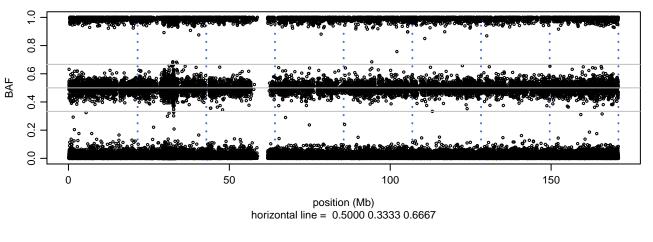
position (Mb) horizontal line = 0.5000 0.3333 0.6667

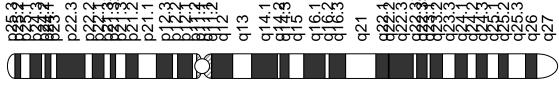


Scan 280 – Chromosome 6 – SNCA_E46K_A02

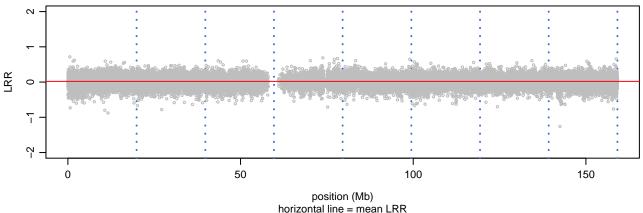


green=AA, orange=AB, purple=BB, black=missing

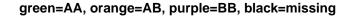


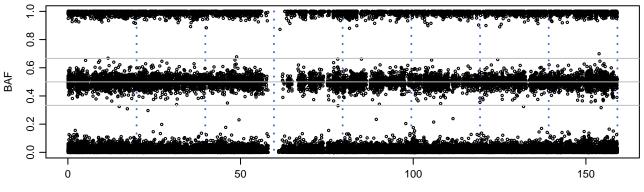


Scan 280 – Chromosome 7 – SNCA_E46K_A02

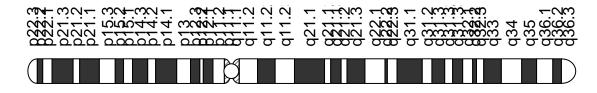


nonzoniai ine = mean LRR

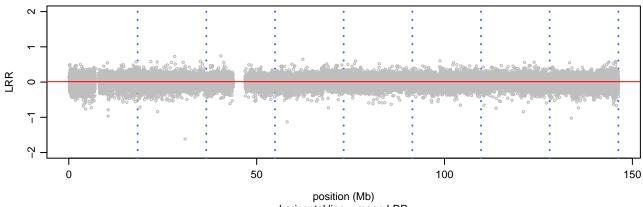




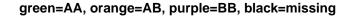
position (Mb) horizontal line = 0.5000 0.3333 0.6667

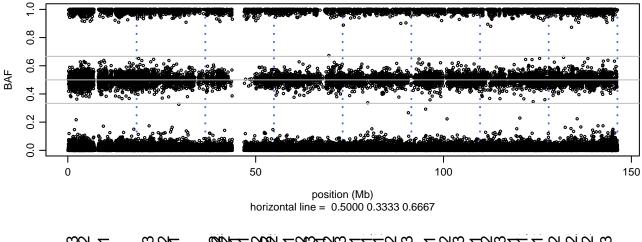


Scan 280 – Chromosome 8 – SNCA_E46K_A02



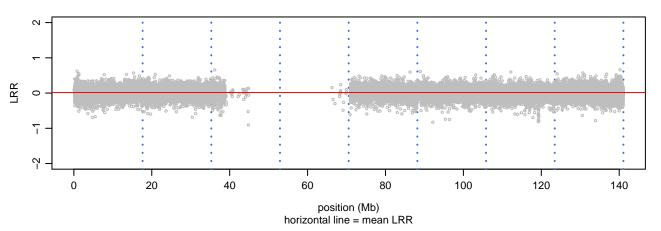
horizontal line = mean LRR

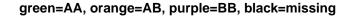


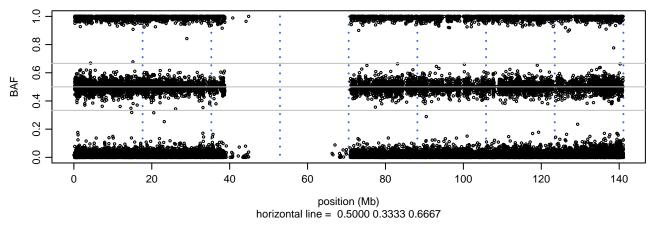


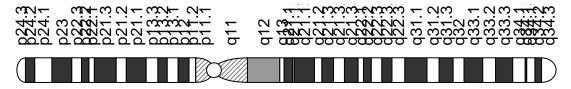


Scan 280 – Chromosome 9 – SNCA_E46K_A02

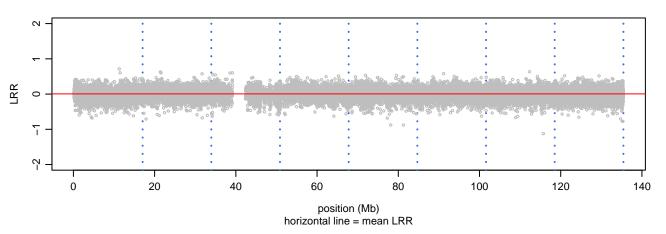


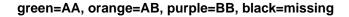


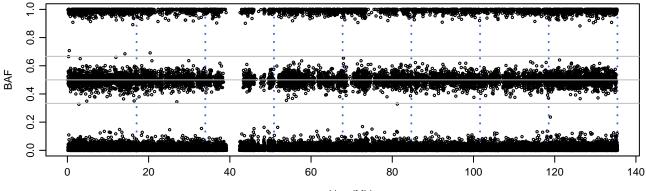




Scan 280 – Chromosome 10 – SNCA_E46K_A02

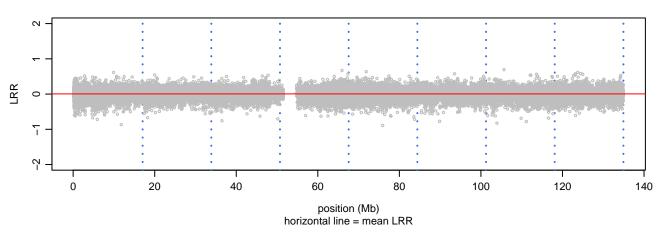




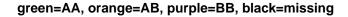


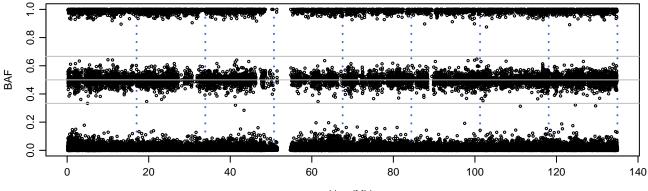
position (Mb) horizontal line = 0.5000 0.3333 0.6667



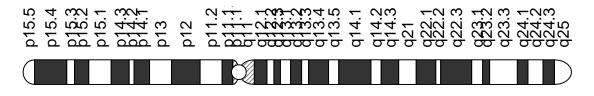


Scan 280 – Chromosome 11 – SNCA_E46K_A02

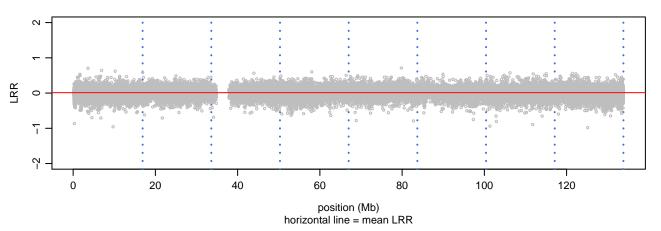


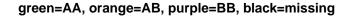


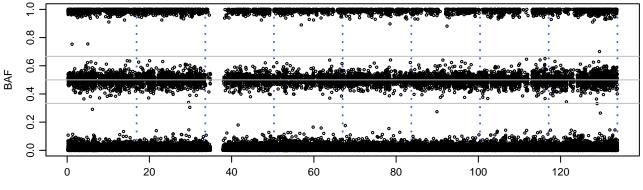
position (Mb) horizontal line = 0.5000 0.3333 0.6667



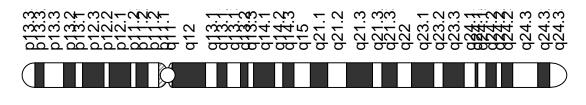
Scan 280 – Chromosome 12 – SNCA_E46K_A02

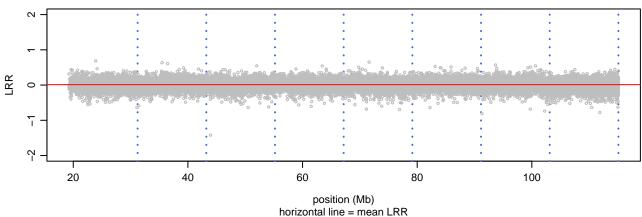


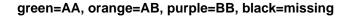


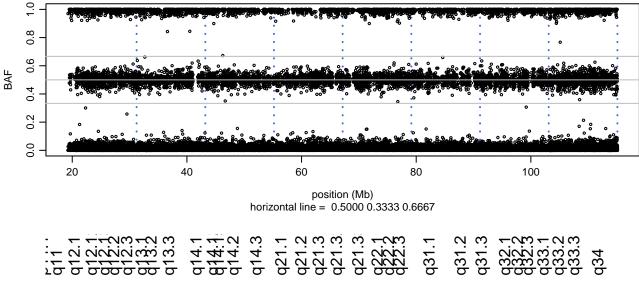


position (Mb) horizontal line = 0.5000 0.3333 0.6667

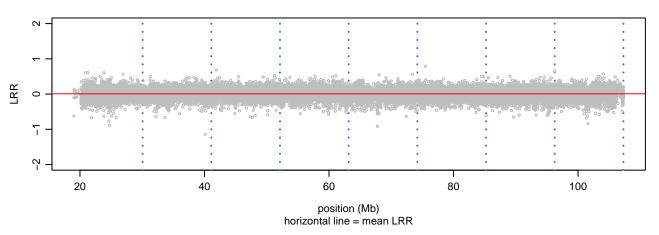


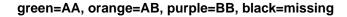


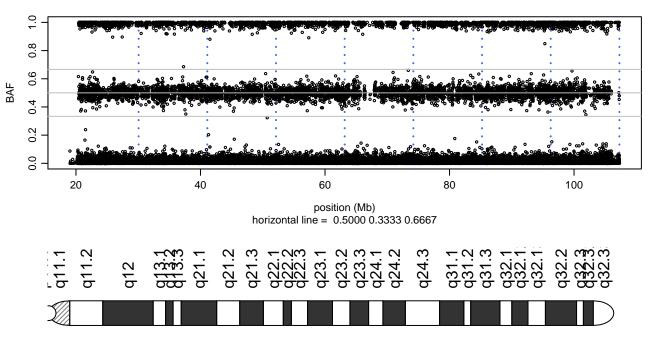




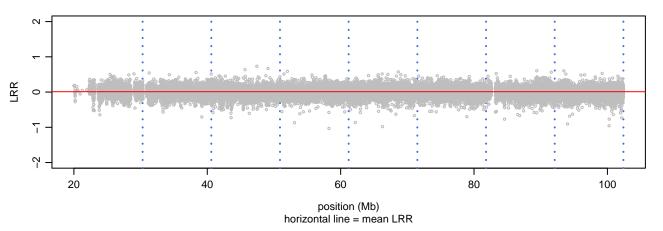


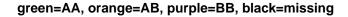


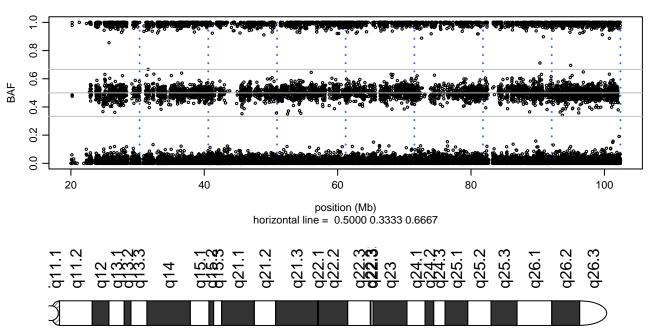




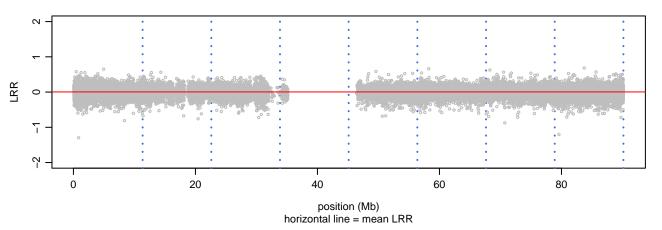
Scan 280 – Chromosome 15 – SNCA_E46K_A02



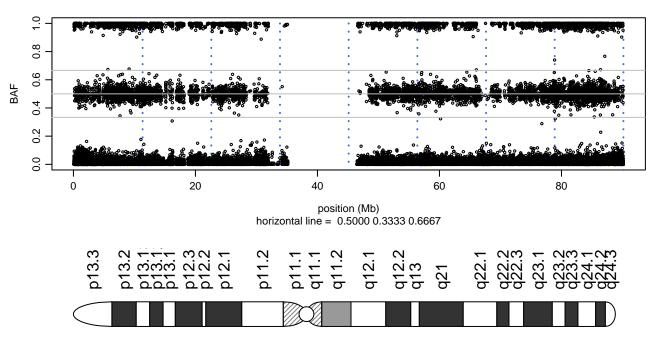


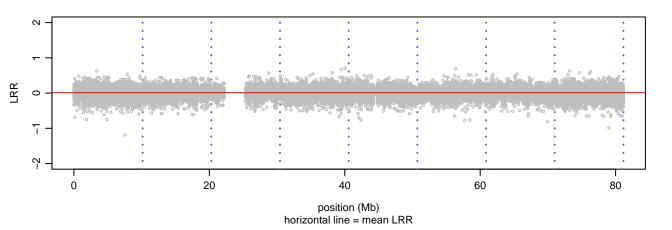


Scan 280 – Chromosome 16 – SNCA_E46K_A02

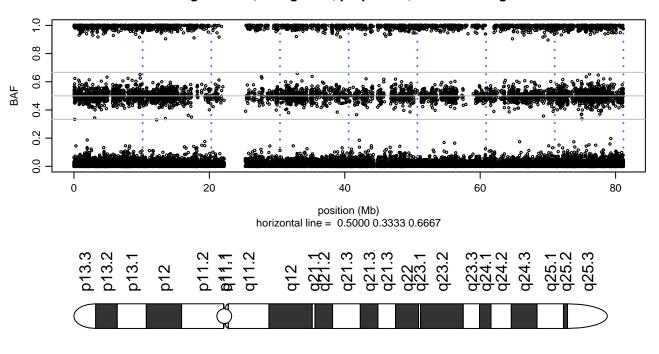


green=AA, orange=AB, purple=BB, black=missing

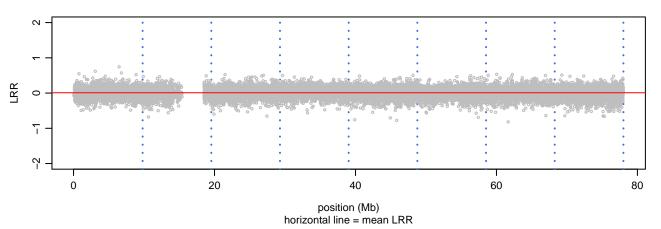




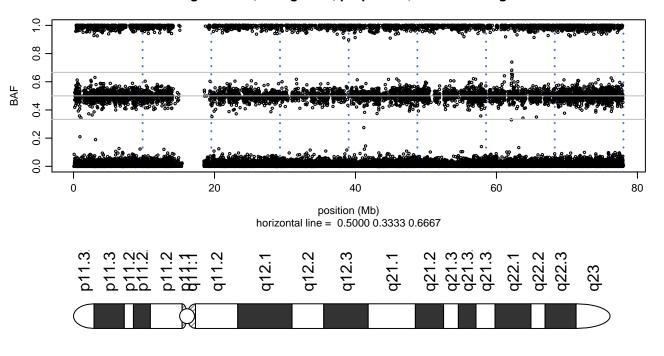
Scan 280 – Chromosome 17 – SNCA_E46K_A02

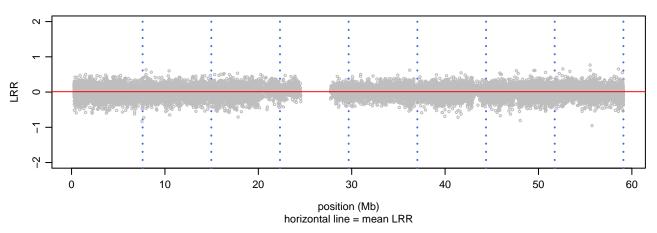


Scan 280 – Chromosome 18 – SNCA_E46K_A02

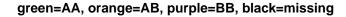


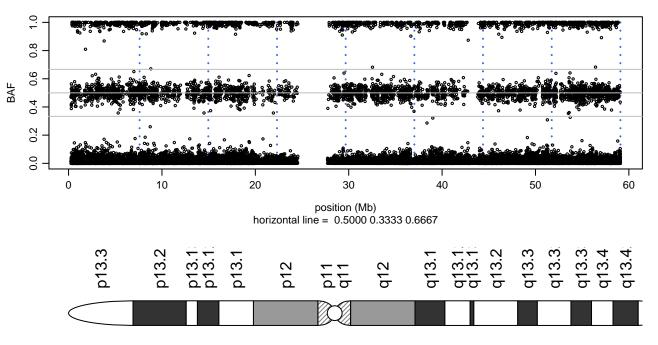
green=AA, orange=AB, purple=BB, black=missing

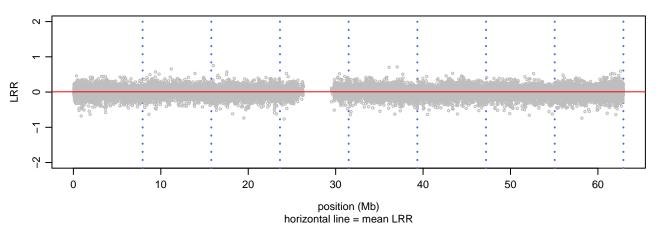




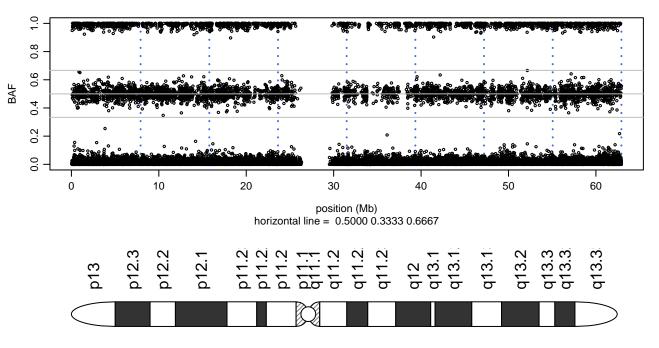
Scan 280 – Chromosome 19 – SNCA_E46K_A02

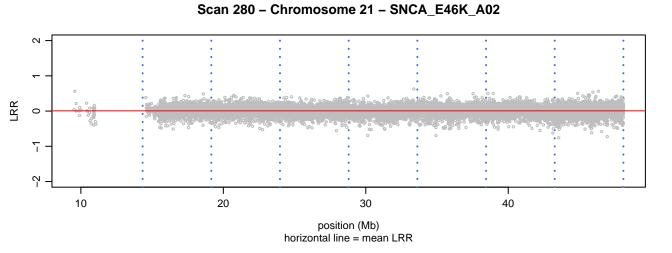


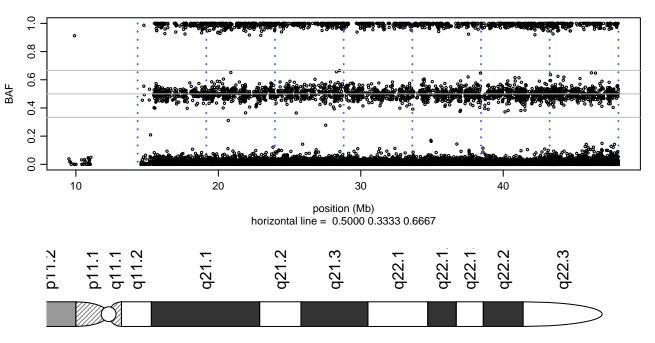


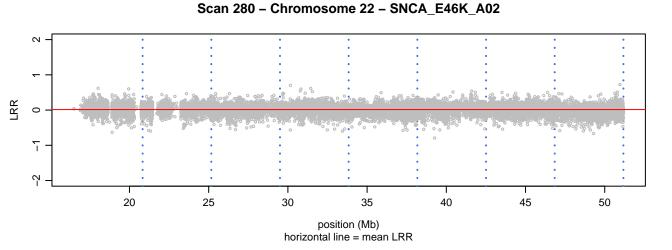


Scan 280 – Chromosome 20 – SNCA_E46K_A02

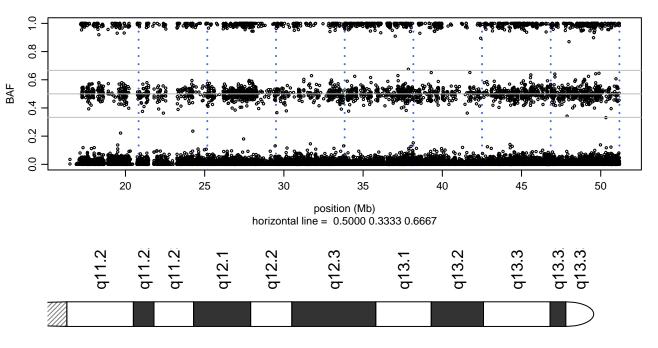




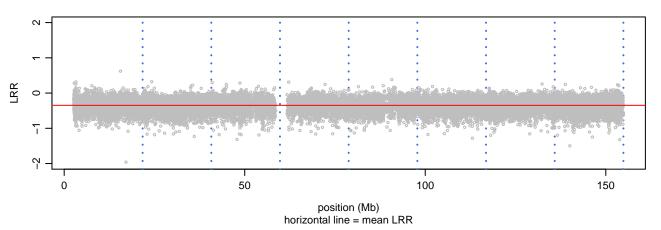


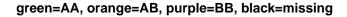


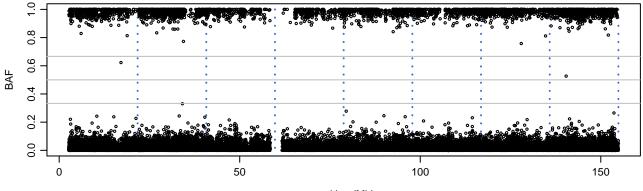
green=AA, orange=AB, purple=BB, black=missing



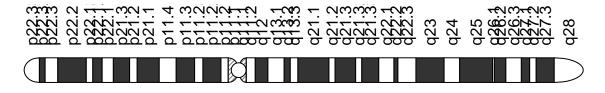
Scan 280 – Chromosome X – SNCA_E46K_A02

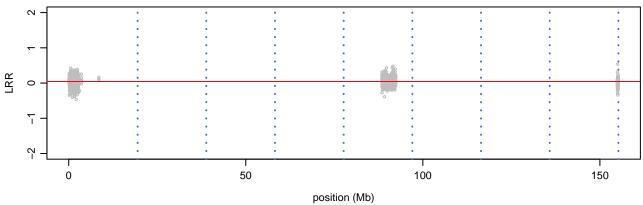






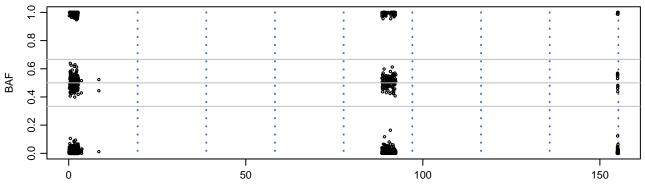
position (Mb) horizontal line = 0.5000 0.3333 0.6667





Scan 280 – Chromosome Y – SNCA_E46K_A02

horizontal line = mean LRR



position (Mb) horizontal line = 0.5000 0.3333 0.6667

