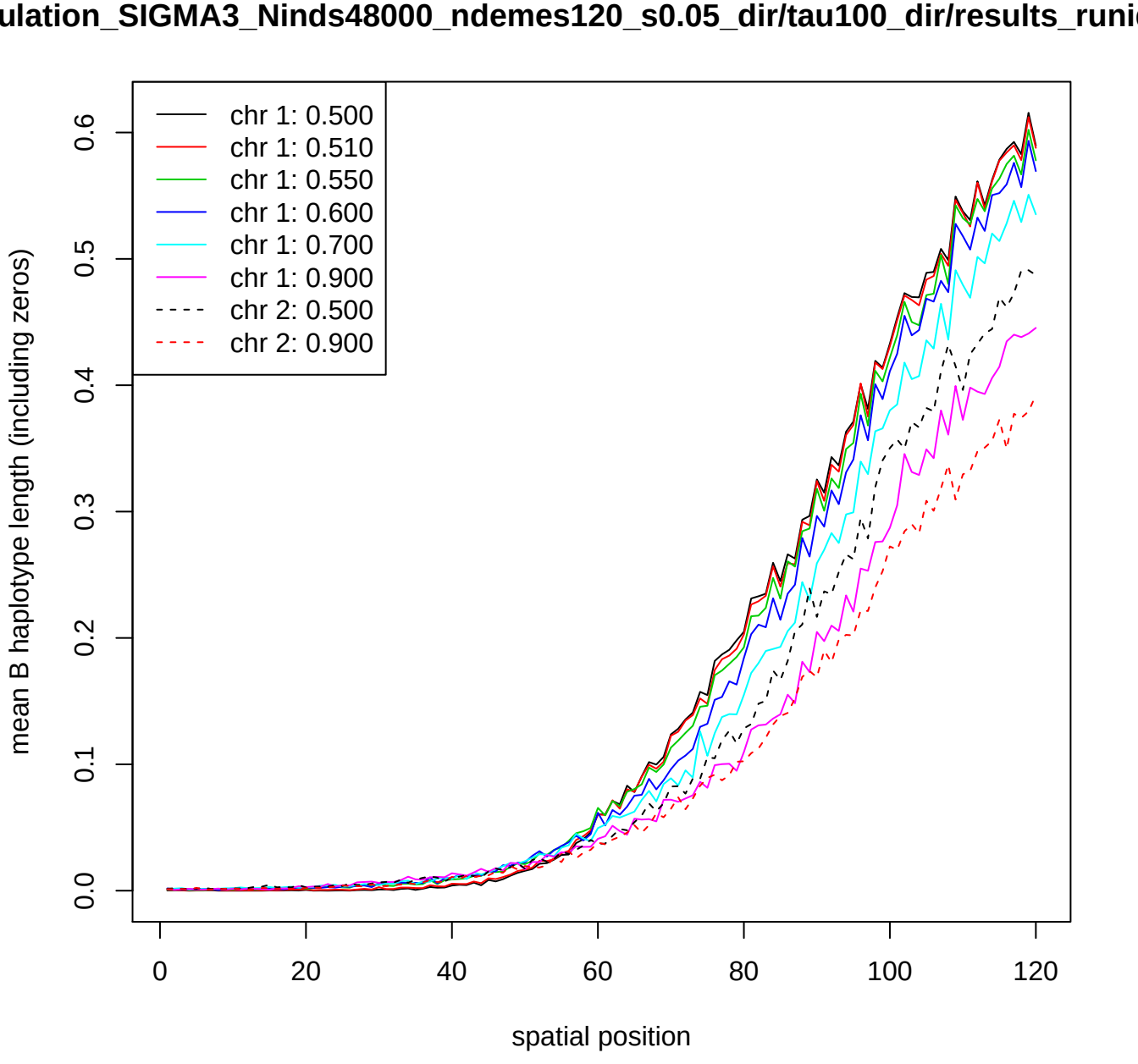
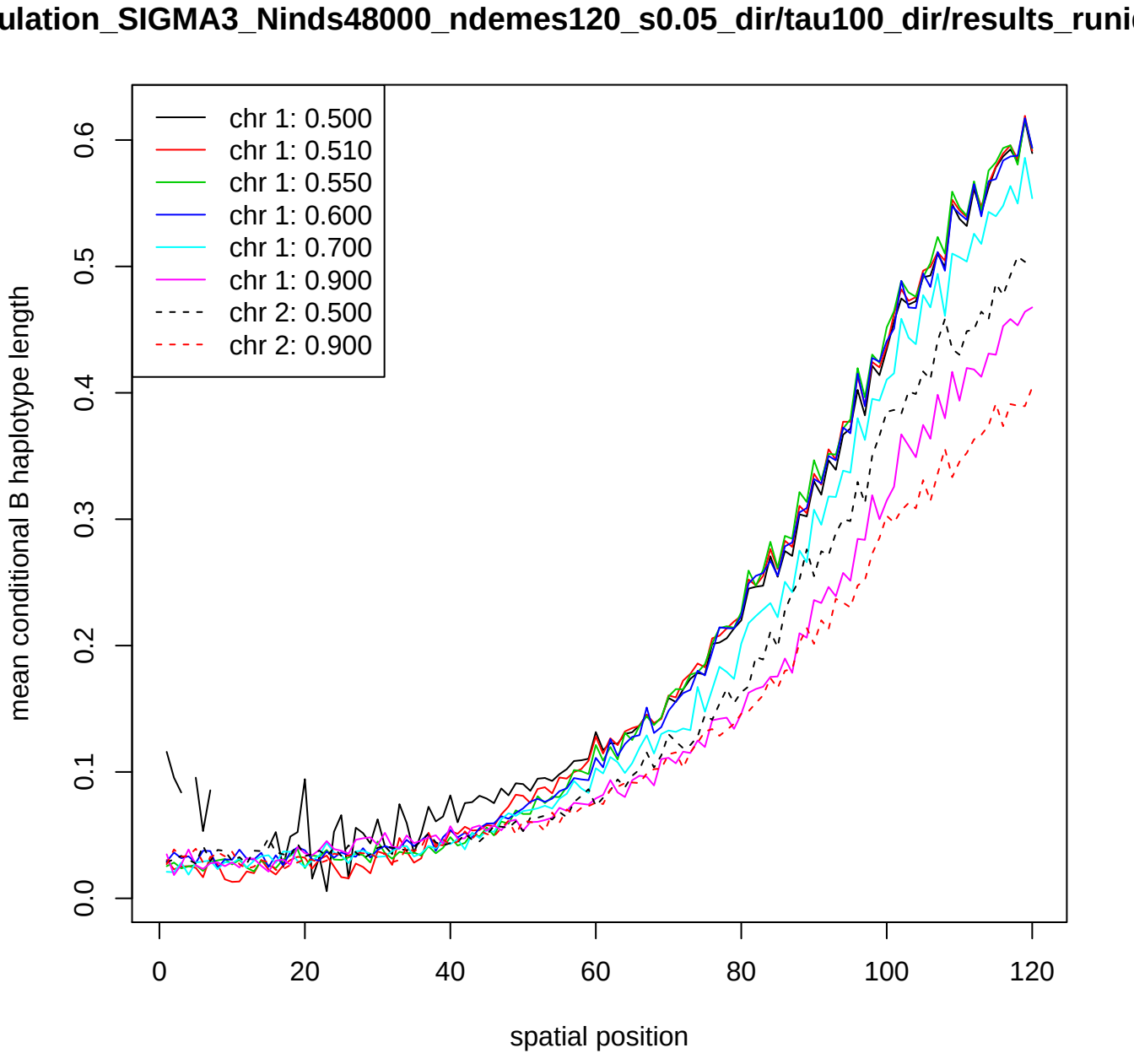
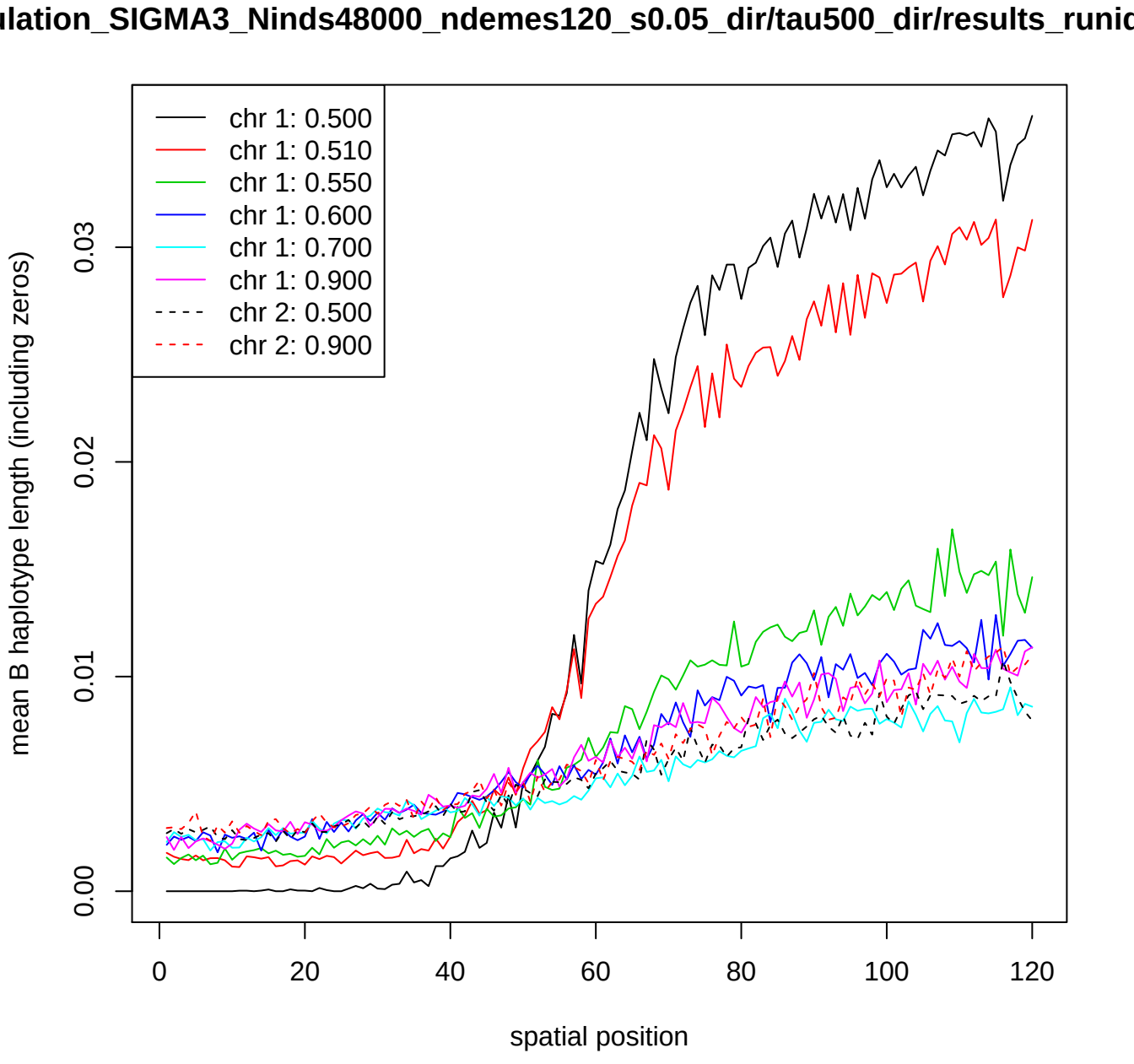
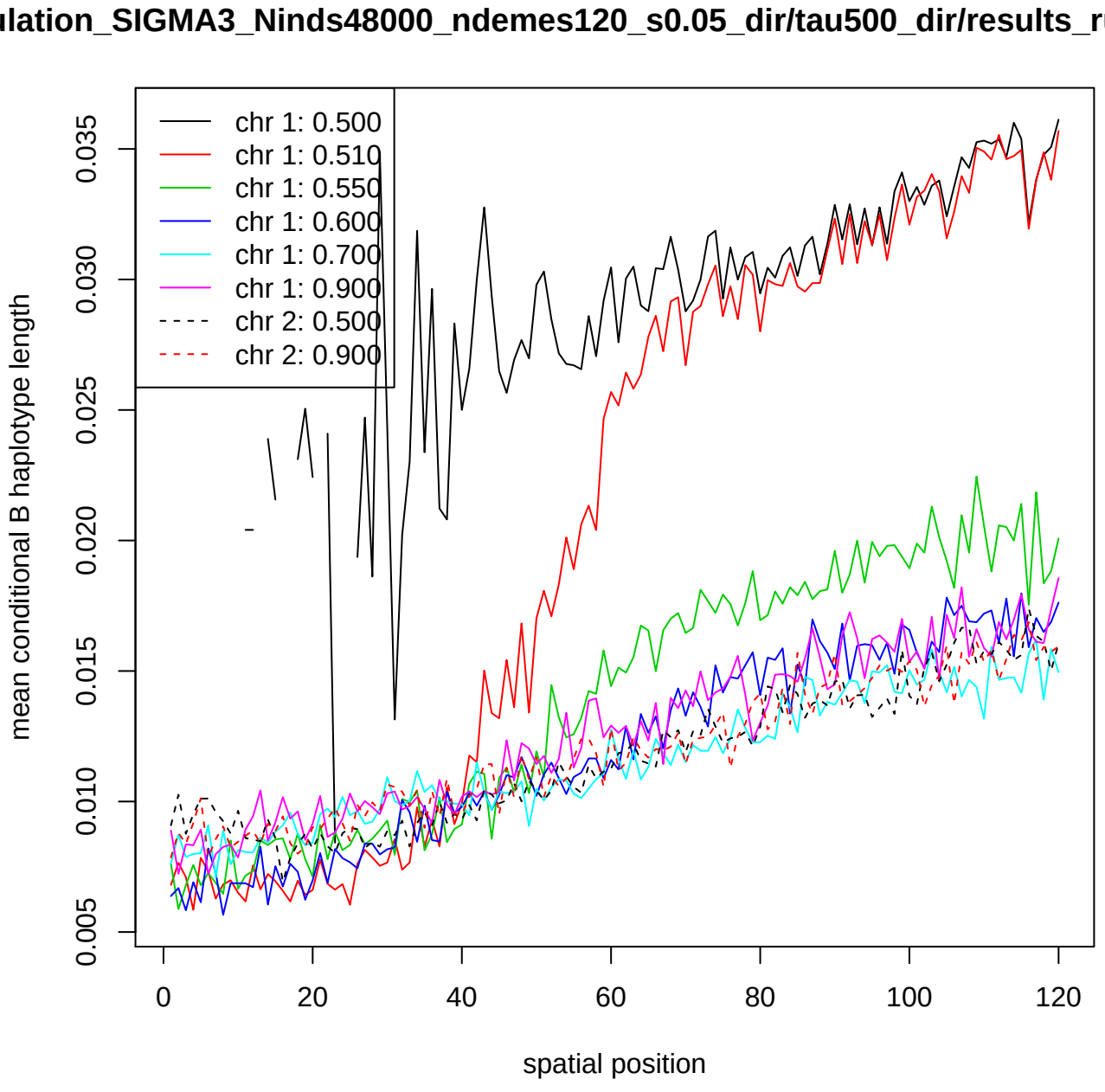
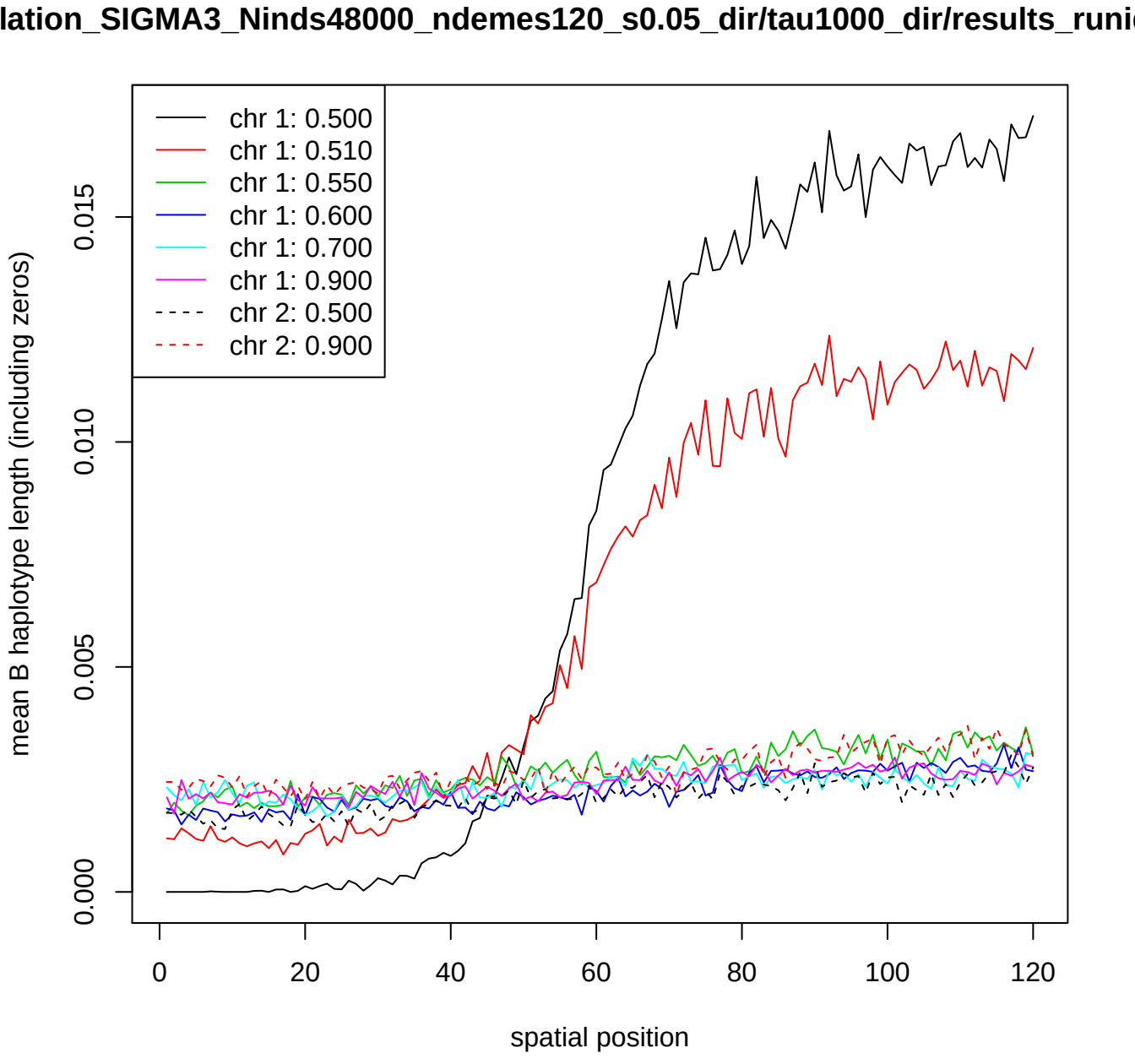
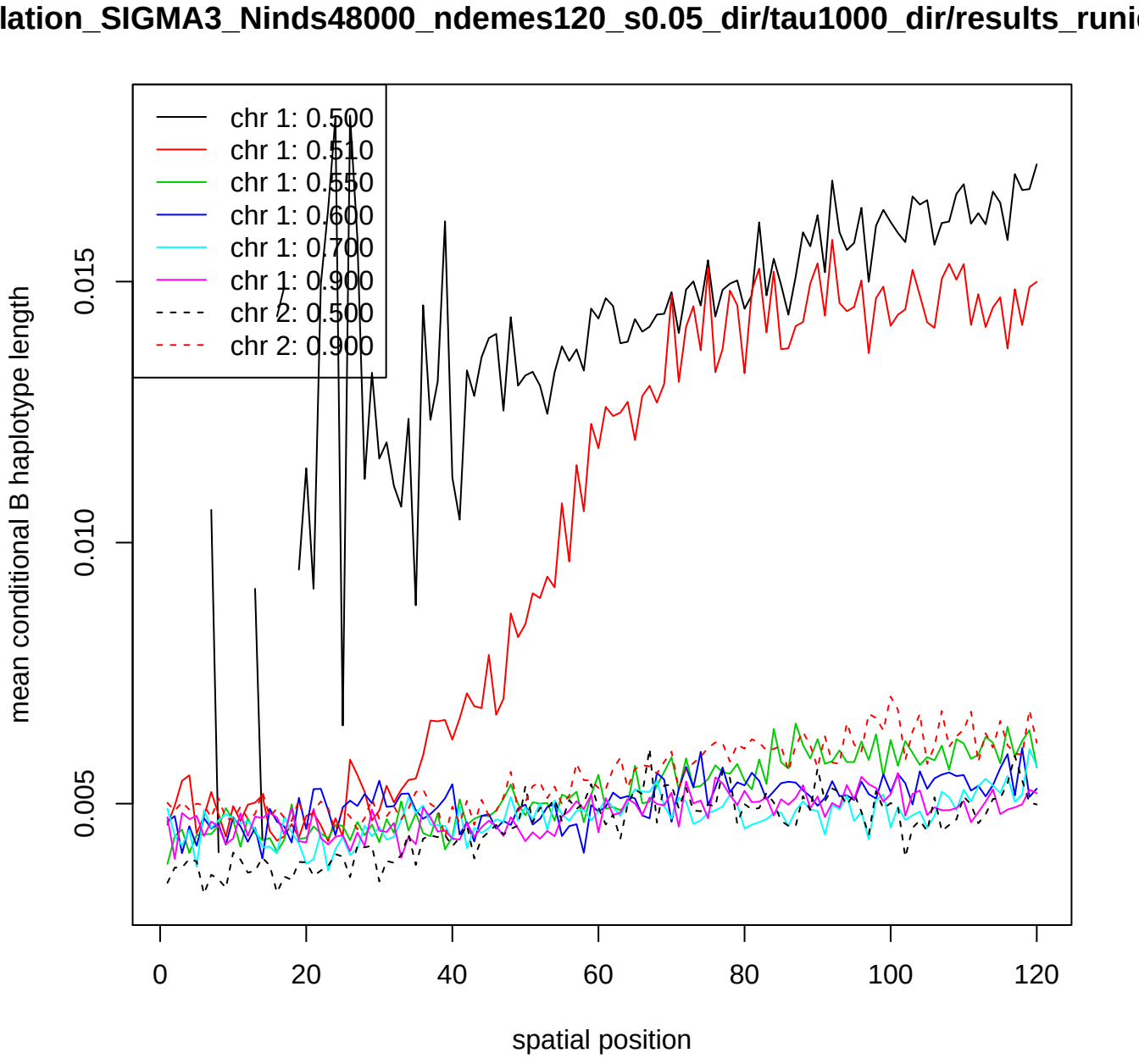

ulation_SIGMA3_Ninds48000_ndemes120_s0.05_dir/tau100_dir/results_runi
chr 1: 0.500
chr 1: 0.510
chr 1: 0.550
chr 1: 0.600
chr 1: 0.700
chr 1: 0.900
chr 2: 0.500
chr 2: 0.900
mean B haplotype length (including zeros)
0.0
0.1
0.2
0.3
0.4
0.5
0.6
0
20
40
60
80
100
120
spatial position


ulation_SIGMA3_Ninds48000_ndemes120_s0.05_dir/tau100_dir/results_runi
chr 1: 0.500
chr 1: 0.510
chr 1: 0.550
chr 1: 0.600
chr 1: 0.700
chr 1: 0.900
chr 2: 0.500
chr 2: 0.900
mean conditional B haplotype length
0.0
0.1
0.2
0.3
0.4
0.5
0.6
0
20
40
60
80
100
120
spatial position


ulation_SIGMA3_Ninds48000_ndemes120_s0.05_dir/tau500_dir/results_runid
chr 1: 0.500
chr 1: 0.510
chr 1: 0.550
chr 1: 0.600
chr 1: 0.700
chr 1: 0.900
chr 2: 0.500
chr 2: 0.900
mean B haplotype length (including zeros)
0.00
0.01
0.02
0.03
0
20
40
60
80
100
120
spatial position


ulation_SIGMA3_Ninds48000_ndemes120_s0.05_dir/tau500_dir/results_runid
chr 1: 0.500
chr 1: 0.510
chr 1: 0.550
chr 1: 0.600
chr 1: 0.700
chr 1: 0.900
chr 2: 0.500
chr 2: 0.900
mean conditional B haplotype length
0.005
0.010
0.015
0.020
0.025
0.030
0.035
0
20
40
60
80
100
120
spatial position


lation_SIGMA3_Ninds48000_ndemes120_s0.05_dir/tau1000_dir/results_runi
chr 1: 0.500
chr 1: 0.510
chr 1: 0.550
chr 1: 0.600
chr 1: 0.700
chr 1: 0.900
chr 2: 0.500
chr 2: 0.900
mean B haplotype length (including zeros)
0.000
0.005
0.010
0.015
0
20
40
60
80
100
120
spatial position


lation_SIGMA3_Ninds48000_ndemes120_s0.05_dir/tau1000_dir/results_runi
chr 1: 0.500
chr 1: 0.510
chr 1: 0.550
chr 1: 0.600
chr 1: 0.700
chr 1: 0.900
chr 2: 0.500
chr 2: 0.900
mean conditional B haplotype length
0.005
0.010
0.015
0
20
40
60
80
100
120
spatial position