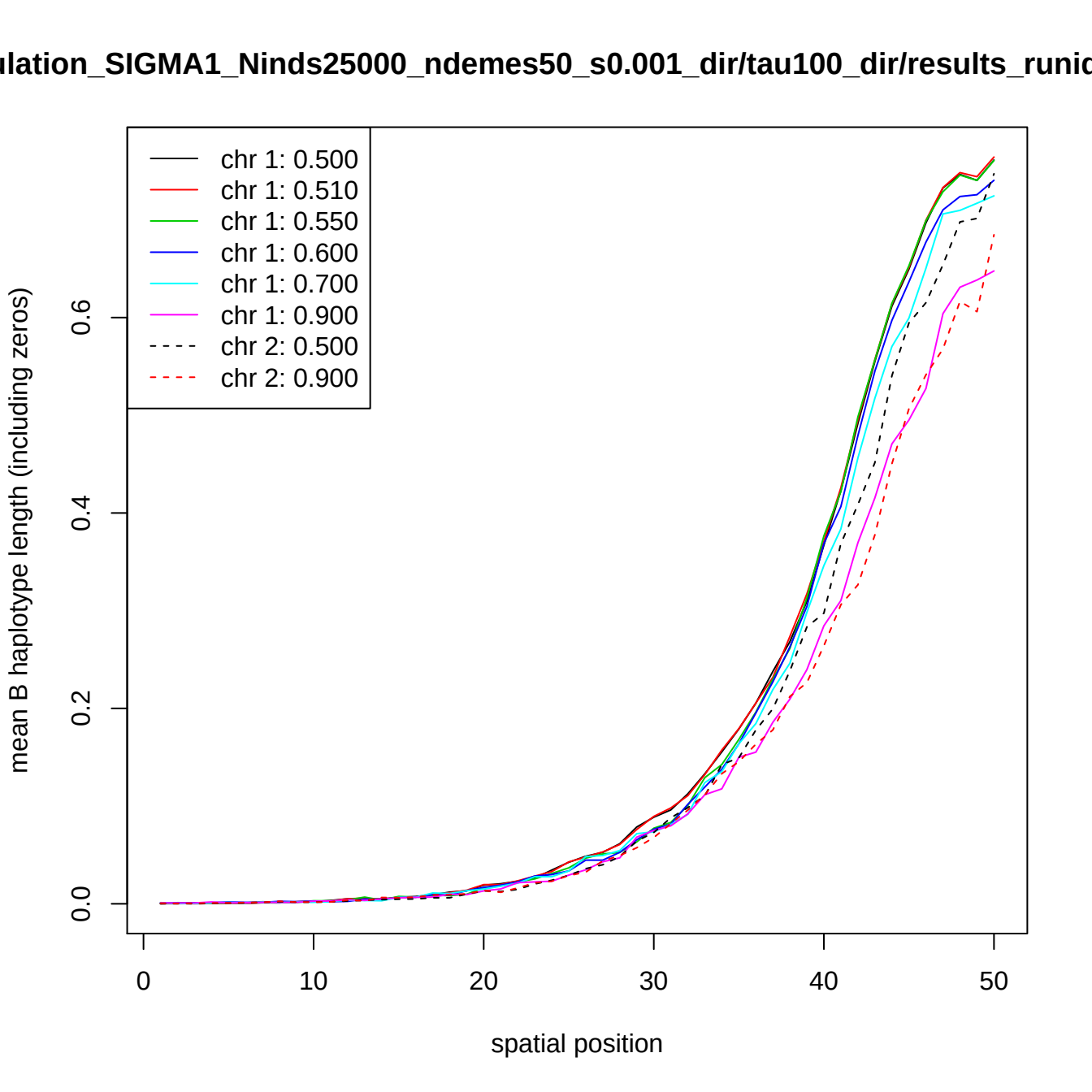
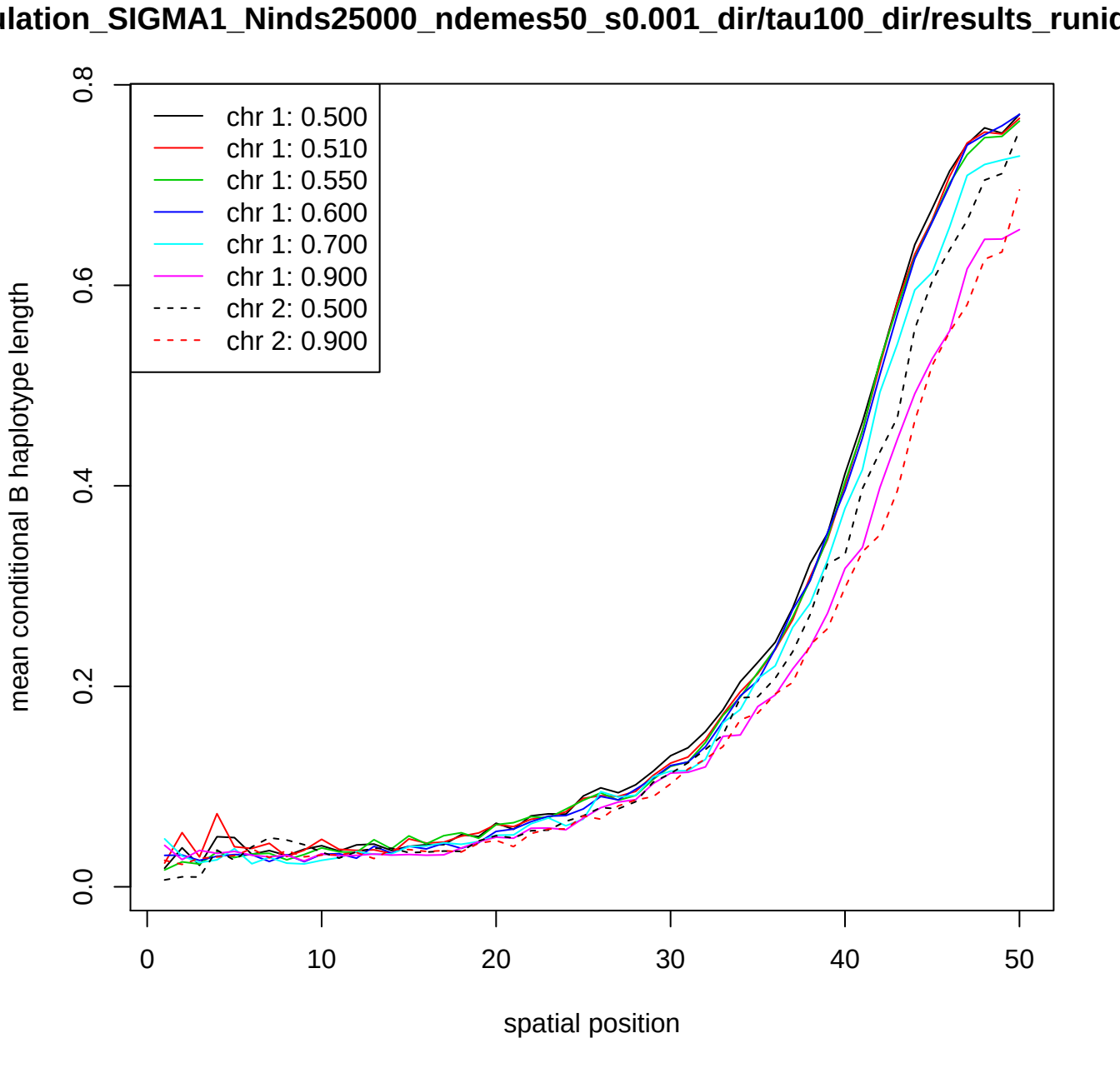
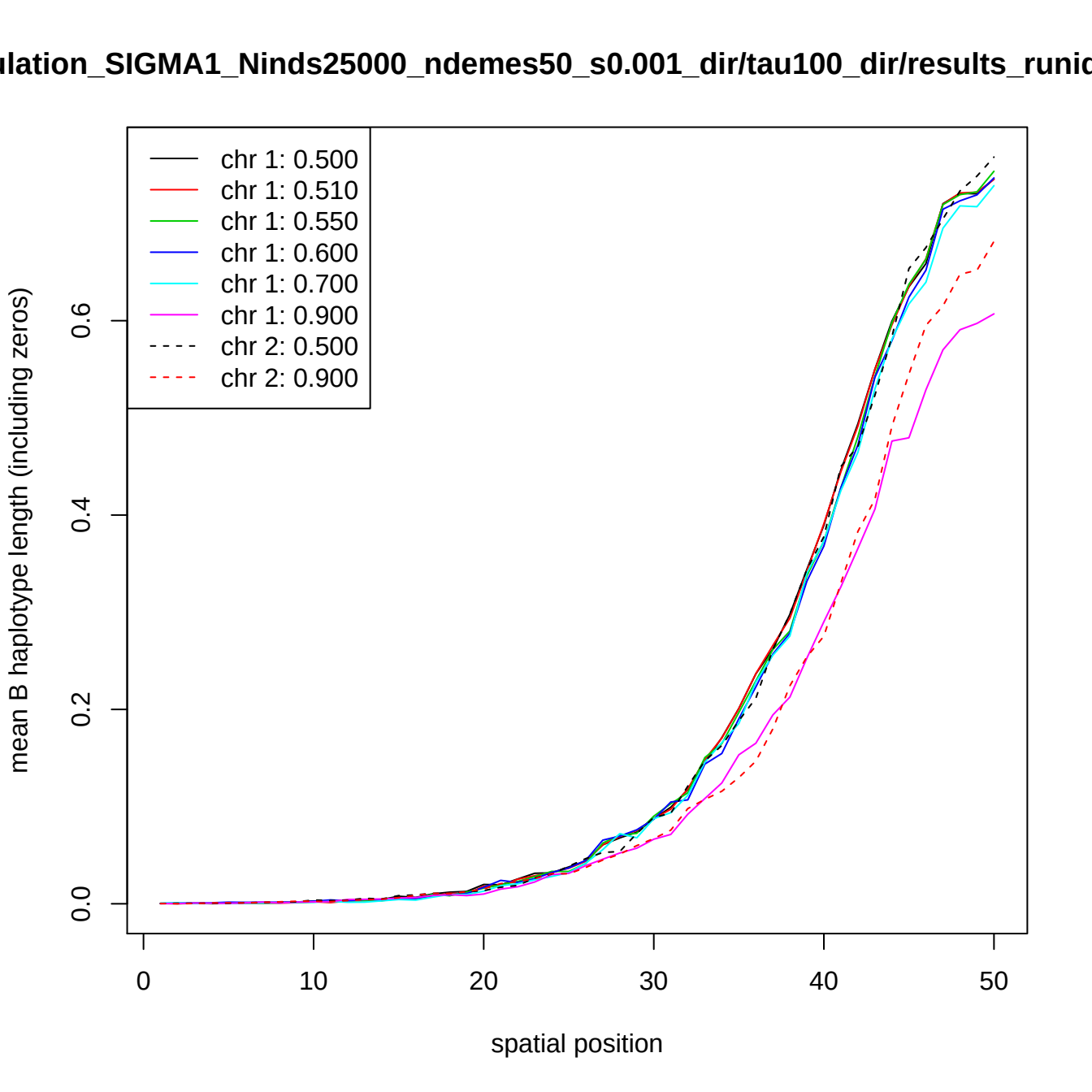
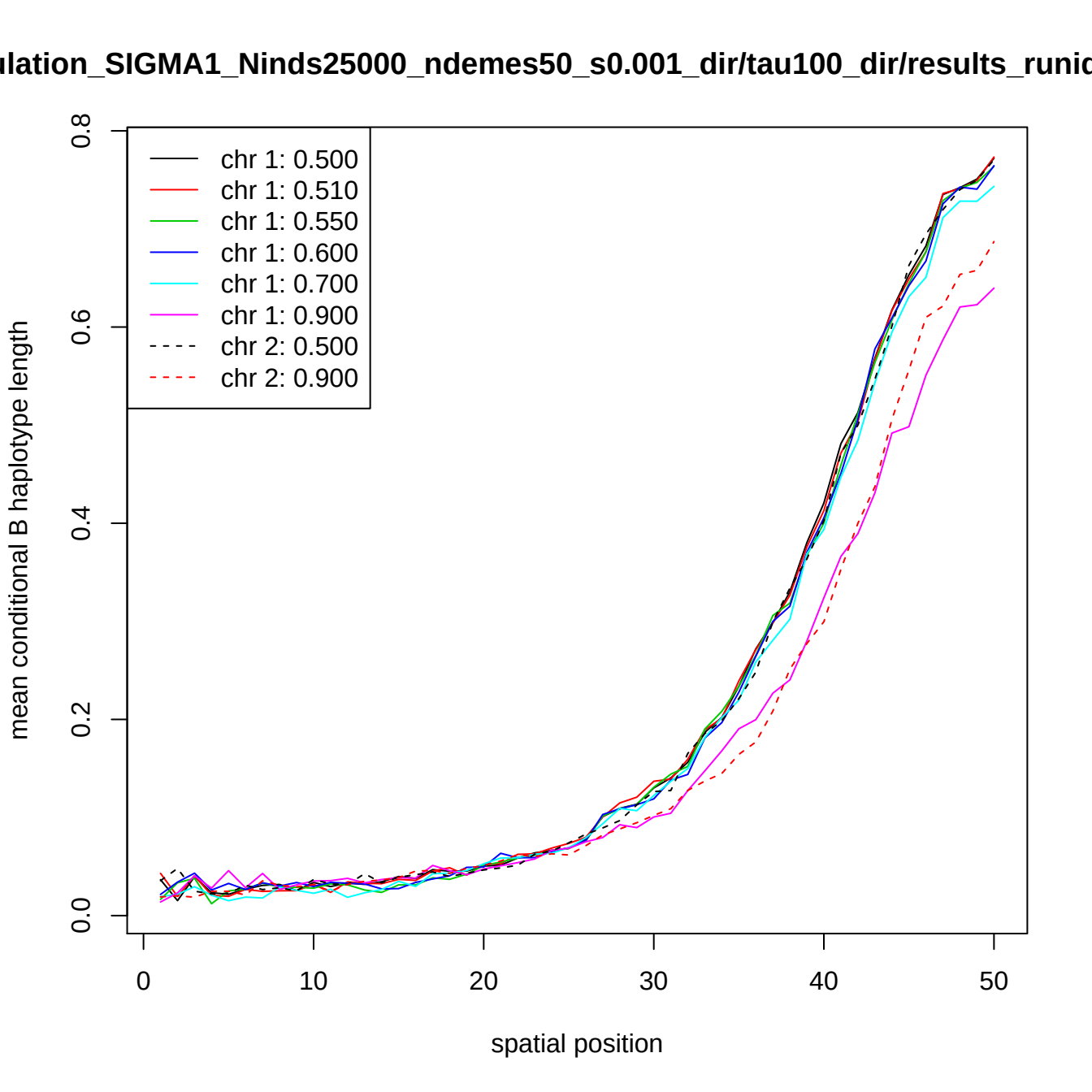
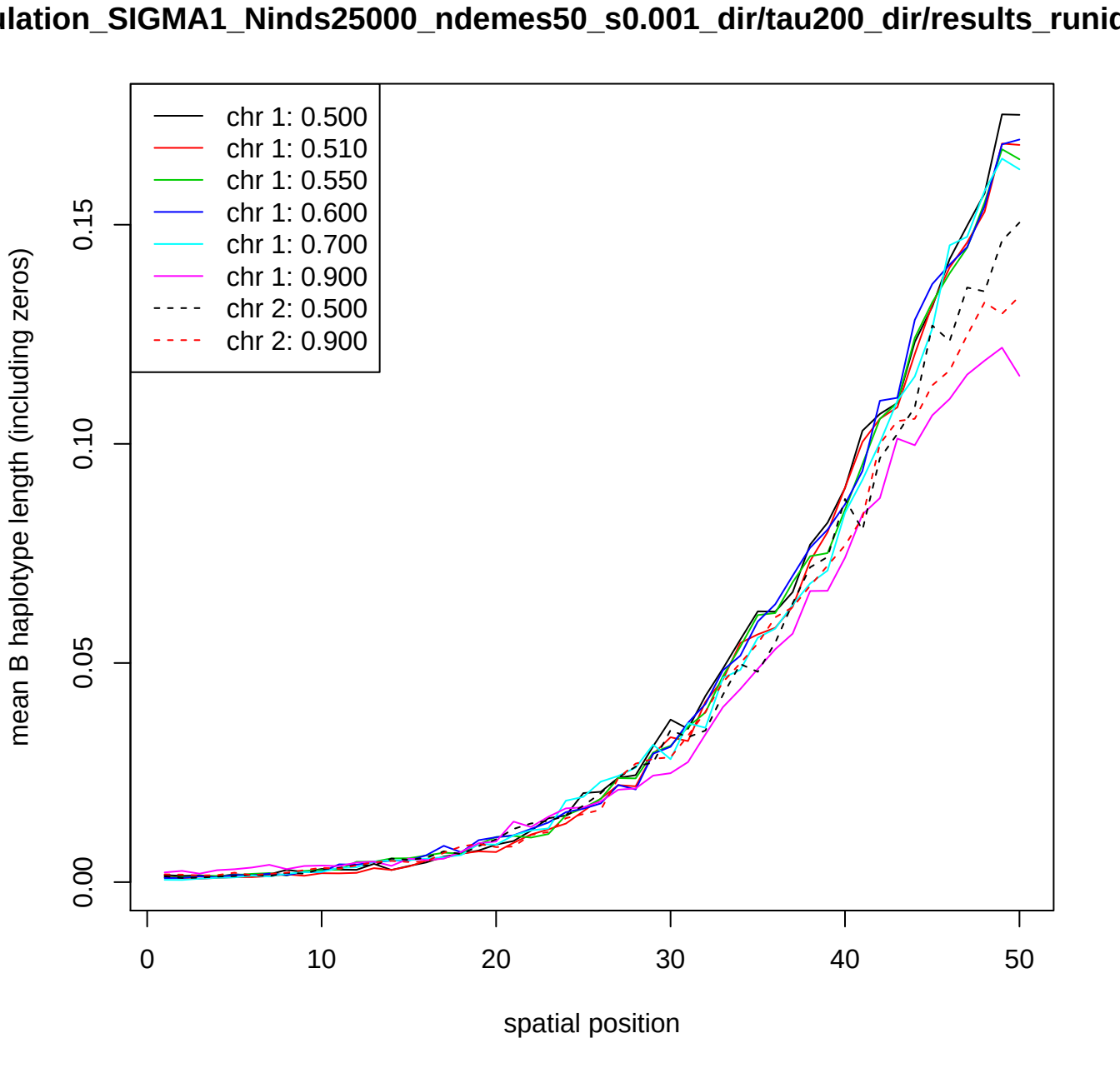

ulation_SIGMA1_Ninds25000_ndemes50_s0.001_dir/tau100_dir/results_runid
chr 1: 0.500
chr 1: 0.510
chr 1: 0.550
chr 1: 0.600
chr 1: 0.700
chr 1: 0.900
chr 2: 0.500
chr 2: 0.900
mean B haplotype length (including zeros)
0.0
0.2
0.4
0.6
0
10
20
30
40
50
spatial position


ulation_SIGMA1_Ninds25000_ndemes50_s0.001_dir/tau100_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
0.0
0.2
0.4
0.6
0.8
0
10
20
30
40
50
spatial position


ulation_SIGMA1_Ninds25000_ndemes50_s0.001_dir/tau100_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)
0.0
0.2
0.4
0.6
0
10
20
30
40
50
spatial position


ulation_SIGMA1_Ninds25000_ndemes50_s0.001_dir/tau100_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
0.0
0.2
0.4
0.6
0.8
0
10
20
30
40
50
spatial position


ulation_SIGMA1_Ninds25000_ndemes50_s0.001_dir/tau200_dir/results_runid
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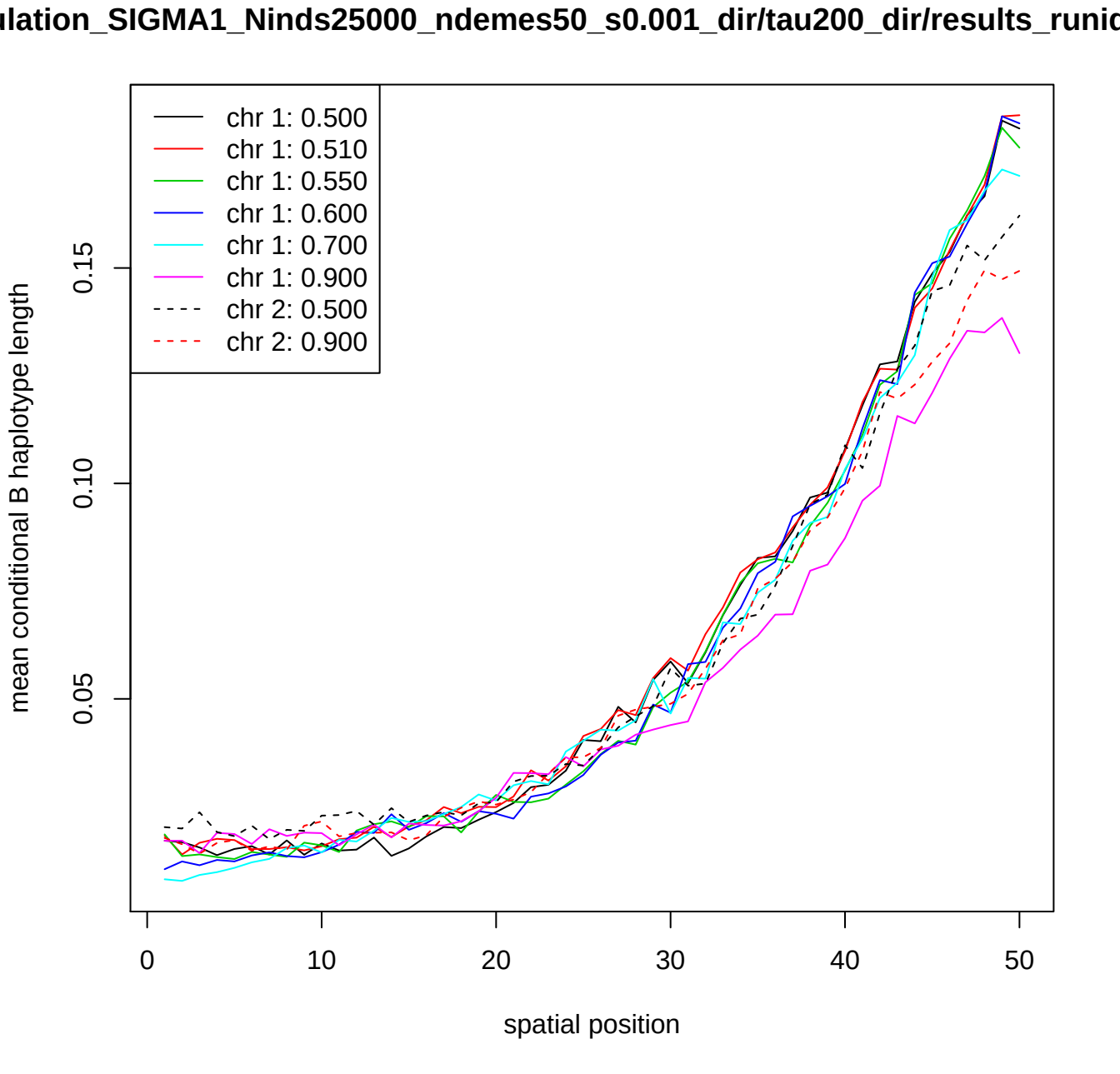
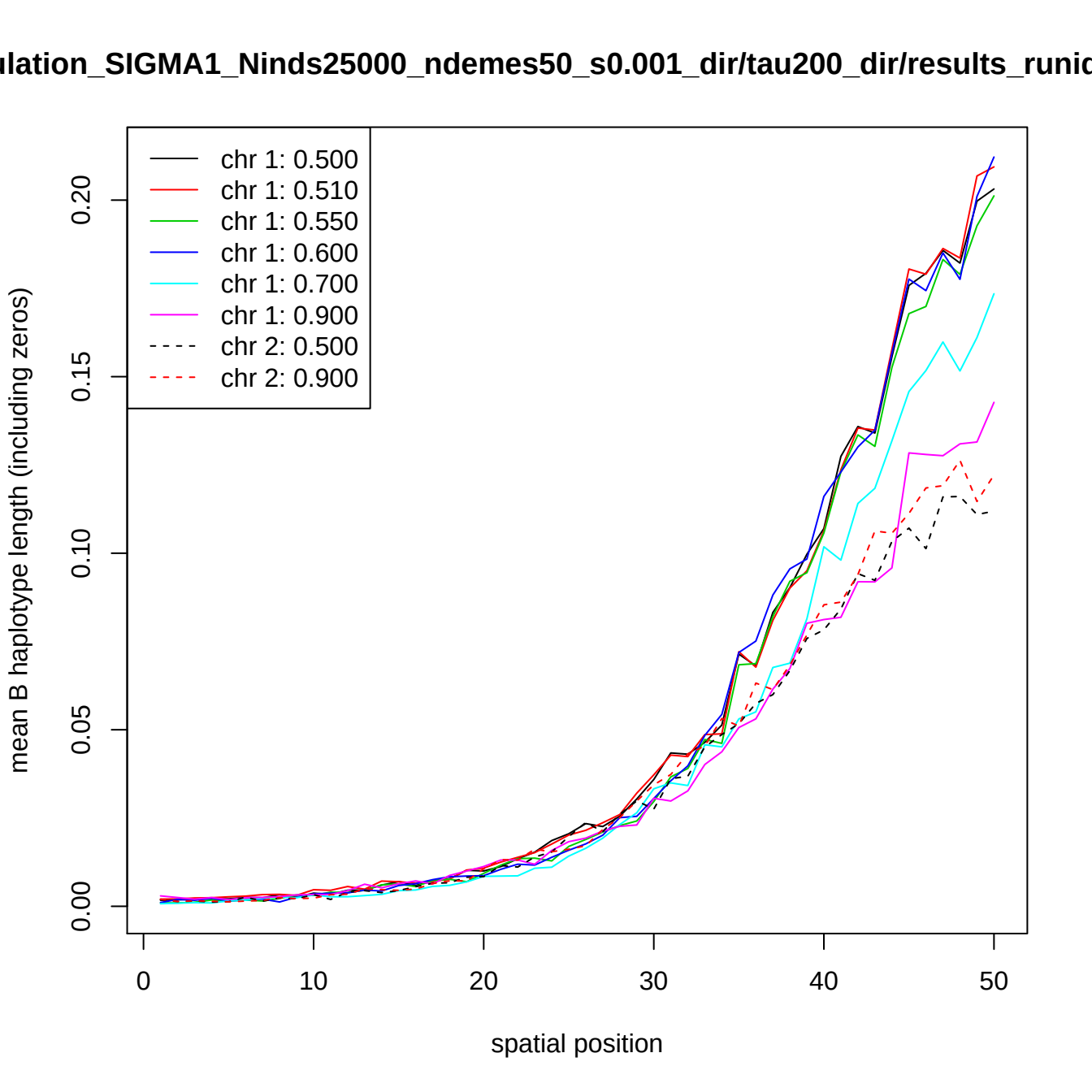
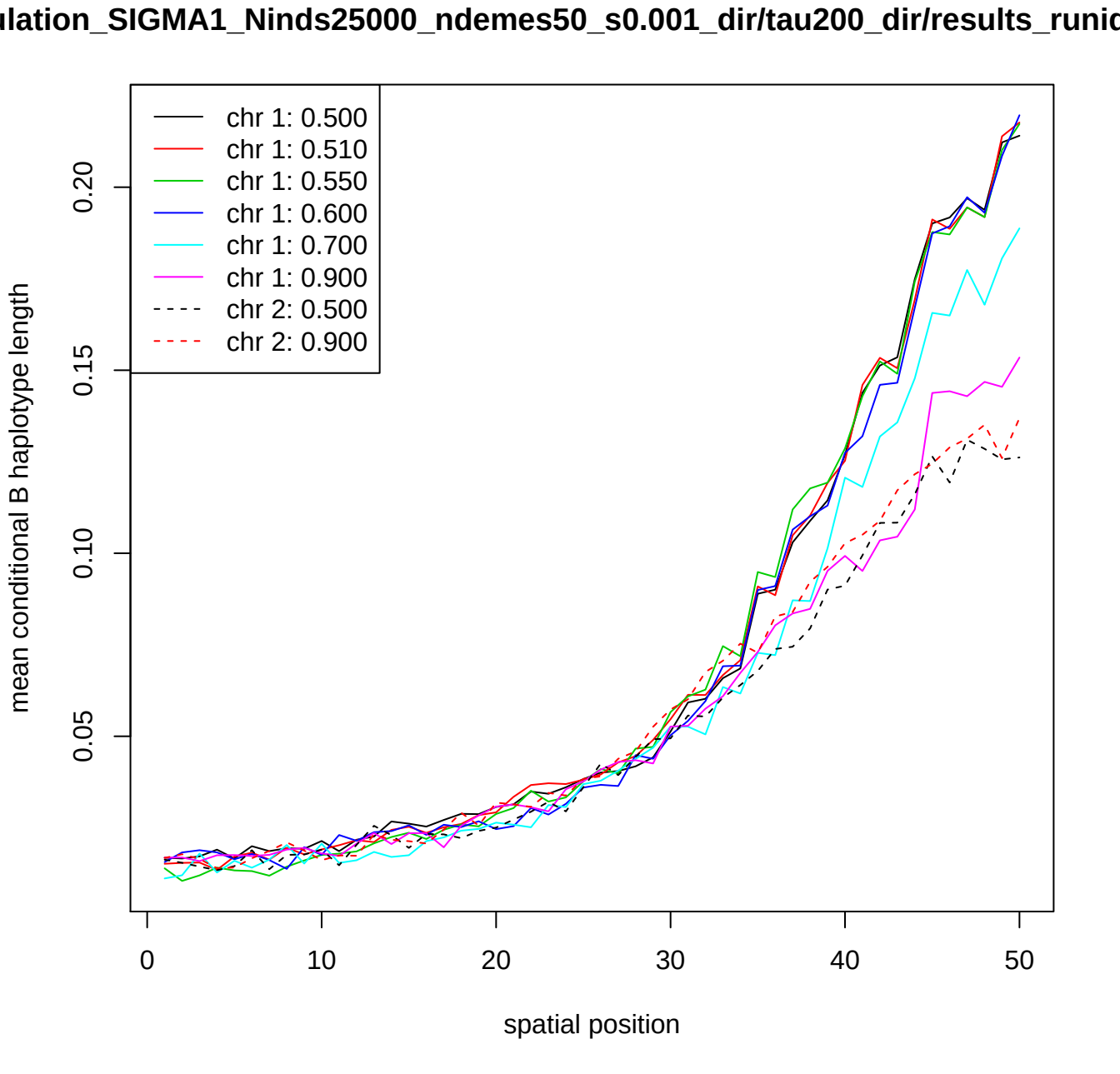
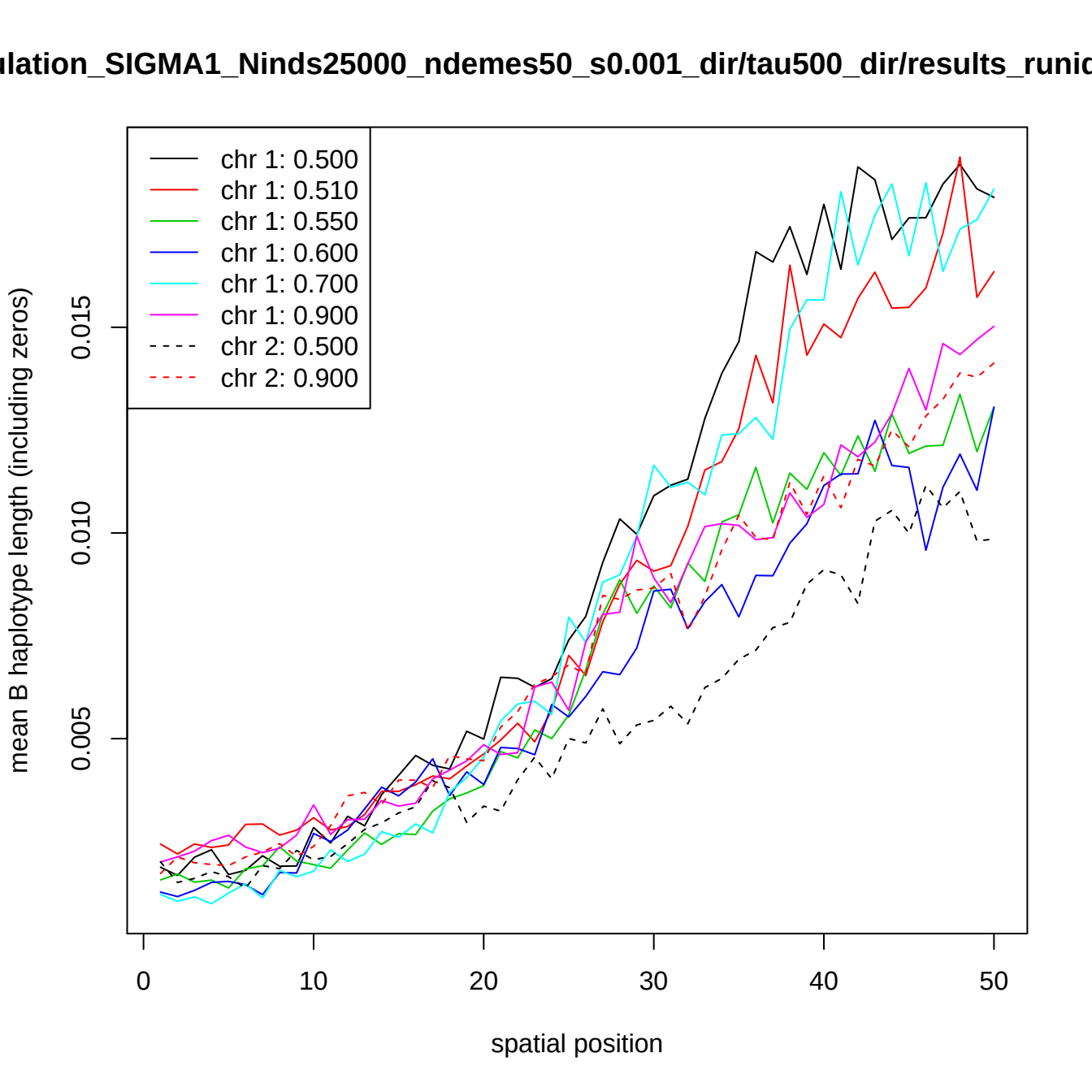
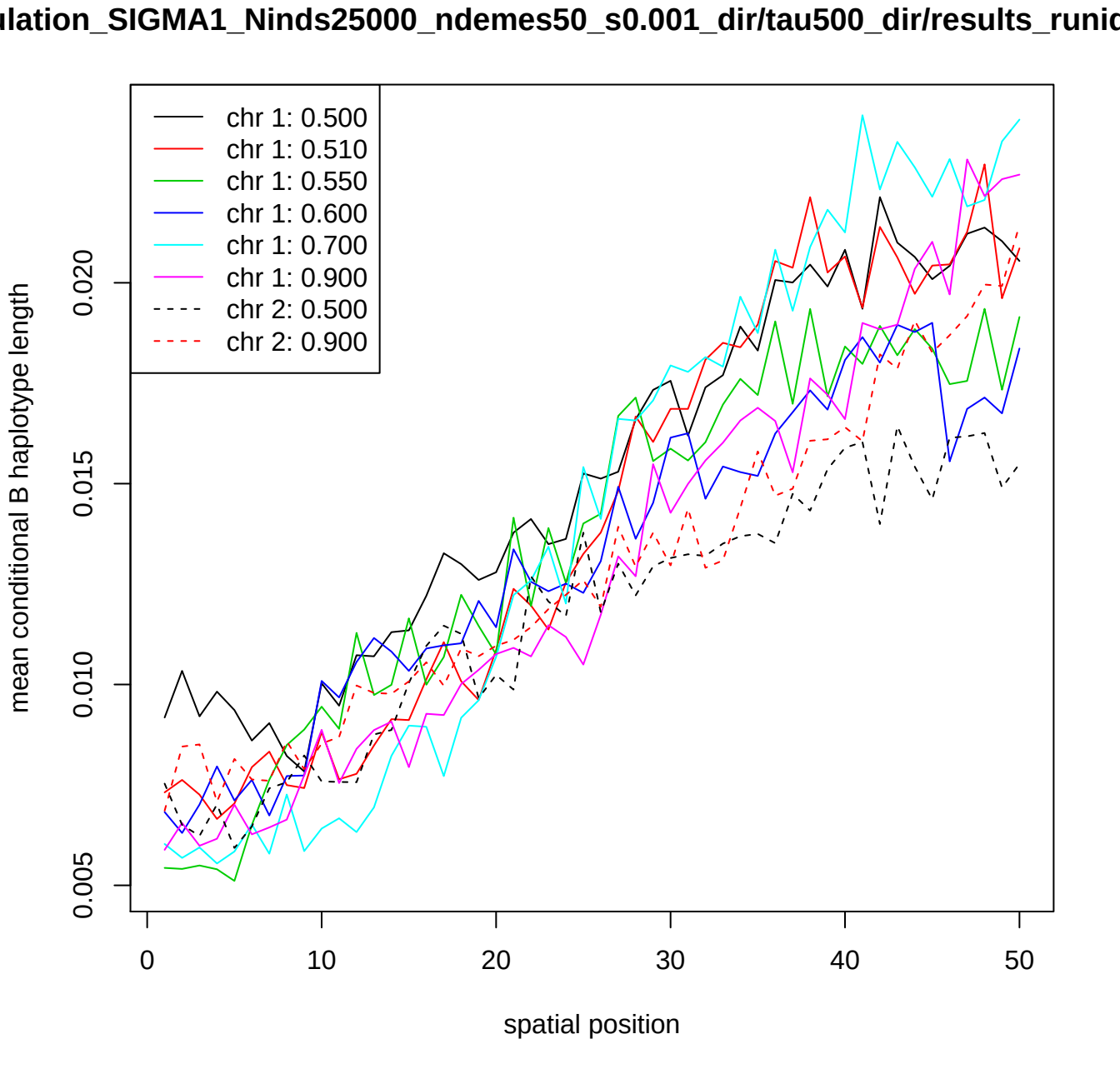
chr 1: 0.510
chr 1: 0.550
chr 1: 0.600
chr 1: 0.700
chr 1: 0.900
chr 2: 0.500
chr 2: 0.900
mean B haplotype length (including zeros)
0.00
0.05
0.10
0.15
0
10
20
30
40
50
spatial position

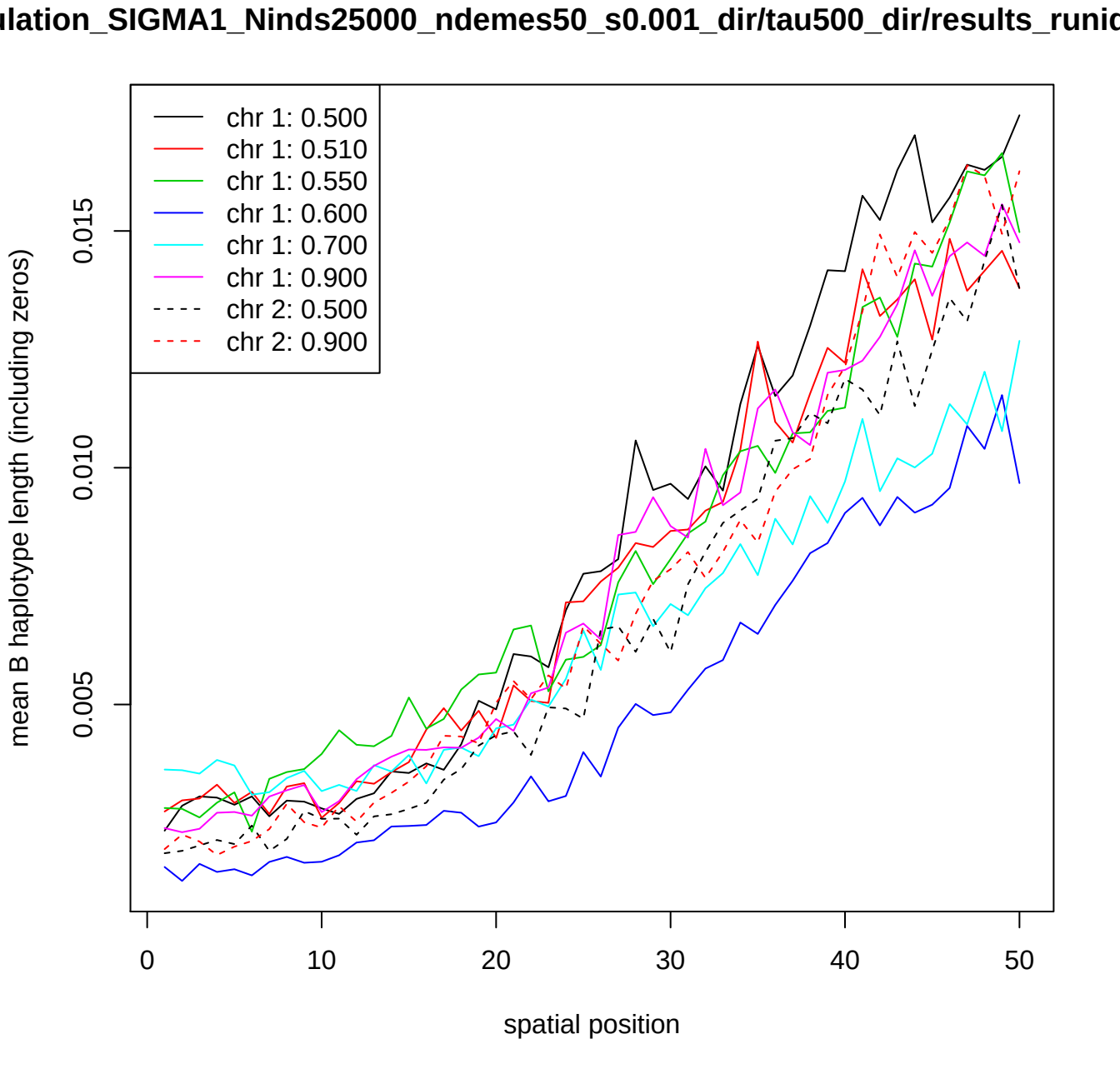

ulation_SIGMA1_Ninds25000_ndemes50_s0.001_dir/tau200_dir/results_runid
chr 1: 0.500
chr 1: 0.510
chr 1: 0.550
chr 1: 0.600
chr 1: 0.700
chr 1: 0.900
chr 2: 0.500
chr 2: 0.900
mean conditional B haplotype length
0.05
0.10
0.15
0
10
20
30
40
50
spatial position

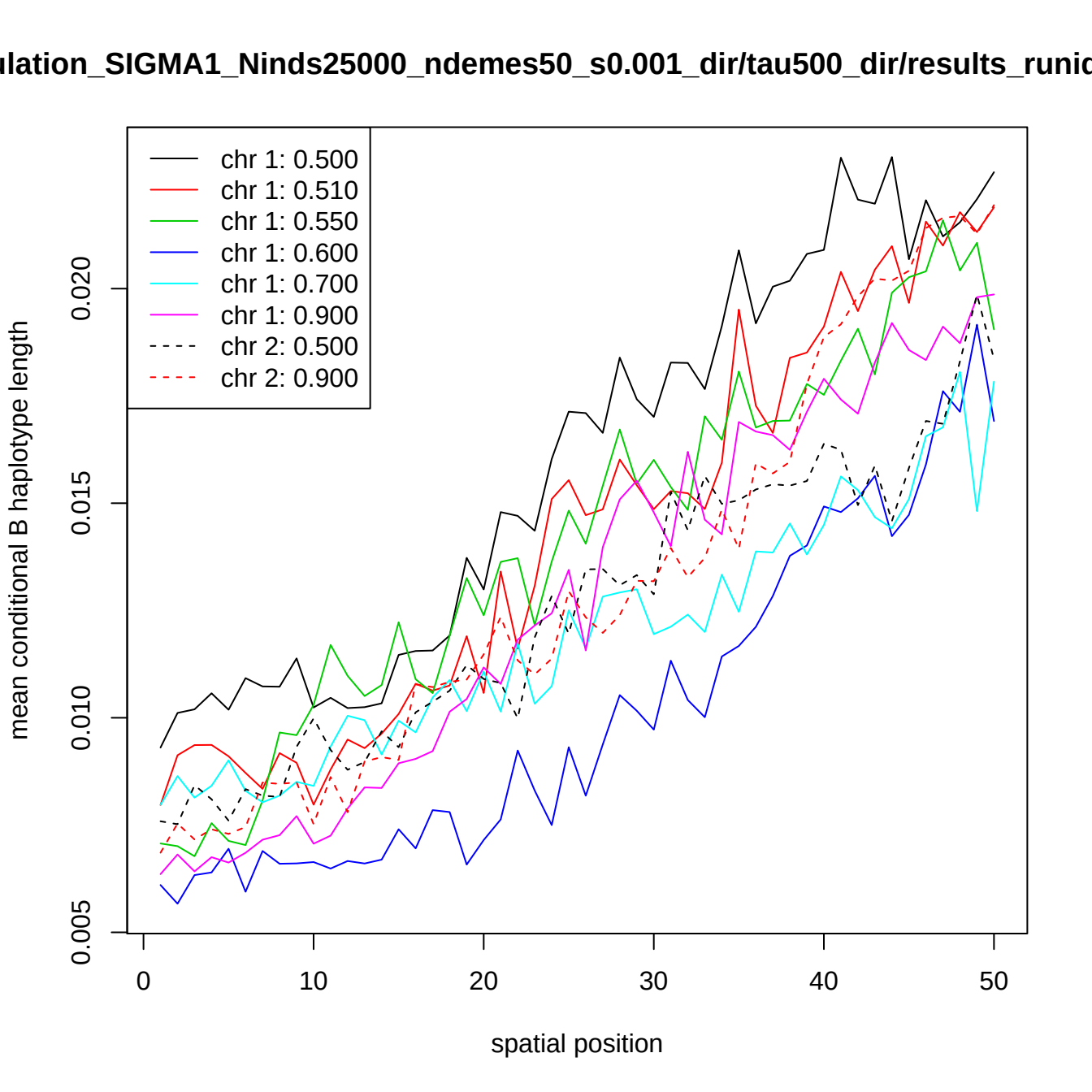

ulation_SIGMA1_Ninds25000_ndemes50_s0.001_dir/tau200_dir/results_runid
chr 1: 0.500
chr 1: 0.510
chr 1: 0.550
chr 1: 0.600
chr 1: 0.700
chr 1: 0.900
chr 2: 0.500
chr 2: 0.900
mean B haplotype length (including zeros)
0.00
0.05
0.10
0.15
0.20
0
10
20
30
40
50
spatial position


ulation_SIGMA1_Ninds25000_ndemes50_s0.001_dir/tau200_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
0.05
0.10
0.15
0.20
0
10
20
30
40
50
spatial position


ulation_SIGMA1_Ninds25000_ndemes50_s0.001_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)
0.005
0.010
0.015
0
10
20
30
40
50
spatial position


ulation_SIGMA1_Ninds25000_ndemes50_s0.001_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
10
20
30
40
50
spatial position


ulation_SIGMA1_Ninds25000_ndemes50_s0.001_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)
0.005
0.010
0.015
0
10
20
30
40
50
spatial position


ulation_SIGMA1_Ninds25000_ndemes50_s0.001_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
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0.015
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10
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spatial position