

Normalized mean length/mean flanking length, ancestry B $\tau=10$, $s=0.100$

Normalized mean block length

distance from center:

— -49.5
— -38.5
— -27.5
— -16.5
— -5.5
— 5.5
— 16.5
— 27.5
— 38.5
— 49.5

-0.02

-0.01

0

0.01

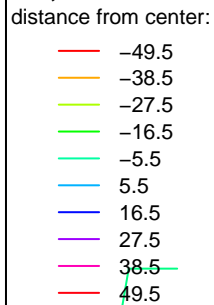
0.02

Distance from selected locus (M)

3.0
2.5
2.0
1.5
1.0
0.5

Normalized mean length/mean flanking length, ancestry B tau=20, s=0.100

Normalized mean block length



-0.02

-0.01

0

0.01

0.02

Distance from selected locus (M)

Normalized mean length/mean flanking length, ancestry B tau=50, s=0.100

Normalized mean block length

distance from center:

— -49.5
— -38.5
— -27.5
— -16.5
— -5.5
— 5.5
— 16.5
— 27.5
— 38.5
— 49.5

-0.02

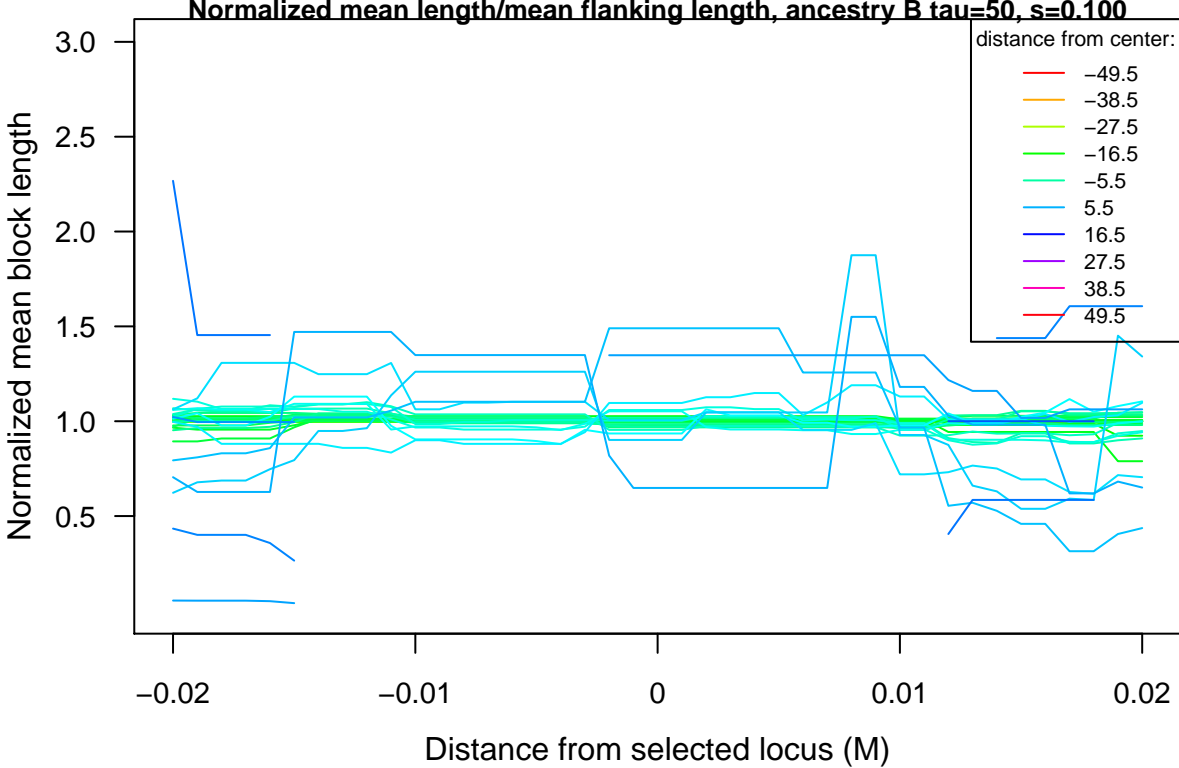
-0.01

0

0.01

0.02

Distance from selected locus (M)



Normalized mean length/mean flanking length, ancestry B $\tau=100$, $s=0.100$

Normalized mean block length

distance from center:

— -49.5
— -38.5
— -27.5
— -16.5
— -5.5
— 5.5
— 16.5
— 27.5
— 38.5
— 49.5

-0.02

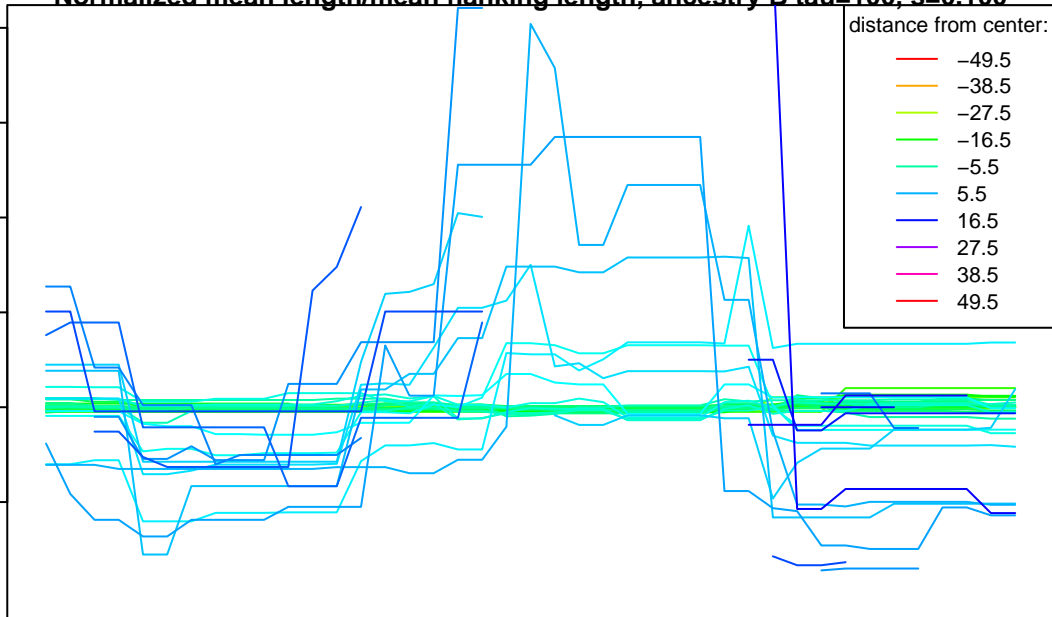
-0.01

0

0.01

0.02

Distance from selected locus (M)



Normalized mean length/mean flanking length, ancestry B tau=200, s=0.100

Normalized mean block length

distance from center:

— -49.5
— -38.5
— -27.5
— -16.5
— -5.5
— 5.5
— 16.5
— 27.5
— 38.5
— 49.5

-0.02

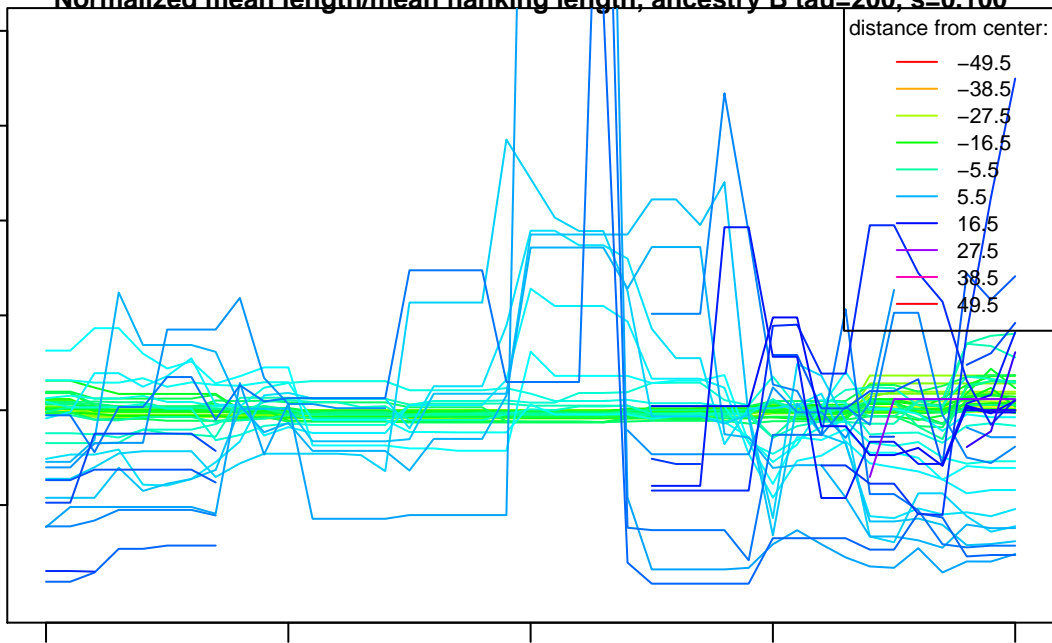
-0.01

0

0.01

0.02

Distance from selected locus (M)



Normalized mean length/mean flanking length, ancestry B $\tau=500$, $s=0.100$

Normalized mean block length

