

Normalized mean length/mean flanking length, ancestry B tau=50, s=0.010

Normalized mean block length

distance from center:

-59.5  
-46.5  
-33.5  
-20.5  
-7.5  
6.5  
19.5  
32.5  
45.5  
59.5

-0.02

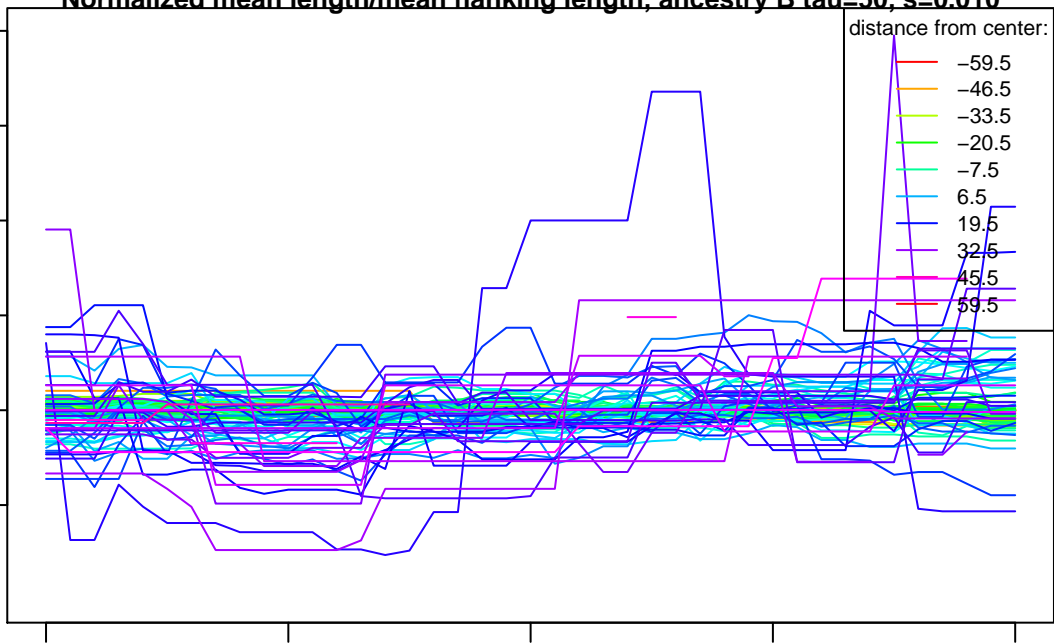
-0.01

0

0.01

0.02

Distance from selected locus (M)



Normalized mean length/mean flanking length, ancestry B  $\tau=100$ ,  $s=0.010$

Normalized mean block length

distance from center:

— 59.5  
— 46.5  
— 33.5  
— 20.5  
— 7.5  
— 6.5  
— 19.5  
— 32.5  
— 45.5  
— 59.5

-0.02

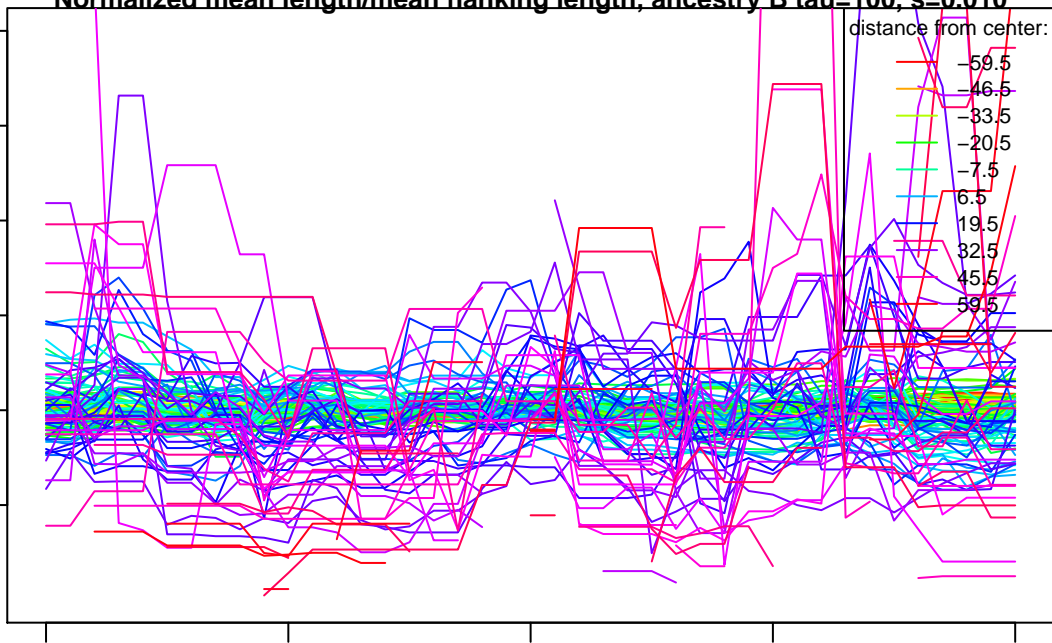
-0.01

0

0.01

0.02

Distance from selected locus (M)



Normalized mean length/mean flanking length, ancestry B tau=150, s=0.010

Normalized mean block length

distance from center:

— -59.5  
— -46.5  
— -33.5  
— -20.5  
— -7.5  
— 6.5  
— 19.5  
— 32.5  
— 45.5  
— 59.5

-0.02

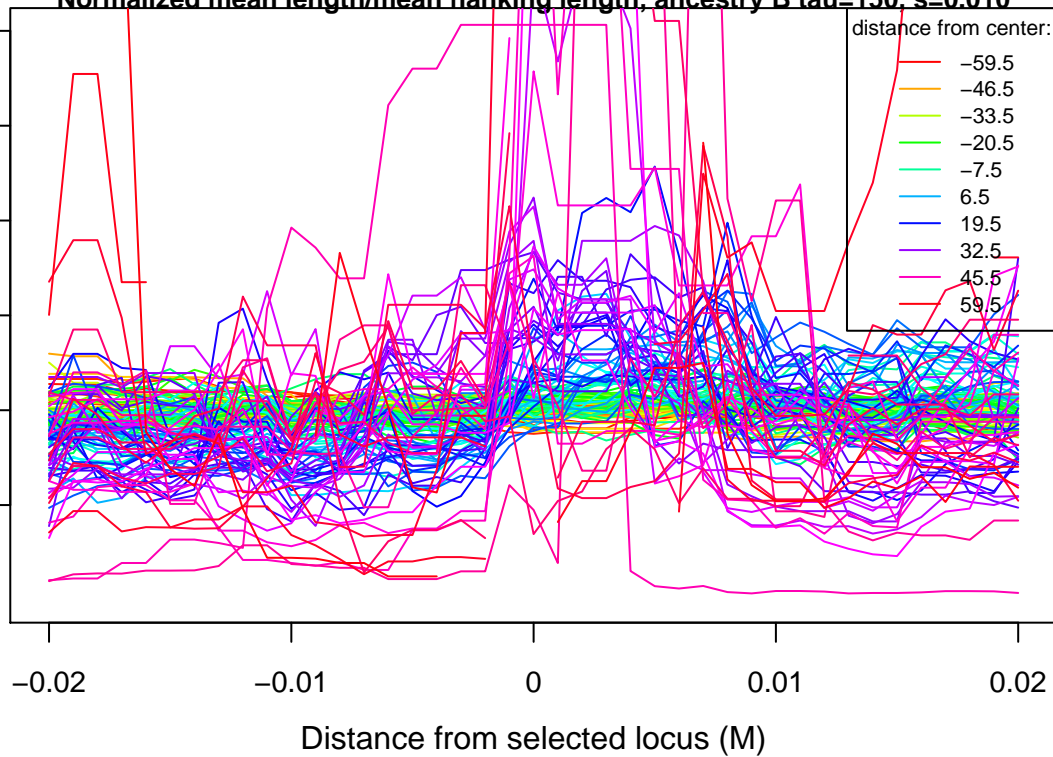
-0.01

0

0.01

0.02

Distance from selected locus (M)



Normalized mean length/mean flanking length, ancestry B tau=250, s=0.010

Normalized mean block length

distance from center:

— -59.5  
— -46.5  
— -33.5  
— -20.5  
— -7.5  
— 6.5  
— 19.5  
— 32.5  
— 45.5  
— 59.5

-0.02

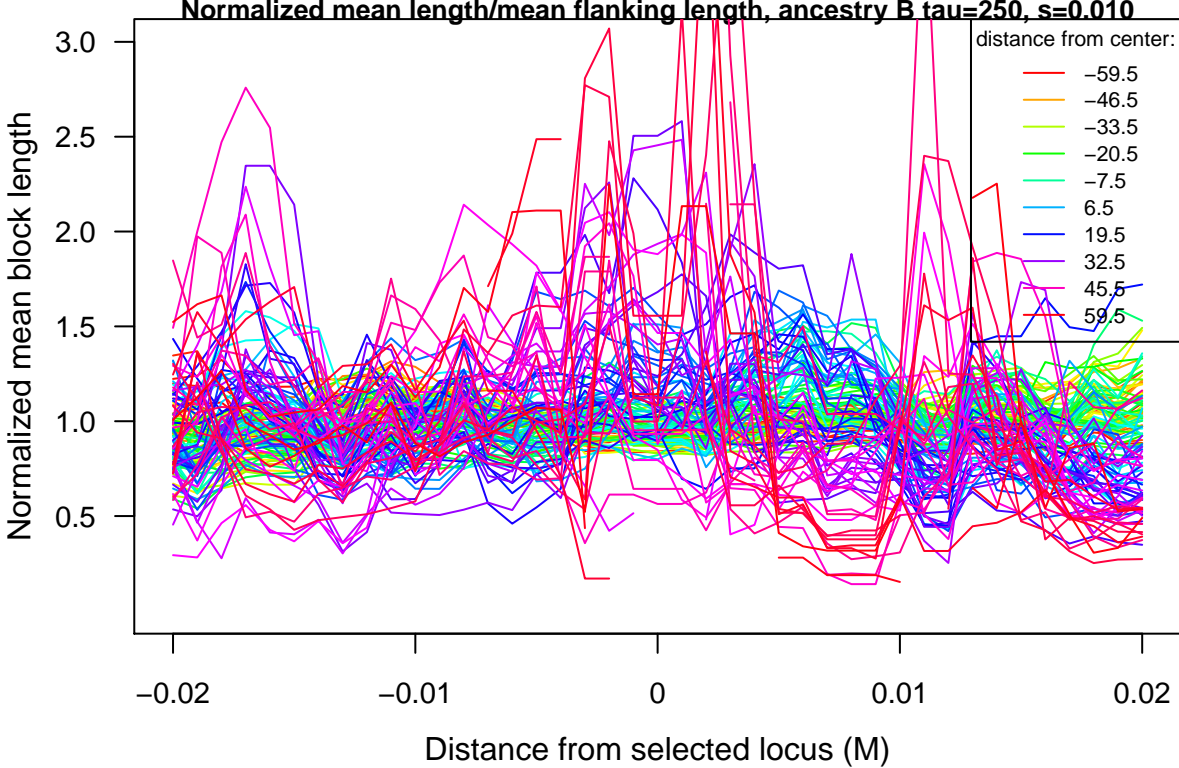
-0.01

0

0.01

0.02

Distance from selected locus (M)



Normalized mean length/mean flanking length, ancestry B tau=750, s=0.010

Normalized mean block length

distance from center:

— -59.5  
— -46.5  
— -33.5  
— -20.5  
— -7.5  
— 6.5  
— 19.5  
— 32.5  
— 45.5  
— 59.5

-0.02

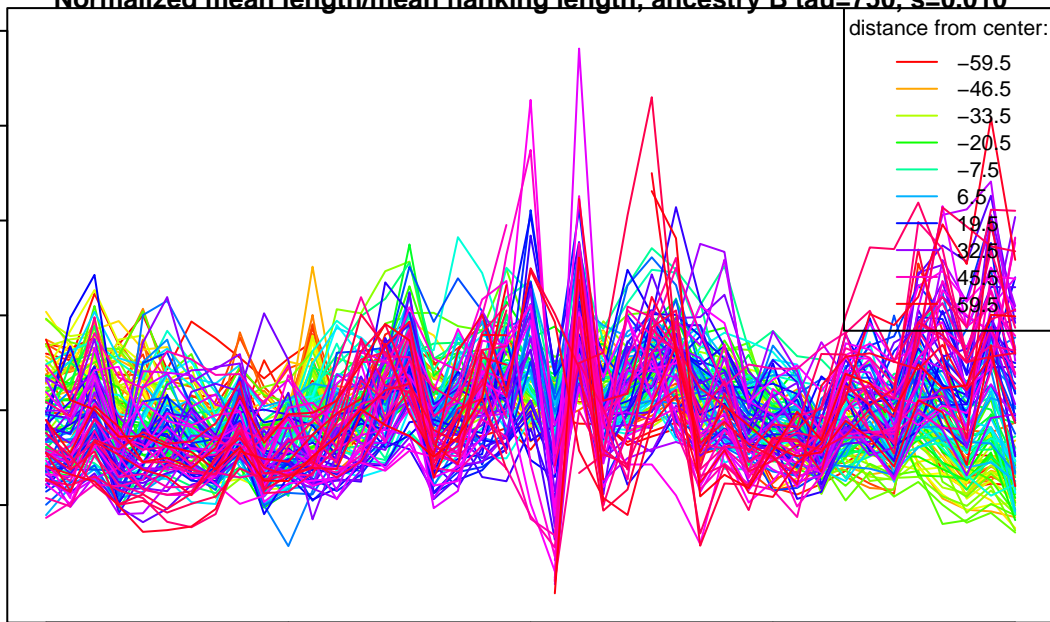
-0.01

0

0.01

0.02

Distance from selected locus (M)



Normalized mean length/mean flanking length, ancestry B tau=1000, s=0.010

Normalized mean block length

distance from center:

-59.5  
-46.5  
-33.5  
-20.5  
-7.5  
6.5  
19.5  
32.5  
45.5  
59.5

-0.02

-0.01

0

0.01

0.02

Distance from selected locus (M)

