

Normalized mean length/mean flanking length, ancestry B  $\tau=100$ ,  $s=0.001$

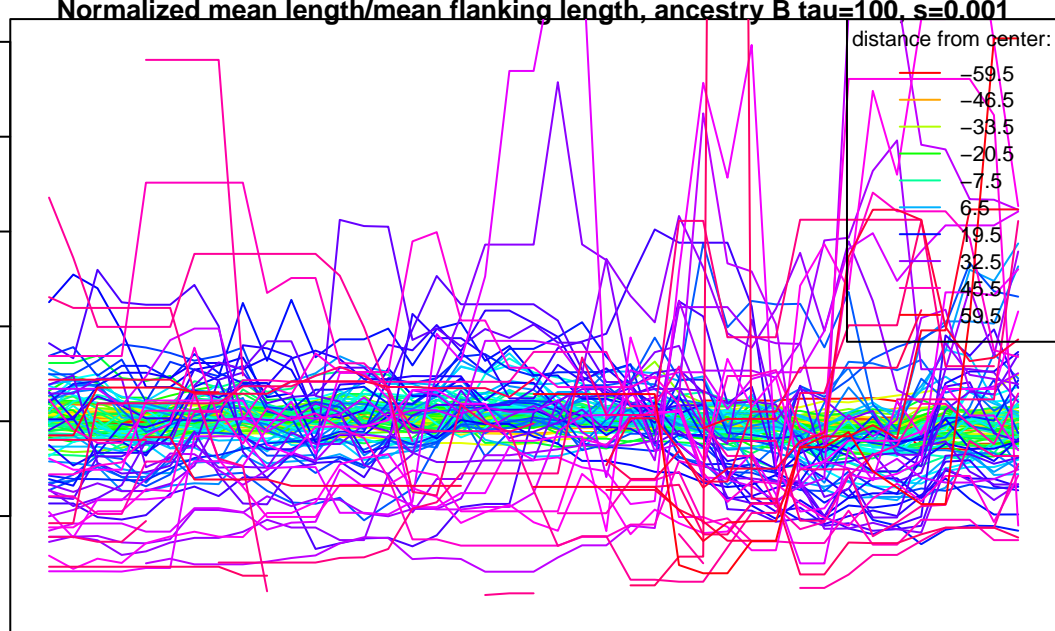
Normalized mean block length

3.0  
2.5  
2.0  
1.5  
1.0  
0.5

distance from center:

59.5  
46.5  
33.5  
20.5  
7.5  
6.5  
19.5  
32.5  
45.5  
59.5

Distance from selected locus (M)



Normalized mean length/mean flanking length, ancestry B tau=100, s=0.001

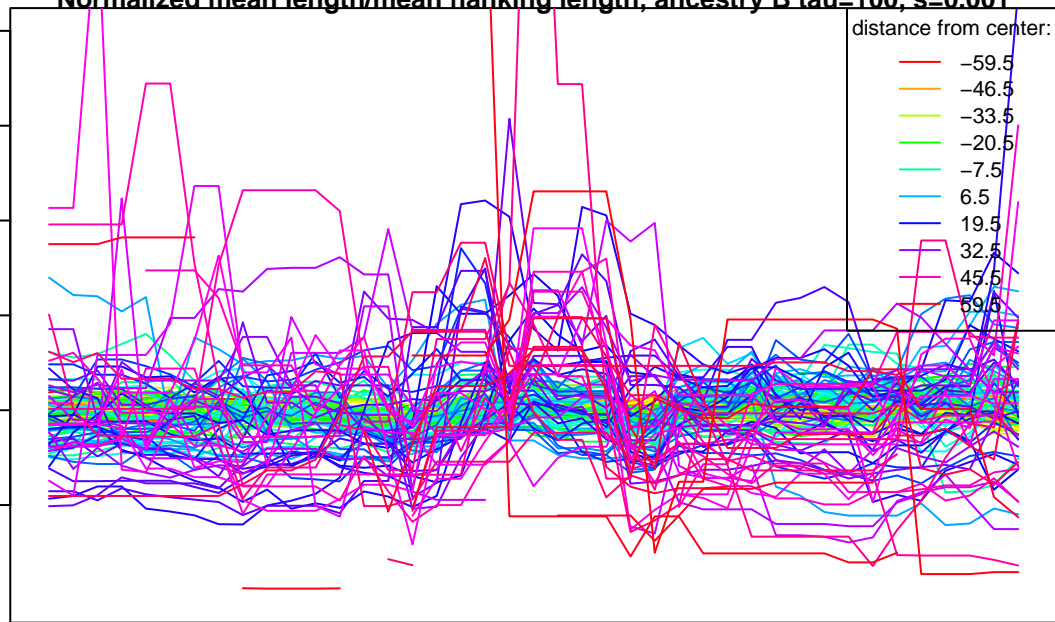
Normalized mean block length

3.0  
2.5  
2.0  
1.5  
1.0  
0.5

distance from center:

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

Distance from selected locus (M)



Normalized mean length/mean flanking length, ancestry B  $\tau=200$ ,  $s=0.001$

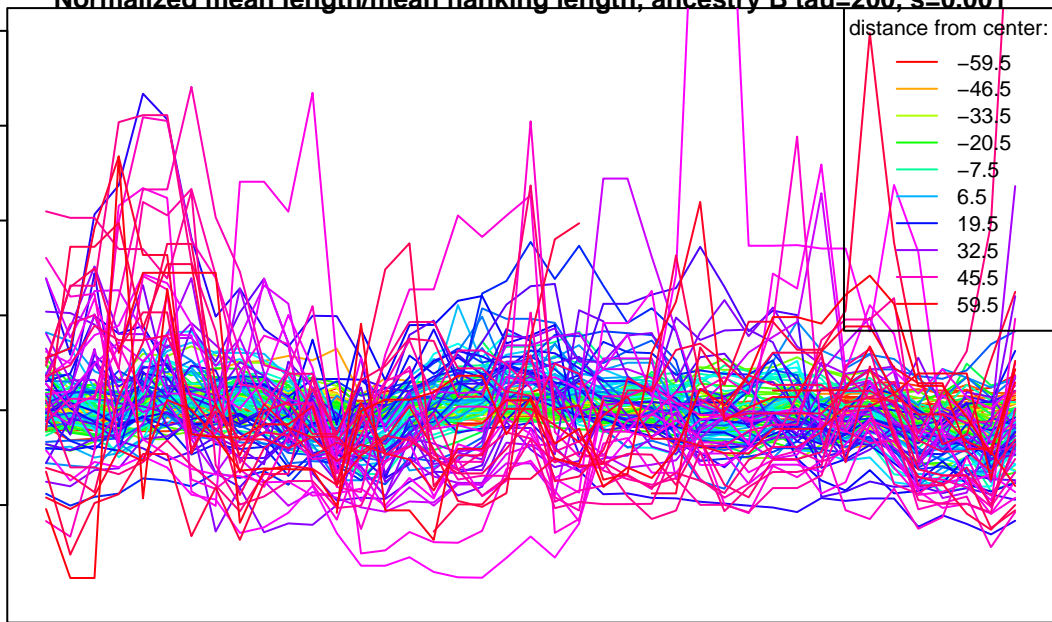
Normalized mean block length

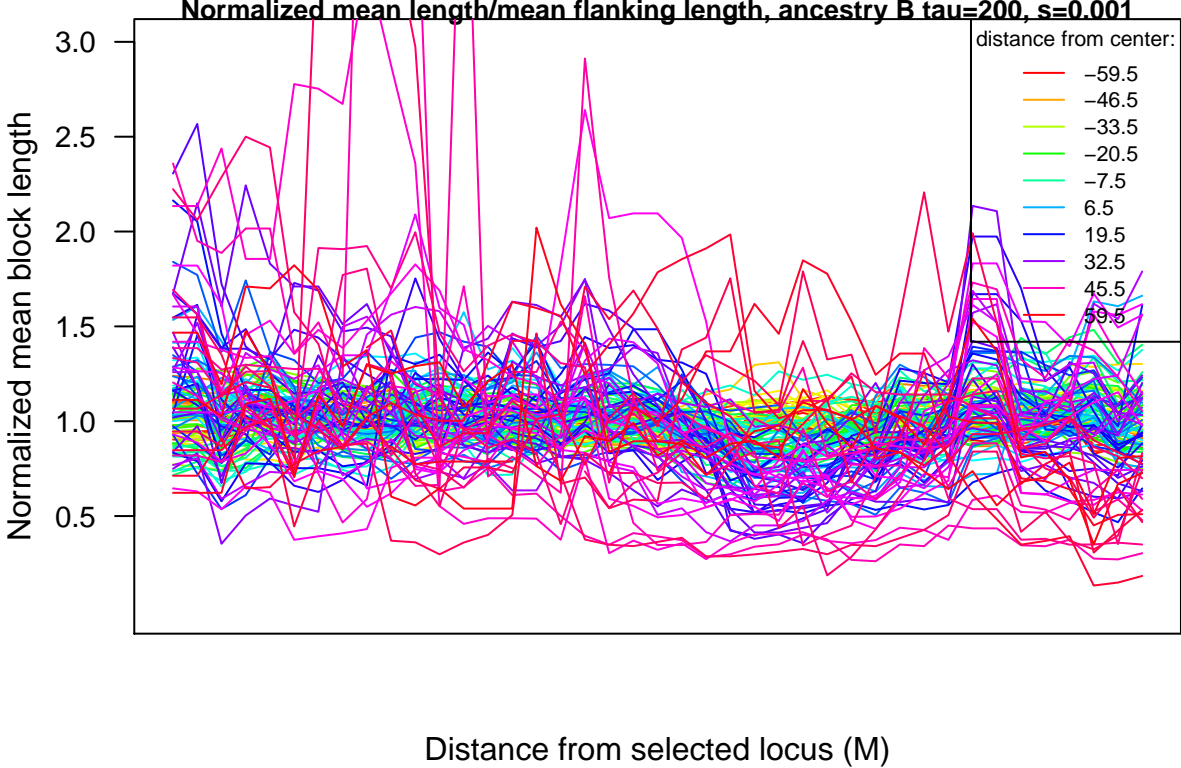
3.0  
2.5  
2.0  
1.5  
1.0  
0.5

distance from center:

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

Distance from selected locus (M)





Normalized mean length/mean flanking length, ancestry B tau=500, s=0.001

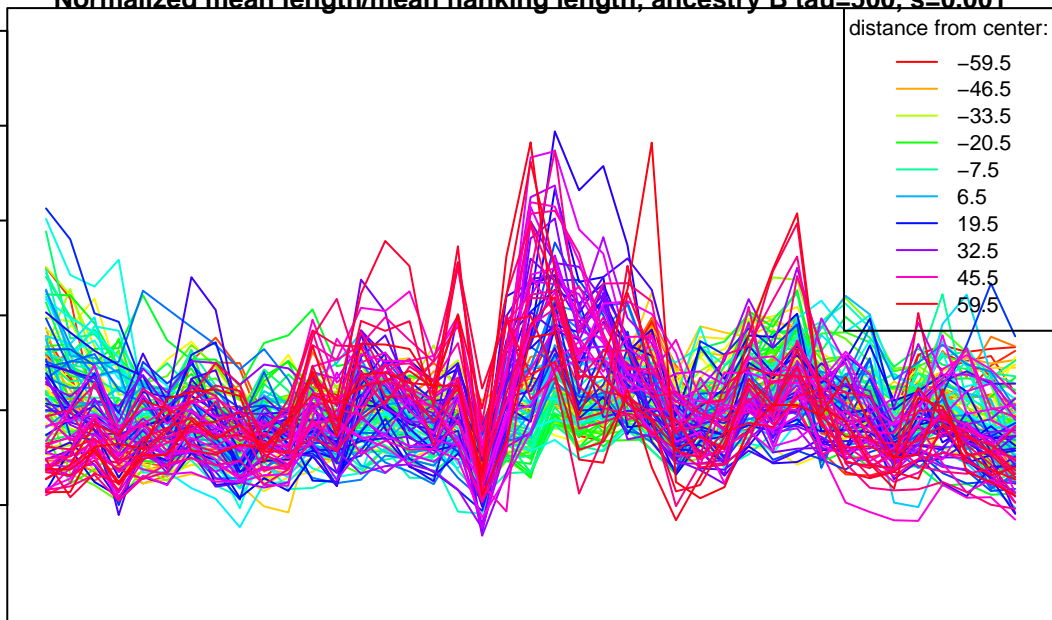
Normalized mean block length

3.0  
2.5  
2.0  
1.5  
1.0  
0.5

distance from center:

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

Distance from selected locus (M)



Normalized mean length/mean flanking length, ancestry B  $\tau=500$ ,  $s=0.001$

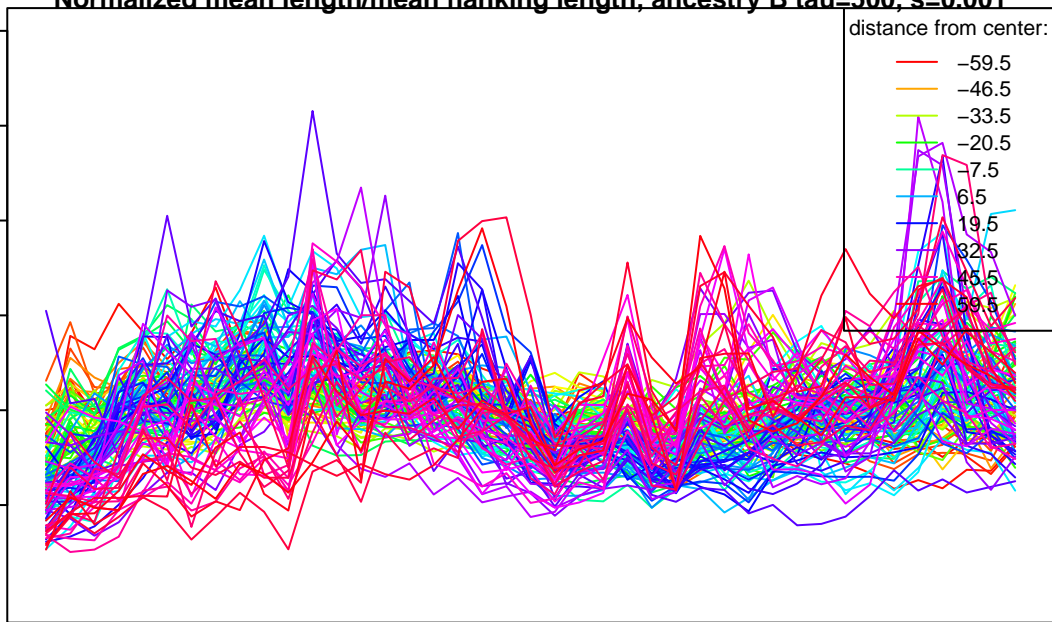
Normalized mean block length

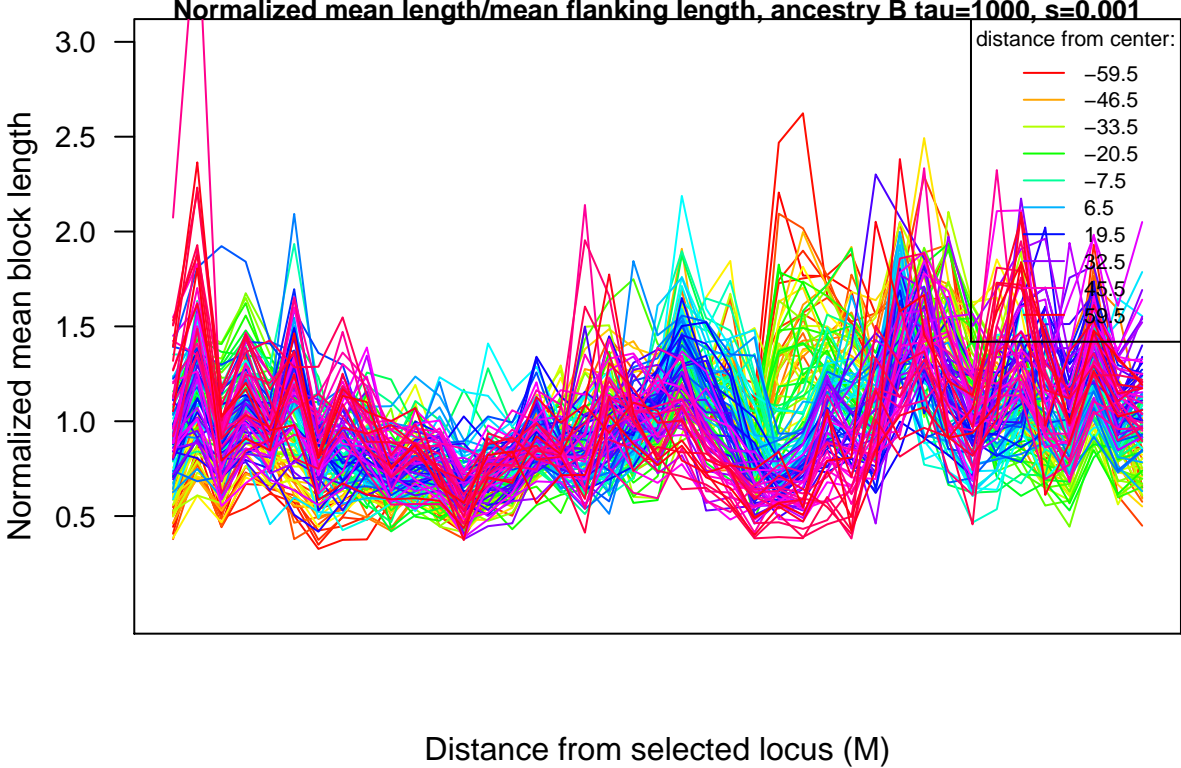
3.0  
2.5  
2.0  
1.5  
1.0  
0.5

distance from center:

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

Distance from selected locus (M)





Normalized mean length/mean flanking length, ancestry B tau=1000, s=0.001

Normalized mean block length

3.0  
2.5  
2.0  
1.5  
1.0  
0.5

distance from center:

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

Distance from selected locus (M)

