

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

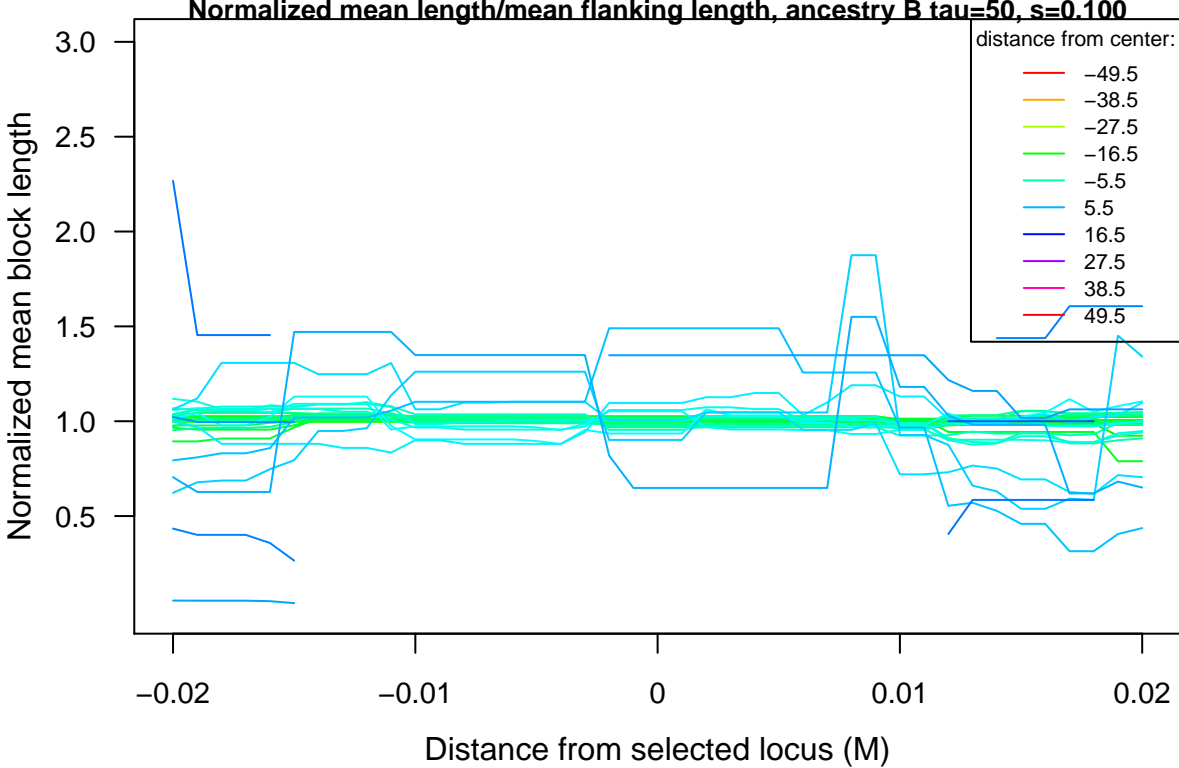
-0.01

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

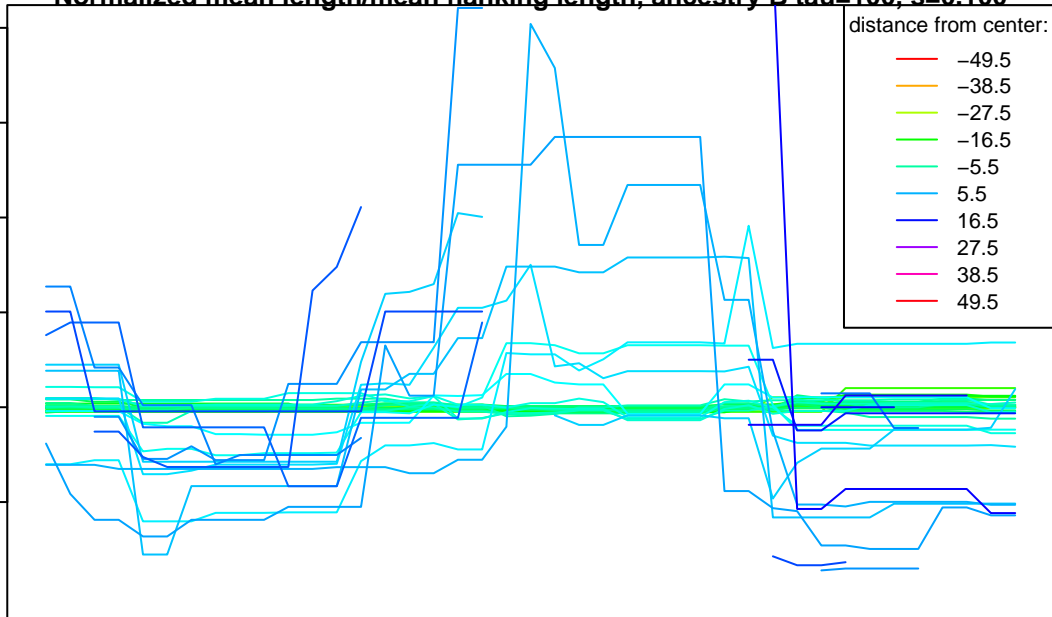
-0.01

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

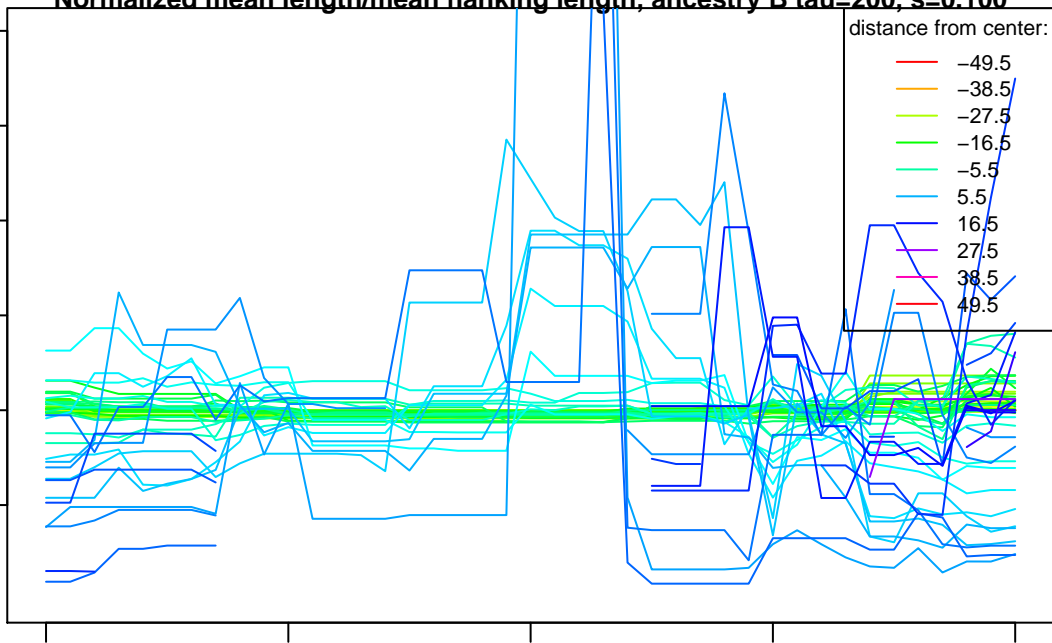
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0

0.01

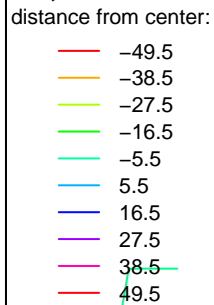
0.02

Distance from selected locus (M)



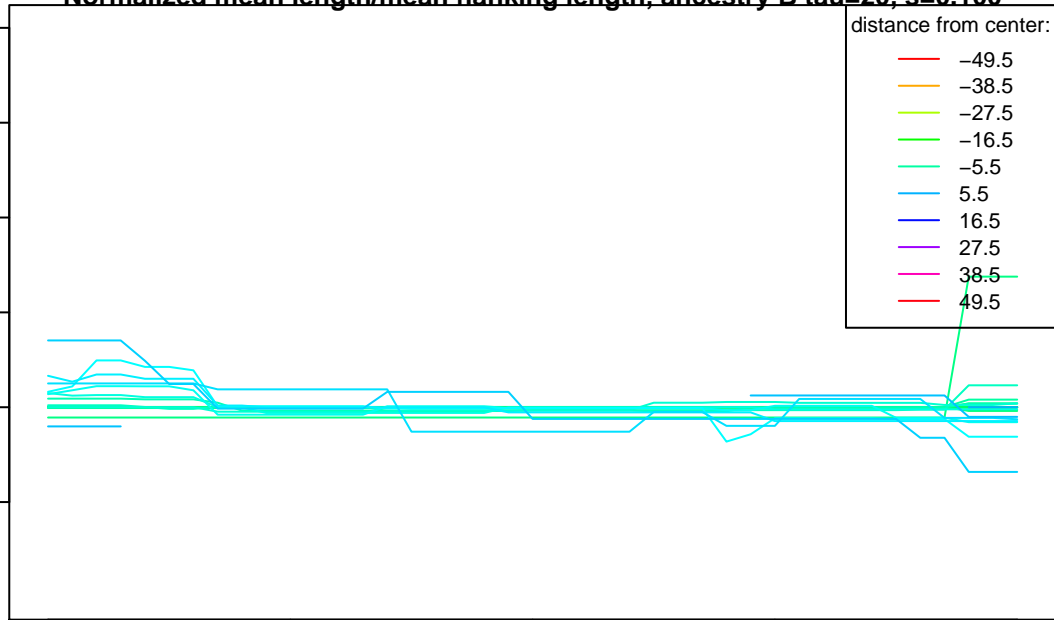
Normalized mean length/mean flanking length, ancestry B tau=20, s=0.100

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=500, 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

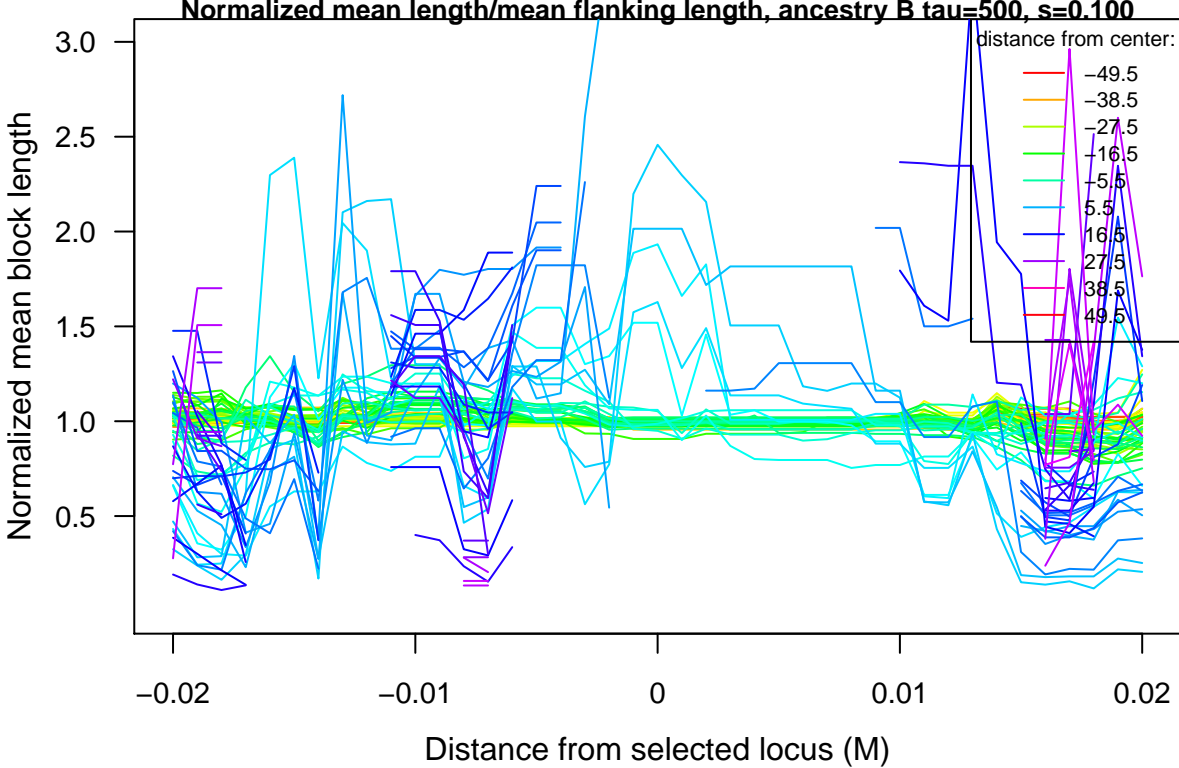
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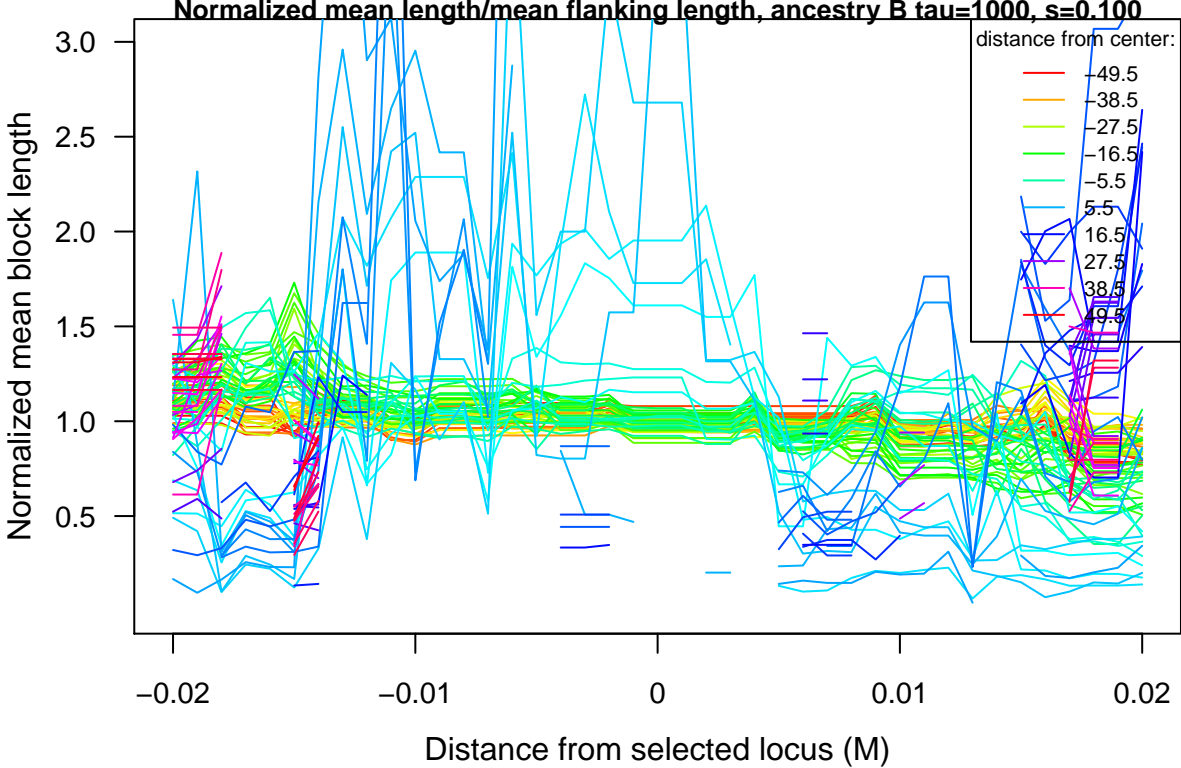
0

0.01

0.02

Distance from selected locus (M)





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