

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

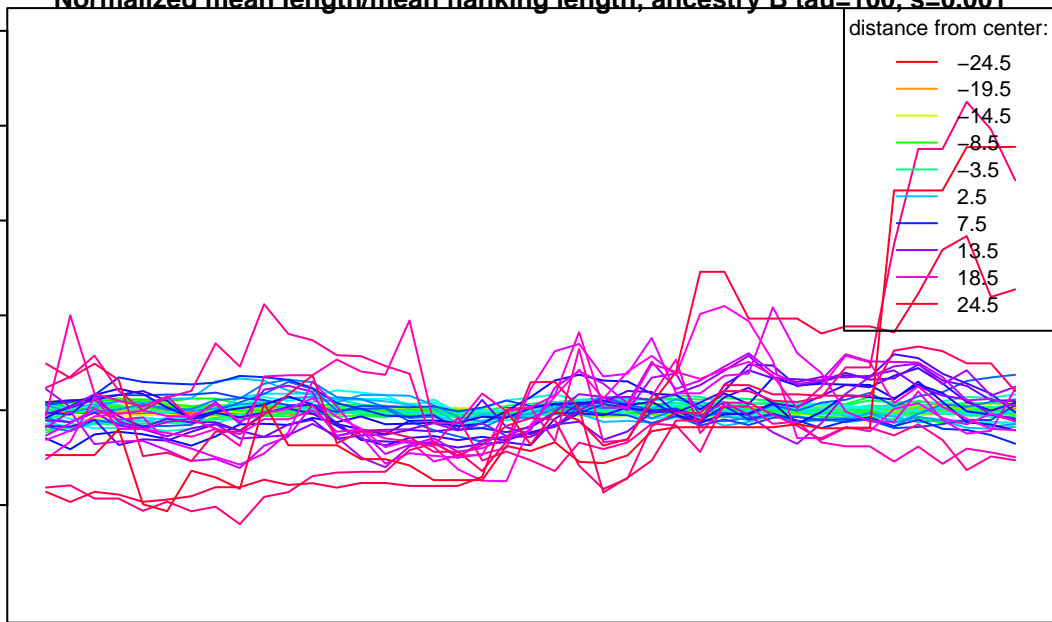
Normalized mean block length

3.0
2.5
2.0
1.5
1.0
0.5

distance from center:

— -24.5
— -19.5
— -14.5
— -8.5
— -3.5
— 2.5
— 7.5
— 13.5
— 18.5
— 24.5

Distance from selected locus (M)



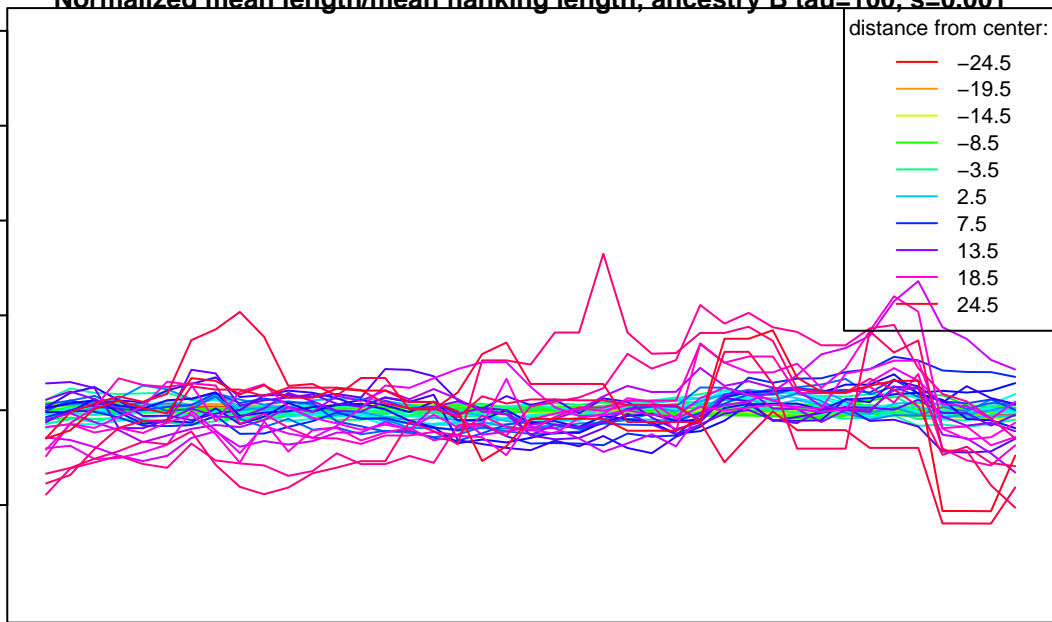
Normalized mean block length

3.0
2.5
2.0
1.5
1.0
0.5

distance from center:

- 24.5
- 19.5
- 14.5
- 8.5
- 3.5
- 2.5
- 7.5
- 13.5
- 18.5
- 24.5

Distance from selected locus (M)



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

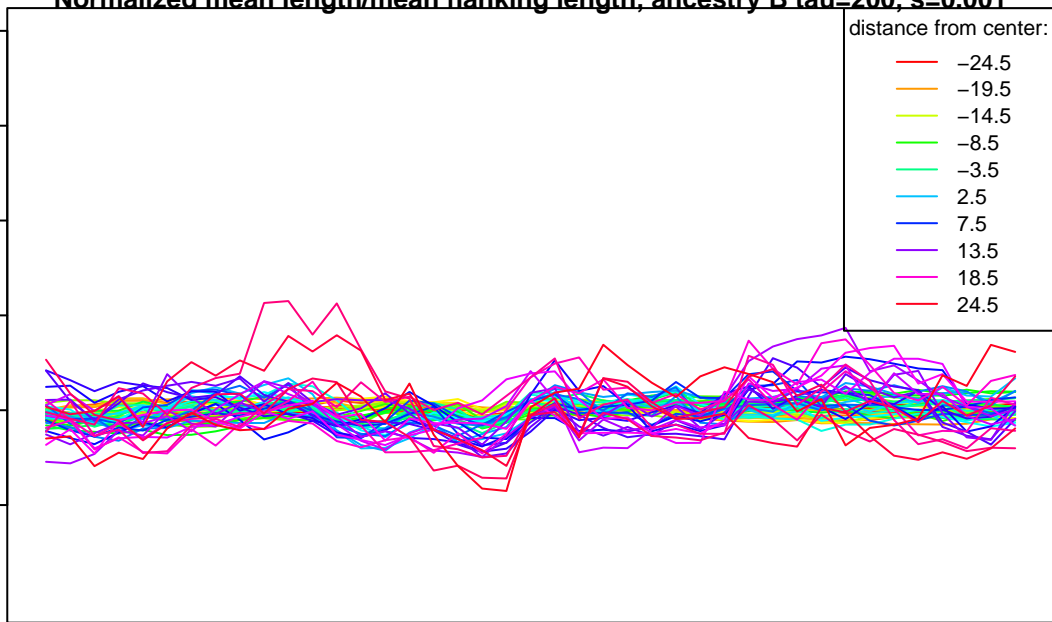
Normalized mean block length

3.0
2.5
2.0
1.5
1.0
0.5

distance from center:

-24.5
-19.5
-14.5
-8.5
-3.5
2.5
7.5
13.5
18.5
24.5

Distance from selected locus (M)



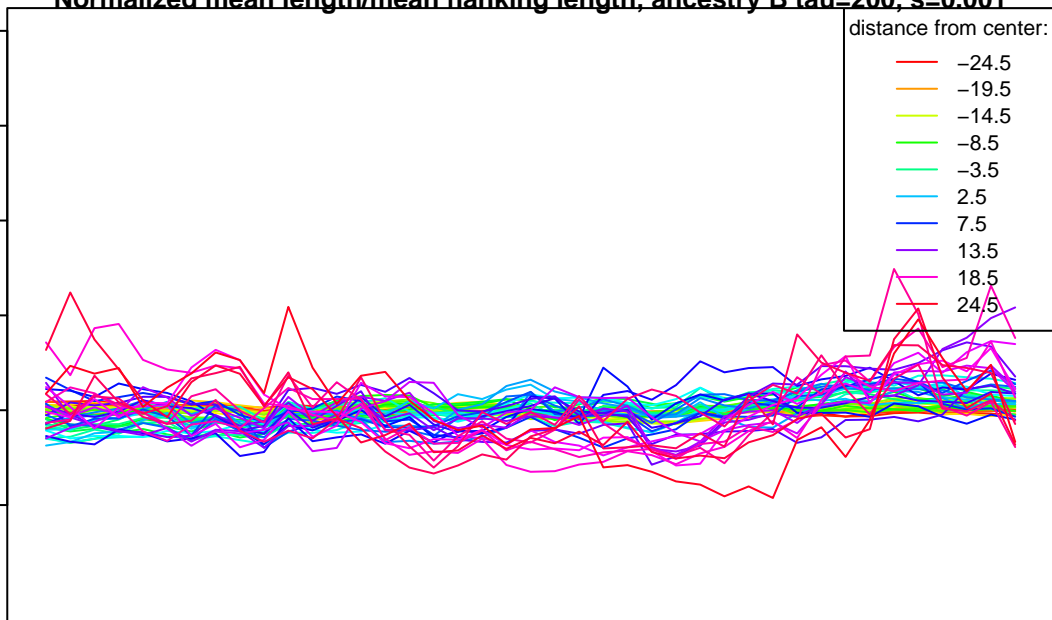
Normalized mean length/mean flanking length, ancestry B $\tau=200$, $s=0.001$

Normalized mean block length

distance from center:

— -24.5
— -19.5
— -14.5
— -8.5
— -3.5
— 2.5
— 7.5
— 13.5
— 18.5
— 24.5

Distance from selected locus (M)



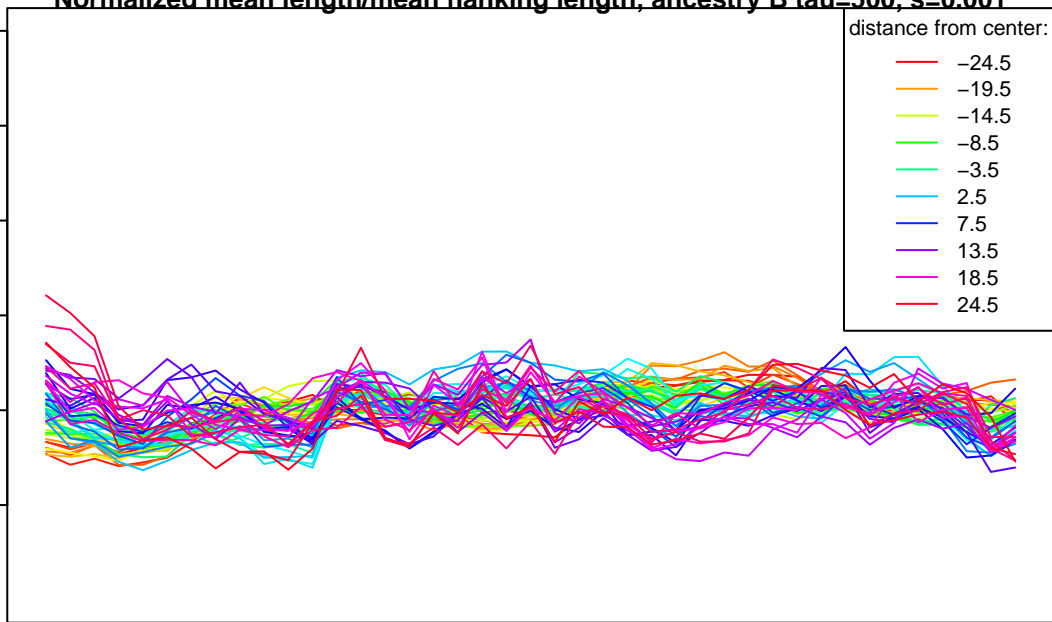
Normalized mean length/mean flanking length, ancestry B tau=500, s=0.001

Normalized mean block length

distance from center:

— -24.5
— -19.5
— -14.5
— -8.5
— -3.5
— 2.5
— 7.5
— 13.5
— 18.5
— 24.5

Distance from selected locus (M)



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

Normalized mean block length

distance from center:

— -24.5
— -19.5
— -14.5
— -8.5
— -3.5
— 2.5
— 7.5
— 13.5
— 18.5
— 24.5

Distance from selected locus (M)

