

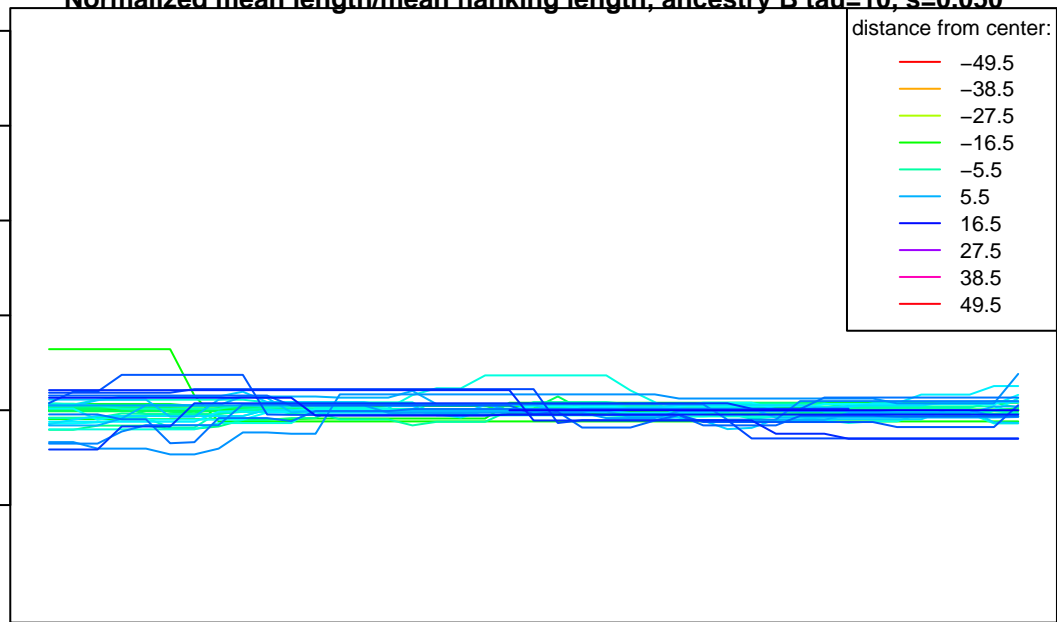
Normalized mean length/mean flanking length, ancestry B tau=10, s=0.050

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

Distance from selected locus (M)



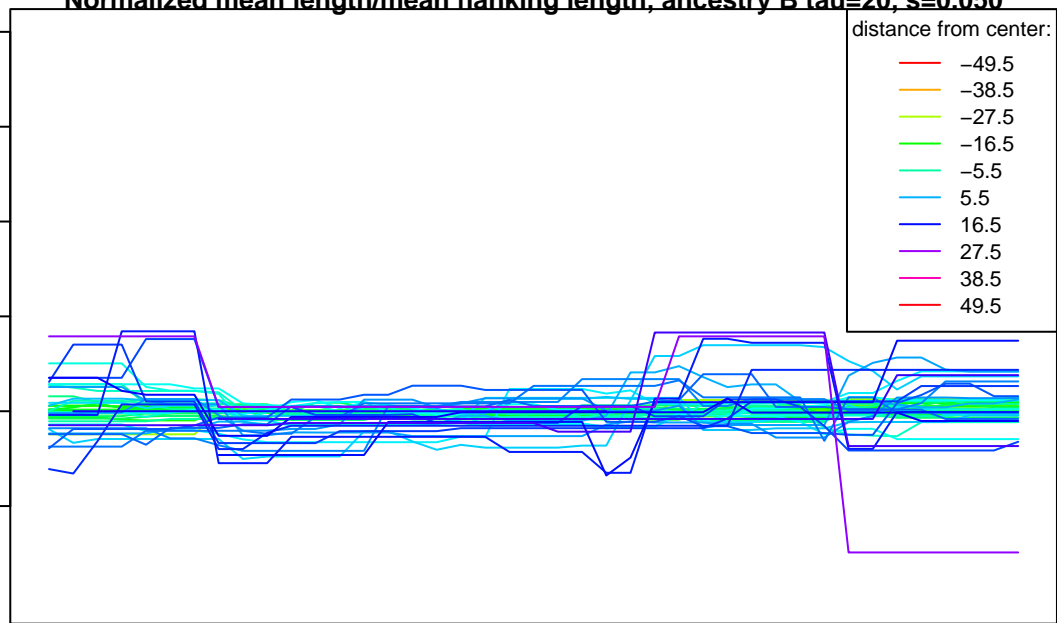
Normalized mean length/mean flanking length, ancestry B tau=20, s=0.050

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

Distance from selected locus (M)



Normalized mean length/mean flanking length, ancestry B $\tau=80$, $s=0.050$

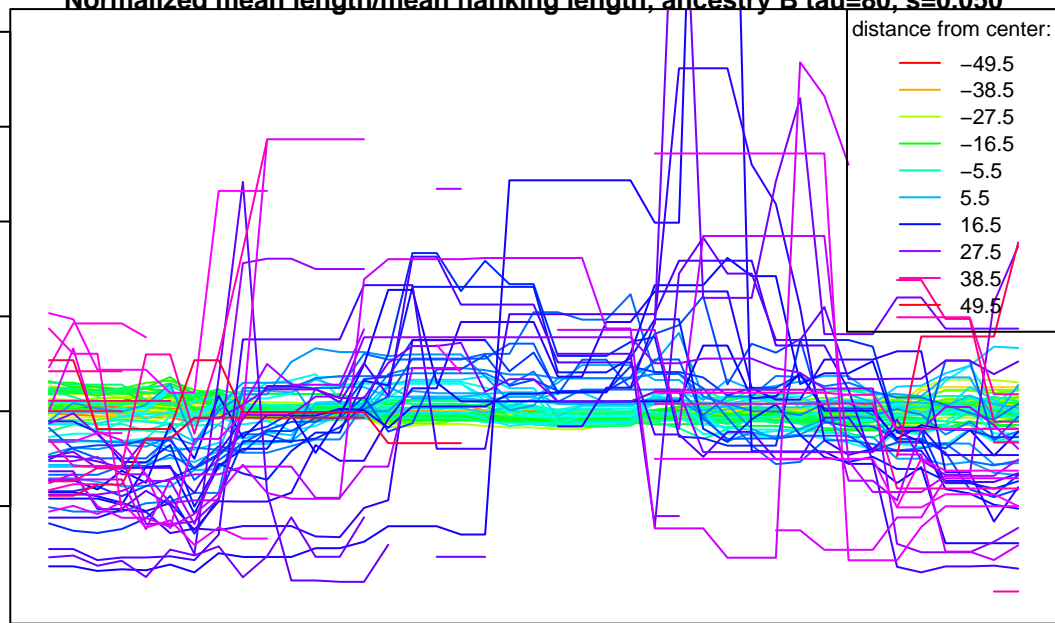
Normalized mean block length

3.0
2.5
2.0
1.5
1.0
0.5

distance from center:

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

Distance from selected locus (M)



Normalized mean length/mean flanking length, ancestry B $\tau=160$, $s=0.050$

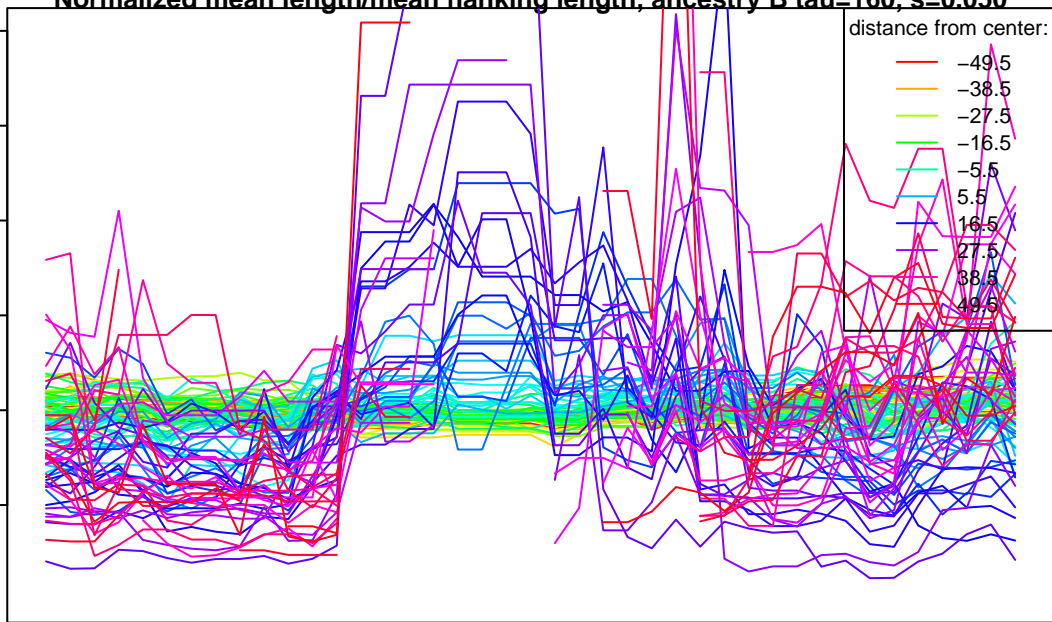
Normalized mean block length

3.0
2.5
2.0
1.5
1.0
0.5

distance from center:

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

Distance from selected locus (M)



Normalized mean length/mean flanking length, ancestry B tau=320, s=0.050

Normalized mean block length

3.0
2.5
2.0
1.5
1.0
0.5

distance from center:

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

Distance from selected locus (M)

