

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

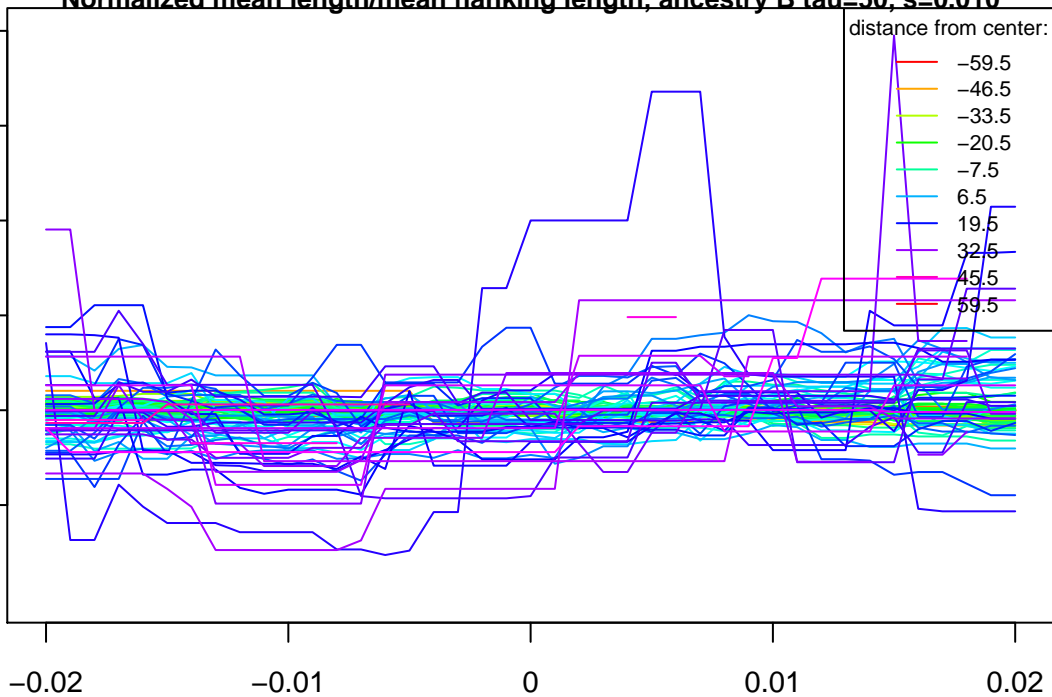
-0.01

0

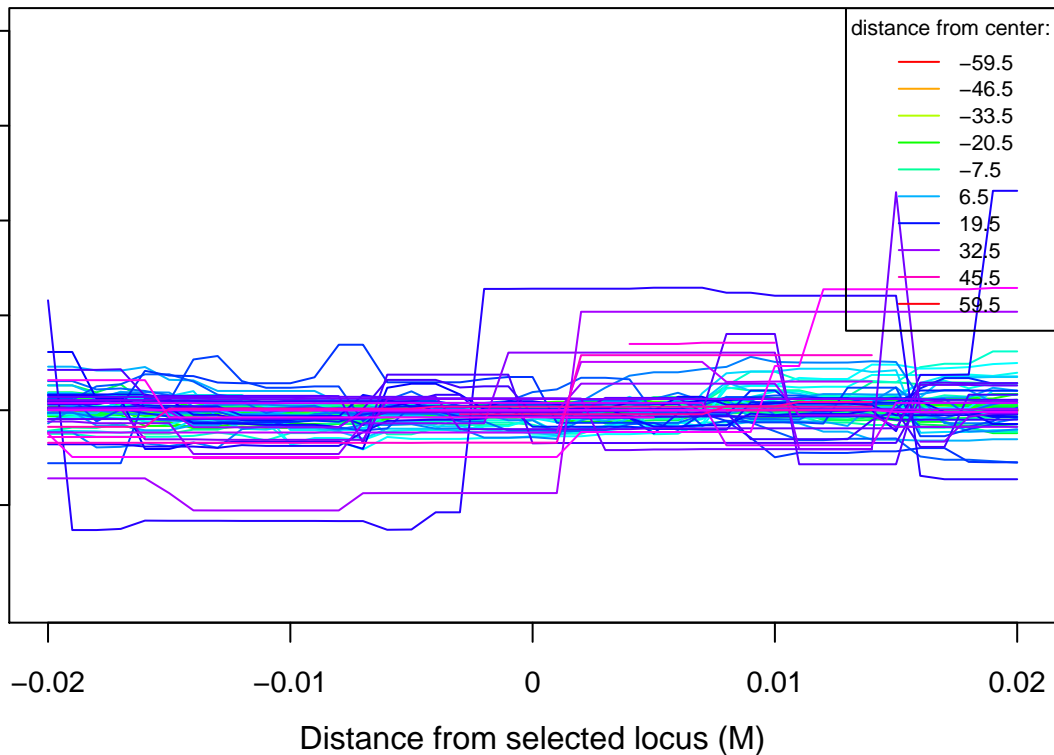
0.01

0.02

Distance from selected locus (M)



Normalized mean block length



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

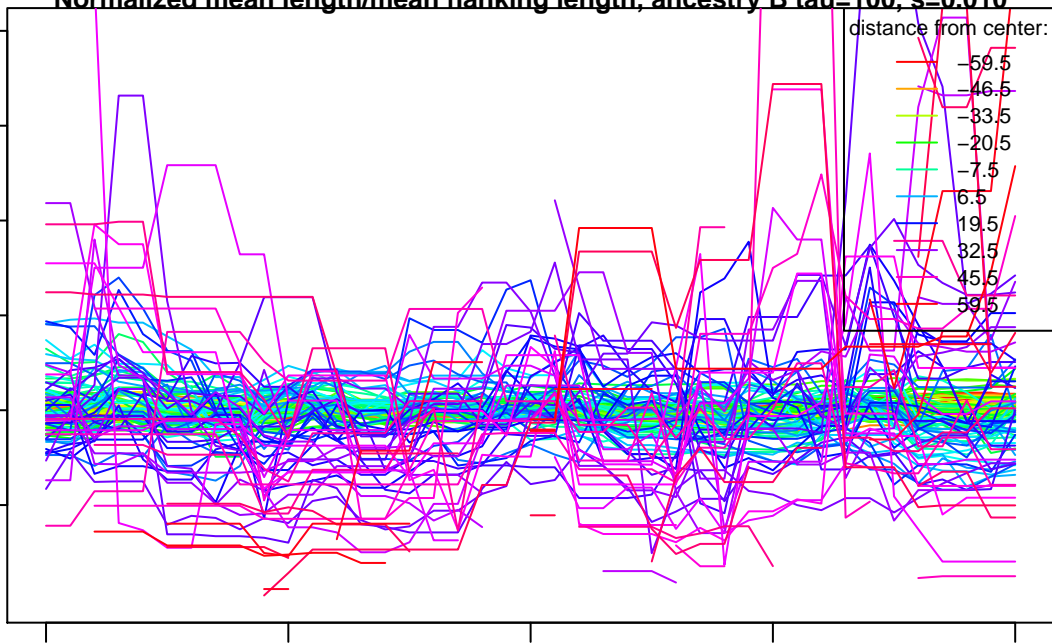
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0

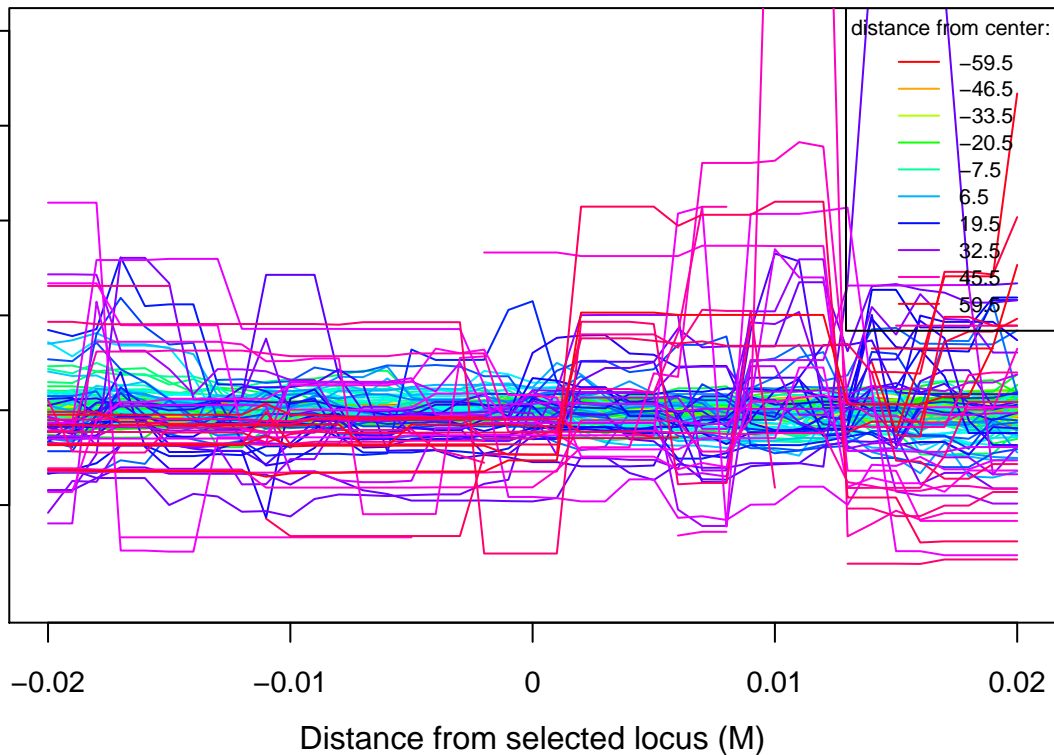
0.01

0.02

Distance from selected locus (M)

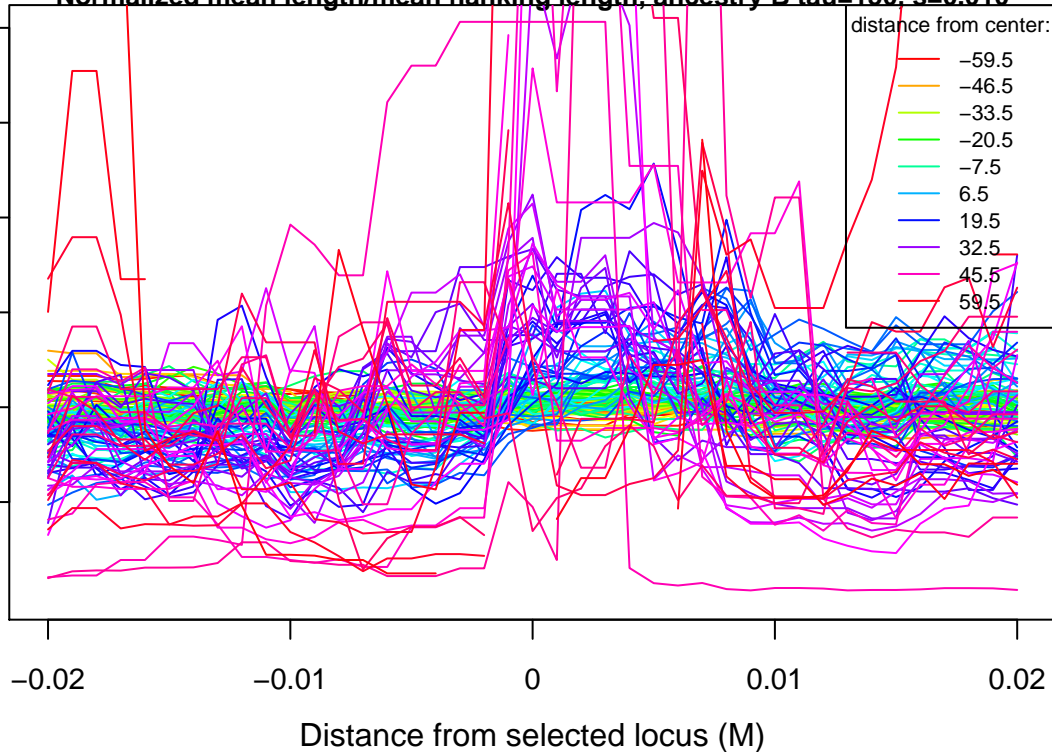


Normalized mean block length

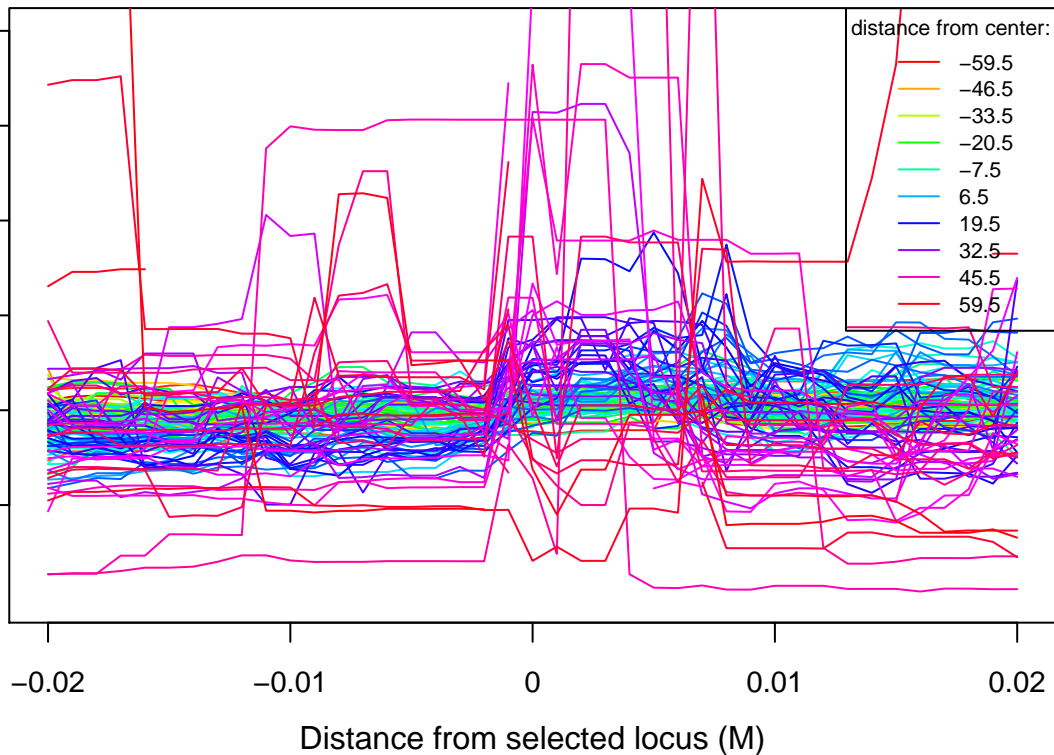


Normalized mean length/mean flanking length, ancestry B tau=150, s=0.010

Normalized mean block length



Normalized mean block length



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

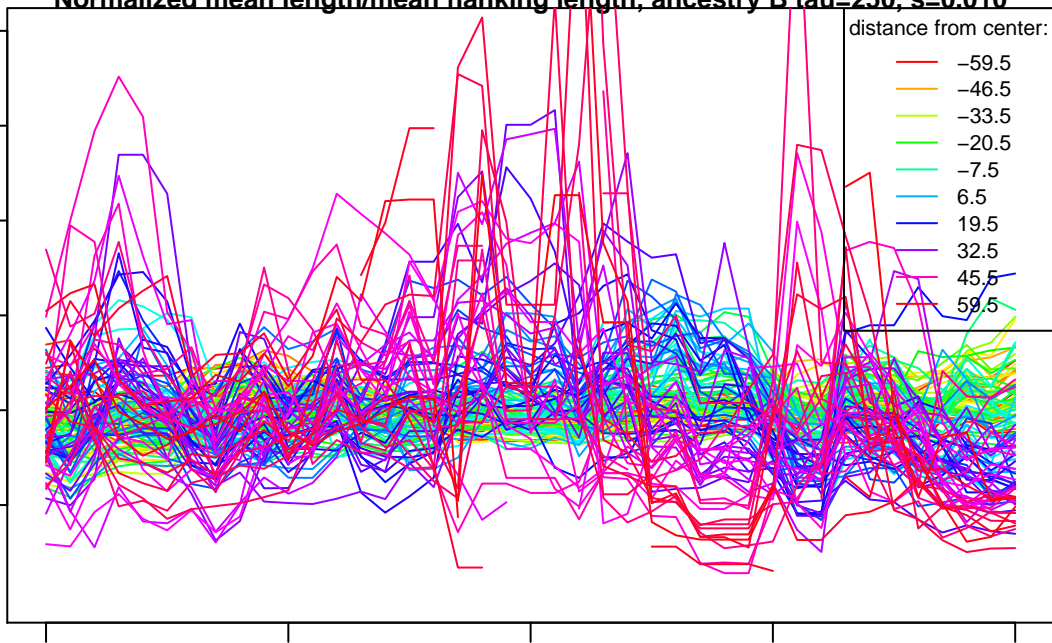
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0

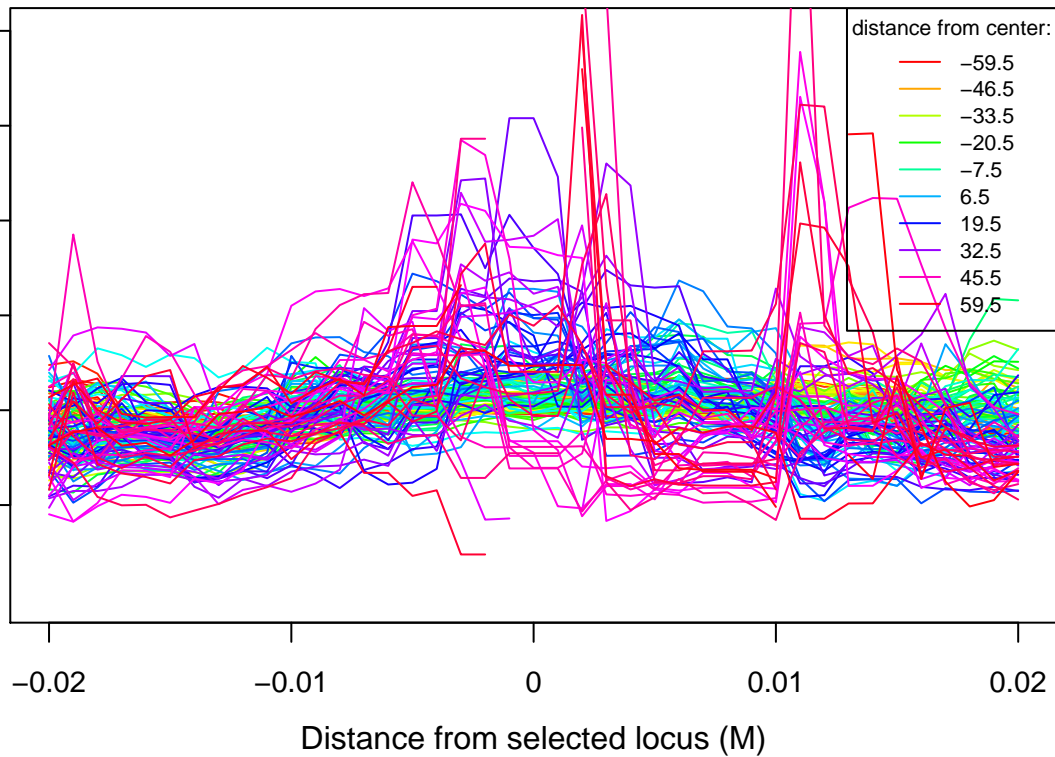
0.01

0.02

Distance from selected locus (M)



Normalized mean block length



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

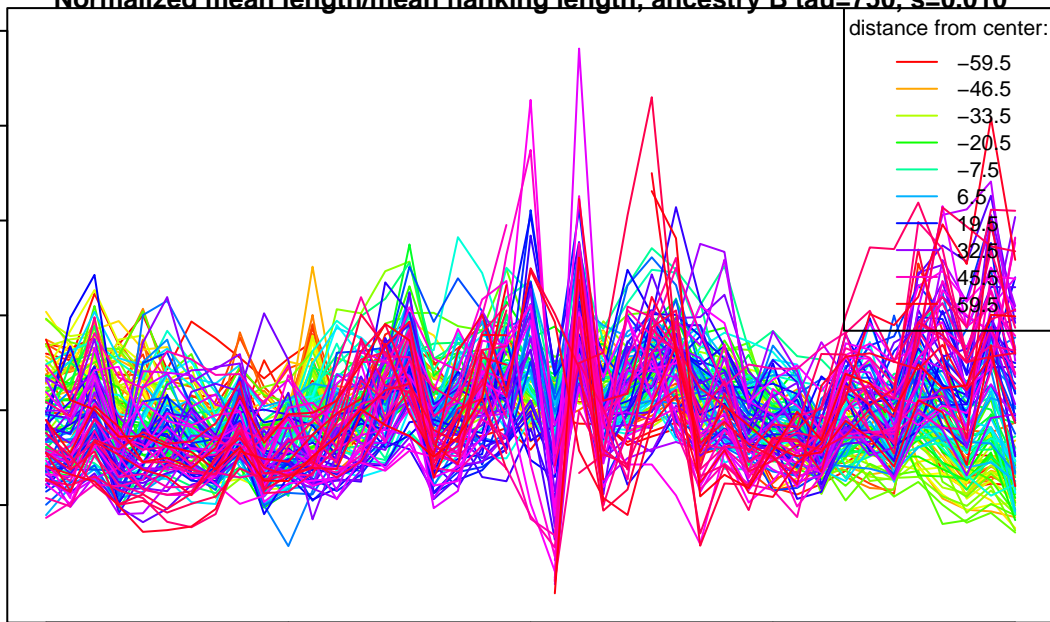
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0

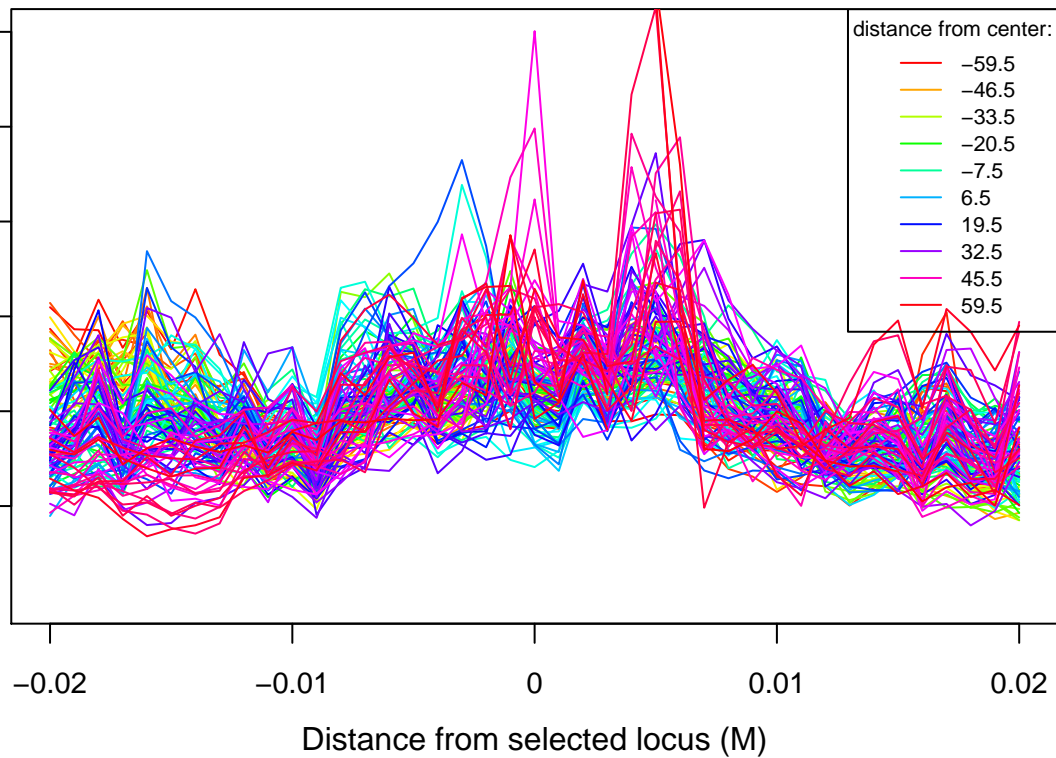
0.01

0.02

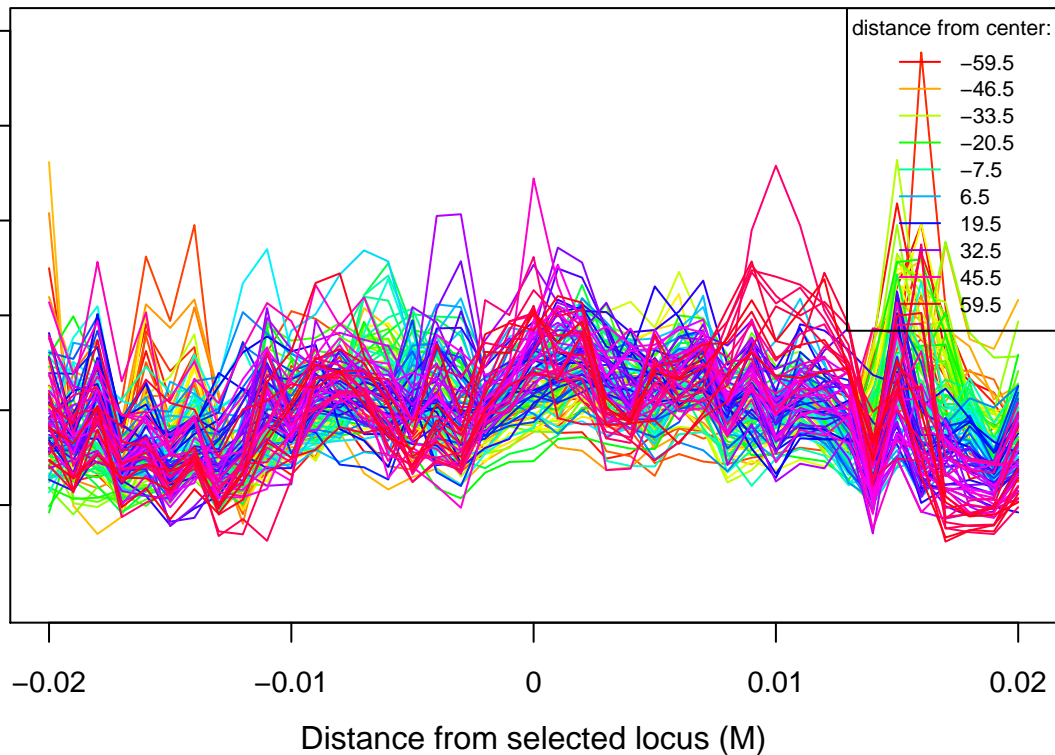
Distance from selected locus (M)



Normalized mean block length



Normalized mean block length



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)

