Functional Mapping Report

Data File:

Report Date: 2018−08−08 02:39:26

1 Data Summary

Cross:	BC
Curve:	Legendre2
Covariance:	AR1
Pheno. file:	/home/zw355/R/x86_64-unknown-linux-gnu-library/3.1/Funmap2/extdata/populus.BC.pheno.c:
Time. file:	
Geno. file:	/home/zw355/R/x86_64-unknown-linux-gnu-library/3.1/Funmap2/extdata/populus.BC.geno.csv
Marker file:	/home/zw355/R/x86_64-unknown-linux-gnu-library/3.1/Funmap2/extdata/populus.BC.marker.
Sample size:	78
Sample times:	11
Marker count:	275
Chr/Group count:	22
Phenotype missing:	0
Marker missing:	4.66200466200466e-05
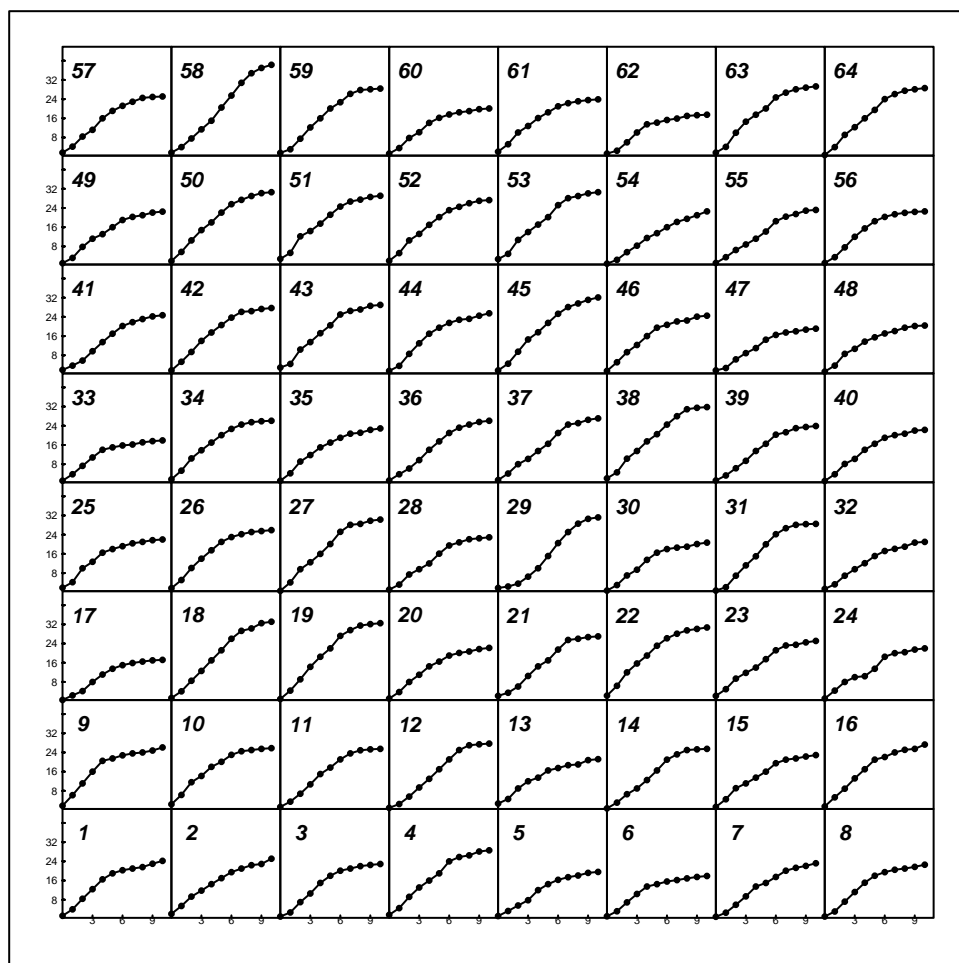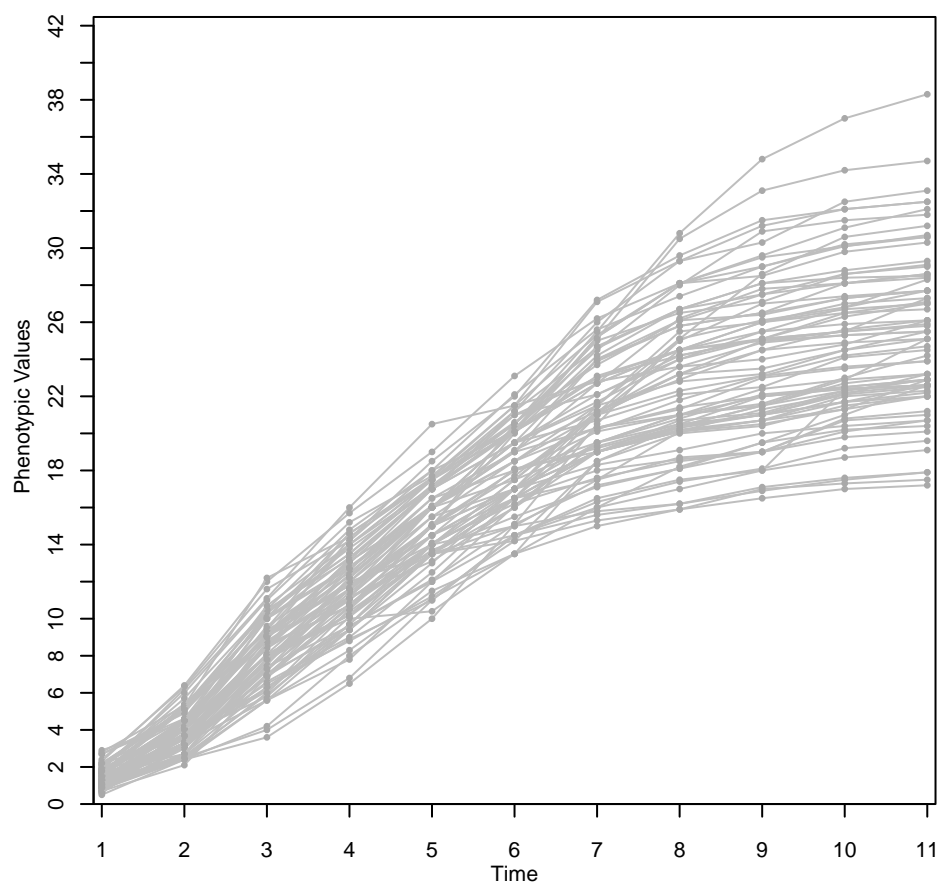
2 QTL Profile
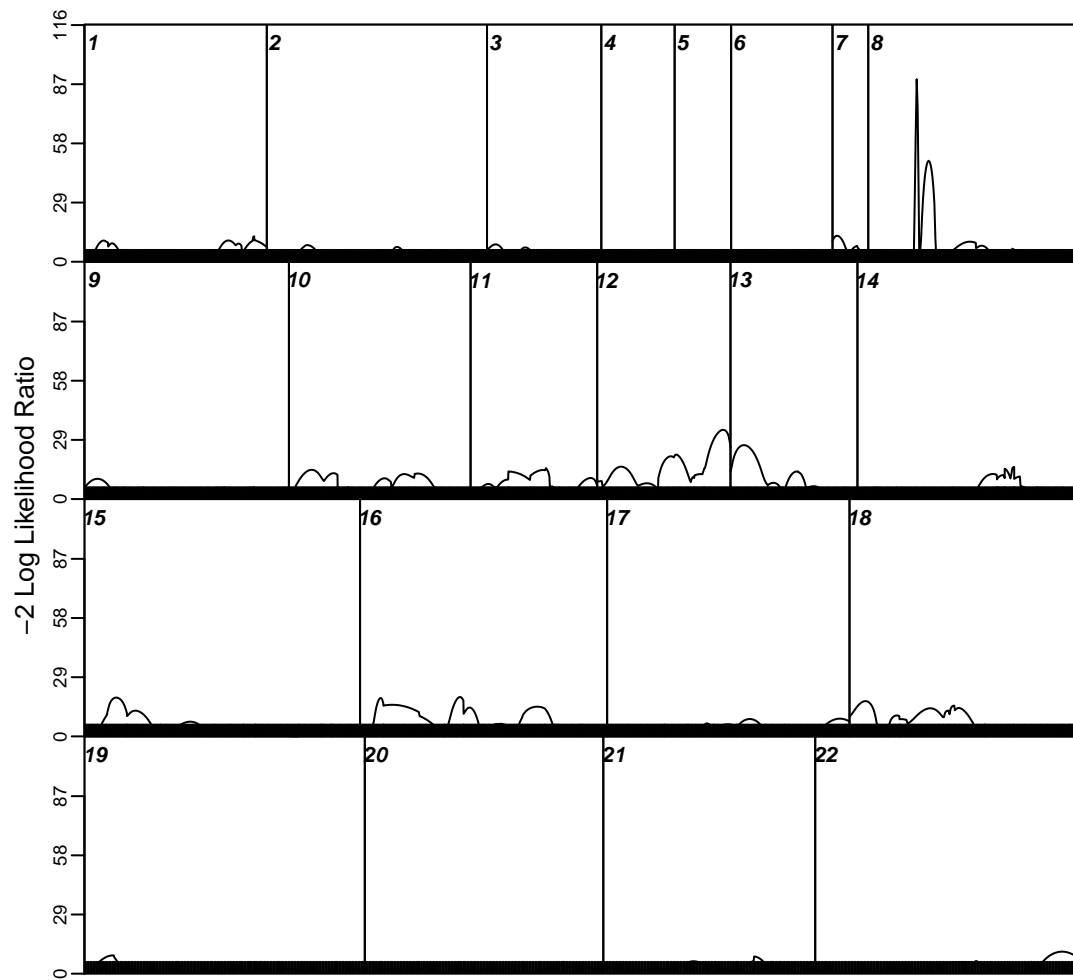
3 QTL Position

3.1 Groupp=8, Postion=51.10
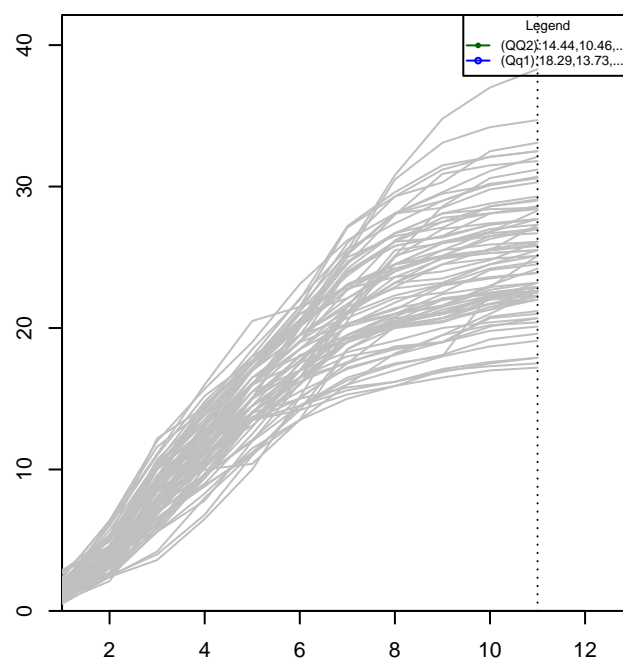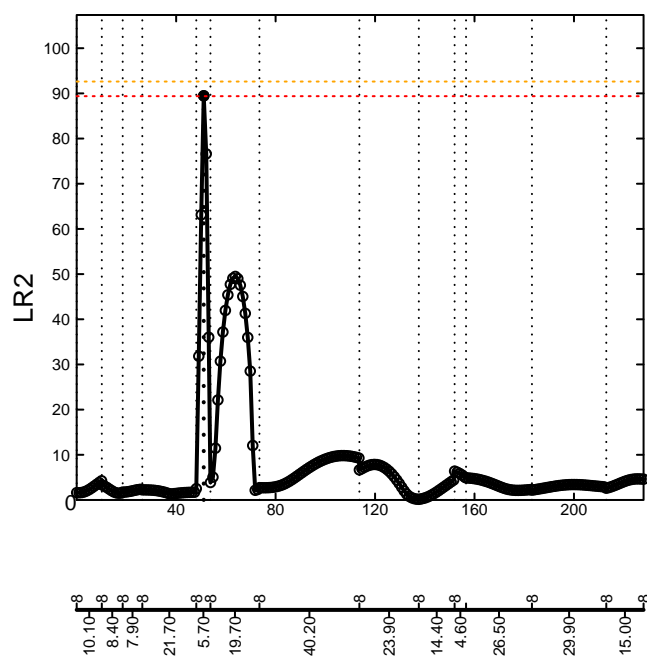
Permutation Cutoff

pv	0.1	0.05	0.01	0.005	0.001	0.0001	0.00001
Cutoff	88.46	89.38	92.62				

Permutation result

Cutoffs
0.050000: 89.38
0.010000: 92.62