

1 Data Summary

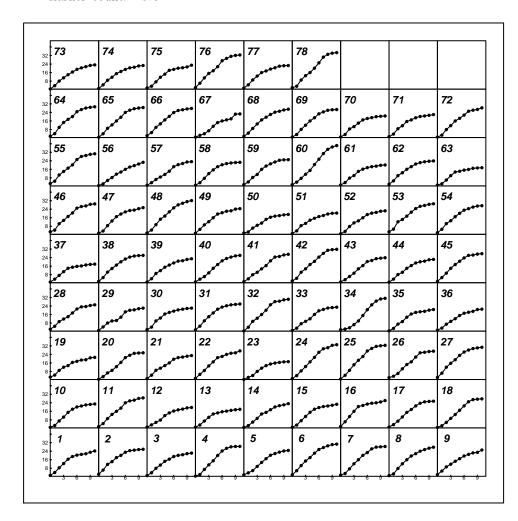
Curve: LC
Cross: BC
Covariance: AR1

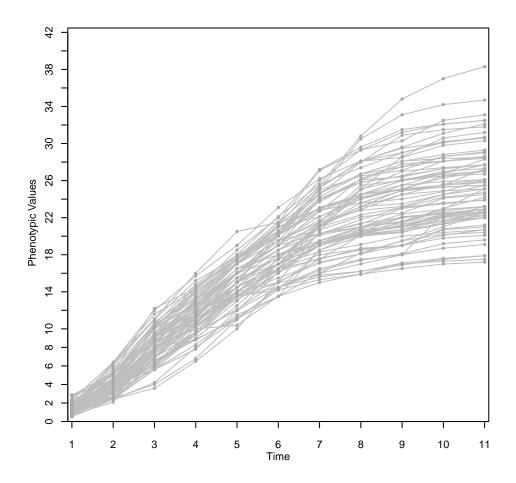
Pheno. file: LC.populus.pheno.csv

Geno. file: LC.populus.geno.csv

Marker file: LC.populus.marker.csv

Sample size: 78
Sample times: 11
Marker count: 275





2 QTL Profile

LR profile

Grp.	Pos.	LR	rho	Sig^2 QQ	Qq	qq
8	50.10	75.05	0.94	2.66	23.19,9.83,0.62,	30.80,11.25,0.54,
12	113.10	59.33	0.96	3.35	24.86,8.78,0.59,	30.19,10.07,0.53,
13	12.00	53.69	0.96	3.42	24.75,8.95,0.61,	29.66,9.89,0.53,
7	4.00	30.75	0.96	3.40	24.93,8.95,0.59,	29.86,10.00,0.53,
18	10.00	27.46	0.96	3.50	25.27,9.02,0.59,	29.40,9.64,0.54,

