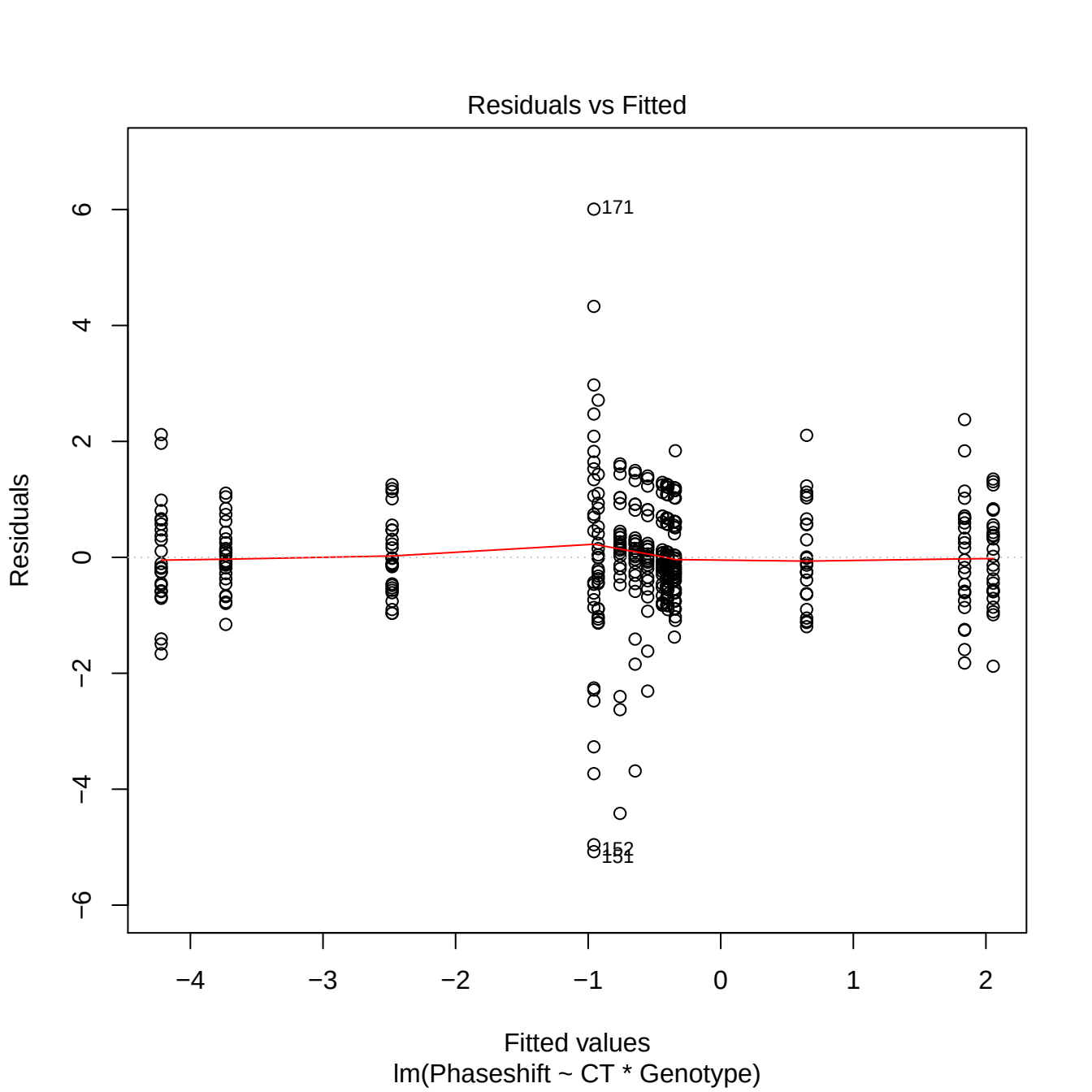
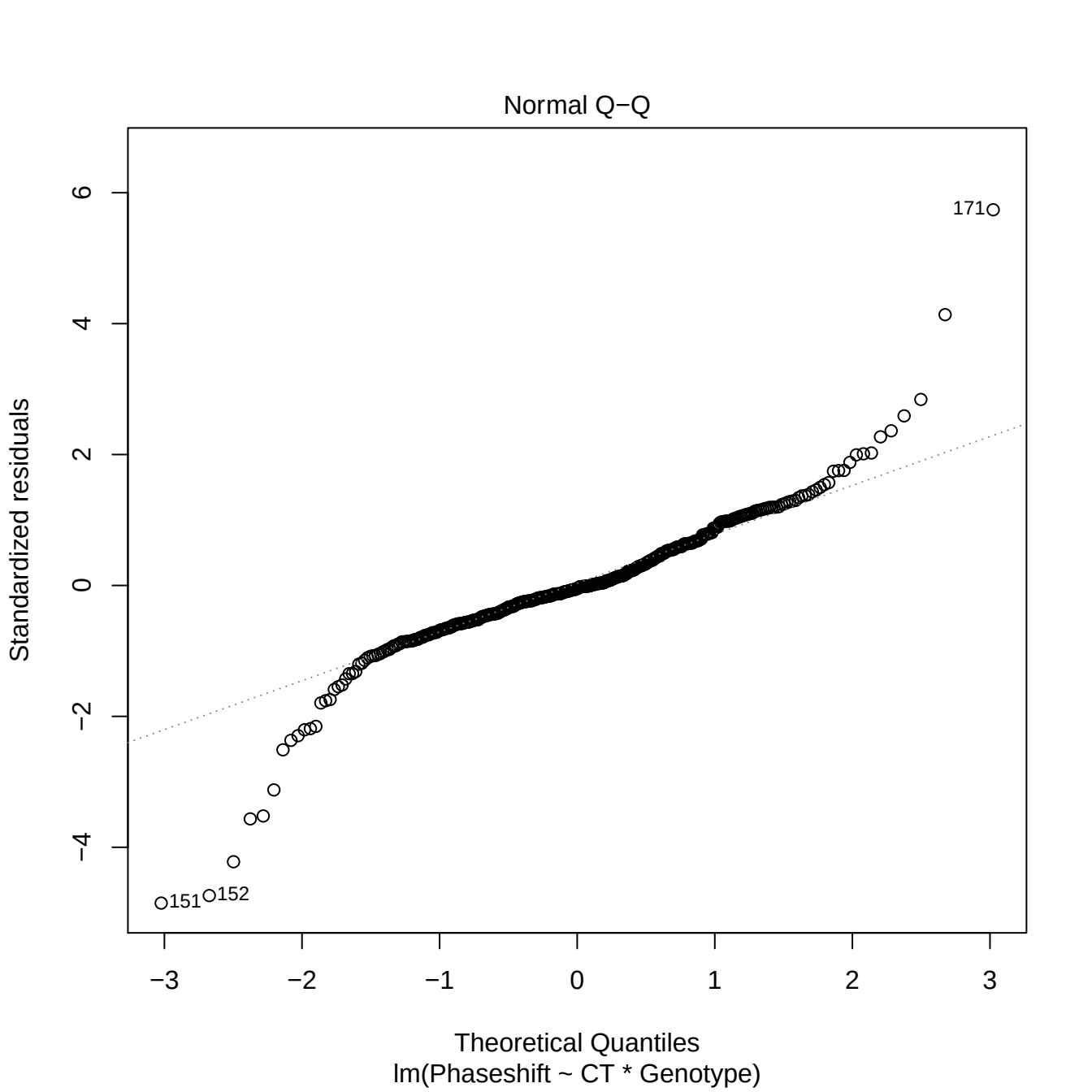
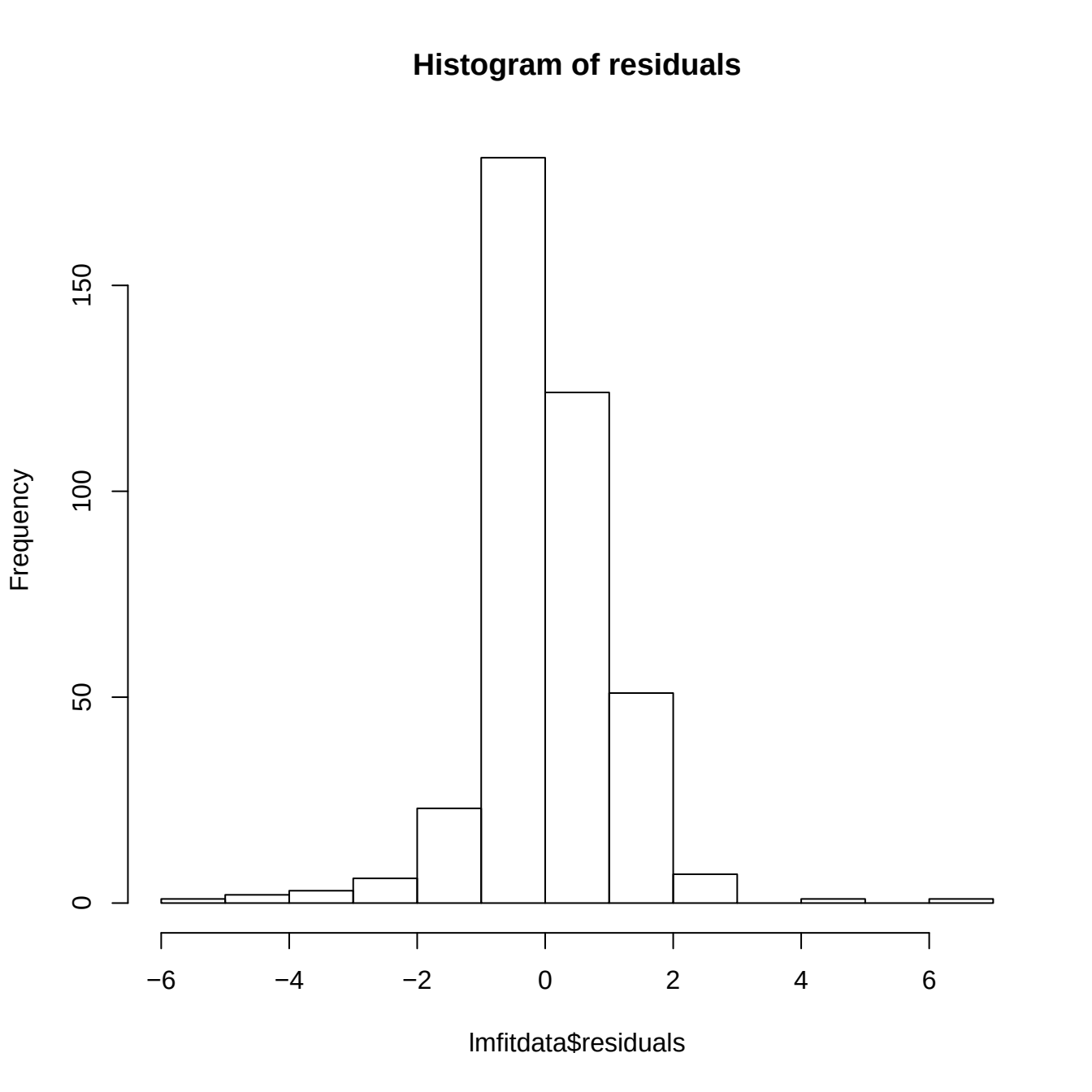

Residuals vs Fitted
Residuals
Fitted values
lm(Phaseshift ~ CT * Genotype)
171
152
151
6
4
2
0
−2
−4
−6
−4
−3
−2
−1
0
1
2


Normal Q−Q
Standardized residuals
Theoretical Quantiles
lm(Phaseshift ~ CT * Genotype)
171
151
152
6
4
2
0
−2
−4
−3
−2
−1
0
1
2
3


Histogram of residuals
Frequency
0
50
100
150
−6
−4
−2
0
2
4
6
lmfitdata$residuals