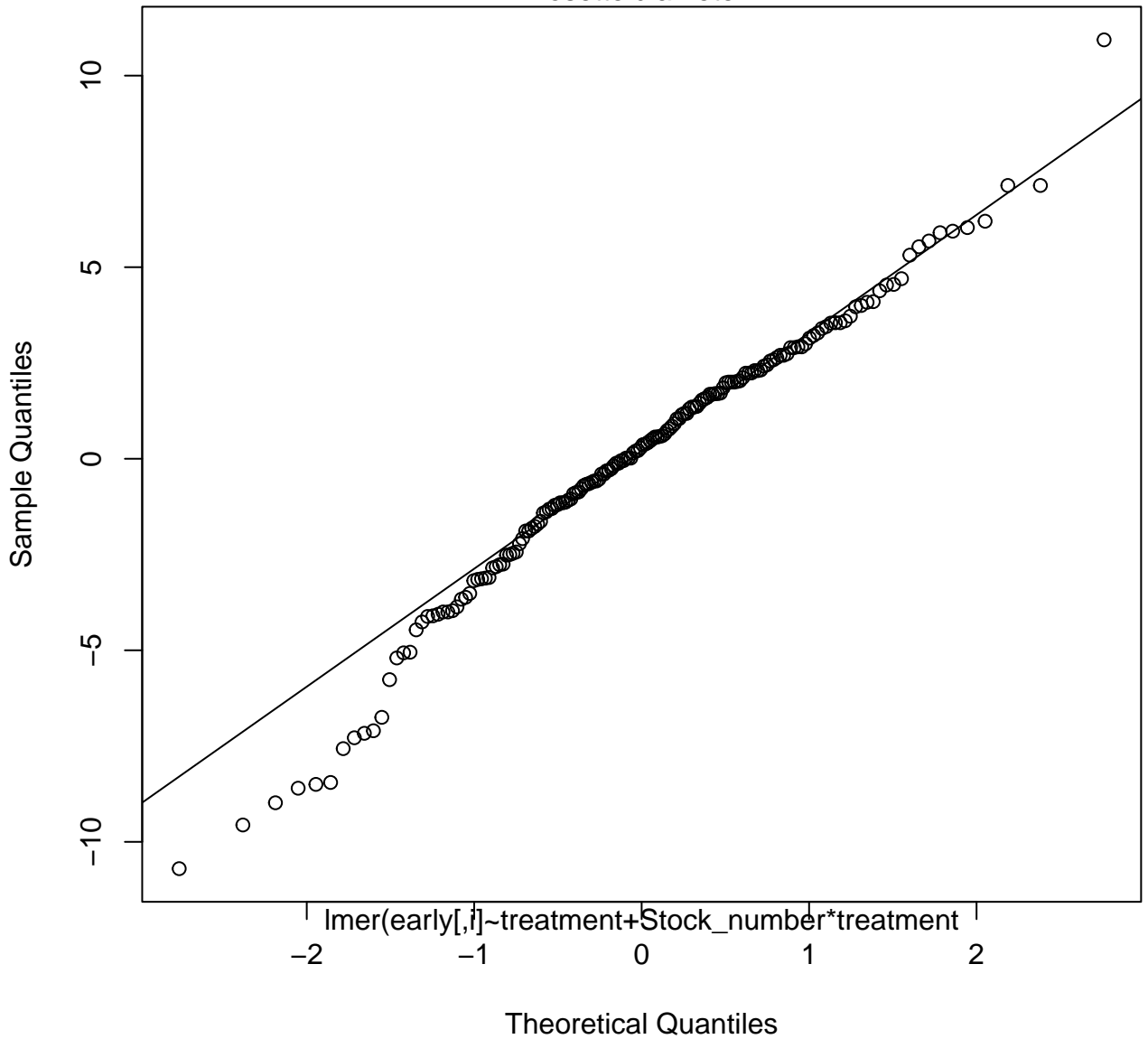
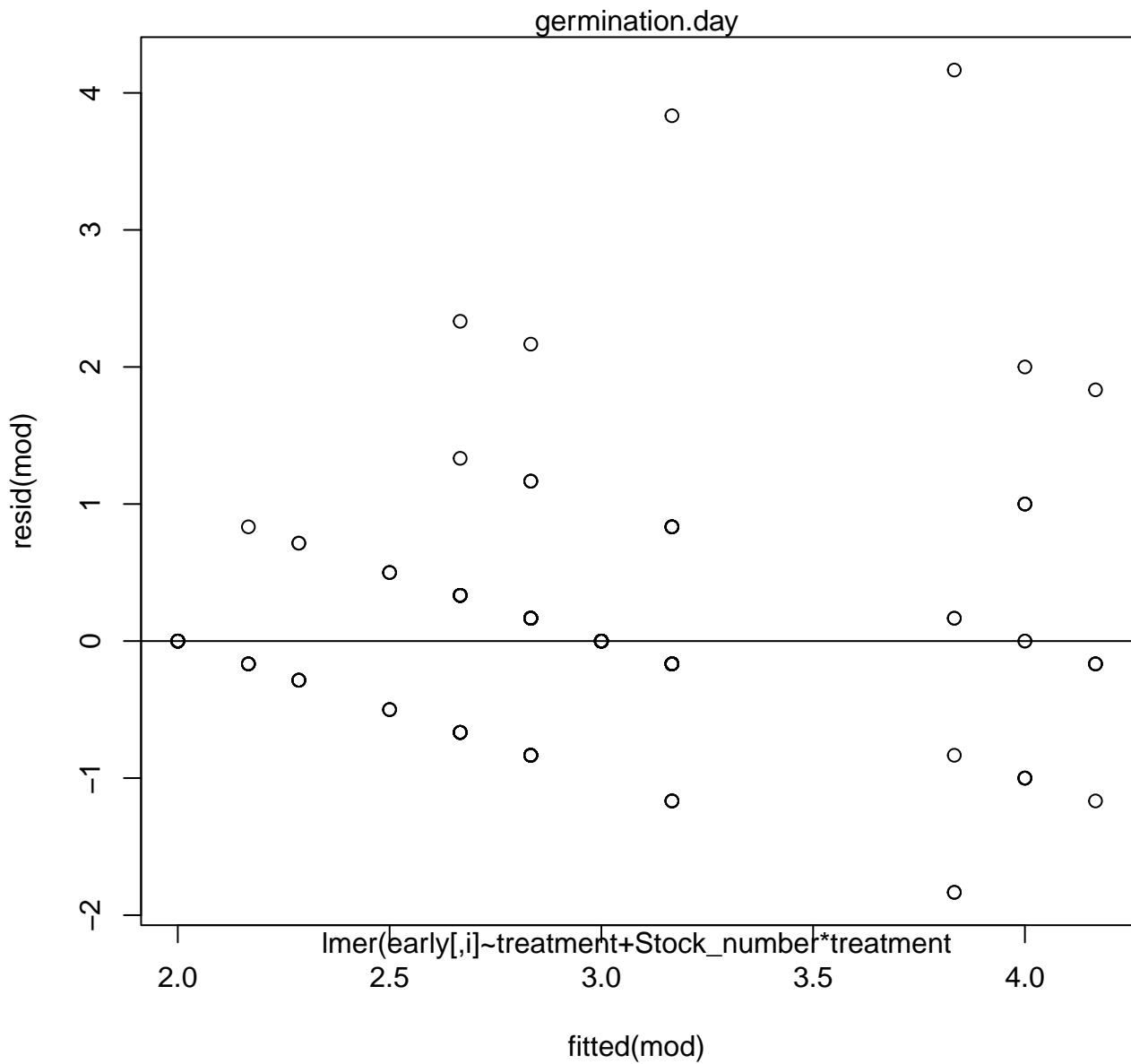


Normal Q-Q Plot

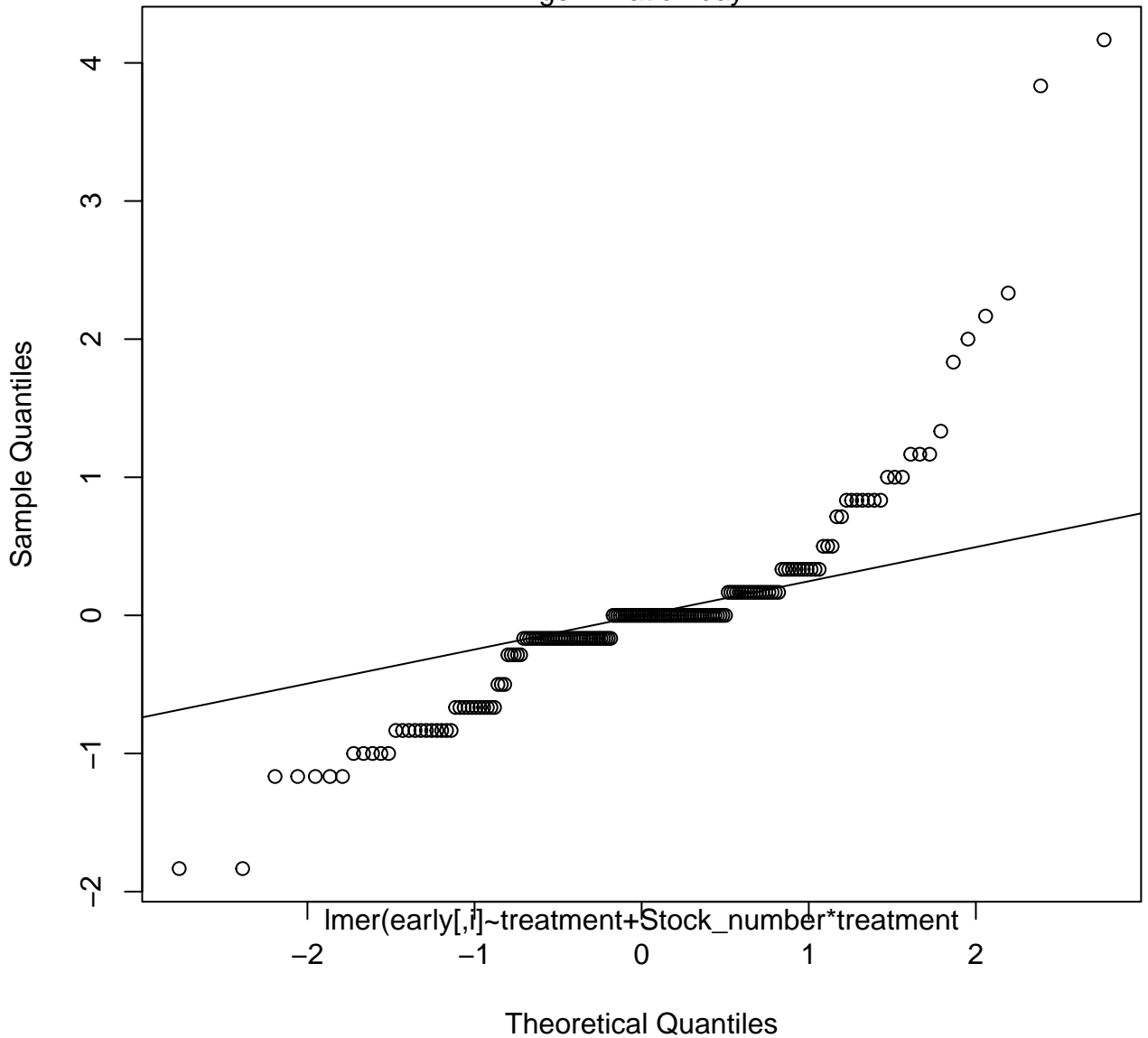
rosette.diameter

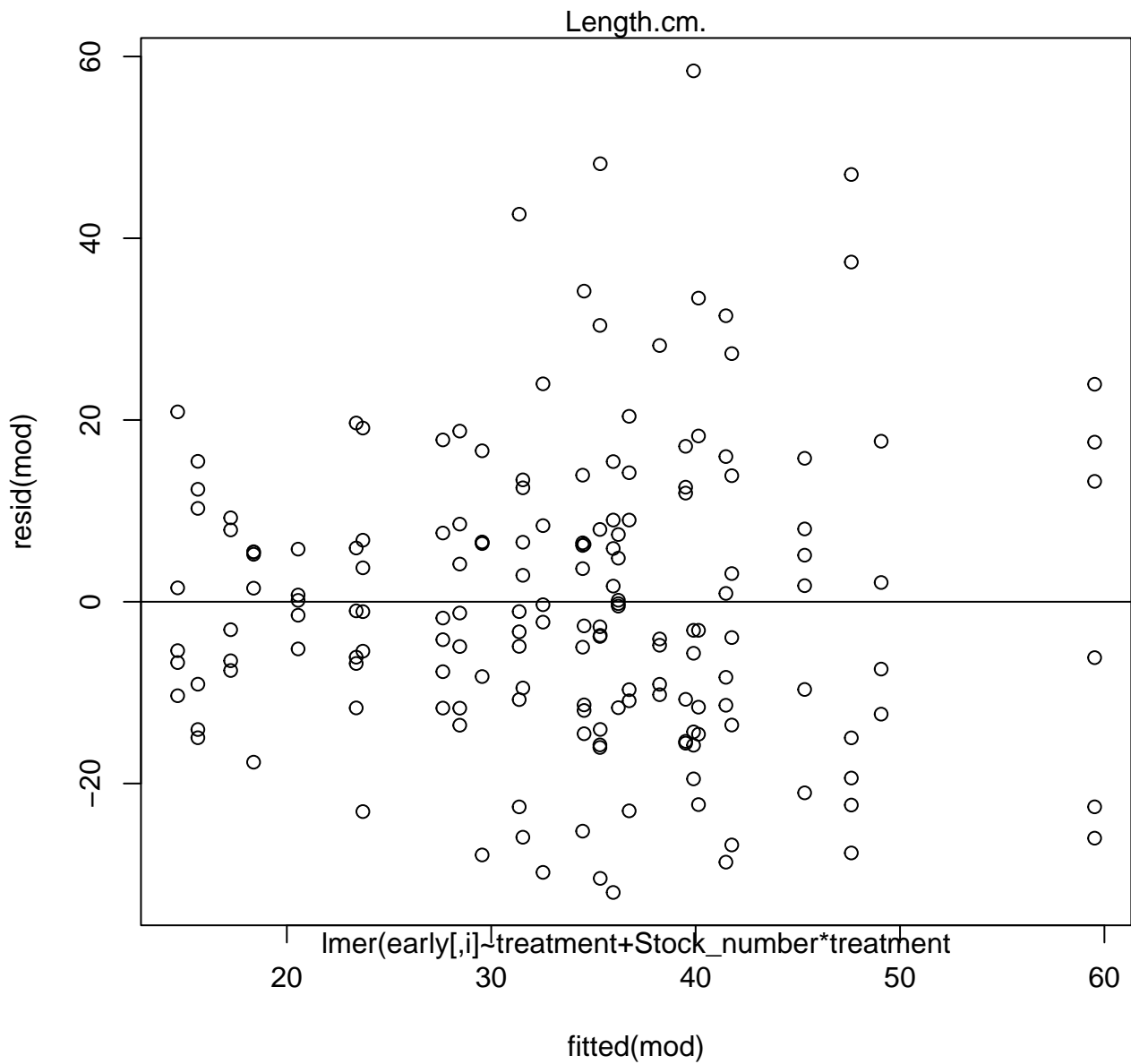




Normal Q-Q Plot

germination.day

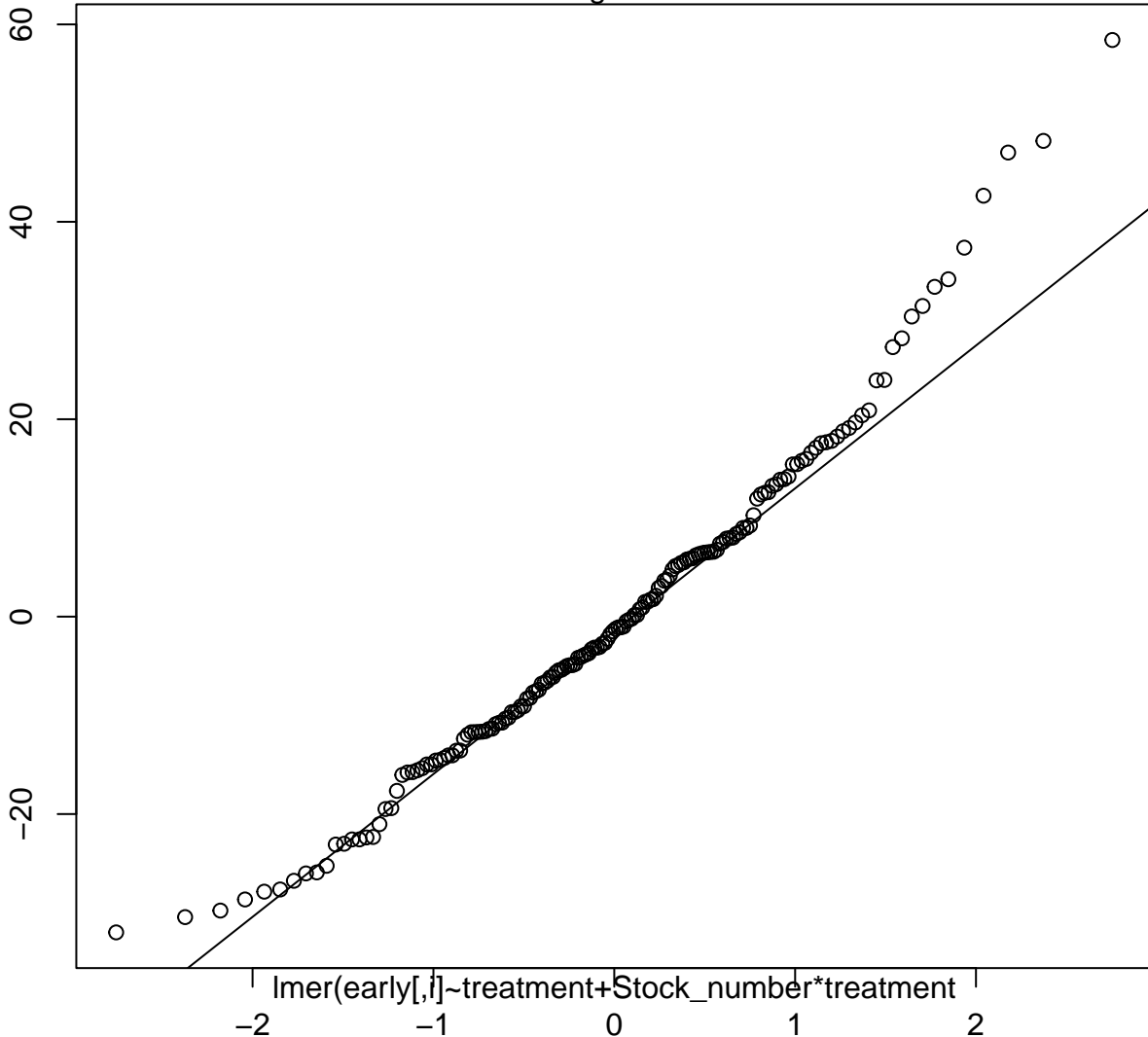




Normal Q-Q Plot

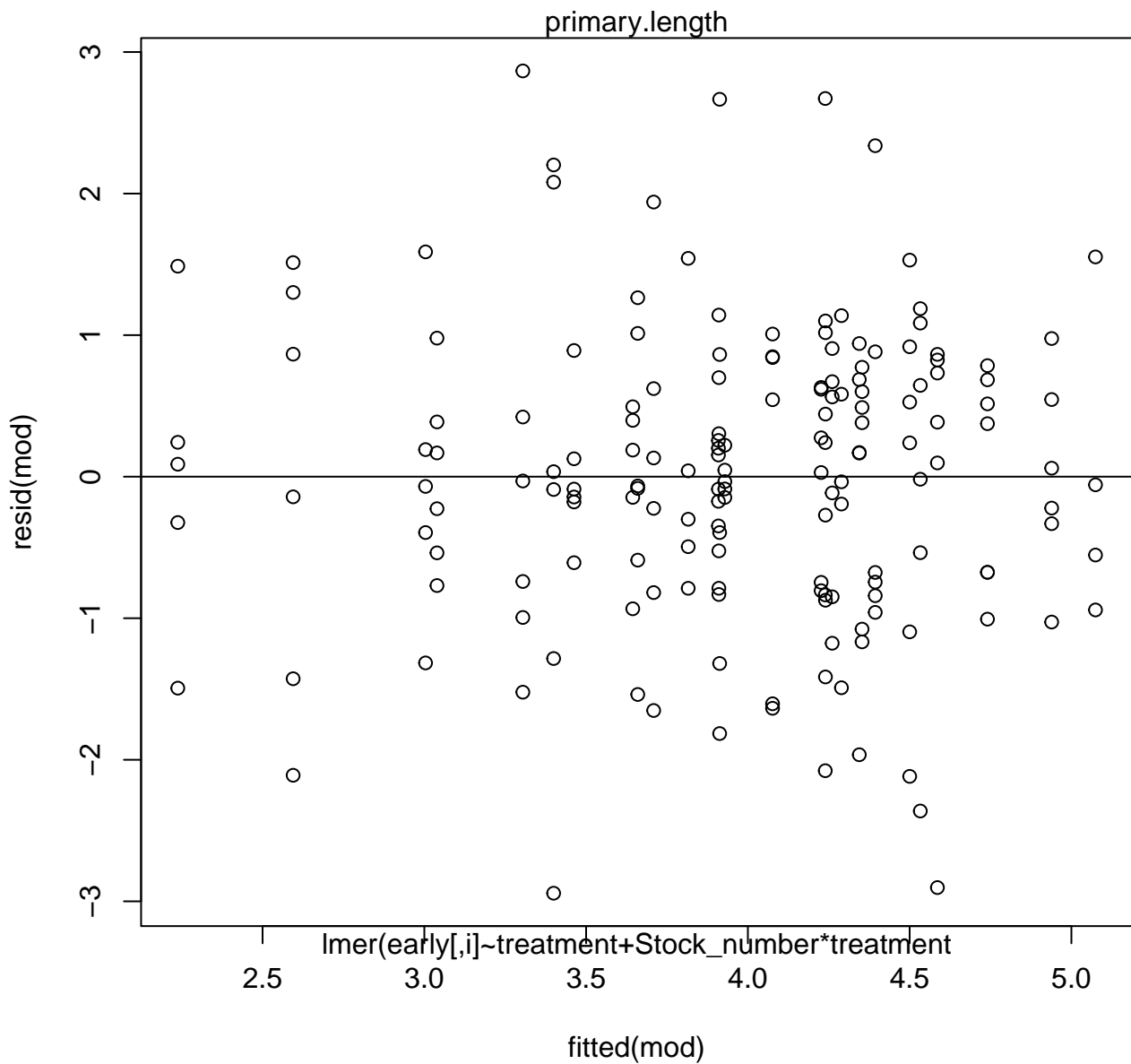
Length.cm.

Sample Quantiles



lmer(early[,i]~treatment+Stock_number*treatment

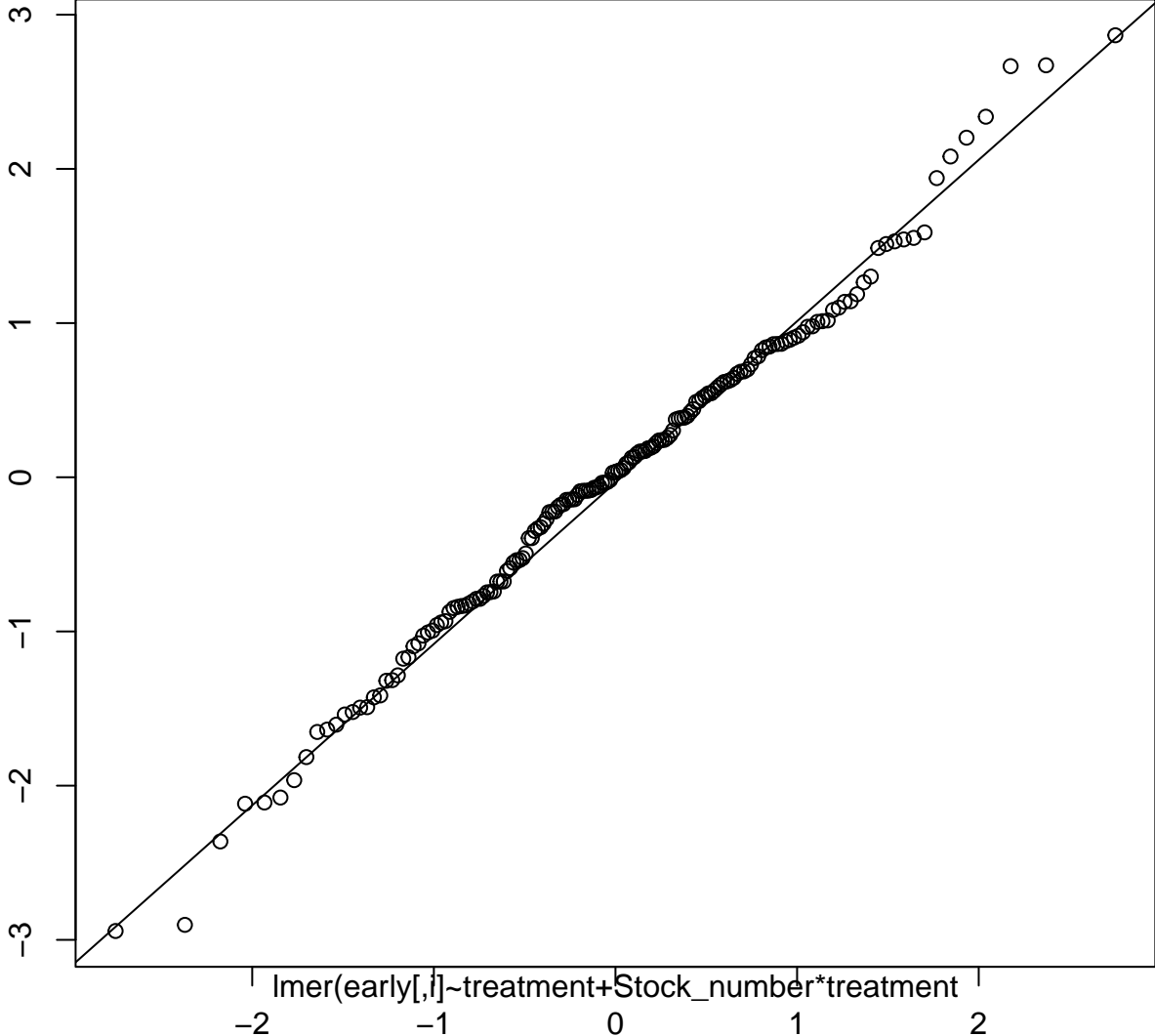
Theoretical Quantiles



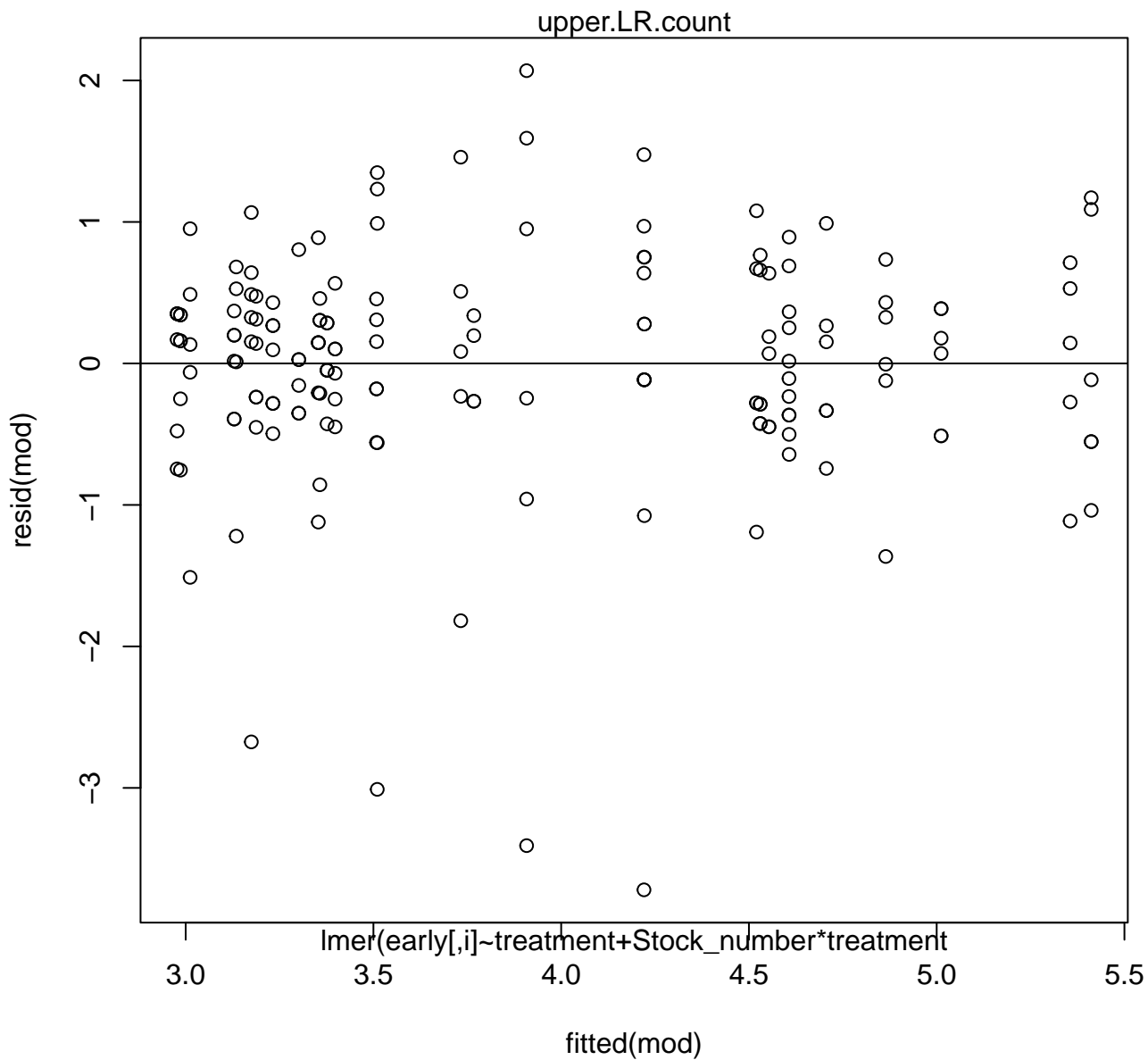
Normal Q-Q Plot

primary.length

Sample Quantiles

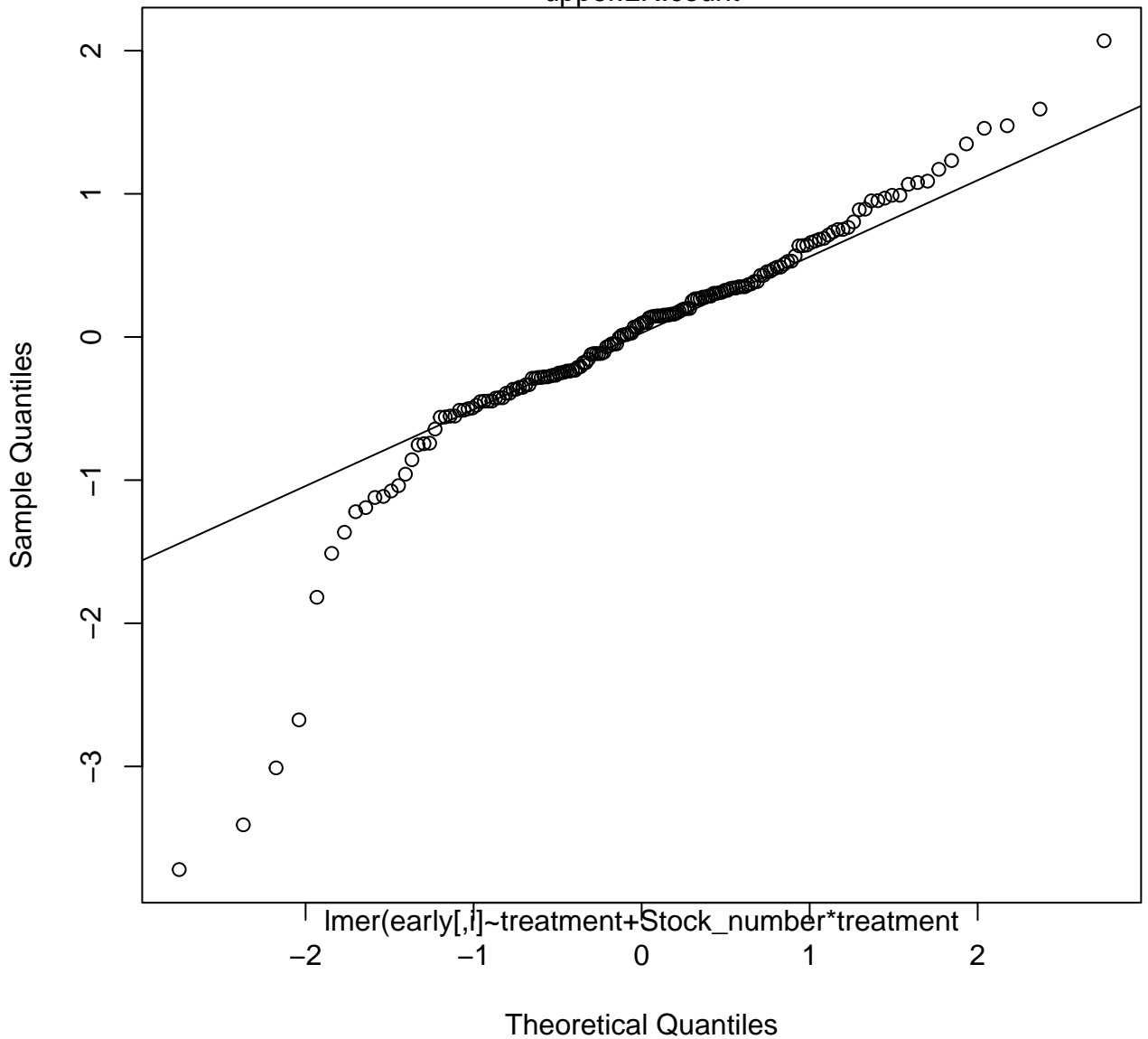


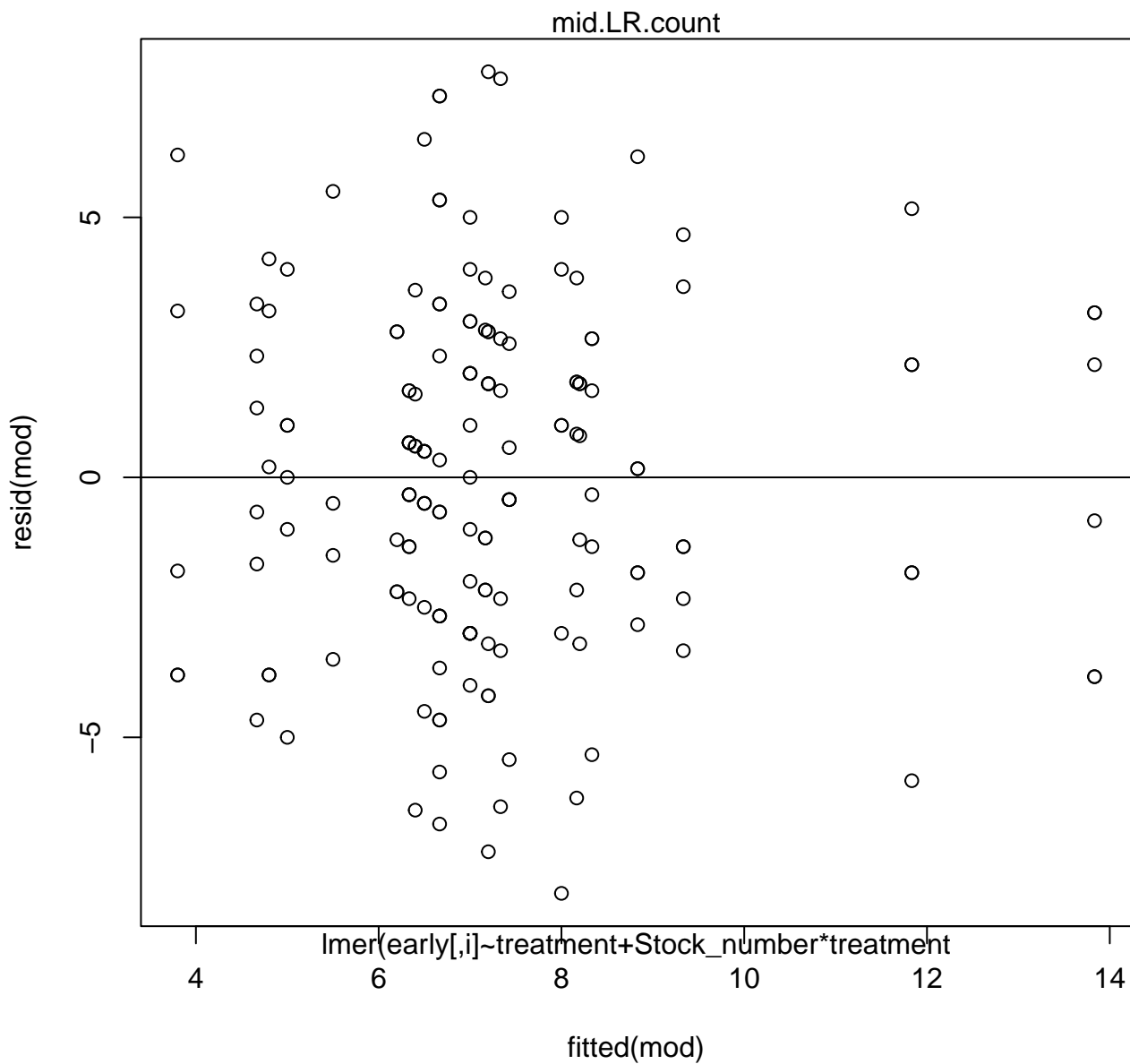
Theoretical Quantiles



Normal Q-Q Plot

upper.LR.count





Normal Q-Q Plot

mid.LR.count

Sample Quantiles

5

0

-5

-2

-1

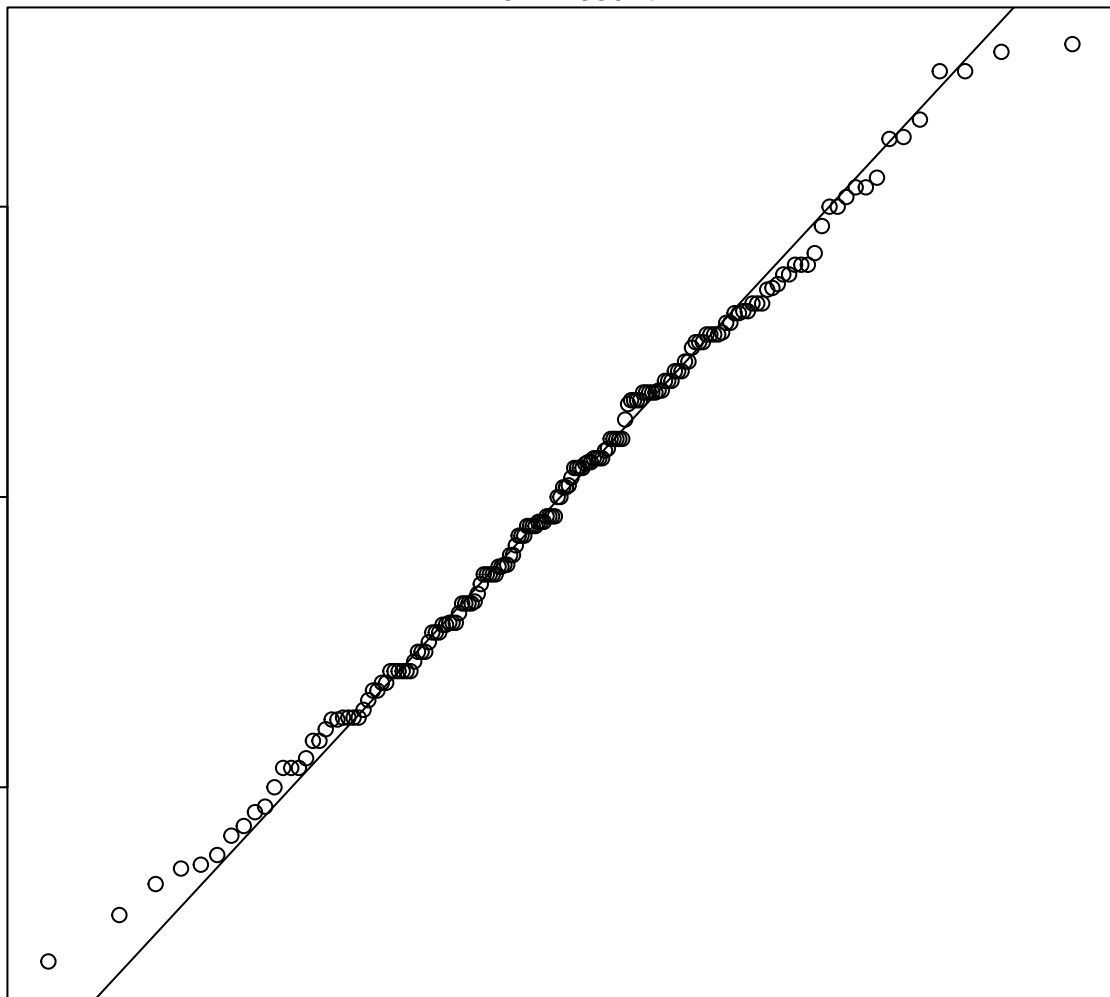
0

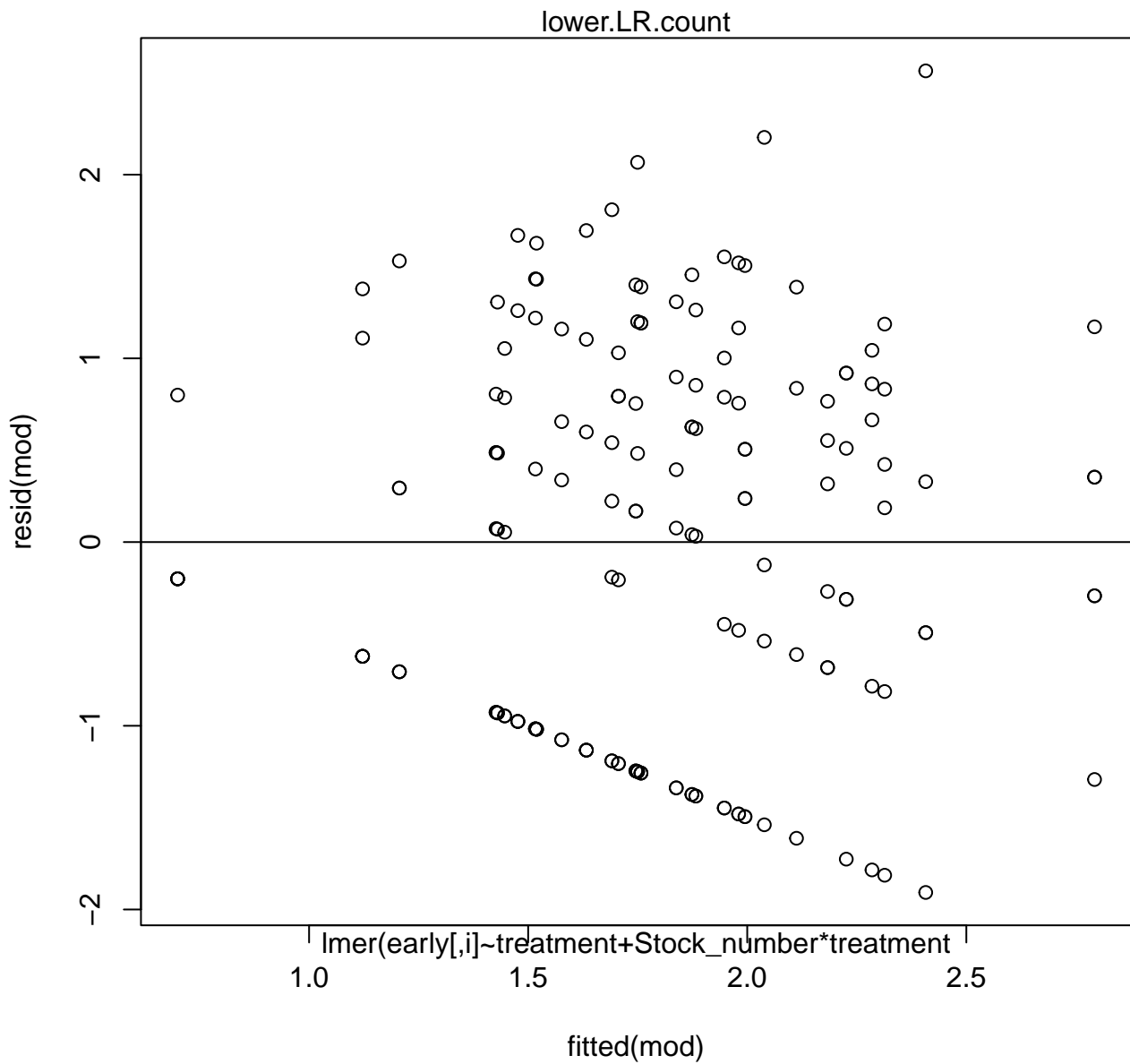
1

2

lmer(early[,i]~treatment+Stock_number*treatment

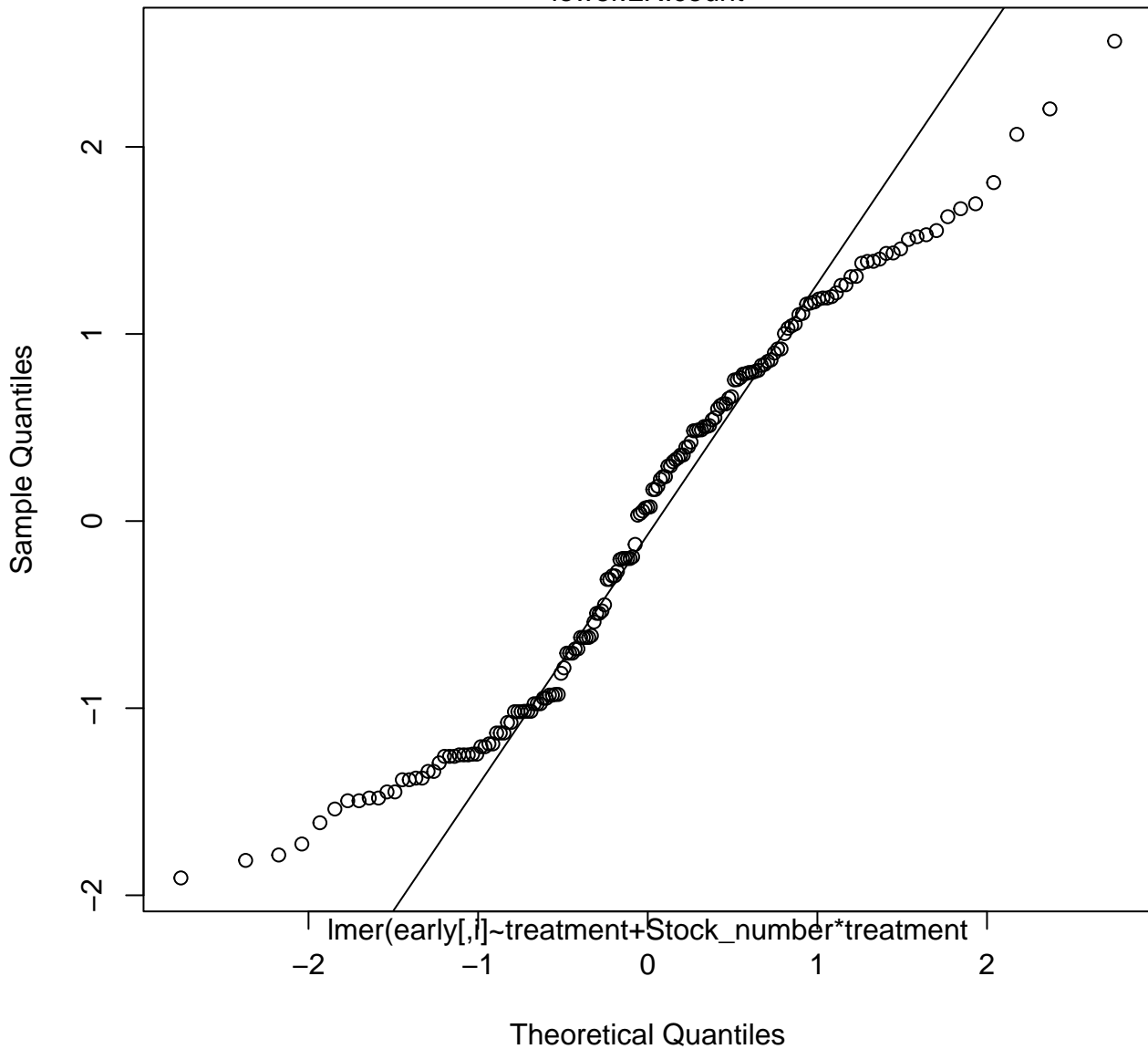
Theoretical Quantiles

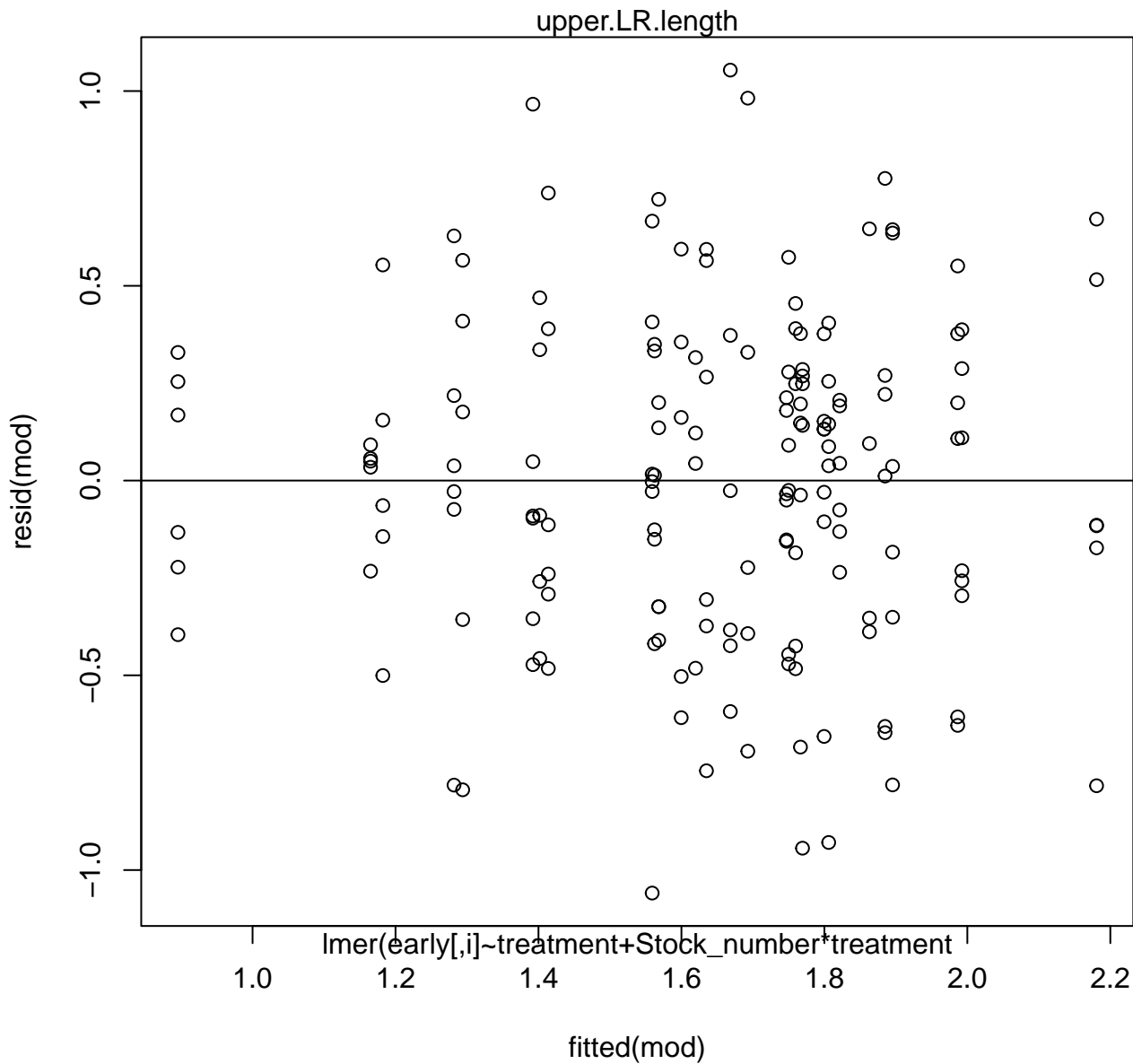




Normal Q-Q Plot

lower.LR.count

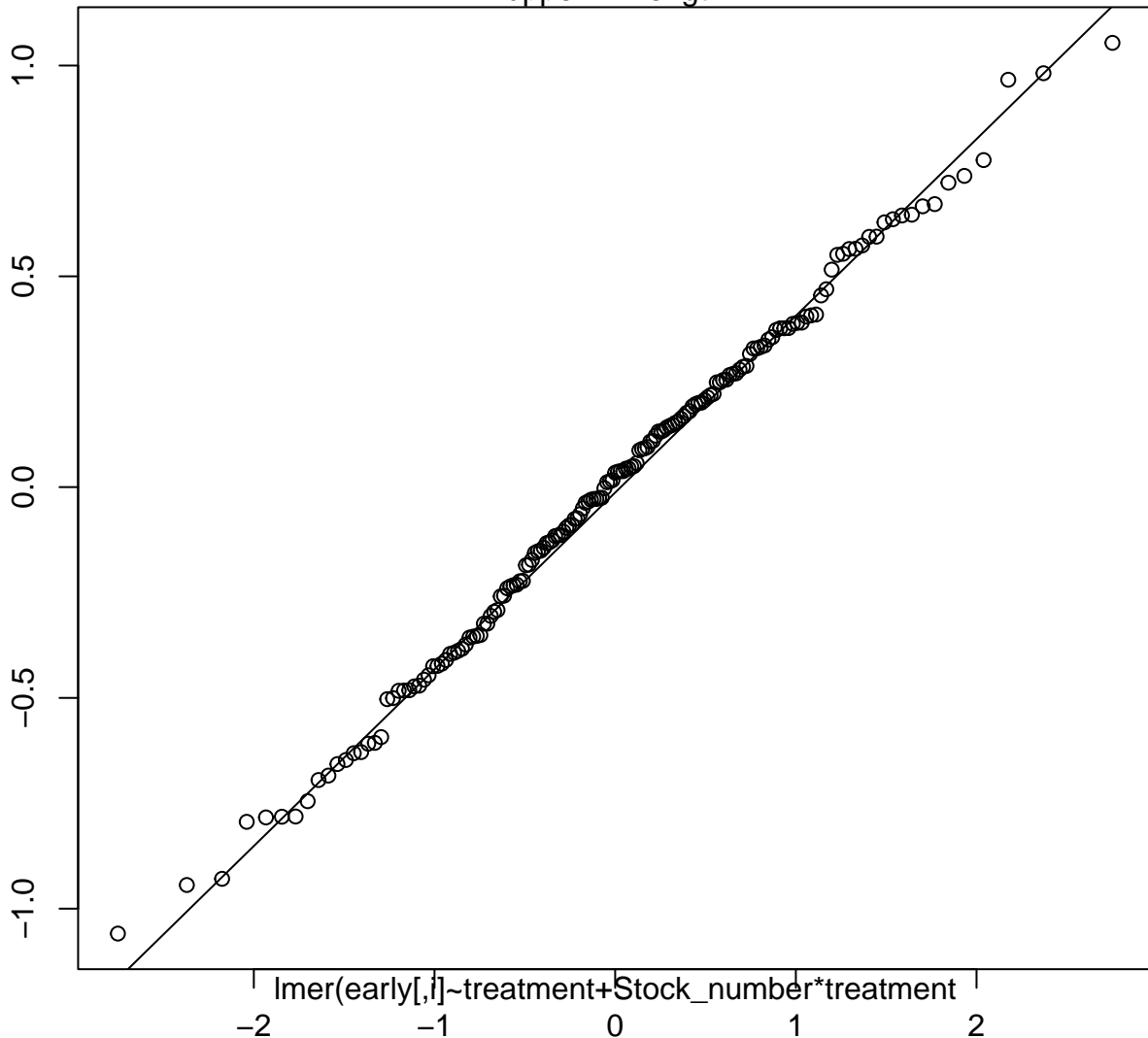




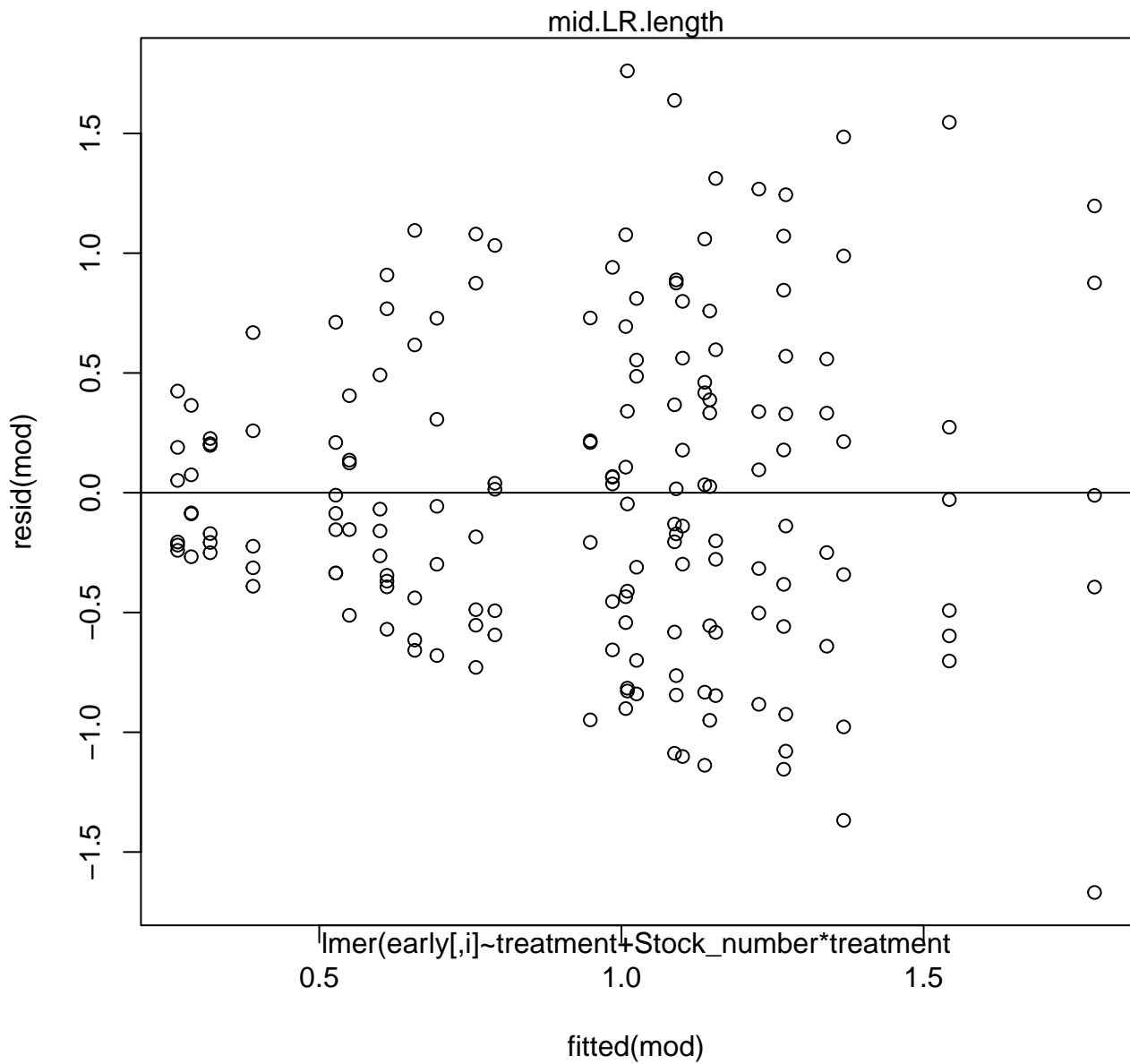
Normal Q-Q Plot

upper.LR.length

Sample Quantiles



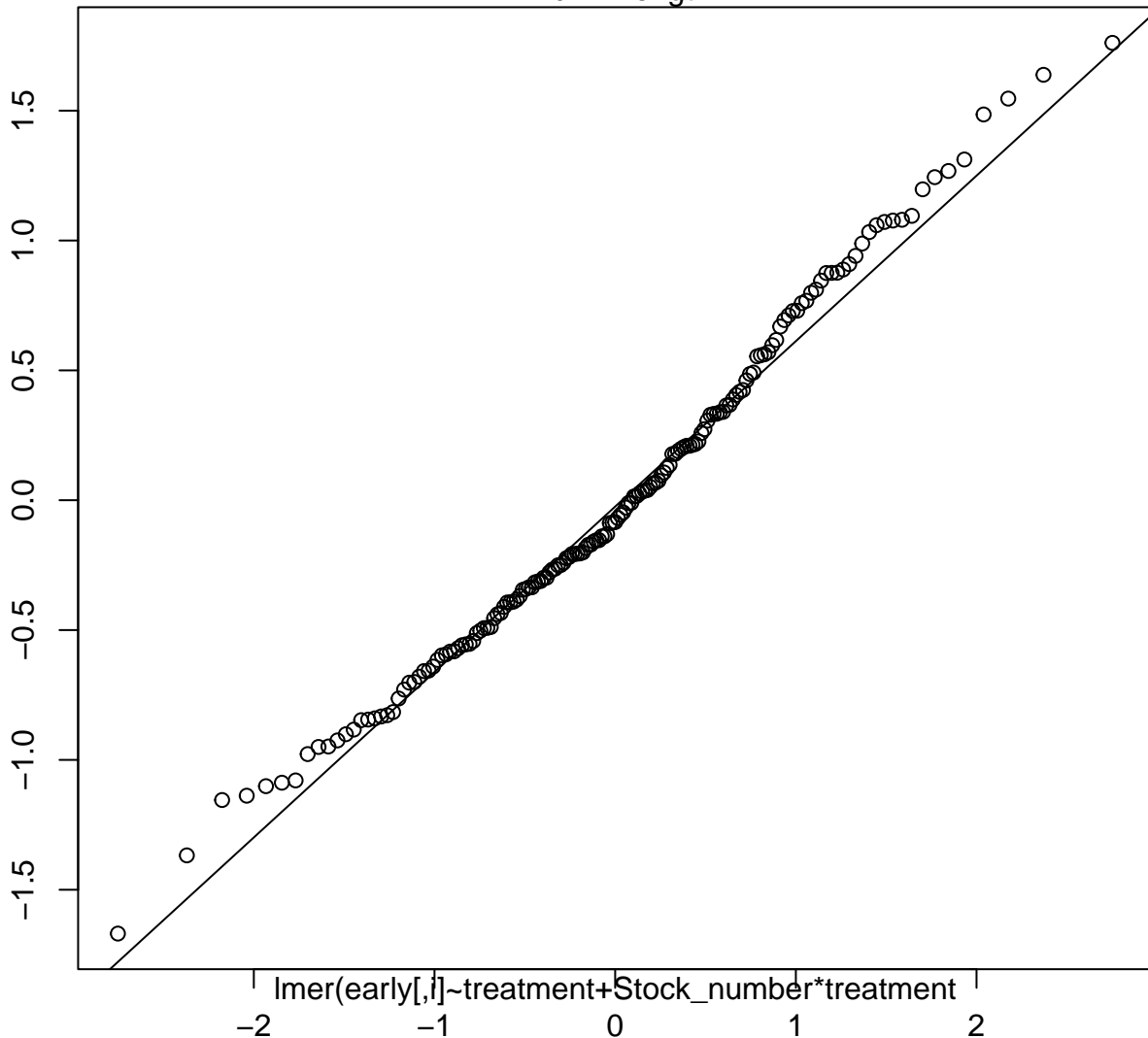
Theoretical Quantiles



Normal Q-Q Plot

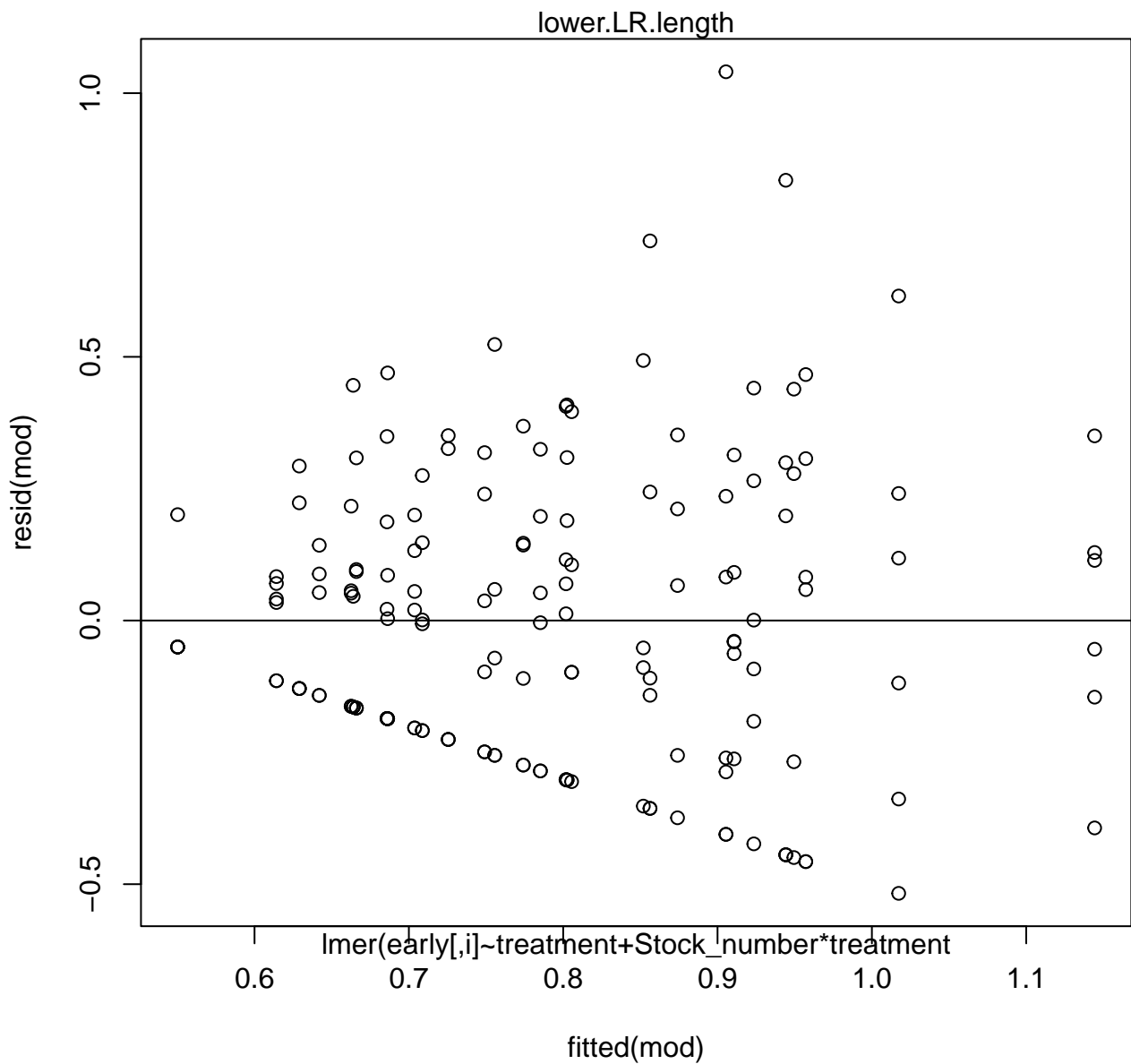
mid.LR.length

Sample Quantiles



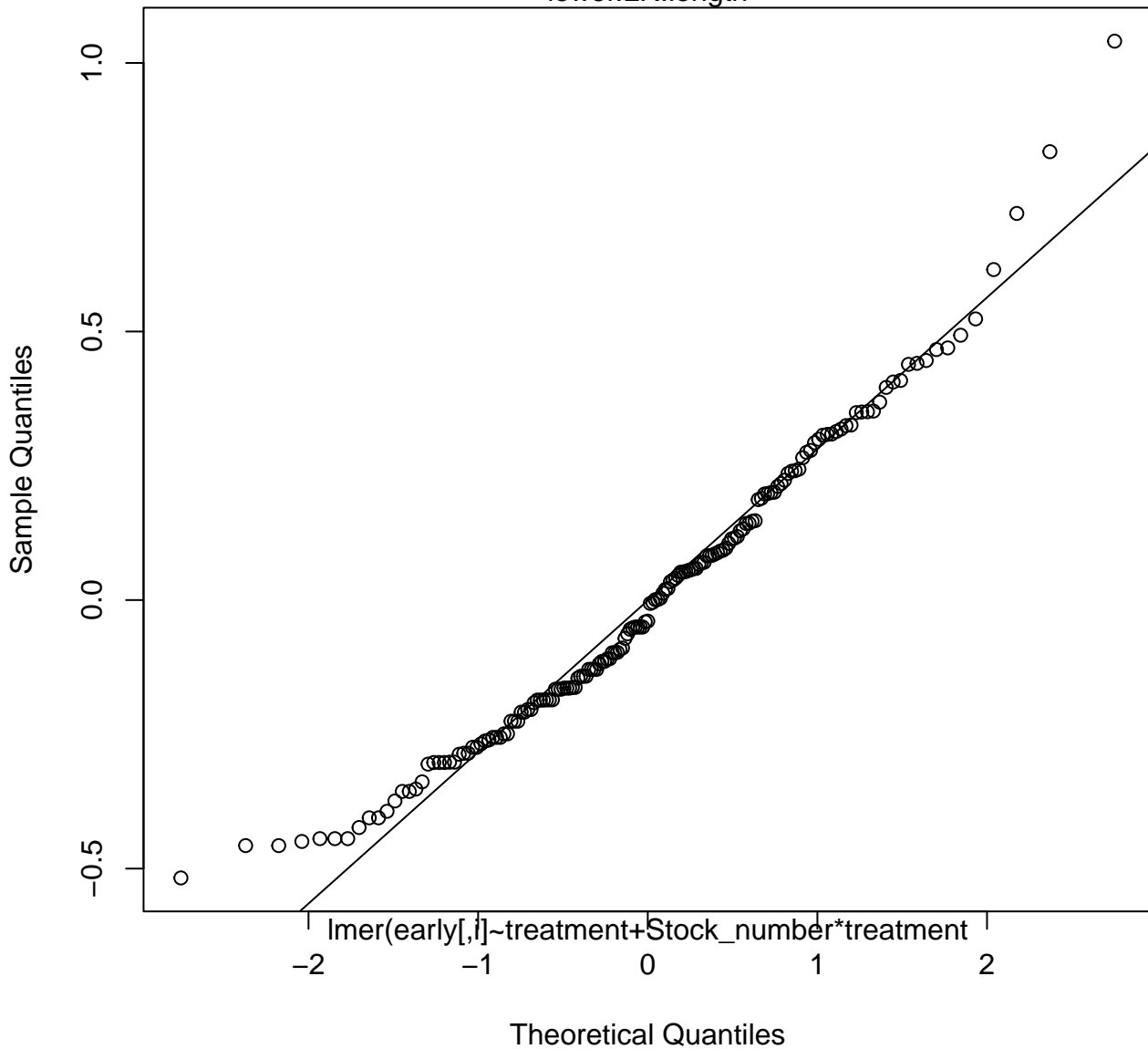
`lmer(early[,i]~treatment+Stock_number*treatment`

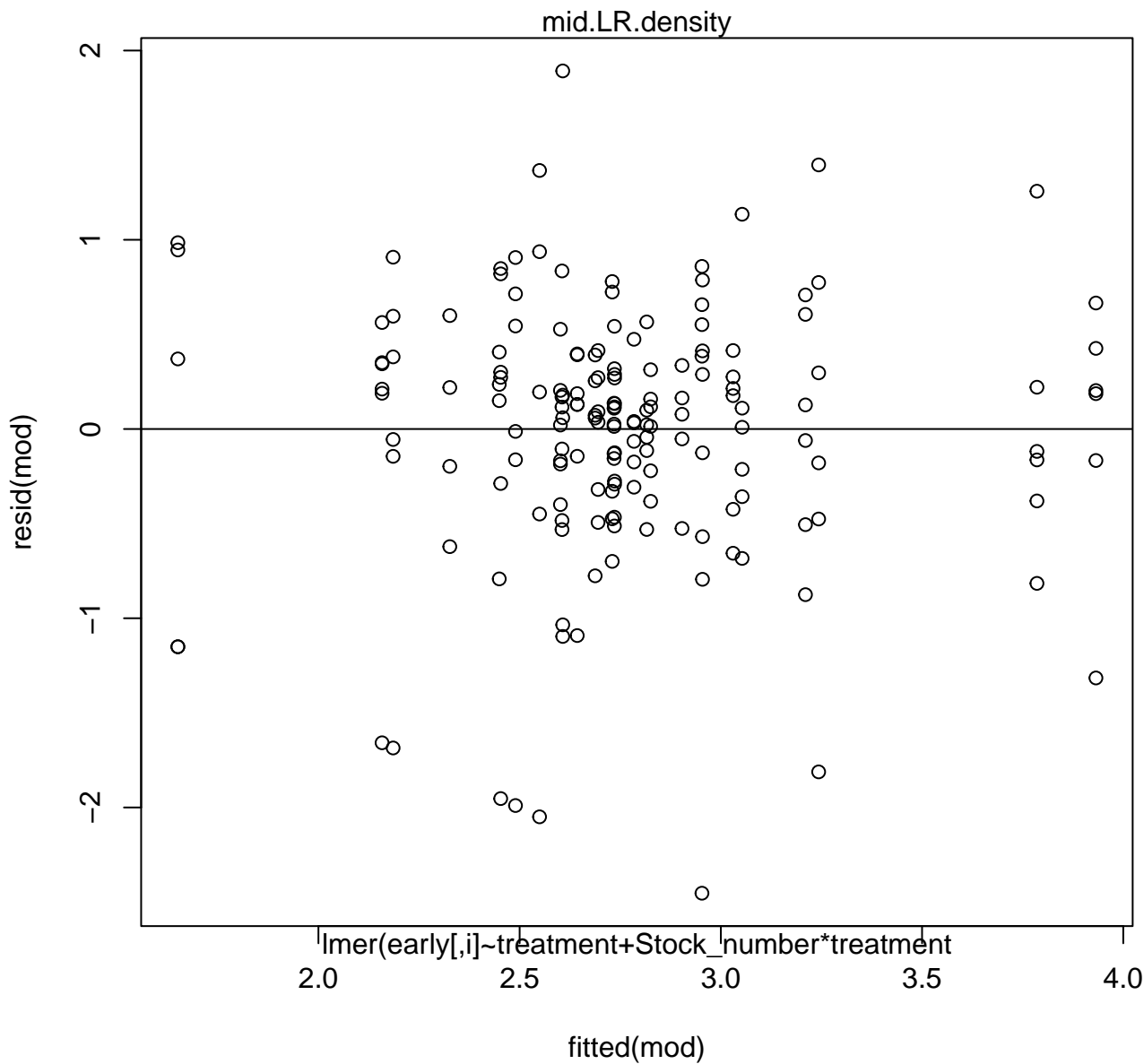
Theoretical Quantiles



Normal Q-Q Plot

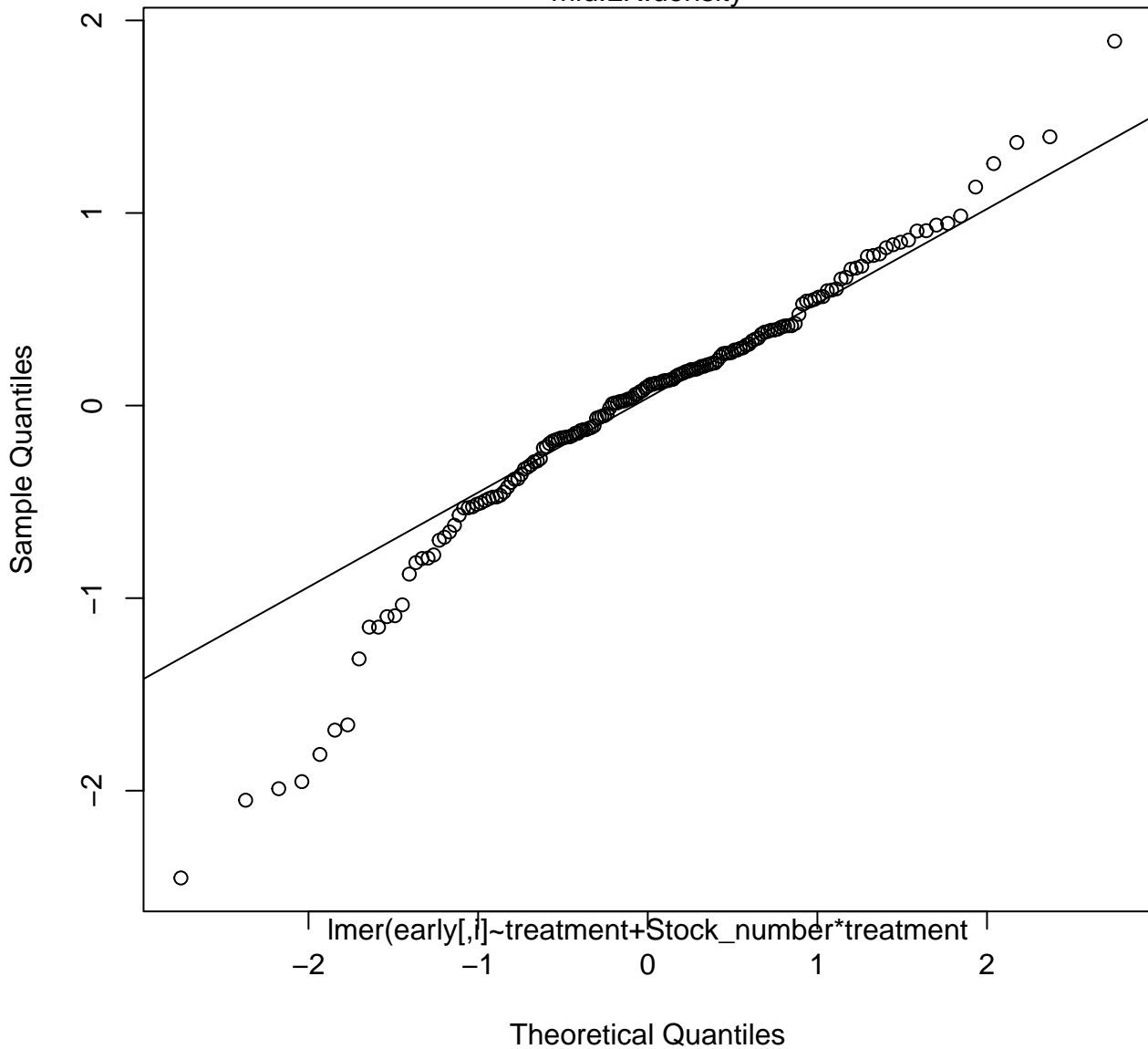
lower.LR.length

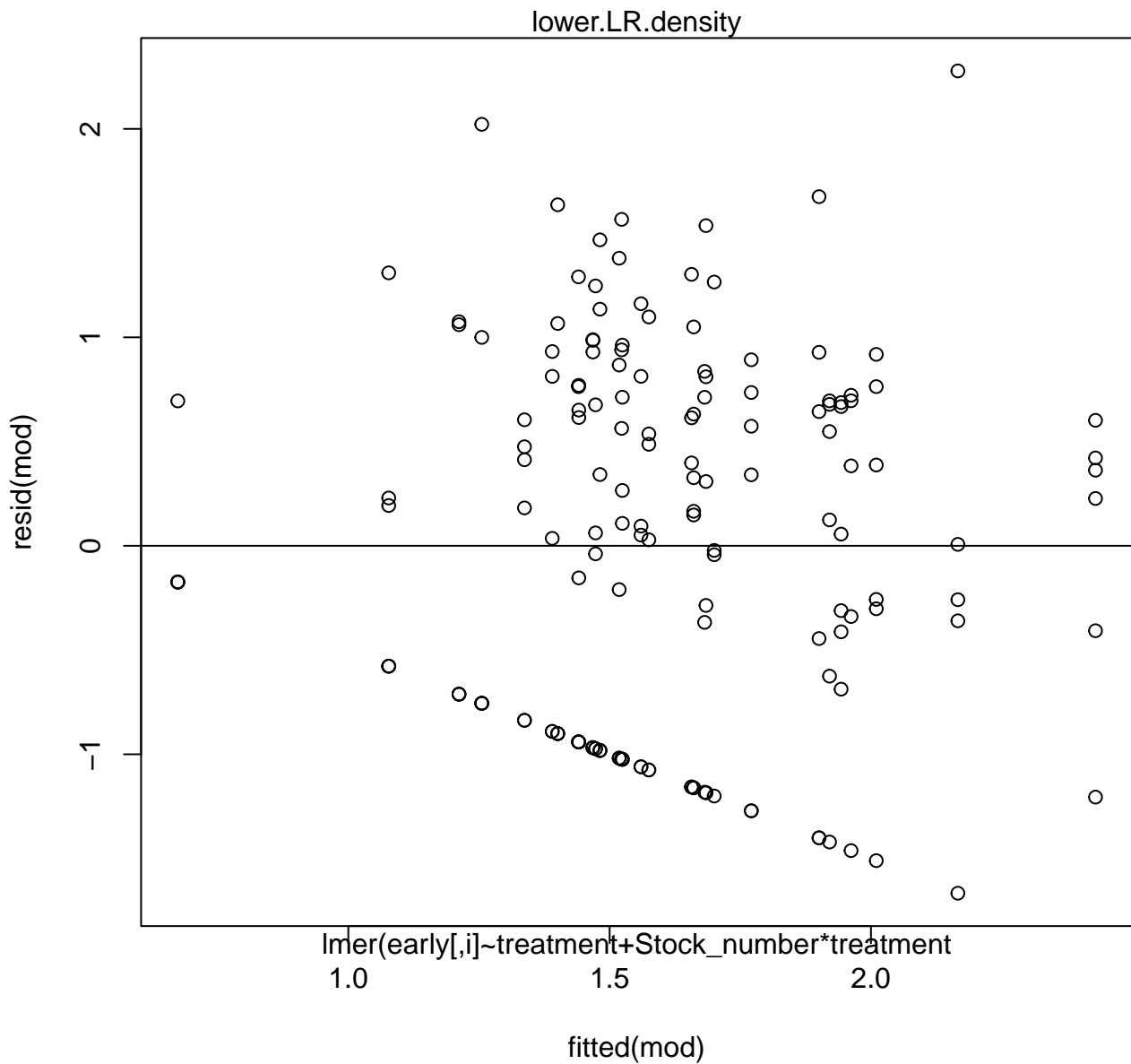




Normal Q-Q Plot

mid.LR.density

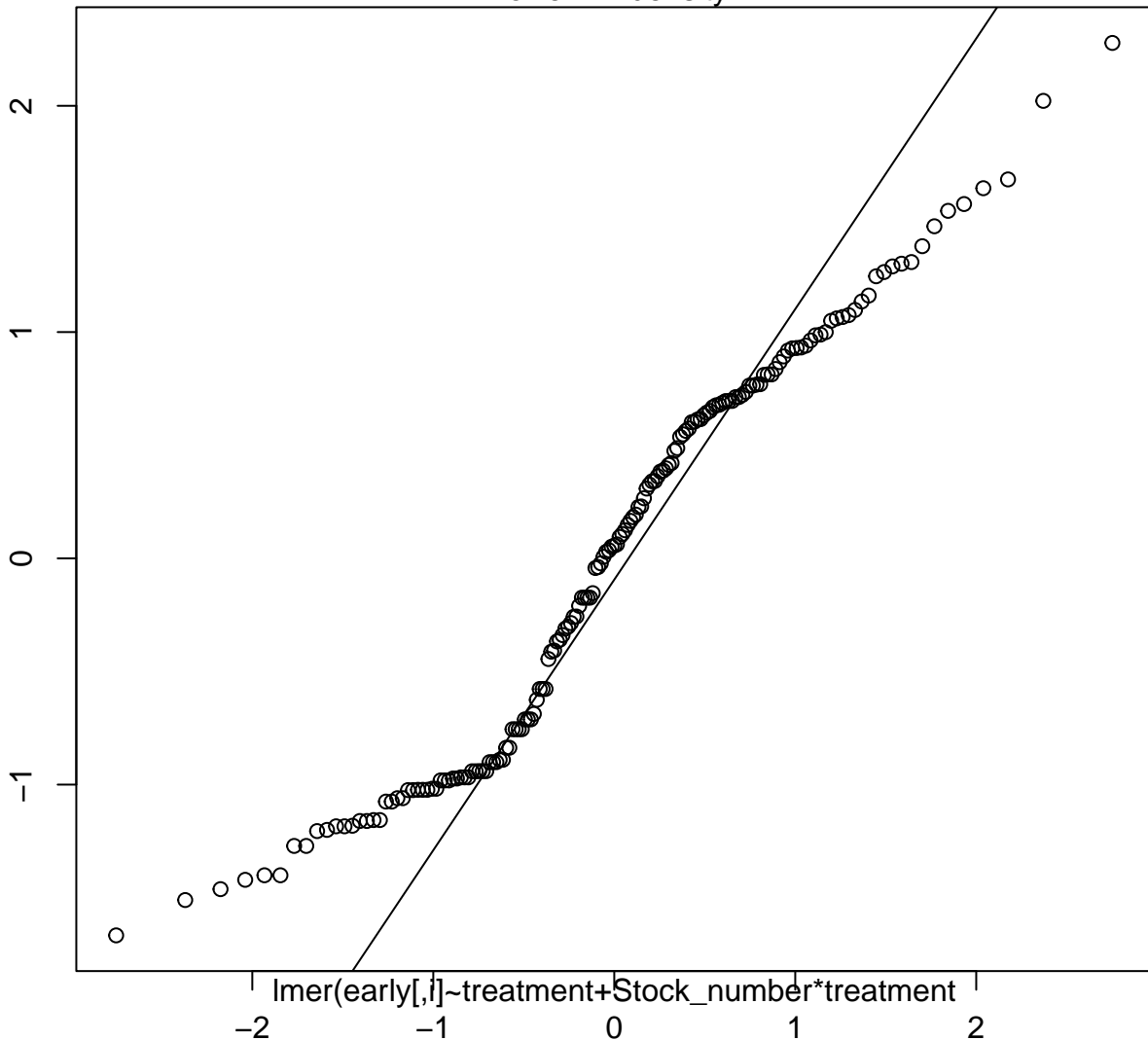




Normal Q-Q Plot

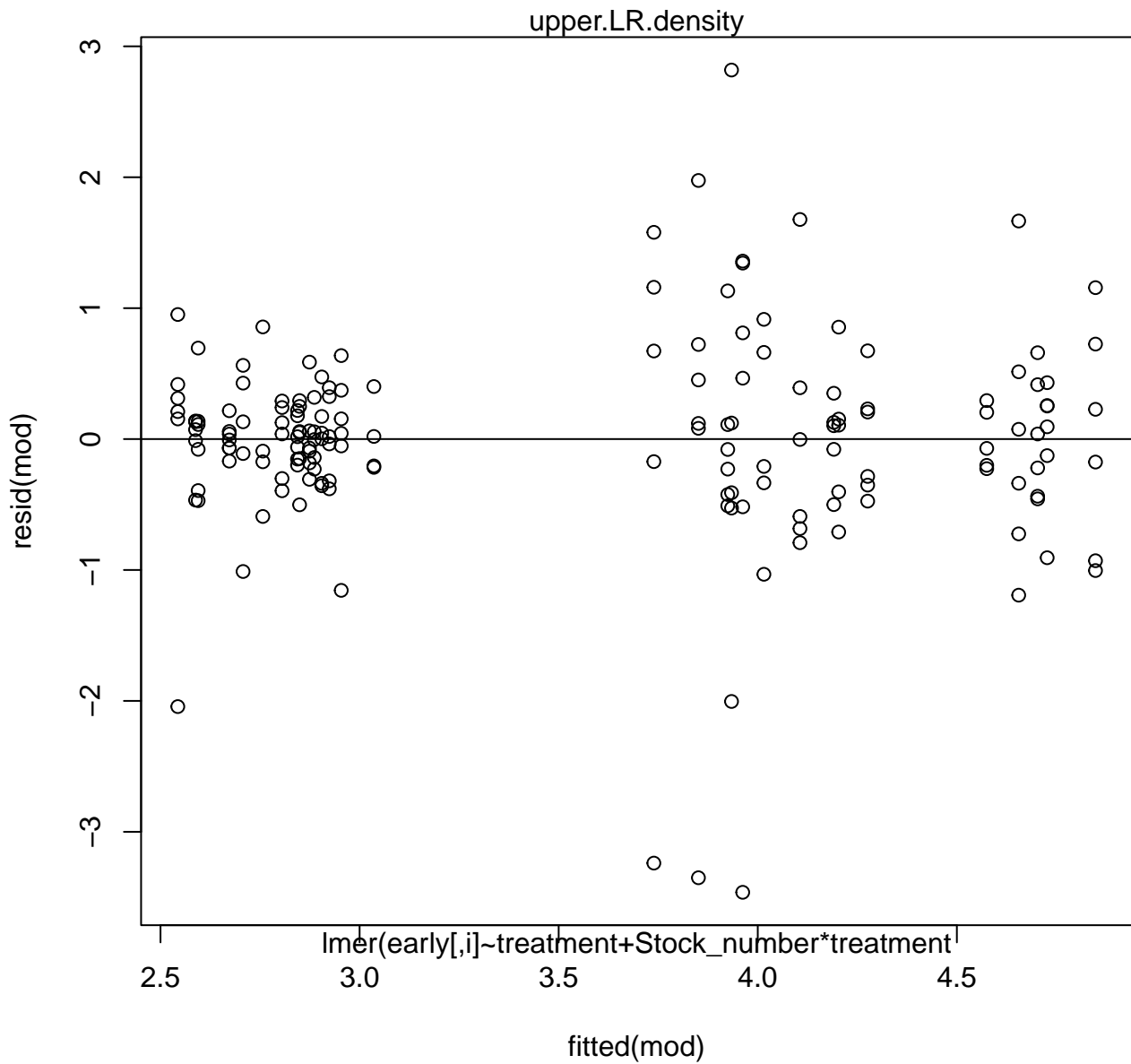
lower.LR.density

Sample Quantiles



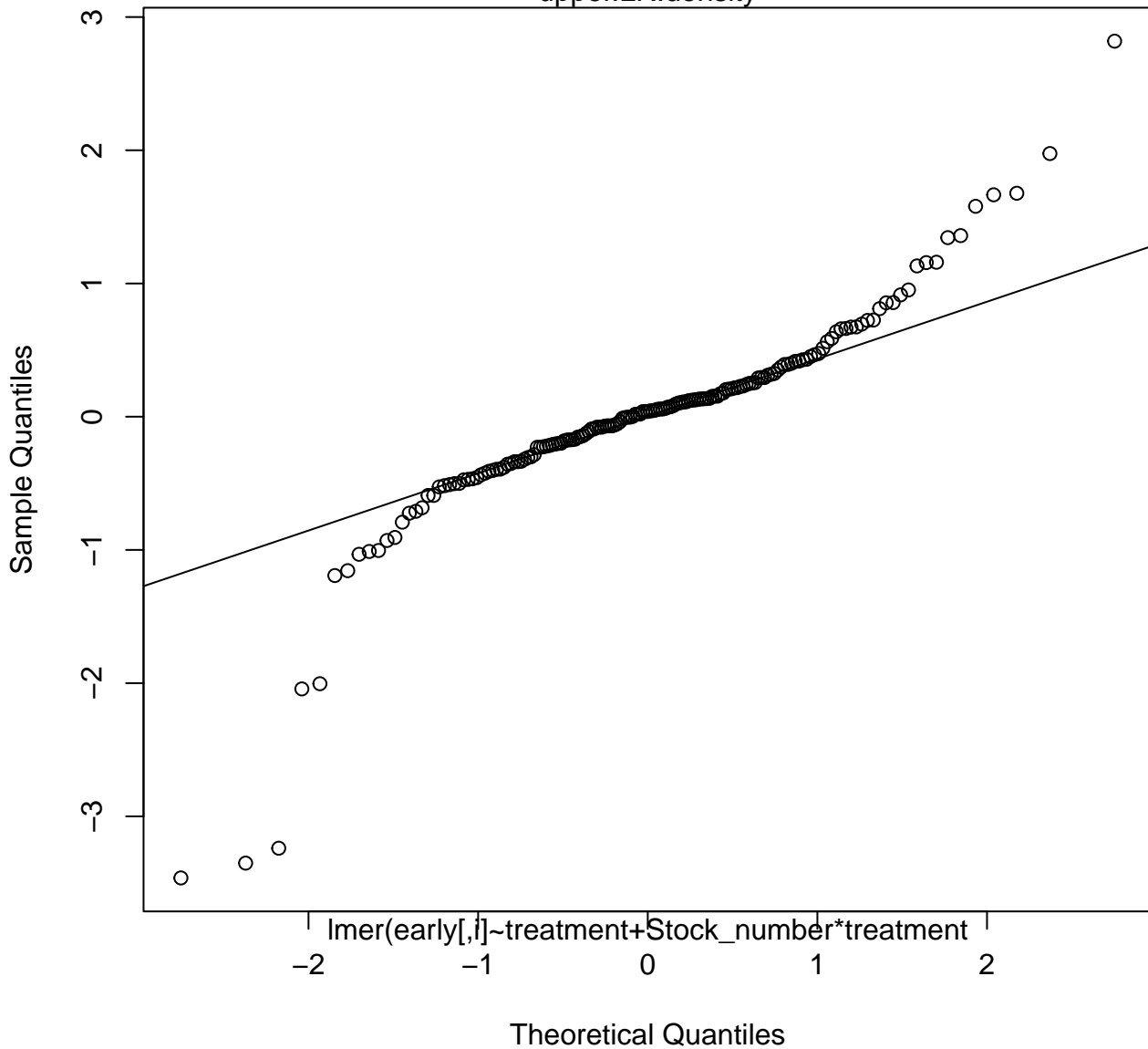
Theoretical Quantiles

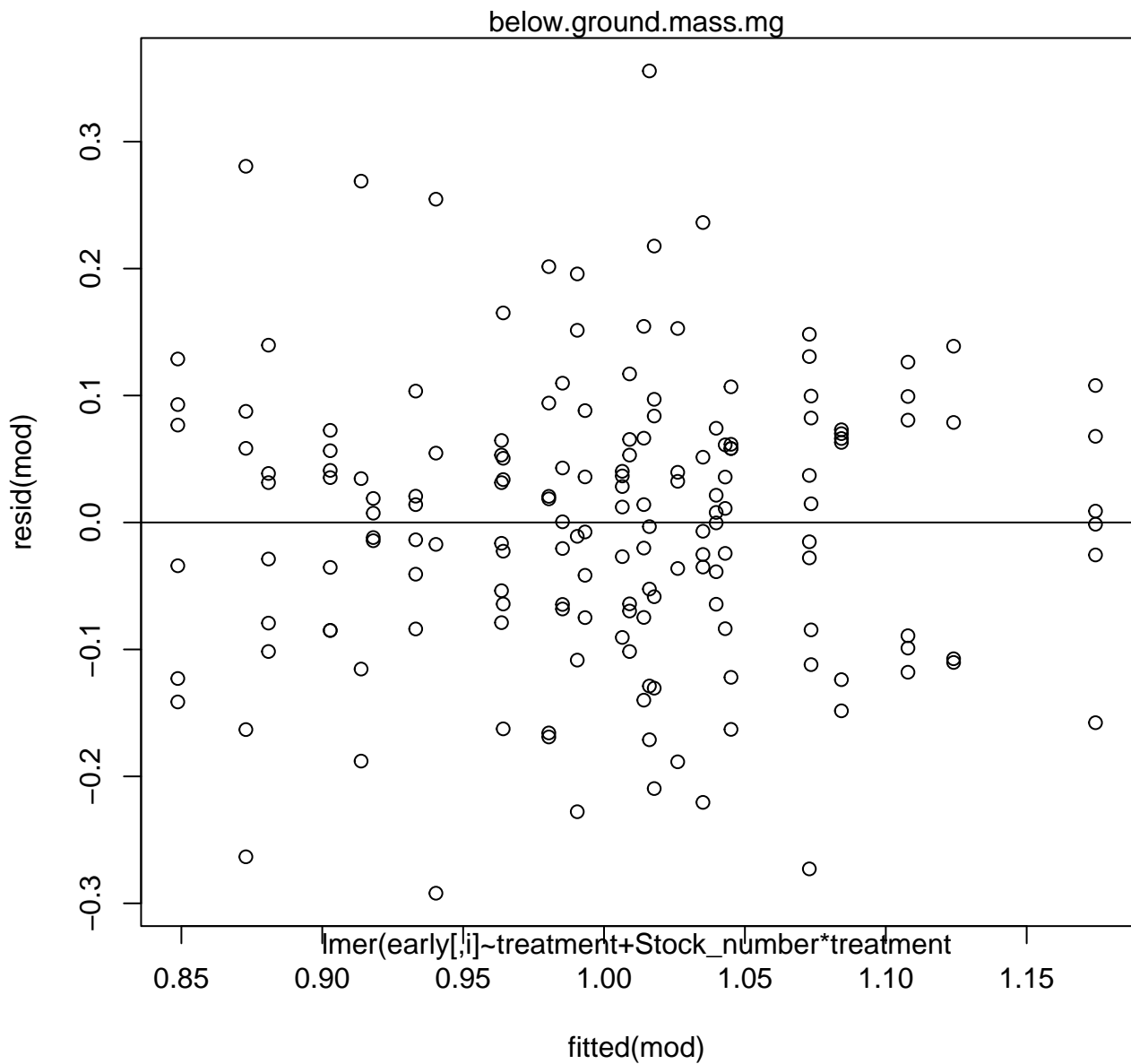
lmer(early[,i]~treatment+Stock_number*treatment



Normal Q-Q Plot

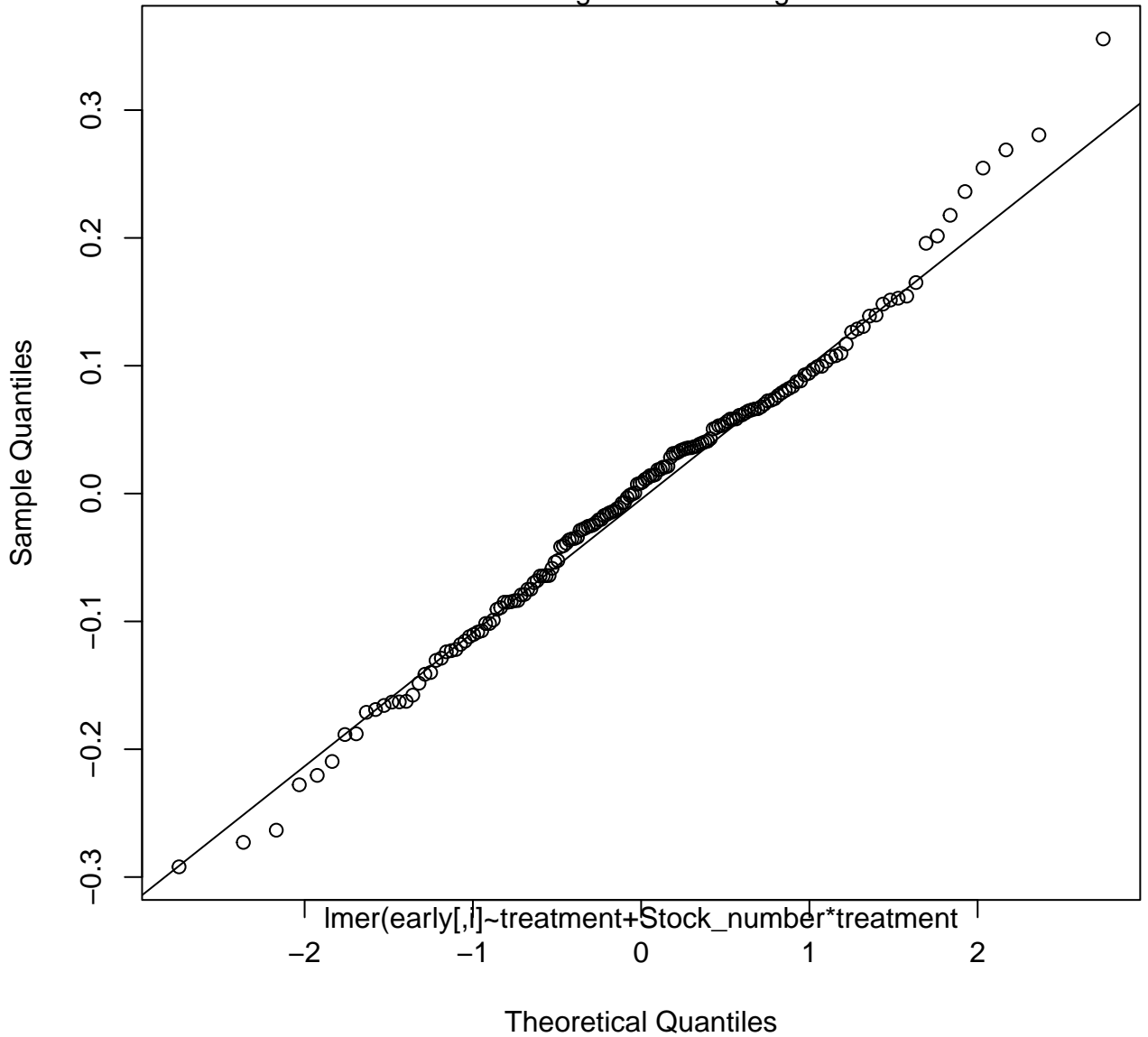
upper.LR.density

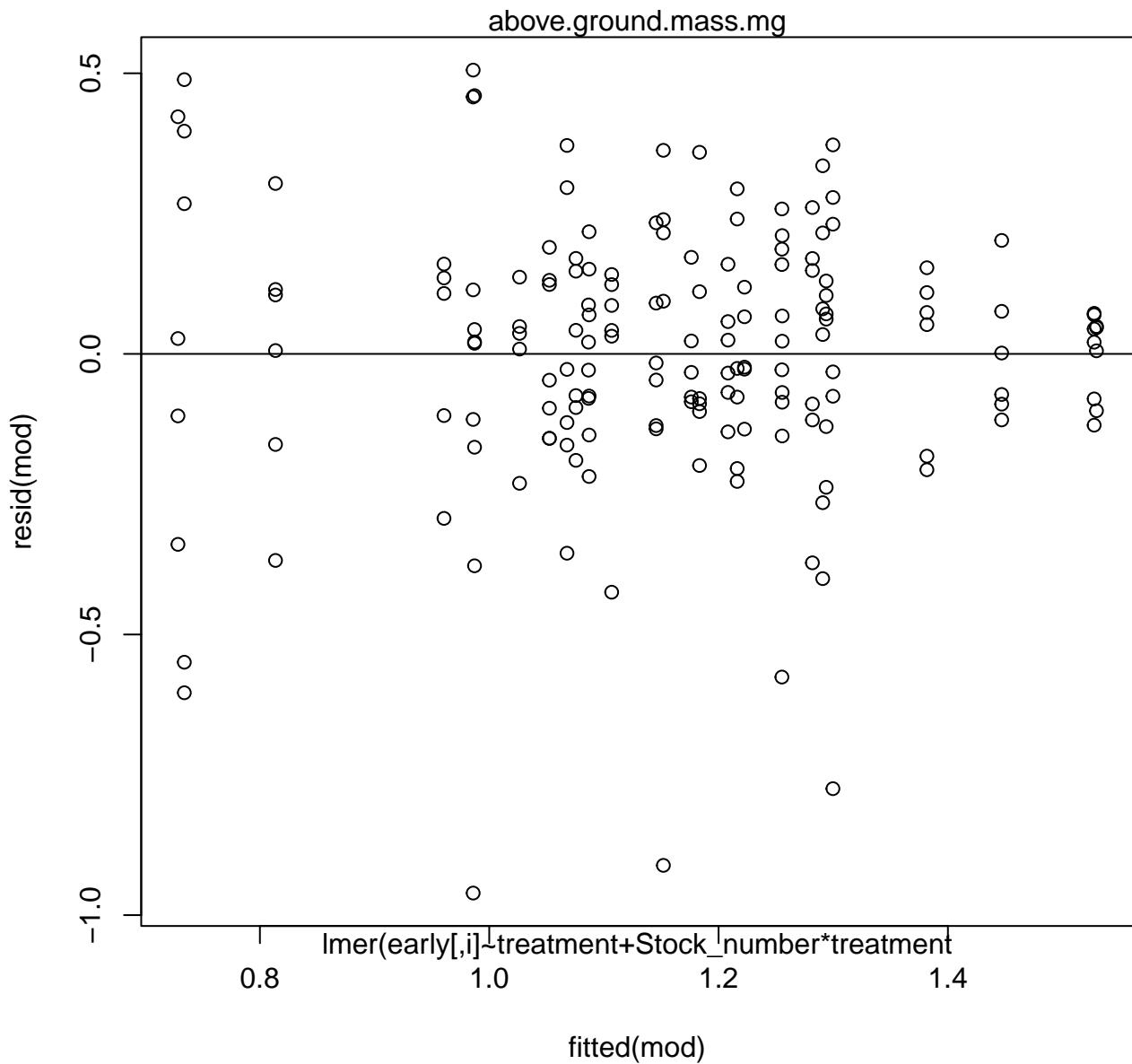




Normal Q-Q Plot

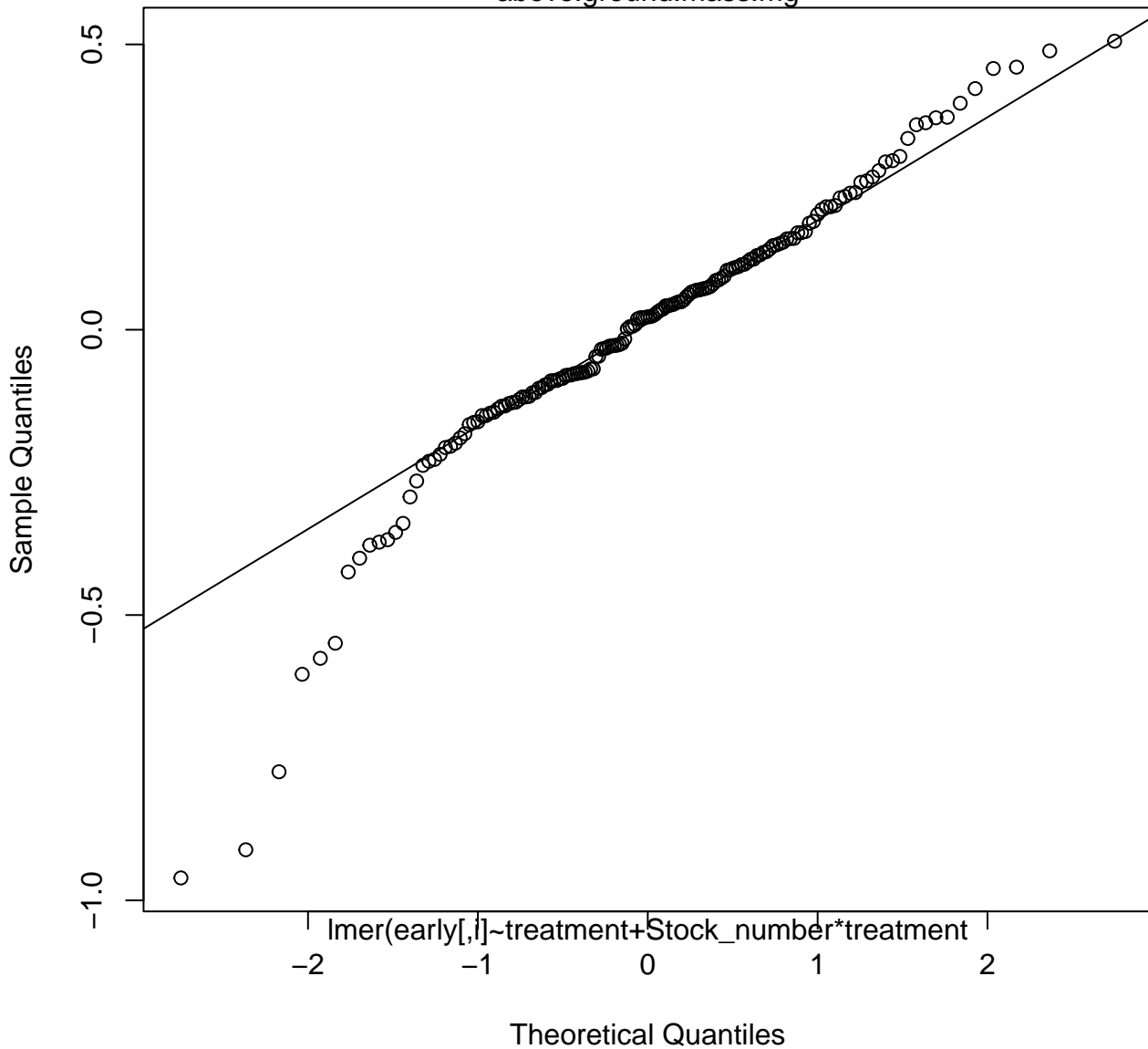
below.ground.mass.mg

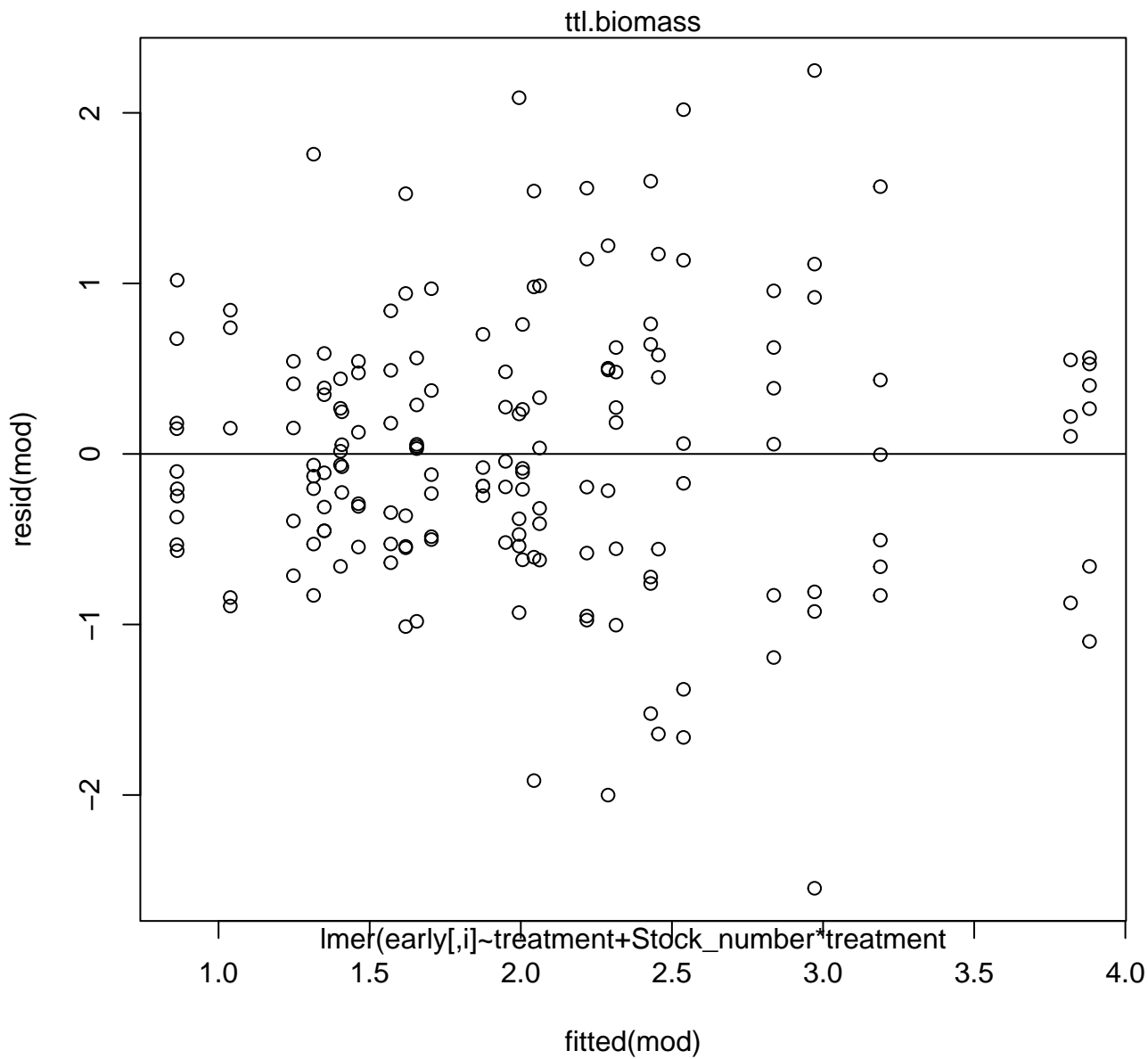




Normal Q-Q Plot

above.ground.mass.mg

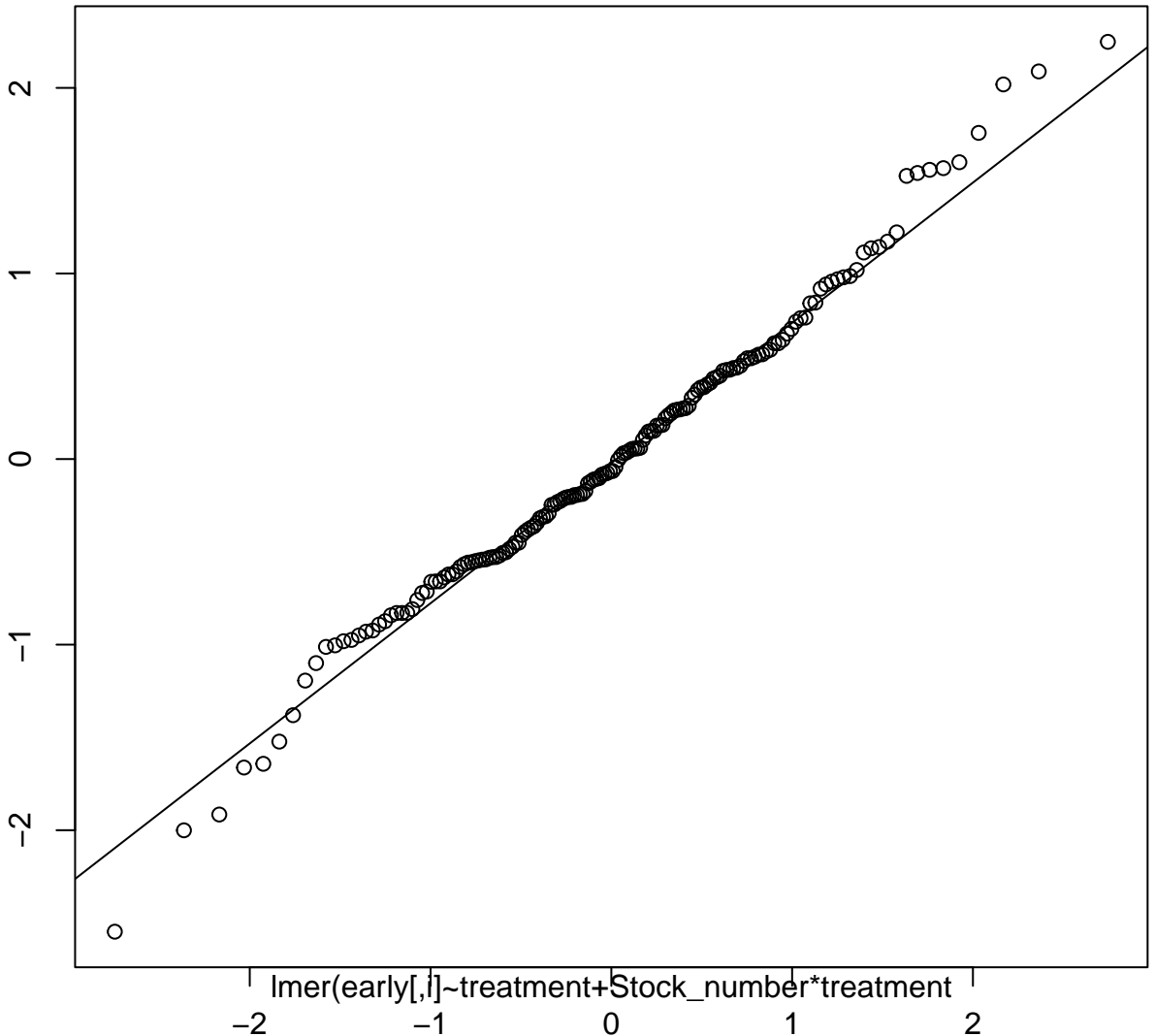




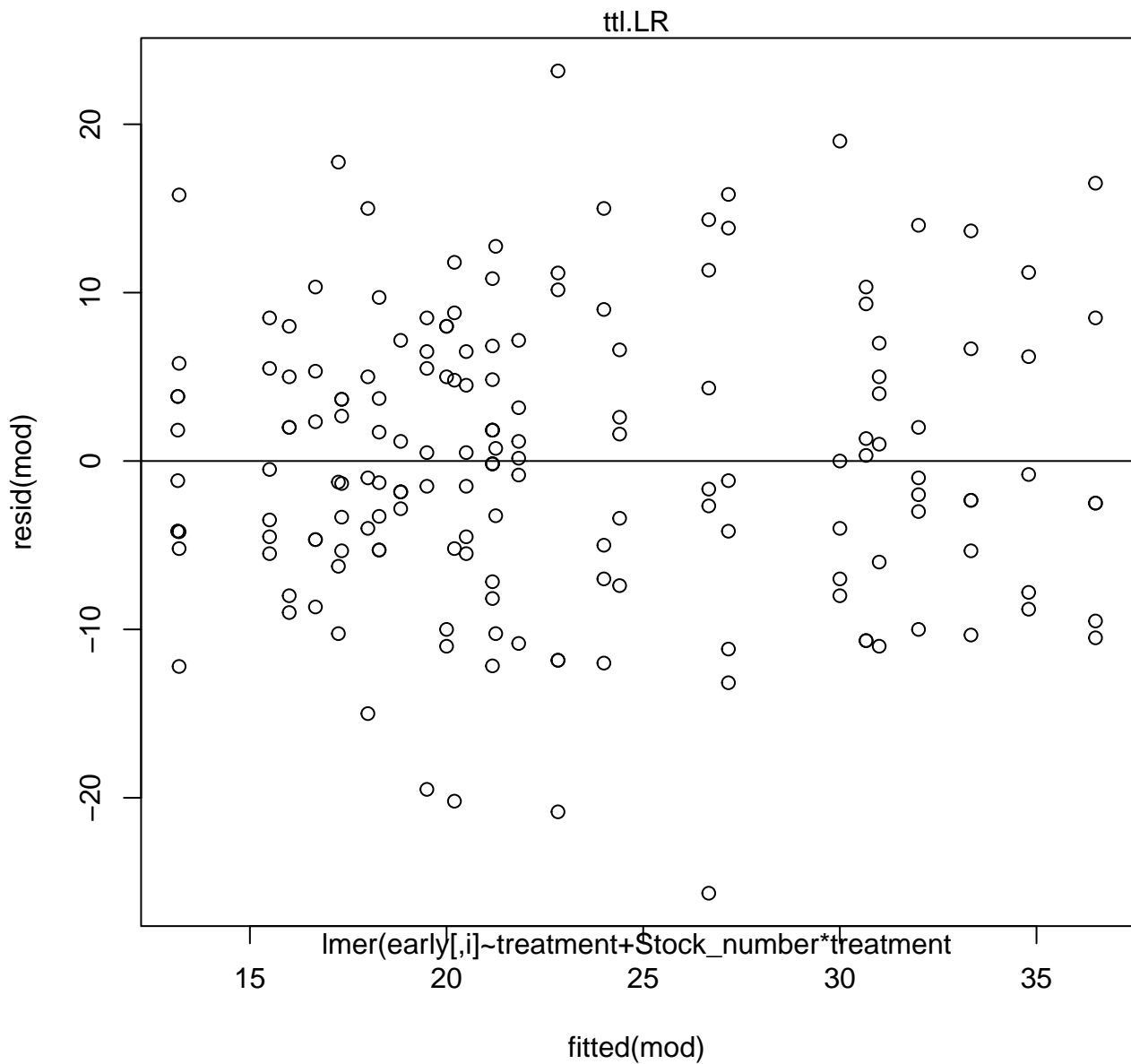
Normal Q-Q Plot

t1.biomass

Sample Quantiles



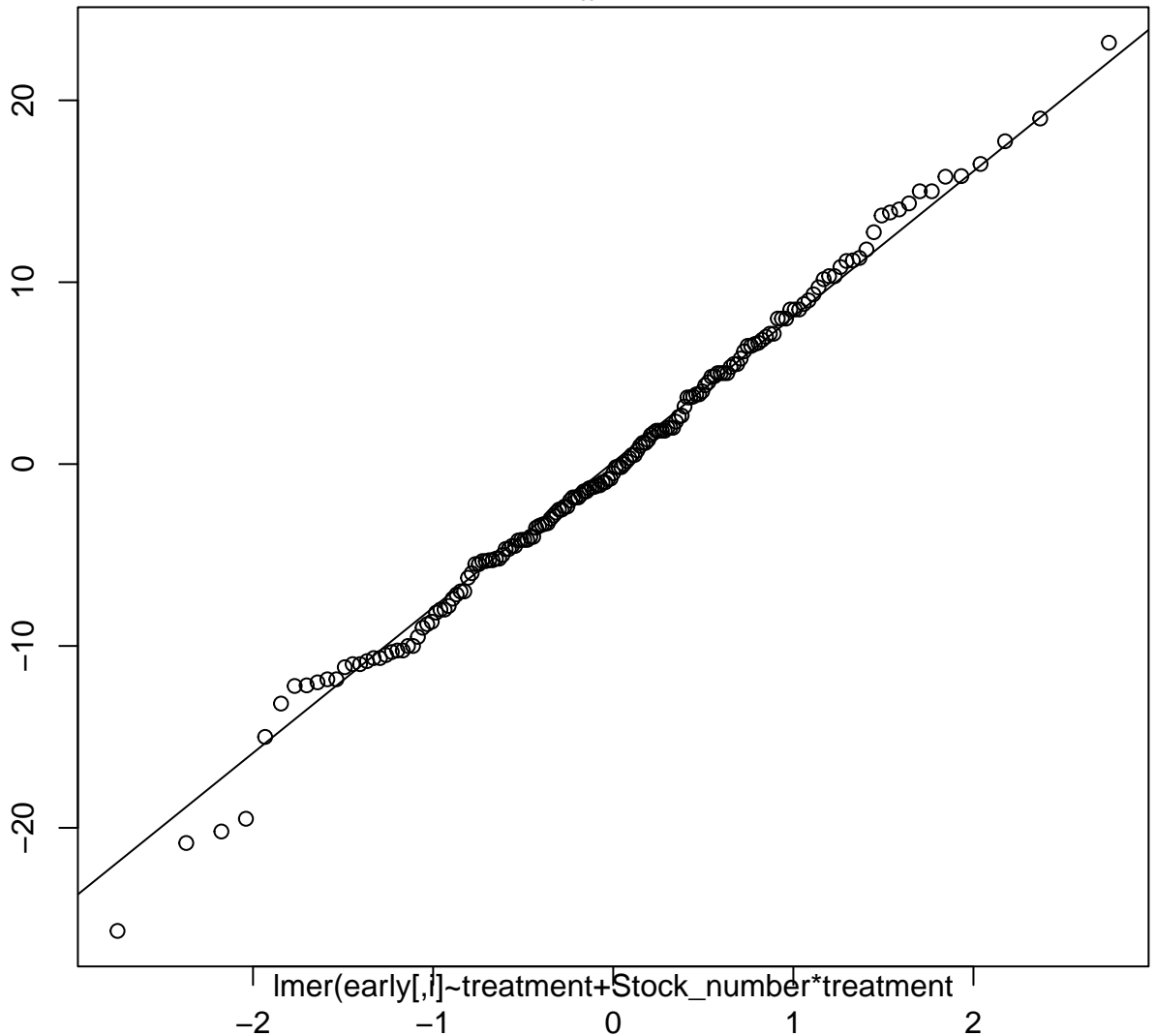
Theoretical Quantiles



Normal Q-Q Plot

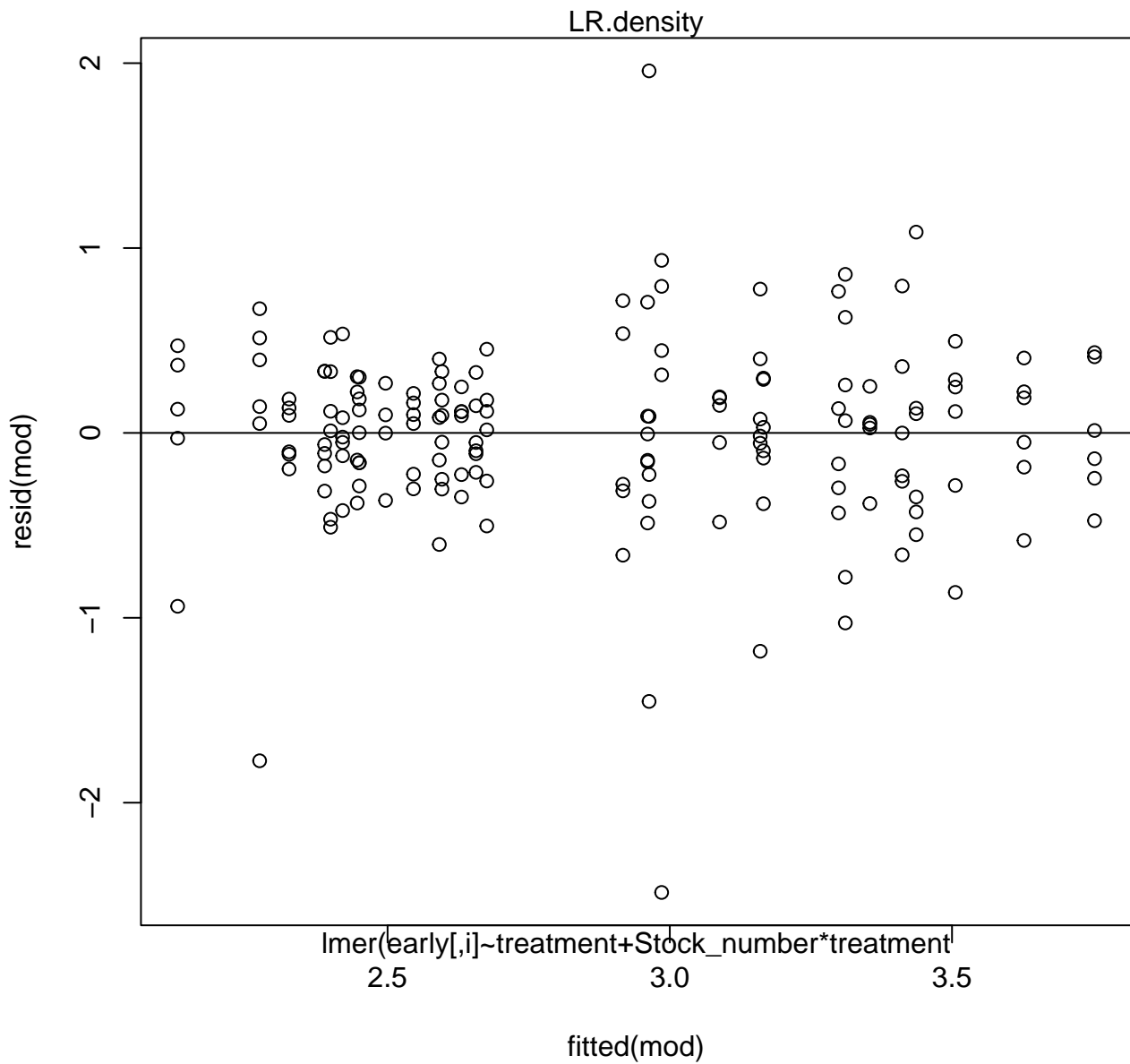
ttl.LR

Sample Quantiles



lmer(early[,i]~treatment+Stock_number*treatment

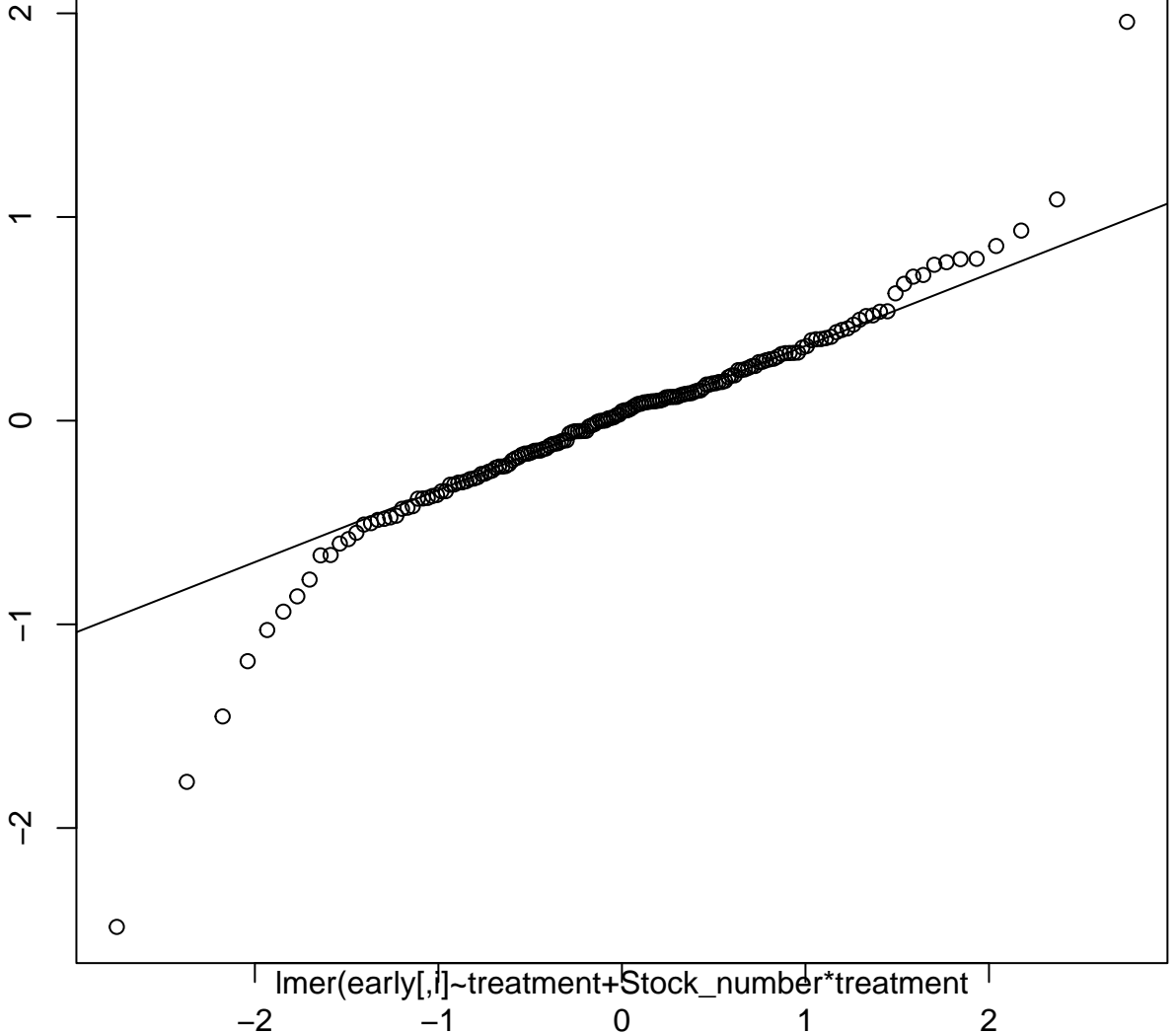
Theoretical Quantiles



Normal Q-Q Plot

LR.density

Sample Quantiles



Theoretical Quantiles