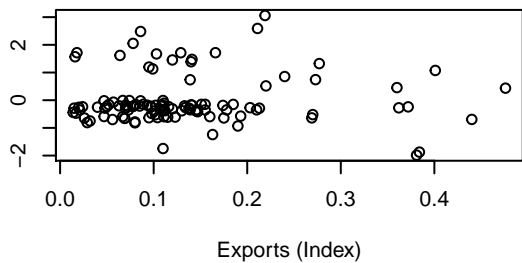
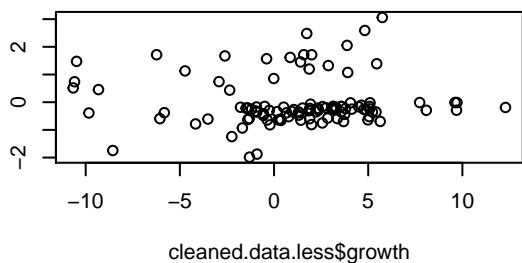


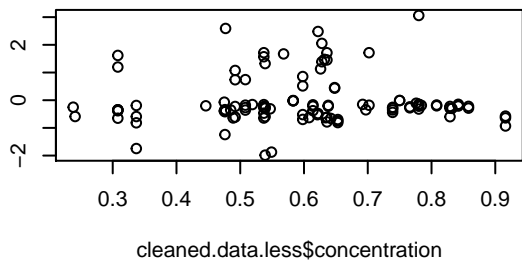
Pearson Residuals



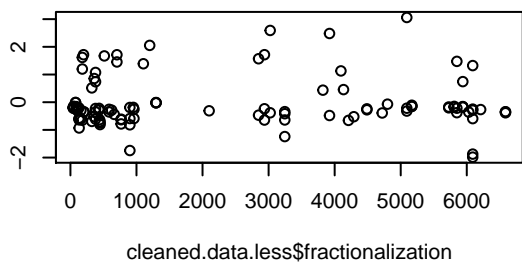
residuals(new.glm.fit, type = "pearson")



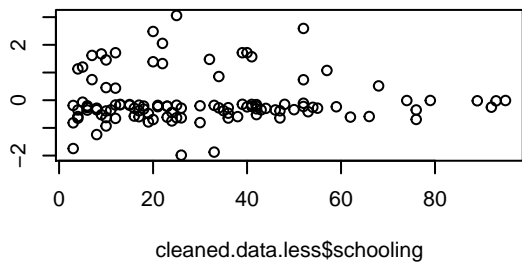
residuals(new.glm.fit, type = "pearson")



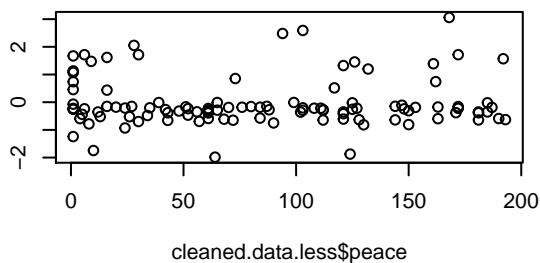
residuals(new.glm.fit, type = "pearson")



residuals(new.glm.fit, type = "pearson")



residuals(new.glm.fit, type = "pearson")



residuals(new.glm.fit, type = "pearson")

