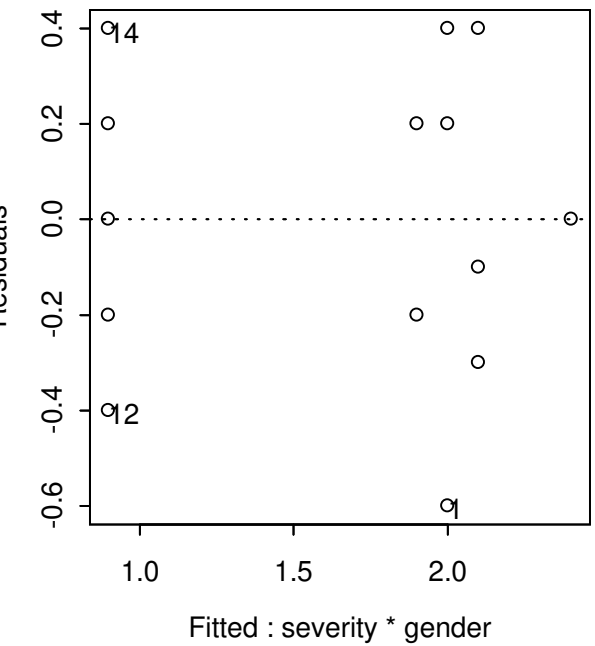
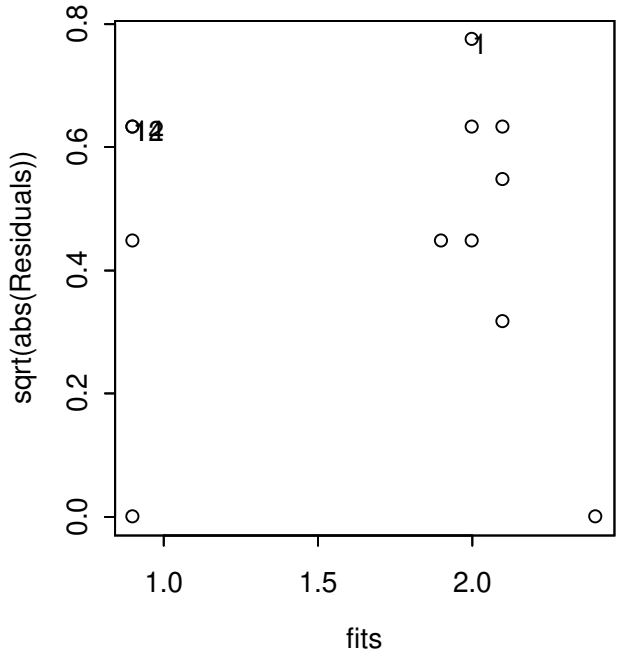
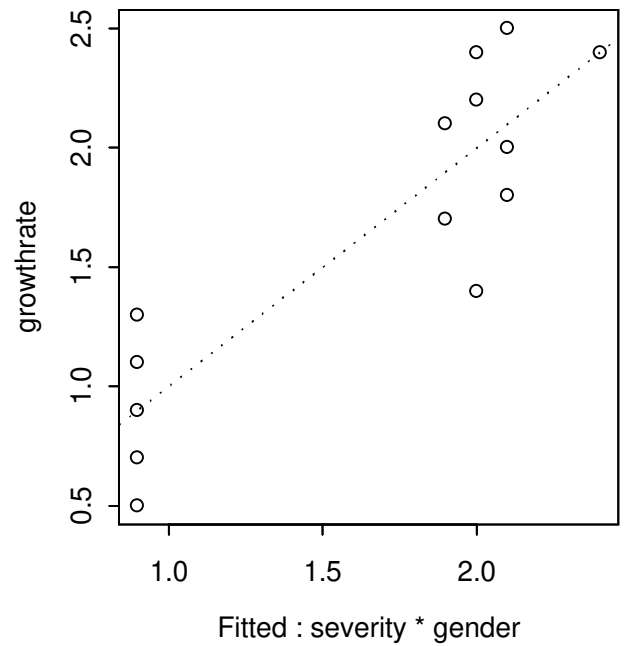
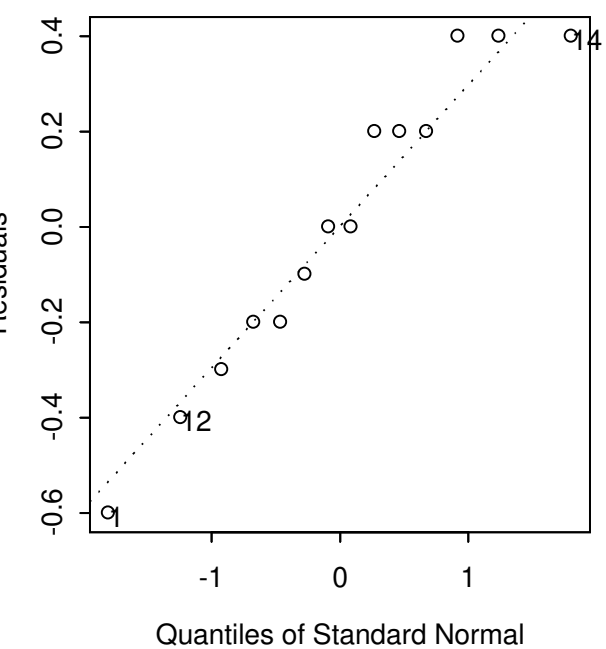
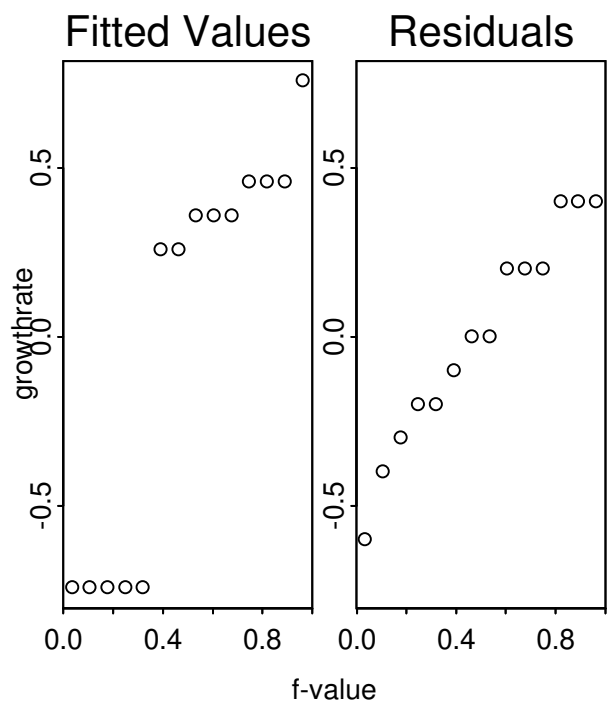
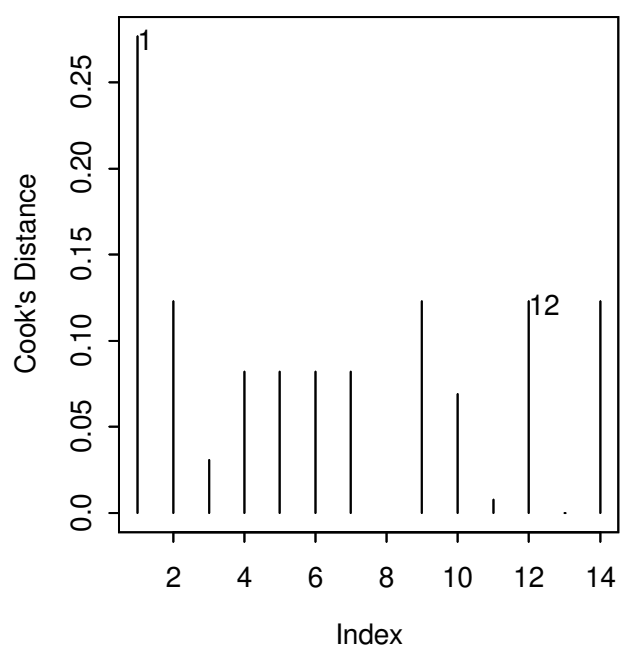

Residuals
14
12
1
0.4
0.2
0.0
-0.2
-0.4
-0.6
1.0
1.5
2.0
Fitted : severity * gender


sqrt(abs(Residuals))
1
12
0.8
0.6
0.4
0.2
0.0
1.0
1.5
2.0
fits


growthrate
2.5
2.0
1.5
1.0
0.5
1.0
1.5
2.0
Fitted : severity * gender


Residuals
14
12
1
0.4
0.2
0.0
-0.2
-0.4
-0.6
-1
0
1
Quantiles of Standard Normal


Fitted Values
Residuals
growthrate
0.5
0.0
-0.5
0.0
0.4
0.8
f-value


Cook's Distance
1
12
0.25
0.20
0.15
0.10
0.05
0.0
2
4
6
8
10
12
14
Index