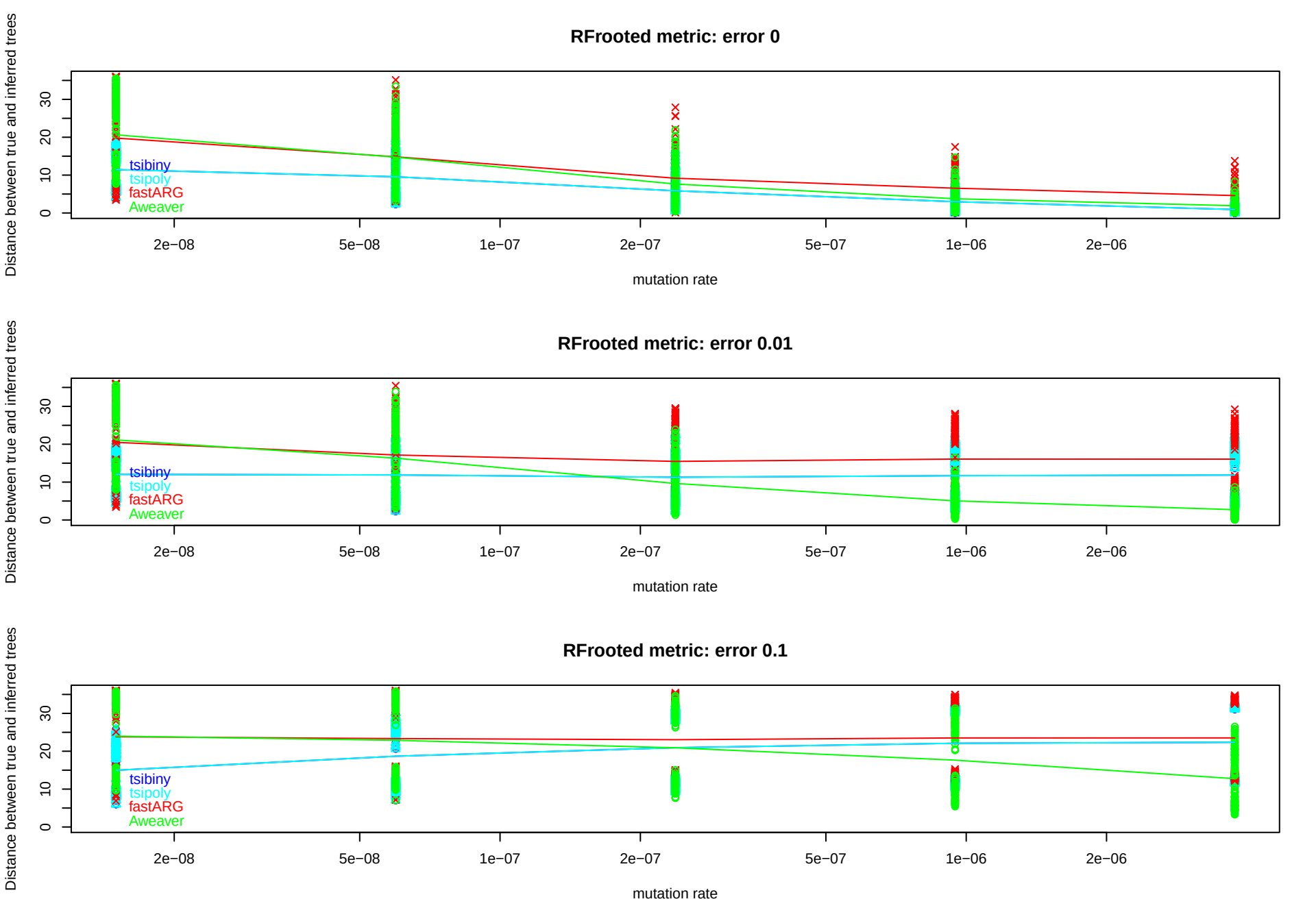
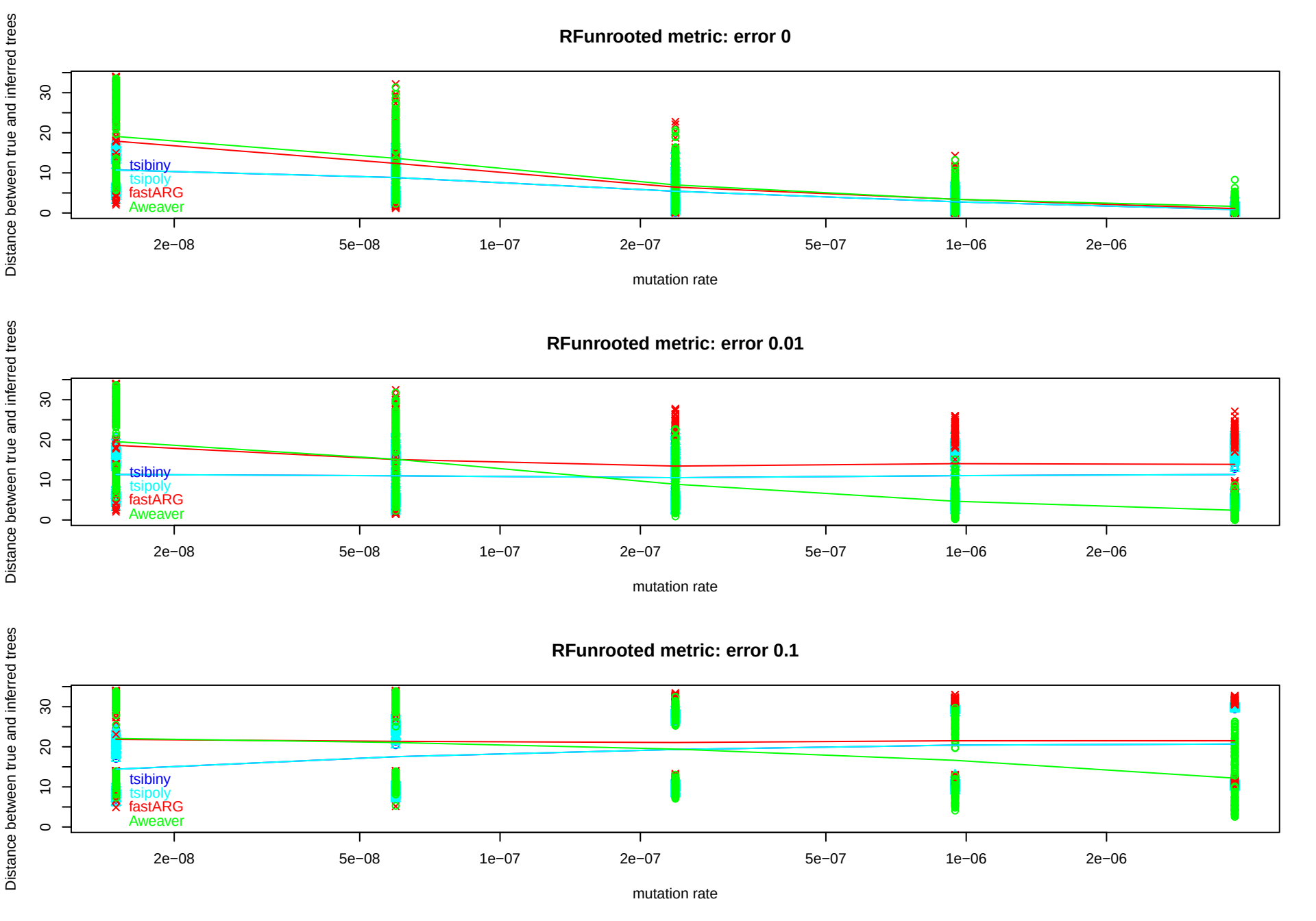
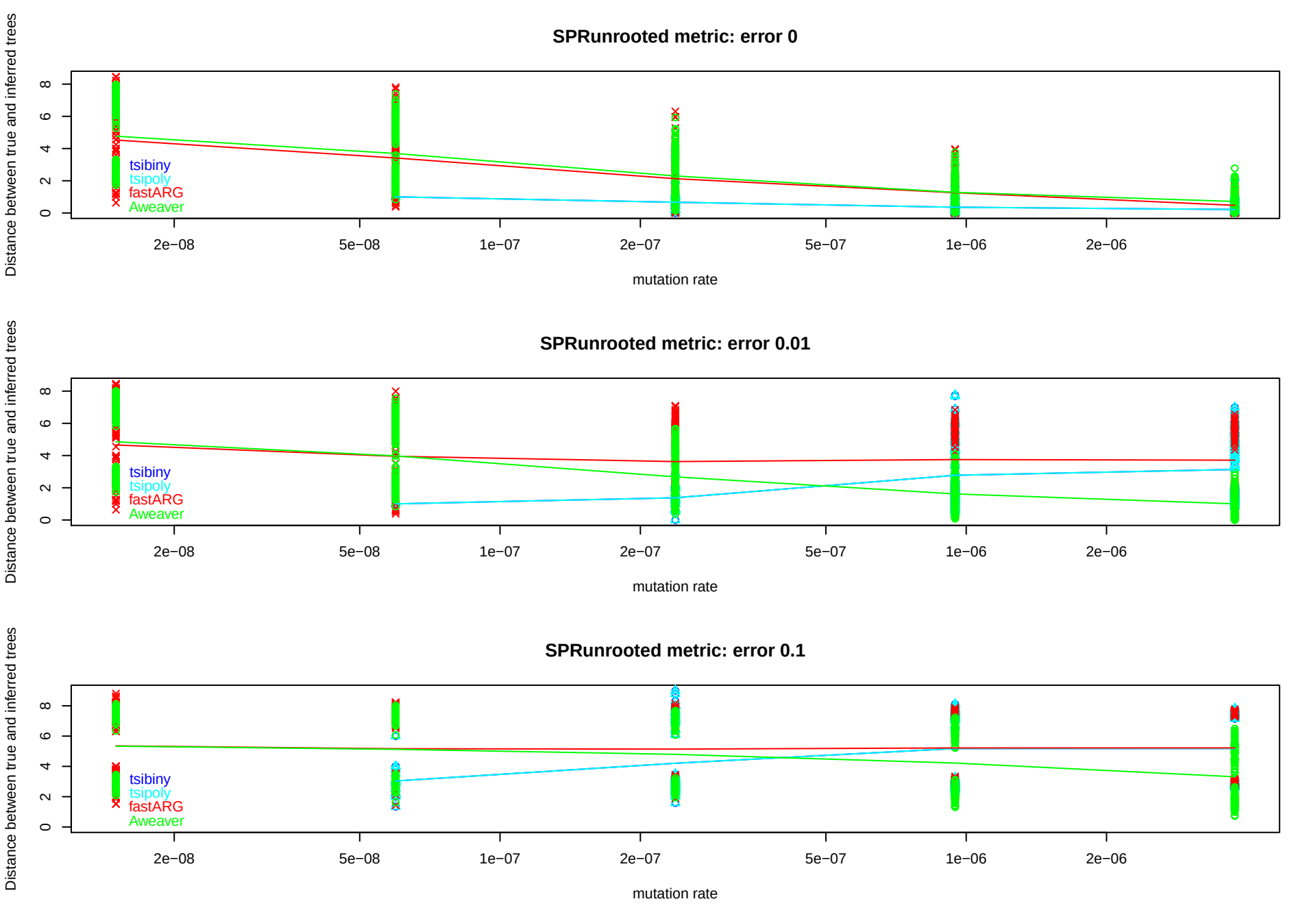
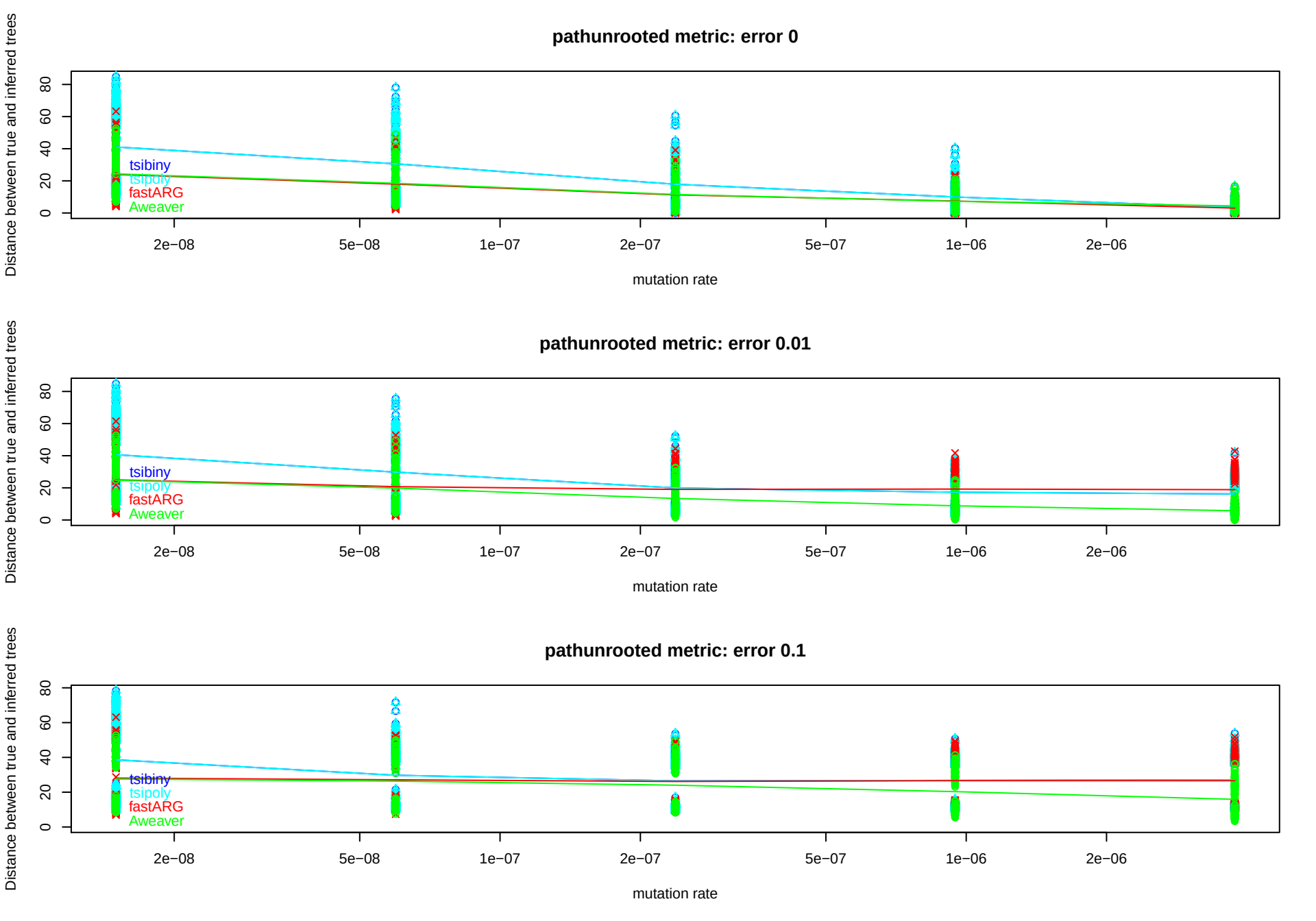

RFrooted metric: error 0
Distance between true and inferred trees
0
10
20
30
tsibiny
tsipoly
fastARG
Aweaver
2e−08
5e−08
1e−07
2e−07
5e−07
1e−06
2e−06
mutation rate
RFrooted metric: error 0.01
Distance between true and inferred trees
0
10
20
30
tsibiny
tsipoly
fastARG
Aweaver
2e−08
5e−08
1e−07
2e−07
5e−07
1e−06
2e−06
mutation rate
RFrooted metric: error 0.1
Distance between true and inferred trees
0
10
20
30
tsibiny
tsipoly
fastARG
Aweaver
2e−08
5e−08
1e−07
2e−07
5e−07
1e−06
2e−06
mutation rate


RFunrooted metric: error 0
Distance between true and inferred trees
0
10
20
30
tsibiny
tsipoly
fastARG
Aweaver
2e−08
5e−08
1e−07
2e−07
5e−07
1e−06
2e−06
mutation rate
RFunrooted metric: error 0.01
Distance between true and inferred trees
0
10
20
30
tsibiny
tsipoly
fastARG
Aweaver
2e−08
5e−08
1e−07
2e−07
5e−07
1e−06
2e−06
mutation rate
RFunrooted metric: error 0.1
Distance between true and inferred trees
0
10
20
30
tsibiny
tsipoly
fastARG
Aweaver
2e−08
5e−08
1e−07
2e−07
5e−07
1e−06
2e−06
mutation rate


SPRunrooted metric: error 0
Distance between true and inferred trees
0
2
4
6
8
tsibiny
tsipoly
fastARG
Aweaver
2e−08
5e−08
1e−07
2e−07
5e−07
1e−06
2e−06
mutation rate
SPRunrooted metric: error 0.01
Distance between true and inferred trees
0
2
4
6
8
tsibiny
tsipoly
fastARG
Aweaver
2e−08
5e−08
1e−07
2e−07
5e−07
1e−06
2e−06
mutation rate
SPRunrooted metric: error 0.1
Distance between true and inferred trees
0
2
4
6
8
tsibiny
tsipoly
fastARG
Aweaver
2e−08
5e−08
1e−07
2e−07
5e−07
1e−06
2e−06
mutation rate


pathunrooted metric: error 0
Distance between true and inferred trees
0
20
40
60
80
tsibiny
tsipoly
fastARG
Aweaver
2e−08
5e−08
1e−07
2e−07
5e−07
1e−06
2e−06
mutation rate
pathunrooted metric: error 0.01
Distance between true and inferred trees
0
20
40
60
80
tsibiny
tsipoly
fastARG
Aweaver
2e−08
5e−08
1e−07
2e−07
5e−07
1e−06
2e−06
mutation rate
pathunrooted metric: error 0.1
Distance between true and inferred trees
0
20
40
60
80
tsibiny
tsipoly
fastARG
Aweaver
2e−08
5e−08
1e−07
2e−07
5e−07
1e−06
2e−06
mutation rate