

Full genetic p-values in 3 datasets (8df)

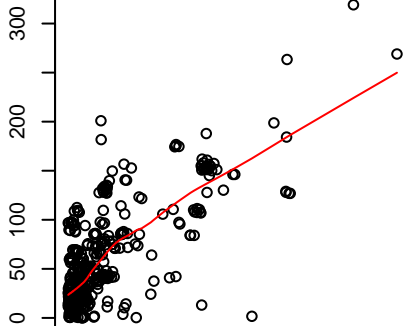
0 50 100 200 300

BSGS

0.73

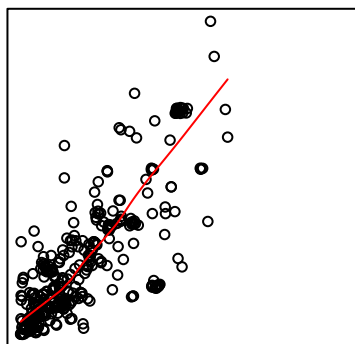
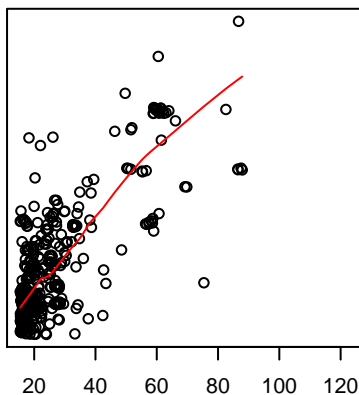
NaN

120
100
80
60
40
20



EGCUT

NaN



Fehrmann

0 50 100 150 200 250