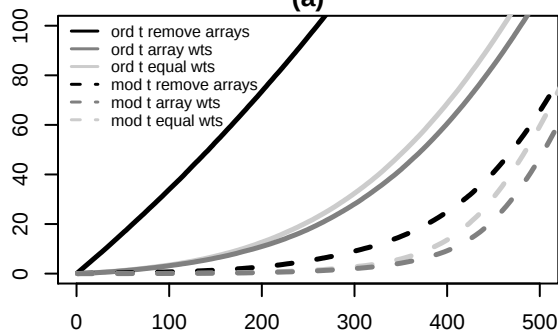


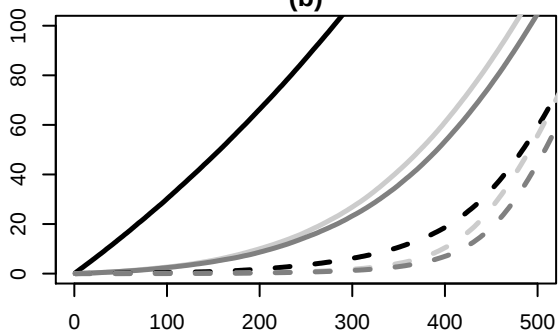
Normal

(a)

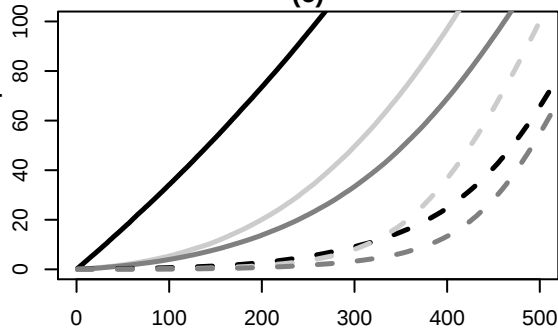


Non-normal

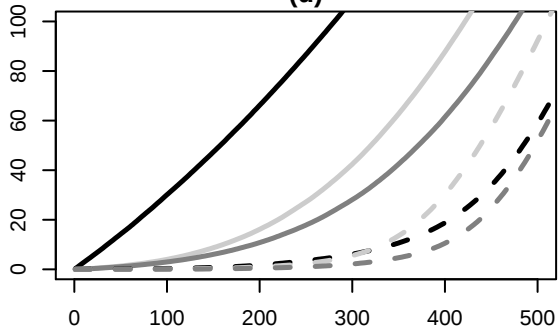
(b)



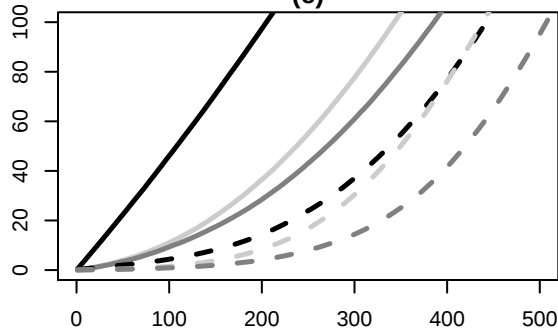
(c)



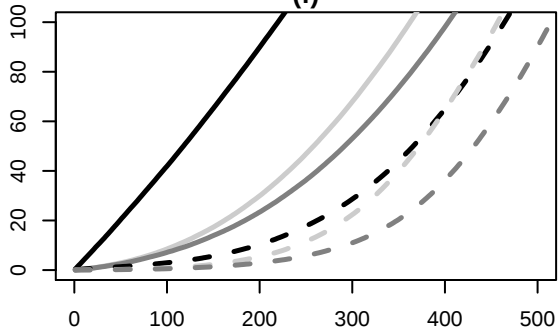
(d)



(e)



(f)



Number of false positives

Number of genes selected