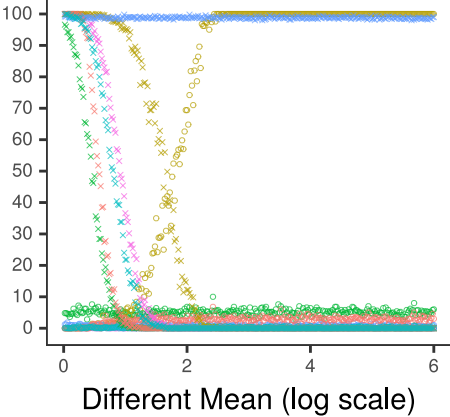
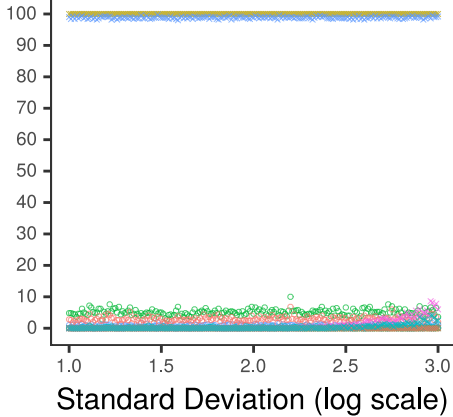
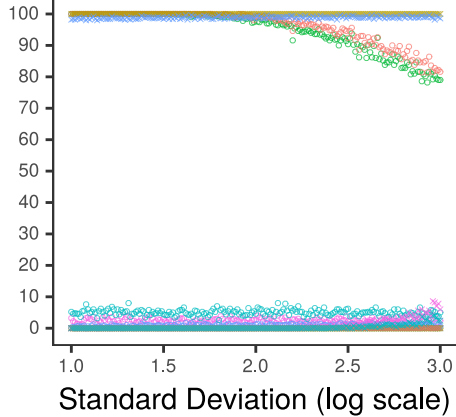


A**B****C****methods**

ANCOM

KW

metagenomeSeq

ANCOMBC

LEfSe

mp_diff_analysis

type

○ FPR × FNR