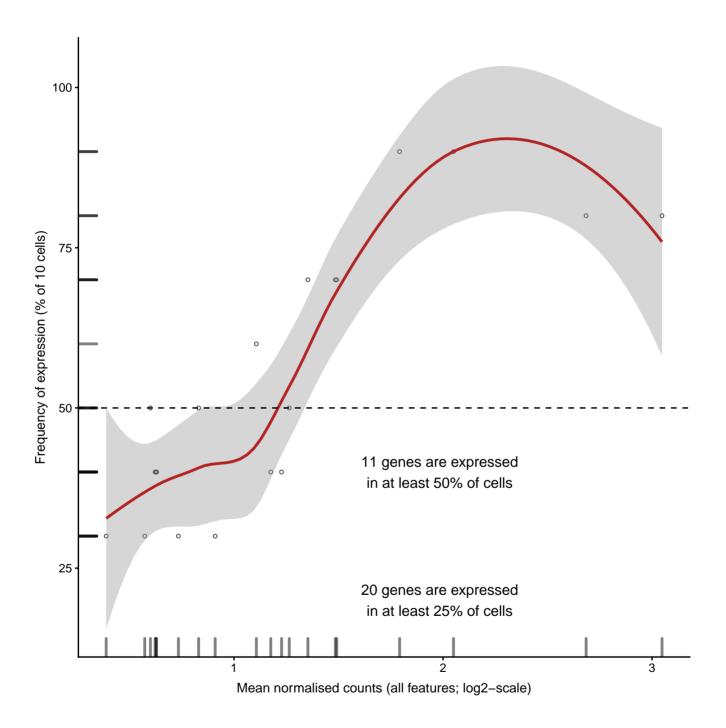


Log-total number of expressed features



Top 20 account for 100% of total RNA18S5 ACTB · GAPDH -| | | | O| | | | | COX1 UBC · CLU · CHI3L1 · COX2 -10 PTPRZ1 -Feature ND4 · FTL · TUBA1A total\_features EEF1A1 -ATP6 | 11 12 ACTG1 13 H3F3B TPI1 18 TMBIM6 -Feature control? FOS. O FALSE 20 80 40 60 % of total counts