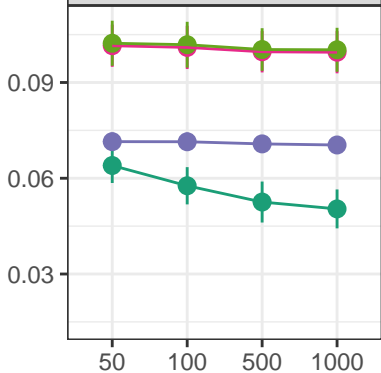
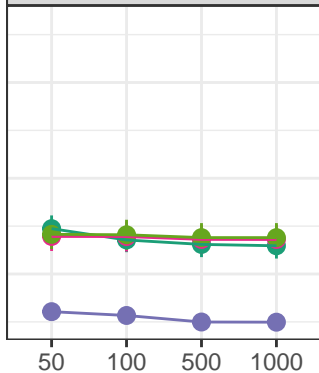


Mean absolute error

High ILS



Low ILS



Number of genes