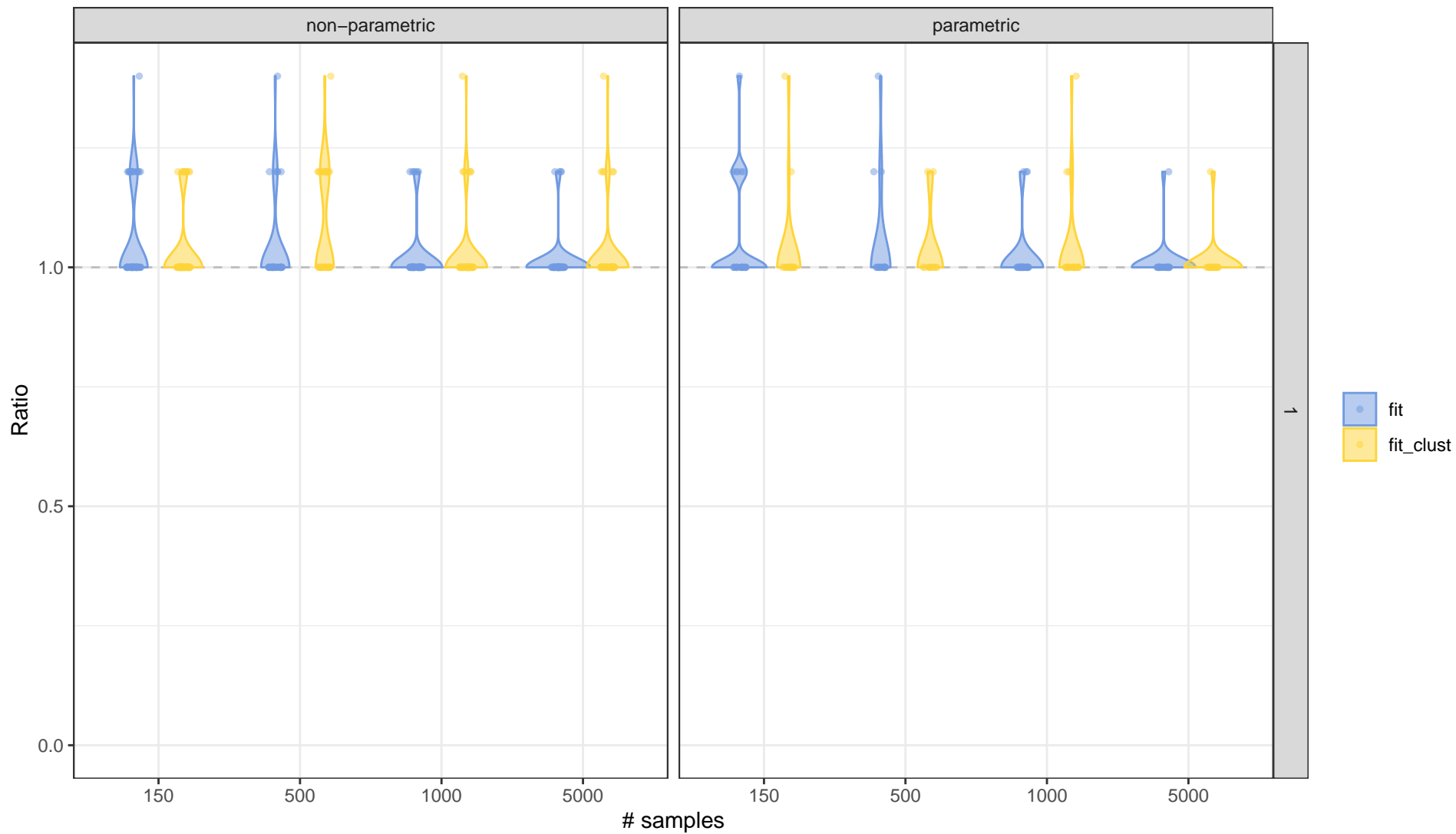
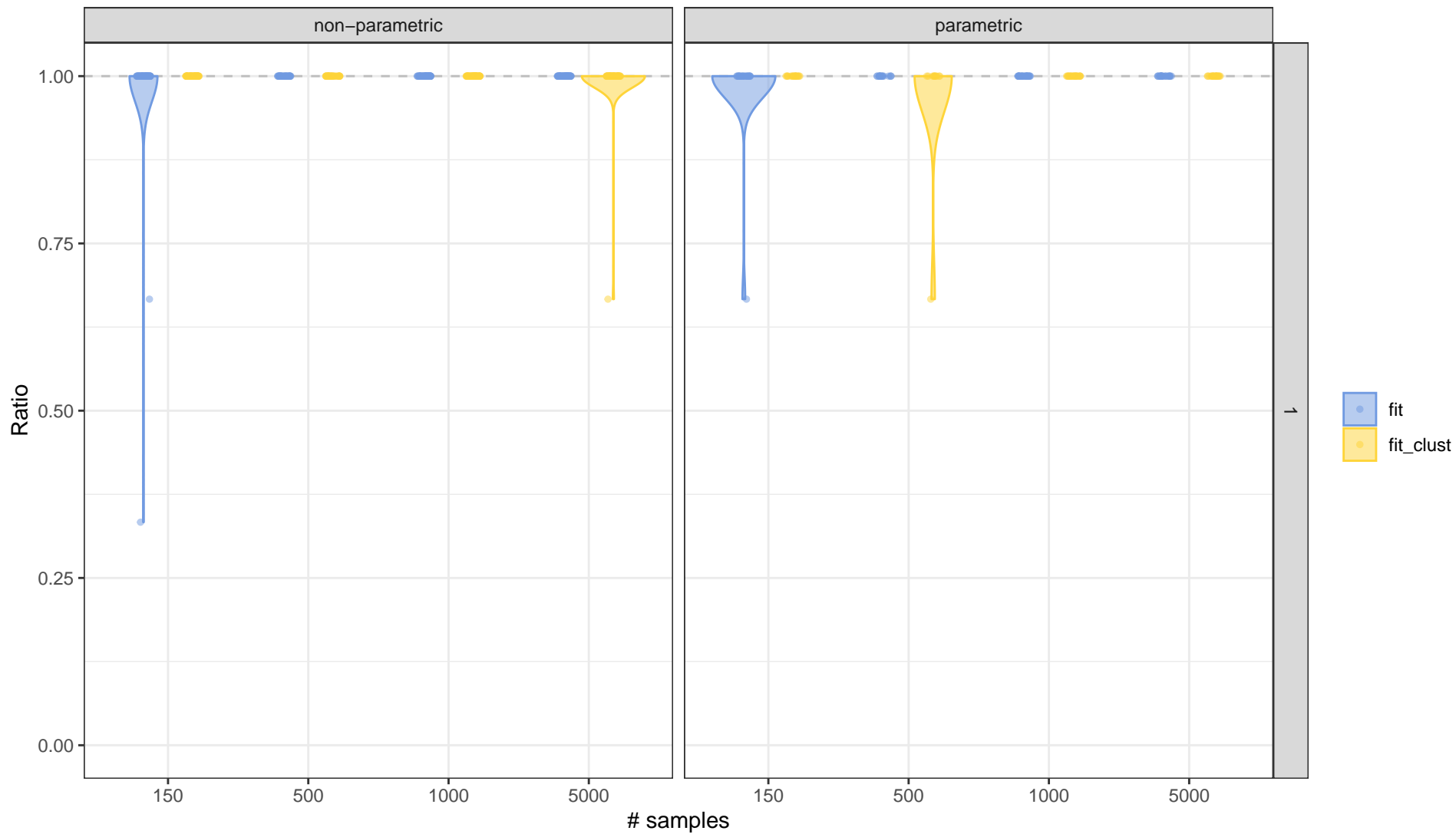


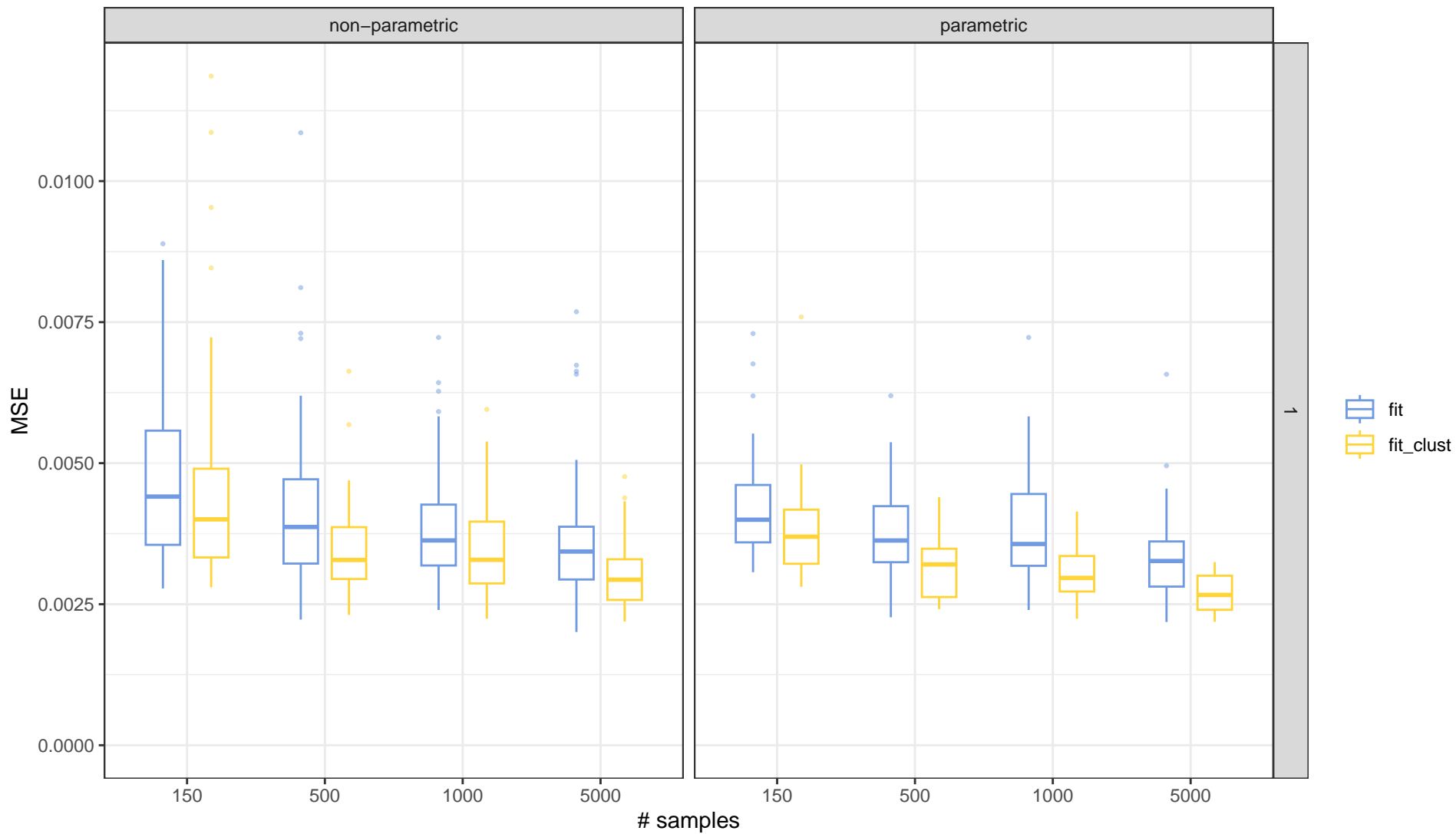
Ratio between number of signatures found and true



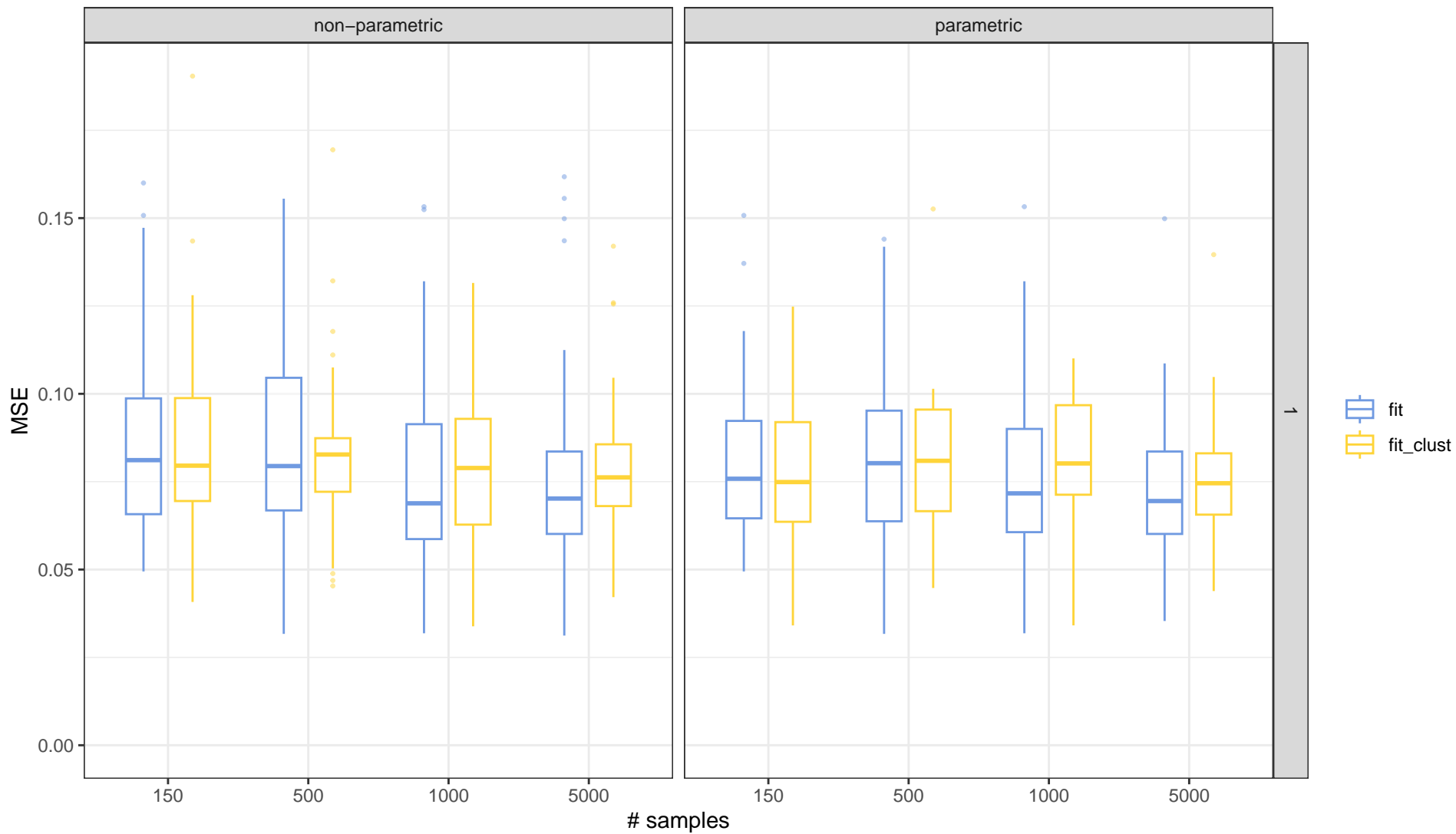
Ratio between number of rare signatures found and true



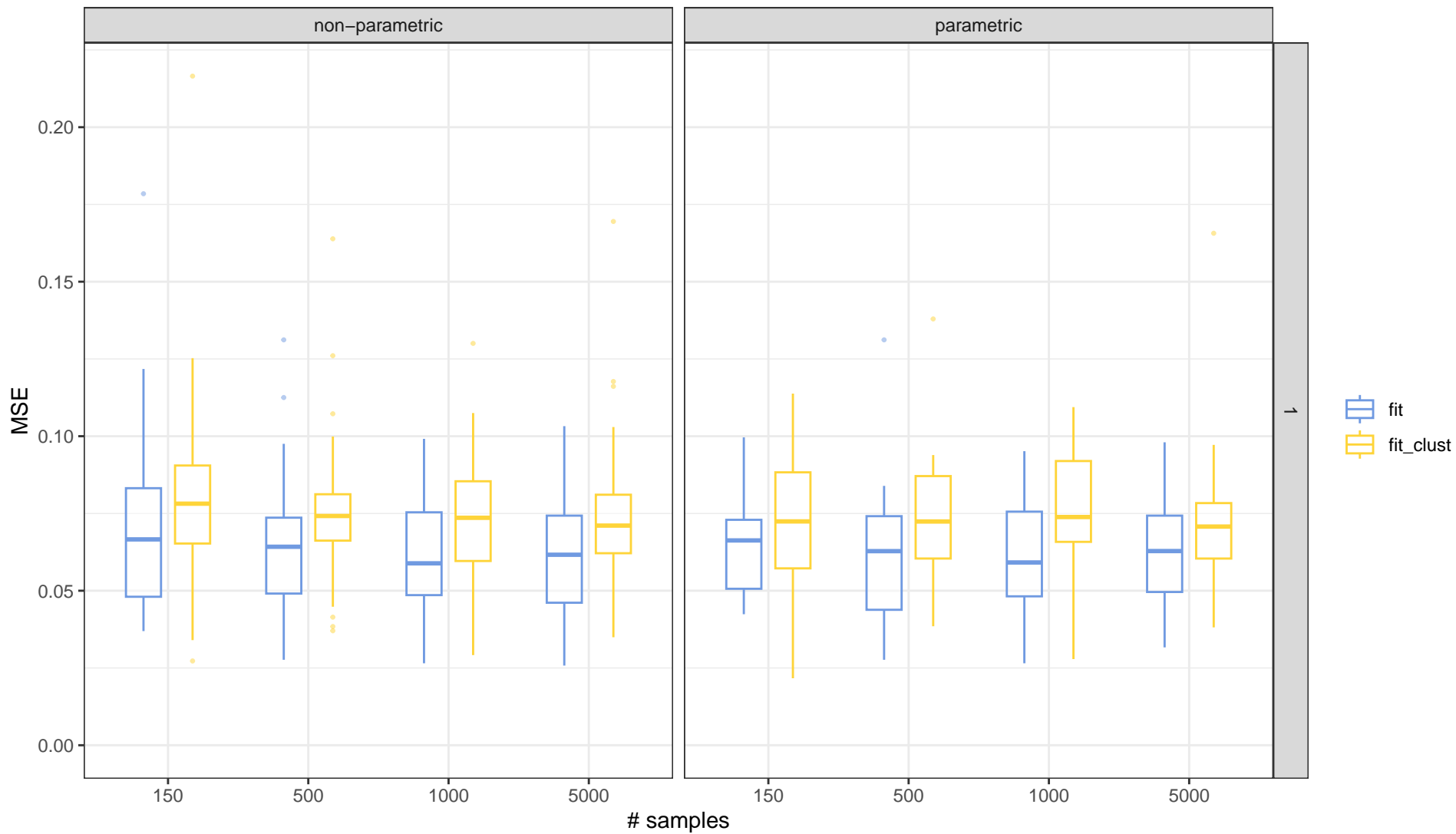
MSE computed between true and inferred counts



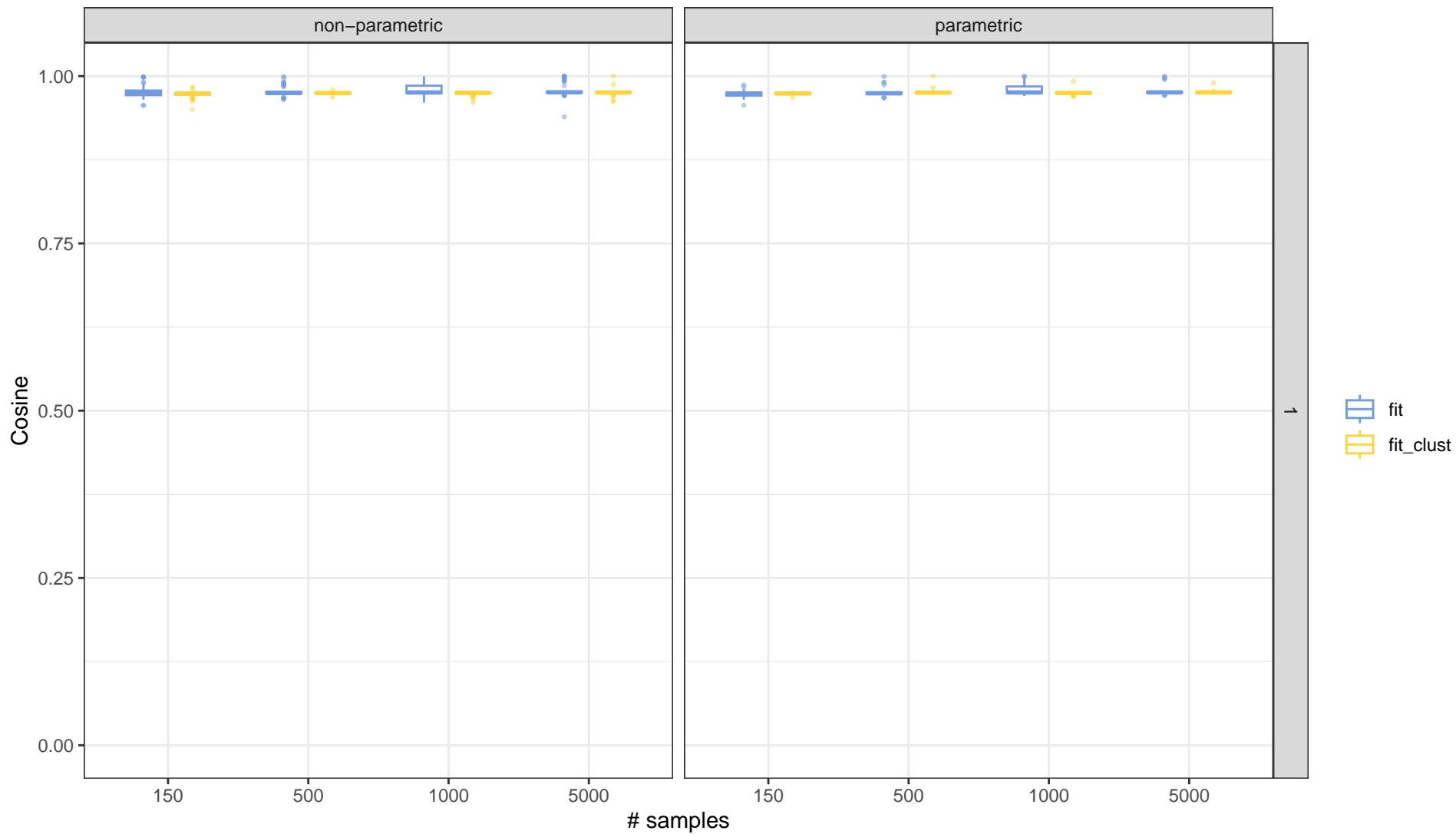
MSE computed between true and inferred exposures



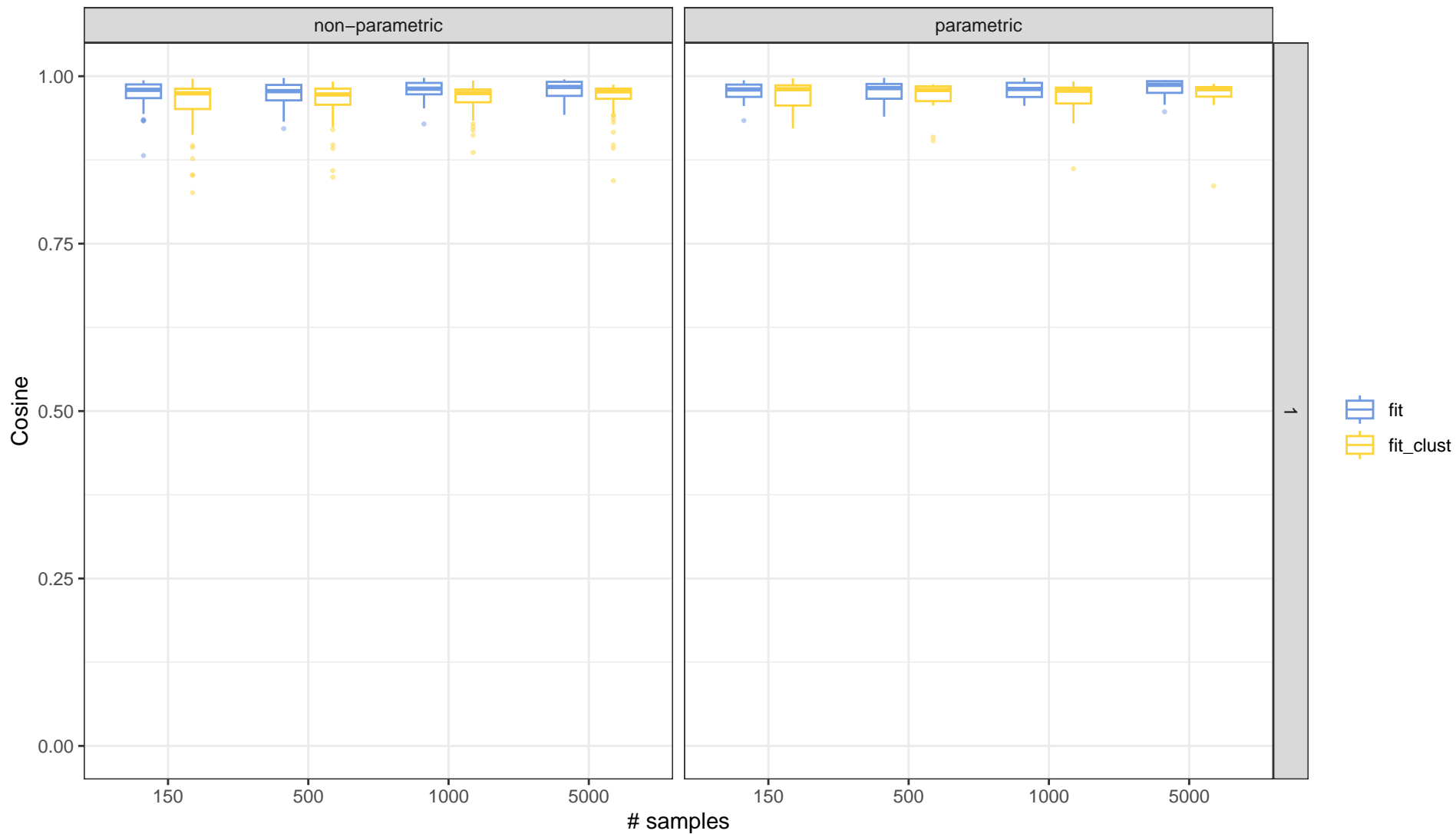
MSE computed between true and inferred rare exposures



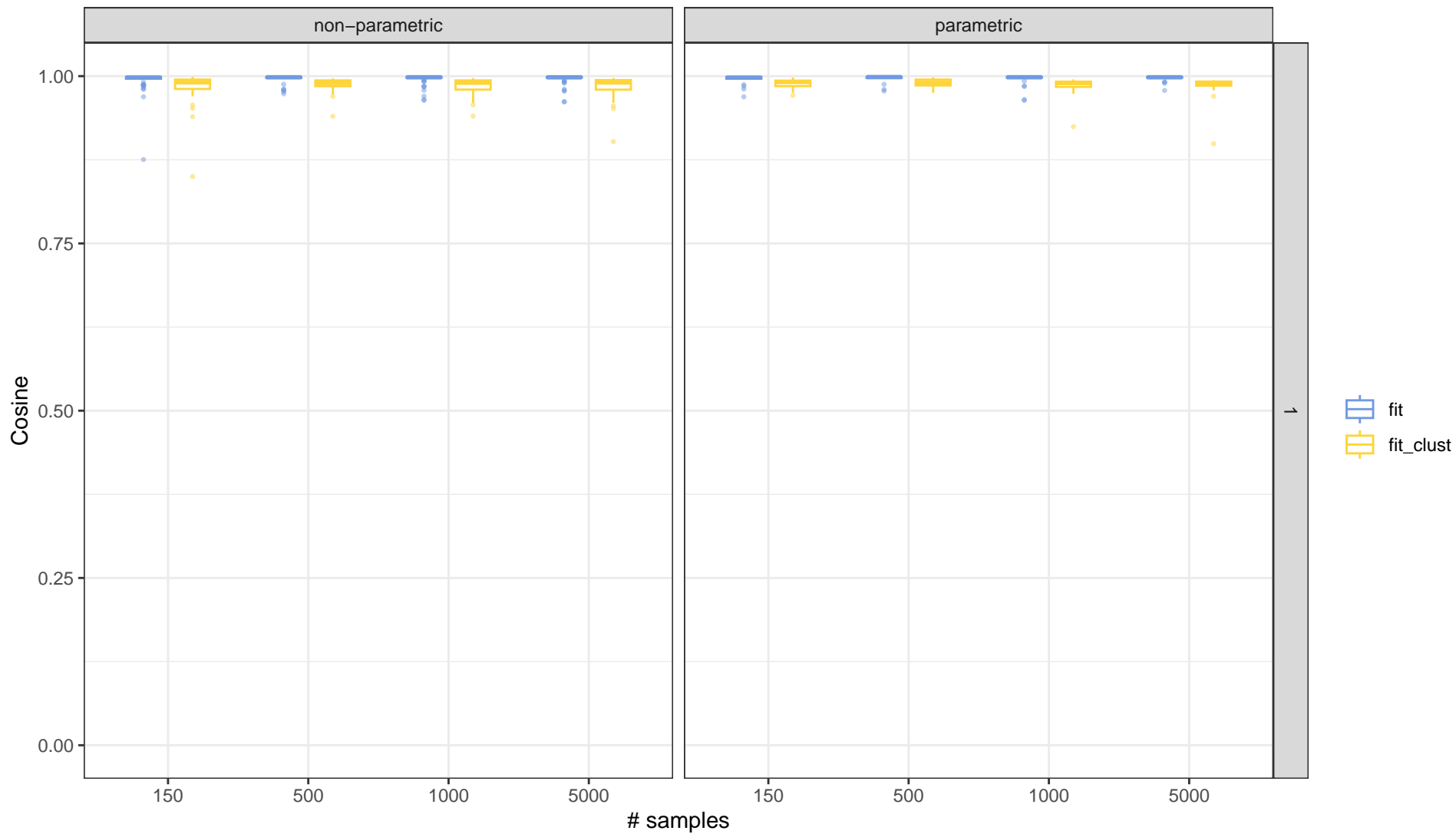
Cosine computed between true and inferred signatures



Cosine computed between true and inferred exposures

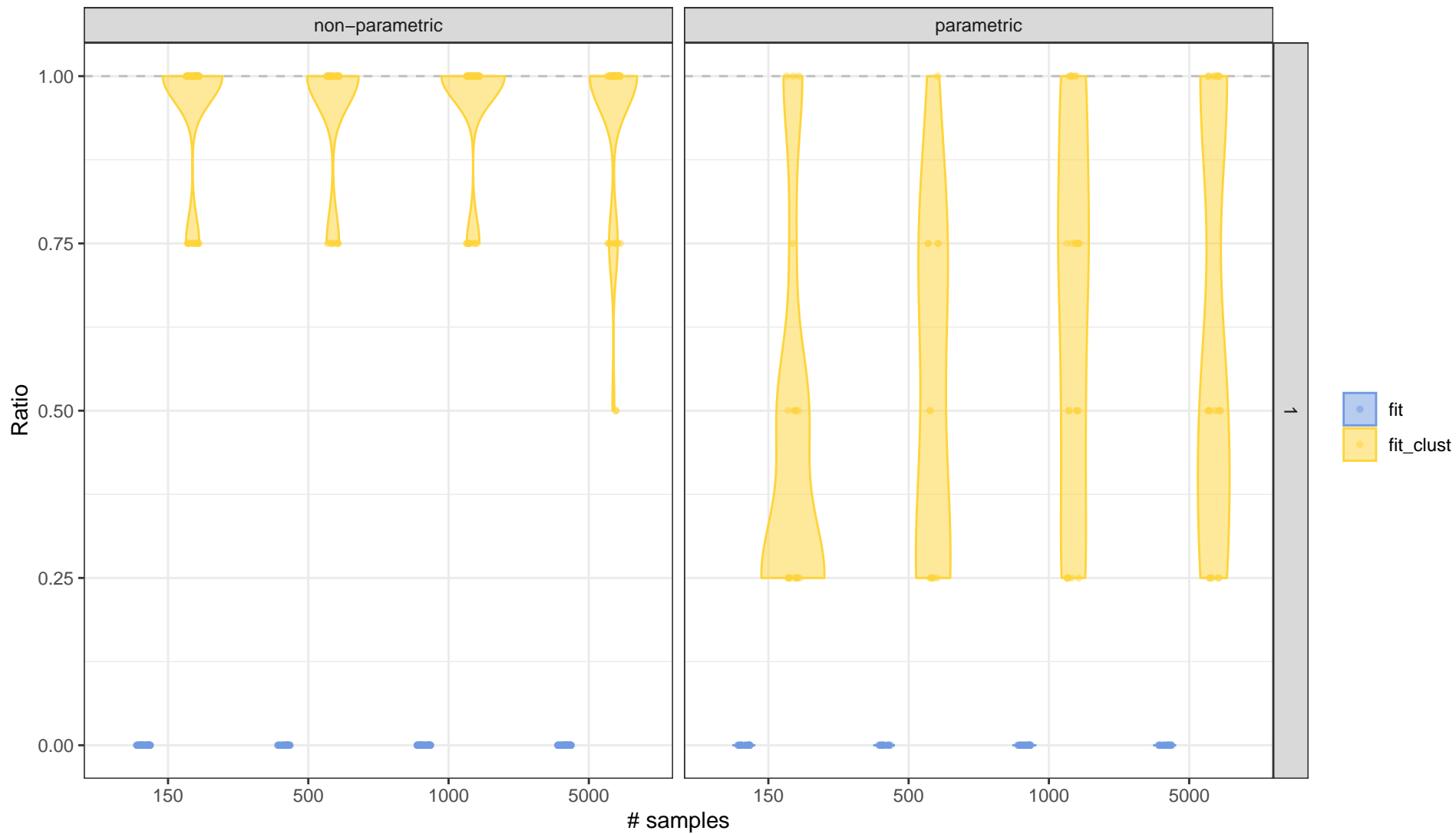


Cosine computed between true and inferred rare exposures

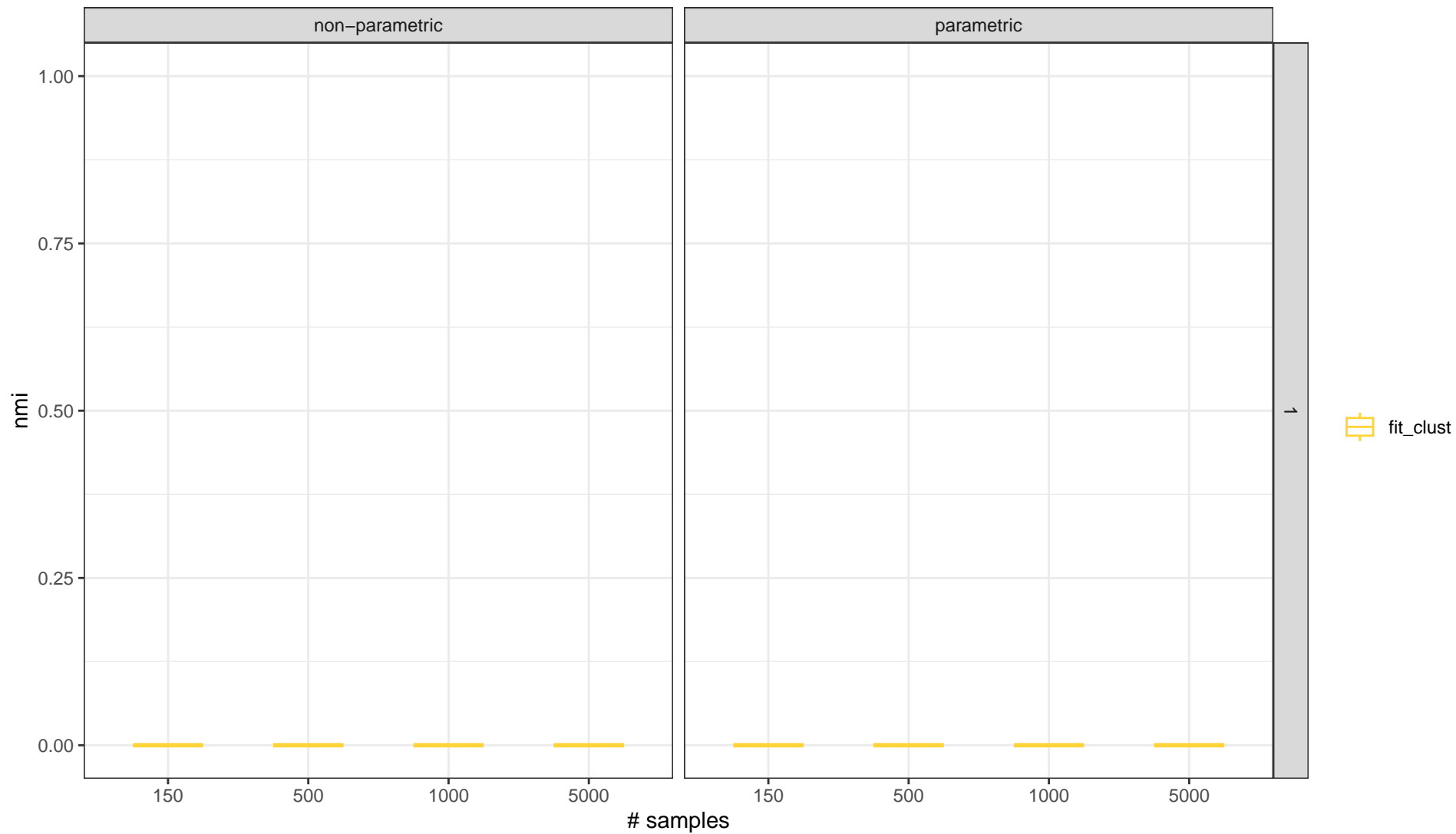




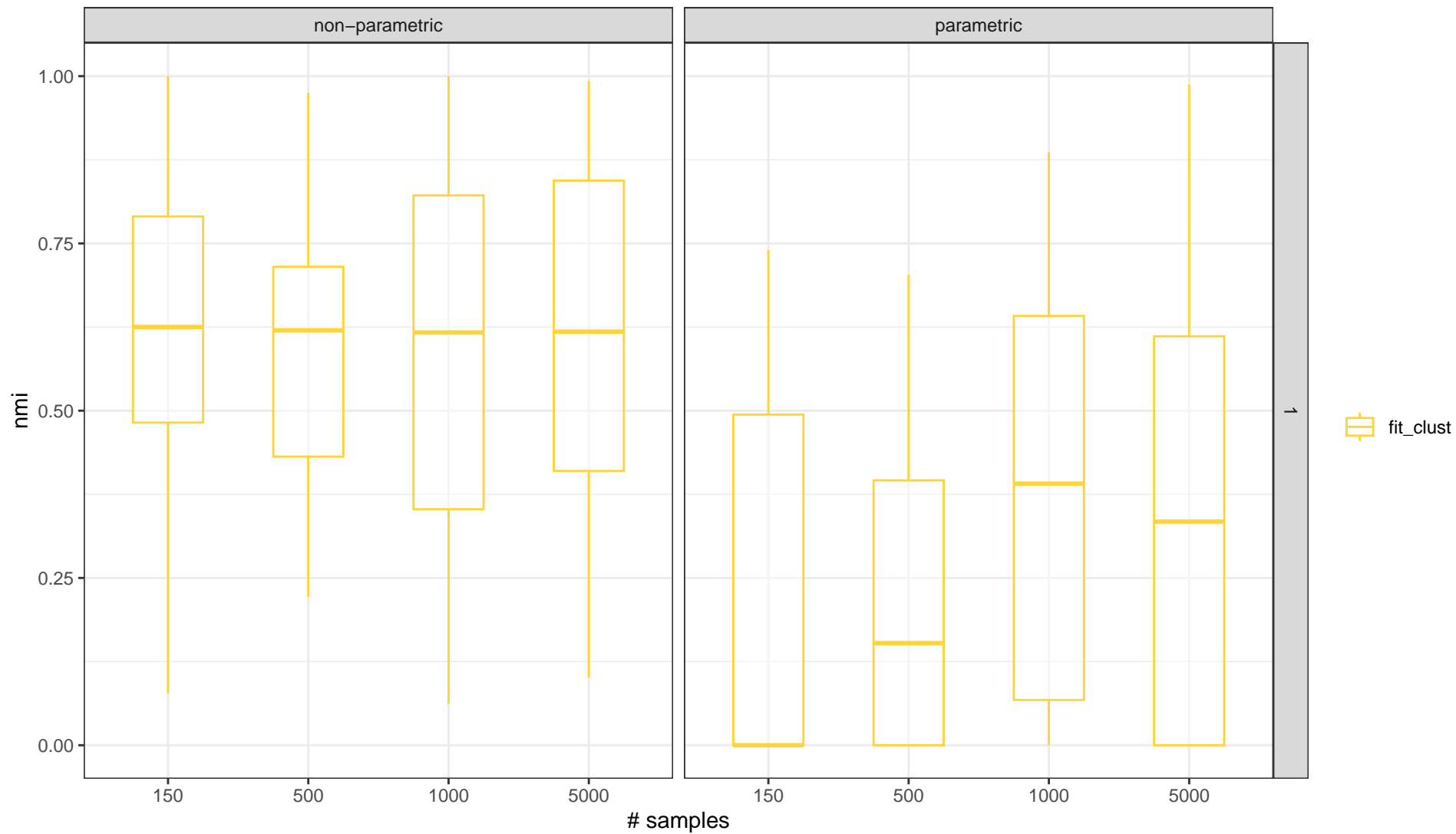
Ratio between number of clusters found and true



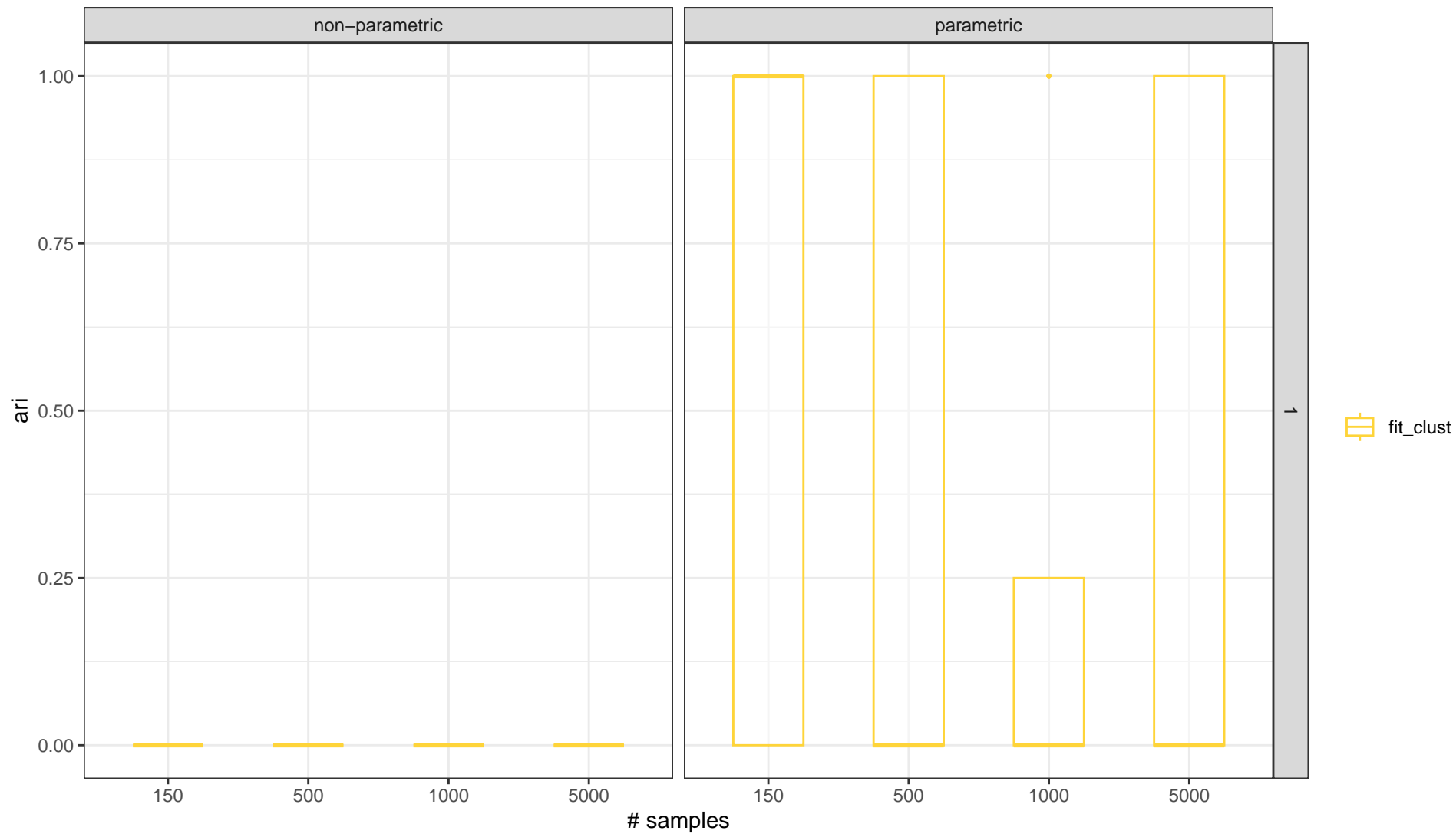
nmi computed between true and inferred NA



nmi computed between true and inferred rare



ari computed between true and inferred NA



ari computed between true and inferred rare

