

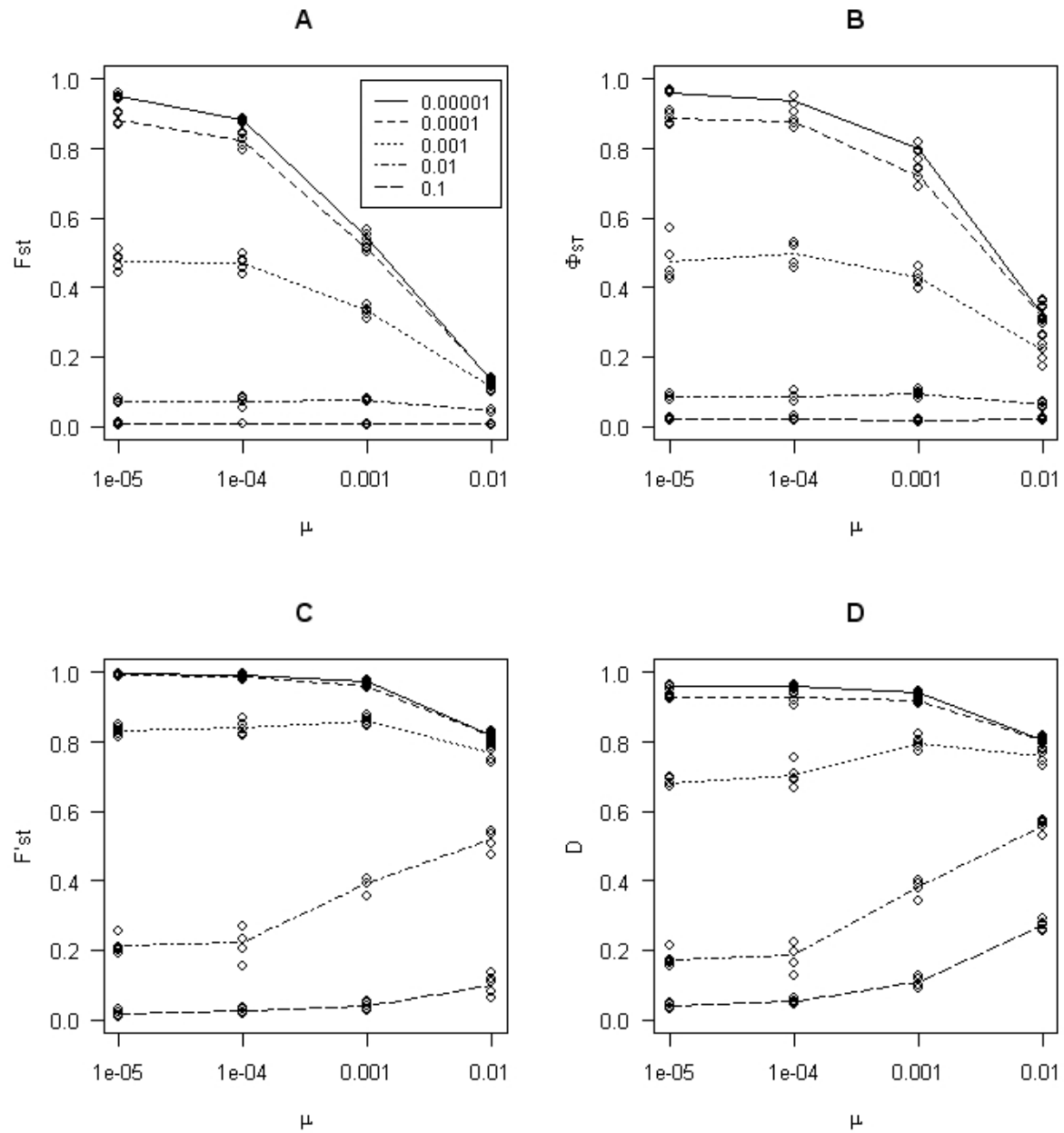


Figure S1. Results of forward simulations for self-fertilisation rate of 0.9. The mixed mutation model was used. Shown are the relationships with different estimators of genetic differentiation to different migration and mutation rates. X-axis is the mutation rate; different types of lines correspond to different migration rates as in the legend in panel A. Different estimators are F_{ST} , Φ_{ST} , F'_{ST} and D in panels A, B, C and D respectively.