Figure S4 legend: Changes in the distributions of $T_0$ with varying recombination rates in different demographic models: A, population bottleneck model with bottleneck severity $c = 0.05$; B, exponential population growth with growth rate $g = 500$; C, population subdivision model with migration rate between two subpopulations $m = 0.1$. Mean correlation coefficient of LD among polymorphic sites (average $\rho^2$) for each model is shown in parenthesis.