

Figure S5

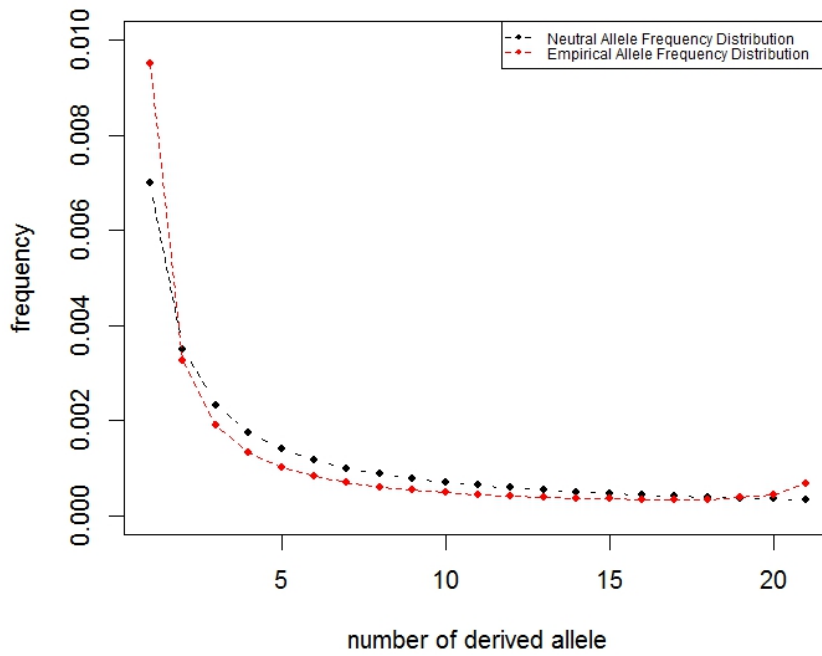


Figure S5 legend: Genome-wide empirical distribution of derived-allele frequency in the Rwanda *D. melanogaster* sample (22 sequences) in comparison with the standard neutral distribution for a sample of same size.