



FIGURE S2.—Tajima's D values for simulations of positive selection on standing variation. Simulations of 10Mb long regions after 50, 100, 200, 500 generations of selection ($s=0.1$). Only the region around the mutation is shown. The thin red lines show the values for 10 simulations of standing selection on an existing allele with a frequency of 0.1 and the thin blue lines are the values for 10 simulations without selection. The values are calculated for windows of size 100kb with a window step size of 10kb. The thick lines are the mean values for the 10 simulations in each scenario and both scenarios are based on the same set of haplotypes.