Figure S6  Signals of selection from various summary statistics in windows containing or flanking a soft sweep from recurrent mutation. For each summary statistic, we examined each individual simulation and located the window exhibiting the most extreme value (in the direction suggestive of a soft sweep). This figure shows the histogram of these locations for each statistic. The total genetic distance of each simulated chromosome ($\rho$) was 2100. The chromosome was subdivided into 21 equally sized windows ($\rho=100$) with a soft selective sweep (with $\alpha=1000$; mutation rate to the adaptive allele ranging from 1 to 2.5) occurring in the central window.