

Table S4 *P*-values from a two-sample *t*-test comparing $N_e s^a$ categories of simulated outcrossing and selfing populations under fixed dominance levels for all mutations

Dominance of mutations (<i>h</i>)	<1	1-10	10-100	>100
0.2	0.257	0.479	0.002**	<0.001***
0.5	0.163	0.415	0.609	0.651
0.8	<0.001***	0.181	0.030*	0.275

^a $N_e s$ is the product of N_e and the selection coefficient (*s*)

*Significant at the 5% level

**Significant at the 1% level

***Significant at the 0.1% level