Table S11: Root mean squared error (RMSE) and mean absolute error (MAE) values when predicting selection coefficient (s), initial frequency (f), and time of selection (T_{sel}) for CEU populations when trained and tested with simulations using recombination rate drawn from an exponential distribution with mean 10^{-8} truncated at three times the mean per site per generation or rate drawn from an empirical human recombination map. The values show RMSE and MAE measured between standardized log-scaled predicted and actual parameters.

Recombination rate	RMSE(s)	RMSE(f)	RMSE(T_{sel})	MAE(s)	MAE(f)	MAE(T_{sel})
10^{-8}	0.88	1.02	0.69	0.68	0.82	0.51
Human map	0.96	1.11	0.84	0.83	0.93	0.68