

Table S13: Root mean squared error (RMSE) and mean absolute error (MAE) values when predicting selection coefficient (s), initial frequency (f), and time of donor-recipient split (T_{split}) under adaptive introgression scenarios for YRI and CEU populations. The values show RMSE and MAE measured between standardized log-scaled predicted and actual parameters in simulated data.

Population	RMSE(s)	RMSE(f)	RMSE(T_{split})	MAE(s)	MAE(f)	MAE(T_{sel})
CEU	0.93	1.00	0.98	0.91	0.96	0.94
YRI	0.97	0.91	0.99	0.93	0.88	0.95