Alpha 2.05 Rho 1 Mu 0.01 -2.5 **-**⊡ ⊡ • ⊡ -5.0 o l log_of_mean_homozygosity -7.5 **-**Simulation Values Linear Fit -10.0 **-**-12.5 **-**2.5 0.0 7.5 5.0 log_of_distance