Alpha 2.05 Rho 100 Mu 0.01 ⊡ **-7.5 -**⊡ • • ⊡ -10.0 log_of_mean_homozygosity Simulation Values –12.5 **-**Linear Fit -15.0 **-**-17.5 **-**0.0 2.5 7.5 5.0 log_of_distance