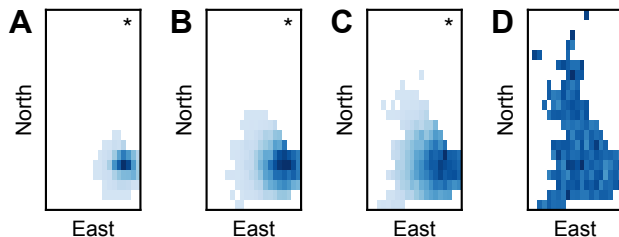
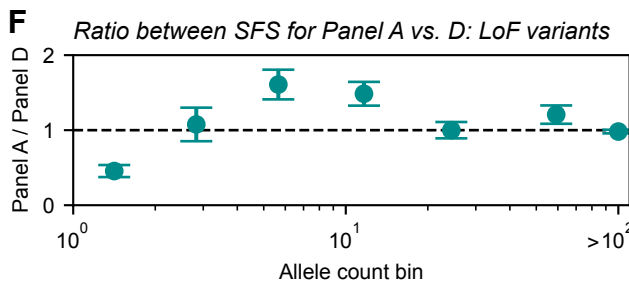


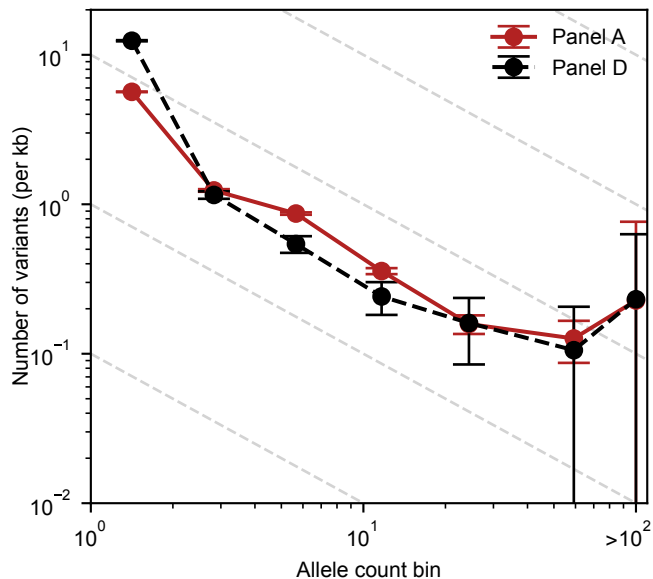
Sampling distributions



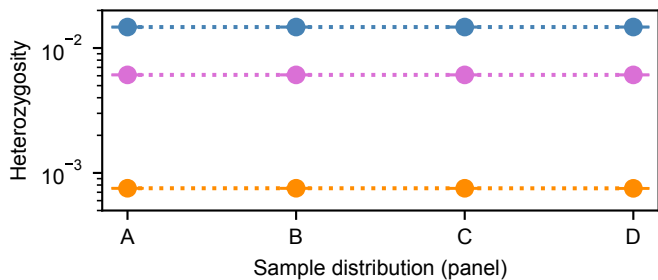
*Results shown below for samples A-C are averaged over three center locations



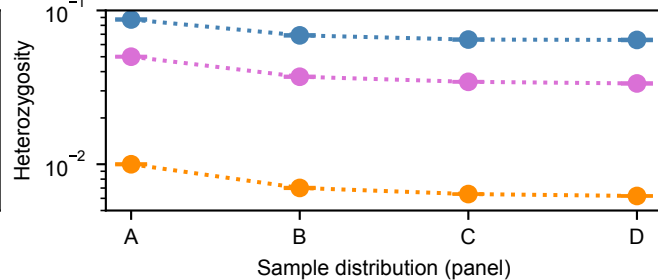
E SFS: LoF variants



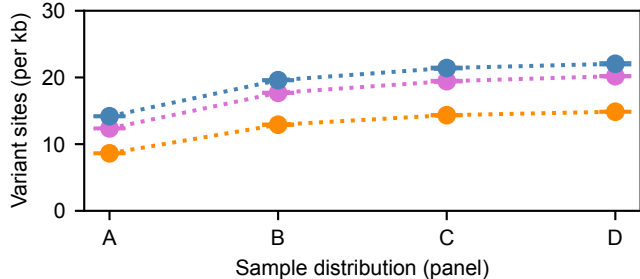
G Heterozygosity: all sites



H Heterozygosity: variant sites



I Variant sites (per kb)



J Singletons (per kb)

