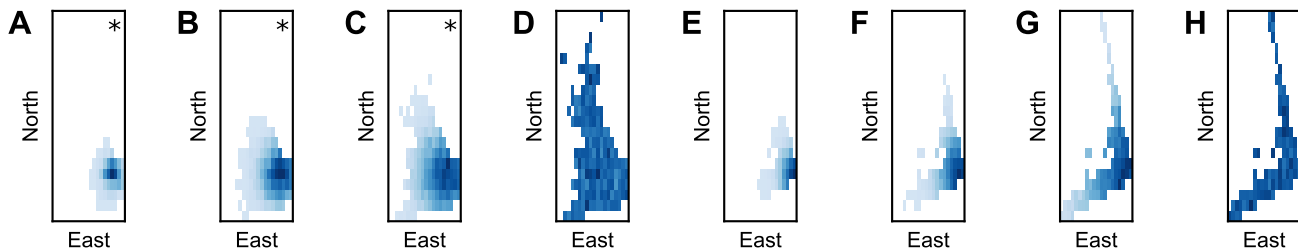
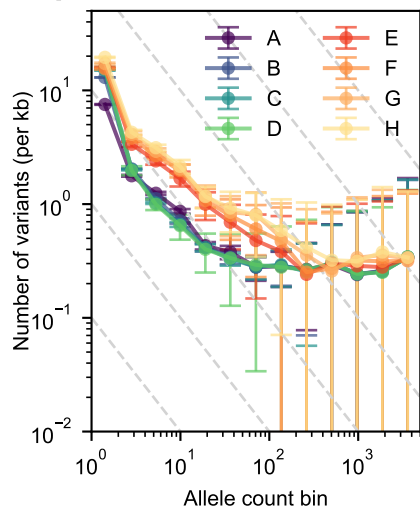


Sampling distributions

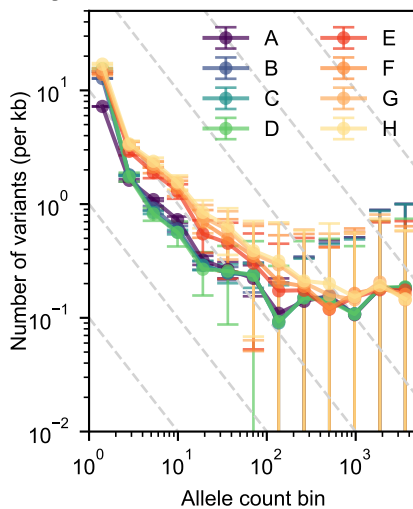


*Results shown below for samples A-C are averaged over three center locations

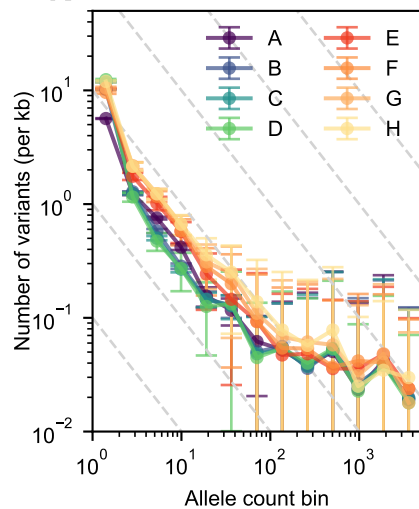
I SFS: Synonymous variants (per kb)



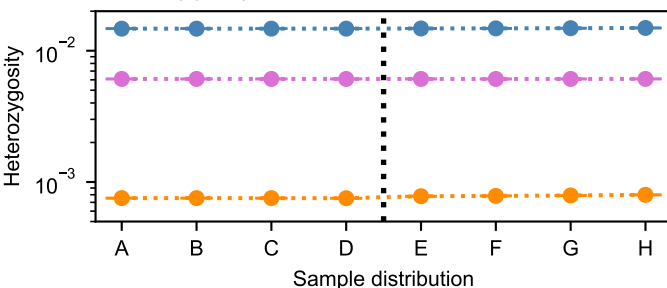
J SFS: Missense variants (per kb)



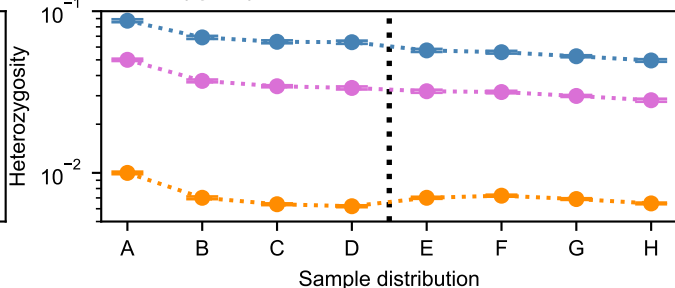
K SFS: LoF variants (per kb)



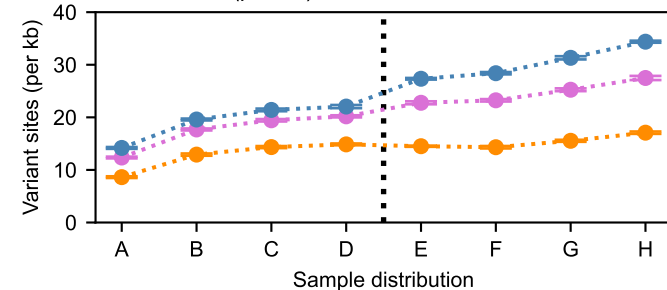
L Heterozygosity: all sites



M Heterozygosity: variant sites



N Variant sites (per kb)



O Singletons (per kb)

