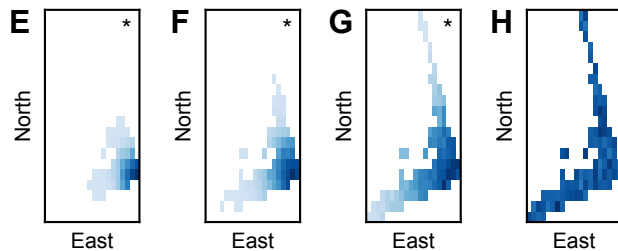
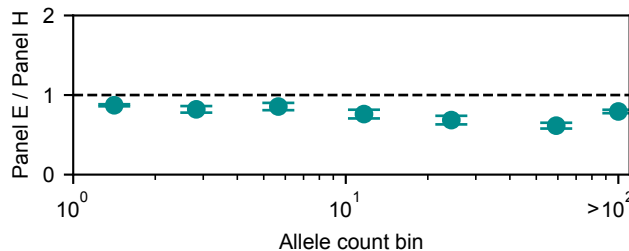


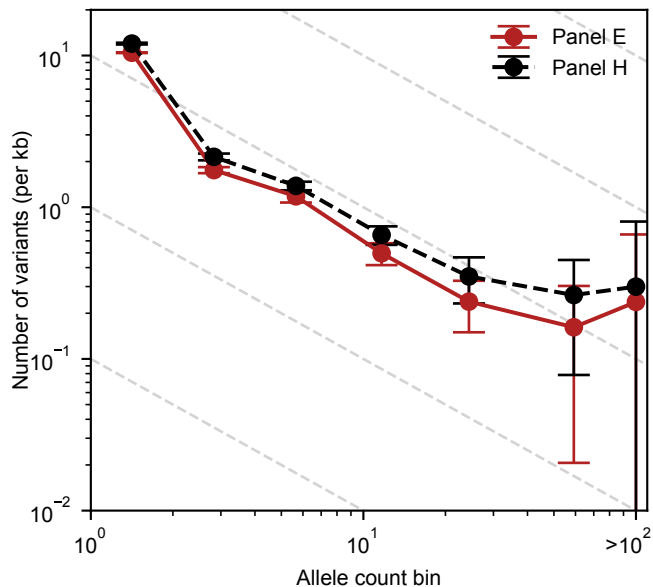
*Sampling distributions*



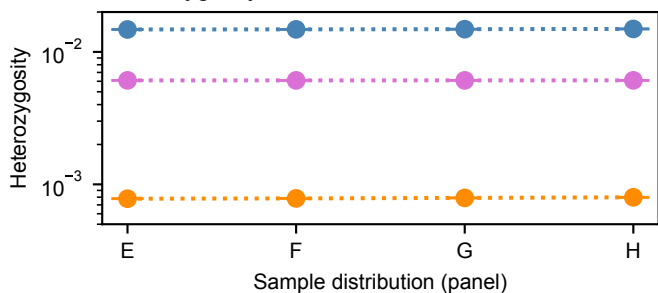
*Ratio between SFS for Panel E vs. H: LoF variants*



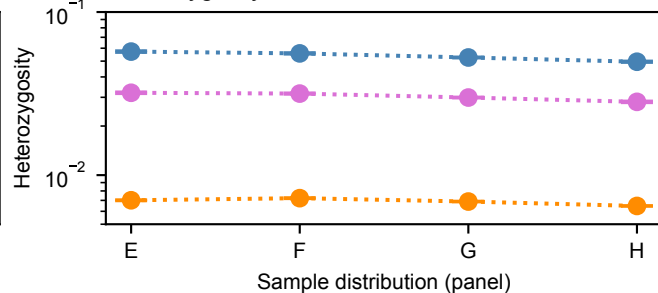
*SFS: LoF variants*



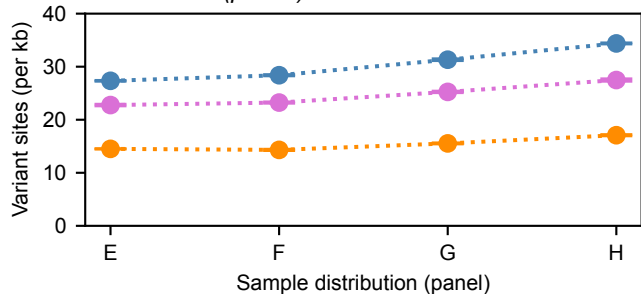
*Heterozygosity: all sites*



*Heterozygosity: variant sites*



*Variant sites (per kb)*



*Singltons (per kb)*

