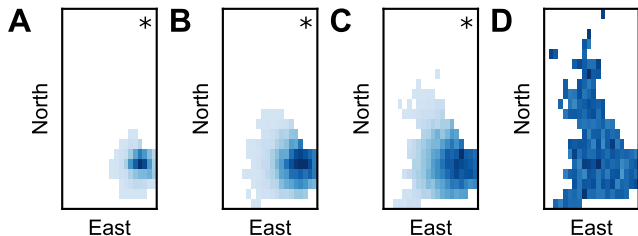
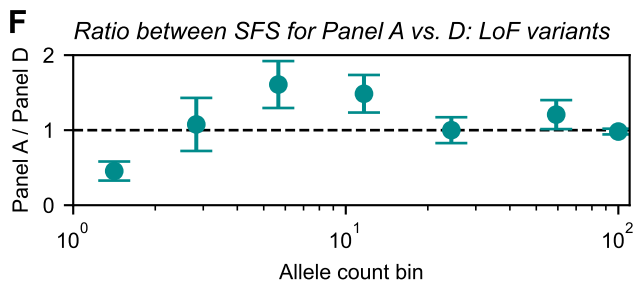


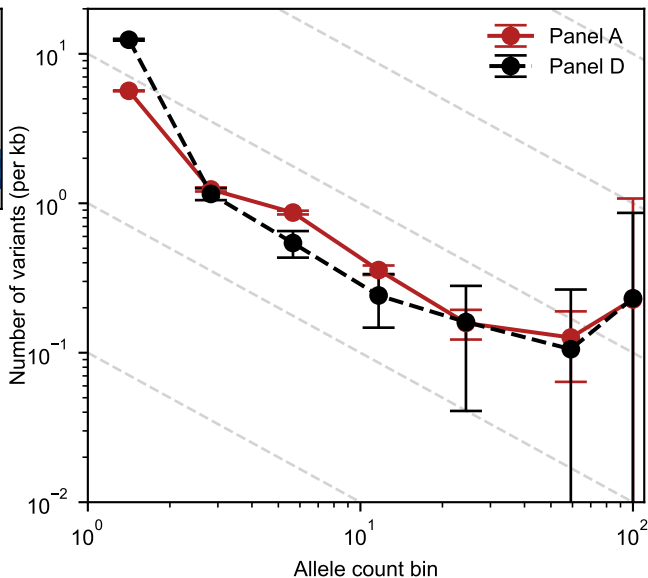
# Sampling distributions



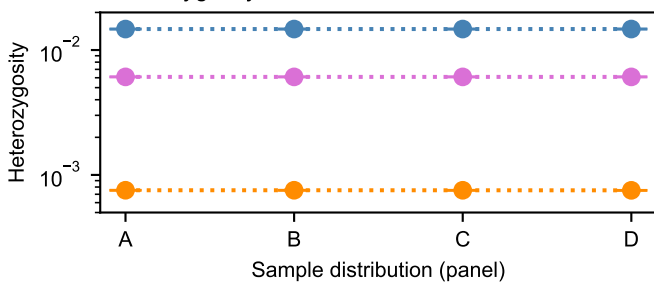
\*Results shown below for samples A-C are averaged over three center locations



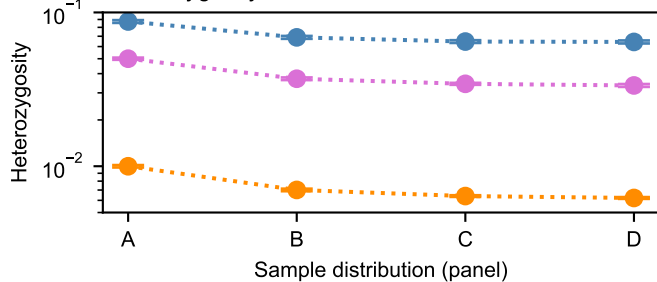
# SFS: LoF variants



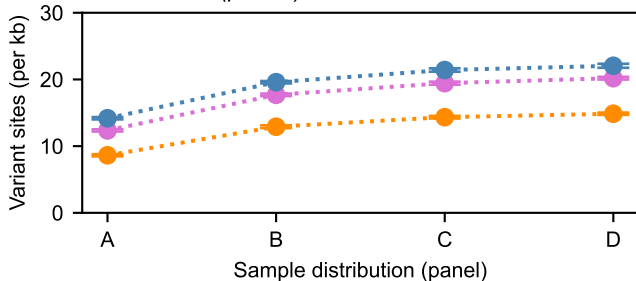
# Heterozygosity: all sites



# Heterozygosity: variant sites



# Variant sites (per kb)



# Singltons (per kb)

