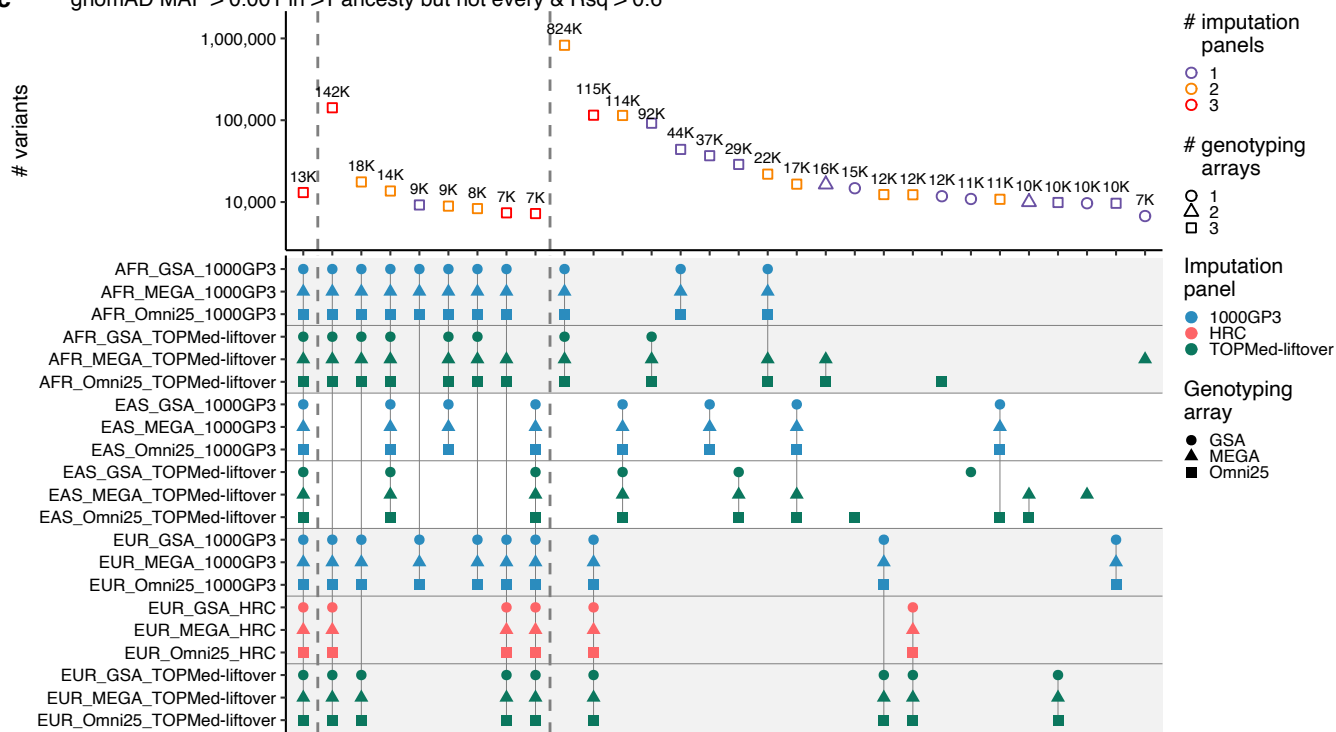


c gnomAD MAF > 0.001 in >1 ancestry but not every & Rsq > 0.6



d per-combination MAF > 0.001 & Rsq > 0.6

