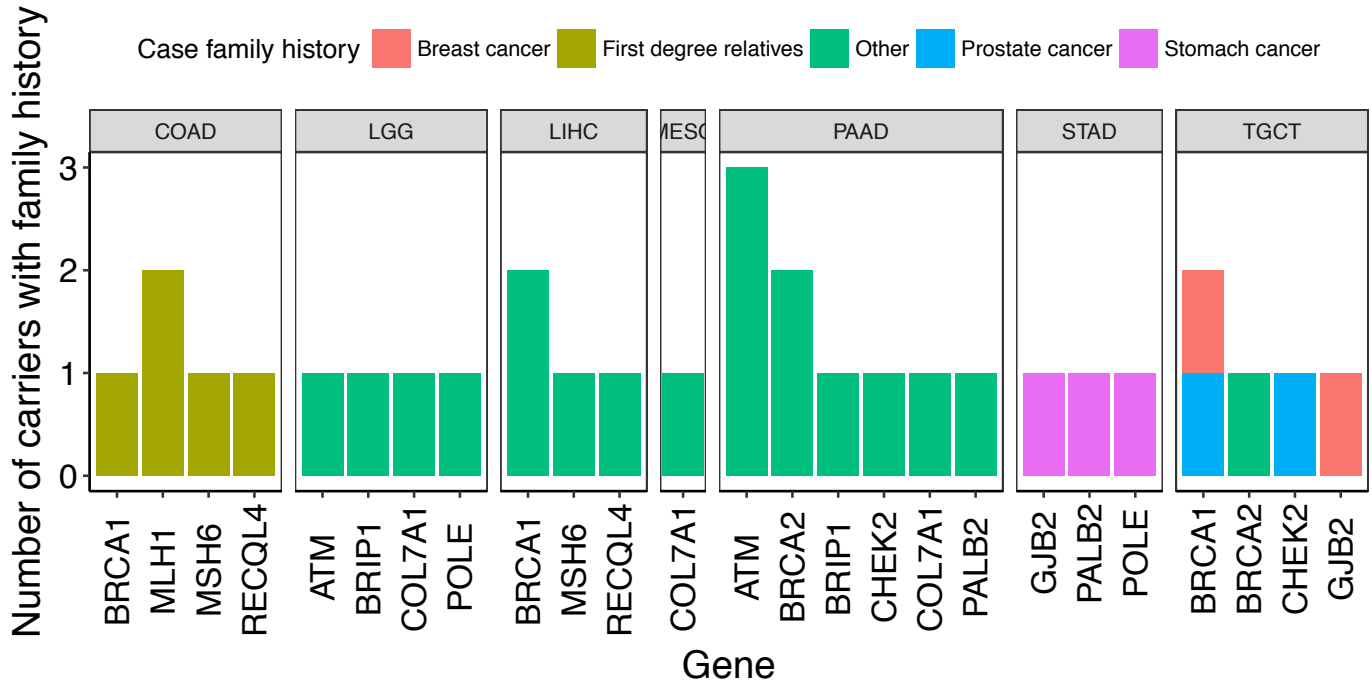


A



B

burden analysis power: collapsing 10 mutations on average
variants are rare, meaning MAF around 0.05%

