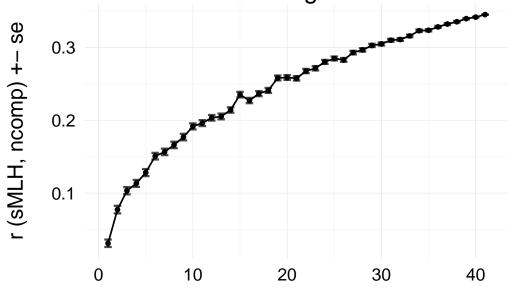
r between number of compounds and sMLH estimated from an increasing number of loci



number of loci