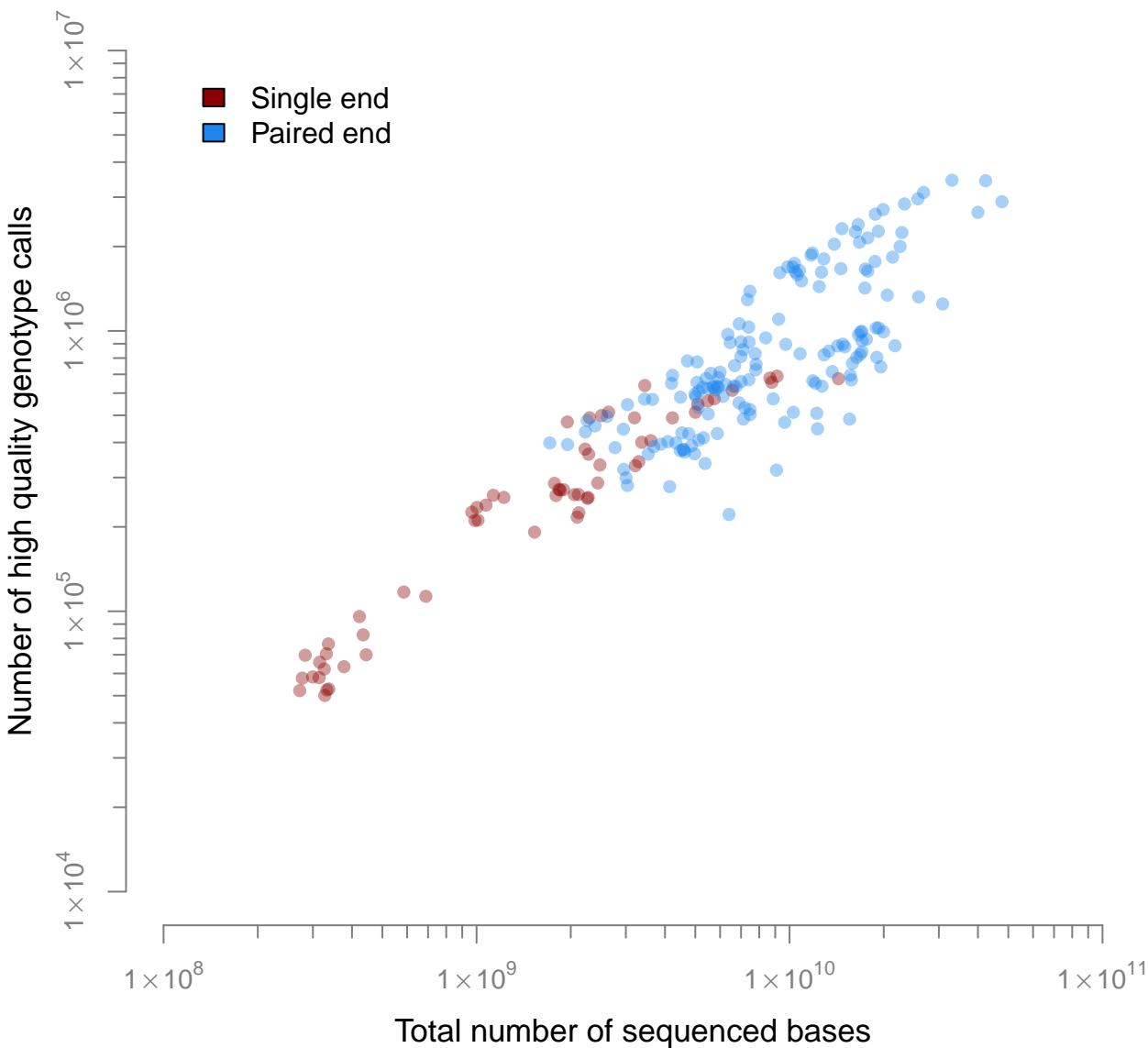


### Sequenced number of bases vs number of high quality genotype calls of 218 hematological samples



### Sequenced number of bases of the 1,262 samples used for eQTL and ASE analyses

