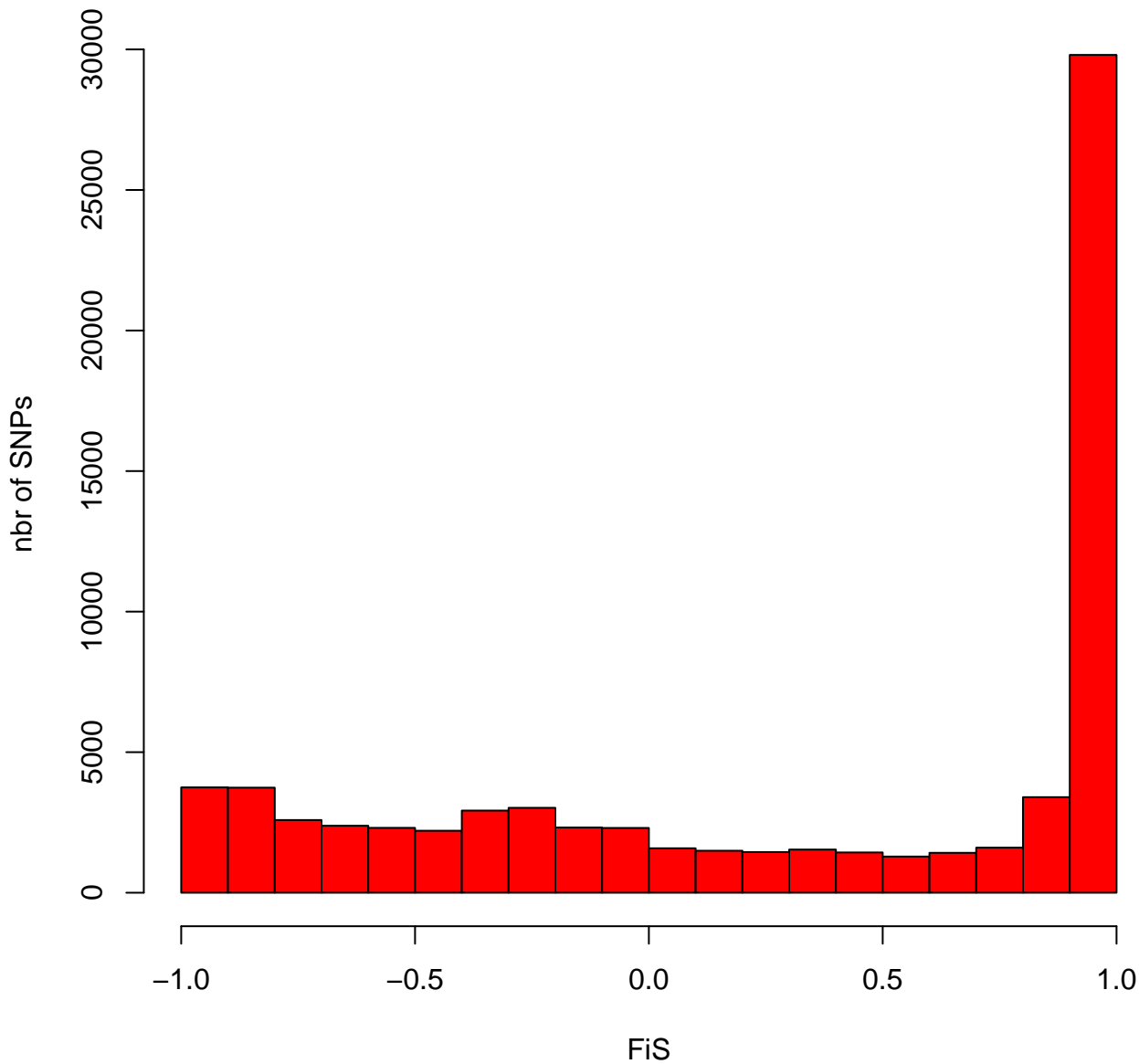
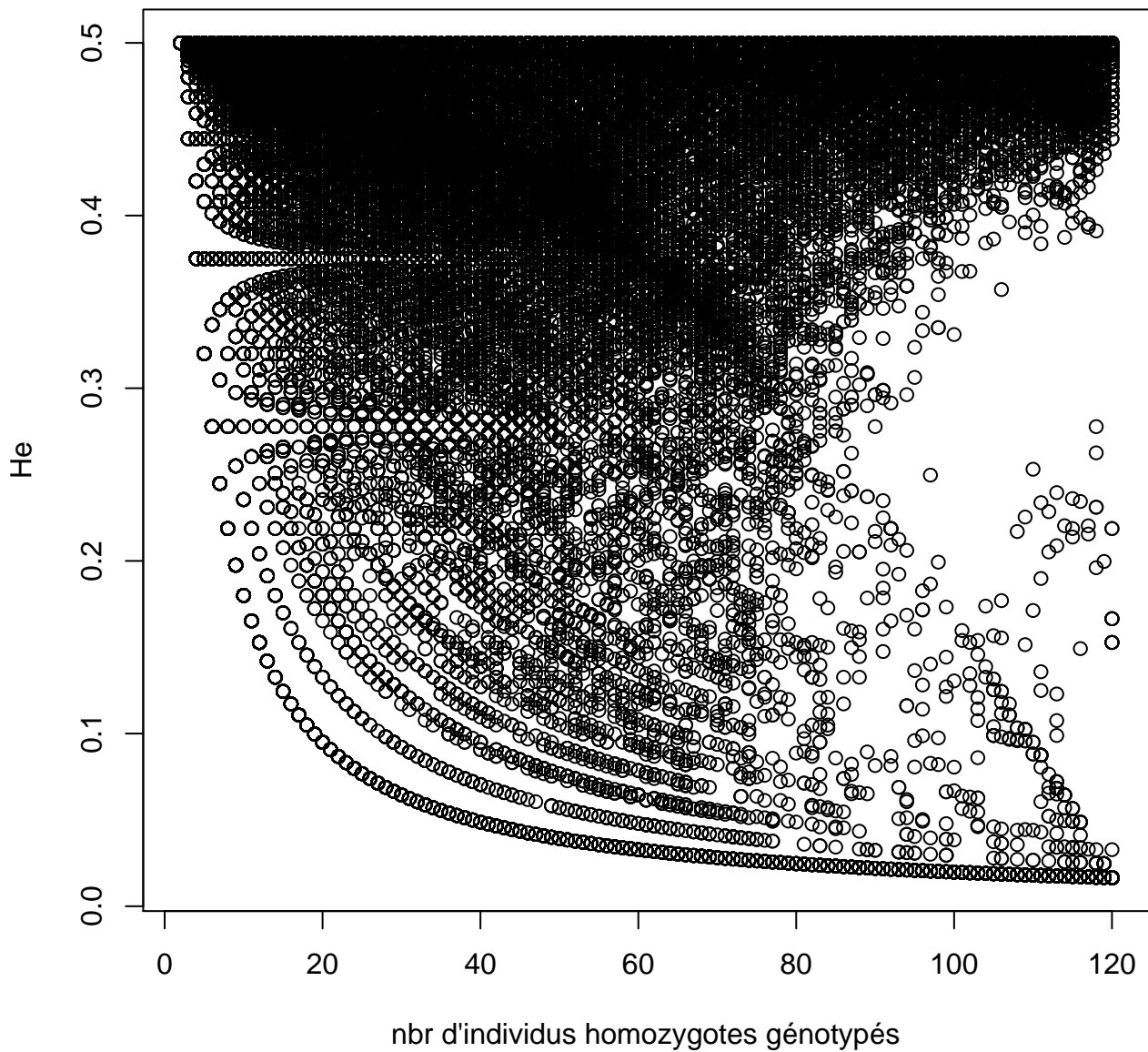
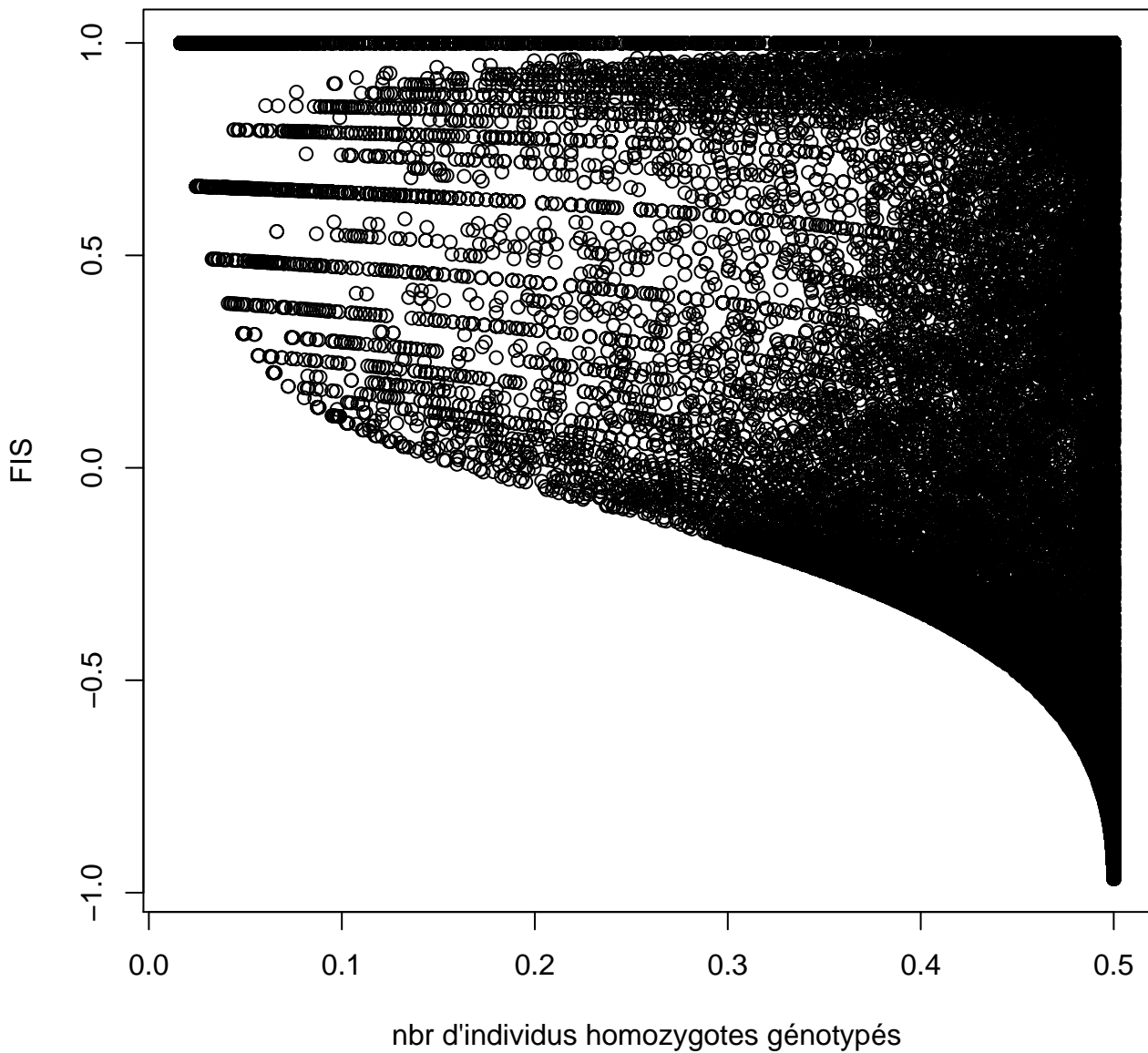
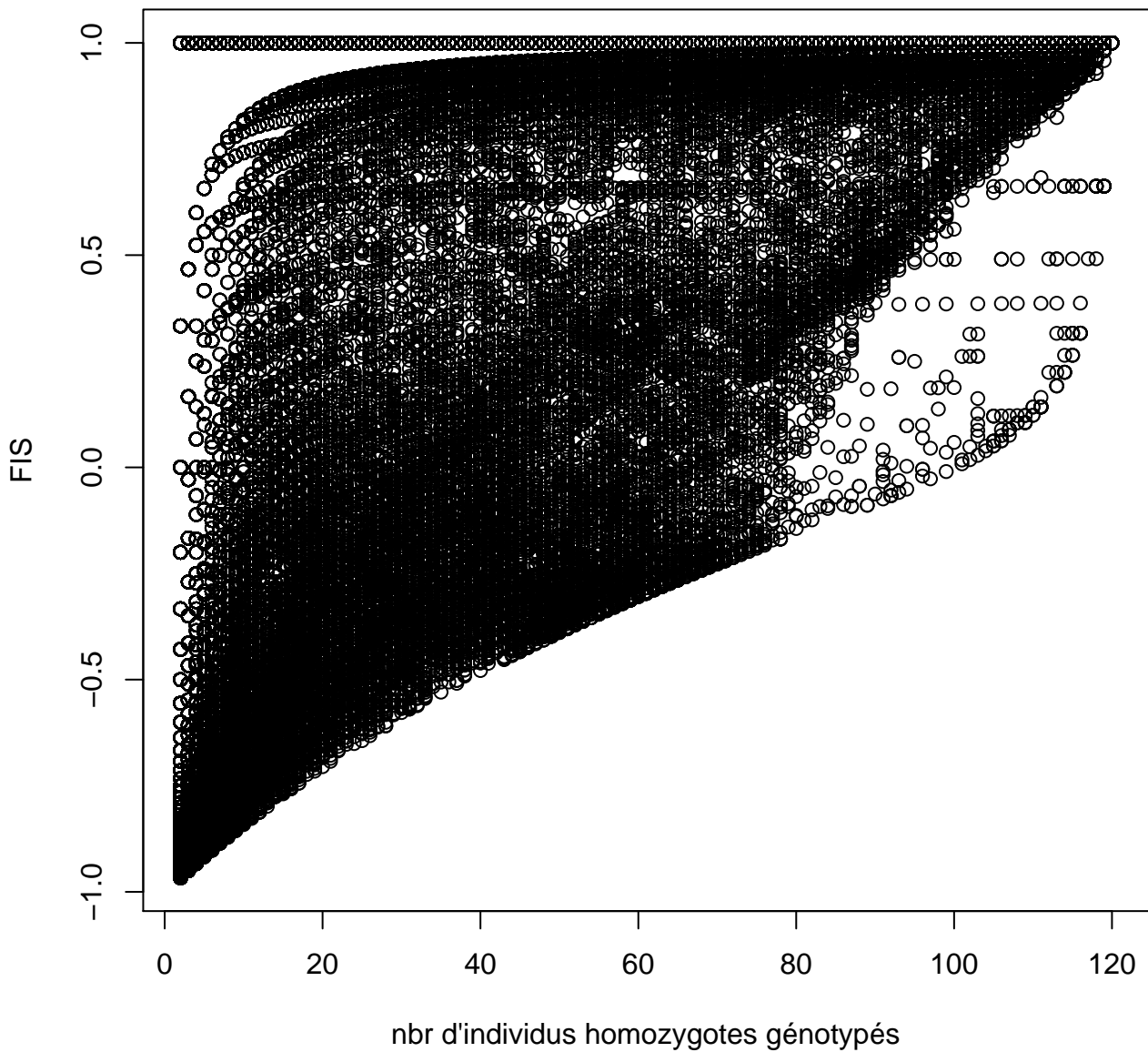


**FIS moyen : 0.3 | nbr of SNP : 73297**

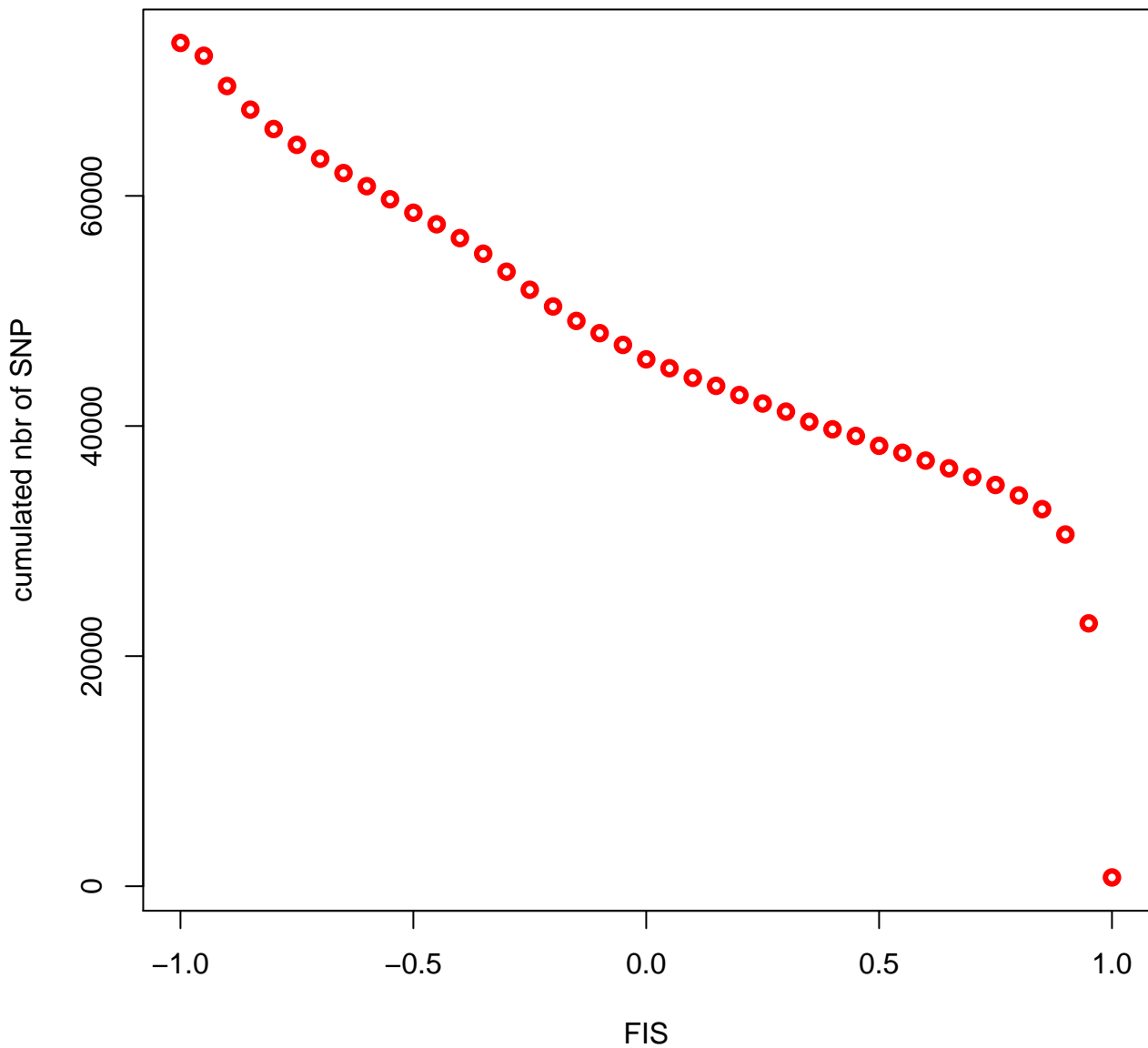




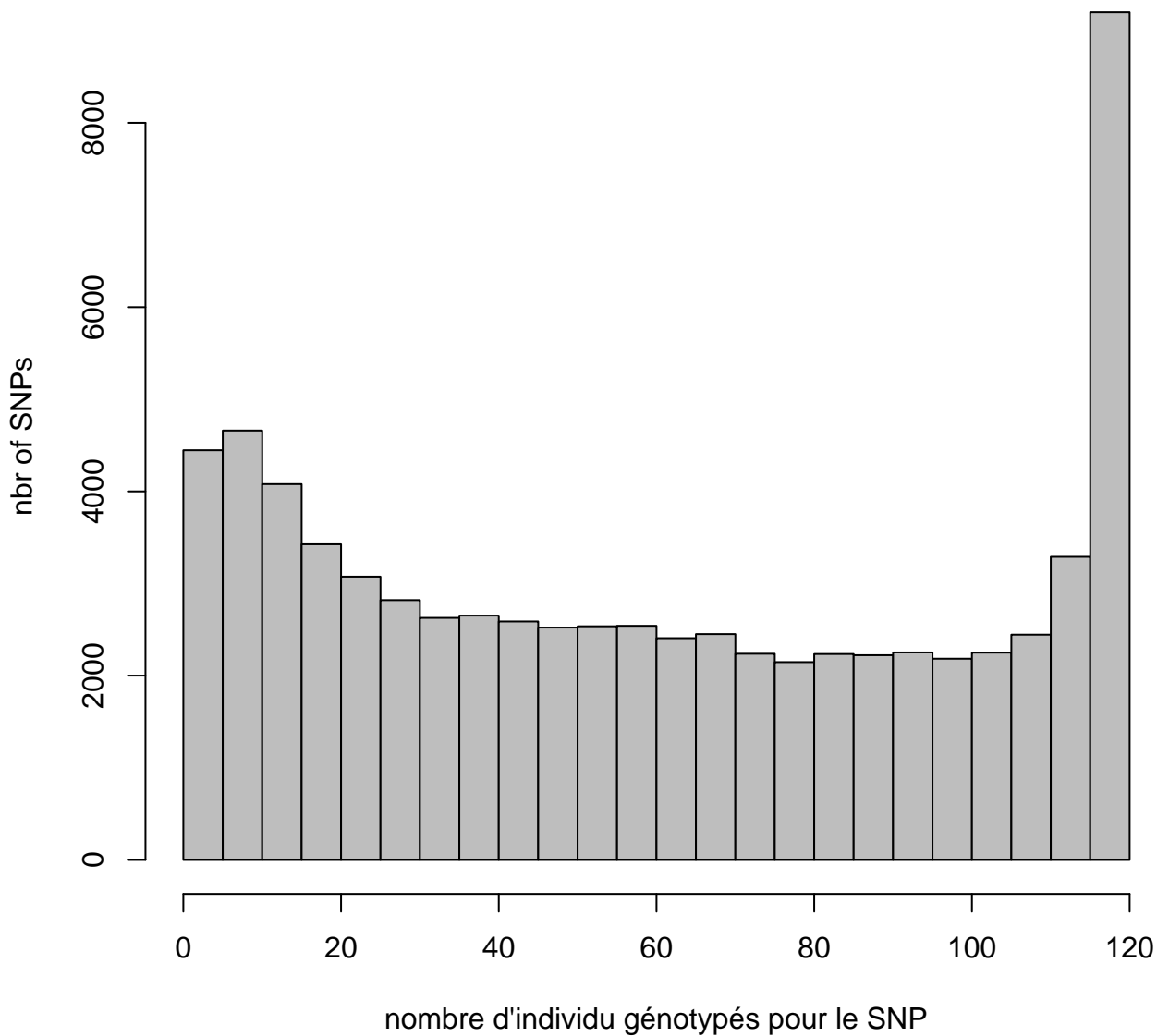




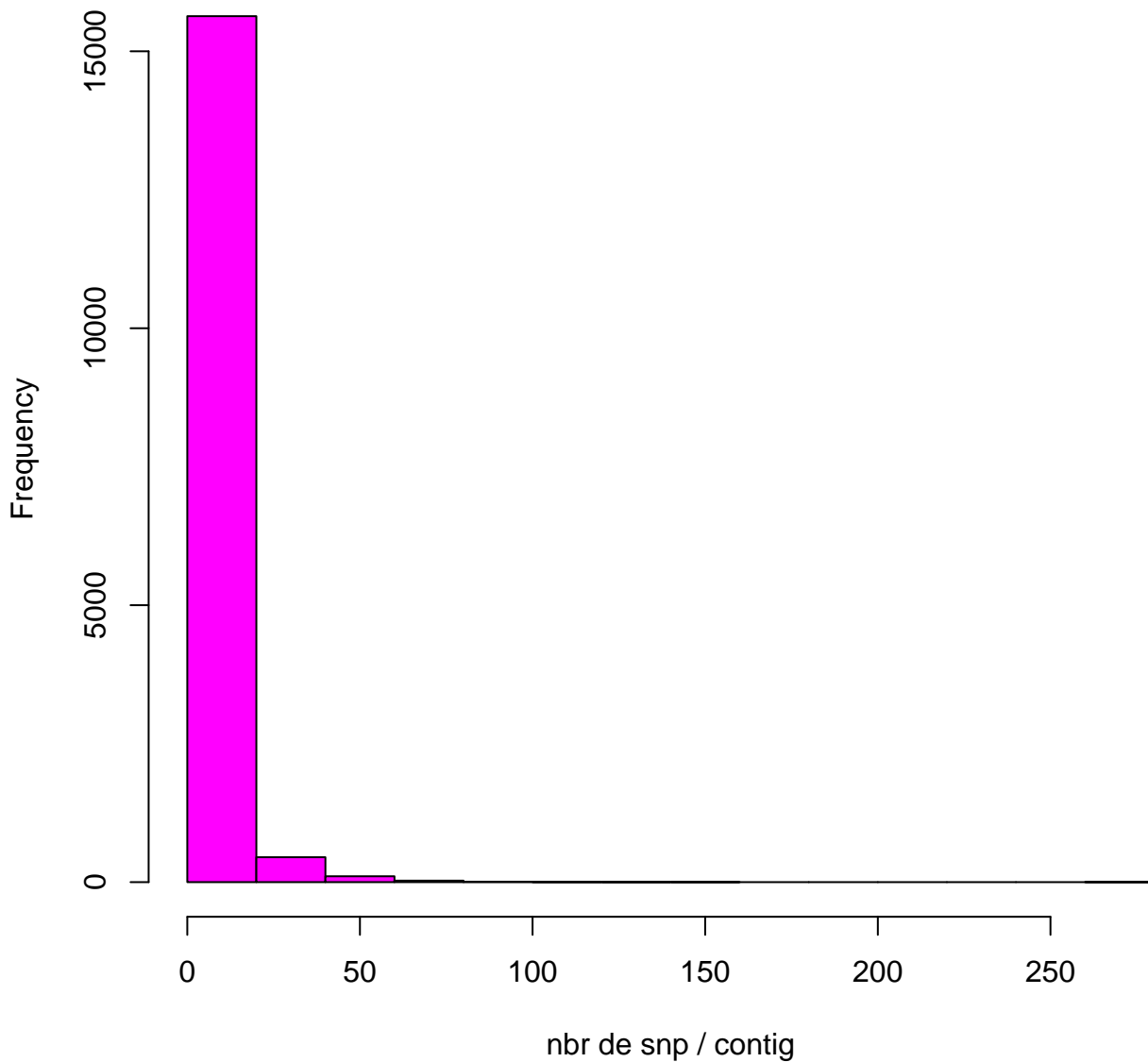
**nbr of SNP available with FIS > x**



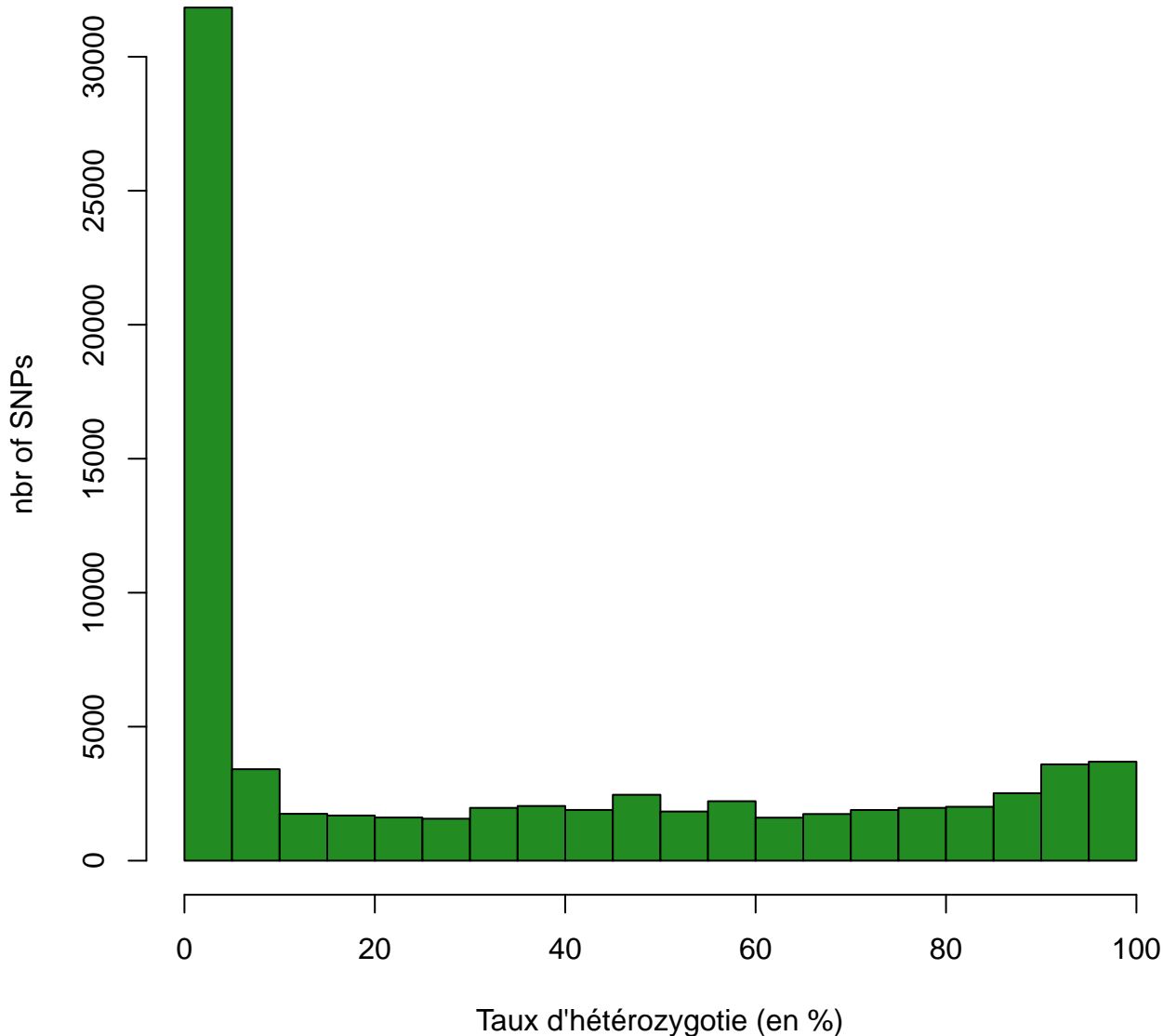
**nbr individu moyen par SNP : 60.5 | nbr of SNP : 73297**



**distribution du nbr de snp par contig**

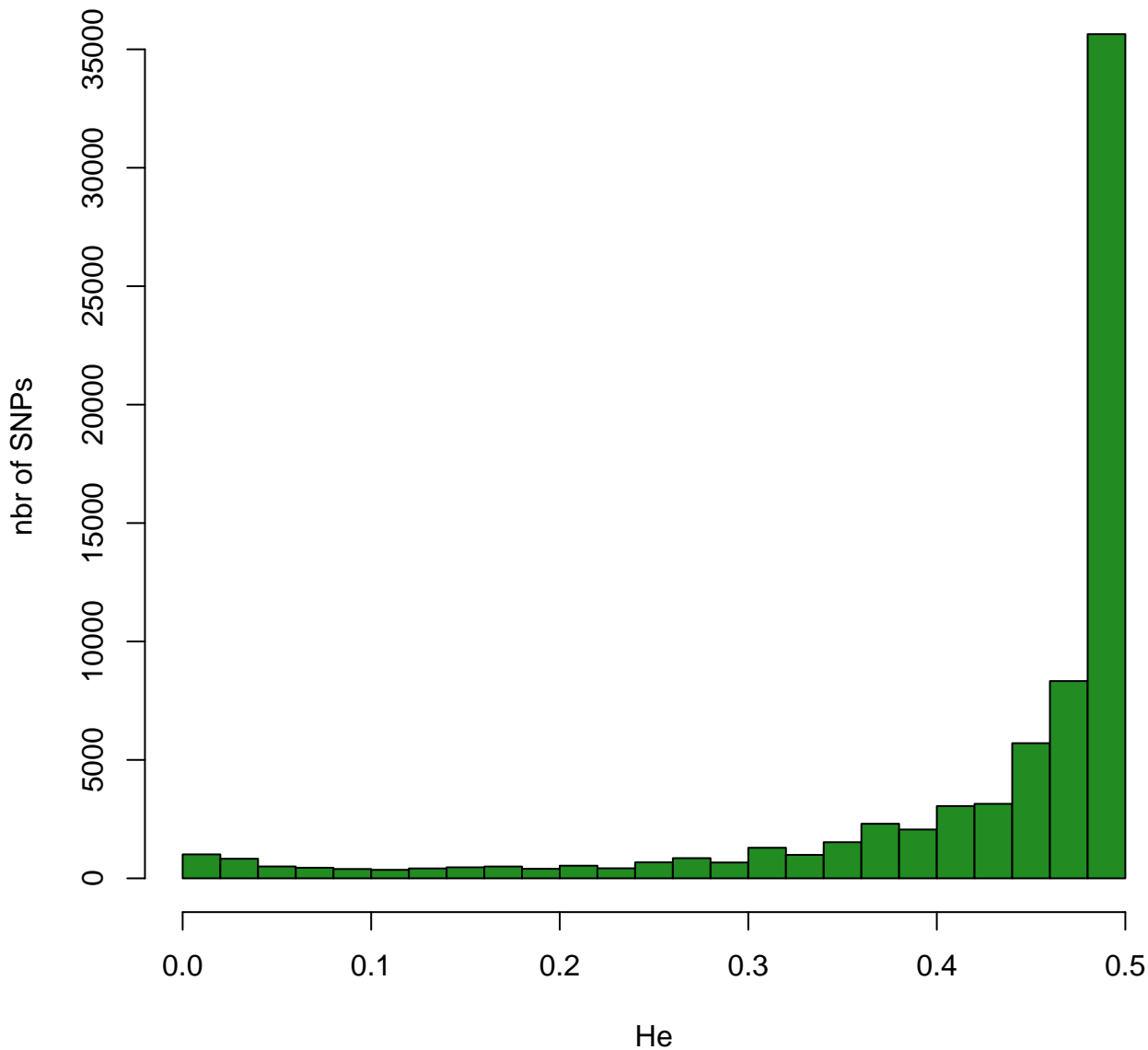


**Hétérozygotie moyenne : 32.11 % | nbr of SNP : 73297**

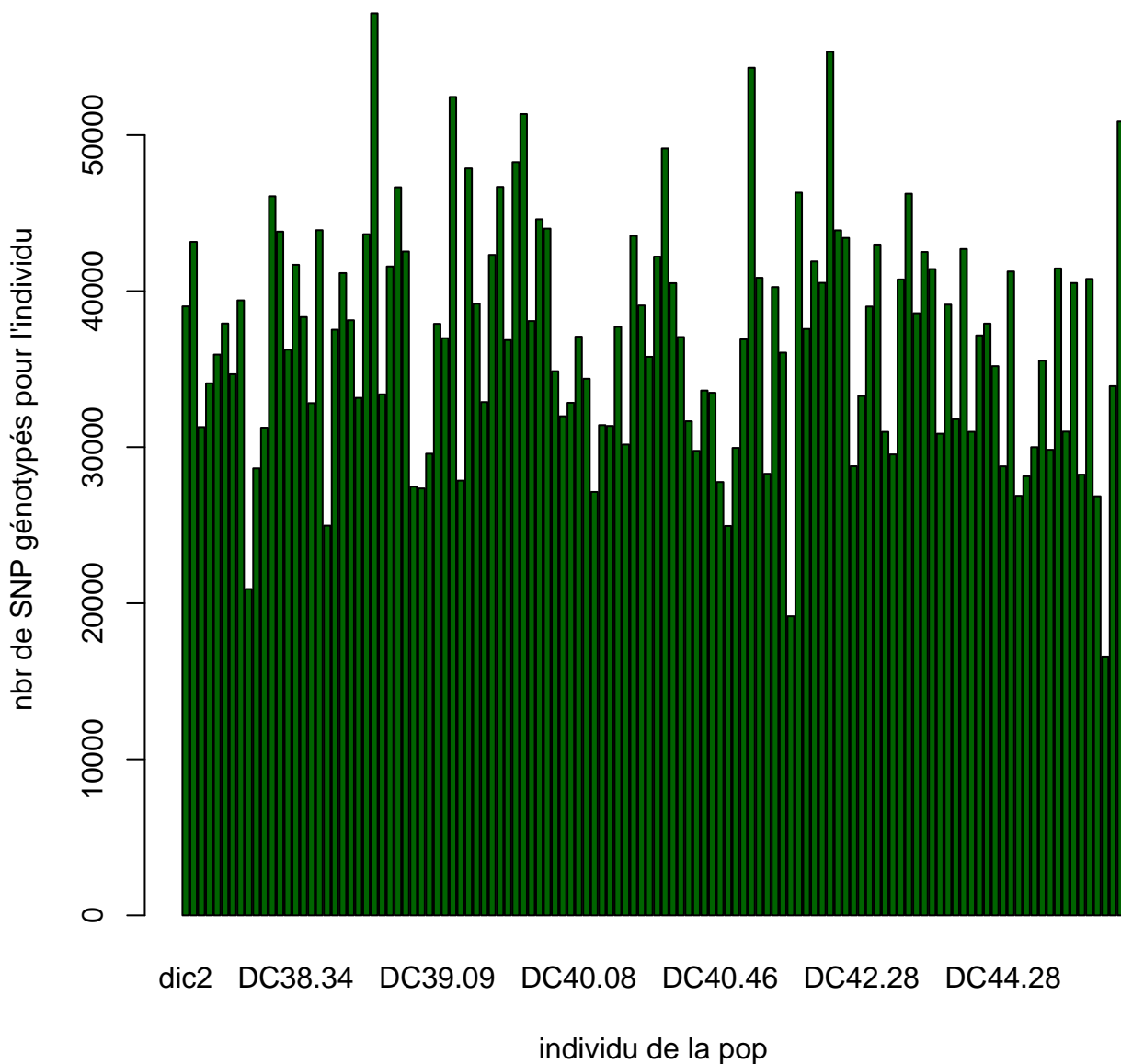




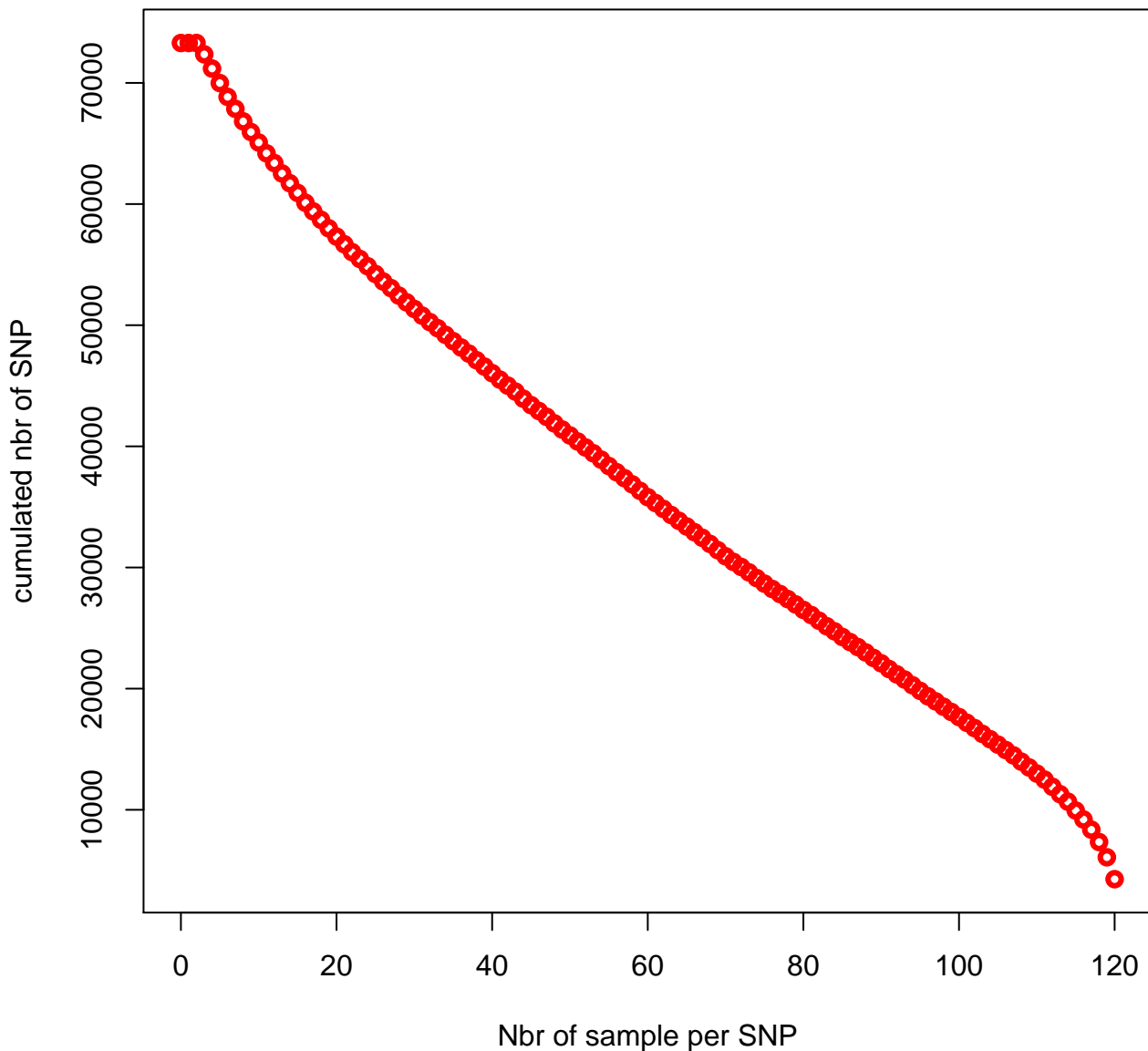
**Hétérozygotie attendue  $H_e$  : 0.43 | nbr of SNP : 73297**



**nbr de SNP genotypés par accession | moyenne = 36951.18**



**nbr of SNP available at least x individual per SNP**



% of heterozygotie for all genotypes | mean = 30.7881010727557

