

population size prediction error

1.0
0.8
0.6
0.4
0.2
0.0

- genotype LD, all SNP
- haplotype LD, all SNP
- genotype LD, MAF 5%
- haplotype LD, MAF 5%
- genotype LD, MAF 10%
- haplotype LD, MAF 10%
- genotype LD, MAF 20%

0 10 100 1,000 10,000 100,000

generations before present (log scale)

