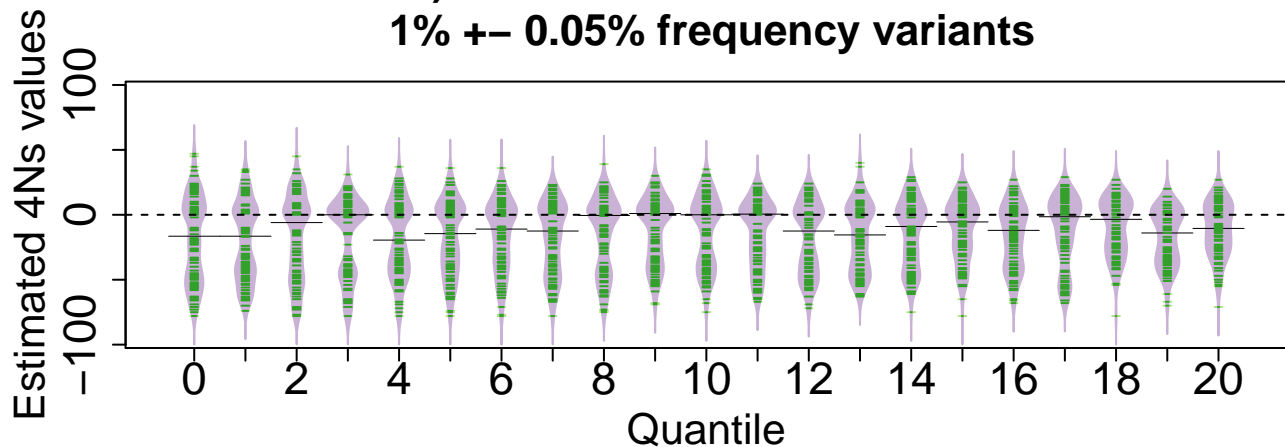
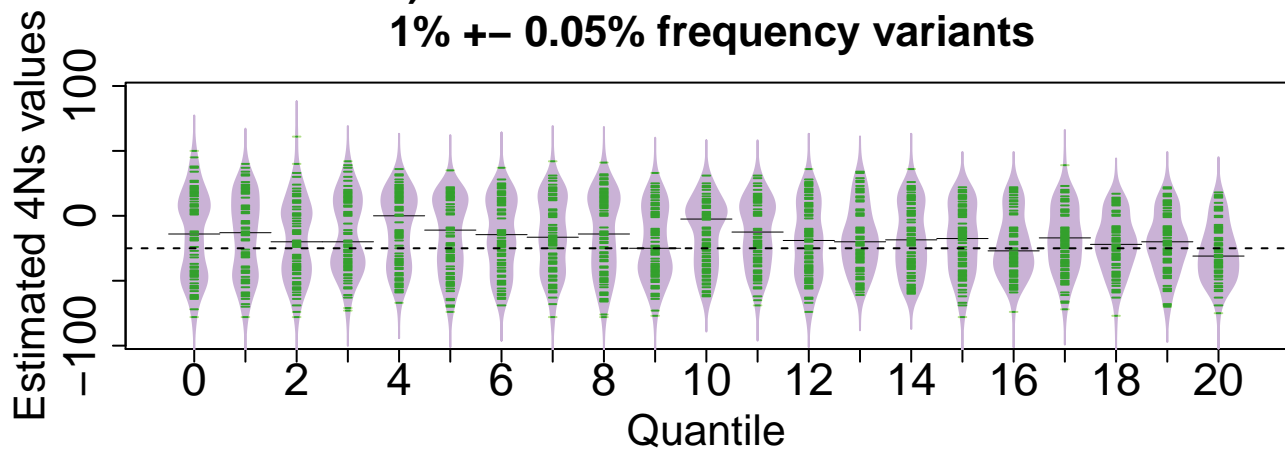


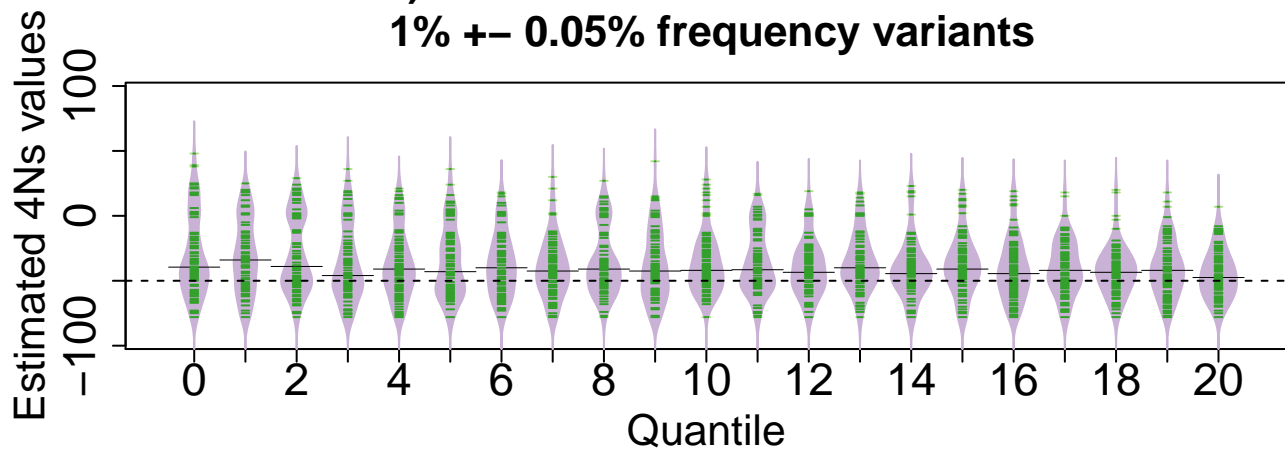
A) Selection Inference 4Ns = 0
1% \pm 0.05% frequency variants



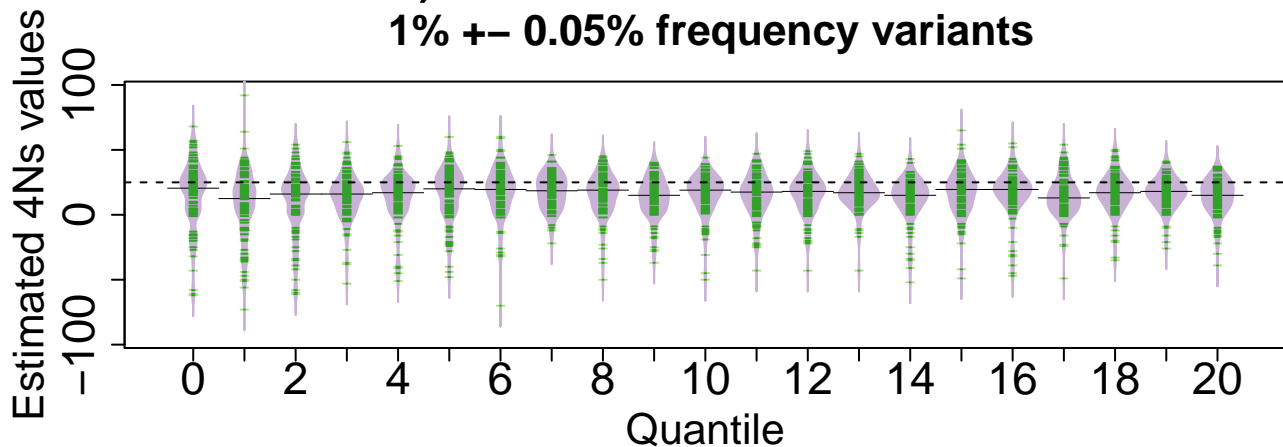
B) Selection Inference 4Ns = -25
1% \pm 0.05% frequency variants



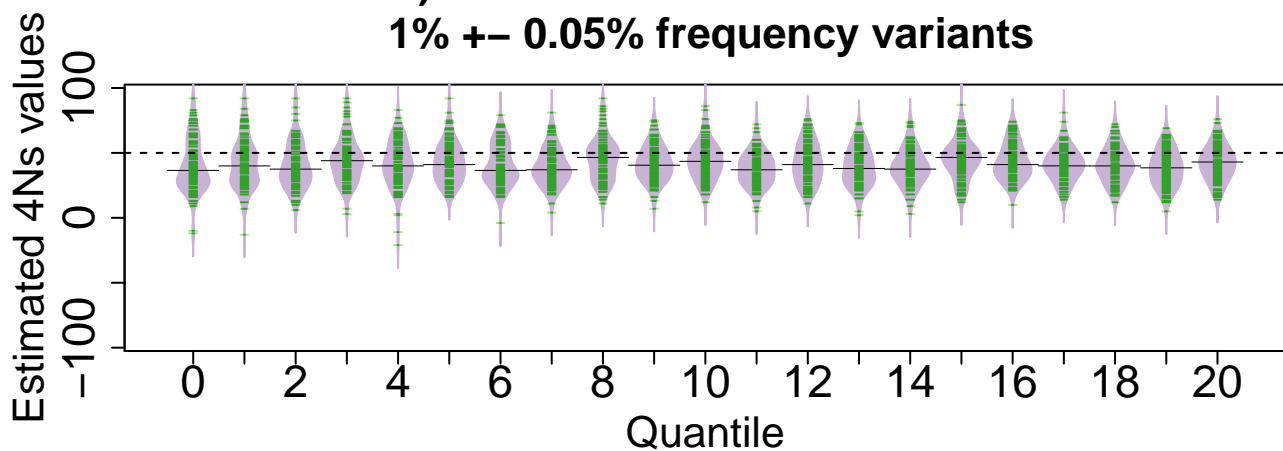
C) Selection Inference 4Ns = -50
1% \pm 0.05% frequency variants



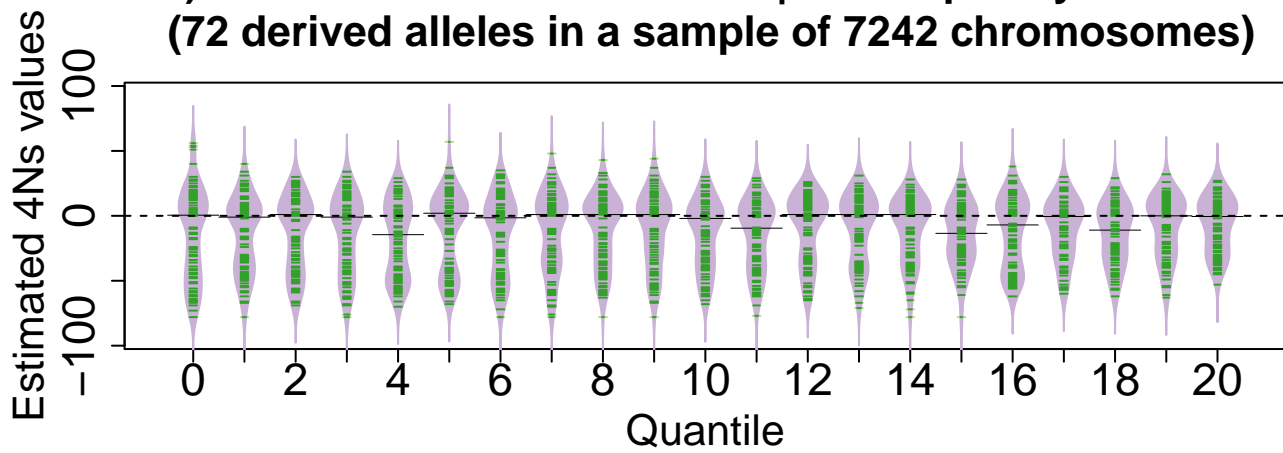
D) Selection Inference 4Ns = 25
1% \pm 0.05% frequency variants



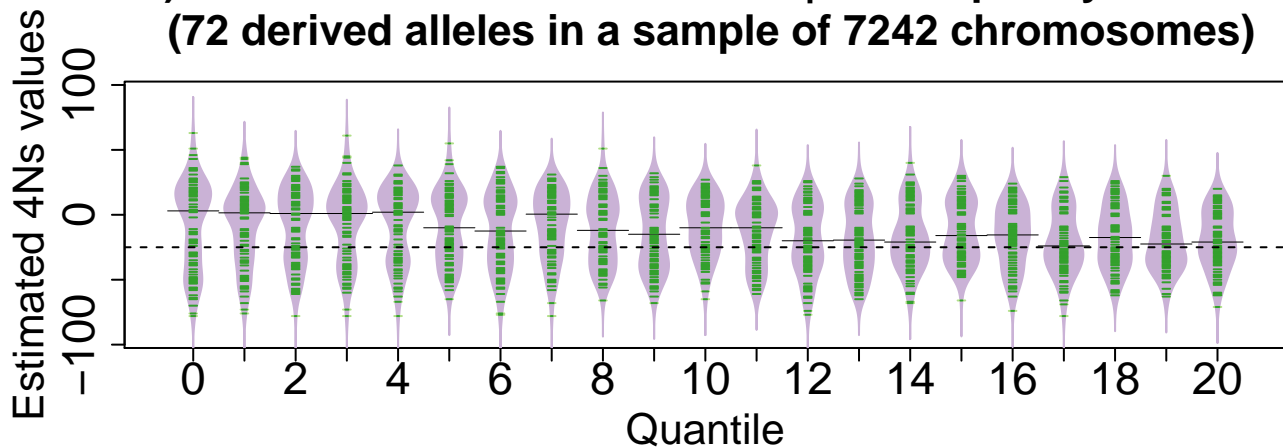
E) Selection Inference 4Ns = 50
1% \pm 0.05% frequency variants



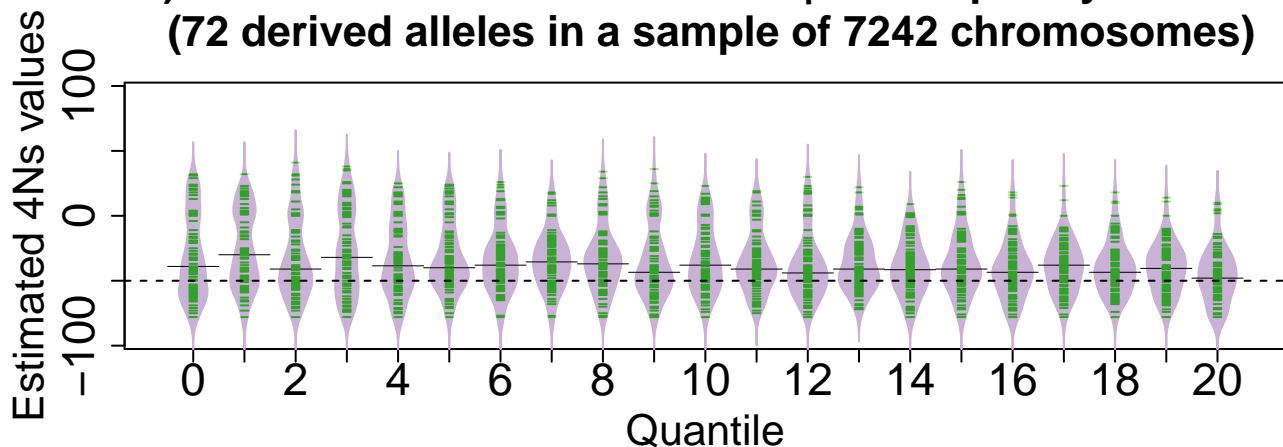
F) Selection Inference 4Ns = 0 | 1% frequency variants
(72 derived alleles in a sample of 7242 chromosomes)



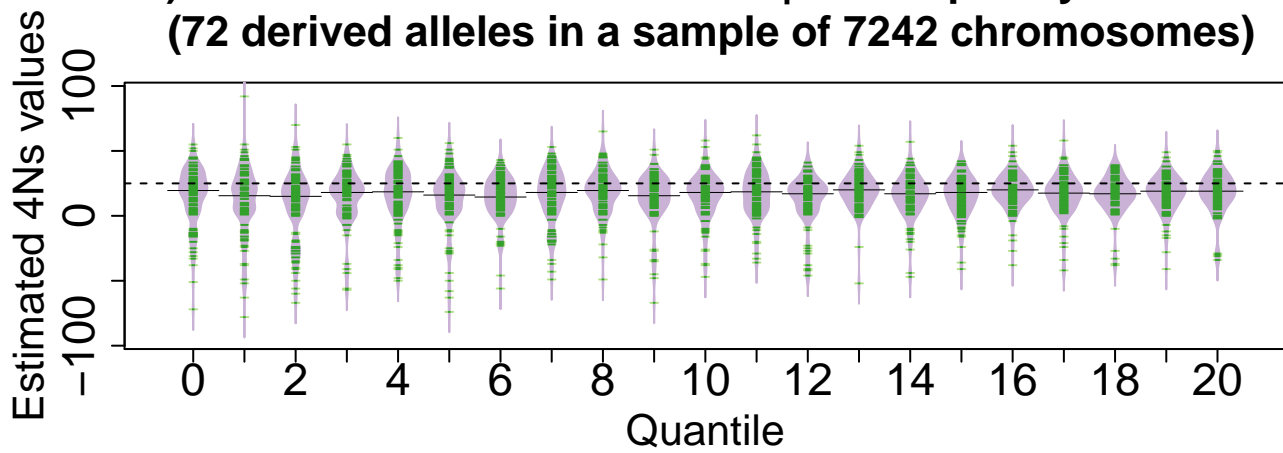
**G) Selection Inference $4N_s = -25$ | 1% frequency variants
(72 derived alleles in a sample of 7242 chromosomes)**



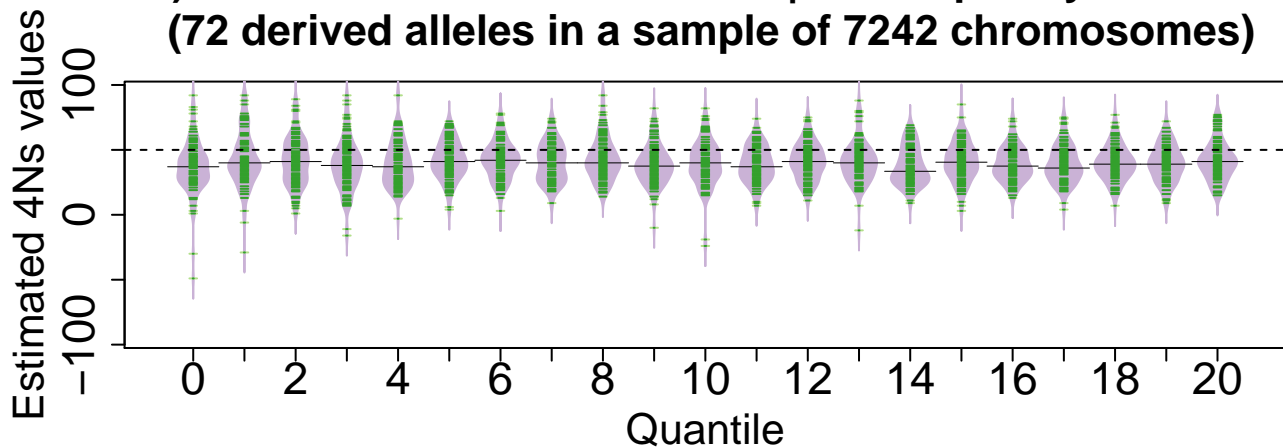
**H) Selection Inference $4N_s = -50$ | 1% frequency variants
(72 derived alleles in a sample of 7242 chromosomes)**



**I) Selection Inference $4N_s = 25$ | 1% frequency variants
(72 derived alleles in a sample of 7242 chromosomes)**



**J) Selection Inference 4Ns = 50 | 1% frequency variants
(72 derived alleles in a sample of 7242 chromosomes)**



K) RMSE

