

Additional file 17

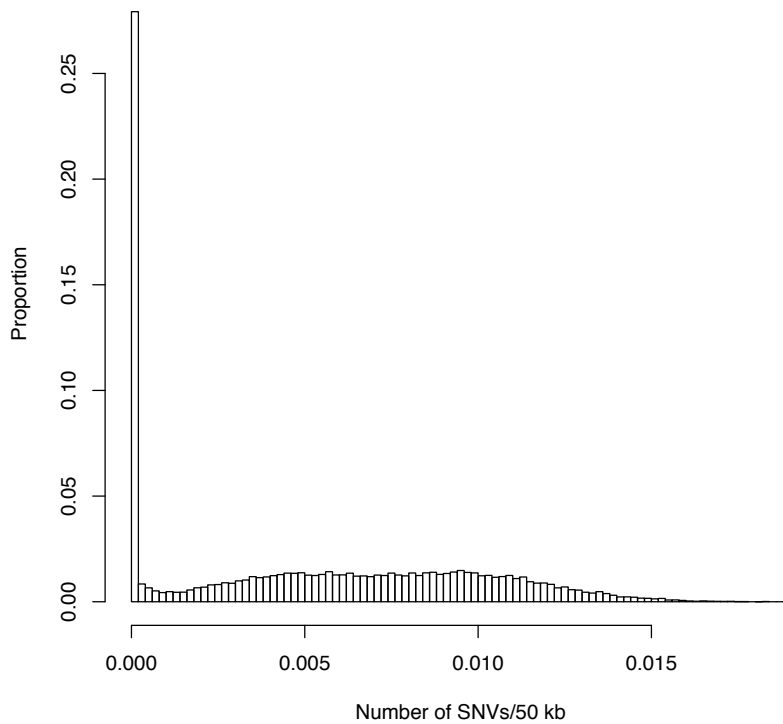


Figure S14. SNV density distribution

A histogram shows the number of SNVs in non-overlapped 50-kb windows across the *P. picta* genome. The distribution indicates that wild individuals of this species have a relatively high heterozygosity (~0.8 % on average), though the heterozygosity is being rapidly lost owing to inbreeding.