



Figure 3: The genotypic clusters of *Hyposoter horticola* based on (from left to right) clustering of one whole population of individuals analysis ($k=16$, $p=0.982$), clustering of five groups of individuals analysis ($k=4$, $p=0.999$), and clustering of 12 spatio-temporal groups of individuals analysis ($k=4$, $p=0.999$) with admixture results. The last two columns present the proportion of individual from each spatio-temporal group that carry the C-mitotype (in grey), or that are infected with *Wolbachia* (in dark grey). Data includes female specimens only ($N=195$).