

**Table S6:** Number of individuals per population assigned to a cluster using *ADEGENET* package<sup>1</sup> performed in R 4.0.2

Population	Cluster 1	Cluster 2	Cluster 3
BNNC	-	1	20
BHMA	-	2	25
CHUH	1	28	1
CHPR	2	27	-
CMRA	2	26	-
YORD	1	29	-
BGMA	23	3	6
LGXM	16	2	6
MADA	20	1	8
TPHU	21	4	6

<sup>1</sup> Jombart T, Devillard S, Balloux F (2010) Discriminant analysis of principal components: a new method for the analysis of genetically structured populations. BMC Genetics 11: 94. <https://doi.org/10.1186/1471-2156-11-94>