

	1.563	1.563	1.563	1.563	1.563	1.563	1.563	1.563	1.563	1.563	1.563	1.563	1.563
ThetaN mode (posterior HDI95)	1.14782 (0.35206- 1.5604)	1.35541 (0.66485- 1.55608)	1.14782 (0.31379- 1.55608)	1.35541 (0.36483- 1.55608)	1,07862 (0.29670- 1.54623)	1,12014 (0.29670- 1.54713)	0,995586 (0.29670- 1.54731)	1,47996 (0.40623- 1.55608)	1,46612 (0.33560- 1.55608)	0.303617 (0.19982- 1.45418)	0.954067 (0.27262 7-1.5284)	1.47996 (0.37571- 1.55608)	1.30005 (0.36165- 1.55608)

all times are given in numbers of years/generations,

* : this model needed to add a time for migration to stop - prior was 20-130 000 and posterior with a mode = 28888.9 (HDI95=21667-12311),

migration matrice 0 allows no migration; **migration matrice 1** allows migration between Alps-Carpathians and Carpathians-North;

migration matrice 2 allows migration between Alps-Carpathians, Carpathians-North and North-Alps; **migration matrice 3** allows migration between Alps-Carpathians and North-Alps; **migration matrice 4** allows migration between Alps-Carpathians; **migration matrice 5** allows migration between Alps-North and Carpathians-North

TDecN: Time of northern population size decrease, **TDecCarp**: Time of Carpathian population size decrease, **TExpN**: time of northern population growth, **TExpCarp**: time of Carpathian population growth,

TDivOE: Divergence time of population from the Alps and the Carpathians,

TDivNS: Divergence time of population from the North and the Carpathians,

Tmigr : timing of gene flow following the migration matrix given for each scenario,

gmigr: intensity of gene flow when existing,

ThetaA, ThetaC, ThetaN: nucleotide diversity given for the COI marker in the Alpine, the Carpathian and the northern populations, respectively.