

Supplementary Table S3. Details per individual of genome skimming and individual clustering analysis of specimens analysed from Tête Grosse (TG) and outside this population (FR).

Individual code	GS	Chrom.	Sequencing	Individual clustering	
	1C (Mb)	number (2n)	run number	No. reads	Coverage (× 1C)
TG84	8,097.84	80	SAMN25529504	1,200,000	0.022 ×
TG137	8,097.84	80	SAMN25529505	1,200,000	0.022 ×
TG237	8,097.84	80	SAMN25529506	1,200,000	0.022 ×
FR475	8,097.84	80	SAMN25529513	1,200,000	0.022 ×
TG337R	6,357	60	SAMN25529507	1,200,000	0.028 ×
TG506	6,357	60	SAMN25529511	1,200,000	0.028 ×
TG507	6,357	60	SAMN25529512	1,200,000	0.028 ×
TG370	4,205.4	40	SAMN25529508	1,200,000	0.043 ×
TG422	4,205.4	40	SAMN25529509	1,200,000	0.043 ×
TG444	4,205.4	40	SAMN25529510	1,200,000	0.043 ×
FR626	4,205.4	40	SAMN25529514	1,200,000	0.043 ×
FR627	4,205.4	40	SAMN25529515	1,200,000	0.043 ×