

Appendix 1.

Table A1. Population genetic parameters obtained after randomly subsampling Conguillío individuals to match the sample size for Nahuelbuta (N=14). We obtain similar results to those reported in Table 1. Note that although we did not change the number of samples from Nahuelbuta, the values of some parameters changed slightly because the missing data filter is now applied to a different total number of individuals (and we only exported loci present in at least 80% of individuals).

Parameters	Population sizes			Divergence time	Gene flow	
	Con	Nah	root		Con ->Nah	Nah -> Con
Median	14025	0	144792.5	0	1.11 E-09	1.17 E-05
L95% CI	14.5	0	133565	0	0	0
U95% CI	101442.5	1.25	156235	0	2.96	0.32