

Lait, L. A., Friesen, V. L., Gaston, A. J. and Burg, T. M. 2012. The post-Pleistocene population genetic structure of a western North American passerine: the chestnut-backed chickadee *Poecile rufescens*. – J. Avian Biol. 43: xxx–xxx.

Supplementary material

Appendix 1

Table A1. Environmental variables used to estimate suitable habitat in the maximum entropy modelling, as described in Carstens et al. (2007). The percent contribution towards the predicted distribution is given.

Temperature variable	%	Precipitation variable	%
Mean temp of wettest quarter	27.1	Precip of coldest quarter	41.2
Mean temp of driest quarter	9.8	Precip seasonality	3.1
Annual mean temp	4.7	Precip of warmest quarter	0.4
Mean temp of warmest quarter	3.9	Annual precipitation	0.2
Mean temp of coldest quarter	3.3	Precip of driest month	0.1
Temp seasonality	3.3	Precip of driest quarter	0.0
Isothermality	1.2	Precip of wettest month	0.0
Max temp of warmest month	0.8	Precip of wettest quarter	0.0
Min temp of coldest month	0.6		
Annual temp range	0.1		
Mean diurnal temp range	0.1		

Table A2. Variable sites table showing the differences between the 16 shared (A to P) and 38 unique haplotypes. Sites 68 to 523 are found in the control region fragment and sites 672 to 1437 are in the ATP fragment of the mtDNA. ‘.’ represents nucleotide identity to haplotype A. Population abbreviations are as in Fig. 1.

	CR			ATP		
	1111111	2222233334	4445	666777	8889001111	11111111
	6791144555	2223312452	2382	788457	5572280112	11222444
	8192337567	2561468661	5753	234237	7812435362	36178003
	CAACAACACA	ACCTCTCACT	TCCT	CGGGTA	CGCTTATCTG	76619047
A
BG.....
CG.....
DGG.....
EGG.....C.
FG.....C
GGG.....	..T.....
HGG.....	..C.....
IGG.....T.
JG.....T.
KGG.....T.C.
LG..G..T.
MGG.....T.....	..T.
NG.....T.....
O	T...G.....T.
P	.G..GG.....A... ..T..
AK102GG.....T.	...T..
AK104G.....T...
AK106GG.....T.	C.....
AK108G.....C	TA..CG
AK110GG.....	.T.....	..T.
AK122GG.....T.C.....
AK123GG.....TT.
TER105	T...G.....T.....	..T.
TER116G....GT.

TER119	T...G.....T.	C...
HAZ124	...GG....	C.T.
HAZ128	...G.....	G..C.....
HAZ129	..G.G....T.
HAZ130GT...T.
VI101	...GG....	T.....
VI102	...GG....T.	C...
VI104	.G...G..T.T.
VI107	...GG....	CT...
VI112	...GG....	..T.....T..
VI114	.G..GG....T.
INT104	...GG....T.G...
INT105	...GG....	C...	..A...	C
INT111	...GG....A...	C
INT112	...G.....T.	..A...	C
INT114	..TGG....T.	..A...	C
INT124	..TGG....	...c.....	..T.	..A...	..A.....	C	..A.....
INT125	...GG....	...C....	...C	..A...	C
NWA105	...G.....	T. ..T.
NWA113	...GG....	...C....	...CG...
NWA114	...GG....	...C....	..T.
NWA116	...GG....	G.. ..T.T..
CWA001	...GG....	T. ..T.
CWA005	...GG....A...T..
CWA128	...GG....	...C....T.....
CWA129	...GG....T.C
WOR001	...GG....	...C..G..A...T..
WOR003	...G.....C..
COR143	...GG..T.T.T.C....

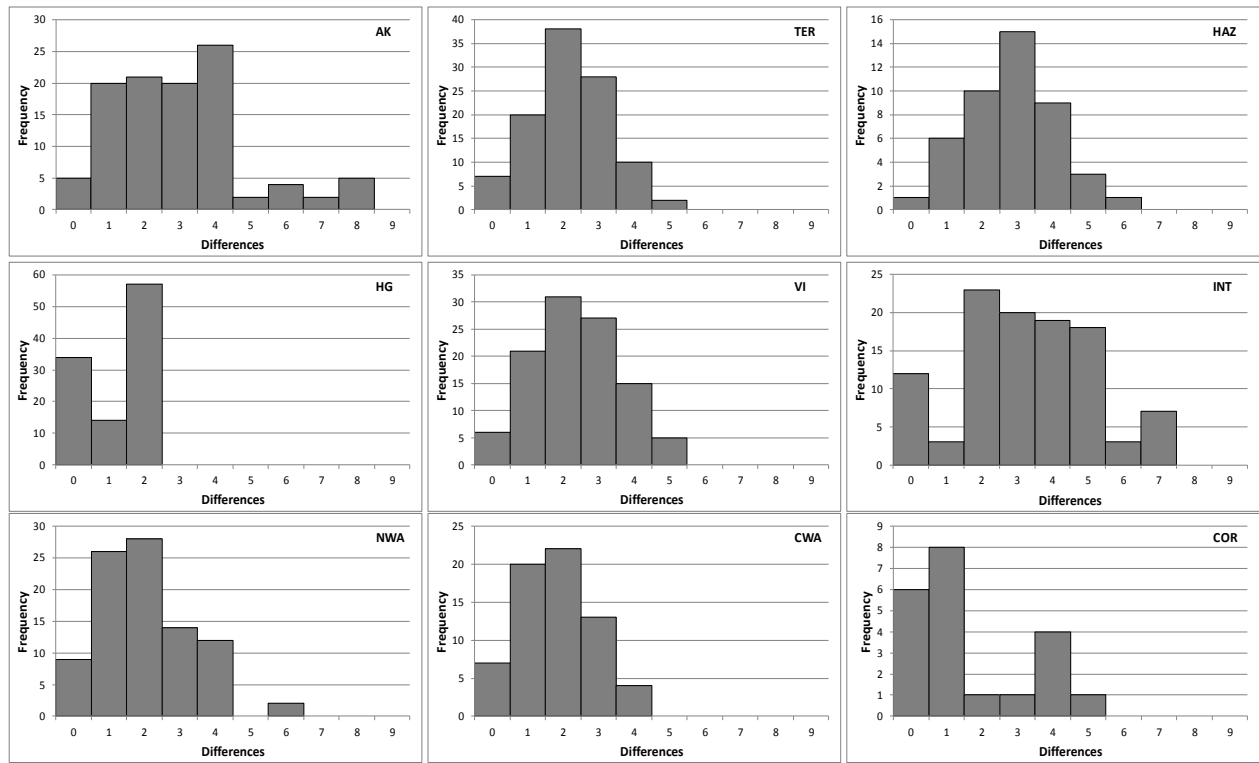


Figure A1. Observed mismatch distributions for nine chickadee populations calculated from pairwise differences.