

SUPPLEMENTARY MATERIAL 1. Genetic diversity estimates for the *Anaxyrus boreas* populations based on microsatellite genotypes. The Dixie Valley toad (DVT) population is highlighted in bold. A = number of alleles, A_{eff} = number of alleles weighted for their frequencies in the populations, H_e = expected heterozygosity.

Population	A	A_{eff}	H_e
Dixie	3.73	2.87	0.60
McCoy	2.27	1.71	0.33
CO1	1.82	1.35	0.18
CO2	3.36	1.97	0.38
CO3	2.18	1.72	0.34
CO4	2.82	2.16	0.36
CO5	2.82	2.30	0.48
CO6	2.27	1.66	0.33
CO7	2.91	2.07	0.40
CO8	2.09	1.78	0.35
CO9	1.64	1.26	0.16
O10 + 11 + WY1 + 2	3.55	1.91	0.40
UT1 + UT11	2.73	1.88	0.37
UT2	1.82	1.45	0.24
UT3	1.91	1.57	0.27
UT4	1.91	1.67	0.29
UT5	2.73	2.00	0.44
UT6	3.73	2.62	0.56
UT7	2.46	1.87	0.38
UT8	4.00	2.64	0.53
UT9	3.55	2.35	0.41
UT10	3.73	2.45	0.53
ID1	3.73	2.45	0.50
ID2	4.73	3.27	0.54
WY3	6.18	3.59	0.57
MT1	10.82	5.13	0.76
ID3	7.73	4.19	0.73
ID4	5.00	3.46	0.67
ID5	5.46	3.02	0.61
ID6	3.55	2.55	0.56
OR1	6.64	4.03	0.66
OR2	7.46	4.26	0.69
OR3	6.55	3.67	0.67
WA1 + 2	9.27	6.22	0.83
WA3 + 4	10.91	7.01	0.81
CA1	7.55	4.41	0.69
CA2	5.82	3.61	0.70
exsul	2.82	2.20	0.46
nelsoni	3.09	2.35	0.48
canorus1 + 2	5.73	3.77	0.63