

SUPPLEMENTARY MATERIAL 3. Numbers of alleles and estimates of observed heterozygosity (H_o), expected heterozygosity (H_e), and inbreeding coefficients (F_{IS}) for 217 camas pocket gophers (*Thomomys bulbivorus*) genotyped at 6 variable microsatellite loci, treated here as a single population. The large inbreeding coefficients and striking differences between observed and expected heterozygosity clearly indicate that significant genetic structure occurs within this data set.

Locus	Number of alleles	H_o	H_e	F_{IS}
Tm1	9	0.29	0.65	0.55
Tm2	4	0.38	0.43	0.12
Tm4	6	0.44	0.60	0.26
Tm5	20	0.41	0.91	0.56
Tm6	7	0.26	0.78	0.67
Tm7	9	0.52	0.66	0.21