Gevaert_SupMat_TableS1 – High genetic diversity in Porter's sunflower (*Helianthus porteri*) – JHered.

Supplementary Table S1. Primer information for the 18 EST-SSR markers used in this study and first published in Pashley et al. (2006; J Hered 97: 381-388). The column titled Locus ID refers to the nomenclature used in this study and previously referenced in Ellis et al. (2006; Mol. Ecol. 15: 2345-2355). The M13 tail used for labeling of all forward primers was CACGACGTTGTAAAACGAC, and is not included in the Forward Primer column. The expected size of each locus is based on the EST sequences originally developed for *Helianthus annuus*, and with the exception of BL0003, BL0010, BL0011 and BL0018, correspond similarly in *H. porteri*. Additional information on these markers is available at http://www.theburkelab.org/datafiles/JHered2006b_Primers.txt.

Locus ID	Forward Primer	Reverse Primer	Repeat Motif	Expected Size
BL0001	CTGCGTCGGCCTCGATCC	GACTTCAACGCCGTATCCAT	(aac)4	292
BL0002	GGCTTTCGTTTCTCGTTGTC	CAGCTCACTCCTAATTGGTTCC	(taat)5	302
BL0003	GCAGTGCTTCCACAGCAAC	AACCATTTCAACTCACTAAATAATCTC	(atg)4	140
BL0004	ATCCTCGGCTCAGCTAAACA	ATGATGGAGCCACCTATGGA	(tcc)4	305
BL0007	CGCAACGGGTCATCTACAG	GGACCGAACCCATCTAACAG	(ata)4	146
BL0008	GTCATGAAACCACGGCCTAC	TCACCACAAACCTCTCATCTTG	(atg)4	436
BL0010	TCCGAGCTTCCAATCATACC	GGAATGGCAATGATGTCTCC	(caa)4	264
BL0011	TGTTTGCTCACCTTGCTGAC	TGACACACAAACACCACTTGC	(gcat)3	116
BL0012	TGCAGTACTCGGGTGATGAG	TGTGATCATACAACACAATGATGA	(atta)4	270
BL0014	GGGTTTGTACCAGGCACTTG	TTCATAGAAATGAGGACCAAAGG	(ggt)4	322
BL0017	TTGAACAAGTTCGCAAGCTG	CAATACTACATCATATAATCGACCAAC	(tga)4	174
BL0018	CATGGTGCATTTGGCTACC	TGCTCCTGAAAGAACCCAAG	(gca)4	212
BL0020	AACCATGTTGCTTCCCAATC	TTCGGGACCGAGAGTATTTG	(cca)4	293
BL0022	ACTTACCGTTGCATTTGGTG	GCTTATCCCTAGAACACGATTACAG	(taa)4	116
BL0023	TTGGGTCAACACAAGCAGTC	AAACTGGCATTGGAAAGGTG	(tta)4	118
BL0025	AGTGGTCGGACGGTAAACAC	AACACACACCAAATCAAATCAC	(tggt)3	208
BL0027	AATGTTTCACCGCCTACAGC	GCCACCGGCTATATCTCAAC	(tgg)5	295
BL0030	GGGTAATGCAAAGTACTAAGATGTG	GCATCATCCAACAAACTAGAAGG	(atgt)3	235