

Supplementary Information for

Variability and nativeness in the Mediterranean taxa: divergence and phylogeography of *Genista etnensis* (Fabaceae) inferred from nuclear and plastid data.

Olga De Castro¹, Gianluigi Bacchetta², Salvatore Brullo³, Emanuele Del Guacchio^{1*}, Emanuela Di Iorio¹, Carole Piazza⁴, Paolo Caputo¹

Supplementary File S1.

Table S1. Information and results on the molecular markers tested (+ = variable sequence; 0 = non-variable sequence; x = amplified with abnormal structure; - = low amplification efficiency) (Pr = plastid DNA; N = nuclear DNA).

Cod.	Marcatore	Ref.	Note	Cod.	Marcatore	Ref.	Note
Pr1	<i>trnQ^(UUG)-psbK</i> IGS	1	+	Pr23	<i>ycf4-ycf10</i> IGS	1	0
Pr2	<i>psbK-trnS^(GCU)</i> IGS	1	x	Pr24	<i>petA-psbJ</i> IGS	1	-
Pr3	<i>trnG^(UCC)-atpA</i> IGS	1	+	Pr25	<i>petL-psaJ</i> IGS	1	-
Pr4	<i>atpF</i> intron	1	0	Pr26	<i>psaJ-rpL20</i> IGS	1	x
Pr5	<i>atpF-atpH</i> IGS	1	0	Pr27	<i>rpL20-rpS12</i> IGS	1	-
Pr6	<i>atpH-atpI</i> IGS	1	-	Pr28	<i>clpP</i> intron 2	1	-
Pr7	<i>rpoC1</i> intron	1	x	Pr29	<i>clpP</i> intron 1	1	-
Pr8	<i>trnS^(UGA)-psbZ</i> IGS	1	+	Pr30	<i>clpP-psbB</i> IGS	1	0
Pr9	<i>psaA-ycf3</i> IGS	1	+	Pr31	<i>psbH-petB</i> IGS	1	-
Pr10	<i>ycf3</i> intron 2	1	+	Pr32	<i>petB-petD</i> IGS	1	-
Pr11	<i>ycf3</i> intron 1	1	+	Pr33	<i>rpS3-rpS19</i> IGS	1	0
Pr12	<i>ycf3-trnS^(GGA)</i> IGS	1	-	Pr34	<i>ccsA-ndhD</i> IGS	1	x
Pr13- 14	<i>rpS4-trnT^(UGU)_ver.1</i>	1	+	Pr35	<i>psaC-ndhE</i> IGS	1	0
Pr15	<i>trnF^(GAA)-ndhJ</i> IGS	1	+	Pr36	<i>ndhE-ndhI</i> IGS	1	x
Pr16	<i>ndhC-trnV^(UAC)</i> IGS	1	-	Pr37	<i>rpS15-ycf1</i> IGS	1	x
Pr17	<i>trnV^(UAC)</i> intron	1	0	Pr38	<i>psbA-trnH^(GUG)</i> IGS	2	+
Pr18	<i>trnV^(UAC)-atpE</i> IGS	1	+	Pr39	<i>trnQ^(UUG)-rps16</i> IGS	3	0
Pr19	<i>atpB-rbcL</i> IGS	1	0	Pr40	<i>trnL^(UAA)</i> intron	4	0
Pr20	<i>rbcL-accD</i> IGS	1	x	Pr41	<i>trnL^(UAA)-trnF^(GAA)</i> IGS	4	0
Pr21	<i>accD-psaI</i> IGS	1	-	Pr42	<i>trnG^(UCC)-trnS^(GCU)</i> IGS	5	+
Pr22	<i>psaI-ycf4</i> IGS	1	0	N43	ITS1-ITS2	6	+

References

- 1 Prince, L.M. Plastid Primers for Angiosperm Phylogenetics and Phylogeography. *Appl. Plant Sci.* **2015**, *3*, 1400085. <https://doi.org/10.3732/apps.1400085>.

- 2 Kress, W.J.; Erickson, D.L. A Two-Locus Global DNA Barcode for Land Plants: The Coding rbcL Gene Complements the Non-Coding *trnH-psbA* Spacer Region. *PLoS ONE* **2007**, *2*, e508. <https://doi.org/10.1371/journal.pone.000050>.
- 3 Calviño, C.I.; Downie, S.R. Circumscription and phylogeny of Apiaceae subfamily Saniculoideae based on chloroplast DNA sequences. *Mol. Phylogenet. Evol.* **2007**, *44*, 175-191. <https://doi.org/10.1016/j.ympev.2007.01.002>.
- 4 Taberlet, P.; Gielly, L.; Pautou, G.; Bouvet, J. Universal primers for amplification of three non-coding regions of chloroplast DNA. *Plant Mol. Biol.* **1991**, *17*, 1105-1109. <https://doi.org/10.1007/BF00037152>.
- 5 Hamilton, M.B. Four primer pairs for the amplification of chloroplast intergenic regions with intraspecific variation. *Mol. Ecol.* **1999**, *8*, 521-523.
- 6 De Castro, O.; Cozzolino, S.; Jury, S.L.; Caputo, P. Molecular relationships in *Genista* L. Sect. *Spartocarpus* Spach (Fabaceae). *Plant Syst. Evol.* **2002**, *231*, 91-108. <https://doi.org/10.1007/s006060200013>.