

Supplementary Material

Table S1. Geographic origin of 40 *Lathyrus* accessions.

Sample ID	Section	<i>Lathyrus</i> spp	Mating system	Origin
1				Syria
2				Ethiopia
3				Ethiopia
4				Ethiopia
5				Ethiopia
6				Ethiopia
7	Lathyrus	<i>Lathyrus sativus</i>	Mainly autogamous	Ethiopia
8				Italy
9				Ethiopia
10				Ethiopia
11				Ethiopia
12				Ethiopia
13				Ethiopia
14				Australia
15				Greece
16				Australia
17				Greece
18				Syria
19				Greece
20				Greece
21	Lathyrus	<i>Lathyrus cicera</i>	Mainly autogamous	Greece
22				Greece
23				Greece
24				Greece
25				Greece
26				Greece
27				Greece
28				Greece
29				Greece
30				Germany
31				Greece
32				Cyprus
33				Greece
34	Clymenum	<i>Lathyrus ochrus</i>	Autogamous	Cyprus
35				Greece
36				Greece
37				Greece
38				Cyprus
39				Greece
40				Cyprus

Table S2. Traits measured in the evaluation of the forty (40) *Lathyrus* accessions from seven diverse origins in 2013/14, 2014/15 and 2015/16 seasons

Traits	Definition	Unit
Days to flowering (DF)	From planting to 50% plants flowering	Days
Biomass production (Biomass)	Aboveground dry biomass weight per hectare	t ha ⁻¹
Grain yield (GY)	Grains weight per hectare at 13% humidity	t ha ⁻¹
seed weight (1000SW)	Weight of 1000 seeds randomly sampled from each plot	g
Harvest index	Ratio of grain yield and total dry above ground biomass	#
Protein content	Estimated according to Kjeldahl method (Tkachuk, 1969)	%

Table S3. Microsatellite primer pairs (SSR), motif and expected size range used for the genetic characterization of *Lathyrus* collection.

SSR	Repeat motif	Forward primer	Revers primer	Size range
G5	(AAC)10	CACAACCAGTTGCATCAGTG	TGGTCACATGATGGTTTGT	200-220 (Wand et al. 2015)
G26	(AC)16	CCACCAAATTTCCCTTTTTG	GGTACGAGAGGTTGACTTTTGTTT	160-197 (this study)
G67	(AC)9	CACCCTCTCACTGCCTAGC	TTGGGGGTTGTAGAAGGAAC	132-150 (this study)
G157	(CAA)6	ACATCCAATCCCCACCATAA	AATGCATGGTTGTTGCTTGA	189-234 (this study)
G245	(TG)6	CGTTGGTTGTTAGTCGGTCA	GAACGAAACAACGACGACAA	220-240 (Wang et al. 2015)
G17922	(CCA)5	CACCACCATAACCACCTCCT	ATGCGATTGAAGGGATGAAC	177-228 (this study)
G18078	(TGT)8	TTCAGATGCAGGTGGTTCAG	AACGGTGC GACTCTTGCTAT	108-154 (this study)
G18109	(CGA)5	GACAGACACACGGCAAACAC	ACGTCGTCGTGTCGTTGTT	170-200 (Wang et al. 2015)
G18549	(GTT)5	TGAGGGTGTGTTGAACGTGAG	CACCACAACAACAACAACCAC	140-170 (Wang et al. 2015)
G19207	(AAG)5	ATCGTAAACCGTGAGGGTCA	AAGCTTGTGGTGGCTACTGC	200-210 (Wang et al. 2015)

Table S4 Three-years means of agro-morphological traits in the accessions of three *Lathyrus* spp.

Specie	Acc.	Origin	1000SW	GY	Biomass	DF	Protein	HI
<i>L. sativus</i>	1	Syria	77.48ab	2.91ab	8.19a	96.22ab	30.96ab	0.36ab
	2	Ethiopia	76.45ab	3.11a	8.37a	96.55ab	30.36b	0.37a
	3	Ethiopia	76.75ab	2.81ab	8.16a	95b	31.17a	0.36ab
	4	Ethiopia	71.32b	3.06a	7.28b	99.22a	30.62ab	0.36ab
	5	Ethiopia	76.35ab	2.58b	6.94c	96.33ab	30.98ab	0.36ab
	6	Ethiopia	74.75ab	2.96a	7.21b	96.66ab	30.52b	0.39a
	7	Ethiopia	80.02ab	2.49b	7.21bc	95.55b	31.23a	0.35b
	8	Italy	72.17b	2.55b	7.19bc	97.88ab	31.08a	0.34b
	9	Ethiopia	76.54ab	2.53b	6.93c	94.33b	30.32b	0.34b
	10	Ethiopia	74.27ab	2.76ab	7.62ab	95b	30.19	0.35b
	11	Ethiopia	81.54a	2.58b	7.55ab	95.77b	30.1b	0.36ab
	12	Ethiopia	74.86ab	2.42b	8.3a	98a	30.52b	0.3c
	13	Ethiopia	72.24b	2.76ab	7.33b	99.11a	30.67ab	0.35b
<i>L. cicera</i>	1	Australia	70.7bc	3.7a	8.7a	99a	26.98b	0.43ab
	2	Greece	71.1bc	3.7a	8.63a	99.66a	28.84a	0.42b
	3	Australia	69.2bc	3.8a	8.85a	98.11ab	27.36b	0.42b
	4	Greece	68.9bc	3.2c	7.93b	98.22ab	28.93a	0.38c
	5	Syria	70.3bc	3.3b	7.59c	98.77a	27.69ab	0.4bc
	6	Greece	76.2ab	3.5ab	8.04b	97.88b	26.23c	0.45a
	7	Greece	73.0abc	3.4ab	8.08b	97.55b	27.08b	0.43ab
	8	Greece	64.2c	3.78a	8.38ab	99.11a	27.62ab	0.44a
	9	Greece	69.4bc	3.3b	8.14b	98.77a	26.64bc	0.45a
	10	Greece	70.7bc	3.36b	7.97b	100.11a	28.8a	0.38c
	11	Greece	71.7bc	3.2c	7.73bc	97.33bc	27.87ab	0.41b
	12	Greece	69.4bc	3.27bc	7.81bc	98.33ab	27.62ab	0.42b
	13	Greece	70.6bc	3.5ab	8.43ab	97.44b	27.33b	0.4bc
	14	Greece	82a	3.5ab	8.51ab	96.33c	27.47b	0.39bc
	15	Greece	75.5bc	3.29bc	7.99b	98.44ab	27.08b	0.41b
<i>L. ochrus</i>	1	Greece	107.72b	2.07b	6.19d	93.88c	29.32b	0.36ab
	2	Germany	101.4c	2.3ab	6.93b	97.77ab	30.36ab	0.32b
	3	Greece	115.89a	2.36ab	6.15d	93.77c	29.75b	0.36ab
	4	Cyprus	95.4cd	2.2b	7.33a	96.33b	29.63b	0.3bc
	5	Greece	106.58b	2.52a	7.46a	95.44bc	29.18b	0.35ab
	6	Cyprus	98.32c	2.09b	7.39a	99.66a	31.4a	0.29c
	7	Greece	106.13b	2.09b	7b	b	30.89ab	0.31b
	8	Greece	92.04d	2.44a	6.38c	92.88c	29.54b	0.38a
	9	Greece	102.07bc	2.24b	6.81b	96.22b	30.72ab	0.33b
	10	Cyprus	103.8bc	2.01c	6.58bc	95.22bc	30.72ab	0.29c
	11	Greece	110.62ab	2.37ab	7.43a	95.55bc	30.37ab	0.33b
	12	Cyprus	102.14bc	2.23b	7.37a	96b	30.57ab	0.3bc

1000SW, 1000 seed weight (g); GY, Grain Yield (t ha⁻¹); Biomass, Biomass production (t h⁻¹); Protein (%); DF, Day to Flowering (day); Protein, protein content (%); HI, Harvest Index.

Table S5. Pearson's correlation between agronomic traits for each *Lathyrus* spp.

		1000SW	GY	Biomass	DF	HI
<i>L. sativus</i>	1000SW					
	GY	-0.2967				
	Biomass	0.1513	0.3060			
	DF	-0.6752	0.2087	-0.0122		
	HI	0.0925	0.7015	-0.2162	-0.1506	
	Protein	-0.0383	-0.1369	-0.0640	0.0921	-0.0452
<i>L. cicera</i>	1000SW					
	GY	-0.1390				
	Biomass	0.0233	0.8722			
	DF	-0.5874	0.1191	0.0006		
	HI	-0.1503	0.4066	0.2096	0.0600	
	Protein	-0.2797	-0.1714	-0.1026	0.3855	-0.7341
<i>L. ochrus</i>	1000SW					
	GY	0.0741				
	Biomass	-0.2019	0.1293			
	DF	-0.2181	-0.3620	0.6572		
	HI	0.1907	0.5927	-0.5891	-0.7457	
	Protein	-0.0954	-0.5488	0.3154	0.7246	-0.7187

1000SW, 1000 seed weight (g); GY, Grain Yield (t ha⁻¹); Biomass, Biomass production (t h⁻¹); DF, Day to Flowering (day); Protein, protein content (%); HI, Harvest Index.

Table S6. Nei genetic distances.

	<i>S.S</i>	<i>S.I</i>	<i>S.E</i>	<i>C.A</i>	<i>C.S</i>	<i>C.G</i>	<i>O.GR</i>	<i>O.G</i>	<i>O.C</i>
<i>S.S</i>	0.000								
<i>S.I</i>	0.367	0.000							
<i>S.E</i>	0.098	0.167	0.000						
<i>C.A</i>	0.269	0.343	0.162	0.000					
<i>C.S</i>	0.267	0.367	0.145	0.124	0.000				
<i>C.G</i>	0.231	0.304	0.127	0.034	0.092	0.000			
<i>O.GR</i>	0.520	0.573	0.360	0.447	0.516	0.417	0.000		
<i>O.G</i>	0.274	0.307	0.147	0.258	0.286	0.218	0.172	0.000	
<i>O.C</i>	0.307	0.385	0.195	0.219	0.289	0.193	0.203	0.059	0.000

The first letters S, C and O indicates *L. sativus*, *L. cicera* and *L. ochrus*, respectively; while the second one S, I, E, A, G, GR and C indicates Syria, Italy, Ethiopia, Australia, Greece, Germany and Cyprus, respectively.

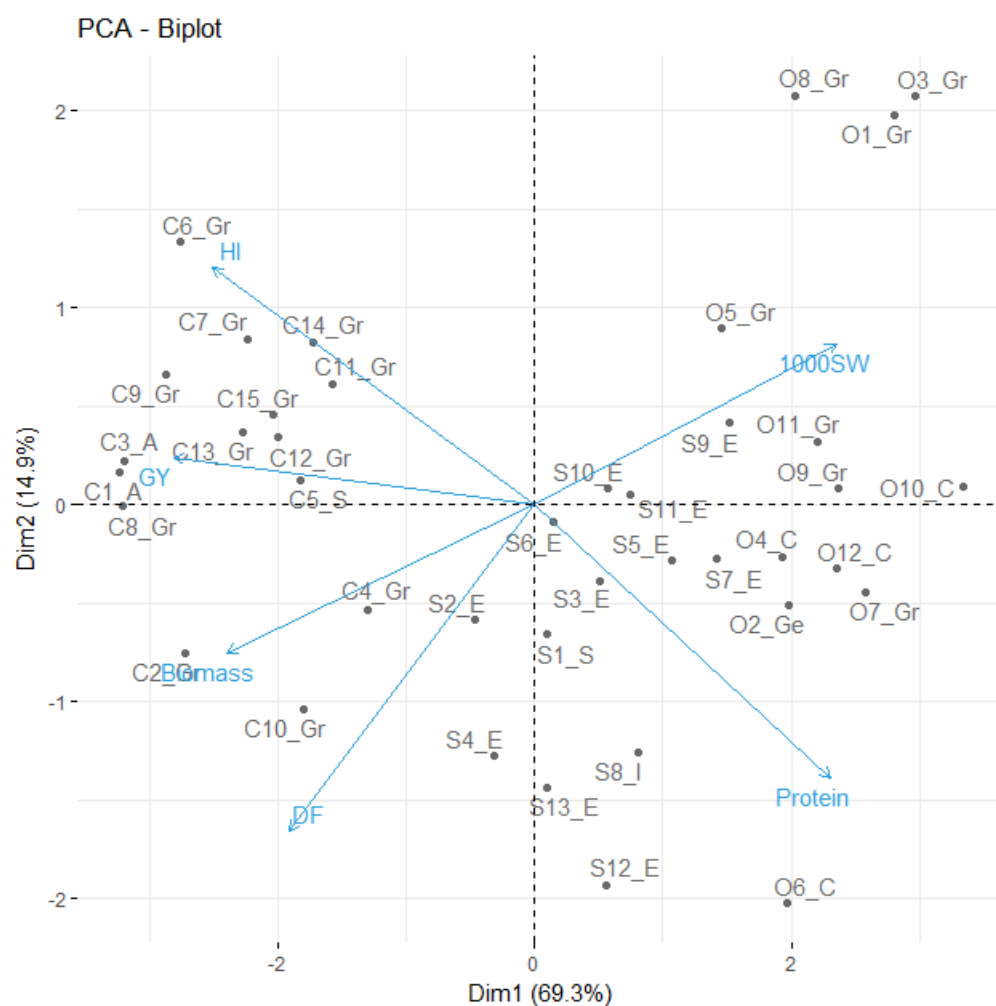


Figure S1. Principal component analysis of forty *Lathyrus* spp. accessions (C, S and O indicates *L. cicera*, *L. sativus* and *L. ochrus*, respectively; Gr, A, S, E, I, C and Ge indicates Greece, Australia, Syria, Ethiopia, Italy, Cyprus and Germany, respectively). 1000SW, 1000 seed weight (g); GY, Grain Yield (t ha⁻¹); Biomass, Biomass production (t h⁻¹); DF, Day to Flowering (day); Protein, protein content (%); HI, Harvest Index.

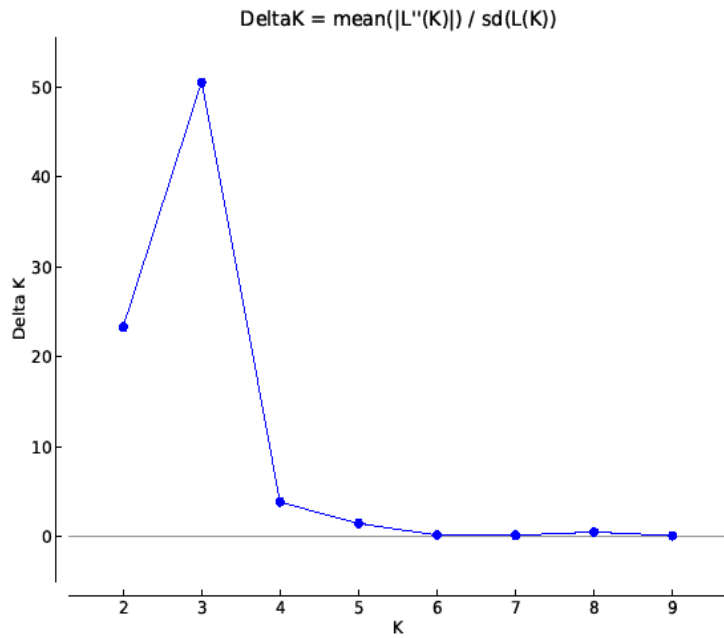


Figure S2 Analysis of the population structure of the 40 *Lathyrus* accessions: ΔK value related to different K.

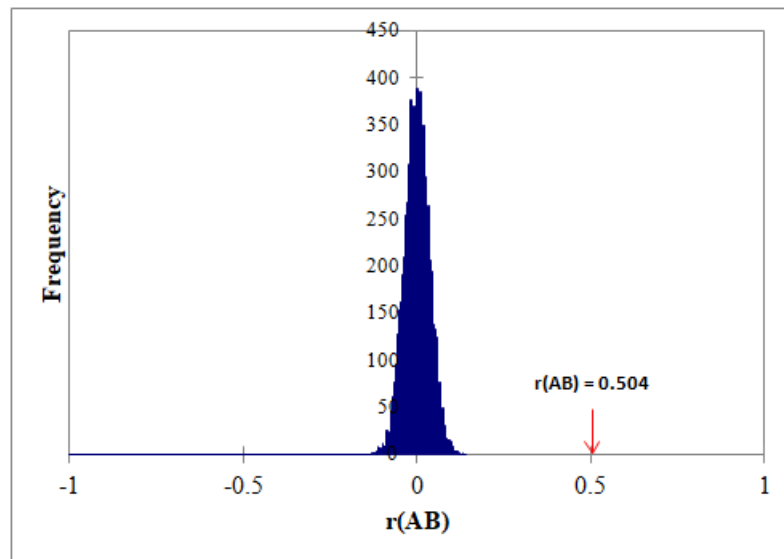


Figure S3. Mantel statistical values for the null hypothesis for the correlation between genetic and agronomic distance matrices. The red arrow shows the position of the statistical value ($r(AB) = 0.504$, in the zone of rejection of the null hypothesis; $p < 0.05$). The data in the frequency distribution curve were generated from 9999 random permutation.