

Erratum to: Genetic diversity of *Patellifolia patellaris* from the Iberian Peninsula, a crop wild relative of cultivated beets

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find in this erratum the correct version of Table 3 that should be regarded as final version by the reader.

Due to an unfortunate turn of events during article processing data in Table 3 was given incorrect. Please

The online version of the original article can be found under doi:[10.1007/s10681-017-1942-0](https://doi.org/10.1007/s10681-017-1942-0).

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Table 3 Pairwise genetic distance values Δ of ten occurrences as well as the complementary compositional differentiation values Δ_{SD} considering elementary genic differences at three levels: (a) gene pool level, (b) mean single locus level and (c) multilocus level

Level of association	Occurrence identifier	Pairwise genetic distance Δ between occurrences										Components of the communities' differentiation					
		AZO	BAL	CGO	CNE	COL	FM	MOR	PCA	PS	STE	Δ_{SD}	Δ_{SD}	Δ_{SD}	Δ_{SD}		
(a) Gene pool (GP)	AZO	0													Δ_{AZO}	0.3772	
	BAL	0.4658	0												Δ_{BAL}	0.3912	
	CGO	0.4541	0.3958	0											Δ_{CGO}	0.3996	
	CNE	0.3614	0.4213	0.4145	0										Δ_{CNE}	0.3358	
	COL	0.2849	0.4182	0.3788	0.2819	0									Δ_{COL}	0.3607	
	FM	0.6128	0.576	0.5975	0.541	0.6146	0								Δ_{FM}	0.5402	
	MOR	0.3151	0.4383	0.3548	0.208	0.1846	0.6042	0							Δ_{MOR}	0.3306	
	PCA	0.459	0.287	0.4375	0.3446	0.4785	0.5956	0.3779	0						Δ_{PCA}	0.409	
	PS	0.5943	0.5441	0.5615	0.4738	0.5729	0.4236	0.5597	0.5485	0					Δ_{PS}	0.5055	
	STE	0.3071	0.354	0.3301	0.3467	0.25	0.6111	0.2792	0.4727	0.6033	0				Δ_{STE}	0.3734	
	Mean														Δ_{SD}	0.4023	
	(b) Mean single locus (MnSL)	AZO	0													Δ_{AZO}	0.3891
		BAL	0.4651	0												Δ_{BAL}	0.4106
		CGO	0.4559	0.3959	0											Δ_{CGO}	0.4123
CNE		0.3639	0.422	0.4184	0										Δ_{CNE}	0.3626	
COL		0.2849	0.4209	0.3797	0.2819	0									Δ_{COL}	0.3725	
FM		0.6302	0.6013	0.6151	0.5708	0.6302	0								Δ_{FM}	0.5534	
MOR		0.3151	0.4391	0.3577	0.208	0.1846	0.6242	0							Δ_{MOR}	0.3434	
PCA		0.459	0.2574	0.4394	0.3456	0.482	0.6217	0.3786	0						Δ_{PCA}	0.4271	
PS		0.6014	0.5493	0.5672	0.4799	0.5773	0.4262	0.5622	0.5523	0					Δ_{PS}	0.516	
STE		0.3135	0.3541	0.333	0.3467	0.25	0.6267	0.28	0.4727	0.6076	0				Δ_{STE}	0.3818	
Mean															Δ_{SD}	0.4169	

Table 3 continued

Level of association	Occurrence identifier	Pairwise genetic distance Δ between occurrences											Components of the communities' differentiation		
		AZO	BAL	CGO	CNE	COL	FM	MOR	PCA	PS	STE	Δ_{SD}	Δ_{SD}		
(c) Multilocus (ML)	AZO	0												Δ_{AZO}	0.4045
	BAL	0.4653	0											Δ_{BAL}	0.415
	CGO	0.4552	0.3934	0										Δ_{CGO}	0.4233
	CNE	0.3631	0.4216	0.4184	0									Δ_{CNE}	0.3663
	COL	0.2841	0.4203	0.3791	0.2827	0								Δ_{COL}	0.3723
	FM	0.637	0.6015	0.6164	0.5702	0.6298	0							Δ_{FM}	0.5642
	MOR	0.3145	0.4407	0.3588	0.2128	0.184	0.6237	0						Δ_{MOR}	0.3575
	PCA	0.4586	0.2633	0.4388	0.3458	0.4823	0.6212	0.3797	0					Δ_{PCA}	0.4305
	PS	0.6019	0.5487	0.5693	0.4807	0.5768	0.4267	0.5625	0.5516	0				Δ_{PS}	0.5789
	STE	0.3129	0.3557	0.3325	0.3467	0.2495	0.6271	0.2803	0.4723	0.6082	0			Δ_{STE}	0.3839
	Mean													Δ_{SD}	0.4236

The mean values of Δ_{SD} are given and also shown as dotted circles in Fig. 7 as rounded values