

Table S1. Population genetic parameters estimated for dolomite and reference flora. Abbreviations: H_s = mean within-population gene diversity. Dif. = population differentiation, the following indicators apply: † = G_{st} or ‡ = F_{st} ; among-population variance from AMOVA)

Citation	Taxon	Data Type	H_s	Dif. (F_{ST} or G_{ST})
[68]	<i>Jurinea pinnata</i>	AFLP	0.135-0.198	0.37‡
[127]	<i>Convolvulus boissieri</i>	AFLP	0.033-0.127	0.395‡
[124]	<i>Viola cazortensis</i>	ISSR	0.215	0.104‡
[125]	<i>Helianthemum apeninum subsp. estevei</i>	SSR	0.47	
[125]	<i>Helianthemum panosum</i>	SSR	0.593	
[129]	<i>Erigeron parishii</i>	Alloenzyme	0.16	0.12†
[130]	<i>Eriogonium ovalifolium var. vineum</i>	Alloenzyme	0.19	0.07†
[131]	<i>Astragalus albens</i>	Alloenzyme	0.139	0.01†
[92]	Gypsum plants		0.196	0.423†, 0.466‡
[117]	Rare			0.212
	Common			0.198