

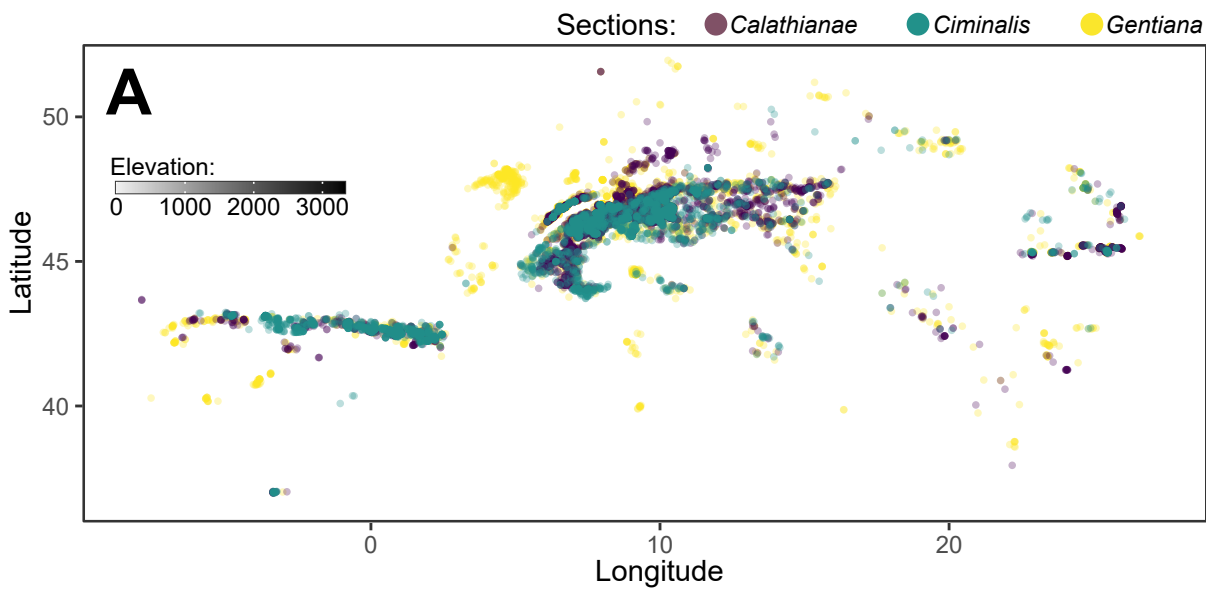
Supplementary Material (Appendix V)

“Phylogenetic incongruences derail the identification of correlates of diversification in *Gentiana* in the European Alpine System”– **Alpine Botany**

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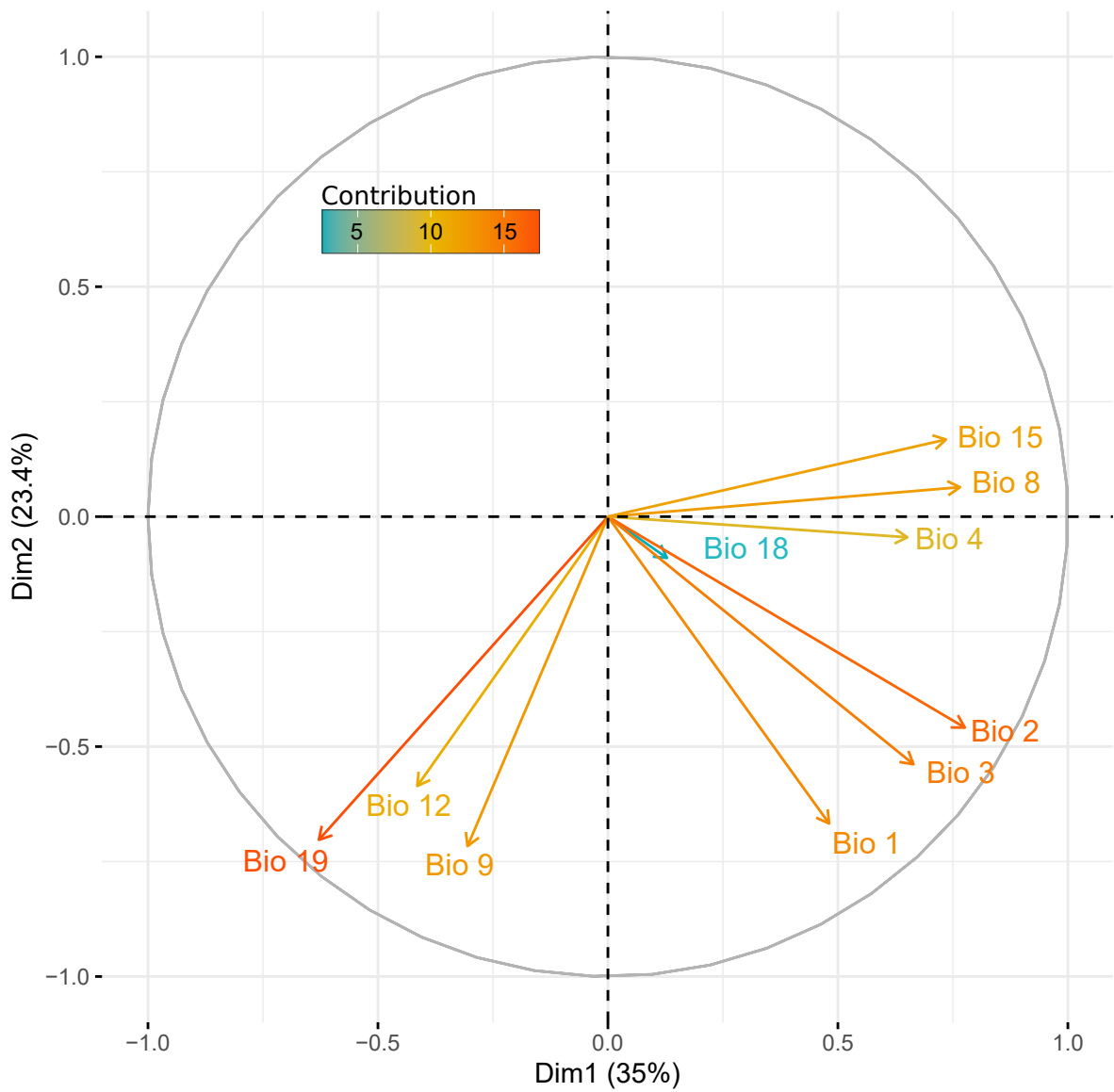
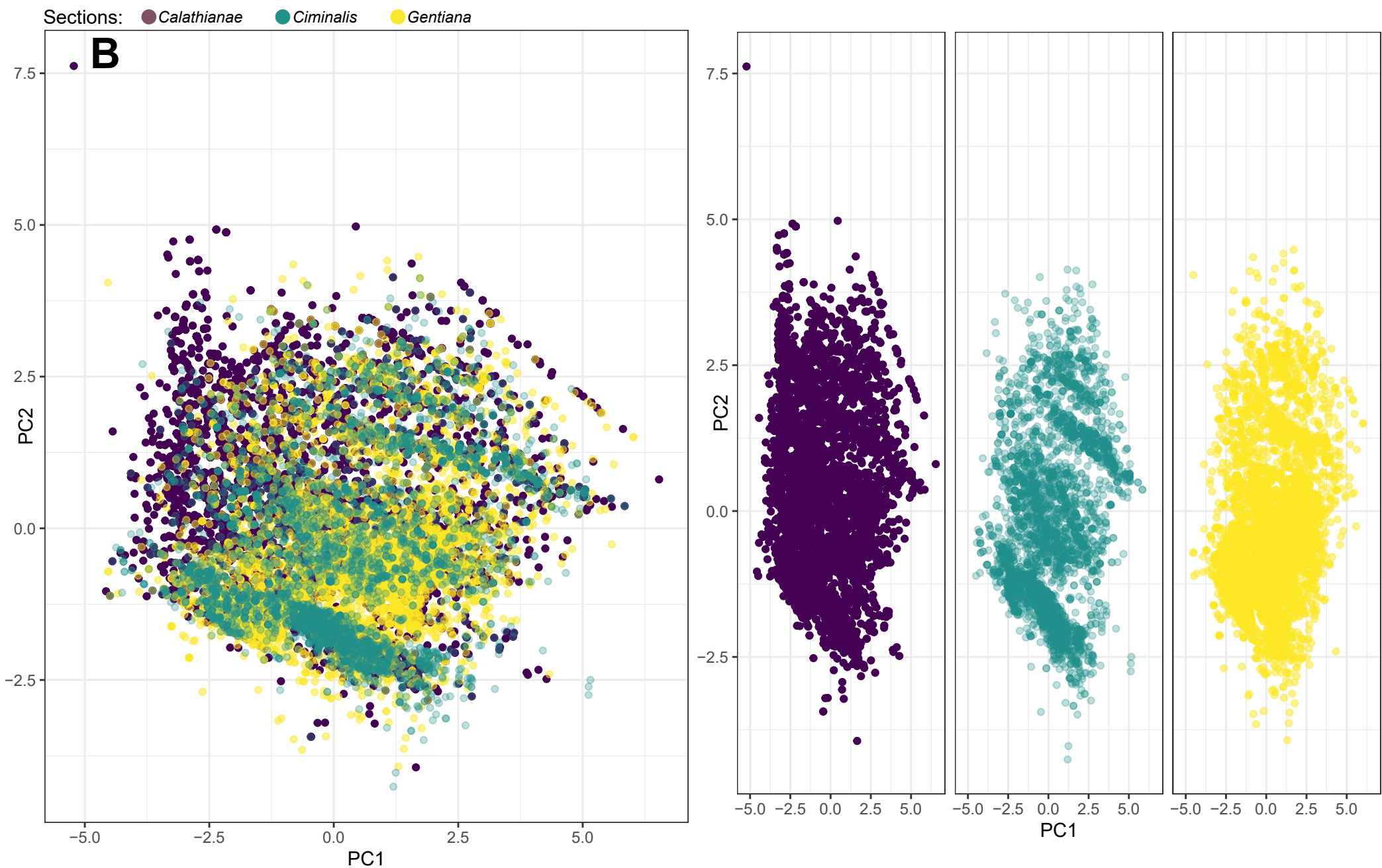
²LOEWE Centre for Translational Biodiversity Genomics (LOEWE-TBG), Senckenberganlage 25, 60325 Frankfurt am Main, Germany



Appendix Va. Comparison of climatic niches of species of three *Gentiana* sections occurring in the European Alpine System. **A.** Distribution map based on occurrence data. **B.** Principal component analysis highlighting *G.* section *Calathianae*, *G.* sect. *Ciminalis* and *G.* sect. *Gentiana* plotted together and separately (C.), based on nine selected WorldClim bioclimatic variables

Note: Incongruences do not affect comparions among sections

Geographic overlap:
Calathianae - *Ciminalis*: 0.65, *Ciminalis* - *Gentiana*: 0.68
Gentiana - *Calathianae*: 0.71



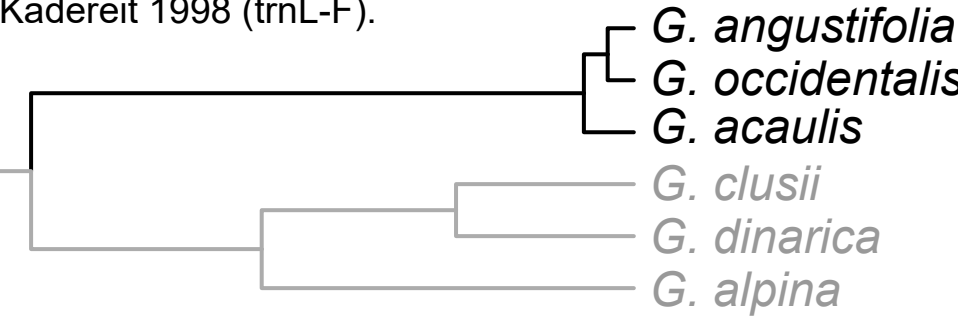
Wordclim Variables

- Bio 19 Precipitation of Coldest Quarter
- Bio 2 Mean Diurnal Range
- Bio 3 Isothermality
- Bio 1 Annual Mean Temperature
- Bio 9 Mean Temperature of Driest Quarter
- Bio 8 Mean Temperature of Wettest Quarter
- Bio 12 Annual Precipitation
- Bio 4 Temperature Seasonality
- Bio 18 Precipitation of Warmest Quarter

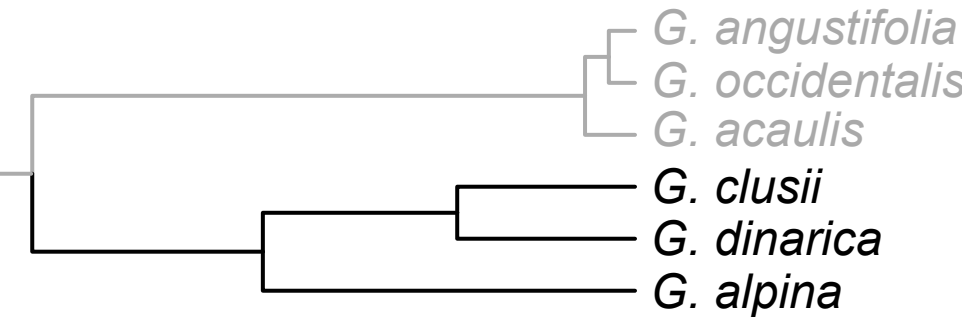
Appendix Vb. Comparison of climatic niches among species of *G.* sect. *Ciminalis* occurring in the European Alpine System. **A.** Comparison among all species. **B.** and **C.** Comparison among species within each main clade. Principal component analysis based on worldclim climate variables.

Note: In *G.* sect. *Ciminalis*, phylogenetic incongruences occur and thus, we present here comparisons based on both plastid and rDNA-cistron phylogenies

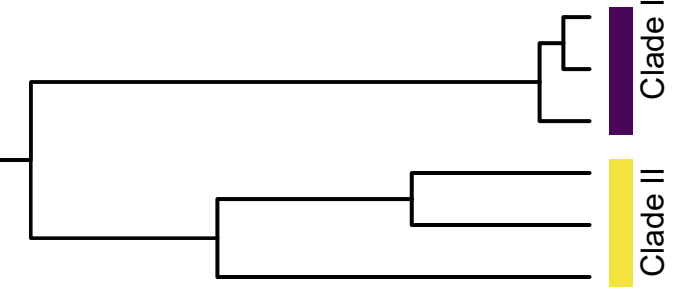
Phylogenetic relationships: plastid tree
We do not include *G. ligustica* in our phylogenetic reconstructions, but we show it here because this species clusters with the clade of *G. acaulis* in Hungerer & Kadereit 1998 (trnL-F).



Phylogenetic relationships: plastid tree
Although *G. ligustica* appears to cluster close to *G. acaulis* in Hungerer & Kadereit 1998 (with ITS), support values were weak and thus we include *G. ligustica* here as well.

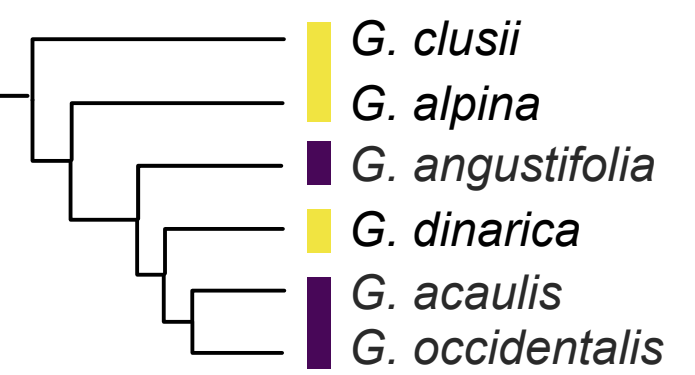


Phylogenetic relationships: plastid tree
Comparison of clade I and II, without *G. ligustica*.

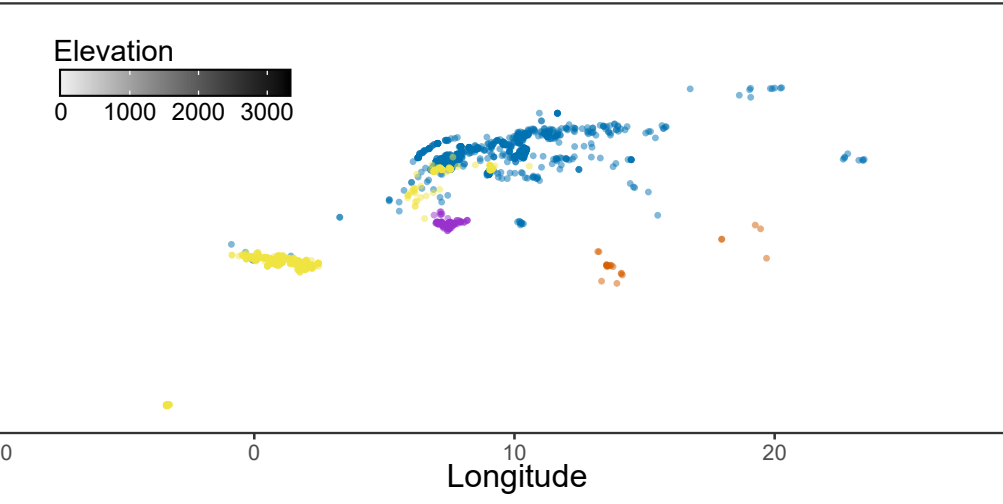
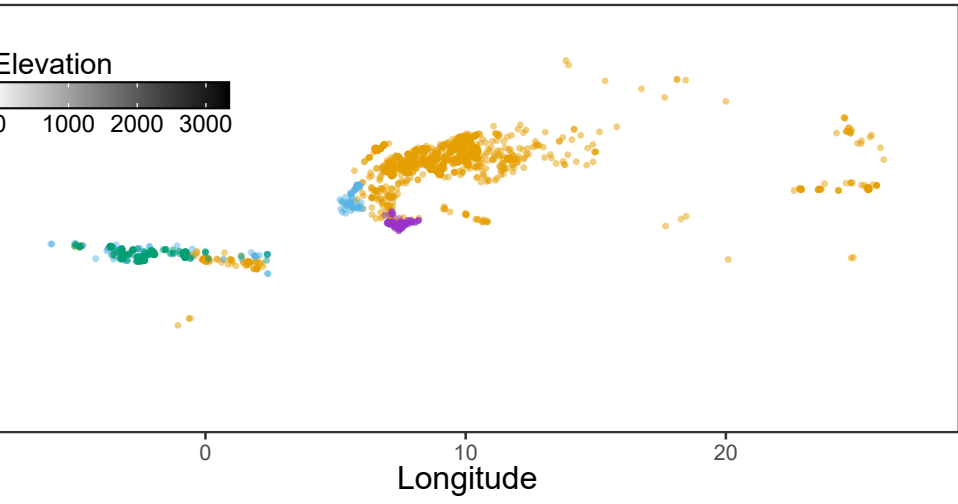
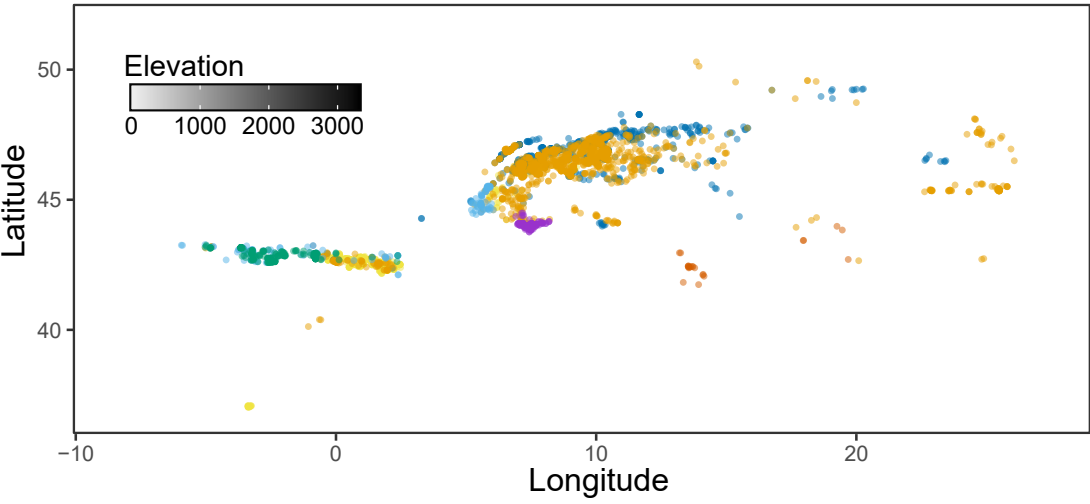
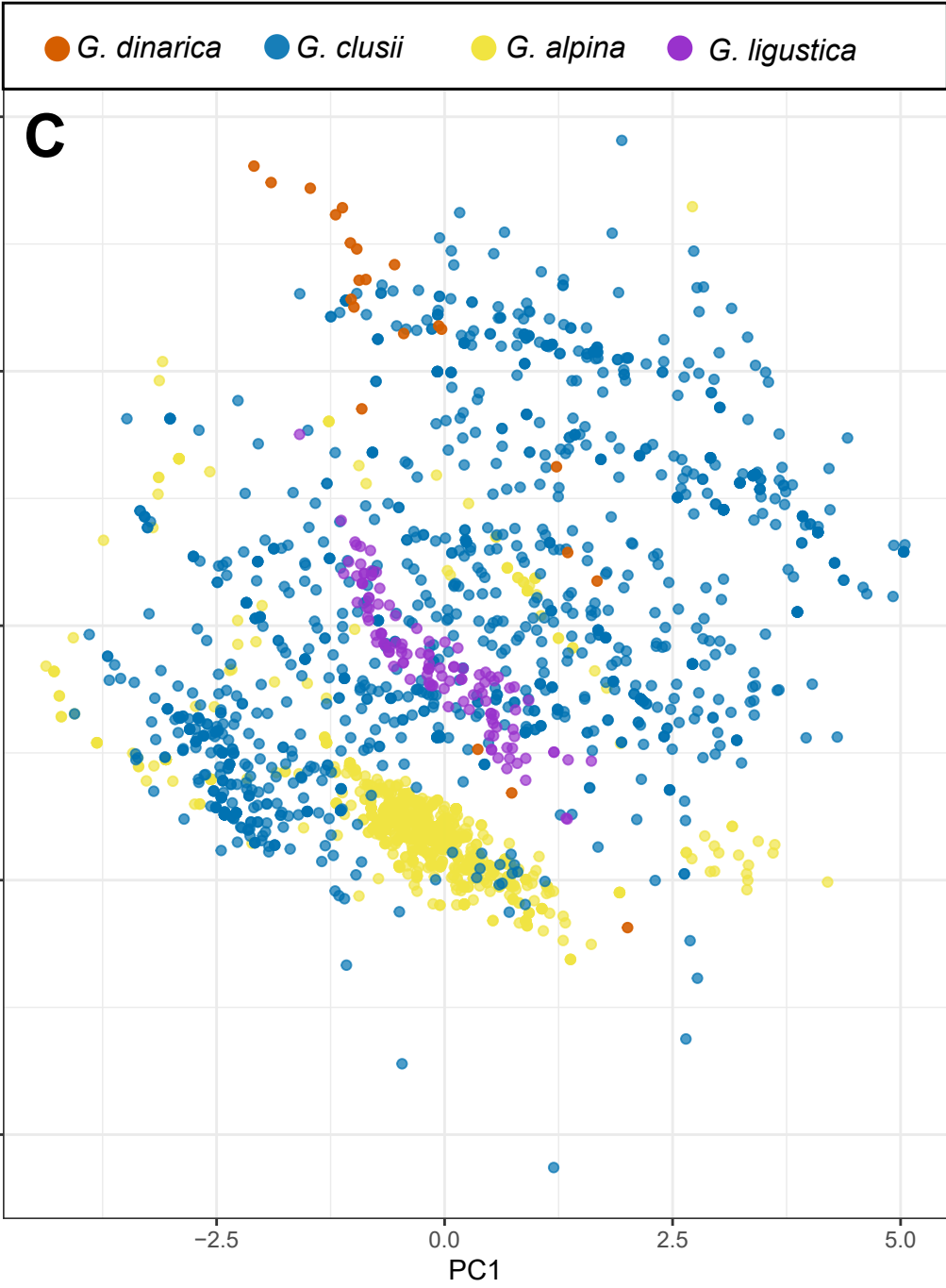
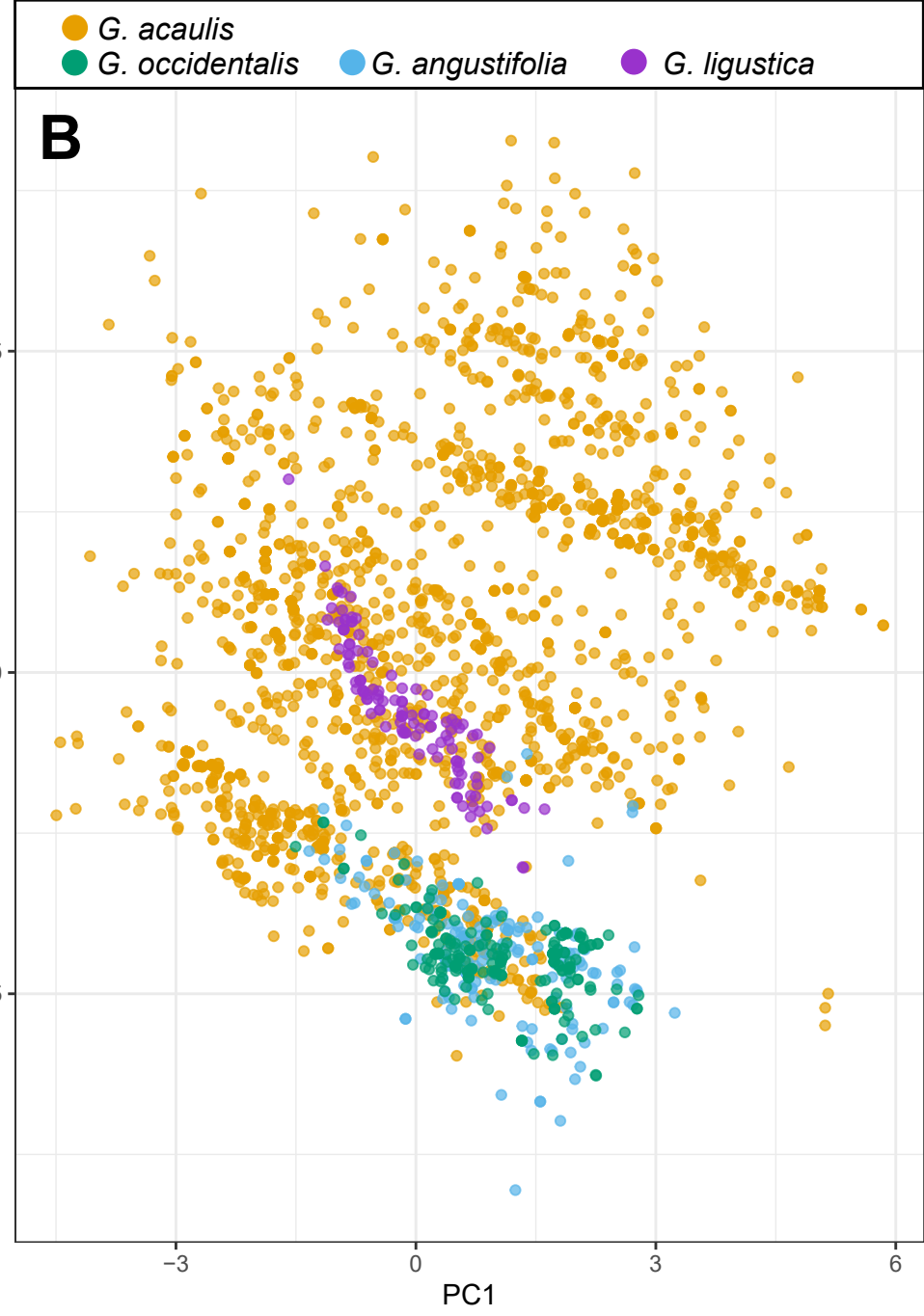
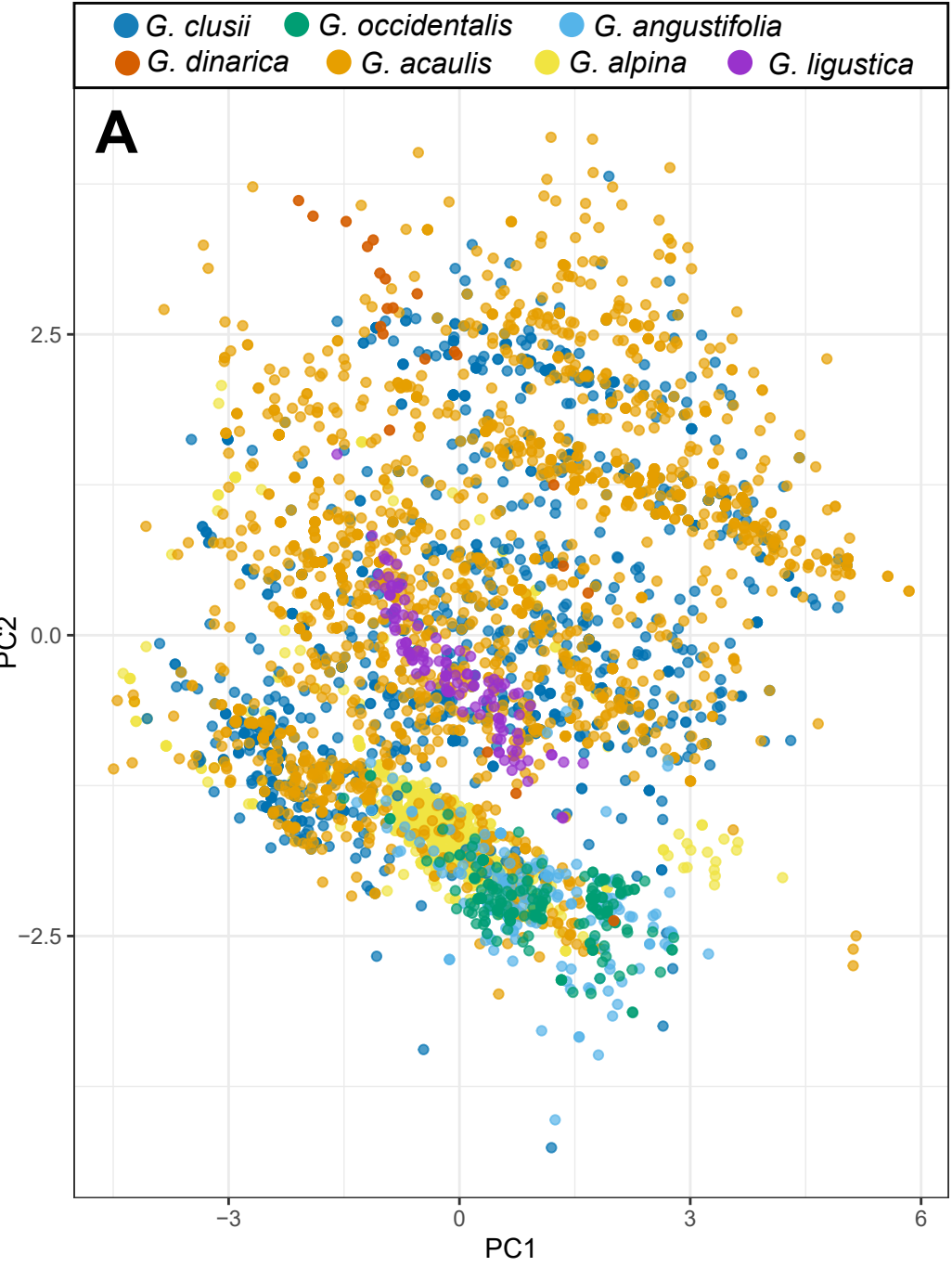


Geographical overlap:
Cp tree, Clade I and II: 0.58

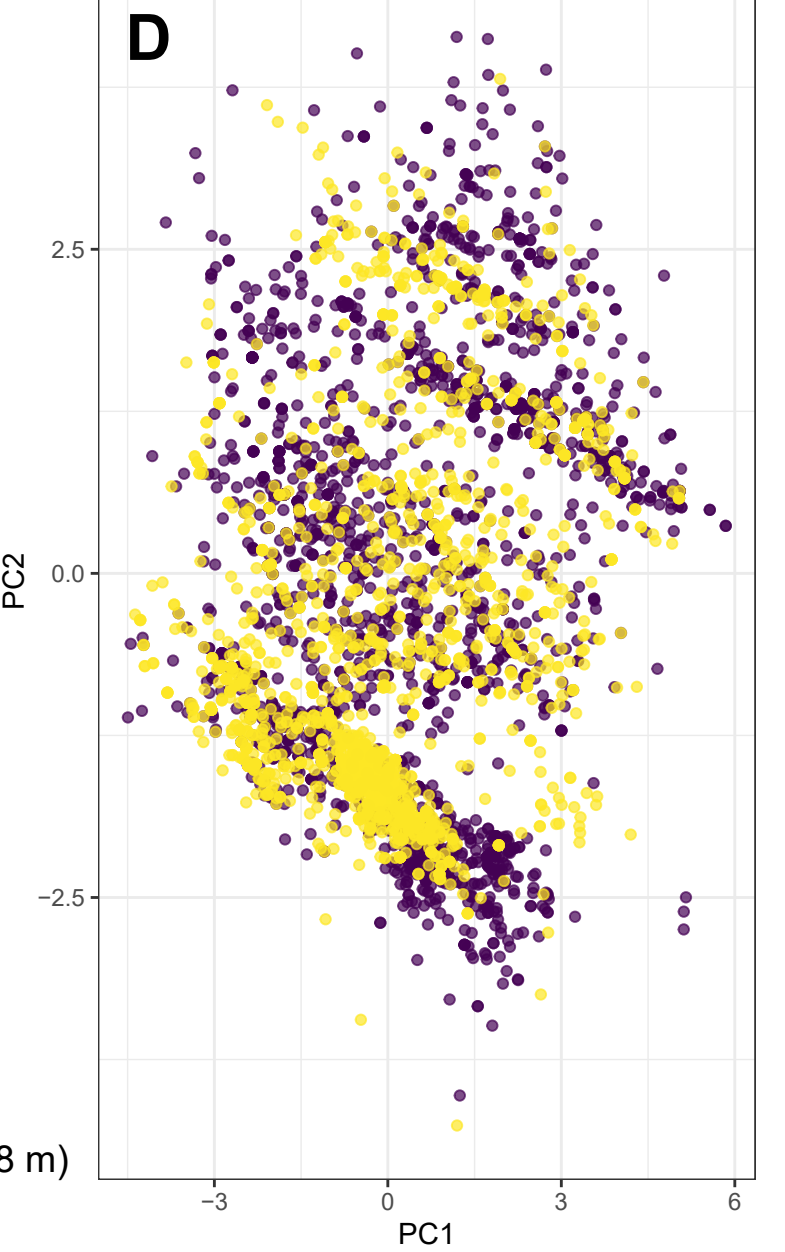
Phylogenetic relationships: rDNA-Cistron
Comparison of clade I and II, without *G. ligustica*. Purple is Clade I, yellow is Clade II.



● *G. acaulis*, *G. occidentalis*, *G. angustifolia*
● *G. clusii*, *G. dinarica*, *G. alpina*

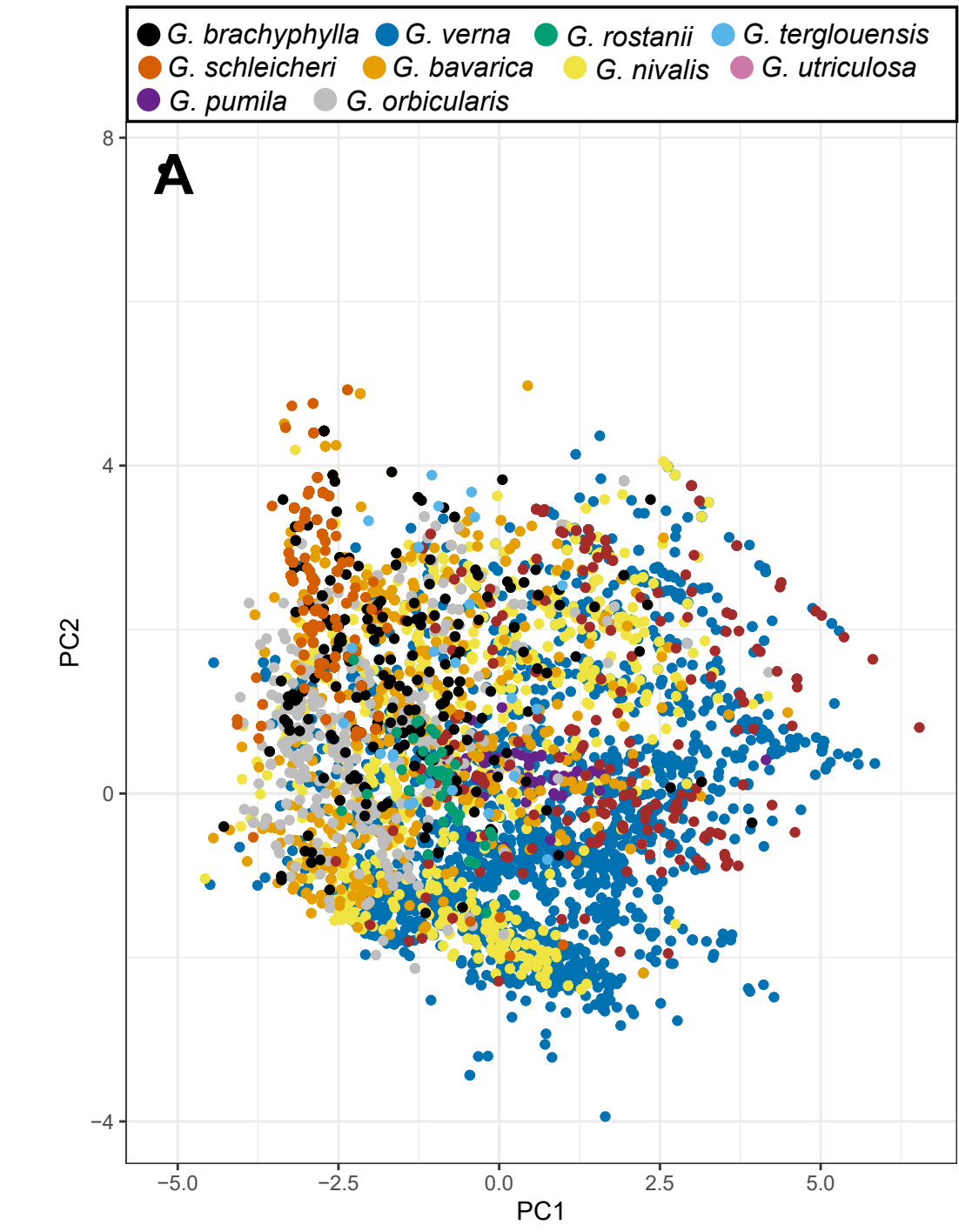


Species mean elevation:
G. acaulis (1793 +/- 493m), *G. alpina* (2094 +/- 315 m), *G. angustifolia* (977 +/- 350 m), *G. clusii* (1575 +/- 530 m), *G. dinarica* (1488 +/- 292 m), *G. ligustica* (1398 +/- 466 m), *G. occidentalis* (1013 +/- 308 m)

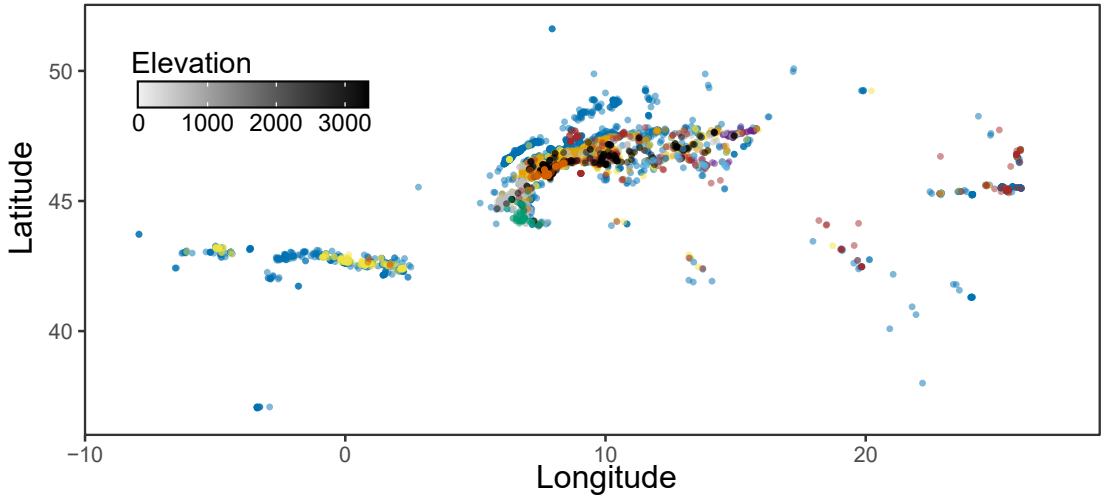


Appendix Vd. Comparison of climatic niches among species of *G.* sect. *Calathianae* occurring in the European Alpine System. **A.** Comparison among all species. **B. to F.** Comparison among selected species. Principal component analysis based on worldclim climate variables.

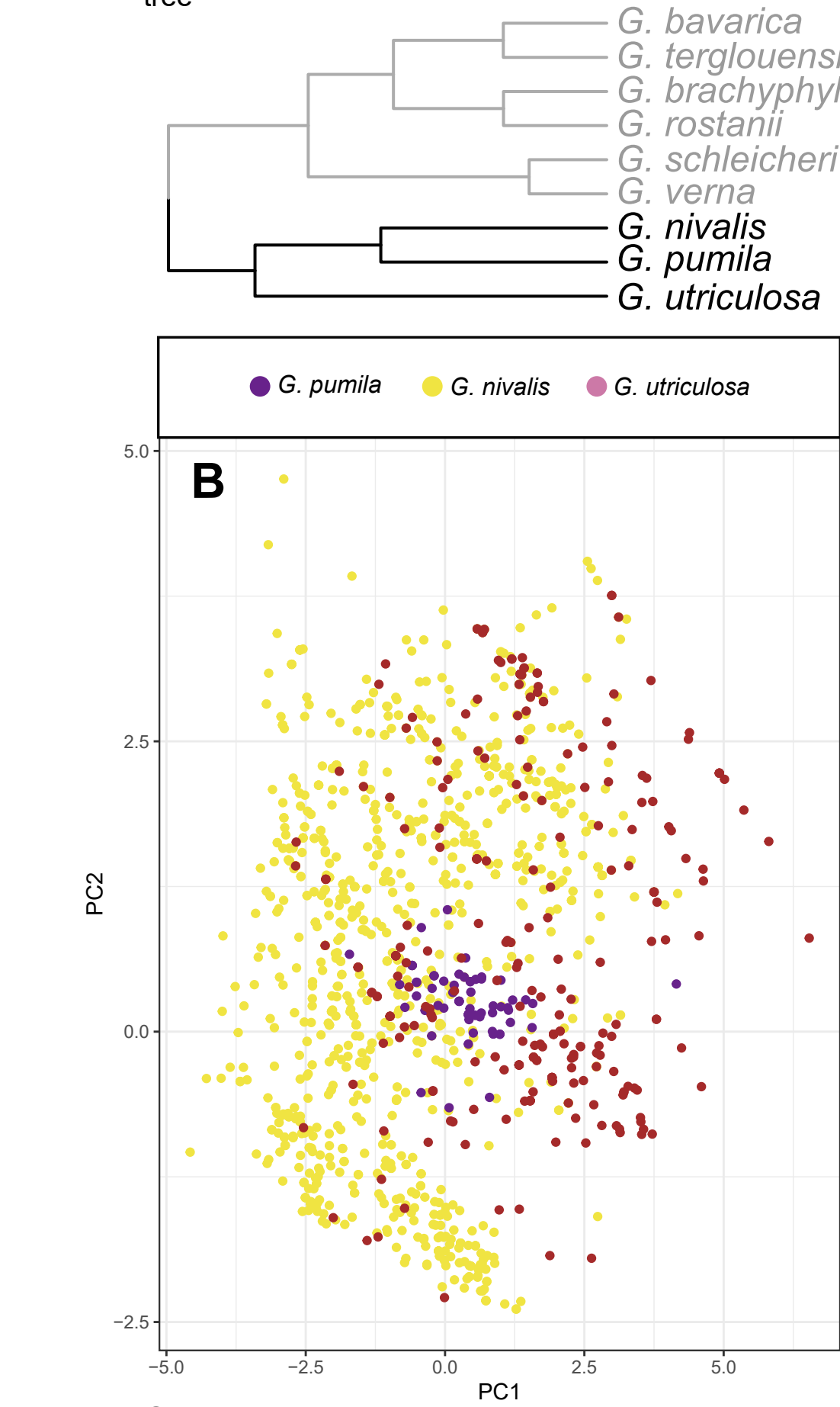
Note: In *G.* sect. *Calathianae*, phylogenetic incongruences occur and thus, we present here comparisons based on both plastid and rDNA-cistron phylogenies



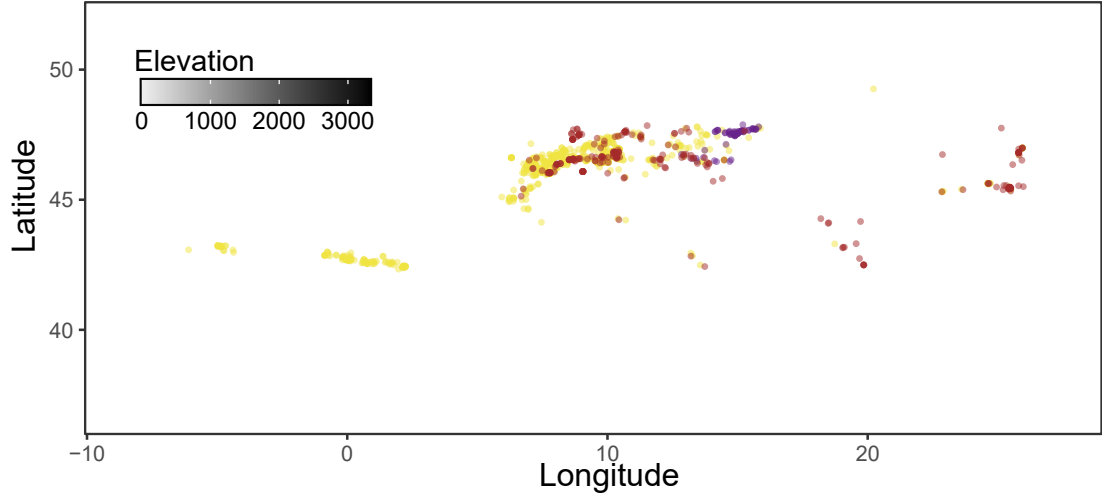
Mean Geographic overlap:
among all species (0.45)



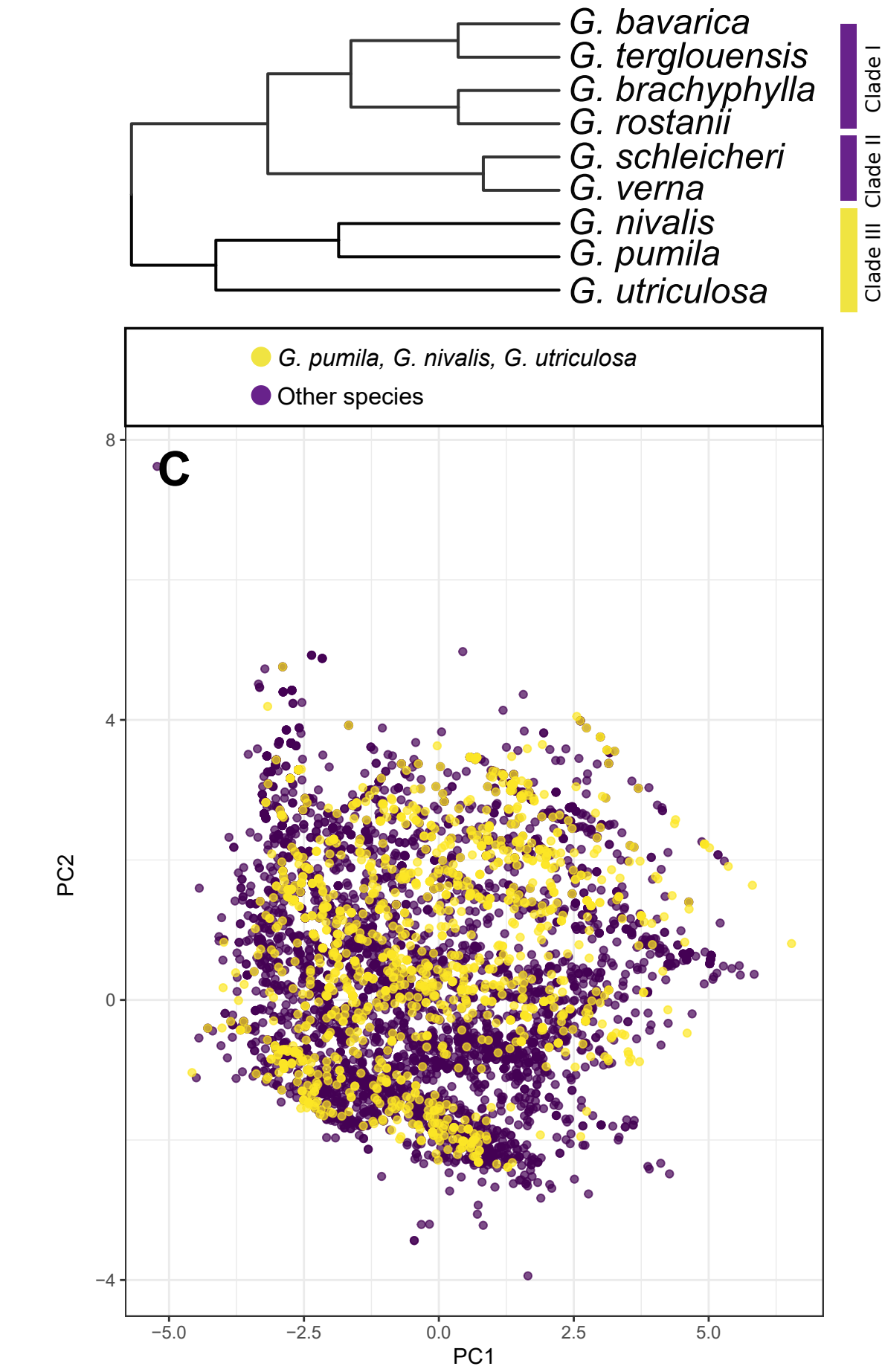
Phylogenetic relationships: plastid tree
Note: phylogenetic relationships among *G. nivalis*, *G. utriculosa* and *G. pumila* are the same in the rDNA-cistron tree



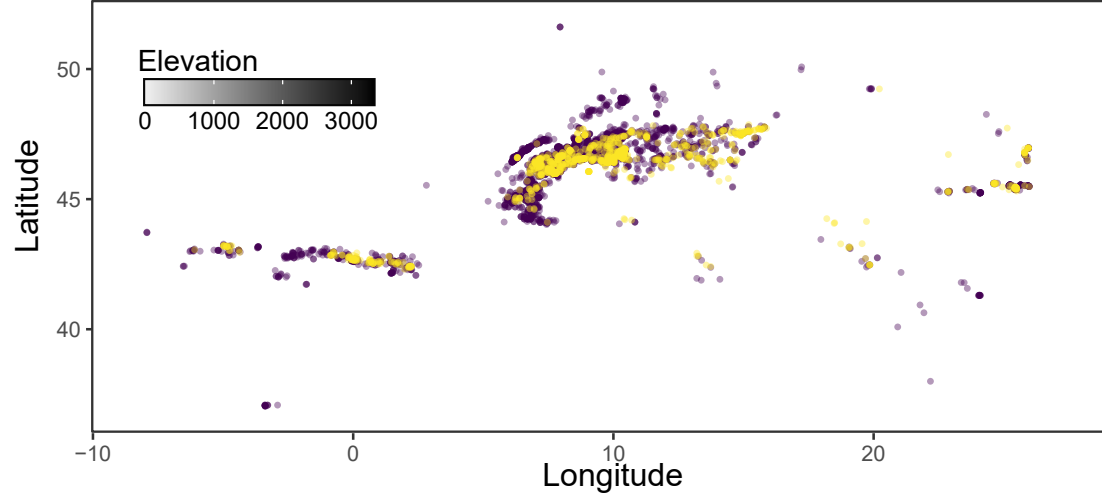
Mean Geographic overlap:
G. utriculosa vs. *G. nivalis* (0.48),
G. utriculosa vs. *G. pumila* (0.20), *G. pumila* vs. *G. nivalis* (0.32)



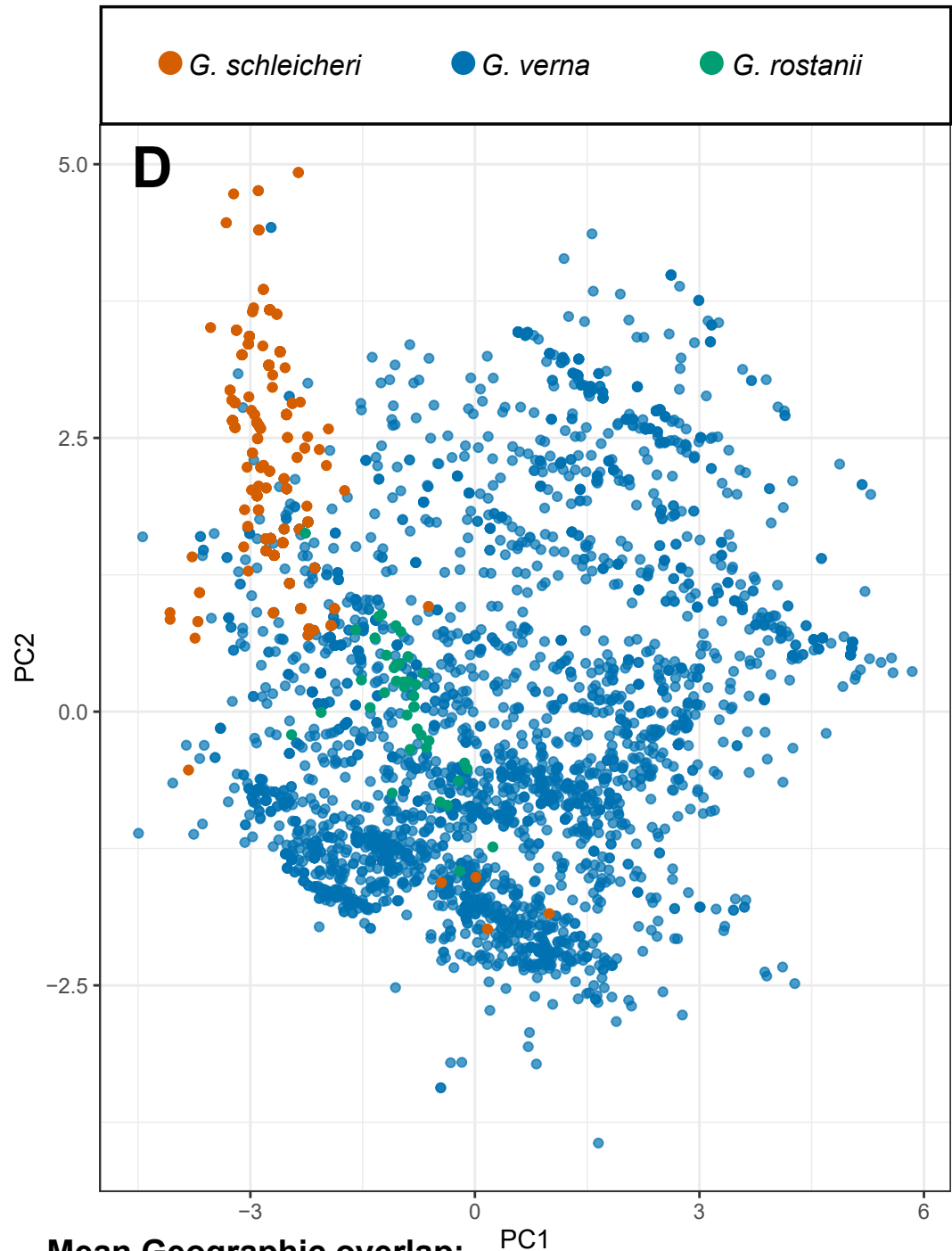
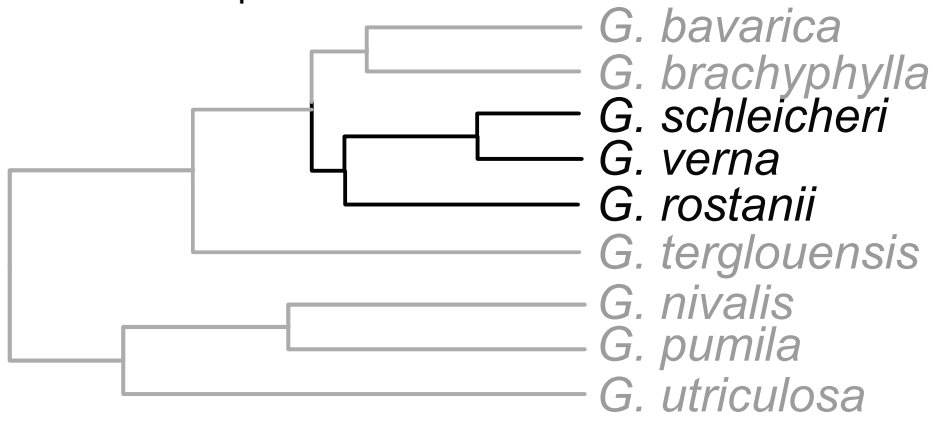
Phylogenetic relationships: plastid tree
Note: Clade I and Clade II are also monophyletic in the rDNA-cistron tree.



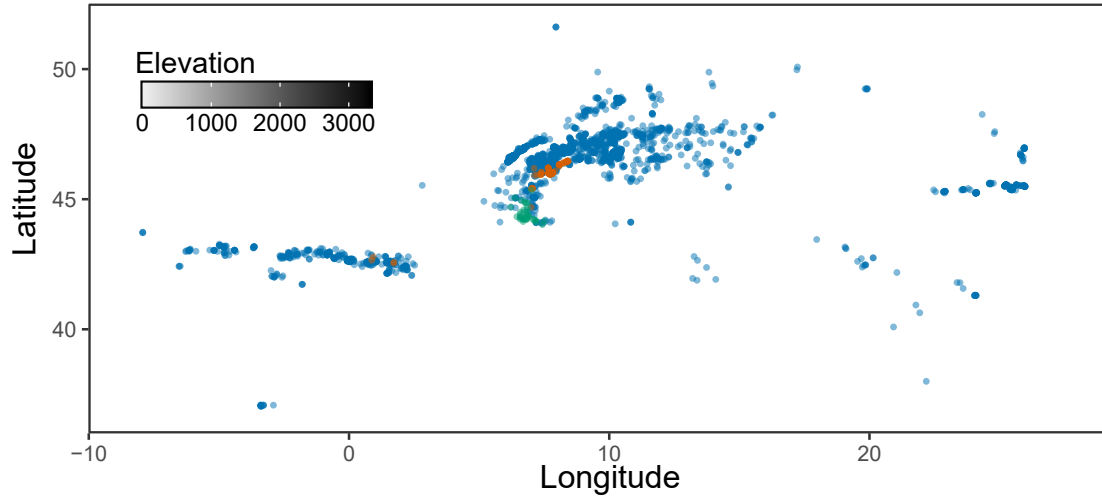
Mean Geographic overlap:
Clade 1 vs. Clade 2 (0.73),
Clade 1 vs. Clade 3 (0.64), Clade 1 vs. Clade 3 (0.64)



Phylogenetic relationships: rDNA-cistron tree
Note: *G. schleicheri* and *G. verna* are sister species also in the plastid tree.



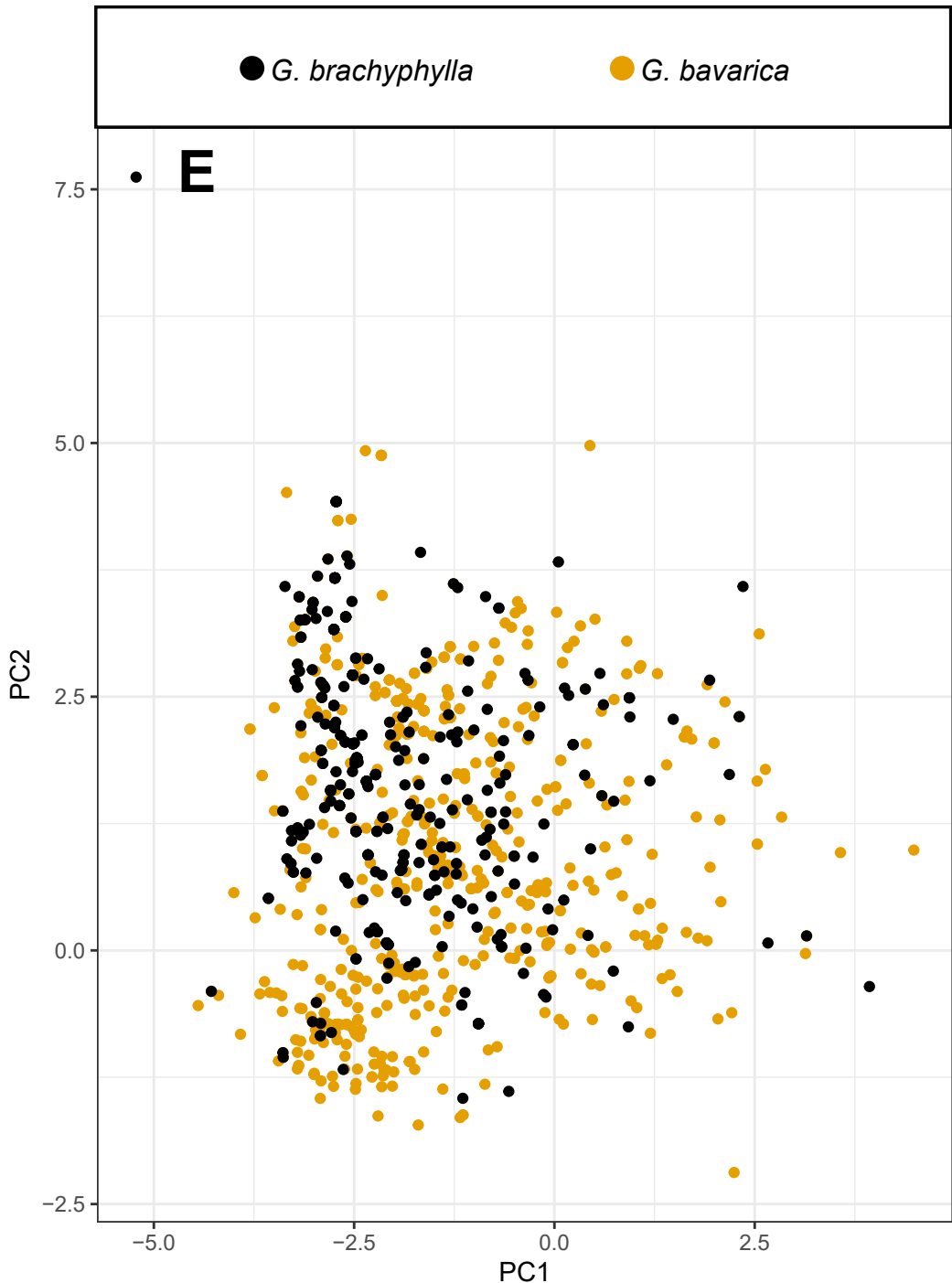
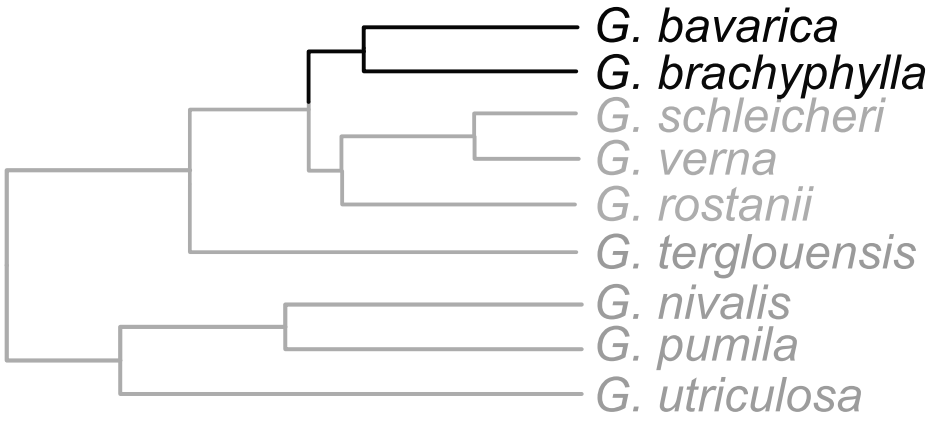
Mean Geographic overlap:
G. rostanii vs. *G. schleicheri* (0.07),
G. verna vs. *G. schleicheri* (1.00), *G. verna* vs. *G. rostanii* (0.59)



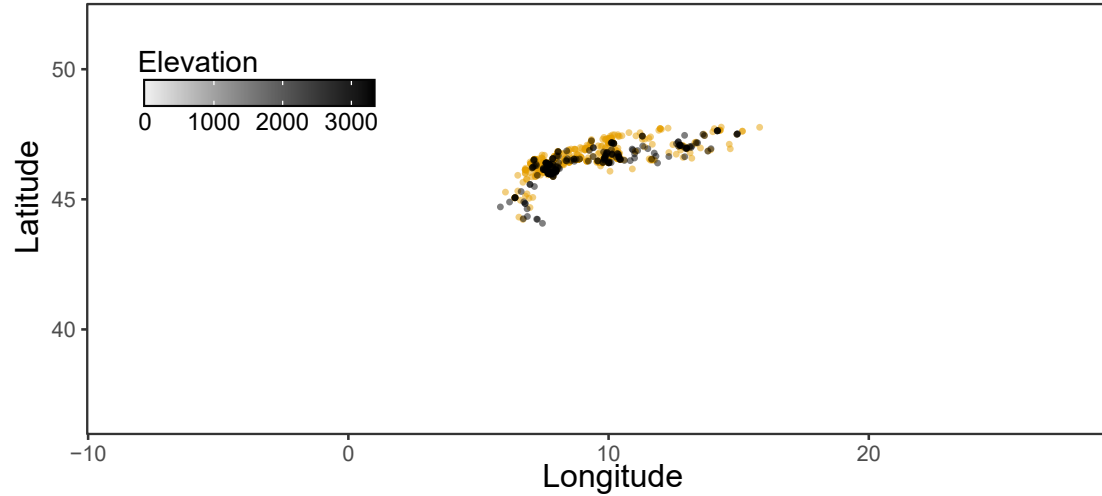
Species mean elevation:

G. bavarica (2100 +/- 468 m), *G. brachyphylla* (2333 +/- 406 m), *G. nivalis* (2022 +/- 461 m), *G. orbicularis* (2194 +/- 405 m), *G. pumila* (1328 +/- 177 m), *G. rostanii* (2161 +/- 198 m), *G. schleicheri* (2648 +/- 310 m), *G. terglouensis* (1651 +/- 383 m), *G. utriculosa* (1567 +/- 589 m), *G. verna* (1586 +/- 593 m)

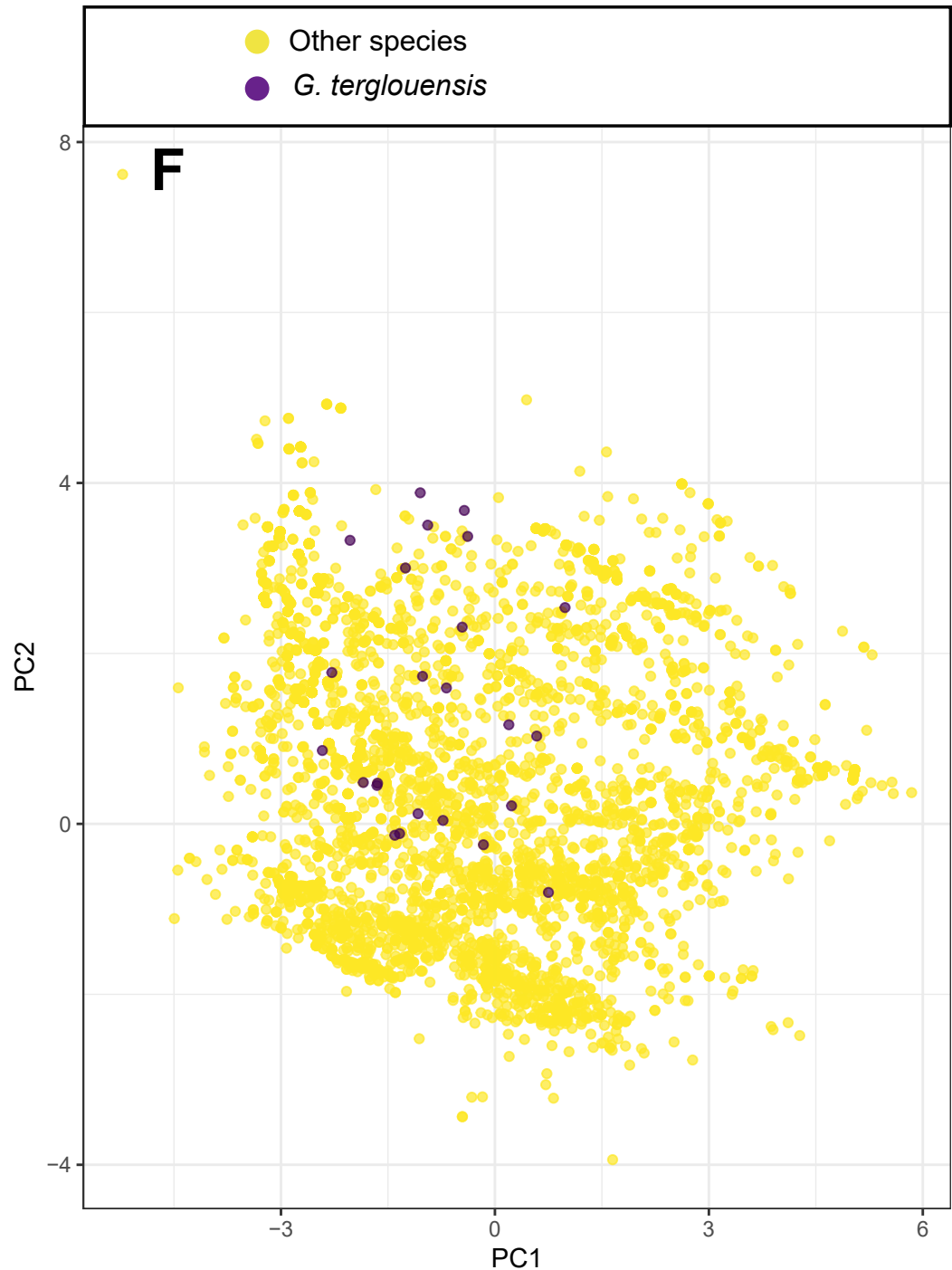
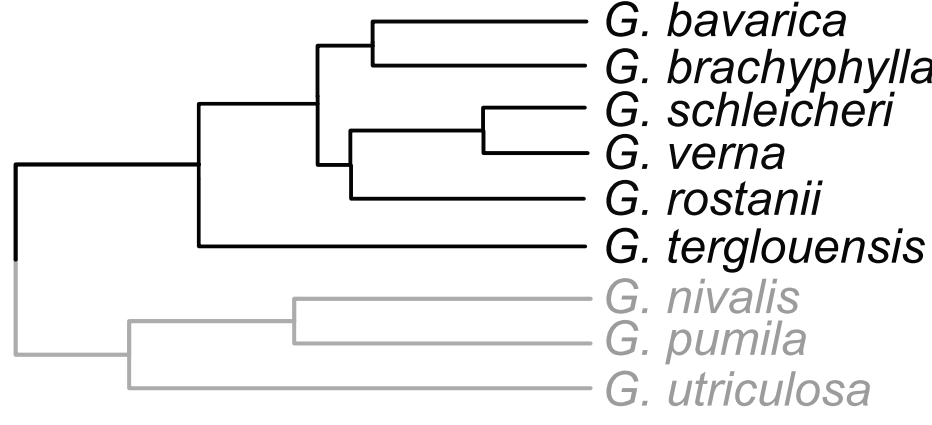
Phylogenetic relationships: rDNA-cistron tree
Note: *G. terglouensis* is the sister species of *G. bavarica* in the plastid tree.



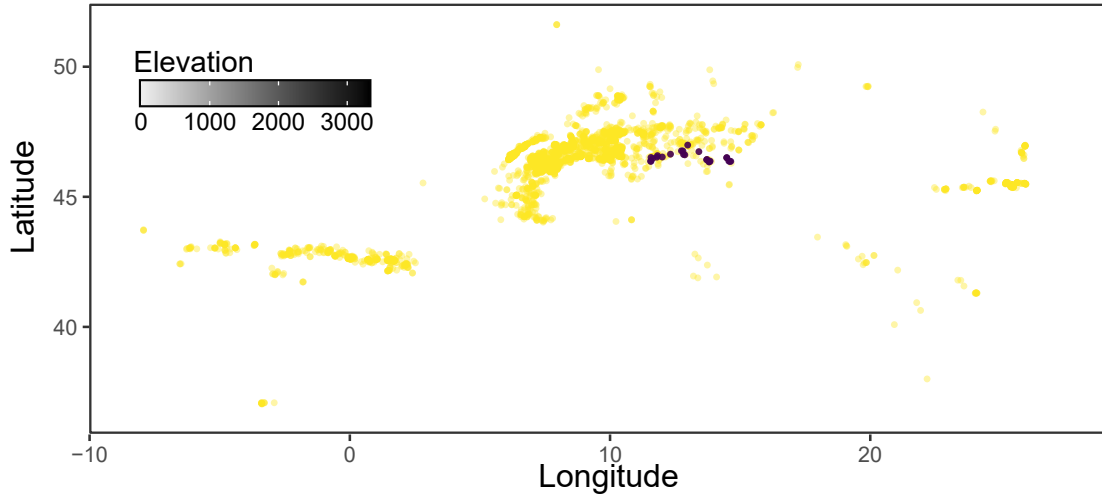
Mean Geographic overlap:
G. bavarica vs. *G. brachyphylla* (0.69)



Phylogenetic relationships: rDNA-cistron tree
Note: *G. terglouensis* is the sister species of *G. bavarica* in the plastid tree.



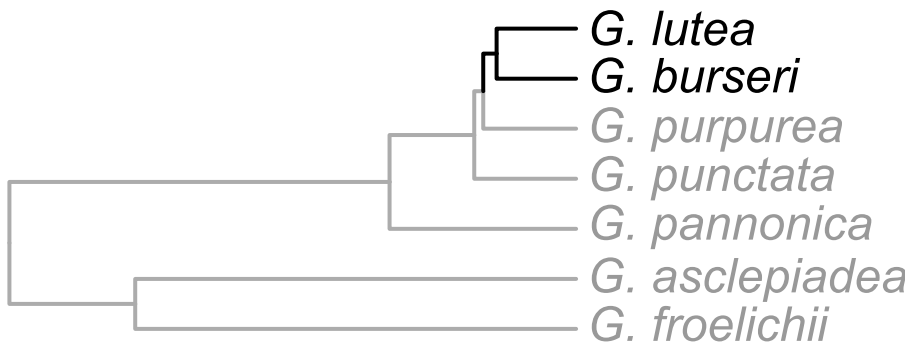
Mean Geographic overlap:
G. terglouensis vs. *G. bavarica* (0.27)



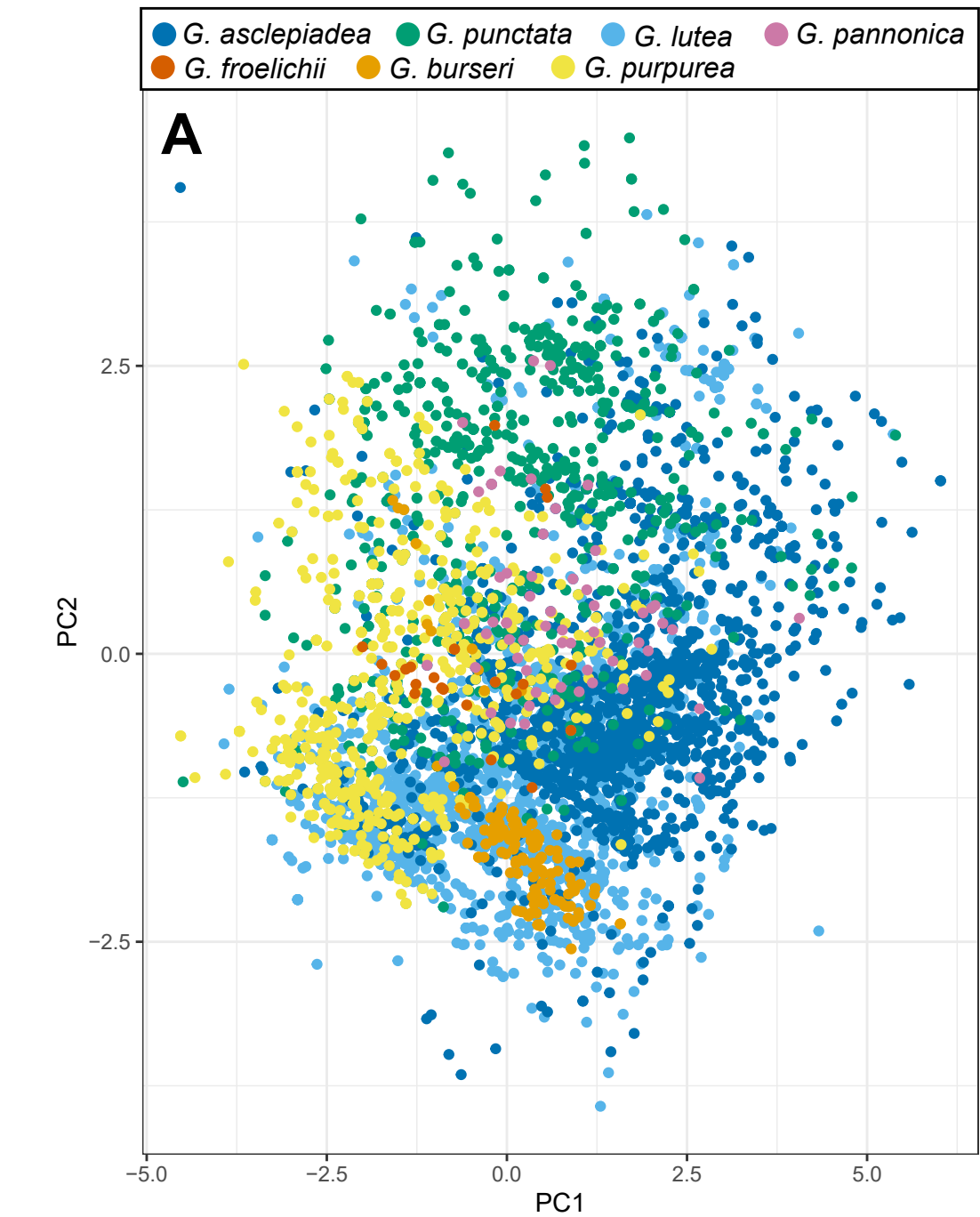
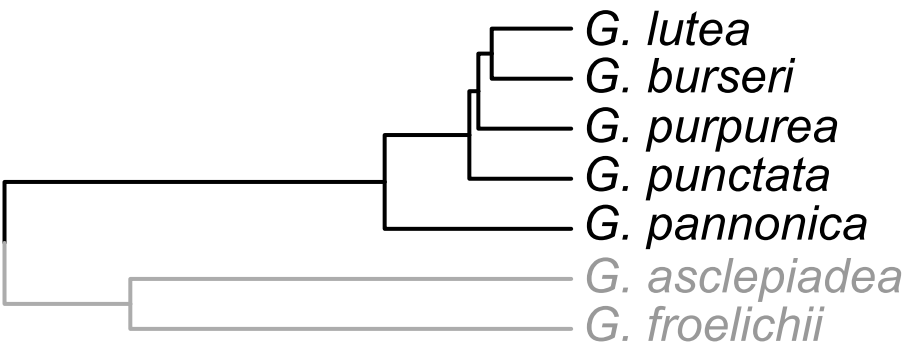
Appendix Vd. Comparison of climatic niches among species of *G.* sect. *Gentiana* occurring in the European Alpine System. **A.** Comparison among all species. **B. to D.** Comparison among species within each main clade. **E. and F.** comparisons among silicolous species and between silicolous and calcicolous species. Silicolous species cluster together only in the rDNA-cistron tree. Principal component analysis based on worldclim climate variables.

Note: In *G.* sect. *Gentiana*, phylogenetic incongruences occur and thus, we present here comparisons based on both plastid and rDNA-cistron phylogenies

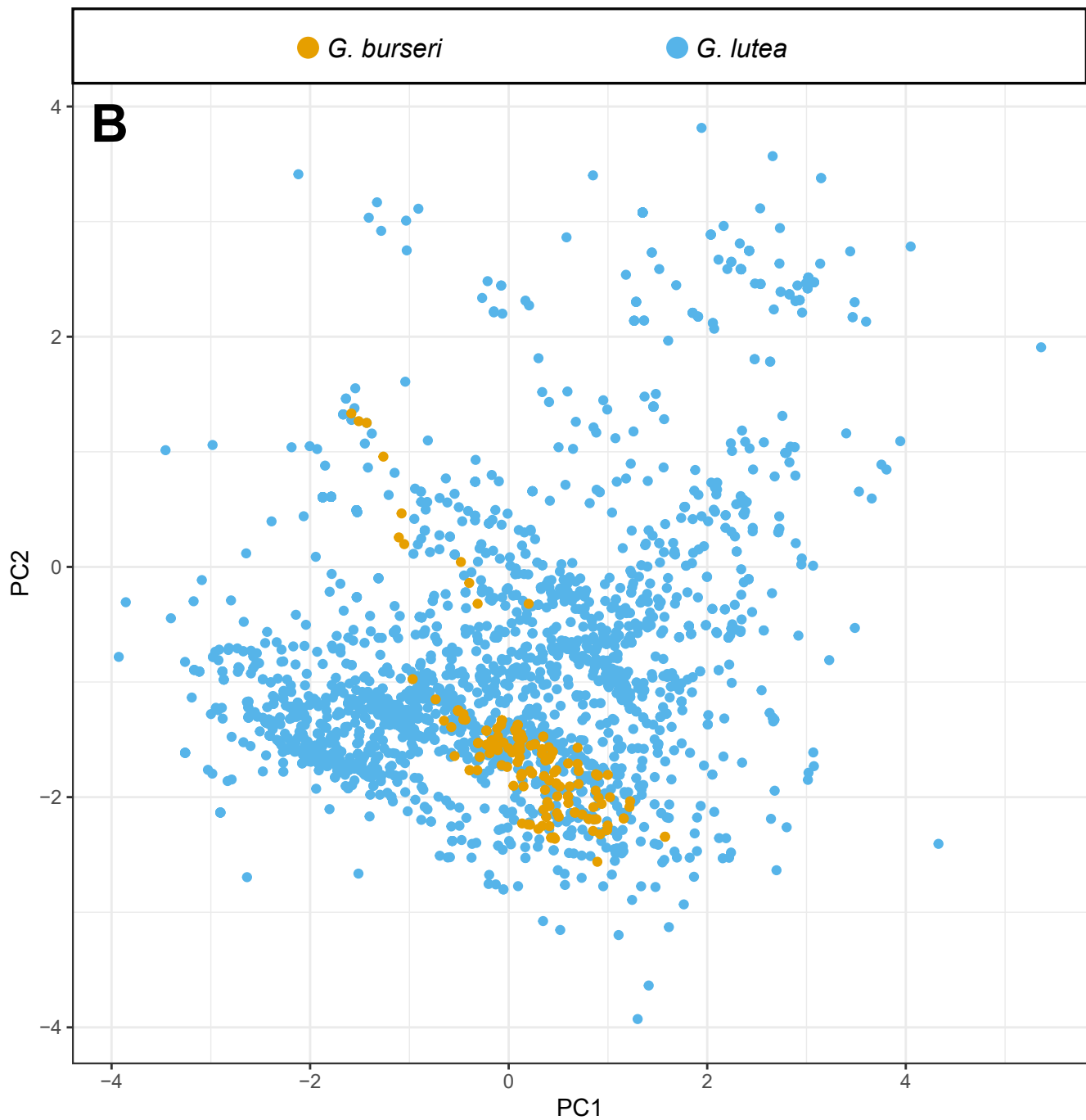
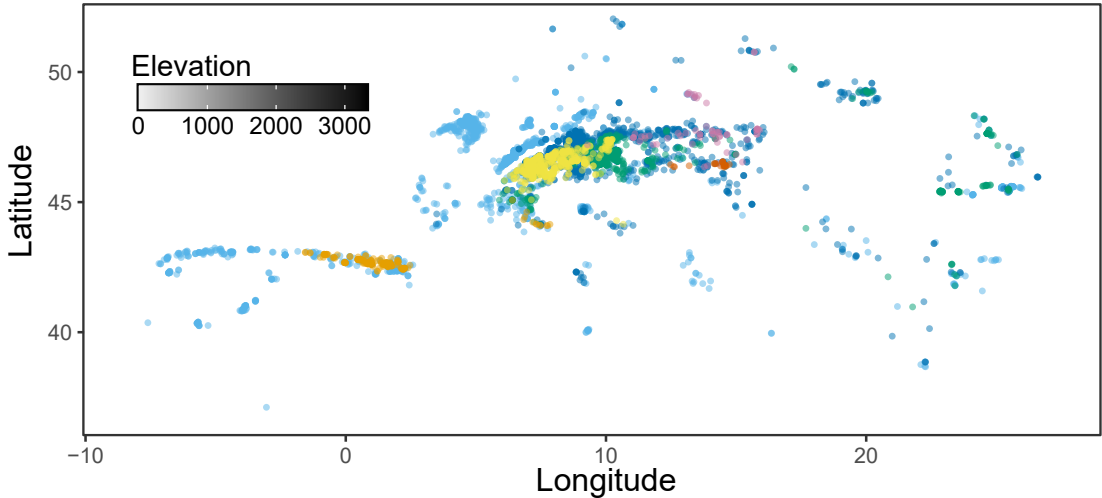
Phylogenetic relationships: plastid tree



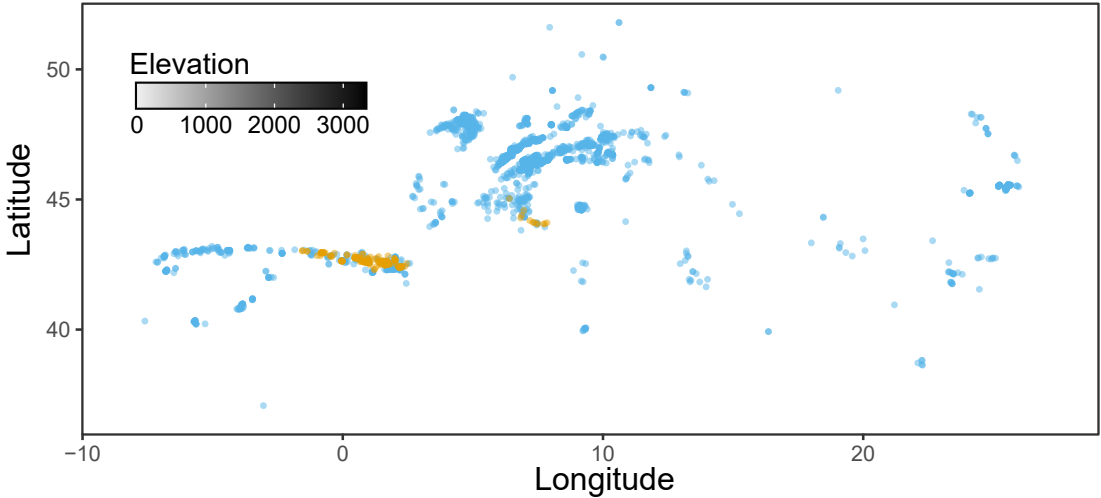
Phylogenetic relationships: plastid tree



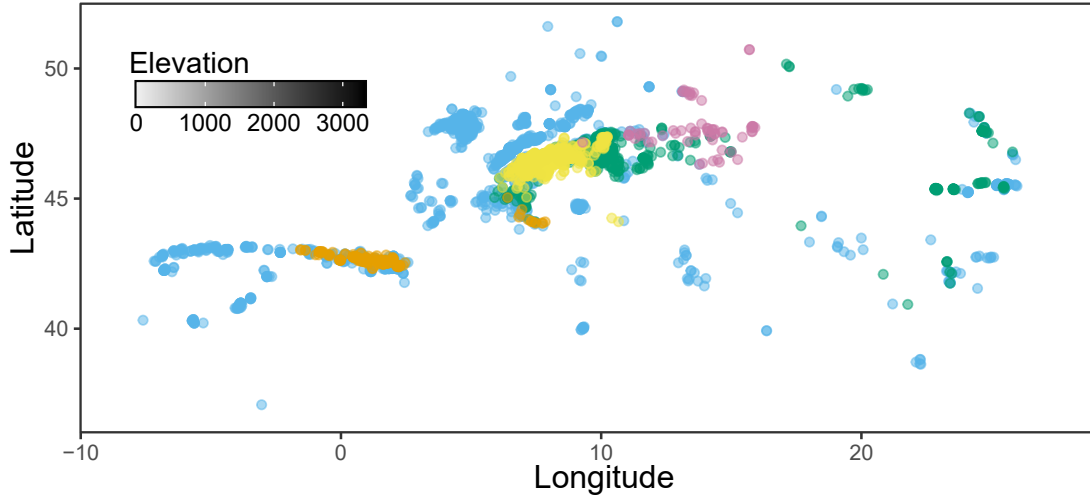
Mean Geographic overlap:
among all species (0.31)



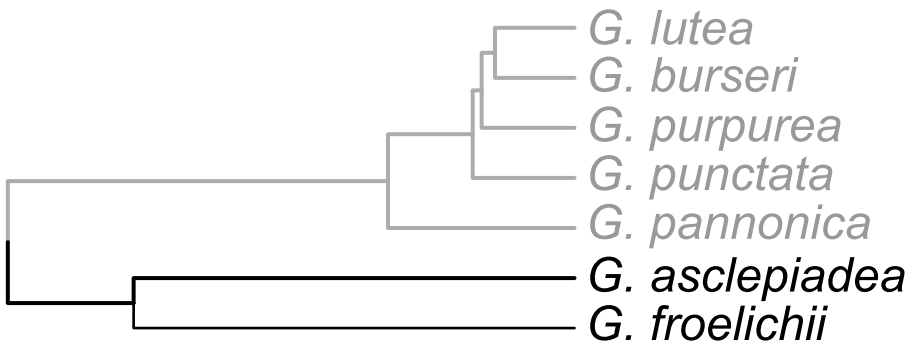
Mean Geographic overlap:
G. lutea vs. *G. burseri* (0.68)



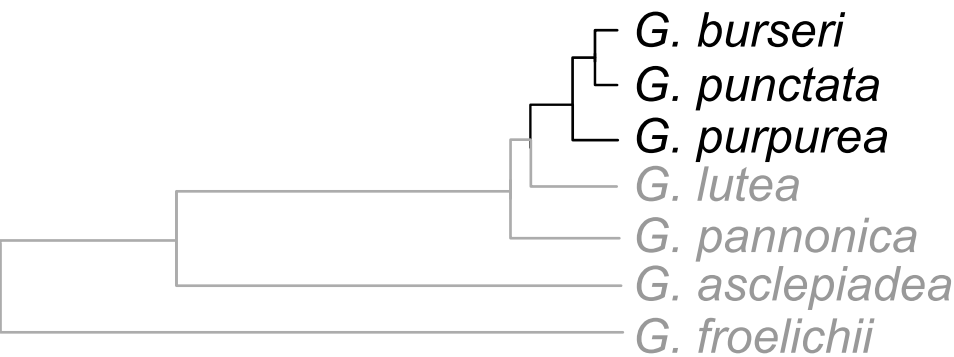
Mean Geographic overlap:
G. pannonica vs. *G. burseri* (0.00), *G. pannonica* vs. *G. lutea* (0.23),
G. punctata vs. *G. lutea* (0.50), *G. purpurea* vs. *G. lutea* (0.64),
G. purpurea vs. *G. pannonica* (0.02)



Phylogenetic relationships: plastid tree

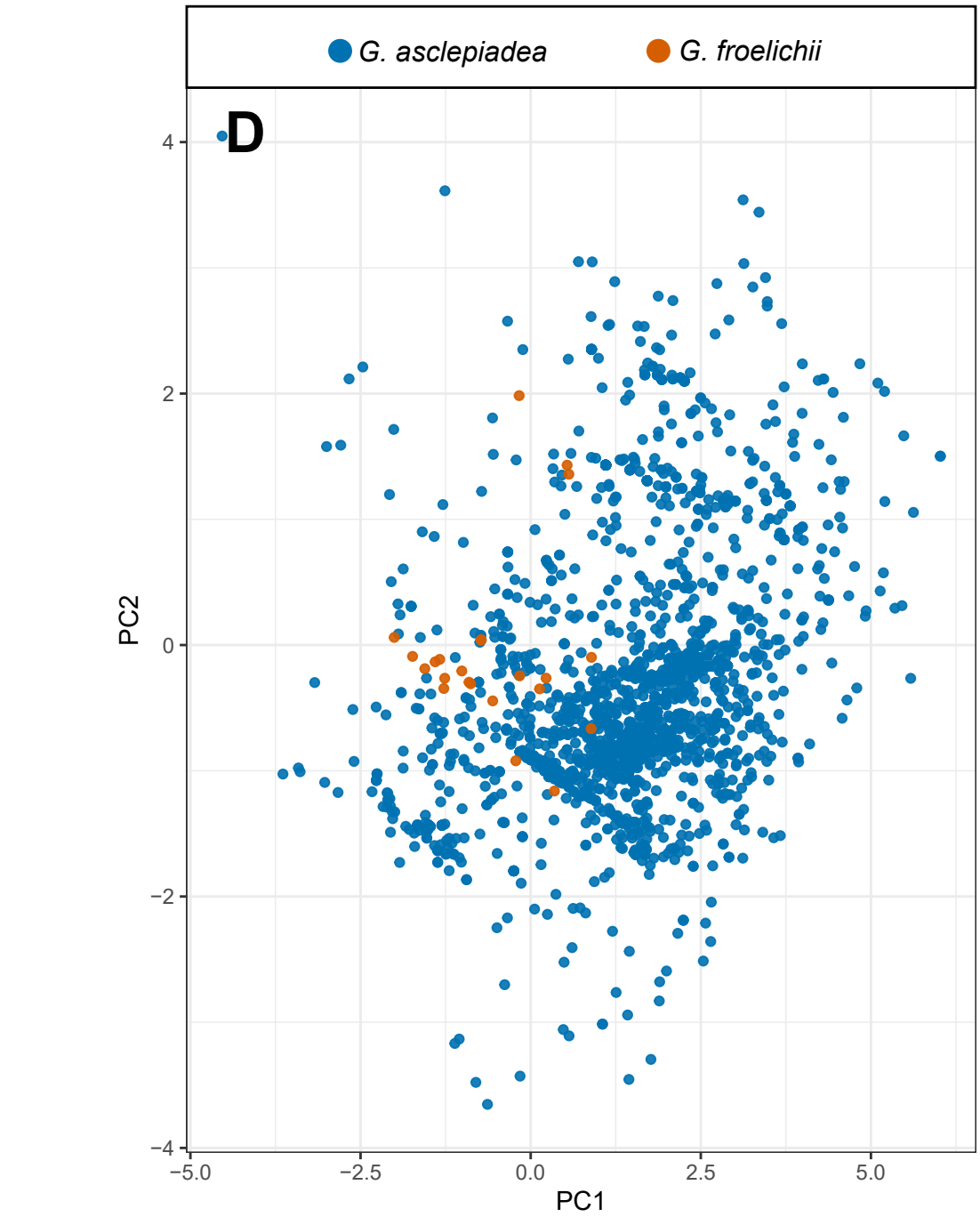
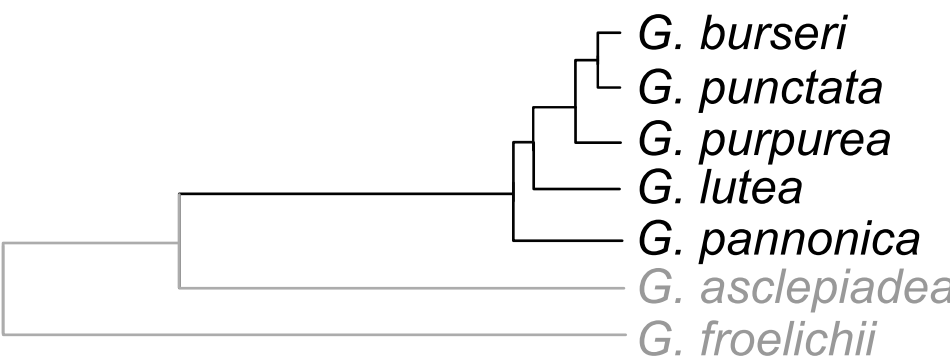


Phylogenetic relationships: rDNA-cistron tree
Comparison among silicolous species

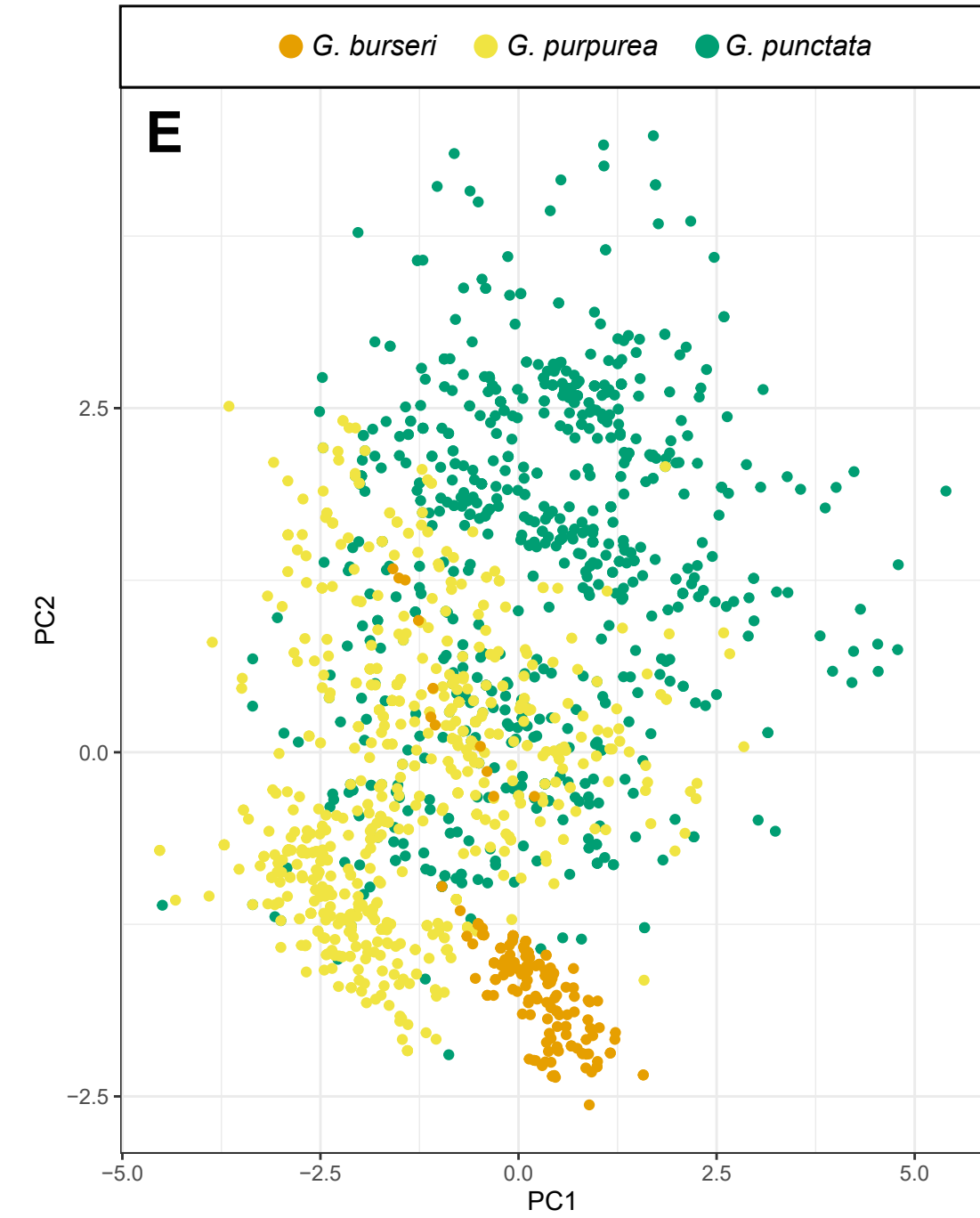
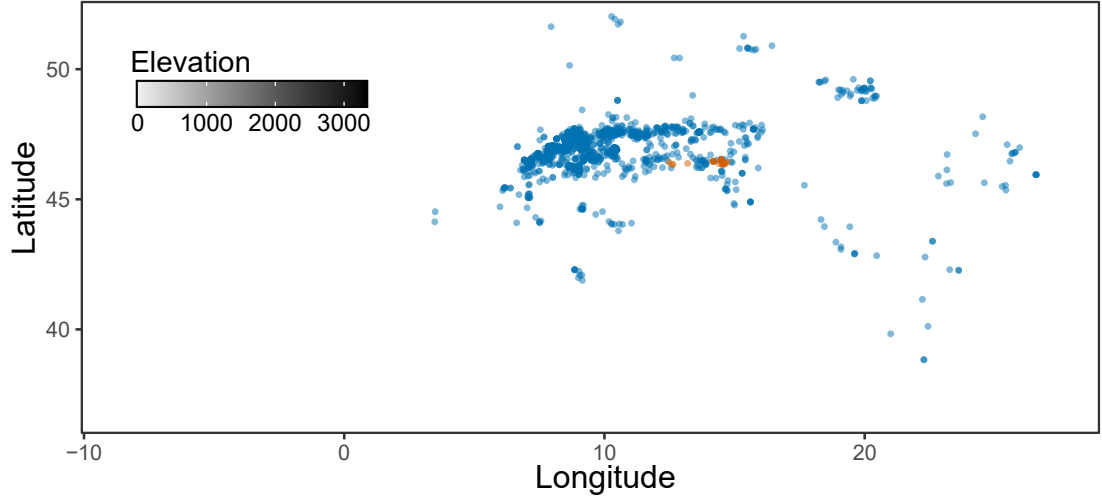


Phylogenetic relationships: rDNA-cistron tree

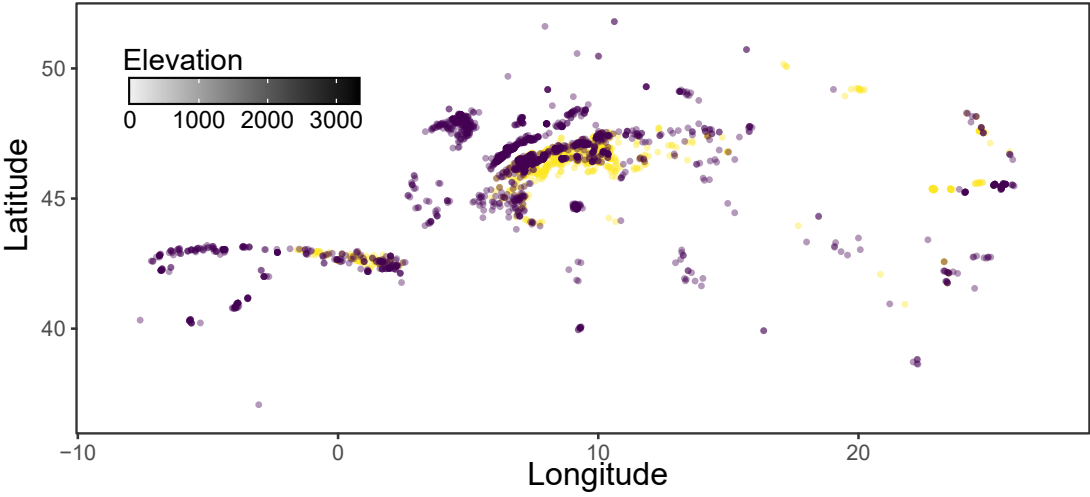
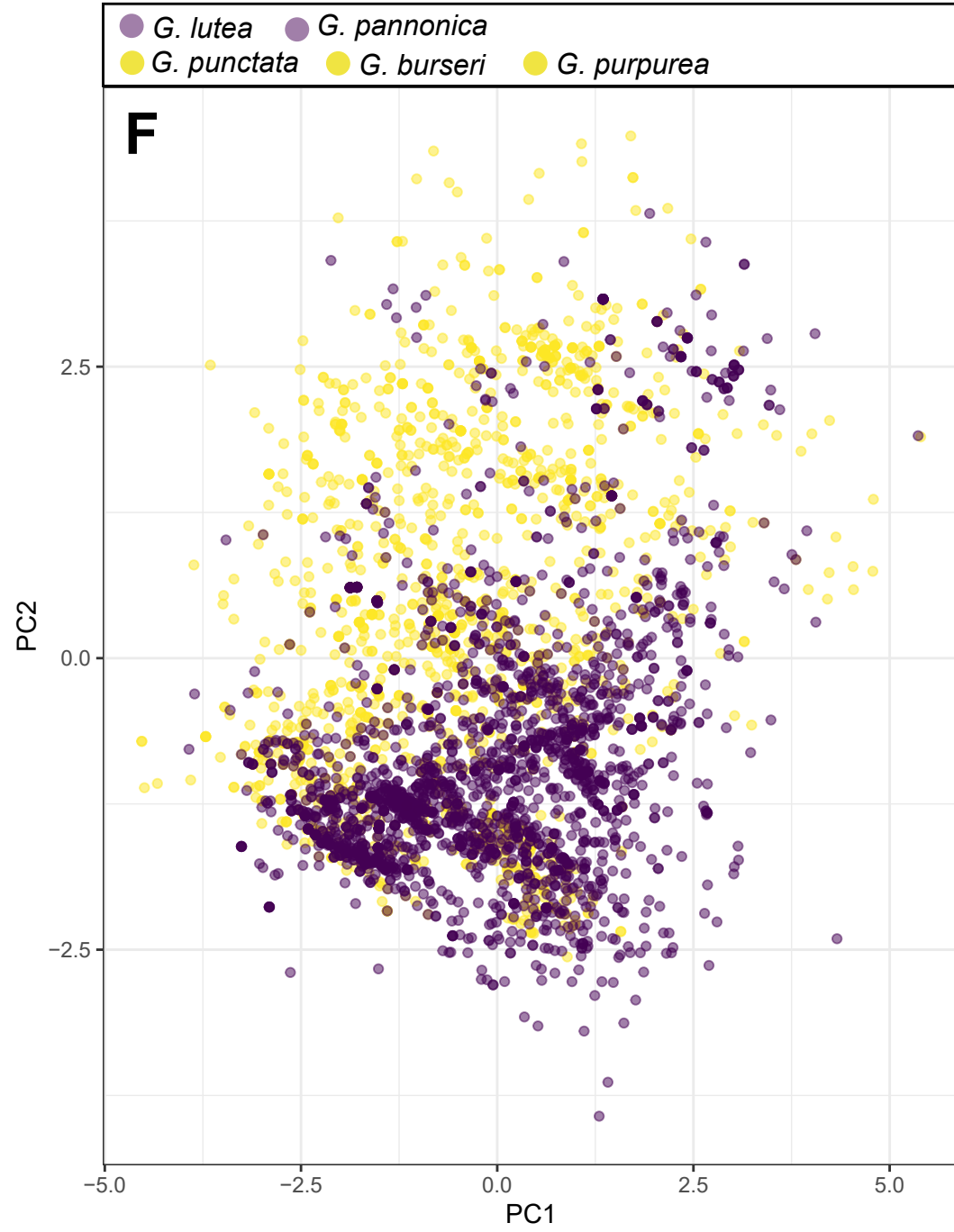
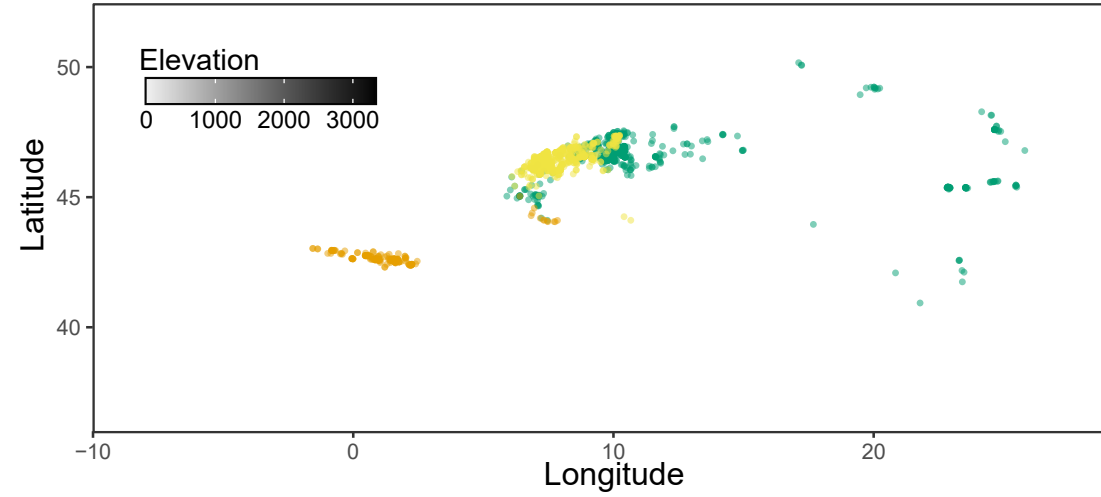
Comparison between calcicolous and silicolous species



Mean Geographic overlap:
G. froelichii vs. *G. asclepiadea* (0.56)



Mean Geographic overlap:
G. punctata vs. *G. burseri* (0.05), *G. purpurea* vs. *G. burseri* (0.00),
G. purpurea vs. *G. punctata* (0.43)



Species mean elevation:

G. asclepiadea (1162 +/- 543 m), *G. burseri* (1956 +/- 367 m), *G. froelichii* (1269 +/- 236 m), *G. lutea* (1277 +/- 531 m), *G. pannonica* (1273 +/- 299 m), *G. punctata* (1987 +/- 423 m), *G. purpurea* (1783 +/- 453 m)

***Gentiana froelichii*:** six additional locations were created for this species near known occurrences. These are listed below.

1. Slovenia, SE of summit of Visoki vrh, lat.: 46.3549111, long.: 14.5266000, elev.: 1956 m.
2. Slovenia, Between Planjava and Kroficka, lat.: 46.3559972, long.: 14.6155750, elev.: 2260 m
3. Slovenia, SW of summit of Velika Baba, lat.: 46.3603694, long.: 14.5520306, elev.: 2170 m.
4. Slovenia, E of summit of Mt Plece, lat.: 46.4350528, long.: 14.1801667, elev.: 2010 m
5. Slovenia, SE of Mt Vajnež, lat.: 46.4378861, long.: 14.1516111, elev.: 1980 m.
6. Slovenia, E of Mt Palec, lat.: 46.4395833, long.: 14.2230861, elev.: 1850 m.

Appendix Ve: Complement to Table 4, including all sister and closely related species comparison of climatic preferences, elevational ranges and geographic distribution ranges between and within three European *Gentiana* sections based on their occurrences in the European Alpine System. Extant climatic preferences of nine bioclimatic variables are summarized via the first two principal components (PC1, PC2). Significance levels: *<0.05, **<0.01, ***<0.001. Geographic overlap is given as the proportion of the smaller range contained in the larger range for each comparison. *Gentiana ligustica* was included in section and clade-wise comparisons (in *G. sect. Ciminalis* clade 1).

Section-wise comparison	Trees		Mean variable differences						Geo.
	rDNA	Plast.	PC1		PC2		Elev. (m)		overlap
<i>G. sect. Ciminalis</i> vs. <i>G. sect. Calathianae</i>	yes	yes	0.740	0.000	0.849	0.000	140.0	0.0	0.65
<i>G. sect. Gentiana</i> vs. <i>G. sect. Calathianae</i>	yes	yes	0.882	0.000	0.902	0.000	472.5	0.0	0.72
<i>G. sect. Gentiana</i> vs. <i>G. sect. Ciminalis</i>	yes	yes	0.141	0.000	0.053	0.153	332.5	0.0	0.68
Intrasectional deep splits									
<i>G. sect. Calathianae</i> : clade 1 vs. clade2	no	yes	1.195	0.000	0.683	0.000	419.2	4E-09	0.73
<i>G. sect. Calathianae</i> : clade 1 vs. clade3	no	no	2.180	0.000	0.274	0.008	530.8	4E-09	0.64
<i>G. sect. Calathianae</i> : clade 2 vs. clade3	no	no	0.985	0.000	0.409	0.000	111.6	3E-03	0.64
<i>G. sect. Calathianae</i> : clade 1 vs. (clade2, clade3)	no	yes	0.300	0.000	0.189	0.000	81.2	3E-05	0.79
<i>G. sect. Calathianae</i> : clade 1 vs. clade2 rDNA	yes	no	1.626	0.000	-0.848	0.000	479.8	3E-12	0.76
<i>G. sect. Calathianae</i> : clade 1 vs. clade3 rDNA	yes	no	1.576	0.000	-0.412	0.000	292.5	3E-12	0.68
<i>G. sect. Calathianae</i> : clade 2 vs. clade3 rDNA	yes	no	0.050	0.691	0.436	0.000	187.2	3E-12	0.72
<i>G. sect. Calathianae</i> : <i>G. terglouensis</i> vs. (clade 1, clade2) rDNA	yes	no	1.576	0.000	-0.412	0.000	292.5	3E-12	0.68
<i>G. sect. Calathianae</i> : clade 3 vs. (clade 1, clade2, <i>G. terglouensis</i>) rDNA	yes	no	0.050	0.691	0.436	0.000	187.2	3E-12	0.72
<i>G. sect. Gentiana</i> : clade 1 vs. clade2	no	yes	1.670	0.000	0.000	1.000	358.2	3E-08	0.49
<i>G. sect. Ciminalis</i> : clade 1 vs. clade2	no	yes	0.426	0.000	0.496	0.000	158.1	4E-09	0.58
Sister and closely related species relationships, recent splits									
<i>G. sect. Calathianae</i>									
<i>G. terglouensis</i> vs. <i>G. bavarica</i>	no	yes	0.706	0.911	0.275	1.000	448.5	4E-03	0.27
<i>G. rostanii</i> vs. <i>G. brachyphylla</i>	no	yes	0.951	0.067	1.603	0.000	172.4	9E-01	0.41
<i>G. verna</i> vs. <i>G. schleicheri</i>	yes	yes	2.949	0.000	2.538	0.000	1062.4	3E-08	1
<i>G. pumila</i> vs. <i>G. nivalis</i>	yes	yes	1.022	0.000	0.449	0.391	694.0	3E-08	0.32
<i>G. utriculosa</i> vs. <i>G. nivalis</i>	yes	yes	2.076	0.000	0.107	1.000	455.1	3E-08	0.48
<i>G. utriculosa</i> vs. <i>G. pumila</i>	yes	yes	1.054	0.000	0.556	0.155	238.9	8E-02	0.20
<i>G. bavarica</i> vs. <i>G. brachyphylla</i>	yes	no	0.366	0.081	4.109	0.000	233.7	3E-08	0.69
<i>G. rostanii</i> vs. <i>G. schleicheri</i>	yes	no	1.782	0.000	2.454	0.000	487.2	1E-06	0.07
<i>G. verna</i> vs. <i>G. rostanii</i>	yes	no	1.167	0.001	0.084	1.000	575.2	3E-08	0.59

<i>G. utriculosa</i> vs. (<i>G. pumila</i> , <i>G. nivalis</i>)	yes	yes	2.009	0.000	0.136	0.185	-410.0	0.000	0.52
<i>G. terglouensis</i> vs. (<i>G. bavarica</i> , <i>G. brachyphylla</i> , <i>G. rostanii</i> , <i>G. schleicheri</i> , <i>G. verna</i>)	yes	no	-0.364	0.377	0.925	0.008	-163.8	0.218	0.73
<i>G. rostanii</i> vs. (<i>G. verna</i> , <i>G. schleicheri</i>)	yes	no	-0.842	0.009	-0.196	0.469	458.2	0.000	0.59
mean cpDNA sister species relationships			1.527		1.000		393.460		0.50
mean cpDNA closely related species			1.538		0.809		380.184		0.46
mean rDNA sister species relationships			0.711		0.922		211.190		0.56
mean rDNA closely related species			1.122		1.116		363.095		0.52
<i>G. sect. Gentiana</i>									
<i>G. lutea</i> vs. <i>G. burseri</i>	yes	yes	0.304	0.736	0.753	0.000	679.0	3.18E-08	0.68
<i>G. pannonica</i> vs. <i>G. burseri</i>	yes	no	0.534	0.832	1.913	0.000	682.7	3.18E-08	0
<i>G. punctata</i> vs. <i>G. burseri</i>	yes	yes	0.069	1.000	2.979	0.000	31.1	1.000	0.05
<i>G. purpurea</i> vs. <i>G. burseri</i>	yes	no	1.578	0.000	1.397	0.000	172.7	0.010	0
<i>G. pannonica</i> vs. <i>G. lutea</i>	yes	no	0.839	0.009	1.160	0.000	3.7	1	0.23
<i>G. punctata</i> vs. <i>G. lutea</i>	yes	yes	0.374	0.000	2.227	0.000	710.1	3E-08	0.50
<i>G. purpurea</i> vs. <i>G. lutea</i>	yes	no	1.274	0.000	0.644	0.000	506.2	3E-08	0.64
<i>G. purpurea</i> vs. <i>G. pannonica</i>	yes	yes	2.112	0.000	0.516	0.206	510.0	3E-08	0.02
<i>G. purpurea</i> vs. <i>G. punctata</i>	yes	no	1.647	0.000	1.583	0.000	203.8	3E-08	0.43
<i>G. froelichii</i> vs. <i>G. asclepiadea</i>	yes	yes	1.964	0.000	0.299	1.000	107.4	1E+00	0.56
<i>G. froelichii</i> vs. all other <i>Gentiana</i>	yes	no	-0.903	0.009	0.297	0.265	-118.4	0.338	0.56
<i>G. asclepiadea</i> vs. (<i>G. burseri</i> , <i>G. punctata</i> , <i>G. purpurea</i> , <i>G. lutea</i> , <i>G. pannonica</i>)	yes	no	1.689	0.000	-0.002	0.940	-359.3	0.000	0.49
<i>G. pannonica</i> vs. (<i>G. burseri</i> , <i>G. punctata</i> , <i>G. purpurea</i> , <i>G. lutea</i>)	yes	no	0.976	0.000	0.674	0.000	-251.8	0.001	0.26
<i>G. lutea</i> vs. (<i>G. burseri</i> , <i>G. punctata</i> , <i>G. purpurea</i>)	yes	no	0.949	0.000	-0.356	0.000	-541.8	0.000	0.53
<i>G. purpurea</i> vs. (<i>G. burseri</i> , <i>G. punctata</i>)	yes	no	-1.634	0.000	-0.999	0.000	-197.7	0.000	0.43
mean cpDNA sister species relationships			0.965		1.355		407.496		0.36
mean rDNA sister species relationships			0.191		0.432		-239.653		0.39
mean rDNA closely related species			0.575		0.836		75.810		0.42
<i>G. sect. Ciminalis</i>									
<i>G. angustifolia</i> vs. <i>G. acaulis</i>	yes	yes	0.625	0.000	2.557	0.000	815.9	3E-08	0.28
<i>G. occidentalis</i> vs. <i>G. acaulis</i>	yes	yes	0.727	0.000	2.600	0.000	780.2	3E-08	0.19
<i>G. occidentalis</i> vs. <i>G. angustifolia</i>	yes	yes	0.101	1.000	0.043	1.000	35.6	1E+00	0.70
<i>G. clusii</i> vs. <i>G. alpina</i>	yes	yes	0.492	0.000	1.616	0.000	519.1	3E-08	0.25
<i>G. alpina</i> vs. <i>G. dinarica</i>	no	yes	0.100	1.000	3.414	0.000	606.3	6E-06	0
<i>G. clusii</i> vs. <i>G. dinarica</i>	no	yes	0.592	0.992	1.798	0.000	87.2	1E+00	0
<i>G. occidentalis</i> vs. <i>G. dinarica</i>	yes	no	1.434	0.015	4.152	0.000	475.3	5E-03	0

<i>G. angustifolia</i> vs. <i>G. dinarica</i>	yes	no	1.333	0.047	4.109	0.000	511.0	2E-03	0
<i>G. dinarica</i> vs. <i>G. acaulis</i>	yes	no	-0.707	0.933	1.552	0.000	-304.9	4E-01	0
<i>G. clusii</i> vs. all other <i>Ciminalis</i>	yes	no	0.017	0.963	-0.584	0.000	-118.4	0.338	0.62
<i>G. alpina</i> vs. (<i>G. acaulis</i> , <i>G. occidentalis</i> , <i>G. dinarica</i> , <i>G. angustifolia</i>)	yes	no	0.713	0.000	1.448	0.000	-359.3	0.000	0.69
<i>G. angustifolia</i> vs. (<i>G. acaulis</i> , <i>G. occidentalis</i> , <i>G. dinarica</i>)	yes	no	-0.562	0.000	2.321	0.000	-251.8	0.001	0.64
<i>G. acaulis</i> vs. (<i>G. occidentalis</i> , <i>G. dinarica</i>)	yes	no	0.586	0.000	-2.192	0.000	-541.8	0.000	0.15
<i>G. alpina</i> vs. (<i>G. clusii</i> , <i>G. dinarica</i>)	no	yes	-0.480	0.000	-1.651	0.000	520.8	0.000	0.25
<i>G. angustifolia</i> vs. (<i>G. occidentalis</i> , <i>G. acaulis</i>)	no	yes	0.554	0.000	-2.302	0.000	-739.4	0.000	0.64
<i>G. dinarica</i> vs. (<i>G. acaulis</i> , <i>G. occidentalis</i>)	yes	no	-0.903	0.009	0.297	0.265	-118.4	0.338	0
<i>G. acaulis</i> vs. (<i>G. occidentalis</i> , <i>G. angustifolia</i>)	no	yes	-0.680	0.000	2.580	0.000	796.7	0.000	0.25
mean cpDNA sister species relationships			0.348		0.111		162.210		0.27
mean cpDNA closely related species			0.339		1.009		328.218		0.29
mean rDNA sister species relationships			-0.002		1.216		-13.518		0.43
mean rDNA closely related species			0.297		1.828		180.399		0.31

Appendix Vf: Complement to Figure 4. Comparisons of rates of niche evolution. Maximum a posteriori density values for estimated evolutionary rates (σ^2) for eleven selected WorldClim variables and the first two principal components of each of the three studied European sections of *Gentiana*. Based on the rDNA and plastid topologies, including *G. ligustica* or not, rates were modelled separately for each section under a Brownian motion model in R.

Variables	Sections	PLASTID TREES						rDNA TREES					
		<i>C. ligustica</i> added		Tree as is		500 trees		<i>C. ligustica</i> added		Tree as is		500 trees	
		value	rescaled	value	rescaled	value	rescaled	value	rescaled	value	rescaled	value	rescaled
Bio 1	<i>Ciminalis</i>	2.4939	1.0000	2.8461	1.0000	2.5940	1.0000	1.8219	1.0000	2.0546	1.0000	2.2100	1.0000
	<i>Gentiana</i>	0.1928	0.0000	0.1928	0.0000	0.1680	0.0000	0.1891	0.0000	0.1891	0.0000	0.1920	0.0000
	<i>Calathianae</i>	1.1898	0.4333	1.1898	0.3758	1.1550	0.4068	0.8979	0.4341	0.8979	0.3799	0.7180	0.2607
Bio 2	<i>Ciminalis</i>	0.4560	1.0000	0.2676	1.0000	0.2300	1.0000	0.4433	1.0000	0.4747	1.0000	0.4040	1.0000
	<i>Gentiana</i>	0.1594	0.0360	0.1594	0.0929	0.1610	0.2333	0.1870	0.2091	0.1870	0.1907	0.1850	0.2935
	<i>Calathianae</i>	0.1483	0.0000	0.1483	0.0000	0.1400	0.0000	0.1192	0.0000	0.1192	0.0000	0.0940	0.0000
Bio 3	<i>Ciminalis</i>	5.1917	1.0000	3.7408	1.0000	4.0530	1.0000	6.0551	1.0000	6.8484	1.0000	6.3470	1.0000
	<i>Gentiana</i>	0.3800	0.0000	0.3800	0.0000	0.3880	0.0000	0.4432	0.0000	0.4432	0.0000	0.4390	0.0000
	<i>Calathianae</i>	0.8022	0.0877	0.8022	0.1256	0.7480	0.0982	0.6804	0.0423	0.6804	0.0370	0.5280	0.0151
Bio 4	<i>Ciminalis</i>	270.5626	1.0000	296.4035	1.0000	324.1240	1.0000	366.4502	1.0000	425.2503	1.0000	522.2760	1.0000
	<i>Gentiana</i>	131.5559	0.1313	131.5559	0.1131	142.2890	0.0890	123.6492	0.1709	123.6492	0.1423	135.7220	0.1306
	<i>Calathianae</i>	110.5387	0.0000	110.5387	0.0000	124.5190	0.0000	73.6055	0.0000	73.6055	0.0000	77.6590	0.0000
Bio 8	<i>Ciminalis</i>	0.8424	0.0000	0.9600	0.0000	0.9090	0.0000	1.0147	0.0000	1.1617	0.0000	1.2630	0.0000
	<i>Gentiana</i>	5.5602	1.0000	5.5602	1.0000	5.7280	1.0000	2.4110	0.4387	2.4110	0.4115	2.3580	0.4917
	<i>Calathianae</i>	4.9439	0.8694	4.9439	0.8660	4.8630	0.8205	4.1976	1.0000	4.1976	1.0000	3.4900	1.0000
Bio 9	<i>Ciminalis</i>	18.1092	1.0000	19.2236	1.0000	18.9120	1.0000	14.4631	1.0000	15.9143	1.0000	17.8250	1.0000
	<i>Gentiana</i>	8.8143	0.3551	8.8143	0.3296	8.9010	0.3348	5.9566	0.3217	5.9566	0.2883	6.3490	0.2716
	<i>Calathianae</i>	3.6968	0.0000	3.6968	0.0000	3.8620	0.0000	1.9221	0.0000	1.9221	0.0000	2.0710	0.0000
Bio 12	<i>Ciminalis</i>	5991.2704	0.4679	6110.3555	0.4869	7120.2080	0.5026	5332.4266	0.5244	5622.7601	0.5784	7389.0080	0.7284
	<i>Gentiana</i>	3058.8923	0.0000	3058.8923	0.0000	2961.5800	0.0000	2513.8904	0.0000	2513.8904	0.0000	2877.3870	0.0000
	<i>Calathianae</i>	9326.1587	1.0000	9326.1587	1.0000	11235.4290	1.0000	7888.6580	1.0000	7888.6580	1.0000	9071.5930	1.0000
Bio 15	<i>Ciminalis</i>	1.7851	0.0000	0.2433	0.0000	0.3030	0.0000	0.9883	0.0000	0.4153	0.0000	0.3260	0.0000
	<i>Gentiana</i>	6.3177	0.5927	6.3177	0.6610	6.3690	0.6033	2.8896	0.3867	2.8896	0.4508	2.9590	0.5103

	<i>Calathianae</i>	9.4328	1.0000	9.4328	1.0000	10.3570	1.0000	5.9045	1.0000	5.9045	1.0000	5.4860	1.0000
Bio 18	<i>Ciminalis</i>	2363.0755	1.0000	2316.9456	0.9944	2244.4120	0.0000	1399.4515	0.1891	1473.6846	0.7778	1835.1030	1.0000
	<i>Gentiana</i>	2319.3337	0.9066	2319.3337	1.0000	2716.7710	1.0000	1375.6084	0.0000	1375.6084	0.0000	1574.7020	0.0000
	<i>Calathianae</i>	1894.7560	0.0000	1894.7560	0.0000	2544.0960	0.6344	1501.7102	1.0000	1501.7102	1.0000	1755.0640	0.6926
Bio 19	<i>Ciminalis</i>	473.7338	1.0000	520.4708	1.0000	531.7590	1.0000	645.0061	1.0000	731.4039	1.0000	941.2920	1.0000
	<i>Gentiana</i>	394.7454	0.3514	394.7454	0.2540	433.3870	0.2279	162.3840	0.0000	162.3840	0.0000	169.9360	0.0000
	<i>Calathianae</i>	351.9425	0.0000	351.9425	0.0000	404.3560	0.0000	258.8405	0.1999	258.8405	0.1695	281.7480	0.1450
Altitude	<i>Ciminalis</i>	49200.3223	1.0000	56398.2901	1.0000	69027.0730	1.0000	38067.5765	1.0000	44370.2258	1.0000	60313.2770	1.0000
	<i>Gentiana</i>	22853.5450	0.0000	22853.5450	0.0000	25632.0780	0.0000	10875.7903	0.0000	10875.7903	0.0000	12451.6440	0.0000
	<i>Calathianae</i>	43195.5885	0.7721	43195.5885	0.6064	54351.3640	0.6618	36215.2792	0.9319	36215.2792	0.7565	43658.8310	0.6520
PC1	<i>Ciminalis</i>	0.0962	0.0000	0.0501	0.0000	0.0430	0.0000	0.0880	0.0000	0.0973	0.0000	0.0860	0.0000
	<i>Gentiana</i>	0.2691	0.7623	0.2691	0.8024	0.2710	0.8507	0.1735	0.4832	0.1735	0.4545	0.1820	0.7164
	<i>Calathianae</i>	0.3230	1.0000	0.3230	1.0000	0.3110	1.0000	0.2649	1.0000	0.2649	1.0000	0.2200	1.0000
PC2	<i>Ciminalis</i>	0.6508	1.0000	0.6795	1.0000	0.6290	1.0000	0.7263	1.0000	0.8409	1.0000	0.8120	1.0000
	<i>Gentiana</i>	0.1315	0.0000	0.1315	0.0000	0.1310	0.0000	0.1108	0.0000	0.1108	0.0000	0.1160	0.0000
	<i>Calathianae</i>	0.2470	0.2224	0.2470	0.2108	0.2430	0.2249	0.1828	0.1170	0.1828	0.0987	0.1440	0.0402