

## 1 SUPPLEMENTAL MATERIALS

2

### 3 TEXT S1

4 The original duckweed family Lemnaceae (Hegelmaier 1895) was shown to be monophyletic  
5 with the family Araceae (Cabrera *et al.* 2008; Cusimano *et al.* 2011; Nauheimer, Metzler & Renner  
6 2012). Although the Angiosperm Phylogeny III suggested therefore to include duckweeds as a  
7 subfamily of Araceae (Lemnoideae), most duckweed researcher consider them as a separate family,  
8 Lemnaceae (Appenroth, Borisjuk & Lam 2013; Appenroth & Crawford 2015; Sree, Bog & Appenroth  
9 2016).

10

### 11 TEXT S2

#### 12 Material and Methods

#### 13 Establishment of a database of fluorescence emission spectra for major *S. polyrhiza* flavonoids - 14 details

15 To test for the linearity of the fluorescence and to compare the fluorescence intensities among  
16 the individual compounds, standard curves of fluorescence intensities of each individual synthetic  
17 flavonoid were recorded on cLSM as described in the main manuscript for final concentrations between  
18 0 and 2.5 mM in DMSO after 1:1 addition of 0.25 % 2-APB solution. To investigate the effect of pH on  
19 the emission spectra, we further recorded the spectra of individual 2-APB conjugated compounds under  
20 varying pH conditions using potassium phosphate buffer with a pH of 5.7, 7.3 and 8. To test for the  
21 effect of DMSO and 2-APB concentration on the flavonoid emission spectra, we additionally recorded  
22 the spectra of compounds dissolved in a lower DMSO concentration (25 vs 5 % of solution drop) as  
23 well as a lower 2-APB concentration (0.25 % vs 0.13 % of stock solution). As the absolute intensity of  
24 a 2-APB stained flavonoid glucoside of a given concentration in a drop of standard solution largely  
25 depended on the focus plane, we standardized the measurements by choosing the focus at the maximal

26 intensity within a defined area of each image (fourth quadrant of each image) for all *in vitro* studies. For  
27 *in vivo* recordings chloroplasts were chosen as the focus plane as these were easily detectable in the  
28 chlorophyll channel and therefore independent of investigating flavonoids within the plant body. Due  
29 to technical limitations the extraction of spectra of a homogenous flavonoid glucoside mixed standard  
30 solution is not possible: The laser excites more than one molecule (fluorophore of flavonoid glucoside/2-  
31 APB) at a time resulting in a “mixed” signal. However, once the spectrum of each individual compound  
32 solution is extracted and stored in the database, channels can be assigned in parallel pixel by pixel *in*  
33 *vivo*. Our chosen cLSM settings of 21 % transmission, 625 gain, pinhole 39  $\mu\text{m}$  were chosen based on  
34 preliminary investigations observing the range of *in vivo* fluorescence intensities of images under  
35 highest (copper treatment) and lowest (AIP -- PAL-inhibitor) treatment.

36

### 37 **TEXT S3**

#### 38 **Material and Methods**

#### 39 **Visualization and quantification of secondary fluorescence across tissue layers of the *S. polyrhiza*** 40 **frond - statistics**

41 To test whether individual compounds are distributed differentially across tissue layers, we  
42 used linear mixed model analysis with tissue layer as fixed effect and the individual frond as random  
43 effect. *P*-values were obtained by likelihood ratio tests that compare the full model with the reduced  
44 model in which tissue layer was removed as a fixed effect. Pairwise comparisons were obtained with  
45 Tukey’s posthoc test.

46

47 **TEXT S4**48 **Material and Methods**49 **Verification of cLSM-based flavonoid visualization *in vivo* – details**

50 Three fronds of very similar appearance (each a single mature mother frond with one emerging  
51 daughter frond attached to it) were placed into 250 mL plastic bowls filled with 180 mL full nutrient  
52 medium without and with 2  $\mu$ M AIP (n=12). After four days, a single mature plant of the second  
53 generation (granddaughters) of half of the replicates of each treatment was used for cLSM analysis  
54 utilizing the ‘online fingerprinting’ feature as described above. The granddaughter generation was used  
55 as these plants fully developed under the specific conditions and as the plants displayed stable flavonoid  
56 levels at this age (see below). The granddaughter fronds of the remaining replicates of each treatment  
57 were briefly dried, weighed and immediately frozen in liquid nitrogen to determine flavonoid  
58 concentration using high pressure liquid chromatography (HPLC). Plant material for HPLC analysis  
59 was stored at -20 °C until extraction. For flavonoid extraction, frozen plant tissue was ground by  
60 vigorously shaking the tubes with three metal beads for 1 min in a paint shaker. Fifty mg powdered  
61 tissue was then extracted with 1 ml 100% methanol (or adjusted volume if less than 50 mg was available)  
62 by vortexing for 8 s. Samples were centrifuged for 10 min at room temperature at 17,000 g. The  
63 supernatant was stored at -20 °C prior to HPLC analysis.

64 Methanol extracts were analysed by HPLC 1100 series equipment (Agilent Technologies)  
65 coupled to a photodiode array detector (G1315A DAD, Agilent Technologies). Analyte separation was  
66 accomplished with a Nucleodur Sphinx RP column (250 4.6 mm, 5  $\mu$ m particle size, Macherey-Nagel).  
67 Injection volume was 10  $\mu$ l. The mobile phase consisted of 0.2 % formic acid in water (solvent A) and  
68 acetonitrile (solvent B) at a flow rate of 1 ml min<sup>-1</sup> using the following gradient: 0 min 10 % (B), 8 min  
69 21 % (B), 17.0 min 55 % (B), 17.1 min 100 % (B), 18.0 min 100 % (B), 18.1 min 10 % (B), 22 min 10  
70 % (B). Peak areas of the four major flavonoids, lut 8-C-glc, lut 7-O-glc, ap 8-C-glc and ap 7-O-glc, were  
71 integrated at 330 nm and quantified based on external standards of lut 7-O-glc and adjusted for the

72 molecular weight of the individual flavonoids (factors for luteolins and apigenins were 1 and 0.96,  
73 respectively).

74 To test the effect of AIP on flavonoid concentrations, we compared mean flavonoid levels  
75 between control and AIP treated plants using Student's *t*-test. Fluorescence intensities were extracted  
76 with ImageJ as described above. To test whether individual compounds were distributed differentially  
77 across tissue layers of fronds grown in the presence of AIP, we used linear mixed model analysis with  
78 tissue layer as fixed effect and the individual frond as random effect. *P*-values were obtained by  
79 likelihood ratio tests that compare the full model with the reduced model in which tissue layer was  
80 removed as fixed effect. Pairwise comparisons were obtained with Tukey's posthoc test. To test whether  
81 AIP had tissue layer effects on flavonoid reduction, we used linear mixed model analysis with tissue  
82 layer, treatment and their interaction as fixed effects and individual frond as random effect for each  
83 metabolite separately. *P*-values were obtained by likelihood ratio tests that compare the full model with  
84 the model in which the interaction of tissue layer and treatment was removed (tissue layer specific effects  
85 of AIP). Wilcoxon Mann Whitney Test were then used to test for differences in individual flavonoid  
86 glucoside fluorescence intensities between treatments within each tissue layer.

87

## 88 **TEXT S5**

### 89 **Material and Methods**

#### 90 **Tissue-specific induction of flavonoids under copper sulphate treatment – statistics**

91 To test the effect of copper sulphate addition on biomass production, we compared mean  
92 biomass of fronds grown in the absence and presence of copper sulphate addition using Student's *t*-test.  
93 To test the effect of copper sulphate addition on flavonoid concentrations, we compared mean flavonoid  
94 levels between control and copper-induced plants using Student's *t*-test. To test for differences in the  
95 magnitude of induction across the four major flavonoids, we divided the metabolite concentration of  
96 each copper-treated plant by the mean of the control plants for each flavonoid individually (relative  
97 induction). Differences in the relative induction across the four flavonoids were analysed using one-way

98 ANOVA. Pairwise comparisons were performed with Tukey's posthoc test. To test whether copper  
99 exposure had tissue layer specific effects on flavonoid induction, we used linear mixed model analysis  
100 of fluorescence intensity data with tissue layer, treatment and their interaction as fixed effects and  
101 individual frond as random effects for each metabolite separately. *P*-values were obtained by likelihood  
102 ratio tests that compare the full model with the model in which the interaction of tissue layer and  
103 treatment was removed (tissue layer specific effects of copper treatment). Wilcoxon-Mann-Whitney-  
104 Tests were then used to test for differences in fluorescence intensities between treatments within each  
105 tissue layer.

106

107

## 108 **TEXT S6**

### 109 **Material and Methods**

#### 110 **Induction of flavonoids under natural UV light – statistics**

111 To test the effect of UV radiation on biomass production, we compared biomass between fronds  
112 grown in the absence and presence of UV light using Student's *t*-test. To test the effect of UV radiation  
113 on flavonoid concentrations, we compared mean flavonoid levels between control and UV treated plants  
114 using Student's *t*-test. To test for differences in the magnitude of induction across the four major  
115 flavonoids, we divided the metabolite concentrations of the UV-treated plants by the mean of the control  
116 plants for each flavonoid individually (relative change). Differences in the relative change across the  
117 four flavonoids were analysed using one-way ANOVA. Pairwise comparisons were performed with  
118 Tukey's posthoc test. To test whether UV radiation had tissue layer-specific effects on flavonoid  
119 induction, we used linear mixed model analysis of fluorescence intensity data with tissue layer, treatment  
120 and their interaction as fixed effects and individual frond as random effects for each metabolite  
121 separately. *P*-values were obtained by likelihood ratio tests that compared the full model with the model  
122 in which the interaction of tissue layer and treatment was removed (tissue layer specific effects of UV

123 stress). Wilcoxon-Mann-Whitney-Tests were then used to test for differences in fluorescence intensities  
124 between treatments within each tissue layer.

125

## 126 **TEXT S7**

### 127 **Material and Methods**

#### 128 **Fitness assay of genotypes exposed to copper sulphate -statistics:**

129 Differences in biomass production between copper sulphate-treated and control plants were  
130 analysed with a paired Student's *t*-test using the mean value of each genotype and treatment. Biomass  
131 production is an excellent predictor for plant fitness in this almost exclusively vegetative reproducing  
132 plant and incorporates both the effect of resistance (reduced damage) and tolerance (growth despite  
133 damage). The correlations among the individual flavonoid concentrations were analysed using Pearson's  
134 moment correlations using the mean values of each clone under control conditions. *P*-values of  
135 Pearson's correlation coefficients were obtained using the Hmisc package (Harrell Jr & Dupont 2015).  
136 To assess the correlation between plant fitness and flavonoid concentrations, we normalized growth  
137 variation among genotypes by expressing the mean biomass accumulation of copper-exposed plants  
138 relative to the mean biomass production under control conditions of each genotype ("relative fitness").  
139 The correlations between relative fitness and constitutive as well as induced flavonoid levels were  
140 analysed with linear models for each metabolite individually as well as for total flavonoid concentration  
141 (sum of the four major flavonoids). To assess the relation between absolute growth and flavonoid  
142 concentrations, the correlations between total fresh mass and individual as well as total flavonoid  
143 concentration in the presence and absence of CuSO<sub>4</sub> were analysed using linear models based on the  
144 mean value of each treatment and genotype.

145

146

147 **TEXT S8**

148 **Material and Methods**

149 **Fitness assay of genotypes exposed to ambient UV light – statistics:**

150           Difference in biomass production between UV-shielded and -exposed conditions was analysed  
151 by a paired Student's *t*-test using the mean value of each genotype and treatment. Analogous to the  
152 correlations of plant fitness and flavonoid concentrations under copper stress, we standardized the  
153 growth between genotypes by expressing the mean biomass accumulation under UV-exposed conditions  
154 relative to the mean biomass production under UV-shielded conditions of each genotype ("relative  
155 fitness"). Correlations between relative fitness and constitutive as well as induced flavonoid levels were  
156 analysed with linear models for each metabolite separately as well as for total flavonoid concentration.  
157 To assess the relation between absolute growth and flavonoid concentrations, the correlations between  
158 total fresh mass and individual as well as total flavonoid concentrations under UV-exposed and -shielded  
159 conditions were analysed using linear models based on the mean value of each treatment and genotype.  
160 The correlation between relative fitness under UV light and under copper sulphate exposure was  
161 analysed with a linear model.

162

163

164 **TABLE S1** *Spirodela polyrhiza* accessions used for fitness experiments (1 = used; 0 = not used).

Accession Number	UV experiment	copper experiment	Continent	Country
8756	1	1	Africa	Ethiopia
8683	1	1	Africa	Kenya
9510	1	1	Africa	Mozambique
9907	1	1	Asia	Bangladesh
9925	1	0	Asia	Bangladesh
0040	1	1	Asia	China
9636	1	1	Asia	China
5523	1	0	Asia	China
0109	1	0	Asia	China
0092	1	1	Asia	China
5521	1	1	Asia	China
0090	0	1	Asia	China
0225	0	1	Asia	China
9668	0	1	Asia	China
9503	1	1	Asia	India
7379	1	1	Asia	India
8442	1	1	Asia	India
9650	1	1	Asia	India
9295	1	1	Asia	India
9305	0	1	Asia	India
9290	0	1	Asia	India
8787	0	1	Asia	Nepal
7674	1	1	Asia	Nepal
9507	1	1	Asia	Russia
9511	1	1	Asia	Russia
9512	0	1	Asia	Russia
9351	1	1	Asia	Vietnam
0013	0	1	Asia	Vietnam
7551	1	1	Australia	Australia
9625	1	1	Europe	Albania
9628	1	1	Europe	Albania
9633	1	0	Europe	Albania
9629	1	0	Europe	Albania
9514	1	1	Europe	Austria
9513	0	1	Europe	Czech Republic
9256	0	1	Europe	Finland
8403	0	1	Europe	France
9509	1	1	Europe	Germany
9500	1	1	Europe	Germany
5513	0	1	Europe	Germany
5501	0	1	Europe	Hungary
9502	1	1	Europe	Ireland
9618	1	1	Europe	Italy
9413	0	1	Europe	Italy
9609	1	1	Europe	Poland
9508	0	1	Europe	Poland
9607	0	1	Europe	Switzerland
8790	1	1	North America	Canada
9505	1	1	North America	Cuba
6613	1	1	North America	USA



8118	1	1	North America	USA
7960	1	1	North America	USA
7498	1	1	North America	USA
6731	0	1	North America	USA
8409	0	1	North America	USA
7003	0	1	North America	USA
7657	0	1	Central America	Mexico
9242	1	1	South America	Ecuador
165				

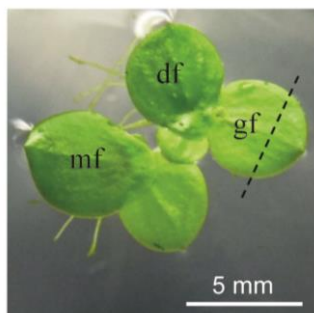
166

167 **TABLE S2** Weather data for outdoor experiment. Radiation data display the cumulative sum of each  
 168 day. Data were provided from a weather station 5 km distant, and were provided by the Thuringian State  
 169 Institute for Environment and Geography, Jena, Germany.

		Average	Global	PAR-			
	date	air temperature	radiation	radiation	UV-A	UV-B	
		[°C]	[kWh/m <sup>2</sup> ]	[kWh/m <sup>2</sup> ]	[Wh/m <sup>2</sup> ]	[Wh/m <sup>2</sup> ]	
genotype 7498	experiment with	19/07/2017	24	6.8	1.2	263.1	8
		20/07/2017	20.9	4.1	0.8	168.5	5.1
		21/07/2017	22.2	6.1	1.1	239.7	7.2
		22/07/2017	20.8	4.2	0.8	180.4	5.7
		23/07/2017	20	5	0.9	202.7	6.1
		24/07/2017	15.4	1.8	0.4	85.6	2.6
		25/07/2017	14.6	2.2	0.5	102.7	2.9
genotypes for fitness assay	experiment with world-wide distributed	05/09/2016	16.6	2.1	0.4	178.8	3
		06/09/2016	17.9	3.1	0.6	220.7	3.7
		07/09/2016	17.6	5.1	0.9	341.2	5.5
		08/09/2016	20.2	5.2	0.9	339.7	5.5
		09/09/2016	20.9	4.4	0.8	293.2	4.8
		10/09/2016	20.5	4.5	0.8	295.7	4.8
		11/09/2016	21.5	4.1	0.7	275.7	4.5
		12/09/2016	22.6	4.7	0.8	306.4	4.9
		13/09/2016	22.9	4.7	0.8	303	4.8
		14/09/2016	22.2	4.4	0.7	278.3	4.4
		15/09/2016	21.3	4.7	0.8	299.2	4.7
		16/09/2016	18.1	2.2	0.4	158.3	2.5
17/09/2016	15.5	0.3	0.1	37.9	0.6		
18/09/2016	14.9	1.1	0.2	93.4	1.6		
19/09/2016	13.7	1.7	0.3	136.9	2.2		

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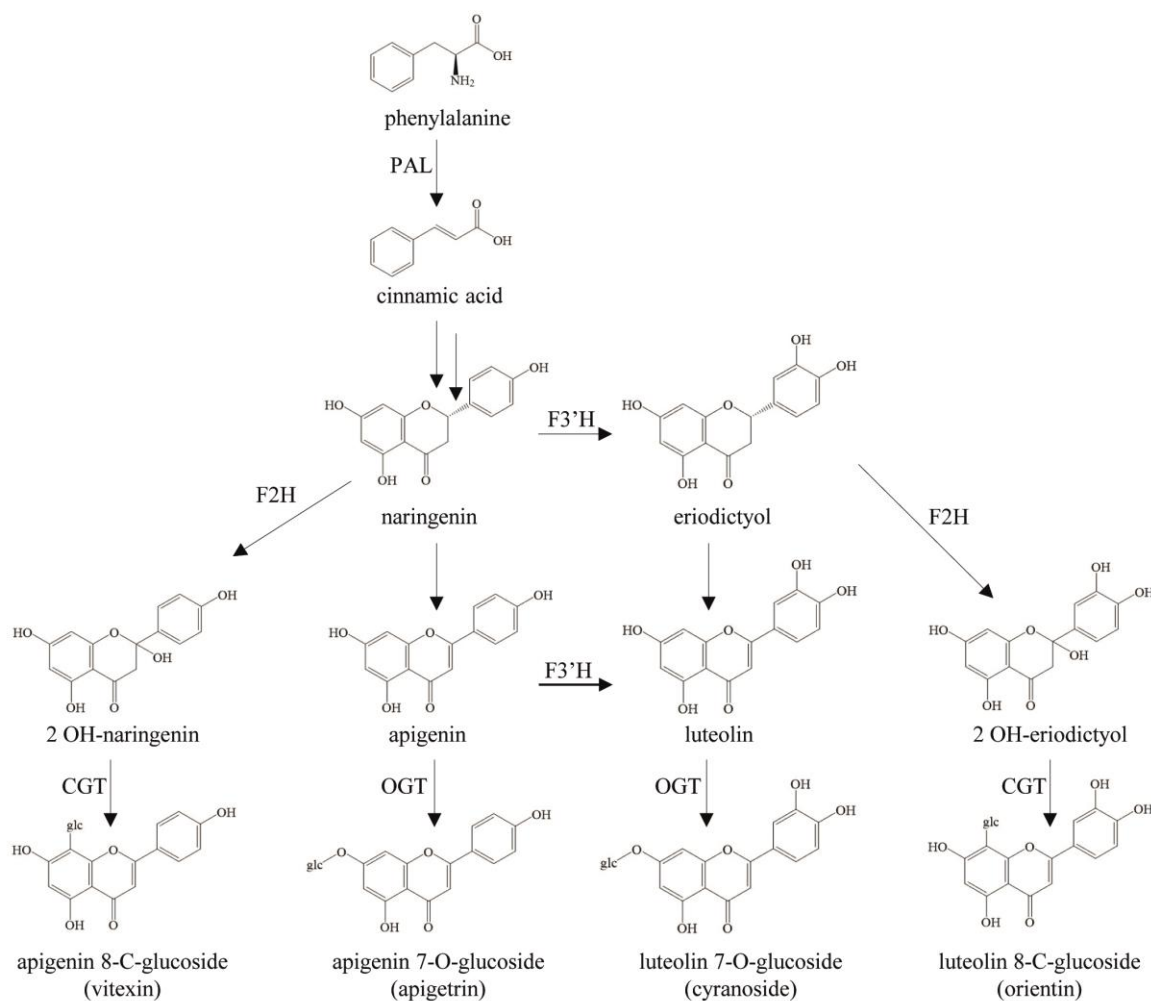
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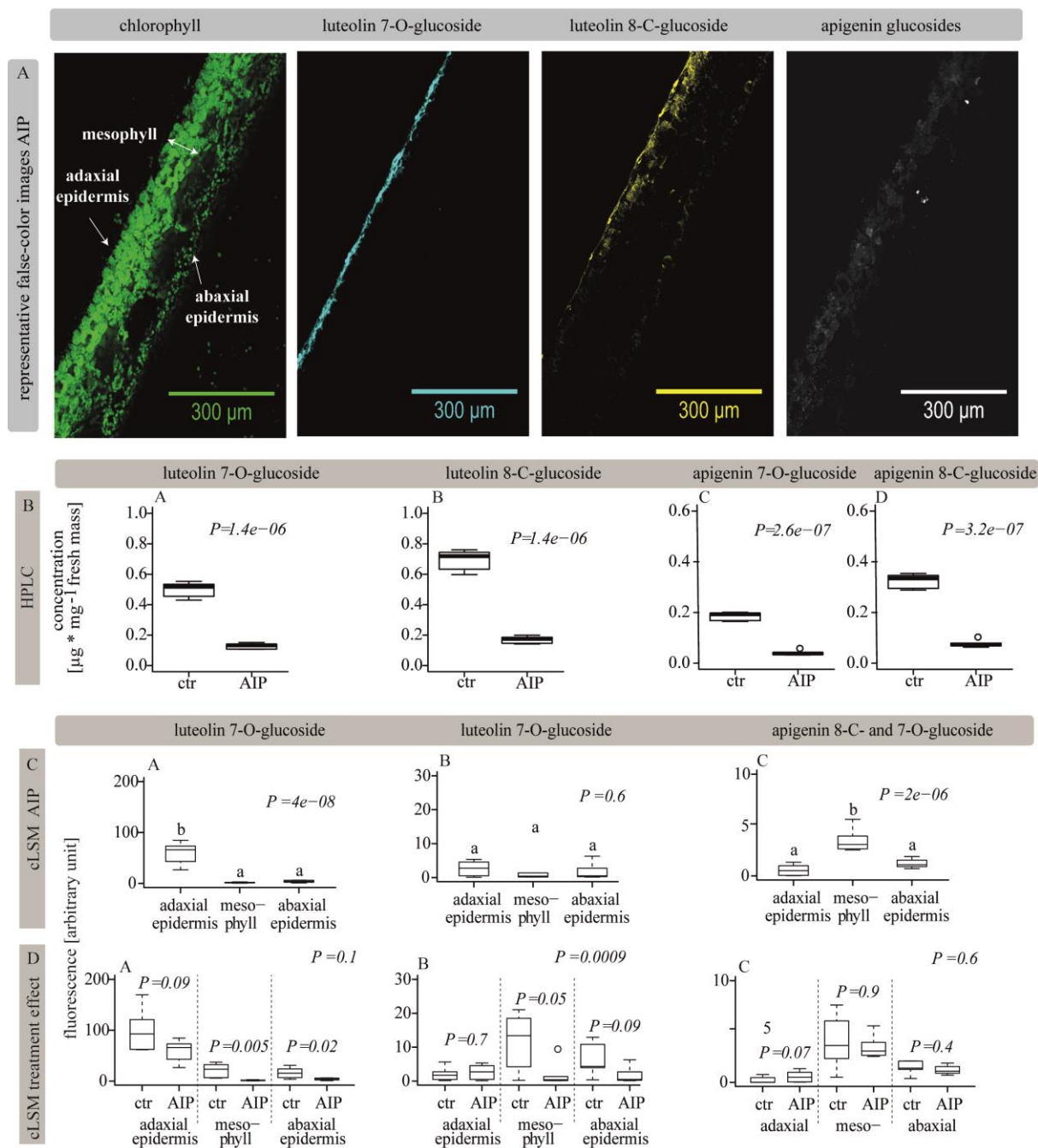
173 **FIGURE S1** *S. polyrhiza* mother fronds (mf), first daughter frond (df) and first granddaughter frond  
174 (gf). Line marks axis where gf frond was hand cross-sectioned to stain and visualize flavonoid  
175 glucosides.

176



177

178 **FIGURE S2** Scheme of the putative biosynthetic pathway of the major flavonoids in *Spirodela*  
 179 *polyrhiza* adapted from *Arabidopsis thaliana* and *Zea mays* (Li, Bonawitz, Weng & Chapple 2010;  
 180 Casas, Duarte, Doseff & Grotewold 2014; Nugroho, Choi & Park 2016). Double arrows represent  
 181 several steps. Dotted arrow represents less favoured pathway. PAL = phenylalanine ammonia lyase.  
 182 F2H = flavanone-2-hydroxylase. F3'H = flavonoid 3'-hydroxylase. CGT = C-glycosyl transferase. OGT  
 183 = O-glycosyl transferase



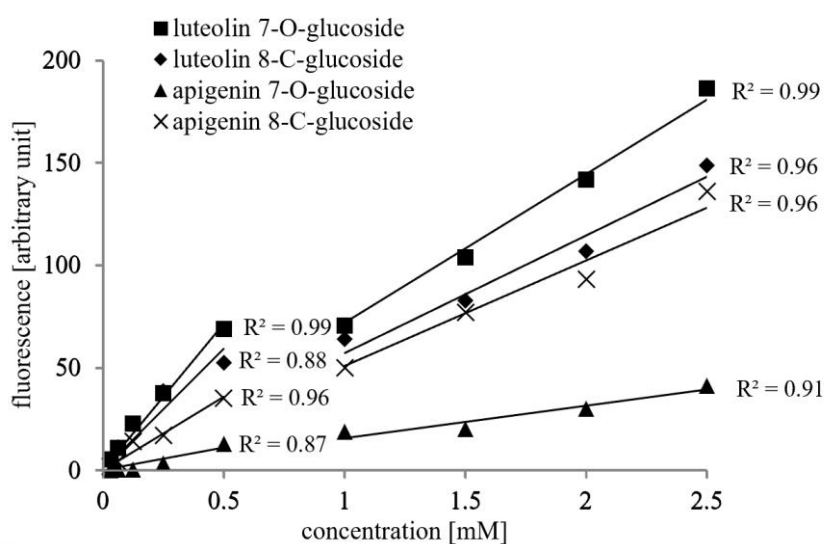
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185 **FIGURE S3** (A) cLSM false-color-images of 2-APB-stained *Spirodela polyrhiza* cross-sections of  
 186 plants growing in the presence of the phenylalanine ammonia-lyase (PAL)-inhibitor 2-aminoidane-2-  
 187 phosphonic acid (AIP). Brightness and contrast were adapted in ImageJ for demonstrative reasons. (B)  
 188 Whole plant flavonoid glucoside content was reduced after exposure to the PAL-inhibitor AIP. Mean  
 189 levels of control and AIP exposed plants were analysed with Student's *t*-test. The second generation

190 (granddaughters) of plants that had been subjected to the individual condition was analysed. (C) Tissue  
 191 distribution of individual flavonoid glucosides via fluorescence-based quantification was altered by AIP.  
 192 Statistics of likelihood ratio tests comparing two linear mixed effect models that differ in the factor  
 193 tissue layer as a fixed effect are shown. Different lower-case letters indicate significant differences in  
 194 fluorescence intensity according to Tukey's HSD test. (D) Comparison of tissue distribution of  
 195 individual flavonoid glucosides (fluorescence-based quantification) by AIP treatment and controls. AIP  
 196 had tissue-specific effects on flavonoid accumulation particularly on luteolin levels. *P*-values of  
 197 likelihood ratio tests comparing two linear mixed effect models that differed in the interaction term of  
 198 treatment and tissue layer are displayed above each figure. Wilcoxon-Mann-Whitney-Tests were then  
 199 used to test for differences of individual flavonoid fluorescence intensities between treatments within a  
 200 tissue layer. Plants were grown in the absence and presence of AIP until the granddaughter generation  
 201 had matured (day 6). ctr = control. AIP = 2-aminoidane-2-phosphonic acid. *n* = 6.

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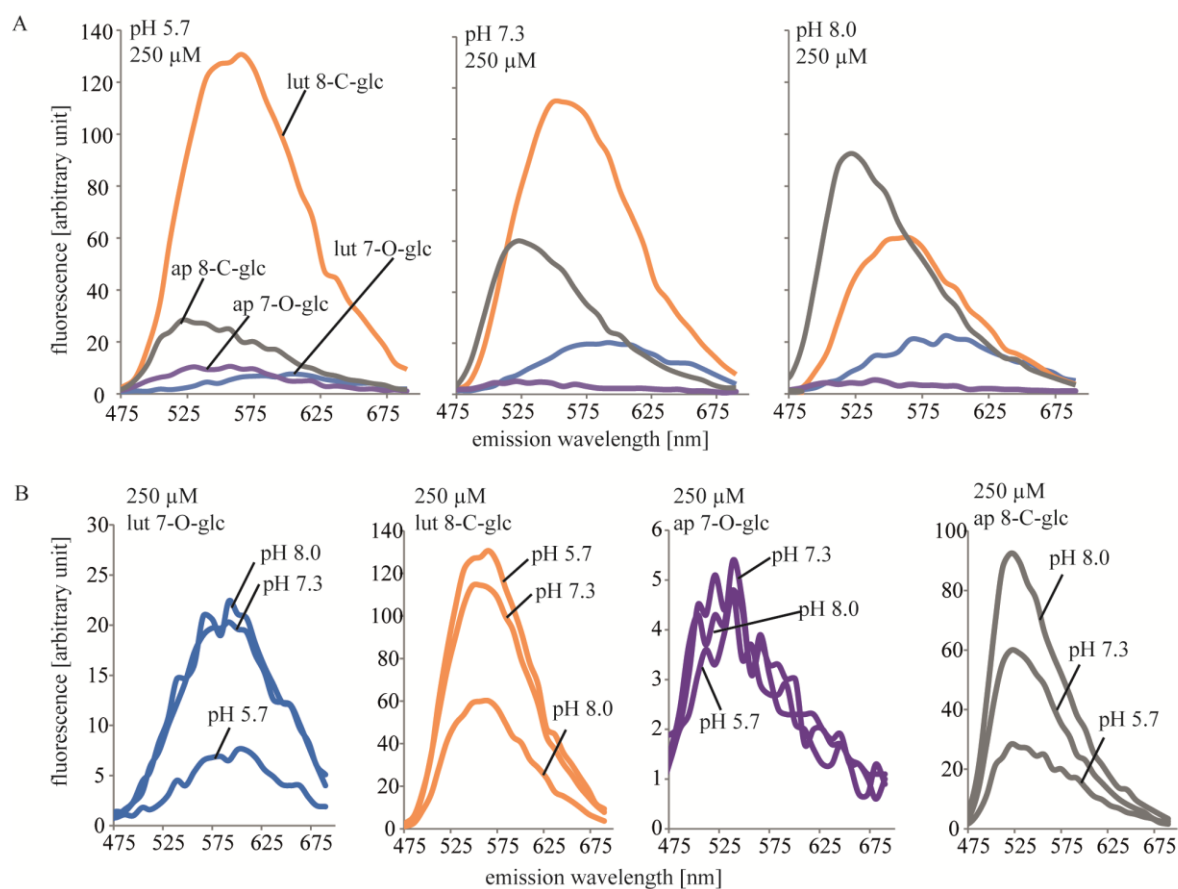


204

205 **FIGURE S4** Fluorescence intensities of individual synthetic flavonoid glucosides for two concentration  
 206 ranges when stained with 2-APB (2-aminoethyl diphenylborinate). Concentration levels refer to final  
 207 flavonoid glucoside concentrations after 1:1 addition of 0.25 % 2-APB solution. Linear regressions of  
 208 two concentration ranges were applied due to non-linearity of the correlations.

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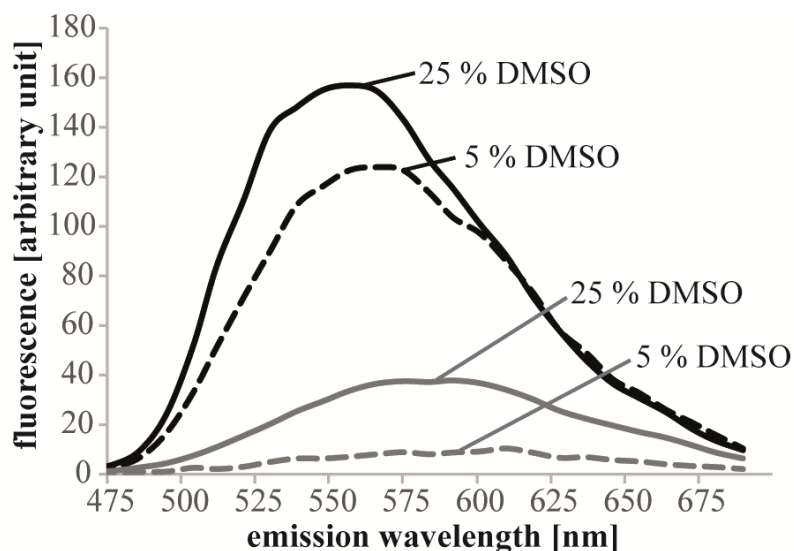


211

212 **FIGURE S5** Effect on pH on fluorescence emission spectra of individual synthetic flavonoid  
 213 glucosides. Individual compounds were dissolved in phosphate buffer of pH 5.7, 7.3 and 8.0 and stained  
 214 with 2-APB (2-aminoethyl diphenylborinate) (1:1) with a final flavonoid glucoside concentration of 250  
 215 μM. (A) Comparison of defined pH value among compounds. (B) Comparison of compounds among  
 216 different pH values. Intensities of apigenin 7-O-glc likely underrepresent true intensities due to  
 217 formation of crystals at acidic conditions.

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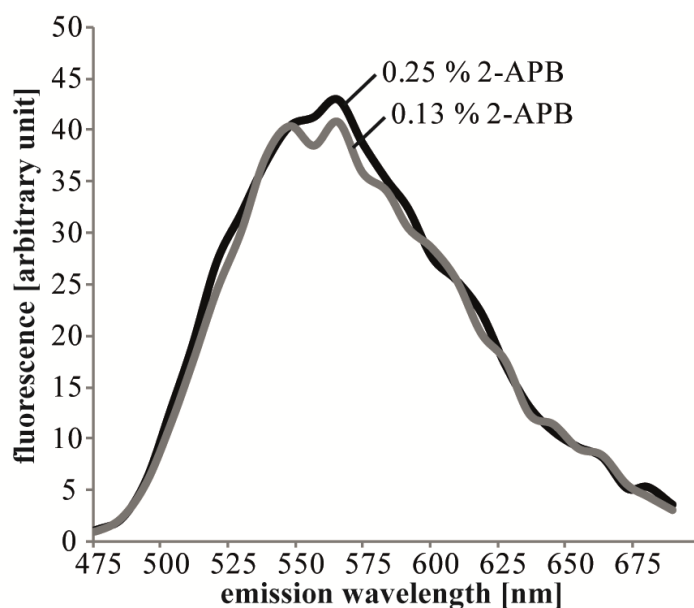
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221 **FIGURE S6** Effect of DMSO concentration on fluorescence emission spectra of luteolin glucosides.

222 Luteolin 8-C-glucoside (black) and luteolin 7-O-glucoside (grey) were dissolved in 25% or 5% DMSO

223 concentration in phosphate buffer (pH=7.3) at a final flavonoid concentration of 500  $\mu$ M.

224



225

226 **FIGURE S7** Effect of 2-APB (2-aminoethyl diphenylborinate) on fluorescence emission spectra of

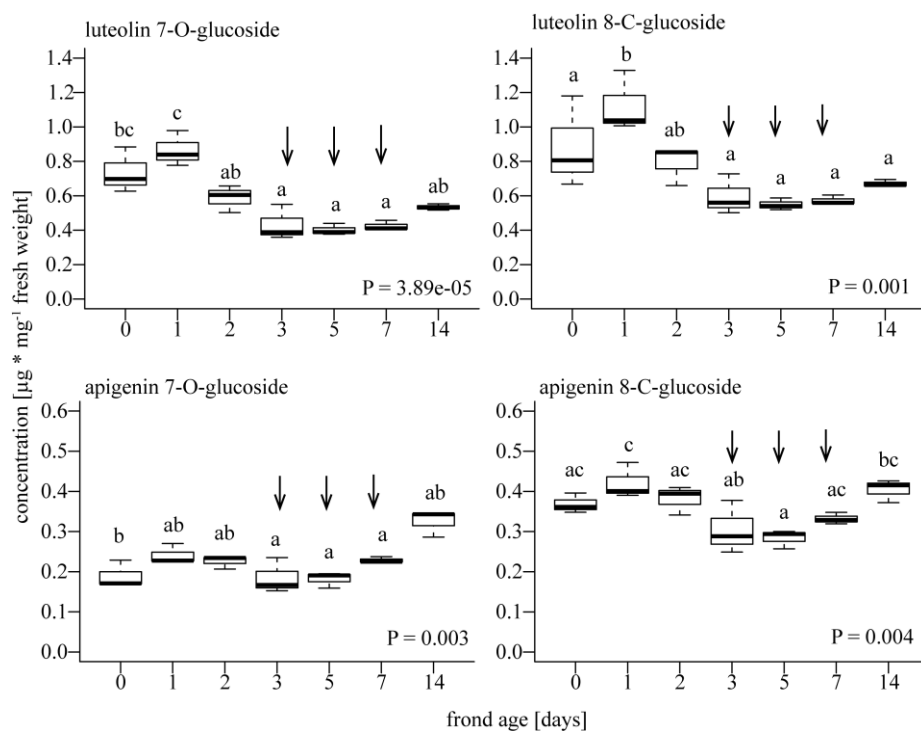
227 luteolin 8-C glucoside (100  $\mu$ M, pH 7.3 in phosphate buffer) when stained 1:1 with either 0.25 % or

228 0.13 % 2-APB solution.

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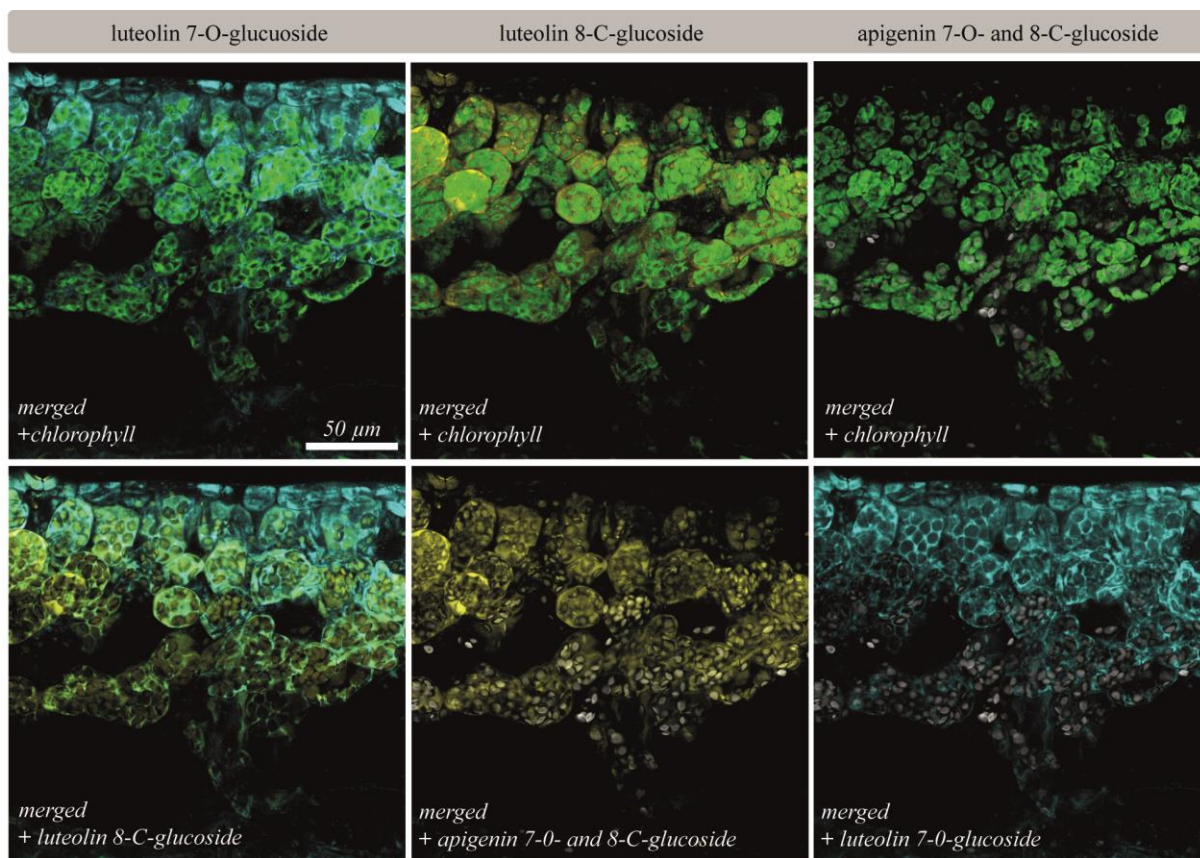


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232 **FIGURE S8** Individual flavonoid concentrations of *S. polyrhiza* fronds over frond lifespan. All  
 233 following experiments were performed with plants between age 3-7 days when flavonoid content was  
 234 most stable (indicated by arrows). Data were analysed with one-way ANOVAs. Different lower case  
 235 letters indicate significant differences according to Tukey's post hoc tests.  $n=3$ .

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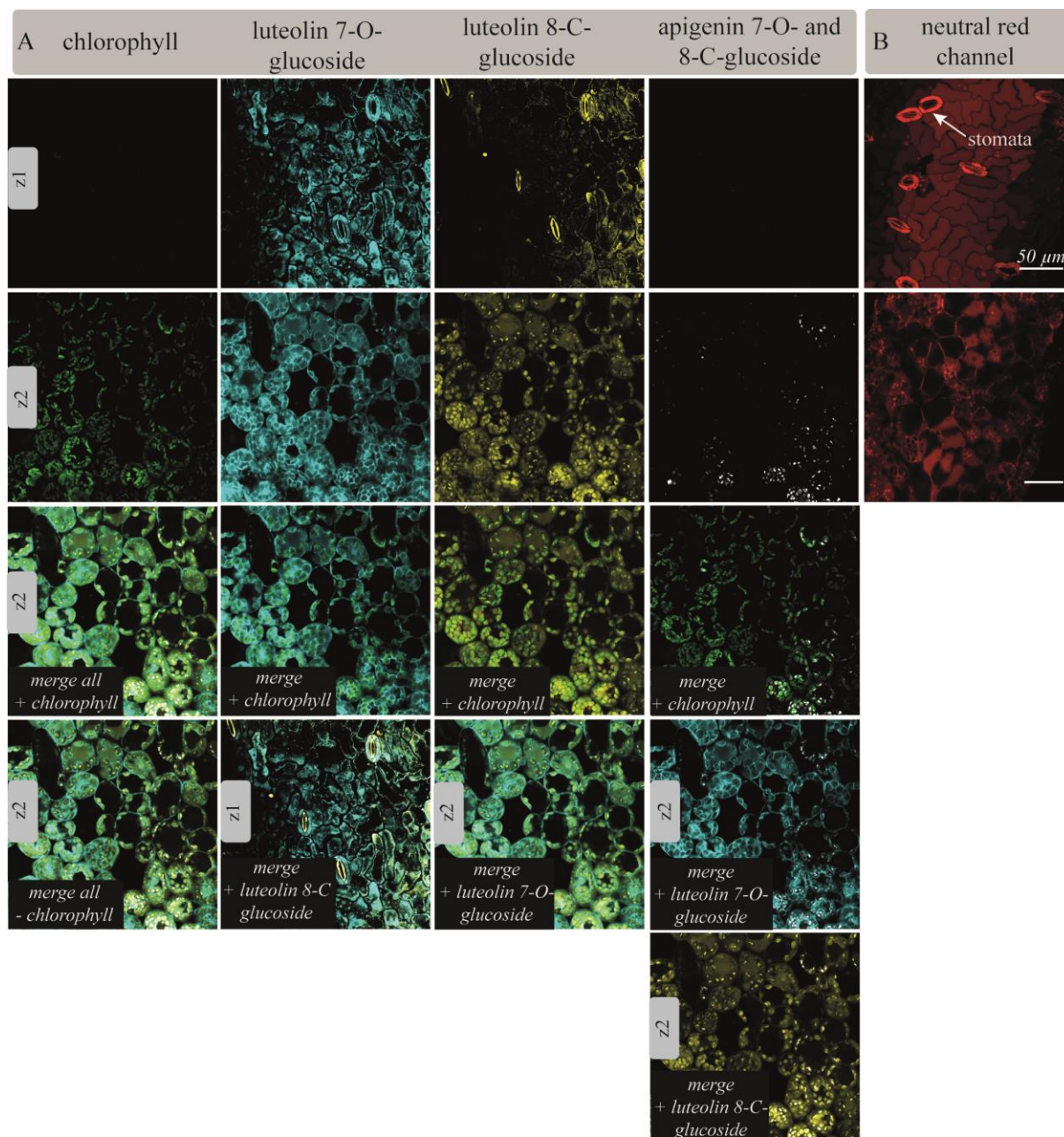


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239 **FIGURE S9** Corresponding merged channels of images of Figure 3. Subcellular distribution of  
 240 individual flavonoid glucosides based on cLSM false-colour-images of a 2-APB stained *Spirodela*  
 241 *polyrhiza* cross-section. Brightness and contrast of all images were adapted in ImageJ for illustrative  
 242 purpose. 2-APB = 2-aminoethyl diphenylborinate.

243

244



245

246 **FIGURE S10** cLSM false-colour-images of DBPA and neutral red-stained *Spirodela polyrhiza* fronds.

247 View on the adaxial epidermis at two different depths (z1- top, z2- deeper) to show the subcellular

248 distribution of individual flavonoid glucosides. Both luteolin glucoside channels showed fluorescence

249 comparable with the neutral red (vacuolar) channel. Luteolin 8-C-glucoside additionally showed overlay

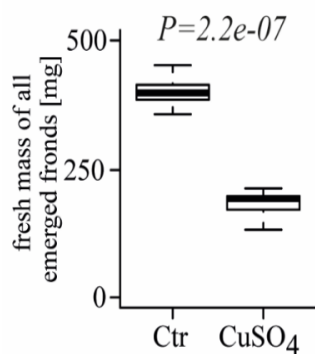
250 with the chlorophyll (chloroplast) channel. Apigenins mostly accumulated in deeper cell layers. cLSM

251 was performed with the ‘online fingerprinting’ feature. 2-APB-stained cross-sections were excited at

252 405 nm; in another cross-section neutral red-stained sections were excited at 543 nm. Brightness and

253 contrast of all images were adapted individually in ImageJ for demonstrative reasons. 2-APB = 2-  
254 aminoethyl diphenylborinate. Scalebars correspond to 50  $\mu\text{m}$ .

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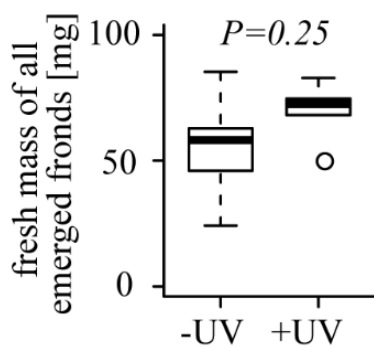


256

257 **FIGURE S11** Reduction of fresh weight of fronds grown in the presence of copper sulphate (*S.*  
258 *polyrhiza* genotype 7498). Mean weights of control and copper sulphate exposed plants were analysed  
259 with Student's *t*-test. n = 6. ctr = control

260

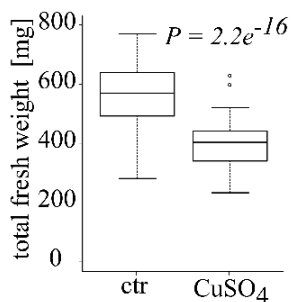
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263 **FIGURE S12** Fresh weight of fronds was not affected by UV treatment (*S. polyrhiza* genotype 7498).  
264 Mean weights of control and UV exposed plants were analysed with Student's *t*-test. n = 5.

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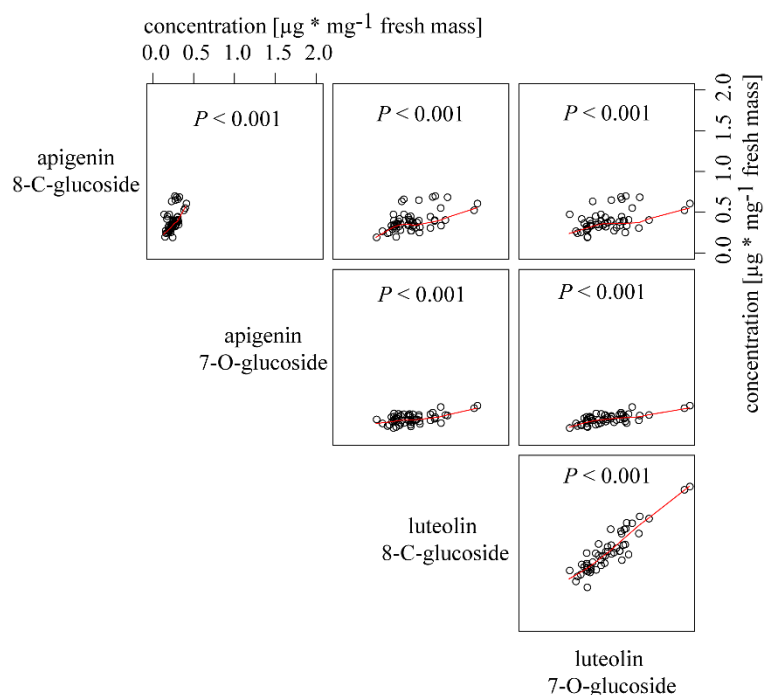


266

267 **FIGURE S13** Reduction of fresh weight of fronds grown in the presence of copper sulphate in 53 *S.*  
 268 *polyrhiza* genotypes. Mean weights of control and copper exposed plants (n=3) were analysed with  
 269 Student's *t*-test. ctr = control

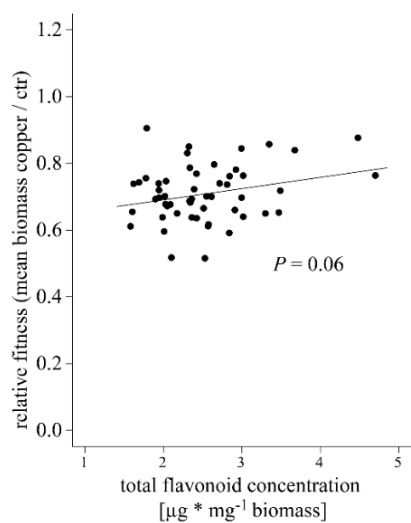
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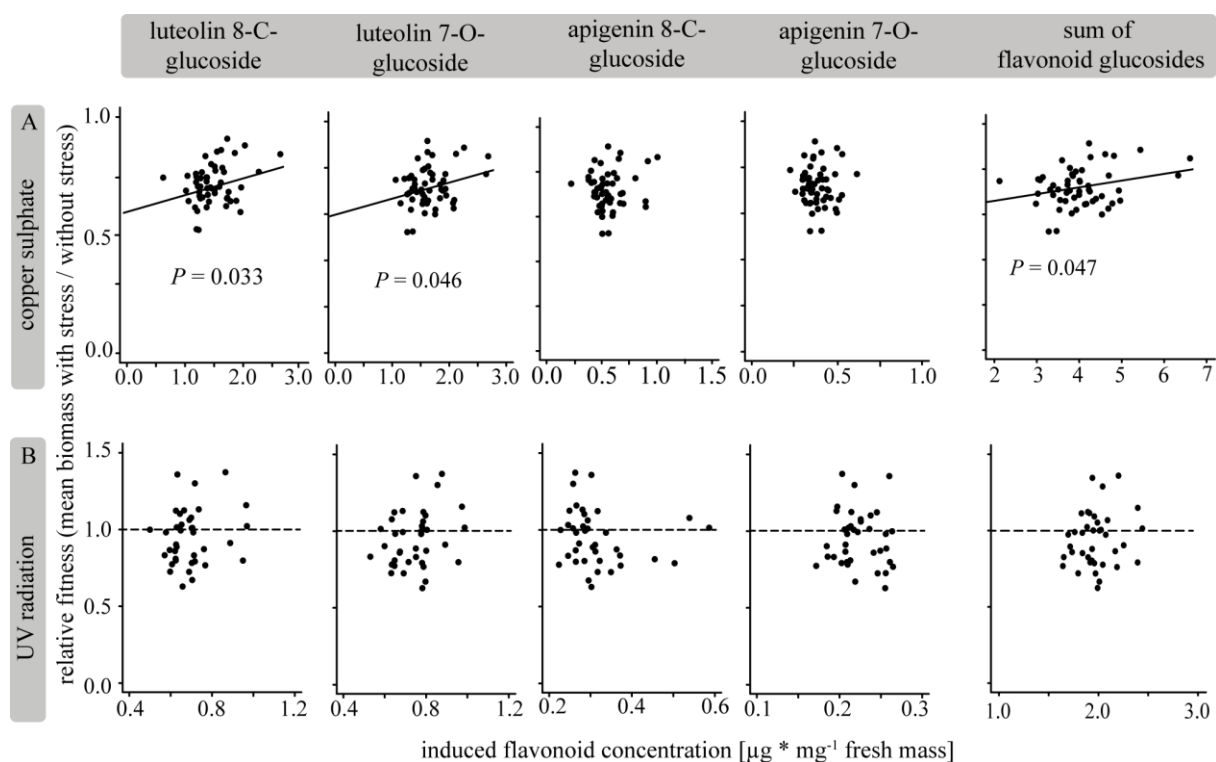
273 **FIGURE S14** Concentrations of individual flavonoid glucosides were highly correlated across 53 *S.*  
 274 *polyrhiza* genotypes exposed to copper sulphate. *P*-values of Pearson's correlation coefficients are  
 275 shown. Each data point indicates the mean flavonoid concentration of one clone growing under control  
 276 conditions.



277

278 **FIGURE S15** Total flavonoid concentration was weakly correlated to relative plant fitness across 53 *S.*  
 279 *polyrhiza* genotypes under copper sulphate treatment. *P*-value of linear model is shown. ctr = control

280



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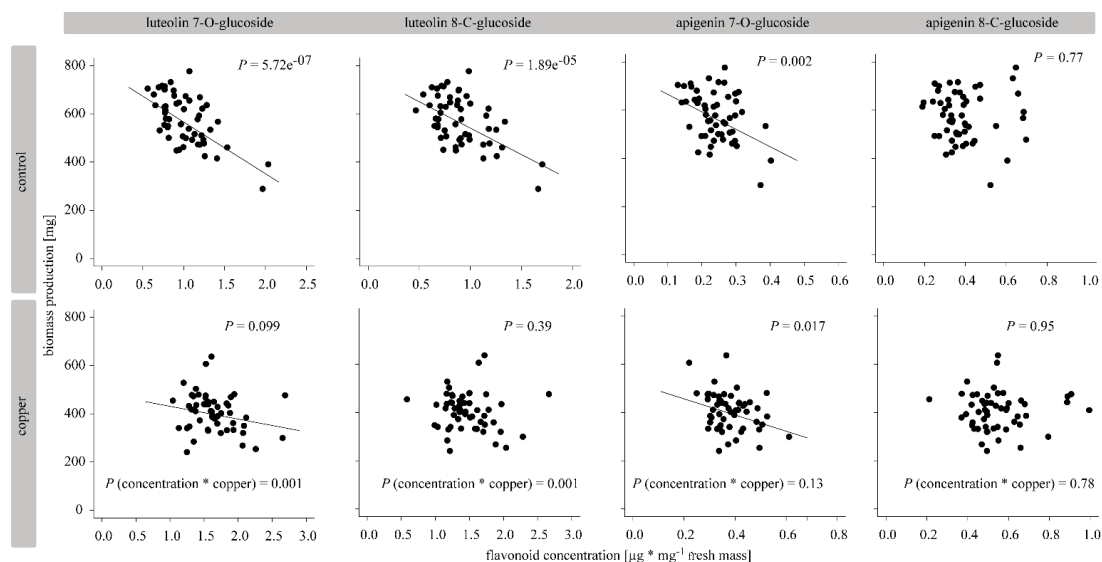
282 **FIGURE S16** Correlation of relative fitness (mean biomass accumulation with stress / without stress)  
 283 and induced flavonoid concentration across of 53 *S. polyrhiza* genotypes under (A) copper sulphate and



284 (B) UV treatment. Each data point represents the mean of one genotype. *P*-values of linear models based  
 285 on mean values per genotype are shown.

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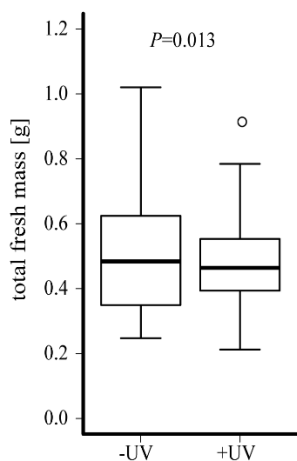


288

289 **FIGURE S17** Correlation of biomass production and individual flavonoid concentration across of 53 *S.*  
 290 *polyrhiza* genotypes under control and copper sulphate treatment. The correlations between biomass  
 291 production and individual as well as total flavonoid concentration in the presence and absence of  $\text{CuSO}_4$   
 292 were analysed using linear models based on the mean value of each treatment and genotype.

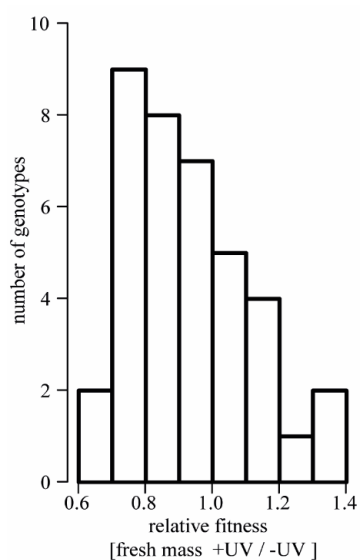
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296 **FIGURE S18** Total fresh mass of fronds across 38 *S. polyrhiza* genotypes was reduced by an average of  
 297 8% in the presence of natural UV radiation. Mean weight of UV shielded (-UV) and UV exposed (+UV)  
 298 plants were analysed with Student's *t*-test.

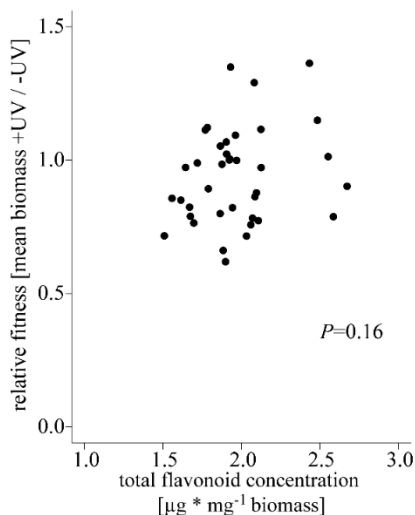


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300 **FIGURE S19** Distribution of 38 *S. polyrhiza* genotypes depending on the relative fitness under UV  
 301 exposure. The majority of genotypes showed reduced fitness but 30% showed increased fitness.

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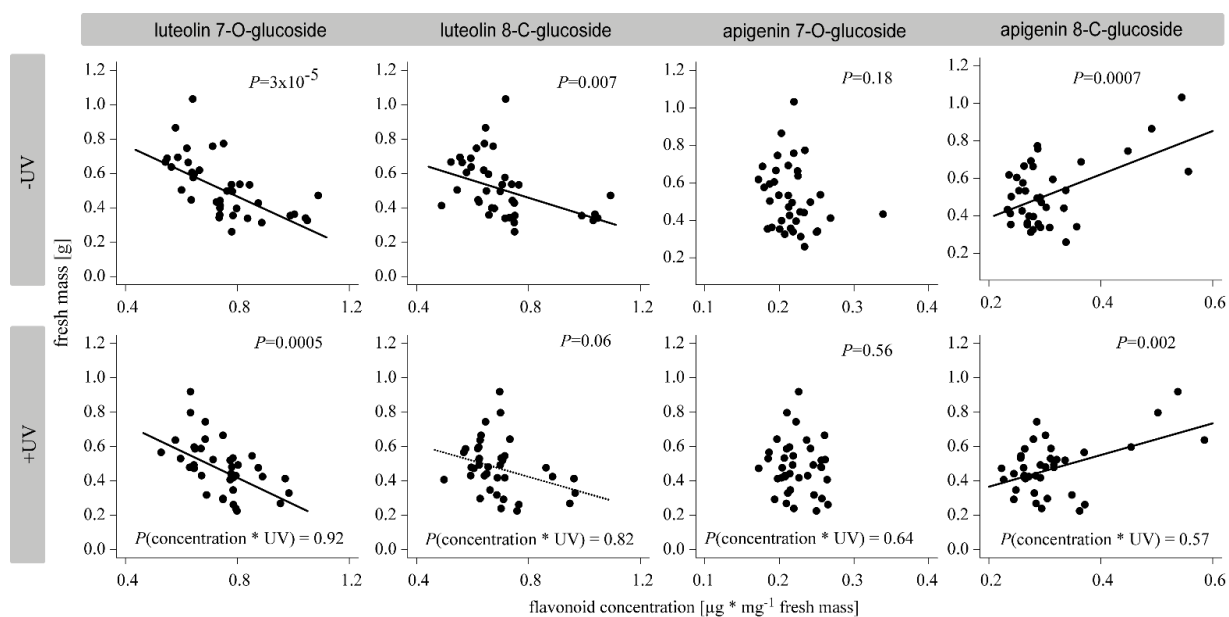


303

304 **FIGURE S20** Total flavonoid concentration did not correlate to relative plant fitness across 38 *S.*

305 *polyrhiza* genotypes under natural UV light exposure. *P*-value of linear model is shown.

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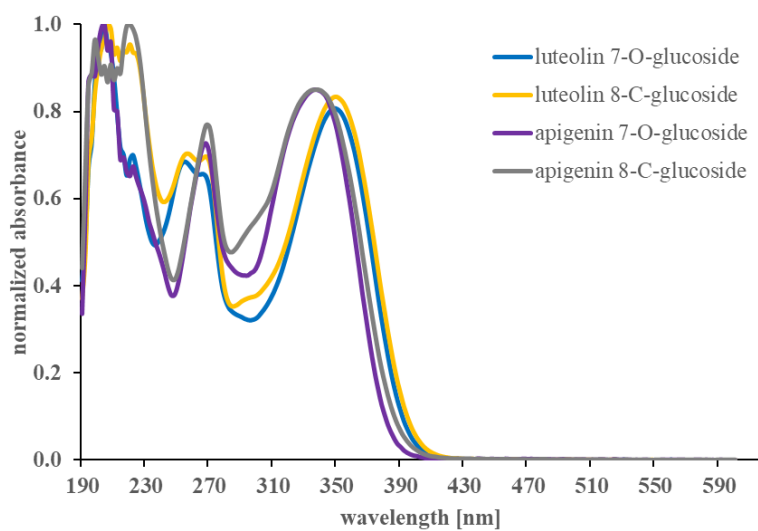
308 **FIGURE S21** Correlation of biomass production and individual flavonoid concentration across 38 *S.*

309 *polyrhiza* genotypes in the presence and absence of natural UV light. The correlations between biomass

310 production and individual as well as total flavonoid concentration in the presence and absence of UV

311 light were analysed using linear models based on the mean value of each treatment and genotype.

312



313

314 **Figure S22** UV spectra of flavonoid glucosides. Absorbance was normalized for each compound

315 separately.

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