

17 ABSTRACT

- 18 Plants compete for photosynthesis light and induce a shade avoidance syndrome (SAS) 19 that confers an important advantage in asymmetric competition for light at high canopy 20 densities. Shade plasticity was studied in a greenhouse experiment cultivating Arabidopsis thaliana plants from 15 populations spread across an altitudinal gradient in the Northeast 21 22 area of Spain that contain a high genetic variation into a reduced geographical range. Plants were exposed to sunlight or simulated shade to identify the range of shade 23 plasticity. Fourteen vegetative, flowering and reproductive traits were measured along to 24 25 the life cycle. Shade plasticity in flowering time and dry mass was significantly associated with the altitude of population origin. Plants from coastal populations showed higher 26 shade plasticity indexes than those from mountains. The altitudinal variation in flowering 27 leaf plasticity adjusted negatively with average and minimum temperatures, while dry 28 29 mass plasticity was better explained by negative regressions with the average, maximum 30 and minimum temperatures, and by a positive regression with average precipitation of the 31 population origin. The lack of an altitudinal gradient for the widest number of traits suggests that shade light could be a driver explaining the distribution pattern of 32 33 individuals in smaller geographical scales than those explored here.
 - 34

37

Key words: Light, phytochromes, phenotypic plasticity, shade avoidance syndrome,
 Arabidopsis thaliana, altitudinal gradient, structured populations, local adaptation.

- A brief summary statement (3-4 sentences maximum) highlighting the importance of the work. 38 39 Low red/far-red (R/FR) ratio of the light environment provides an early and unambiguous warning 40 of the presence of competing vegetation. This paper focuses on the study of the shade avoidance 41 syndrome in structured populations of Arabidopsis thaliana originated in an altitudinal gradient of 42 the Northeastern of Spain. Plasticity to shade of two important fitness traits, like flowering and 43 aerial dry biomass, were associated with the altitudinal gradient of population origin being plants 44 collected in the coast more plastic than those from the mountains. The clinal variation of the 45 shade plasticity index was negatively associated with temperature for both traits and positively with precipitation for dry biomass, suggesting that these climatic parameters could be relevant for 46 47 light adaptation in these populations.
 - 48

Accepted

49 INTRODUCTION

50 Plants need resources such as water, nutrients and light to grow. Under dense vegetation, 51 light is a limiting resource and competition for light can strongly influence the success of a 52 plant (Pierik and de Wit, 2013). Plants have evolved sophisticated mechanisms mediated 53 by phytochromes that allow them to detect the early presence of neighboring plants and 54 to initiate developmental adaptive developmental strategies that avoid shading before the canopy is closed (Ballaré et al., 1990). The most significant changes in the red/far-red 55 (R/FR) ratios occur when daylight is reflected or transmitted by green vegetation. 56 Absorption of red (R) and blue photons by chlorophyll and carotenoids results in a 57 58 selective enrichment of far-red (FR) photons, reducing the R/FR ratios perceived by the plant tissues. As a result of changes in the light spectrum, plants display the shade 59 60 avoidance syndrome (SAS), a set of physiological responses that increase vegetative 61 structures like stems, petioles and hypocotyls, accelerates flowering, and reduce seed 62 number and size (Casal, 2012).

63 In Arabidopsis thaliana and other species, phytochrome B (phyB) is the main phytochrome, and phyD and phyE contribute secondarily, mediating the SAS. In open 64 65 environments, the Pr, the inactive form of the phytochromes located in the cell cytoplasm, 66 migrates to the nucleus when it absorbs photons of R light and photo-transforms to Pfr. In 67 the nucleus, the accumulated Pfr form interacts and degradates PIFs (Phytochrome 68 Interacting Factors) through the proteosome leading to growth inhibition by the deactivation of gene expression (Lorrain et al. 2008). In opposition, the shade light 69 70 converts Pfr to Pr form that no longer interacts with PIFs. These proteins will thus rapidly

This article is protected by copyright. All rights reserved.

re-accumulate promoting the expression of early shade genes such as *PIL1, ATHB2, HFR1,*and *PAR1* inducing cell elongation responses (Lorrain et al., 2008, Hornitschek et al.,
2012). In addition, the full expression of SAS requires other photomorphogenic regulators
like COP1 (McNellis et al., 1994, Pacín et al., 2013), SPA (Rolauffs et al., 2012) double BBox proteins (Crocco et al., 2010; Gangappa et al., 2013) and bHLH/HLH transcription
factors (Hao et al., 2012).

The hypothesis of adaptive plasticity predicts that the phenotype of shade avoidance 77 induced by low R/FR ratios has a better fitness in dense canopies but is penalized at low 78 79 densities (Schmitt et al., 1995). Because the light is a critical resource for plants, the SAS 80 confers an important advantage in asymmetric competition for light at high densities. However in the absence of competition, allocation of resources to height at the expense 81 82 of leaves, roots and branches may reduce growth and reproduction, and elongated stems 83 may have a greater risk of mechanical damage (Casal and Smith, 1989; Schmitt and Wulff, 84 1993). Natural variation is a pre-requisite for the evolution of phenotypic plasticity (Via, 85 1985). At molecular level, nucleotide polymorphism at photoreceptor genes underlying natural variation in light responses (Aukerman et al., 1997; Balasubramanian et al., 2006; 86 87 El-assal et al., 2001; Maloof et al., 2001). PhyB is the principal photoreceptor responsible 88 for red light and shade avoidance responses, and is proposed to be the gene responsible for several QTL found when plants grow under shade (Borevitz et al., 2002; Botto et al., 89 2003; Botto and Coluccio, 2007; Kasulin et al., 2013). By the analysis of the phyB sequence 90 in 33 A. thaliana accessions, Filiault et al. (2008) found 14 non synonymous 91 92 polymorphisms with at least one of them responsible for the phenotypic variation

observed in seedlings exposed to red light. PIF4 polymorphisms are also supposed to be 93 94 associated with internode length of inflorescence and reproductive timing and fitness 95 under shade (Brock et al., 2010). Natural variation at *ELF3*, a gene involved in the circadian 96 clock, was clearly associated with a function for shade avoidance in A. thaliana. In fact, a single amino acid change in the ELF3 gene is responsible for the natural variation 97 98 mediating cell elongation growth (Coluccio et al., 2011) and flowering time (Jiménez-Gómez et al., 2010) between two contrasting accessions originated in Bayreuth (Bay, 99 Germany) and Shahdara (Sha, Tajikistan). Genetic diversity and structure analysis in more 100 101 than 6000 wild genotypes from different world regions, at global and regional scales, suggest several major events in A. thaliana demographic history in Europe (Nordborg et 102 al., 2005; Platt et al., 2010). In particular, high diversity has been described in the 103 104 Mediterranean Peninsulas compared to Central and Northern Europe (Beck et al., 2008; 105 Picó et al., 2008). The largest diversity has been found in the Iberian Peninsula, whose 106 strong geographic structure has prompted the hypothesis of multiple Iberian glacial 107 refuges with differential contribution to the colonization of Europe (Picó et al., 2008). The 108 structure of northeastern Iberian populations is important and contains huge genetic 109 variation across an altitudinal gradient suggesting that they may be locally adapted 110 (Montesinos-Navarro et al., 2009; Montesinos-Navarro et al., 2011; Wolfe and Tonsor, 2014). These A. thaliana populations grow in two contrasting climatic environments: 111 112 maritime lowland coastal area characterized by cool temperatures and moderate rainfall 113 in the winter, low rainfall and high maximum temperatures in spring and summer; and 114 higher altitude in mountainous areas, with higher rainfalls and lower minimum

115 temperatures in winter and a prolonged cool and wet spring. Interestingly, Montesinos-116 Navarro et al. (2011) found that the phenotypes of these populations are associated with 117 a climatic gradient defined by altitudinal clines. Working with northeastern Iberian 118 populations, Tonsor's group showed that biomass, leaf number, flowering time, and seed weight increase, whereas translocation of resources to the root, vegetative growth and 119 120 number of seeds decrease with the altitude of the genotype origin. These life strategies 121 favor the selection of individuals for rapid life cycle in Mediterranean regions near the sea avoiding typical warm dry summer periods, and long life cycles in individuals growing in 122 123 the mountains that help to maximize growth, cold tolerance in the winter and late 124 flowering.

The eco-physiological basis of the shade plasticity variation remains obscure. Some studies 125 126 have found a significant correlation between light sensitivity (Maloof et al., 2001; Stenøien 127 et al., 2002) and shade elongation response (Kasulin et al., 2013) with the latitude of 128 accession location, suggesting that light phenotypic variation could be a result of genotype 129 adaptation to a latitudinal gradient. Stenoien et al. (2002), working with 10 Norwegian 130 populations of A. thaliana collected in a narrow geographic range, found a latitudinal cline 131 in response to light: the northern genotypes are more responsive than southern 132 populations to R or FR continuous light during seedling de-etiolation. Furthermore, the hypocotyl elongation response to a FR pulse at the end of the day, a laboratory treatment 133 134 that simulates shade avoidance, was positively associated with the increase of latitude for European accessions collected between 15 and 65° (Kasulin et., 2013). However, in other 135 136 studies, hypocotyl and flowering shade response correlations with latitude are missing

(Botto and Smith, 2002; Filiault and Maloof, 2012). The strong structure of northeastern
Iberian populations, containing a wide range of genetic diversity within a narrow
geographical range, is an ideal system for testing hypotheses associated with the SAS. To
have a better understanding about the drivers of shade avoidance plasticity, we designed
a greenhouse experiment using the northeastern Iberian populations of *A. thaliana*. We
evaluated the range of variation of vegetative, flowering and reproductive traits in
response to simulated shade to answer the following questions:

a) What is the range of phenotypic variation to R/FR ratios in structured populations?
b) Is the expression of shade plasticity traits associated with an altitudinal gradient?
c) If the previous question is yes, what are the climatic drivers explaining this variation?

147

148 MATERIALS AND METHODS

149 Genetic material

Sixty genotypes from 15 populations of *A. thaliana* originated from the Northeast area of Spain were used in this study. These populations were collected in different locations defined by an altitudinal gradient (Montesinos-Navarro et al., 2009). In this area of collection, the rainfall increases and high spring temperatures and minimum winter temperatures decrease with the altitude (Montesinos-Navarro et al., 2011).

155

156 Culture conditions and light treatments

157 Seeds were sown in transparent plastic boxes on an agar solution of 0.8%. The boxes were 158 placed in darkness at 5 °C for one week to break dormancy. After that, the boxes with

159 seeds were placed in a chamber with continuous white light for another week to induce uniform germination and the development of seedlings with well-developed green 160 161 cotyledons and radicle. Then, seedlings were vernalized for two weeks in a light chamber 162 under non-inductive short-day conditions (8+16h light, dark) at 5 °C before transplanting. The seedlings were transplanted on 27^{th} August of 2011 to 7 x 4 cm pots (height x 163 164 diameter) with a substrate of vermiculite, perlite and peat in a ratio of 30:30:10. After that, the plants were grown with natural radiation and controlled temperature in a 165 greenhouse at IFEVA, Faculty of Agronomy, University of Buenos Aires (34º35'S, 58º29'W), 166 167 Buenos Aires, Argentina. The pots with plants were watered with Hoagland solution (20 milliliters of Hakaphos Compo Red solution in 5 liters of water). 168

After a week of transplanting, the plants were exposed to sunlight or simulated shade, a 169 treatment that consisted in sunlight plus lateral FR light mimicking neighboring plants 170 (Rondanini et al. 2014). The FR light was provided by two banks equipped with 9 171 172 incandescent reflector lamps of 40W each, and a red acetate filter with two filters of blue 173 acrylic Paolini (1 x 0.25 m long x wide) of 2 mm thick placed in front of the plants. To avoid 174 the increase of temperature by the lamps, transparent bottles with water were placed 175 between lamps and filters along with two fans that allowed ventilation. Plant grown in 176 simulated shade received R/FR ratios ranged between 0.07 and 0.12 (Suppl. Table 1). Two lines of plants were located in front of the Paolini filters. The pots with plants were 177 rotated every week to randomize light differences into the treatment. The sunlight 178 179 treatment consisted of a similar experimental design without the addition of FR light. The 180 lateral R/FR ratio was 0.65 (Suppl. Table 1). The sunlight on the top of the plants was similar between both light conditions: the average photosynthetically active radiation
(PAR) was near to 500 μmol m⁻² s⁻¹ at noon for sunny days and the R/FR ratios ranged
between 0.94 and 1.02 (Suppl. Table 1). During the experiment, the daily average
temperature ranged between 20.5 °C and 30.4 °C (Suppl. Fig. 1). PAR and R/FR ratios were
measured with a model Spectroradiometer SPECTROSENSE2 / 2 + Meter, Skye Instruments
Ltd (UK). The temperature was measured with a digital maximum and minimum
thermometer (Thermometer, Germany).

188

189 Traits and shade plasticity index

Fourteen vegetative, flowering and reproductive traits were measured during the 190 experiment: length and width of leaf, petiole length, leaf angle, rosette diameter and 191 192 height, flowering time as number of leaves or days at flowering, length and diameter of primary axis, number of basal axes, number of secondary axes on the primary 193 194 inflorescence, seed weight (100 seeds) and above-ground dry mass. The leaf angle was 195 taken in the first and second week after starting the light treatments with a goniometer 196 consisting of a protractor and a weight to mark the normal. The angle formed between 197 the normal and the tallest petiole leaf was estimated. Vegetative traits were measured 198 every week during the 28 d after the beginning of light treatments. Above-ground dry mass was assessed at the end of the trial (23/12/11) by placing the harvested aerial parts 199 200 (including flowering axes) in an oven at 80 °C for 72 hours and then dried material was weighed with a precision balance. Seed yield, as the total weight of seeds, was not 201 202 included in the analysis because heat stress increased towards the end of the experiment

- 203 (Suppl. Fig. 1) producing higher flower mortality at the extreme of inflorescences in the204 late flowering individuals compared with the earlier flowering individuals.
- 205 A shade plasticity index for each individual and trait was estimated as the difference 206 between sunlight and simulated shade relative to sunlight as follow:
- 207 Shade plasticity index = 1+ [(simulated shade sunlight)/sunlight]

Shade plasticity indexes higher than 1 indicate that simulated shade increases the response, and values lower than 1 indicate that simulated shade reduces the response with respect to sunlight. Values close to 1 mean that the individuals have low shade plasticity in opposition to higher or lower indexes that mean the individuals display strong shade plasticity.

213

214 Experimental design and statistical analysis

The experimental design was a randomized-block factorial design of two factors: 215 216 population (P) consisted in 15 populations distributed along an altitudinal gradient, and 217 light (L) consisted in two light conditions: sunlight and simulated shade. For each light 218 condition, 8 replicates for population were established and each population was 219 represented by 4 genotypes. Data were statistically analyzed by two-factor ANOVA 220 including P, L, P x L and B (block) factors. Paired comparisons by Bonferroni test were included when the P X L interaction factor was significant. Genetic correlations for 221 222 vegetative and reproductive traits within and between light treatments were estimated. 223 Univariate regression analyses were done to evaluate clinal population differentiation 224 between shade plasticity indexes and altitude or climatic parameters.

This article is protected by copyright. All rights reserved.

225 Because the SAS includes several morphological and developmental traits rather than any single factor, multivariate analysis was done. The suite of SAS traits were treated as a 226 227 group testing the effect of light of all the measured traits (F= 14.57, P < 0.0001). The 228 overall clinal population differentiation between shade plasticity indexes and altitude was evaluated using a multivariate analysis of variance MANOVA of all the measured traits as 229 230 dependent variables and altitude of population origin as the independent variable (F= 231 1.96, P < 0.0001). In addition, a principal component analysis (PCA) was conducted to represent the complexity of data matrix in two principal axes. All measured traits for each 232 233 individual were included in the analysis as dependent variables. For graphical representation, population and light were introduced as classification factors. Statistical 234 analyses performed Infostat 235 were using the statistical program 236 (http://www.infostat.com.ar/).

237

238 **RESULTS**

239 Time-course responses to simulated shade for vegetative traits

Six vegetative traits (length and width of lamina, petiole length and leaf angle, and diameter and height of rosette) were measured during the four weeks after the beginning of light treatments to study the effect of simulated shade on the time-course of vegetative growth. To increase the robustness of the analysis, the average response was estimated for the 15 populations in each light condition and date. The time-course growth was affected by simulated shade in four traits: leaf angle, petiole length, and height and diameter of rosette (Fig. 1). Simulated shade increased the erect position of the young leaves and rosette height at the starting of the experiment (Fig. 1, first and second weeks).
Furthermore, the shade light increased significantly the petiole length and rosette
diameter during the first month of the experiment (Fig. 1). The length and the width of the
leaves also increased systematically during the first month but no significant differences
were found between light treatments (Fig. 1).

253 **Reaction norms to simulated shade**

252

The average expression of the six vegetative traits with the exception of leaf length 254 255 differed significantly among populations and light treatments (Fig. 2, Suppl. Table 2, see 256 population and light factors). Simulated shade altered the vegetative phenotype in different intensities increasing the petiole length and the rosette height, and reducing the 257 leaf angle with respect to the normal. In addition, simulated shade reduced marginally the 258 259 leaf width and increased the rosette diameter in most of the populations (Fig. 2, Suppl. 260 Table 3). No significant effects were detected for the population by light interaction factor 261 in any of the six vegetative traits (Fig. 2, Suppl. Table 2).

Flowering was affected by population and light factors. As expected, the simulated shade accelerated flowering. The light factor was more sensitive for the number of leaves than for the number of days at flowering (Fig. 3, Suppl. Table 2). The population by light interaction effect was not significant for leaves and days at flowering. After flowering, six reproductive traits were measured. Population and light effects were significant for the number of basal axes, length of the principal axis, above-ground dry mass and seed weight. The population by light interaction factor was significant for the length of principal

axis and seed weight (Fig. 3, Suppl. Table 2). In these traits, plants from the ARU 269 population showed significantly longer axes under simulated shade compared with 270 271 sunlight (45.6 vs. 31.4 cm, respectively), and the individuals from the VDM population 272 produced significantly lighter seeds under sunlight compared with simulated shade (2.7 273 vs. 3.6 mg/100 seeds, respectively). These differences disappeared in other populations, 274 suggesting different light sensitivities to the same light signal (Suppl. Table 3). Although 275 the population factor was significant for the principal axis diameter and the number of secondary axes, the light factor did not affect significantly these reproductive traits (Fig. 3 276 277 and Suppl. Table 2).

278

279 Genetic correlations for traits within and between light treatments

280 Least-squares means of vegetative, flowering and reproductive traits were used to 281 estimate Pearson's product correlations and genetic variance-covariance matrices within 282 each light environment. Independently of the light factor, stronger positive correlations 283 were found among leaf length with other vegetative traits like as leaf width, petiole length 284 and rosette diameter; and also between flowering time (leaves or days) with leaf angle, 285 number of secondary axes and above-ground dry mass (Table 1). Some positive 286 correlations were only found in plants cultivated in simulated shade. For example, plants with higher above-ground dry mass produced wider flowering axes and heavier seeds 287 288 (Table 1).

A lower number of negative correlations were also found. For both light conditions, the rosette height showed a negative correlation with flowering (days or leaves) and

This article is protected by copyright. All rights reserved.

inflorescence length (Table 1). Interestingly, the rosette height was negatively correlated
with fitness traits such as dry biomass and seed weight, specifically in plants cultivated
under simulated shade (Table 1).

294

295 **Population origin is the principal driver of the phenotypic variation**

296 Multivariate analysis was applied for all the measurements of individuals corresponding to 297 15 populations and 14 traits in sunlight and simulated shade. The principal component analysis (PCA) reduced the variability of the data in a lower dimensional space than the 298 299 original space of variables. The first and the second dimensions of the PCA analysis (CP1 and CP2) explained 34 % and 19.6 % of the data variability (Fig. 4). The first axis ordered 300 the cases following a pattern that was very similar to the geographical gradient of the 301 302 population origin. The biplot representation allowed the identification of the variables 303 (arrows in the graph) that determine the location of the cases in this gradient: with some 304 exceptions, vegetative traits and coastal populations were grouped together on the left 305 side of the first axis, and flowering and reproductive traits together with populations 306 originating from the mountains were grouped on the other side. Light was the secondary 307 factor explaining the observations principally on the second axis. On the upper side of the 308 graph appeared the cases associated with plants cultivated in simulated shade while in the bottom were grouped the cases of plants exposed to sunlight with the exception of BOS 309 310 population (see 8: shade).

311

312 Shade plasticity for flowering and dry mass is associated with an altitudinal cline

This article is protected by copyright. All rights reserved.

313 The shade plasticity was estimated in six vegetative and eight flowering and reproductive 314 traits for each population as the average response of the individuals. To test whether the 315 shade plasticity is associated with the altitude of the place of population origin, regression 316 analyses were done for each trait. In those traits associated significantly with an altitudinal 317 pattern, climatic parameters were examined in order to explain this variation. Number of 318 leaves or days at flowering and above-ground dry mass plasticity indexes showed a clear 319 and significant regression with the altitude of origin (Fig. 5). The distribution of shade plasticity indexes adjusted to a regression line that differed from the horizontal (Fig. 5, P = 320 321 0.003 for leaves at flowering, P = 0.037 for days at flowering and P= 0.001 for above-322 ground dry mass). Individuals from coastal areas showed a higher plasticity to shade than those plants from mountain locations that displayed null or reduced shade plasticity (Fig. 323 5). Shade plasticity indexes for other vegetative and reproductive traits were not 324 325 associated with the altitude of population origin (Suppl. Fig. 2 and 3). Furthermore, 326 flowering leaf plasticity index showed a significant regression with average and minimum 327 temperatures (Fig. 6), but not with maximum temperature neither average precipitation 328 of the place of population origin (Suppl. Fig. 4). In other words, individuals from coastal 329 areas that experience higher average and minimum temperatures showed higher 330 flowering shade plasticity than those individuals from mountains sites. The shade plasticity of above-ground dry mass showed a negative regression with temperatures 331 332 (average, minimum and maximum) and a positive regression with the average 333 precipitation (Fig. 6), but not with the distribution of precipitations in autumn and spring 334 (Suppl. Fig. 4). It means that higher dry mass plasticity index was associated with plants

- from coastal areas growing with higher temperatures and lower average precipitation than those from mountainous areas. No significant regressions were found for the flowering day plasticity index and the six climatic parameters evaluated (Suppl. Fig. 5).
- 338

339 DISCUSSION

340 Shade response in vegetative, floral and reproductive traits was studied in 15 populations 341 of A. thaliana spread across an altitudinal transect in the northeastern area of Spain. These 342 populations contain a high genetic diversity and are characterized by a strong population 343 structure (Montesinos-Navarro et al., 2011; Picó et al., 2008). Vegetative traits such as petiole length, leaf width, diameter and height of rosette, and angle of insertion of the 344 leaf were significantly affected by changes in R/FR ratios. The simulated shade produced 345 346 plants with narrower lamina, larger petioles and rosette diameter, and also more erected 347 leaves at early developmental stages compared with those plants cultivated under 348 sunlight (Fig. 1 and 2). In other species close to Arabidopsis, like rapeseed plants, it has 349 also been observed that low R/FR ratios induce dramatic shade avoidance responses. 350 Shade signals increase the leaf length but not the leaf width, and produce elevated leaf 351 angles in early stages of the development of a spring rapessed hybrid (Rondanini et al., 352 2014). It is well known that the low R/FR ratios of the reflected light provide early warnings of the presence and proximity of neighboring plants, allowing the initiation of 353 354 development adaptive strategies to avoid shading before the canopy is closed (Ballaré et al., 1990). Furthermore, a huge natural variation for shade avoidance responses was 355 356 documented in a representative panel of Arabidopsis accessions (Botto and Smith, 2002).

Interestingly, natural variation at the *ELF3*, a circadian clock gene, is responsible for the
shade avoidance variation for hypocotyl length elongation, leaf angle movement (Coluccio
et al., 2011) and flowering response (Jiménez-Gómez et al., 2010) between Bay and Sha
accessions of *A. thaliana*. The altered shade elongation response and leaf movement in
Sha accession was associated to a rare Alanine by Valine substitution that alters ELF3-Sha
circadian rhythms of leaf movements and clock gene expression (Coluccio et al., 2011,
Anwer et al., 2014).

Shade light accelerates flowering response in A. thaliana plants. Some studies show that 364 365 the number of leaves at flowering is a more sensitive trait than bolting time (Fig. 3, Callaghan and Pigliucci, 2002, Botto and Coluccio, 2007). In Arabidopsis, low R/FR ratios 366 accelerate flowering by enhancing the expression of FLOWERING LOCUS T (FT), the gene 367 368 involved in the induction of flowering by long days (Halliday et al., 2003). Callaghan and Pigliucci (2002) found that flowering time was accelerated by shade under field 369 370 conditions, but not when Arabidopsis plants where grown in a greenhouse with the presence of grass neighbors. However, working with a wide range of natural variation, 371 372 flowering time in response to low R/FR ratios was accelerated either when plants were 373 cultivated in a light chamber (Botto and Smith, 2002) or in a greenhouse (Botto and 374 Coluccio, 2007).

375 Reproductive traits were also affected by shade. Low R/FR ratios produced taller 376 inflorescences and reduced the number of flowering axes, plant biomass and seed weight 377 (Fig. 3). The effect of light on the inflorescence length was dependent on the population. 378 Interestingly, the ARB population produced longer axes under simulated shade (P X L

This article is protected by copyright. All rights reserved.

interaction, P= 0.0022) but these differences disappeared in other populations suggesting 379 380 that shade sensitivity depends on the origin of the population. Although cell elongation is 381 stimulated by shade, Brock et al. (2010) found that most of the accessions of A.thaliana 382 cultivated in low density in a greenhouse developed taller inflorescences than those 383 growing in crowded stands. The authors interpreted these odd results as a limitation of 384 translocation resources from leaves to fruits. Furthermore, branching is inhibited by low R/FR ratios. Loss of phyB function leads to a reduced branching by a down-regulation of 385 the expression of auxin genes (Reddy and Finlayson, 2014; Su et al., 2011). Molecular and 386 387 pharmacological assays suggest that the active form of phyB suppresses auxin signaling to promote branching (Reddy and Finlayson, 2014). Furthermore, in the present study, the 388 simulated shade reduced above-ground dry mass and seed weight (Fig. 3), and these 389 390 results are in accordance with previous evidence demonstrating that environments with 391 resource limitation, such as low R/FR ratios, reduce plant growth and productivity (Sultan, 2000). 392

393 It is well known that climatic variables have important consequences for the geographical 394 distribution of individuals and species. Shade plasticity indexes of flowering time and 395 above-ground dry mass were significantly associated with the altitude of collection place 396 (Fig. 5). However, other plasticity indexes of vegetative and reproductive parameters did not show clinal variation associated with the altitude (Suppl. Fig. 2 and 3). Interestingly 397 plants from coastal populations showed higher plasticity to shade than mountainous 398 399 populations, suggesting different light sensitivities according with the population origin. In 400 fact, low R/FR ratios accelerated flowering and reduced the plant biomass more

401 dramatically in coastal populations than in mountainous populations. Furthermore, some 402 climatic parameters were significantly associated with shade plasticity indexes of some 403 traits. For example, the flowering leaf index showed a significant correlation with the 404 average temperature and the minimum temperature of population origin, while the shade plasticity index for dry mass was better explained by the pattern of variation in 405 406 temperatures and the average precipitation (Fig. 6). Northeastern Iberian populations of 407 A.thaliana show strong demographic and genetic patterns defined by the altitude of origin, with mountain populations less genetically diverse than coastal populations 408 409 (Montesinos-Novarro et al., 2009; Picó, 2012). The drivers of this altitudinal cline are associated with colder winter temperatures and wetter and longer springs in mountain 410 areas (Montesinos-Navarro et al., 2009; Gomaa et al., 2011). Accordingly, the patterns of 411 412 evolutionary diversification in these structured populations can be influenced by the 413 plasticity to light. As predicted by the ecological theory, adaptation through natural 414 selection will not occur as readily for genetically distinct coastal populations because 415 individuals are more plastic and produce phenotypes more appropriate to different local 416 environments. Conversely, mountain ecotypes, in which individuals express limited 417 plasticity, would be predicted to show greater response to local selection regimes and 418 therefore greater genetic divergence (Sultan, 2000).

The altitudinal patterns in shade plasticity found for flowering leaf and biomass were obtained from plants cultivated in optimal growth conditions. We should be cautious in generalizing the conclusion of this work to other suboptimal environmental conditions. In fact, the expression of the shade avoidance plasticity can be limited by

This article is protected by copyright. All rights reserved.

423 microenvironmental variation in water availability in seedlings from natural populations of
424 *Impatiens capensis* (Huber et al., 2004). The authors found that local seedling density was
425 a poor predictor of selection on shade-avoidance traits as a consequence of the
426 unpredictability of water availability, particularly in dry microsites that may affect the
427 costs and benefits of expressing shade avoidance (Huber et al., 2004).

428 The results of this work illustrate clearly that the shade plasticity for flowering and dry 429 biomass show clinal variation associated with altitude in structured populations of 430 Arabidopsis originating in the Northeast area of Spain. Ecotype differences in response to 431 shade signals were also documented for Stellaria longipes adapted to two ecological 432 environments. In concordance with the results showed here, the prairie ecotype responds quickly to low R/FR ratios elongating their ramets as an adaptation to growth in dense 433 434 vegetation stands, in contrast to the alpine ecotype that displays dwarf phenotypes with 435 resistance to wind but unresponsive to shade signals allowing adaption to areas of sparse 436 vegetation where abiotic stresses predominate (Sasidharan et al., 2008). Furthermore, the 437 shade light, changing also across a micro-environmental context, may be a driver 438 explaining the distribution patterns of individuals in smaller geographical scales than those 439 explored here. New experimental approaches should be undertaken to test this 440 hypothesis. To evaluate this idea it is necessary to work with a bigger collection of populations correctly described both in their geographical positions and types and 441 442 environments (woodland, scrubland, anthropic, prairie, etc.); as well as having detailed 443 descriptions of the environmental conditions of the collection site (radiation, light quality, 444 etc.). The atlas of ecological and climatic information along with genetic databases of each

This article is protected by copyright. All rights reserved.

individual and their corresponding phenotype may help to identify the underlying genes
that express the enormous plasticity documented in the SAS. Deciphering the genetic and
molecular basis of phenotypic plasticity is a challenge to understand how plants function,
and it is essential to understand the evolutionary forces operating in the adaptation of
species to a changing environment (Alonso-Blanco and Méndez-Vigo, 2014).

451 ACKNOWLEDGEMENTS

450

452 I thank Felipe Kleine for his technical support in greenhouse experiments. I also thank
453 Stephen Tonsor for the gift of the seeds used in this work, Susana Perelman for statistical
454 discussions, and Xavier Picó, Stephen Tonsor and Carlos Alonso-Blanco for the stimulating
455 talks. This research was supported by Foncyt/SECyT and UBACyT grants to JFB.

- 457 **REFERENCES**
- 458 Alonso-Blanco C., Méndez-Vigo B. 2014. Genetic architecture of naturally occurring
 459 quantitative traits in plants: an updated synthesis. *Current Opinion in Plant Biology*460 18:37-43. DOI: http://dx.doi.org/10.1016/j.pbi.2014.01.002.
- Anwer M.U., Boikoglou E., Herrero E., Hallstein, M., Davis, A. M., Velikkakam J.G, Nagy F.,
 Davis, S.J. 2014. Natural variation reveals that intracellular distribution of ELF3
 protein is associated with function in the circadian clock. *eLife* 10.7554/eLife.02206
 Aukerman M.J., Hirschfeld M., Wester L., Weaver M., Clack T., et al. 1997. A deletion in
 the *PHYD* gene of the Arabidopsis Wassilewskija ecotype defines a role for
 phytochrome D in red/far-red light sensing. *Plant Cell* **9**:1317-1326.
- Balasubramanian S., Sureshkumar S., Agrawal M., Michael T.P., Wessinger C., et al. 2006.
 The PHYTOCHROME C photoreceptor gene mediates natural variation in flowering
 and growth responses of *Arabidopsis thaliana*. *Nature Genetics* 38:711-715.
- Ballaré C.L., Scopel A.L., Sánchez R.A. 1990. Far-red radiation reflected from adjacent
 leaves: an early signal of competition in plant canopies. *Science* 247:329-332.
- Beck J.B., Schmuths H., Schaal B.A. 2008. Native range genetic variation in Arabidopsis
 thaliana is strongly geographically structured and reflects Pleistocene glacial
 dynamics. *Molecular Ecology* 17:902-915. DOI: 10.1111/j.1365-294X.2007.03615.x.
 Borevitz J.O., Maloof J.N., Lutes J., Dabi T., Redfern J.L., et al. 2002. Quantitative trait loci
 controlling light and hormone response in two accessions of *Arabidopsis thaliana*. *Genetics* 160:683-696.

This article is protected by copyright. All rights reserved.

478 Botto J.F., Smith H. 2002. Differential genetic variation in adaptive strategies to a common
479 environmental signal in Arabidopsis accessions: phytochrome- mediated shade
480 avoidance. *Plant, Cell and Environment* 25:53-63.

481 Botto J.F., Coluccio M.P. 2007. Seasonal and plant-density dependency for quantitative
482 trait loci affecting flowering time in multiple populations of *Arabidopsis thaliana*.
483 *Plant, Cell and Environment* **30**:1465-1479.

- Botto J.F., Alonso-Blanco C., Garzarón I., Sánchez R.A., Casal J.J. 2003. The Cvi allele of
 cryptochrome 2 enhances cotyledon unfolding in the absence of blue light in *Arabidopsis. Plant Physiology* 133:1547-1556.
- Brock M.T., Maloof J.N., Weinig C. 2010. Genes underlying quantitative variation in
 ecologically important traits: PIF4 (PHYTOCHROME INTERACTING FACTOR 4) is
 associated with variation in internode length, flowering time, and fruit set in
 Arabidopsis thaliana. *Molecular Ecology* 19:1187-1199. DOI: 10.1111/j.1365294X.2010.04538.x.

492 Casal J.J. 2012. Shade Avoidance. *The Arabidopsis Book*:**e0157**. DOI: 10.1199/tab.0157.

- 493 Casal J.J., Smith H. 1989. The "end-of-day" phytochrome control of internode elongation
 494 in mustard: kinetics, interaction with the previous fluence rate and ecological
 495 implications. *Plant, Cell and Environment* 12:511-520.
- Coluccio M.P., Kasulin L., Yanovsky M.J. and Botto J.F. 2011. Genetic mapping of natural
 variation in a shade avoidance response: ELF3 is the candidate gene for a QTL in
 hypocotyl growth regulation. *Journal of Experimental Botany* 62:167-176.

This article is protected by copyright. All rights reserved.

- 499 Crocco C.D., Holm M., Yanovsky M.J., Botto J.F. 2010. AtBBX21 and COP1 genetically
 500 interact in the regulation of shade avoidance. *The Plant Journal* 64:551-562. DOI:
 501 10.1111/j.1365-313X.2010.04360.x.
- 502 El-assal S.E., Alonso-Blanco C., Peeters A.J., Raz V., Koornneef M. 2001. A QTL for flowering
 503 time in *Arabidopsis* reveals a novel allele of *CRY2. Nature Genetics* 29:435-439.
 504 Filiault D.L., Maloof J.N. 2012. A Genome-Wide Association Study Identifies Variants
 505 Underlying the *Arabidopsis thaliana* Shade Avoidance Response. *PLoS Genet*506 8:e1002589. DOI: 10.1371/journal.pgen.1002589.
- Gangappa S.N., Crocco C.D., Johansson H., Datta S., Hettiarachchi C., et al. 2013. The
 Arabidopsis B-BOX Protein BBX25 Interacts with HY5, Negatively Regulating BBX22
 Expression to Suppress Seedling Photomorphogenesis. *The Plant Cell* 25:1243 1257. DOI: 10.1105/tpc.113.109751.
- Gomaa N.H., Montesinos-Navarro A., Alonso-Blanco C., Picó F.X. 2011. Temporal variation
 in genetic diversity and effective population size of Mediterranean and subalpine
 Arabidopsis thaliana populations. *Molecular Ecology* 20:3540-3554. DOI:
 10.1111/j.1365-294X.2011.05193.x.
- Halliday K.J., Salter M.G., Thingnaes E., Whitelam G.C. 2003. Phytochrome control of
 flowering is temperature sensitive and correlates with expression of the floral
 integrator FT. The *Plant Journal* 33:875-885. DOI: 10.1046/j.1365313X.2003.01674.x.



This article is protected by copyright. All rights reserved.

Maloof J.N., Borevitz J.O., Dabi T., Lutes J., Nehring R.B., et al. 2001. Natural variation in 541 light sensitivity of Arabidopsis. Nature Genetics 29:441-446 doi:10.1038/ng777 542 McNellis T.W., von Arnim A.G., Araki T., Komeda Y., Miséra S., Deng X.W. (1994) Genetic 543 544 and molecular analysis of an allelic series of cop1 mutants suggests functional roles for the multiple protein domains. *The Plant Cell Online* **6**:487-500. DOI: 545 546 10.1105/tpc.6.4.487. 547 Montesinos-Navarro A., Tonsor S.J., Alonso-Blanco C., Picó F.X. 2009. Demographic and Genetic Patterns of Variation among Populations of Arabidopsis thaliana from 548 549 Environments. Contrasting Native PLoS ONE **4**:e7213. DOI: 550 10.1371/journal.pone.0007213. Montesinos-Navarro A., Wig J., Xavier Pico F., Tonsor S.J. 2011. Arabidopsis thaliana 551 552 populations show clinal variation in a climatic gradient associated with altitude. 553 *New Phytologist* **189**:282-294. DOI: 10.1111/j.1469-8137.2010.03479.x. 554 Nordborg M., Hu T.T., Ishino Y., Jhaveri J., Toomajian C., et al. 2005. The pattern of 555 polymorphism in Arabidopsis thaliana. PLoS Biology 3:1289-1299. 556 Pacín M., Legris M., Casal J.J. 2013. COP1 re-accumulates in the nucleus under shade. The 557 Plant Journal 75:631-641. DOI: 10.1111/tpj.12226. 558 Picó F.X. 2012. Demographic fate of Arabidopsis thaliana cohorts of autumn- and springgerminated plants along an altitudinal gradient. Journal of Ecology 100:1009-1018. 559 560 DOI: 10.1111/j.1365-2745.2012.01979.x.

This article is protected by copyright. All rights reserved.

- 561 Picó F.X., Méndez-Vigo B., Martínez-Zapater J.M., Alonso-Blanco C. 2008. Natural Genetic
 562 Variation of Arabidopsis thaliana Is Geographically Structured in the Iberian
 563 Peninsula. *Genetics* 180:1009-1021. DOI: 10.1534/genetics.108.089581.
- 564 Pierik R., de Wit M. 2013. Shade avoidance: phytochrome signalling and other
 565 aboveground neighbour detection cues. *Journal of Experimental Botany* 65:2815566 2824. OI: 10.1093/jxb/ert389.
- Platt A., Horton M., Huang Y.S., Li Y., Anastasio A.E., Mulyati N.W., Ågren J., et al. 2010.
 The Scale of Population Structure in *Arabidopsis thaliana*. *PLoS Genet* 6:e1000843.
 DOI: 10.1371/journal.pgen.1000843.
- 570 Reddy S.K., Finlayson S.A. 2014. Phytochrome B Promotes Branching in Arabidopsis by
 571 Suppressing Auxin Signaling. *Plant Physiology* 164:1542-1550. DOI: 10.1104/pp.113.234021.
- 573 Rolauffs S., Fackendahl P., Sahm J., Fiene G., Hoecker U. (2012) Arabidopsis COP1 and SPA
 574 Genes Are Essential for Plant Elongation But Not for Acceleration of Flowering
 575 Time in Response to a Low Red Light to Far-Red Light Ratio. *Plant Physiology*576 160:2015-2027. DOI: 10.1104/pp.112.207233.
- 577 Rondanini D.P., Vilariño M.P., Roberts M.E., Polosa M.A., Botto J.F. 2014. Physiological 578 responses of spring rapeseed (Brassica napus L.) to red/far-red ratios and 579 irradiance on pre and post flowering stages. *Physiologia Plantarum* 580 DOI: 10.1111/ppl.12227.
- 581 Sasidharan R., Chinnappa C.C., Voesenek L.A.C.J., Pierik R. 2008. The Regulation of Cell
 582 Wall Extensibility during Shade Avoidance: A Study Using Two Contrasting Ecotypes

This article is protected by copyright. All rights reserved.



This article is protected by copyright. All rights reserved.



Fig. 1: Time-course response to simulated shade for vegetative traits.

- Response to R/FR ratios was calculated as the average of 15 Iberian populations at each
 date. The lines outside the box plot graphs indicate the minimum and maximum values for
 sunlight and simulated shade between 7 and 28 days after the starting of light treatments.
 Means were compared by Tukey test (P < 0.05) after analysis of One-Way ANOVA.
 - 608

609 Fig. 2: Reaction norms to simulated shade for vegetative traits.

- Each point represents the average response in the control and low R/FR treatments for
 each population. On the top of each graph is indicated the output of the ANOVA analysis
 for each independent variable (L= light, P= population) and the interaction between L x P.
 *, ** and *** indicate P < 0.05, 0.01 and 0.001, respectively. ns: not significant.
- 614

615 Fig. 3: Reaction norms to simulated shade for flowering and reproductive traits.

- Each point represents the average response in the control and low R/FR treatments. On the top of each graph is indicated the output of the ANOVA analysis for each independent variable (L= light, P= population) and the interaction between L x P. *, ** and *** indicate P < 0.05, 0.01 and 0.001, respectively. ns: not significant.
- 620

Fig. 4: Multivariate analysis for all the traits from plants of 15 populations cultivated insunlight and simulated shade.

Principal component analysis (PCA) was conducted to reduce the complexity of the data
matrix in two eigenvectors. All the measured traits for each individual were included in
the analysis as dependent variables and population and light as classification factors.
Numbers indicate populations as 1: PIN, 2: RAB, 3: SAL, 4: BAR, 5: HOR, 6: ARU, 7: COC, 8:
BOS, 9: MUR, 10: VDM, 11: ALE, 12: PAL, 13: BIS, 14: VIE, and 15: PAN. For additional
references on traits names see Table 1.

629

Fig. 5: Clinal variation associated with altitude for flowering and above-ground massplasticity to simulated shade.

Regression fitted between plasticity shade indexes for flowering and above-ground mass and altitude of the population origin. Fitting regression lines are presented with R² and P values indicating significant regression with respect to zero. Each point represents the average plasticity response estimated as the difference between simulated shade and sunlight relative to sunlight. Values closed to 1 mean that the population has null or low plasticity in opposition to higher or lower indexes than 1 indicating that populations display strong shade plasticity.

639

Fig. 6: Climatic parameters associated with clinal variation for leaf flowering and aboveground mass plasticity to simulated shade.

Plasticity traits are represented as function of the climatic parameters associated to each population. Fitting regression lines are presented with R² and P value indicating significant regression with respect to zero. For other references see Fig. 5.

This article is protected by copyright. All rights reserved.

- 645 Suppl. Fig. 1: Average, minimum and maximum temperature during the experiment.
- Daily temperatures (minimum, maximum and average) measured during the experiment
 into the greenhouse. The experiment started on 28th of August and finished on 23rd of
 December 2011.
- 649

650 Suppl. Fig. 2: Shade plasticity in vegetative traits regressed with altitude.

- Plasticity traits are represented as function of the altitudinal gradients. Each point represents the average plasticity response estimated as the difference between simulated shade and sunlight relative to sunlight. Values closed to 1 mean that the population has null or low plasticity in opposition to higher or lower indexes than 1 indicating that populations display strong shade plasticity.
- 656

657 Suppl. Fig. 3: Shade plasticity for flowering and reproductive traits regressed with 658 altitude.

- Plasticity traits are represented as function of the altitudinal gradients. For otherreferences see Suppl. Fig. 2.
- 661

666

- Fig Suppl. 4: Climatic parameters not associated with clinal variation for leaf flowering and above-ground dry mass to simulated shade.
- Plasticity traits are represented as function of the climatic parameters associated to eachpopulation.

- 667 Fig Suppl. 5: Climatic parameters not associated with clinal variation for flowering day to
- 668 simulated shade.
- 669 Plasticity traits are represented as function of the climatic parameters associated to each
- 670 population.

A C C

671 Table 1: Genetic correlation matrix of vegetative, flowering and reproductive traits from *A. thaliana* plants exposed to solar and low R/FR

672 ratios. Sunlight (above the diagonal) and low R:FR conditions simulating foliar shade (below the diagonal). Significant correlations are indicated

by *** (P<0.001), ** (P<0.01) and * (P<0.05). LL, leaf length; LW, leaf width; PL, petiole length; LA, leaf angle; RD, rosette diameter; RH, rosette
height; FD, flowering in days; FL, flowering in leaves; PAL, principal axe length; PAD, principal axe diameter; SAN, secondary axe number; AN,

675 basal axe number; DW, above dry weight; and SW, seed weight.

676

		LL	LW	PL	LA	RD	RH	FD	FL	PAL	PAD	SAN	AN	DW	SW
	LL		0.55***	0.51***	-0.16	0.78***	0.4***	-0.23*	-0.08	-0.18	0.2*	-0.11	0.31**	0.42***	0.15
	LW	0.69***		0.37***	-0.09	0.44***	0.16	0.001	0.03	-0.19	0.19*	-0.01	0.12	0.16	0.2
	PL	0.64***	0.59***		-0.06	0.58***	0.05	0.1	0.14	-0.21*	0.12	0.009	0.31**	0.28**	0.07
	LA	-0.08	-0.01	0.05		-0.13	-0.58***	0.34***	0.39***	0.1	-0.02	0.28***	-0.31***	0.03	0.13
	RD	0.91***	0.71***	0.82***	-0.05		0.29**	-0.04	0.06	-0.15	0.15	0.004	0.22*	0.42***	0.1
	RH	0.06	0.09	-0.16	-0.46***	-0.03		-0.57***	-0.48***	-0.2*	-0.04	-0.31***	0.27**	-0.01	-0.08
	FD	-0.13	-0.11	0.06	0.38***	-0.06	-0.47***		0.9***	0.17	0.15	0.66***	-0.34**	0.34***	0.3*
	FL	-0.02	-0.04	0.12	0.4***	0.04	-0.43***	0.92***		0.18	0.25**	0.67***	-0.26**	0.42***	0.35***
	PAL	0.04	-0.1	0.14	0.17	0.11	-0.28**	0.16	0.21*		0.22*	0.25**	-0.11	0.21*	-0.08
	PAD	0.13	0.13	0.3**	0.09	0.25**	-0.19	0.24*	0.3**	0.4***		0.31**	-0.06	0.018	0.03
	SAN	-0.08	-0.01	0.13	0.09	0.01	-0.11	0.55***	0.54***	0.2*	0.46***		-0.23*	0.34***	0.26*
	AN	0.2	0.13	0.28**	-0.02	0.25**	-0.04	-0.1	-0.1	0.04	0.02	-0.22*		0.28**	-0.15
	DW	0.1	0.09	0.14	0.19	0.17	-0.35***	0.68***	0.68***	0.28**	0.34***	0.39***	0.19		0.18
	SW	-0.17	-0.19	-0.16	0.44***	-0.15	-0.49***	0.35**	0.35**	0.29*	0.17	0.22	0.09	0.39***	

677

Acceb



Fig. 1

U S



Fig. 2



Fig. 3

0 Acce



Fig. 4





Fig. 5



Fig. 6