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Evolutionary responses of invasive grass species to variation in precipitation and soil nitrogen

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Summary

1. Global climate models suggest that many ecosystems will experience reduced precipitation over the next century and the consequences for invasive plant performance are largely unknown. Annual invasive species may be able to quickly evolve traits associated with drought escape or tolerance through rapid genetic changes.

2. We investigated the influence of five years of water and nitrogen manipulations on trait values in a southern California grassland system. Seeds from two annual grass species (*Avena barbata*, *Bromus madritensis*) were collected from experimental plots and grown in a common environment over two generations. We measured 14 physiological, morphological, phenological, and reproductive traits.

3. Both species displayed phenotypic differences depending on the water treatment from which they were collected, but not depending on the nitrogen treatment. Both species displayed trait values characteristic of drought escape (e.g., earlier flowering in *A. barbata* and *B. madritensis*, lower water-use efficiency in *B. madritensis*) when grown from seeds collected from plots that experienced five years of reduced precipitation. Furthermore, *A. barbata* individuals grown...
from seeds collected from drought plots had higher reproductive output and higher photosynthetic performance than individuals grown from water addition plots, with individuals grown from ambient plots displaying intermediate trait values. Notably, we found no phenotypic variation among treatments for six root traits.

4. Synthesis. Trait differences were observed following two generations in a common garden, suggesting that treatment differences were genetically based. This suggests that populations were responding to selection over the five years of water manipulations, a remarkably short time period. The rapid evolutionary responses observed here may help these two widespread invasive grass species thrive under reduced precipitation scenarios, which could have important implications for fire dynamics, invasive species management, and native plant restoration in communities invaded by annual grasses.

Key-words: annual species, *Avena barbata, Bromus madritensis*, drought escape, invasion ecology, invasive species, phenology, photosynthesis, root traits, water-use efficiency
Introduction

Global change, including increases in temperature, atmospheric CO$_2$ concentration, and nitrogen (N) input from anthropogenic activities, is one of the most significant threats to natural ecosystems. Further, global climate models have predicted alterations in inter- and intra-annual precipitation in many regions (Melillo et al., 2014). Biological invasion is another important agent of global change with significant effects on global biodiversity (Millennium Ecosystem Assessment, 2005). A meta-analysis of 1041 cases of invasion found that alien invasive plants decrease the abundance and diversity of native resident plant species by 43.5% and 50.7%, respectively (Vilà et al., 2010). Significant impacts on native resident species’ abundance and diversity are more likely to occur if the invading species is an annual grass (Pyšek et al., 2012). Refining our understanding of how invasive plants respond to environmental variation will improve our ability to predict their effects on natural plant communities in the face of global change. In this study, we explore the adaptive responses of two annual invasive grass species to declines in precipitation and increased N deposition in a southern California grassland.

There are two major strategies a plant can employ to cope with reduced water availability: drought tolerance and drought escape. Drought tolerance, characterized by low photosynthetic activity, high water-use efficiency (WUE; the rate of carbon assimilation relative to transpiration), and slow growth, allows an individual to conserve resources while continuing to be active during the drought period (Sherrard and Maherali, 2006, Franks, 2011). Conversely, drought escape involves the completion of an individual’s life cycle before the drought reaches its most extreme state. Drought escape is characterized by high photosynthetic rates, low WUE, and early flowering (Sherrard and Maherali, 2006, Franks, 2011). An increase in N allocation to photosynthetic enzymes and chlorophyll content in leaves may act to increase light harvesting
(Terashima and Evans, 1988, Evans, 1989) and, consequently, carbon fixation over a shorter period of time. Thus, increased N availability resulting from human activities may enhance growth in species displaying a drought escape strategy.

Will invasive species benefit from projected arid, N-rich conditions? High soil N concentrations can increase invasive species performance (Huenneke et al., 1990, Ostertag and Verville, 2002, Gross et al., 2005, Davis et al., 2000); however, the effects of N addition are complex and will depend on soil water availability, competition, and the form of available N (Everard et al., 2010, Ross et al., 2011, Eskelinen and Harrison, 2014). Furthermore, an analysis of several U.S. databases suggests that invasive species tend to initiate leaves and flower earlier in the growing season compared to native species (Wolkovich and Cleland, 2011). This phenological pattern, consistent with drought escape, suggests that many annual invasive species may be well-suited to increasingly arid conditions.

Many studies have shown that invasive species can be phenotypically plastic in response to changes in water, light and nutrient availability (e.g., Funk, 2008, Davidson et al., 2011); however, fewer studies have examined the potential for invasive species to adapt to environmental variation through genetic changes. In a study comparing flowering times of annual Brassica rapa before and after a four year drought in southern California, Franks et al. (2007) found that seeds collected from wet and dry environments post-drought flowered significantly earlier than those collected before the drought, which is characteristic of a drought escape response. Further, Franks (2011) suggested that populations of B. rapa escape drought through reduced WUE, which allows for rapid development and earlier flowering. A study of the annual invasive grass Avena barbata grown in wet and dry environments also found strong evidence that earlier flowering was adaptive under drought (Sherrard and Maherali, 2006).
Despite evidence of rapid evolution in the flowering times of *B. rapa* and *A. barbata*, adaptive responses to climate change may be slower or lacking in other traits and for other species (Franks et al., 2007). Anatomical and physiological characteristics of roots will influence both water and N acquisition, yet few studies have examined root adaptation along a precipitation gradient. Studies using congeneric and conspecific pairs occurring in high and low rainfall sites suggest that low specific root length (SRL, m g\(^{-1}\)), large root diameter, or slow root elongation rates may be favored under drier conditions, reflecting an investment in thick, longer-lived roots that more efficiently transport water (Wright and Westoby, 1999, Nicotra et al., 2002, Heschel et al., 2004).

We searched for evolutionary responses to N and water manipulations in two invasive grass species (*A. barbata* and *Bromus madritensis*) that are widespread throughout much of the west coast of North America. We subjected populations of both species to altered precipitation and N availability for five years, collected seeds from multiple maternal plants, and grew them in a common environment over two generations. We measured a suite of above- and below-ground traits that are associated with water and N use to address this question: Did invasive species exhibit trait differences in response to the 5-year environmental manipulations that persisted after two generations in a common environment? If so, our results would suggest that the trait differences were genetic and, thus, the result of rapid evolution in response to environmental change. Answering this question is a critical first step in understanding how invasive species may respond to future climate change.
Materials and Methods

Study site & seed source

The study site was located in a grassland community within the Irvine Ranch Conservancy in Orange County, California (Kimball et al., 2014, Allison et al., 2013, Potts et al., 2012). Orange County has a Mediterranean climate characterized by hot, dry summers and cool, wet winters. We focused on two invasive grass species, *A. barbata* and *B. madritensis*, which were abundant at our site. These species are native to the Mediterranean Basin and, due to repeated introductions, have widespread global distributions. In California, where our study was conducted, *A. barbata* and *B. madritensis* have overlapping but slightly different geographic distributions, with *B. madritensis* occurring in more arid environments including the Mojave Desert (DeFalco et al., 2003, Steers et al., 2011).

Twenty-four experimental plots (6.7 m × 9.3 m) extending over an area roughly one acre in size were imposed on existing vegetation, with each plot randomly exposed to one of three precipitation treatments (ambient, water addition, or water reduction, n=8 per treatment, Fig. 1). Each plot was divided lengthwise with each half receiving one of two N treatments (ambient or N addition). The water reduction treatment received approximately 51% less water than the ambient water treatment while the water addition treatment received approximately 33% more over the five year manipulation period (March 2007-May 2012) (Kimball et al., 2014). Clear, retractable roofs were deployed during a subset of winter storms to control water input into the reduction plots. Rainfall collected from reduction plots was stored in opaque polyethylene tanks and used to supplement water addition plots through a system of gasoline-powered pumps connected to drip tubing (Kimball et al., 2014). Baseline N deposition at our study site is
Nitrogen addition plots received 6 g N m$^{-2}$ year$^{-1}$, a value chosen to simulate a site approaching N saturation after years of continuous deposition (Kimball et al., 2014). Supplemental N was given in two applications: 2 grams of quick-release calcium nitrate following the first storm of the season and 4 grams of slow-release calcium nitrate following the first month of the growing season (Kimball et al., 2014).

In May 2012, seeds were collected from randomly selected maternal plants of A. barbata and B. madritensis and transported to Chapman University, which is located 16 km from the field site and has similar weather conditions. To ensure representation of the entire population, and minimize the effect of genetic drift, we tried to sample individuals from all plots (n=8) per watering treatment. However, some plots did not contain individuals of both species. When a species was present in a given plot, we collected seeds from one to eight maternal plants in that plot. Only seeds mature enough to be removed with a gentle pull of the spikelet were collected. Separate envelopes were used to ensure offspring from each mother plant remained isolated. On December 3, 2012, we germinated seeds on moist filter paper at 4°C for 96 hours. After germination, only one seedling from each mother plant was used; thus, each replicate had a different mother. Because later analyses showed that there was no significant effect of N on any of the traits measured, individuals from the two N treatments were grouped within each water treatment. We had at least nine replicates per water treatment for B. madritensis and at least 20 replicates per water treatment for A. barbata (Table 2). Following germination, seedlings were transferred to 4.1L pots filled with moistened potting soil (Sunshine Mix #1, Sun Gro Horticulture). Plants were grown in full sun at Chapman University and watered daily to saturation.
Because adaptation cannot be assumed based on the phenotypes displayed in one generation (Turner et al., 2014), common garden experiments must occur over multiple generations in order to differentiate between genetic and epigenetic effects, including maternal effects (Moran and Alexander, 2014). Maternal effects refer to a situation in which an individual’s phenotype is determined not only by its genotype and current environment, but by the environment experienced by its mother (Lacey, 1998). We controlled for maternal effects by growing seeds collected from the field (F1) and their offspring (F2) in a common environment (see Fig. 1). Any maternal effects present in the F1 generation should have been eradicated in the F2 generation because F1 plants were grown under identical environmental conditions. Thus, treatment differences observed in the F2 generation should be the result of genetic differences rather than maternal effects. We collected seeds from each F1 mother plant and germinated them as described above so that each F2 replicate had a different mother. Because of some mortality during germination and establishment, we had slightly fewer replicates in the F2 generation (n=67 A. barbata, n=42 B. madritensis) than in the F1 generation (n=72 A. barbata, n=43 B. madritensis) across treatments.

Above-ground measurements

Collection of physiological measurements began 70 days after germination of the F2 generation (February 2014). We conducted gas exchange measures with a LI-6400 portable photosynthesis system (LI-COR, Lincoln, NE, USA). We maintained constant chamber conditions that approximated growing conditions, including CO$_2$ at 400 $\mu$L L$^{-1}$, light at 1800 $\mu$mol photon m$^{-2}$ s$^{-1}$, chamber temperature at 25 °C, and relative humidity at 52-62%. Measures

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included photosynthetic rate, stomatal conductance, transpiration rate, and instantaneous WUE (photosynthetic rate/transpiration rate). When leaves were too small to fill the chamber, the leaf area was determined and used to correct gas exchange data.

We measured leaf chlorophyll content with a portable chlorophyll SPAD-502 meter (Spectrum Technologies, Plainfield, IL, U.S.A.). Three leaves were harvested following gas exchange and chlorophyll measures to determine leaf dry mass per unit area (LMA) and leaf N concentration. The leaves were scanned to determine total leaf area, dried at 60 °C for 72 hours, and weighed. The dried leaves were then ground in a Wiley mill with a 40 mesh screen and leaf N was determined with an elemental analyzer (Costech 4010, Pioltello, Italy).

We assessed plant size by measuring canopy cover prior to flowering, estimated as the product of plant width along two perpendicular axes. We checked plants daily and recorded date of flowering, defined as the first appearance of florets. While both species largely reproduce by selfing in natural populations (Johansen-Morris and Latta, 2006, Grossman and Rice, 2014), outcrossing does occur; thus, plants from different treatments were grown in the same area but with physical barriers between them to avoid cross-fertilization. Though spatially separated, all treatments received similar light and water levels daily. From March to June 2014, mature seeds were collected, air dried, and weighed to determine total seed biomass as a metric of reproductive fitness.

Root measurements

To obtain root measures for the F2 generation, seeds from a subset of F1 plants from the aboveground trait survey were germinated using the procedure described above and transferred...
to 4.1L pots filled with a moistened sand-perlite-vermiculite (1:1:1) mixture. Plants were randomly selected within plots and, as with above-ground measures, we sampled from all plots (n=8) per watering treatment where possible. We had eight replicates per watering treatment for each species. Seedlings were grown for five weeks, receiving a total of 0.05 g N, 0.05 g P, and 0.05 g K (MiracleGro) over the first two weeks. Plants were watered daily to saturation except on days when pots received fertilizer.

After five weeks, plants were harvested and roots were washed and scanned using the WinRHIZO image analysis system (Regent Instruments Inc., Quebec, QC, Canada) to determine total root length. Root growth rate (cm per day), root length density (root length per volume of soil), and fine root ratio [the ratio of fine root length (diameter <0.5mm) to total root length] were calculated. We separated below and above ground biomass, dried them at 60°C for 72 hours, and weighed them to determine root to shoot biomass ratio (R:S) and specific root length (SRL, m g⁻¹). Biomass was combined, ground and analyzed for N concentration as described above. Plant N uptake rate (g N per day) was calculated as [(plant biomass x plant N)/growing day number].

Statistical analyses

Box-Cox transformations of the data were used where necessary to meet the assumptions of normality for statistical analyses. We performed analysis of variance (ANOVA) with water and N treatment as fixed effects and plot as a random effect to compare trait values between treatments within each species. The significance of fixed effects was assessed using Satterthwaite approximated degrees of freedom. We used type III sums of squares ANOVA for
a mixed effects model in the LME4 package in R (http://www.r-project.org, v.3.2.0). Following Moran (2003), all P-values are reported and sequential Bonferroni corrections for multiple statistical tests were not conducted. Post hoc analyses were performed using Tukey’s honestly significant difference.

Results

There was no significant effect of N on any trait; thus, data from the two N treatments were grouped within a water treatment and N as a main effect was excluded from all subsequent analyses. Water availability affected several aboveground traits in A. barbata as well as first flowering date and water-use efficiency in B. madritensis (Table 1). With respect to physiological traits, photosynthetic rate and leaf chlorophyll content in A. barbata differed across water treatments with a trend toward higher trait values in the water reduction treatment relative to the ambient and added precipitation treatments (Table 1, Fig. 2). Photosynthetic rate and chlorophyll content did not differ across treatments in B. madritensis (Table 1, Fig. 3). Water-use efficiency was lower in plants from the reduced precipitation plots in B. madritensis compared to the ambient and water addition treatments (Table 1, Fig. 3). There was no difference in WUE across treatments in A. barbata (Fig. 2). The pattern observed in B. madritensis was driven by higher transpiration rates in plants from reduced precipitation plots (data not shown). Water had no significant effect on leaf N content in either species (Table 1, Fig. 2).

With respect to growth and reproductive traits, flowering date was earlier in the water reduction and ambient water treatments relative to the water addition treatment in both A. barbata and B. madritensis (Fig. 2,3; Table 1). Plants from the water reduction plots in A. barbata and B. madritensis.
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earlier flowering time or a shift in allocation from vegetative to reproductive structures (Wolfe and Tonsor, 2014), and could increase the number of successful offspring in arid environments, where establishment is difficult.

Notably, we found that field manipulations did not alter root traits, which is largely unstudied outside of model organisms (reviewed in Comas et al., 2013). Previous studies of root adaptation in natural systems have focused on conspecific or congeneric comparisons of species from different rainfall environments, which reflect evolution over potentially long timescales (Wright and Westoby, 1999, Nicotra et al., 2002, Heschel et al., 2004). It is possible that our five-year experiment was too short to observe differences in root traits. In the absence of rapid genetic responses, plasticity in root traits may be an important mechanism for responding to environmental changes on short time scales. In response to reductions in water availability, individual plants can increase root biomass, root elongation rate, and SRL to maximize water acquisition (e.g., Williams and Black, 1994, Padilla et al., 2009, Drenovsky et al., 2012, Larson and Funk, 2016). Several of these responses (rapid elongation, high SRL) are contrary to strategies found in species adapted to low rainfall environments (Wright and Westoby, 1999, Nicotra et al., 2002). More data are needed to understand the relative role of plasticity and genetic adaptation in belowground trait responses to variation in water availability.

In contrast to water manipulation, nitrogen addition over a five-year period did not result in detectable trait differences. Previous work from these field plots suggested that N addition only increased grassland productivity when combined with water addition, and only in some years (Goulden et al., unpublished data). It is possible that the proximity of the subplots (ambient and N addition within a single plot) may explain why N addition failed to alter traits in either species. Cross-pollination or seed dispersal may have contributed to gene flow among...
subplots. While many studies have found that exotic annual grasses respond positively to N addition (e.g., Huenneke et al., 1990, Davis et al., 2000, Gross et al., 2005), growth rates may saturate or decline at high soil N (e.g., Padgett and Allen, 1999, Mattingly and Reynolds, 2014). Our site is relatively fertile (0.19% N and 628.3 mg P kg\(^{-1}\); Funk, unpublished data), likely resulting from historic cattle grazing and N deposition in the Orange County area. Thus, N manipulations may yield different results if conducted in more N-limited environments.

One novel implication of our study is that it may be possible for invasive plant species to adapt to environmental change over just five years or five generations. Although there are examples of rapid evolution in other weed species (Franks et al., 2007, Frenck et al., 2013, Sultan et al., 2013, Grossman and Rice, 2014), five years is a remarkably short time period based on data from other studies. In a review of evolutionary responses of invasive species to novel environmental conditions, Moran and Alexander (2014) suggested that significant evolutionary changes in relevant traits are not likely to be observed in less than 25 generations. For example, Turner et al. (2014) confirmed that rapid evolution in growth and reproductive output has occurred in populations of Centaurea diffusa in the invaded range, which were separated from native populations by 100 years, or approximately 50 generations. Similarly, Dlugosh and Parker (2008) found evidence for rapid evolution of growth rate, date of first flowering, and size of Hypericum canariense in what they predicted to be 25 generations following introduction.

Our results suggest that populations of annual invasive grasses can show significant changes in response to environmental conditions in as little as five generations.

Higher rates of carbon assimilation, earlier phenology, and higher reproductive output in weedy species under conditions of reduced precipitation could have important implications at the community level. In Mediterranean climate ecosystems, such as our southern California site,
rainfall occurs almost exclusively during the winter months (November to May) and drought often brings reduced rainfall and/or a shorter rainfall period. Invasive species that are able to complete their life cycle during a shorter wet season while maintaining or, in our case, enhancing reproductive output (A. barbata) will have an advantage over species that have longer life cycles and are adversely affected by drought. Although A. barbata may be more evolutionarily responsive to water manipulation than B. madritensis (displayed more phenotypic responses), differences in absolute trait values suggest that B. madritensis was better at executing a drought escape strategy than A. barbata (lower LMA and WUE, higher total reproductive biomass and leaf N content compared to A. barbata; Fig 2,3). Consequently, B. madritensis was one of the most common species in reduced precipitation plots after five years (Goulden et al., unpublished data). While we did not measure trait values in co-occurring native species, the plot-level community composition data suggest that native grassland species may have limited potential to respond to drought on short timescales. Future research should test for adaptive evolution in co-occurring native and invasive species to assess the potential impact that rapid evolution may have on community-level processes and ecosystem function.

**Acknowledgements**

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Data Accessibility

-F2 generation root traits figure: Uploaded as online Supporting Information

-F2 above- and below-ground trait data: Zenodo doi: 10.5281/zenodo.46049

References


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**Supporting Information**

Additional Supporting Information may be found in the online version of this article:

**Figure S1.** Effects of water availability on root traits.

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Table 1. F-values calculated from one-way ANOVA for second generation (F2) plants of two invasive grass species with water treatment as a fixed factor and plot as a random factor.

Numerator and denominator degrees of freedom for F-values are presented in parentheses.

Significant effects ($P < 0.05$) are in boldface type.

<table>
<thead>
<tr>
<th></th>
<th>Avena barbata</th>
<th>Bromus madritensis</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>$F$</td>
<td>$P$</td>
</tr>
<tr>
<td><strong>Aboveground traits</strong></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Photosynthetic rate</td>
<td>3.82 (2,18)</td>
<td><strong>0.041</strong></td>
</tr>
<tr>
<td>Leaf chlorophyll</td>
<td>5.09 (2,67)</td>
<td><strong>0.009</strong></td>
</tr>
<tr>
<td>WUE</td>
<td>0.18 (2,24)</td>
<td>0.835</td>
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<tr>
<td>Leaf N content</td>
<td>2.62 (2,64)</td>
<td>0.081</td>
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<tr>
<td>LMA</td>
<td>0.23 (2,66)</td>
<td>0.793</td>
</tr>
<tr>
<td>Canopy cover</td>
<td>3.32 (2,67)</td>
<td><strong>0.042</strong></td>
</tr>
<tr>
<td>Flowering date</td>
<td>8.04 (2,67)</td>
<td>&lt;0.001</td>
</tr>
<tr>
<td>Seed biomass</td>
<td>8.92 (2,67)</td>
<td>&lt;0.001</td>
</tr>
<tr>
<td><strong>Belowground traits</strong></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Root growth rate</td>
<td>0.49 (2,24)</td>
<td>0.617</td>
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<tr>
<td>R:S</td>
<td>0.30 (2,24)</td>
<td>0.747</td>
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<tr>
<td>Root length density</td>
<td>0.47 (2,24)</td>
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<tr>
<td>Fine root ratio</td>
<td>0.93 (2,12)</td>
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<tr>
<td>Specific root length</td>
<td>0.57 (2,17)</td>
<td>0.576</td>
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<tr>
<td>N uptake rate</td>
<td>0.62 (2,24)</td>
<td>0.545</td>
</tr>
</tbody>
</table>

Abbreviations: Water-use efficiency (WUE), leaf mass per area (LMA), root to shoot biomass ratio (R:S)
Table 2. The number of individuals of *A. barbata* and *B. madritensis* sampled for each water treatment. The number of different plots that maternal plants originated from is given in parentheses. Because there was no significant effect of N on any trait, individuals from the two N treatments were grouped within their respective water treatments.

<table>
<thead>
<tr>
<th></th>
<th><em>A. barbata</em></th>
<th><em>B. madritensis</em></th>
</tr>
</thead>
<tbody>
<tr>
<td>Reduction</td>
<td>26 (7)</td>
<td>18 (8)</td>
</tr>
<tr>
<td>Ambient</td>
<td>21 (8)</td>
<td>15 (5)</td>
</tr>
<tr>
<td>Addition</td>
<td>20 (8)</td>
<td>9 (3)</td>
</tr>
</tbody>
</table>
Figure 1. Following five years of water manipulations in the field (Irvine, CA), seeds were collected from experimental plots and grown in a common environment over two generations. Physiological, morphological, and phenological traits were measured on the second generation grown in a common environment.
Figure 2. Photosynthetic rate (a), water-use efficiency (b), leaf nitrogen content (c), and leaf chlorophyll content (d), first flowering date (e), leaf mass per area (f), seed biomass (g), and canopy cover (h) for *A. barbata* from three watering treatments grown in a common environment (F2 generation). Means and standard errors of water addition, ambient, and reduction treatments are shown. Different letters denote significant differences among means as determined through post-hoc Tukey HSD tests. Where applicable, significant effects of water are noted.
Figure 3. Photosynthetic rate (a), water-use efficiency (b), leaf nitrogen content (c), and leaf chlorophyll content (d), first flowering date (e), leaf mass per area (f), seed biomass (g), and canopy cover (h) for B. madritensis from three watering treatments grown in a common environment (F2 generation). Means and standard errors of water addition, ambient, and reduction treatments are shown. Different letters denote significant differences among means as determined through post-hoc Tukey HSD tests. Where applicable, significant effects of water are noted.