Life history evolution and phenotypic plasticity in parasitic eyebrights (*Euphrasia*,

Orobanchaceae)

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ABSTRACT

Premise of the study: Parasite lifetime reproductive success is determined by both genetic variation

and phenotypically plastic life history traits that respond to host quality and external environment.

Here, we use the generalist parasitic plant genus *Euphrasia* to investigate life history trait variation, in

particular whether there is a trade-off between growth and reproduction, and how life history traits are

affected by host quality.

Methods: We perform a common garden experiment to evaluate life history trait differences between

eleven *Euphrasia* taxa grown on a common host, document phenotypic plasticity when a single

Euphrasia species is grown on eight different hosts, and relate our observations to trait differences

recorded in the wild.

Key results: *Euphrasia* exhibit a range of life history strategies that differ between species that

transition rapidly to flower at the expense of early season growth, and those that invest in vegetative

growth and delay flowering. Many life history traits show extensive phenotypic plasticity in response

to host quality and demonstrate the costs of attaching to a low-quality host.

Conclusions: Common garden experiments reveal trait differences between taxonomically complex

Euphrasia species that are characterised by postglacial speciation and hybridisation. Our experiments

suggest life history strategies in this generalist parasitic plant genus are the product of natural

selection on traits related to growth and flowering. However, host quality may be a primary

determinant of lifetime reproductive success.

Keywords: flowering time; host range; life history evolution; parasitic plants; phenotypic plasticity

BACKGROUND

Parasitism is a ubiquitous feature of the natural world, with parasitic organisms present in every ecosystem and found to exploit all free-living organisms (Price, 1980; Windsor, 1998). The life history of parasitic organisms, in particular their developmental timing, must be synchronised with that of the host to maximise a parasite's individual growth, survival and fecundity. However, the life history of external parasites (exoparasites) must not only be timed to their host, but to external environmental conditions (Fain, 1994). Changes in the developmental timing of a parasite will primarily be plastic when responding to local conditions or fluctuating environments, or genetic in response to more predictable or spatially structured variation (Reed et al., 2010). Understanding how genetic and plastic differences contribute to life history trait variation in parasite species and populations is essential for understanding parasite speciation and host-parasite interactions (Birget et al., 2017).

Parasitic plants are a group of c. 4500 species of at least 12 separate evolutionary origins that have evolved a modified feeding organ, the haustorium, which allows them to attach to a host plant and extract nutrients and other compounds (Westwood, Yoder, and Timko, 2010; Twyford, 2018). Parasitic plants are morphologically diverse and present a broad range of life history strategies (Schneeweiss, 2006; Těšitel, Plavcová, and Cameron, 2010) with the host species range a critical trait that determines parasite interactions and performance in the wild. Generalist parasitic plants can often attach to a broad range of hosts, with the well-studied grassland parasite *Rhinanthus* found to attach to over 50 co-occurring grass and herbaceous species (Cameron, Coats, and Seel, 2006). All generalist parasitic plants are exoparasites whose leaves, stems, roots and flowers grow outside the host and only the haustorium invades and grows within the host (Twyford, 2017).

The range of hosts and their visible external growth make hemiparasitic plants ideal for investigating life history trait evolution and host-parasite interactions. To date, research has largely focused on three aspects of life history variation in parasitic plants. Firstly, a body of research has looked to

understand variation for specific traits between populations and related species. For example, work on

the hemiparasite *Pedicularis* has shown how investment in male reproductive organs primarily

depends on extrinsic environmental conditions rather than intrinsic resources (Guo, Mazer, and Du,

2010a), and how seed mass depends on intrinsic factors rather than elevation (Guo, Mazer, and Du,

2010b). Secondly, researchers have investigated how life history traits are affected by host quality. In

the widespread and weedy obligate hemiparasite *Phelipanche ramosa*, the duration of the lifecycle

differs between 14 weeks and 40 weeks depending on host (Gibot-Leclerc et al., 2013), with evidence

of local host adaptation. Finally, a number of studies have looked at life history variation between

species studied in a phylogenetic context (Schneeweiss, 2006; Těšitel et al., 2010). For example,

broad-scale analyses of the Rhinantheae clade in the Orobanchaceae has shown a shift from a

perennial ancestor to annuality, with correlated shifts to a reduced seed size (Těšitel et al., 2010).

Despite the diversity of this research, there are still considerable gaps in our knowledge as to how life

history trait variation is maintained (e.g. how common are trade-offs between life history traits), how

much of this variation is genetic and how much is plastic, and which traits are the targets of natural

selection.

In this study, we explore life history trait evolution in generalist parasitic eyebrights (*Euphrasia*,

Orobanchaceae). *Euphrasia* is the second largest genus of parasitic plants with c. 350 species, and is

characterised by recent transoceanic dispersal and rapid species radiations (Gussarova et al., 2008). In

the United Kingdom, the 21 *Euphrasia* species demonstrate recent postglacial divergence with many

species indistinguishable at DNA barcoding loci (Wang et al., 2018), with complex morphological

variation (Yeo, 1968; Metherell and Rumsey, 2018), and found to readily hybridise (Stace, Preston,

and Pearman, 2015). Despite this shallow species divergence, *Euphrasia* species demonstrates

substantial ecological divergence, with many taxa restricted to specific habitats such as coastal turf,

mountain scree or open grassland. Habitat differences would be expected to exert strong selection on

life history traits, and this may include selection on growth to match seasonal water availability and to

exploit local hosts, or selection on flowering time in response to local competition from surrounding

plants, or in response to mowing or grazing.

Our research builds on a large body of experimental work with *Euphrasia*, which has been used in

common garden studies for over 125 years (Koch, 1891). Early growth experiments revealed

phenotypic differences observed in the field between the related species *E. rostkoviana* and *E.*

montana are maintained in a common garden (Wettstein, 1895). Experimental work in the 1960s

showed the growth of various *Euphrasia* species differs depending on the host species (Wilkins, 1963;

Yeo, 1964). More recent replicated pot experiments of individual *Euphrasia* species or populations in

a common garden (Matthies, 1998; Zopfi, 1998; Lammi, Siikamäki, and Salonen, 1999; Svensson and

Carlsson, 2004) or in experimental field sites (Seel and Press, 1993; Hellström et al., 2004), have

shown the effect of commonly encountered hosts such as grasses and legumes on parasite biomass,

mineral accumulation, plant architecture and reproductive output. Despite this extensive experimental

work, studies in *Euphrasia* have yet to compare life history strategies of different species, and the

extent of phenotypic plasticity in life history traits. This work is critical for understanding the nature

of species differences in a taxonomically complex group, as well as for evolutionary studies of

parasite diversity. It also unclear whether *Euphrasia* are restricted to growing on hosts such as grasses

and herbaceous species, or can parasitize a broad range of taxa including novel hosts rarely

encountered in the wild. To address these questions requires simultaneously investigating the growth

of multiple *Euphrasia* species and multiple host species with replication to enable suitable statistical

comparisons.

Our experiments first assess the extent of life history trait variation among several *Euphrasia* species

and their hybrids when grown on a single host species in standardised common garden conditions.

This experiment addresses whether there is life history trait divergence among recently diverged

parasite species. We then inspect plasticity of a single focal *Euphrasia* population grown on many

different hosts. This experiment tests whether they are truly generalist parasites by growing them on a

wide range of hosts as well as growing them without a host. Finally, we relate our trait observations

made in a common garden to recordings made on herbarium specimens collected in the wild. This

comparison will help us understand whether life history trait differences hold in both the common

garden and in nature. Overall, our joint observations of phenotypic variation between closely related

taxa, and the extent of host-induced plasticity within a species, both in an experiment and in the wild,

provide new insights into life history variation in parasitic taxa. These results provide valuable

comparisons to other generalist parasitic plants such as *Rhinanthus* and *Melampyrum*, to see

similarities in life history across grassland parasitic plants.

MATERIALS AND METHODS

Experimental design and plant cultivation

We performed two common garden experiments to investigate phenotypic variation in parasitic

Euphrasia. Our species differences experiment observed life history trait differences of twenty four

populations from five species and six natural hybrids when grown on clover (*Trifolium repens*), which

is often used as a host in parasitic plant experiments because the parasite grows vigorously and has

high survival (Zopfi, 1998). This experiment includes multiple populations of three widespread and

closely related species, *E. arctica*, *E. confusa* and *E. nemorosa*, and sparse population sampling of

two more distinct taxa, *E. micrantha* (one population) and *E. pseudokerneri* (two populations). Our

phenotypic plasticity experiment assessed the impact of eight potential hosts (*Arabidopsis thaliana*,

Equisetum arvense, *Festuca rubra*, *Holcus lanatus*, *Marchantia polymorpha*, *Pinus sylvestris*,

Plantago lanceolata, and *Trifolium repens*), and growing without a host, on the phenotype of a single

focal taxon, *E. arctica*. These hosts were chosen to represent a broad representation of functional

groups and phylogenetic diversity, with species encountered in the wild as well as novel hosts (full

details in Table S1). The novel hosts were included to see the limits to which parasitic *Euphrasia* can

benefit, namely with a woody tree (*Pinus*), a pteridophyte that produces adventitious roots

(*Equisetum*), and a liverwort that produces rhizoids (*Marchantia*). Both common garden experiments

took place in parallel in 2016. Wild-collected open-pollinated *Euphrasia* seeds were contributed by

plant recorders as part of the 'Eye for Eyebrights' (E4E) public engagement project and as such

included a scattered geographic sample across Great Britain (Table S2). Species were identified from

herbarium specimens by *Euphrasia* referee Chris Metherell. Host seeds were sourced from

commercial suppliers and from wild-collected material (Table S1).

Reliable cultivation of *Euphrasia* can be challenging due to low seed germination, variation in time to

establishment, the requirement of seed stratification, and high seedling mortality when transplanted

(Yeo, 1961; Zopfi, 1998). Here, we develop cultivation protocols that combine winter germination

cues that improve germination and mimic nature, but also use highly standardised and replicated pot

conditions that avoid transplanting *Euphrasia* and thus maximise survivorship. We planted a single

Euphrasia seed into 9 cm plastic pots filled with Melcourt Sylvamix Special growing media in

December, with plants left outside overwinter to experience natural seed stratification. Hosts were

planted in seed trays in April. *Euphrasia* plants were moved to an unheated and well-ventilated

greenhouse at the Royal Botanic Garden Edinburgh (RBGE) in the spring once the cotyledons were

fully expanded, and a single seedling from each host (or a 1cm^2 clump of *Marchantia*) introduced.

Hosts that died within ten days of planting were replaced. Twenty or more replicates were grown for

each host-parasite combination. Plants were subsequently grown to flowering with regular watering,

randomised at weekly intervals, and foreign weed seedlings removed.

Trait measurements and statistical analyses

We measured a suite of seven morphological traits at first flowering. These traits are characters

related to life history variation, indicators of plant vigour, or characters used in taxonomy. In addition

to date of first flowering, we measured: corolla length, cauline leaf length:internode length ratio

('internode ratio'), number of leaf teeth on the lower floral leaf, number of nodes to flower, number of

branches and plant height. All length measurements were made to the nearest millimetre, and

followed Metherell and Rumsey (2018). For the phenotypic plasticity experiment, we also recorded

early season growth (height six weeks after transplantation of potential host) and height at the end of

season after senescence. We did not make direct observations of below-ground attachment due to the

fine root structure of *Euphrasia*. Instead, we inferred that attachment is likely to have taken place

based on observations of height, following Yeo (1964).

We analysed data for the two experiments separately but with similar statistical approaches. For the

species differences experiment we used generalised linear mixed effects models to fit species as a

fixed effect and population as a random effect. Population was implemented as a random effect

because we were interested in the variability between populations, as well as to account for it for

estimating species parameters. For the phenotypic plasticity experiment we used generalised linear

mixed effects models with presence or absence of host as a fixed effect and host species identity as a

random effect. We used a Gaussian distribution for all models except where the response were

positive integer counts, where we used a Poisson distribution. To calculate the significance of a fixed

or random effect we used Likelihood Ratio Tests where the full model was compared with a nested

model where one effect had been dropped. When a random effect term was dropped for each variable,

we implemented generalised linear models using R base functions. For the trait number of branches

the generalised linear mixed effects models failed to converge so we used MCMCglmm (Hadfield,

2010), with the best fitting of either a full or nested model chosen based on a lower Deviance

Information Criterion (DIC) value. For the phenotypic plasticity experiment, models for number of

branches failed to converge in every model run, so this model was omitted. Using the flexible

variance structures accommodated by MCMCglmm we were able to calculate "variance explained"

by random effects in each model. In the case of a Gaussian model, variance explained by an

explanatory variable is the fraction of the total variance accounted for by the variable. In the case of a

Poisson model the calculation is more complex due to the mean-variance relationship (Nakagawa and

Schielzeth, 2010). The models were run for 91,000 iterations with a burn-in of 21,000 and a thinning

interval of 70. The priors were standard inverse-Wishart for the (co)variances and parameter

expanded where there was poor mixing. Overdispersion was accounted for in each case by either

implementing quasipoisson models or in mixed effect models by implementing an observation level

random effect. We calculated r^2 values from Pearson Correlation Coefficients between all traits and

used Principal Component Analyses to quantify phenotypic clustering of individuals. All analyses

were done in R version 3.4.3, with the packages lme4 (Bates et al., 2014) and MCMCglmm for

generalised linear mixed effects models, Hmisc for correlations (Harrell Jr and Harrell Jr, 2018) and

ggplot2 for data visualisation (Roff, 1992).

Trait variation in the wild

We tested how phenotypes in the experiments relate to phenotypes in nature, by comparing results

from the species differences experiment to phenotypic measurements made on herbarium specimens

of the same populations sampled in the wild. Pressed plants submitted by field collectors varied in

quality, and these sometimes missed the base of the plant, therefore we were unable to measure

height. We measured traits on three individuals from each collection sheet for a given population. We

used generalised linear mixed effects models to fit traits as a response variable, with where

individuals were grown (i.e. common garden or wild-collected) as an explanatory variable. This two-

levelled categorical factor was implemented as a fixed effect. Adding species as a random effect

shows how much of the variability of the trait was explained by species given where individuals were

measured. Count data were analysed with a Poisson distribution, in all other cases a Gaussian

distribution was used. R-values were calculated using Pearson's correlations of the population level

means between the common garden and the wild samples. Analyses were performed in MCMCglmm.

RESULTS

Species differences

Our species differences experiment reveals extensive morphological trait variation across *Euphrasia*

species when compared at first flowering. From the 222 individuals surviving to flowering on their

clover host, we see a 3-fold difference in mean height, while other traits also proved variable (Fig. 1e-

h; Table S3). A large degree of this variation is separated by species and by population (Table 1). The

species with the most distinct lift-history strategy is *E. micrantha*, which rapidly transitions to flower

from a low node on the plant (8.3 ± 0.2 nodes), and flowers while it is short (70.6 mm ± 8.1). It also

forms a distinct cluster in the PCA analysis (Fig. S1). *E. pseudokerneri* is relatively distinct, flowering

once it has grown tall (176.4 mm ± 15.6) and from a high node on the plant (13.2 ± 0.4 nodes), but

shows little separation in the PCA analysis. The morphologically similar *E. arctica*, *E. confusa* and *E.*

nemorosa are partly distinct, with *E. nemorosa* initiating flowering 14 days later and from 3.3 nodes

higher than *E. arctica*, but overlaps in many other traits and in overall multi-trait phenotype (Fig. S1).

In most cases hybrids combine morphological characters of their parental progenitors, for example

hybrids involving *E. nemorosa* flower later in the season and initiate flowering from a higher node

than *E. arctica* hybrids (Fig. 1a-d).

Correlation analyses across species reveal clear suites of traits that are related. Significant correlations

were found between 14 of the 21 pairwise comparisons, with 5 of these correlations with an R > 0.6

(Table 2a). Plants flowering at a late node are more likely to be tall, more highly branched, as well as

having many teeth on the lower floral leaf. The relationship of traits is also supported in the PCA

analysis, with many traits contributing to multiple principal components (Table S4). These height- and

flowering node-related traits are largely uncorrelated with cauline internode-leaf ratio and corolla

length.

Phenotypic plasticity

Our phenotypic plasticity experiment shows substantial morphological variation across 194 *E. arctica*

plants grown with 8 different host species, and the 22 plants grown without a host. Plants growing on

clover transitioned to flower quickly, grew tall by the time of first flowering, and produced large

flowers (Fig. 1e-h, Table S5). This contrasts with *Euphrasia* with no host, which flowered on average

53 days later, were extremely short at first flowering, and produced small flowers. Despite this, the

presence or absence of host was not significant (Table 1), due to *Euphrasia* grown with *Marchantia*

and *Pinus* growing similarly to the no host treatment (Fig. 1e-h). The other hosts, *Arabidopsis*,

Equisetum, *Festuca*, *Holcus* and *Plantago* conferred some benefit for *Euphrasia* and fell between

clover and no host. Our comparison of growth across host treatments measured through the year

showed that height at the end of the season is weakly predicted from height 6-weeks after introducing

a host (R = 0.47), but strongly correlated with height at first flowering (R = 0.82; Fig. S2). Plants that

flowered early were more likely to grow larger by the end of season (R = -0.55) and become more

highly branched (R = -0.57).

Across host treatments, there was a significant negative correlation between Julian days to flower and

most other traits (Table 2b). We find that late flowering individuals are likely to be smaller at first

flowering, are less highly branched, have less highly toothed leaves, and have smaller flowers (Table

S5). While these traits were strongly correlated, there were substantial differences in the amount each

trait changed. Days to flower differed considerably depending on host, with a 3.8x greater difference

than seen between means of different *Euphrasia* species (Fig. 1d & 1h). In contrast corolla length and

node to flower proved less variable depending on host, with a 1.4x and 1.2x change between means,

respectively.

Variation in the wild

The comparison between the species differences common garden experiment and wild-collected

herbarium specimens revealed a single trait, nodes to flower, is strongly correlated (R=0.79) and not

significantly different (p_{MCMC}=0.71) between environments (Fig. 2). All other traits did differ

significantly between environments ($p_{MCMC} < 0.05$), with *Euphrasia* plants in the common garden

having corollas on average 1.4mm longer, with 0.2 more teeth on the lower floral leaves, an increase

in cauline:internode ratio of 1.0mm, and 4 more pairs of branches. Despite differing between

environments, trait values were highly correlated for corolla length (R= 0.93, $p_{MCMC} < 0.001$) and

cauline internode:leaf length ratio (R=0.65, $p_{MCMC} < 0.001$), but not for number of leaf teeth (R=0.07,

$p_{MCMC} = 0.034$) and number of branches (R=0.29, $p_{MCMC} < 0.001$).

DISCUSSION

Our study sheds new light on the life history evolution and phenotypic plasticity of the generalist

parasitic plant *Euphrasia*. We find different life history strategies between recently divergent species,

with some species rapidly transitioning to flower at the expense of growth-related traits, while others

delay flowering and invest in early-season vegetative growth. However, many traits related to life

history differences are phenotypically plastic and show major changes in response to host quality. In

addition to our experimental observations, the consistency between life history traits in a common

garden and in the wild suggest our experimental observations generalise to patterns observed in

nature, and that currently delimited *Euphrasia* species are, at least in part, distinct. Overall our study

highlights the value in integrating multiple common garden experiments and field collections to study

life history strategies in parasitic plants, and demonstrates the rapid evolution of life history

differences in a postglacial radiation of parasites. We contrast our results with other well-studied

hemiparasitic taxa to look for general patterns of life history evolution in grassland parasites.

Life history variation in a parasitic plant

Our study finds evidence for different life history strategies between *Euphrasia* species in Britain. *E.*

arctica, *E. micrantha* and hybrids such as *E. arctica* x *E. confusa*, transition rapidly to flower, flower

while they are short, and produce their first flower from a low node on the plant. This contrasts with

E. pseudokerneri, *E. nemorosa* and hybrids involving *E. nemorosa* that delay flowering until later in

the season, grow tall before flowering, and produce their first flower from a late node on the main

axis. These different life history strategies correspond to the known of ecology of these species, with

E. nemorosa flowering late in tall mixed grassland, while *E. micrantha* flowers early in patchy

heathland where competition is less intense (Metherell and Rumsey, 2018). A similar relationship

between internode number and habitat have been observed in populations of *E. rostkoviana* in Sweden, where plants from intensely grazed pasture flower at a lower node in a common garden (Zopfi, 1998). Overall, these observations within and between populations are consistent with the classic life history trade-off between growth and reproduction (Roff, 1992; Stearns, 1992). For *Euphrasia* growing in the wild, early reproduction allows the plants to reliably complete their lifecycle before summer competition, herbivory, mowing, summer drought and other seasonal abiotic and biotic stresses. However early flowering involves reproducing at the expense of early season growth and at a time when the resource budget is constrained by relatively few haustorial connections. These trait trade-offs pose an interesting comparison to the well-studied *Mimulus guttatus* (*syn. Erythranthe guttata*), a non-parasitic relative in the Lamiales that shares the same basic plant architecture. In *M. guttatus* multiple traits related to growth and reproduction are correlated both within and between populations, due to genetic trade-offs between time to flower and fecundity (Mojica et al., 2012; Friedman et al., 2015). In *Euphrasia* the genetics underpinning this life history trade-off have yet to be characterised, and may be a consequence of multiple independent loci or trade-offs at individual loci (Hall, Lowry, and Willis, 2010).

While much life history variation is captured by differences in time to flower and growth-related traits, we also see evidence for flower size representing a separate axis of variation across *Euphrasia* species. In our common garden *E. micrantha* has small corollas, while *E. arctica* and *E. nemorosa* have larger corollas, and corolla size is not strongly correlated with other traits. *Euphrasia* species are well-known to have flower size variation, with a continuum between small flowered species that are highly selfing (e.g. *E. micrantha*, corolla size = 4.5 - 6.5mm, inbreeding coefficient $F_{IS} > 0.88$; Stone, 2013), and large flowered species that are highly outcrossing (e.g. *E. rostkoviana* flower size 8 - 12mm, $F_{IS} = 0.17 - 0.25$; French et al., 2005). Such continuous variation in outcrossing rate has been documented in species of *Datura* (Motten and Stone, 2000), *Mimulus* (Karron et al., 1997) and *Nicotiana* (Breese, 1959). Small flowers have shorter anther-stigma separation and thus increased potential for autogamous selfing (Karron et al., 1997), while also having reduced attractiveness to pollinators and thus receiving less outcross pollen (Mitchell et al., 2004). In addition to differences in

corolla size between *Euphrasia* species, corolla size also shows a change of up to two millimetres in

response to host quality, with this change of a magnitude that may potentially affect the mating

system (Luo and Widmer, 2013). This suggests host quality represents a previously unaccounted

factor affecting the mating system of parasitic plants.

Our comparisons of *Euphrasia* species in a common garden also sheds light on the distinctiveness of

these recently divergent species. *Euphrasia* is one of the most taxonomically complex plant genera,

with the 21 currently described British species presenting complex and often overlapping

morphological variation (French et al., 2008; Metherell and Rumsey, 2018; Wang et al., 2018). Our

study suggests varying degrees of morphological distinctiveness of *Euphrasia* species. We see *E.*

micrantha is clearly morphologically distinct and *E. pseudokerneri* mostly distinct, while the closely

related species *E. arctica*, *E. confusa* and *E. nemorosa* differ in life history traits such as nodes to

flower, but overlap in many other traits and are not clearly separated in the PCA. Moreover, our study

is likely to overestimate the distinctiveness of taxa by only including a subset of UK species and by

choosing populations that could be identified to species-level in the field. We suspect adaptive

divergence between closely related *E. arctica*, *E. confusa* and *E. nemorosa* is a consequence of

differential natural selection for local ecological conditions such as soil water availability or mowing.

Selection appears to be operating at a fine spatial scale, with significant life history trait differences

evident between populations within species. These taxa may be genetically cohesive, either showing

genome-wide divergence, or divergence in genomic regions underlying life history differences

(Twyford and Friedman, 2015). Alternatively, these taxa may be polytopic and not genetically

cohesive (Hollingsworth, Neaves, and Twyford, 2017). Genomic sequencing of natural populations is

hoped to resolve this issue.

Phenotypic plasticity in response to host quality

Our phenotypic plasticity experiment shows *Euphrasia* benefit from growing with a range of hosts.

Specifically, *E. arctica* with a high quality host such as clover rapidly transitions to flowering.

Differences in growth are established early in the season, and early flowering plants go on to grow the

tallest, are more highly branched and have the potential to produce many extra flowers. At the other

extreme, growing with a poor host or without a host results in late-flowering minimally vegetative

plants. Most hosts fall between these extremes and offer moderate return to the parasite. Two

surprising results were that *E. arctica* parasitizing *Arabidopsis* grew relatively tall despite the host

senescing early in the growth season, and that *Euphrasia* on *Equisetum* performed similarly to the

commonly encountered grass *Holcus lanatus*, suggesting surprising attachment or indirect benefits

through association with *Equisetum* fungal symbionts (Bouwmeester et al., 2007). Differences in host

quality are complex, but may be attributed to root architecture, germination time and resource

availability, as well as the presence of mechanisms to defend against parasite attack, such as cell wall

thickening, localised host dieback, and chemical defence (Cameron, Coats, and Seel, 2006; Twyford,

2018). Many of these attributes of host quality appear to be shared between related hemiparasites in

the Orobanchaceae, though contrasting results from *Rhinanthus* suggest there are not universal 'good'

hosts for all parasitic plants (Matthies, 2017). While *Euphrasia* is generally thought to have low

reliance on host resources, deriving only ~30% carbon heterotrophically (Těšitel, Plavcová, and

Cameron, 2010), at least under our experimental conditions a good host is required in order to

produce multiple flowers. Overall our results point to *E. arctica* being a true generalist parasite, but

one that only grows better with a subset of hosts that it may encounter.

We observe considerable variation in the plasticity of traits in response to host quality. Only three

pairs of trait correlations are consistent across both *Euphrasia* common garden experiments (between

height, number of branches and leaf teeth), with the positive correlation between node to flower,

height, and number of branches seen between species breaking down when *Euphrasia* are grown on

different hosts. The most notable plasticity is seen in flowering time, with plants on the best hosts

rapidly transitioning to flower within ~100 days of germination, while plants with a more typical host

(e.g. *Holcus lanatus*) flower a month later. Phenotypic plasticity in flowering time in response to

resource availability is well documented in many plant groups, particularly *Arabidopsis* (e.g. Zhang

and Lechowicz, 1994), but has received less attention in studies of parasitic plants, which are more

likely to look at growth-related traits such as biomass (Ahonen, Puustinen, and Mutikainen, 2005;

Matthies, 2017). However date of first flowering has been show to differ by up to 10 weeks in

populations of *Rhinanthus glacialis* across Switzerland (Zopfi, 1995). Overall we expect date of first

flowering to be critical for the life time reproductive success of parasitic plants in the wild.

In contrast to traits showing extensive plasticity, we also see evidence of developmental constraint in

number of nodes to flower. This trait showed the least plasticity with different hosts, is consistent

between populations within species, and between the common garden and the field. This suggests that

the developmental event of transitioning to flower is genetically determined, with changes in

flowering time altered by plasticity in internode length and not nodes to flower. This may explain why

nodes to flower is such an important diagnostic trait for species identification in *Euphrasia* and related

species in the Rhinantheae (Jonstrup, Hedrén, and Andersson, 2016). Despite nodes to flower

changing little in response to host quality, our overall impression is that *Euphrasia* show considerable

plasticity and little developmental constraint in many aspects of growth, or that trait values are

affected by complex interactions related to host-parasite attachment. In particular, the variation

between individuals on a given host, suggests other variation, such as genetic background in host and

parasite, as well as the timing of attachment, may be crucial in determining performance.

CONCLUSIONS

Despite over a century of experimental studies in parasitic plants, our understanding of the evolution

of life history strategies in these diverse organisms is extremely limited. Our results in *Euphrasia*

provide strong support for the rapid evolution of distinct life history strategies in response to local

ecological conditions, with phenotypic plasticity further altering plant growth in response to host

availability. We anticipate that future studies that test life time reproductive success of many parasitic

plant species grown on many different host species will give further insight into the complex nature of

host-parasite interactions.

Data accessibility

Phenotypic data from both common garden experiments and from herbarium collections, as well as the R scripts used for data analysis, are deposited in Dryad XXXXXX (accession to be given on acceptance).

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Author contributions

A.D.T conceived and designed the research. A.D.T., N.F. and E. L. Y.W. carried out the experiments. C.M. identified the plants. A.D.T. and M.B. analysed the data. A.D.T. and M.B. wrote the manuscript. All authors read and approved the manuscript.

Data accessibility

R scripts for analysis are included in the Online Supporting Information. All phenotypic data are available from Dryad <Details given on acceptance>.

Legends to figures and tables

Figure 1. Trait variation in a common garden experiment of diverse *Euphrasia* species and hybrids grown on clover (A-D) and *E. arctica* grown on many different hosts (E-H). The edges of the boxplots show the first and third quartiles, the solid lines the median, the whiskers the highest and lowest values within 1.5-fold of the inter-quartile range and the dots each individual measurement.

Figure 2. Relationship between morphological trait measurements made in the common garden and on wild-collected herbarium specimens for (A) nodes to flower, (B) corolla length (mm), (C) number of leaf teeth, (D) cauline leaf:internode ratio. Points are for populations, with bars representing the standard error of measurements. The line of best fit was calculated using coefficients from linear regression models on the means of each *Euphrasia* population.

Table 1. Summary of generalised linear mixed effects models of the influences on life history, plant vigour, and taxonomic traits for *Euphrasia* in a common garden. Table summarises model outputs for the species differences and the phenotypic plasticity experiment. The percentage variance explained by random effects are reported in brackets along with the 95% credibility interval. †Models for number of branches were implemented with a different statistical approach in MCMCglmm (see methods). *** $P < 0.001$, ** $P < 0.01$, * $P < 0.05$

Table 2. Pearson's correlation coefficients for seven phenotypic traits measured in a common garden experiment for (a) Five *Euphrasia* species and 6 hybrids, (b) *Euphrasia arctica* grown with 8 hosts and without a host. *** $P < 0.001$, ** $P < 0.01$, * $P < 0.05$. Asymptotic p-values values are reported from the Hmisc package in R using the rcorr() function.

Supporting information

Figure S1. Principal components analysis of morphological variation of *Euphrasia* in a common

garden for (A) five species and six hybrids, (B) five species, (C) *Euphrasia arctica* with nine host

treatments.

Figure S2. Relationship between early and midseason growth and flowering related traits and end of

season height. (A) Height at first flowering, (B) height 6-weeks after germination, (C) Julian days to

flower, (D) number of branches. Length measurements are reported in mm.

Table S1. Species and collection details for hosts used in the common garden experiment.

Table S2. Collection details for *Euphrasia* species used in the common garden experiment.

Table S3. Summary of trait values for many *Euphrasia* species and hybrids grown on a clover host.

Values are mean +/- one standard error. Length measurements are in mm.

Table S4. Factor loadings for the principal components analyses of (A) five species and six hybrids,

(B) five species, (C) *Euphrasia arctica* with nine host treatments.

Table S5. Summary of trait values for *Euphrasia arctica* grown on many different hosts. Values are

mean +/- one standard error. Length measurements are in mm.

Table 1

	Species differences		Phenotypic plasticity	
	Species	Population	Host/no host	Host
Branches†	DIC_{full} = 867.36; DIC_{-spp} = 869.11	DIC_{full} = 867.36; DIC_{-pop} = 905.68 (27.7%, 16.4 - 61.2%)	n/a	n/a
Corolla length	$\chi^2(1) = 22.38^{*}$	$\chi^2(1) = 42.89^{***}$ (49.1%, 28.8 - 75.5%)	$\chi^2(1) = 3.19$	$\chi^2(1) = 28.79^{***}$ (13.1%, 5.6 - 52.3%)
Height	$\chi^2(1) = 20.10^{*}$	$\chi^2(1) = 67.68^{***}$ (57.2%, 37.3 - 80.2%)	$\chi^2(1) = 0.9033$	$\chi^2(1) = 91.15^{***}$ (54.1%, 35.9 - 85.5%)
Internode ratio	$\chi^2(1) = 24.5^{**}$	$\chi^2(1) = 27.39^{***}$	$\chi^2(1) = 1.38$	$\chi^2(1) = 6.94^{**}$ (4.6%, 0.8 - 23.0%)
Julian days to flower	$\chi^2(1) = 8.34$	$\chi^2(1) = 1.34$ (21.2%, 7.7 - 64.6%)	$\chi^2(1) = 2.34$	$\chi^2(1) = 205.2^{***}$ (37.4%, 13.4 - 66.3%)
Node to flower	$\chi^2(1) = 26.66^{**}$	$\chi^2(1) = 0.71$ (10.1%, 0.4 - 31.6%)	$\chi^2(1) = 0.38$	$\chi^2(1) = 0$ (1%, 0.18 - 9.8%)
Number of leaf teeth	$\chi^2(1) = 23.43^{*}$	$\chi^2(1) = 0$	$\chi^2(1) = 1.49$	$\chi^2(1) = 5.46^{*}$

Table 2

(a) Species differences experiment

	Corolla length (mm)	Height (mm)	Internode ratio	Julian days to flower	Leaf teeth	Nodes to flower
Branches	0.260 ***	0.609***	-0.116	0.057	0.658***	0.775***
Corolla length (mm)		0.319***	-0.161*	-0.127	0.197 ***	0.049
Height (mm)			0.246***	0.292***	0.563***	0.628***
Internode ratio				0.204*	-0.120	0.076
Julian days to flower					0.053	0.249***
Leaf teeth						0.651***

(b) Phenotypic plasticity experiment

	Corolla length (mm)	Height (mm)	Internode ratio	Julian days to flower	Leaf teeth	Nodes to flower
Branches	0.524***	0.834***	-0.299***	-0.572***	0.694***	-0.572***
Corolla length (mm)		0.503***	0.098	-0.406***	0.536***	-0.166*

Height (mm)	0.477***	-0.481***	0.692***	-0.186**
Internode ratio		-0.034	0.168*	-0.009
Julian days to flower			-0.691***	0.530***
Leaf teeth				-0.239***

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Species differences experiment

Phenotypic plasticity experiment

A — Height (mm)

B — Nodes to flower

C — Corolla length (mm)

D — Julian days to flower

E

F

G

H

E. arctica, *E. confusa*, *E. micrantha*, *E. nemorosa*, *E. pseudokerneri*, *E. anglica x nemorosa*, *E. anglica x rostkoviana*, *E. arctica x confusa*, *E. arctica x nemorosa*, *E. confusa x nemorosa*, *E. confusa x tetraquetra*

No host, *M. polymorpha*, *E. arvense*, *P. sylvestris*, *F. rubra*, *H. lanatus*, *A. thaliana*, *P. lanceolata*, *T. pratense*