1	Adapted dandelions increase seed dispersal when they are attacked by
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#### 27 Abstract

28 Plants allow their offspring to escape unfavourable local conditions through seed 29 dispersal. Whether plants use this strategy to escape herbivores is not well understood. Here, 30 we explore how different Taraxacum officinale populations modify seed dispersal in response 31 to root herbivore attack by *Melolontha melolontha* in the field. Root herbivore attack increases 32 seed dispersal potential through a reduction in seed weight in populations that have evolved 33 under high root herbivore pressure, but not in populations that have evolved under low 34 pressure. This increase in dispersal potential is associated with reduced germination, 35 suggesting that adapted plants trade dispersal for establishment. Analysis of vegetative growth 36 parameters suggests that increased dispersal is not the result of stress flowering. These results 37 suggest that root herbivory selects for genotypes that increase their dispersal ability in response 38 to herbivore attack.

#### 39 Introduction

40 As sessile organisms, plants are bound to grow and develop where they germinate. 41 Local conditions are therefore important determinants of plant survival and reproductive 42 success (Kawecki & Ebert 2004, Walter et al. 2018). Although the ability of plants to move is 43 limited, they can influence the displacement capacity of their offspring by modifying dispersal 44 traits (Nathan & Muller-Landau 2000, Martorell & Martínez-López 2014, Teller et al. 2014). 45 Theory suggests that whenever local conditions are unfavorable, increased seed dispersal is 46 advantageous, as it allows the next generation to escape from the unsuitable environment and 47 increases their chance to colonize new habitats with better conditions (Levin et al. 2003, Ronce 48 2007). Major dispersal strategies of plants include dispersal by wind (anemochory), water 49 (hydrochory) and animals (zoochory) (Poschlod et al. 2013).

50 Herbivores can have strong impacts on plant development and fitness and thus act as 51 a force of natural selection (Rutter & Rausher 2004, Lau et al. 2008, Agrawal et al. 2012, Züst 52 et al. 2012). To date, the impact of herbivores on plant dispersal remains poorly understood 53 (de la Peña et al. 2011). Compared to aboveground herbivores, belowground herbivores are 54 generally more limited in their mobility due to the dense nature of soil and are therefore often 55 distributed in a patchy way (Van der Putten et al. 2001). Hence, seed dispersal may allow plants 56 to escape belowground enemies, which may lead to selection for high dispersal genotypes by 57 root herbivores (Van der Putten 2003). Experimental evidence for this hypothesis is currently 58 lacking. Understanding how herbivores influence seed dispersal and establishment traits would 59 help to elucidate their impact on plant population dynamics. In general, herbivores may affect 60 plant dispersal strategies either by selecting for genetically fixed high dispersal genotypes or 61 by favoring genotypes that are phenotypically plastic and increase dispersal in response to 62 herbivore attack. Especially in stochastic environments with fluctuating herbivore pressure, 63 inducible increase of seed dispersal distance as plastic response to herbivory may be 64 advantageous (Jakobsson & Eriksson 2003).

Trade-offs between dispersal capacity and seed viability are frequent in nature. For distribution processes of wind-dispersed plant species, seed mass is often reported as a critical trait that influences both dispersal and seedling establishment. Lower seed mass typically increases dispersal (Meyer & Carlson 2001; Greene & Quesada 2005; Skarpaas et al. 2011),

69 but reduces germination and early growth (Leishman & Westoby 1994; Turnbull et al. 2004). 70 Thus, by varying seed mass, wind-dispersed plant species may trade dispersal distance for 71 establishment or vice versa. Studying seed dispersal capacity together with establishment traits 72 is, therefore, essential to understand how seed-dispersal influences plant population dynamics. 73 Here, we studied the ecological and evolutionary impact of root herbivory on seed 74 dispersal and fitness in the common dandelion, Taraxacum officinale agg. (Asteraceae). T. 75 officinale is a species complex native to Eurasia that consists of sexual, obligate outcrossing 76 diploids and asexual, apomictic triploids. T. officinale produces seeds with characteristic 77 parachutes (pappus), which enhance uplift by the formation of separated vortex rings after 78 detachment (Cummins et al. 2018). The seeds can be dispersed efficiently by wind and allow 79 the plant to rapidly colonize new habitats (Tackenberg et al. 2003). A recent study found that 80 T. officinale exposed to above ground herbivory produced seeds with longer pappus and thus 81 increased dispersal ability (de la Peña & Bonte, 2014). However, as locust exposure also 82 results in higher biomass production, it is unclear if the change in seed morphology is a direct 83 plastic response to herbivore exposure or an indirect consequence of changes in plant growth 84 (de la Peña & Bonte, 2014). Furthermore, whether herbivory-induced changes in seed traits 85 are related to the plant's evolutionary history, i.e. whether they are the result of natural selection 86 by herbivore attack, is unclear. The larvae of Melolontha melolontha (Coleoptera: 87 Scarabaeidae), also called white grubs, are the major native belowground herbivores of T. 88 officinale. Although the larvae are polyphagous, they prefer to feed on T. officinale (Hauss 1975, 89 Hauss & Schütte 1976). In a recent study, we showed that *T. officinale* populations that evolved 90 under high *M. melolontha* pressure over the last decades produce higher concentrations of 91 deterrent root secondary metabolites, suggesting that M. melolontha exerts positive selection 92 pressure on T. officinale defenses (Huber et al. 2016b). Here, we tested the ecological and 93 evolutionary impact of M. melolontha root herbivory on seed dispersal and fitness in T. 94 officinale. We planted offspring of sympatric and allopatric T. officinale populations into the field 95 and selectively infested them with *M. melolontha* larvae. We then assessed the morphological 96 seed characteristics that determine T. officinale seed dispersal capacity by linking vertical seed 97 fall rate with seed morphology. We compared dispersal and fitness related traits of seeds 98 collected from plants under grub attack with traits of seeds collected from non-attacked plants.

- 99 By comparing grub-attack induced changes in seed dispersal and fitness of *T. officinale*
- 100 populations with different evolutionary histories, we present evidence that *M. melolontha* affects
- 101 plant fitness and dispersal through induced phenotypic plasticity that is shaped by natural
- 102 selection.

## 103 Material and Methods

#### 104 Morphological seed characteristics underlying T. officinale seed dispersal capacity

105 Average horizontal dispersal distance (x) of wind-dispersed seeds can be calculated 106 using a common ballistic model by multiplying the shedding height of the seed (H) with the 107 horizontal wind speed (u), divided by the vertical fall rate of the seed (F): x = Hu/F (Pasquill & 108 Smith 1983; Brock et al. 2005). Seed fall rate is determined by seed traits such as seed mass 109 and seed shape (Matlak 1987; Meyer & Carlson 2001; Skarpaas et al. 2011). To identify the 110 morphological seed traits that contribute most strongly to seed dispersal potential of T. 111 officinale, we measured the fall rate of individual T. officinale seeds along with morphological 112 characteristics of individual seeds and used this data to construct a model that predicts 113 dispersal potential from seed morphology. To cover a broad range of genotypes and 114 environments, seeds of a total of 20 plants from the Botanical Garden of the University of Bern, 115 Switzerland (46.95 °N, 7.43 °E, 520 m above sea level), a semi-natural grassland in Boltigen, 116 Switzerland (46.38 °N, 7.23 °E, 1038 m above sea level), and the greenhouse (genotype A34, 117 Verhoeven et al. 2010) were collected. Five seeds per plant were randomly selected for 118 experiments.

119 To measure seed fall rate, we constructed a semi-automated time-of-flight system (Fig. 120 1a). The system consists of a plexiglass tube (2 m height, 20 cm diameter, ABC-Kunststoff-121 Technik GmbH, Germany) that is coated with a fine metal wire grid on the inside to prevent 122 static charges (Fig. 1a). A custom-built sampler on top of the tube ensures standardized, serial 123 release of individual seeds into the time-of-flight tube. The sampler consists of a conveyor belt 124 with 10 seed-containers, an electromotor to move the belt and a series of controllers to the 125 sampler (components from Tinkerforge, Germany). A remote control allows the user to release 126 a seed and simultaneously start a digital timer by pressing the start button. As soon as the seed 127 crosses the finish line at the bottom of the tube, the user presses the stop button, which stops 128 the timer and saves the time-of-flight in a small memory module. The source code of the control 129 software and a list of the individual components can be found on Github (to be inserted at a 130 later time). Using this system, 100 test seeds were measured three times, and average fall 131 rates for each seed were calculated. As physiological parameters that may determine fall rate, 132 seed weight, length and width, achene length, pappus stem length and pappus hair length were

determined. To standardize measures, a glass plate was put on top of the seed, and traits were measured two-dimensionally from the top. Linear regression models were then built using falling rate as response variable and combinations of seed traits as explanatory variables. We then compared the models using ANOVAs to select the best-performing model. Major variables contributing to fall rate (seed mass and pappus hair length, see results), were selected as measures of seed dispersal potential in subsequent experiments.

#### 139 Effect of root herbivory on seed dispersal and fitness

140 To examine how root herbivory affects seed dispersal and fitness of allopatric and 141 sympatric T. officinale plants, we performed a one-year experiment in a semi-natural grassland 142 using seeds from 18 T. officinale populations evolved in the presence or absence of M. 143 melolontha over the last 20 years (Table S2). A detailed description of the evolutionary history 144 of the majority of these populations can be found in Huber et al. (2016b). In brief, nine 145 populations originated from areas in Switzerland and Germany with high documented M. 146 melolontha densities over the last 20 years (sympatric populations) and nine populations came 147 from areas with no or only low M. melolontha abundance over the last 20 years (allopatric 148 populations). Within each type of evolutionary history, six populations consisted of sexual, 149 outcrossing diploids, and three populations consisted of apomictic, clonal triploids. Triploid 150 seeds were collected directly from their respective mother plants. Diploid seeds were obtained 151 by hand-pollination within populations. For each population, we created a seed mixture by 152 mixing 15 seeds per plant from six randomly selected plants, giving a total of 90 seeds per 153 population. Seed mixtures were germinated in the greenhouse and after two weeks, seedlings 154 were individually transplanted into plastic pots (5 x 5 x 5 cm) filled with soil (Selmaterra, Bigler 155 Samen AG, Thun, Switzerland). Three months after germination, in March 2016, the plants 156 were introduced to the field site in Boltigen (46.38 °N, 7.23 °E, 1038 m above sea level). The 157 field site was established in a semi-natural, organically managed grassland. T. officinale occurs 158 naturally at the field site. *M. melolontha* is present in low abundance in this region.

For the experiment, a plot of 50 m x 15 m was fenced in to prevent cattle from entering. Plants were transplanted individually into bigger pots (13 x 13 x 13 cm) filled with soil (Selmaterra, Bigler Samen AG, Thun, Switzerland). To enable exchange of exudates, a hole was cut into the bottom of the pots (5 cm diameter) and a permeable root barrier (15 x 15 cm; 163 Trenn-Vlies, GeoTex Windhager, Switzerland) was put into the pots before transplanting. Then 164 we excavated soil blocks (approximately 14 x 14 x 14 cm) in the field plot in seven parallel rows, 165 with 40 cm space between the holes and 210 cm space between the rows and inserted the 166 plants into the field in a fully randomized design (Fig. S1). This setup allowed us to embed the 167 plants into their natural environment while controlling root herbivore exposure. To infest plants 168 with *M. melolontha*, individual second instar larvae were introduced to half of the pots in a 169 randomized fashion in the beginning of June 2016. M. melolontha larvae were collected from a 170 field in Uri, Switzerland (46.45 °N, 8.42 °E, 822 m above sea level) and reared on carrots for 2 171 weeks prior to the start of infestation.

172 Measurement of seed dispersal and fitness

173 On the northern hemisphere, T. officinale has its main flowering peak in spring and a 174 secondary flowering peak in late summer / early autumn (Stewart-Wade et al. 2002). We 175 continuously monitored flower development during both flowering peaks (summer 2016 and 176 spring 2017). Every 3-4 days, all fully ripened seeds were collected for analysis of dispersal-177 related seed traits. For collection, seeds of inflorescences were carefully transferred into a small 178 paper bag without cutting or damaging the flower and flower stem heights were measured. If 179 seeds of an inflorescence were missing from the flower heads, we estimated the percentage of 180 remaining seeds. Seeds were stored in darkness at 4 °C until analysis.

181 For seed phenotyping, one seed head from summer 2016 and one seed head from 182 spring 2017 were randomly chosen from each plant. Five to twelve seeds with no visible signs 183 of damage were randomly selected from the different seed heads and phenotyped. Average 184 seed weight and average pappus hair length per seed head were then determined. Seed fall 185 rate was calculated as a function of seed mass and length of pappus hairs. Seed fall rate, 186 vertical wind speed and shedding height of the seed then allowed for an estimation of the 187 dispersal potential for each capsule. A fixed value of 23.88 cm, corresponding to the average 188 stem height in summer 2017, was chosen as a release height for our calculations. We used a 189 constant release height, as the variation in stem height between genotypes recorded in summer 190 2017 was minimal compared to the variation in seed traits (Table S1, Fig. S2). For wind speed, 191 we used 2 m/s, which is the yearly average wind speed in 2016 recorded by the nearest 192 SwissMetNet surface weather station (Boltigen, 46.37 °N, 7.23 °E, 820 m above sea level).

193 To determine if the dispersal potential and the number of viable offspring per plant are 194 correlated, we measured germination rates of collected seeds in a climate chamber in August 195 2017 (temperature at day 22 °C and at night 18 °C; 16 h light and 6 h darkness; 65% humidity; 196 light intensity approximately 120 µmol\*m<sup>-2</sup>\*s<sup>-1</sup>). From each phenotypic analysed seed capsule, 197 ten seeds were randomly selected and equally distributed in a small pot (5.5 cm diameter) filled 198 with moist seedling substrate. The pots were watered regularly and checked daily. Each day 199 we recorded the number of germinated seeds per pot and after 14 days, germination rate was 200 calculated by dividing the number of emerged seedlings through the number of seeds. By 201 multiplying germination rate with the number of flowers per plant and number of seeds per head 202 per plant, we then estimated the number of viable offspring per plant as a proxy for plant fitness 203 (Erb 2018).

## 204 Measurement of vegetative growth

205 For each of the 797 plants in the field experiment, we determined plant vegetative 206 growth by counting the number of leaves and measuring the length of longest leaf as a proxy 207 for plant performance (Huber et al. 2016a). Plant performance was determined on a monthly 208 basis from June 2016 to June 2017, with the exception of January-March 2017, when snow 209 cover prevented measurements. In mid-July 2016, the field was mown according to normal 210 agricultural practice. Before mowing, we individually collected the aboveground biomass of 211 each plant. Shoots were cut 5 cm above ground, put into paper bags, dried in an oven at 80 °C 212 until constant mass was reached and weighted. In June 2017, the experiment was terminated, 213 and aboveground biomass was measured again.

214 Statistical analyses

All statistical analyses were performed in R 3.3.3 (R Core Team 2017). To predict seed fall rate, 'Seed mass' and 'Pappus hair length' were used in a linear regression model. No interactions were included in the model. The predictive power of the model was evaluated using k-fold cross validation. For the linear regression model, the function 'lm' of the stats package was used (R Core Team 2017) and the validation of the predictive power was done with the function 'train' of the 'caret' package (Kuhn 2008).

For the analysis of seed traits, mean values per population were calculated separately for plants with and without grub infestation. Mean population values were then examined with

223 linear models. Multiple linear models were built with 'Herbivory' x 'Evolutionary history' x 'Ploidy 224 level' as explanatory variables and each recorded seed parameter as response variable 225 ('Flower heads', 'Seeds per head', 'Pappus hair length', 'Seed mass', 'Dispersal potential', 226 'Germination rate' and 'Offspring per plant'). For analysis of vegetative growth, a linear model 227 was employed on 'Biomass' as response variable with 'Herbivory' x 'Evolutionary history' x 228 'Ploidy level' as explanatory variables. Further, 'Length of longest leaf' was analysed as 229 response variable in two linear models with 'Herbivory' x 'Evolutionary history' x 'Time' as 230 explanatory variables separately for diploid and triploid plants. All models were examined using 231 the function 'plotresid' (package 'RVAMemoire' (Hervé 2018)). Significance of explanatory 232 variables was tested using the function 'Anova' (package 'car' (Fox & Weisberg 2011)) and 233 results were plotted using the package 'ggplot2' (Wickham 2009). Similar results were obtained 234 with models for values of individual plants, which included 'Population' as random factor, 235 compared to models for mean values per population.

To investigate if changes in seed mass are linked to changes in biomass, correlation strength between 'Seed mass' and Biomass' was analysed separately for seeds from summer 2016 and spring 2017. Further, to examine if there is a trade-off regarding the plant's investment into seed production optimized for dispersal or germination, correlation strength between 'Dispersal potential' and 'Germination rate' was analysed separately for seeds from summer 2016 and spring 2017.

242 To investigate direct and indirect relationships between and within analysed seed and 243 growth traits, we used structural equation modelling (SEM) (Shipley 2016). We developed an 244 a priori model based on the results of our data analysis and on physiological and ecological 245 knowledge, which includes all measured plant traits except the estimated variables 'Dispersal 246 potential' and 'Offspring per plant', as those were calculated out of other seed traits. We 247 hypothesized direct effects of season (summer 2016 vs. spring 2017) on the measured seed 248 and flower traits and on biomass production. Further, we assumed direct effects of biomass on 249 the measured seed and flower traits and an indirect effect of biomass on germination rate 250 through seed mass and pappus hair length. In addition, we hypothesized an indirect effect of 251 number of flower heads per plant and number of seeds per head on germination rate through 252 seed mass and pappus hair length. We assumed bidirectional links between seed mass and

- 253 pappus hair length and between number of flower heads per plant and number of seeds per
- head. To avoid large differences in the variances among the variables we first scaled all
- variables to an equal range by dividing through powers of 10 if necessary. We then used the R
- package 'lavaan' (Rosseel 2012) for fitting and testing the SEM. Model fit indices (CFI = 0.999,
- TLI = 0.996, RMSEA = 0.018) indicated good model fit, so maximum likelihood was used to
- 258 estimate standardized path coefficients.

# 259 Results

260 Seed weight and pappus hair length determine dispersal potential of T. officinale seeds

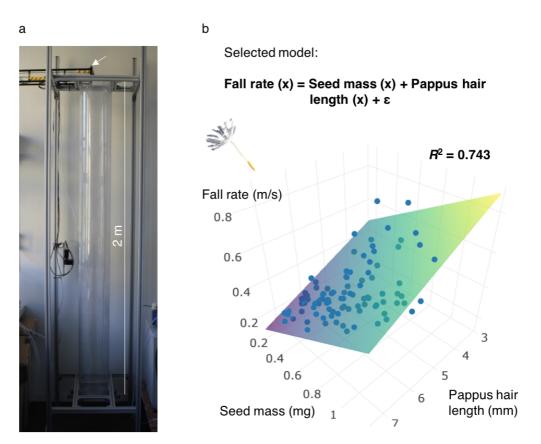
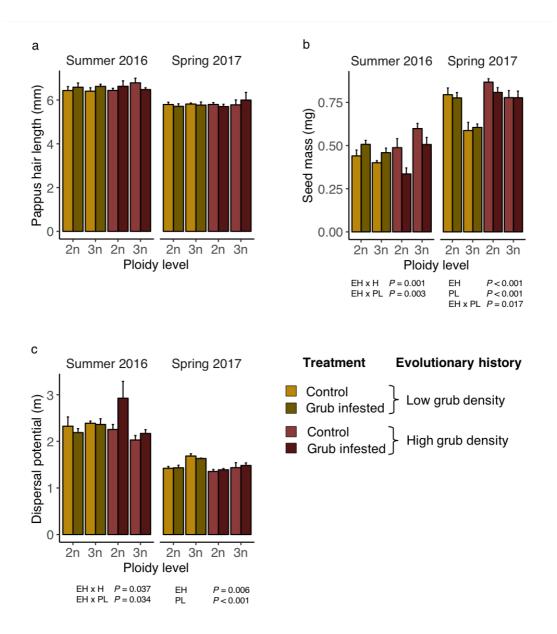


Fig. 1 Seed mass and pappus hair length determine dispersal potential of dandelion seeds. (a) Photograph of a custom-built plexiglass time-of-flight tube with semi-automatic seed dispenser (white arrow) to determine fall rates of individual dandelion seeds. (b) A linear model predicts seed fall rate using seed mass and pappus hair length. The adjusted  $R^2$  of the modelfit is displayed.

- 266 Comparisons of multiple linear regression models revealed that vertical seed fall rate 267 of T. officinale – and subsequently seed dispersal potential - is predicted well by seed mass 268 combined with pappus hair length. The final model (Fall rate (x) = Seed mass (x) + Pappus hair 269 length (x) +  $\varepsilon$ ) showed high predictive power using these two parameters (R<sup>2</sup> = 0.743, Fig. 1b). 270 While seed mass increases fall rate and thus decreases dispersal potential, the length of the 271 pappus hair decreases fall rate and thus increases dispersal potential. 272 Root herbivory attack increases seed dispersal potential in sympatric populations 273 During the summer flowering period, *M. melolontha* infestation did not affect pappus
- hair length (Fig. 2a), but reduced the mass of *T. officinale* seeds (Fig. 2b), resulting in increased
- 275 dispersal potential compared to control plants (Fig 2c). This response was observed in

- sympatric, but not allopatric populations (LM, factor 'EH x H', P = 0.001, Fig. 2b; respectively
- 277 LM, factor 'EH x H', P = 0.037, Fig. 2c). During the spring flowering period, no effect of root
- herbivory on seed traits was detected (Table S1).
- 279





281 Fig. 2 Root herbivore attack increases seed dispersal potential in sympatric dandelion 282 populations. Compared to allopatric populations (vellow bars), sympatric populations (red 283 bars) developed lighter seeds with increased dispersal potential under root herbivory. Seed 284 mass (a), pappus hair length (b) and dispersal potential (c) of seeds collected from plants with 285 (hatched bars) and without (plain bars) root herbivory by M. melolontha are shown. Mean 286 values of population means (N = 3-6 populations) and standard errors (± SE) are indicated. P-287 values from linear models are shown (significant effects only). EH: Evolutionary history of 288 population. H: Root herbivory treatment. PL: Ploidy level. 2n: Diploid populations. 3n: Triploid 289 populations.

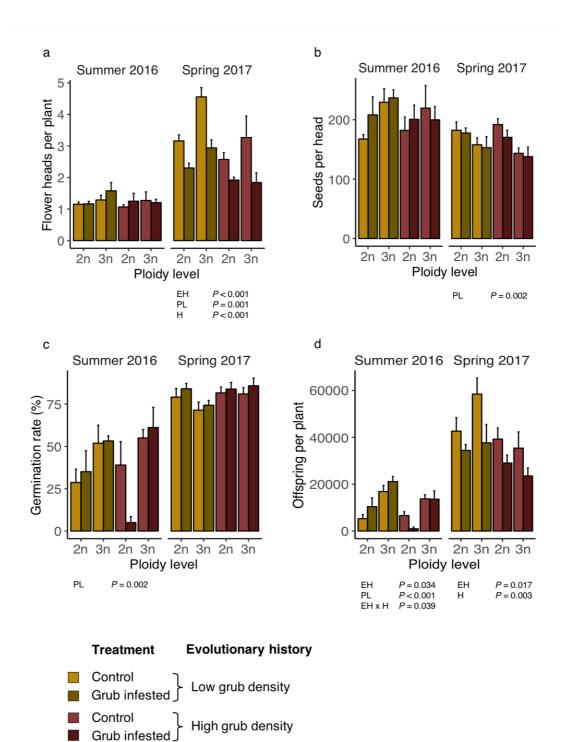
#### 291 Evolutionary history and cytotype interact to determine seed dispersal potential

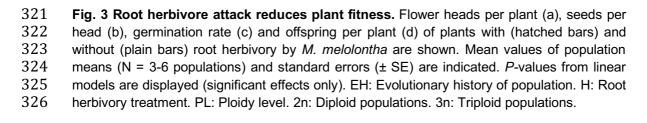
292 We detected significant effects of evolutionary history and cytotype on seed mass and 293 dispersal potential (Fig. 2). During the summer flowering period, seeds of sympatric triploids 294 were heavier and had a lower dispersal potential than seeds of sympatric diploids, while the 295 opposite was the case for allopatric populations (LM, factor 'EH x PL', P = 0.003, Fig. 2b; 296 respectively LM, factor 'EH x PL', P = 0.034, Fig. 2c). During the spring flowering period, seeds 297 of sympatric populations were heavier and had a lower dispersal potential than seeds of 298 allopatric populations (LM, factor 'EH', P < 0.001, Fig. 2b; respectively LM, factor 'EH', P = 299 0.006, Fig. 2c), and triploid seeds were generally lighter and thus had a higher dispersal 300 potential than diploid seeds (LM, factor 'PL', P < 0.001, Fig. 2b; respectively LM, factor 'PL', P

301 < 0.001, Fig. 2c).

### 302 Root herbivory reduces plant fitness

303 During the summer flowering period, root herbivory did not significantly affect the 304 number of flower heads, the number of seeds or the percentage of germinating seeds per plant 305 (Fig. 3a-c). The total number of offspring per plant, which was calculated by multiplying flower 306 heads, seeds per head and germination rate, however, was reduced by *M. melolontha* attack 307 in sympatric, but not in allopatric populations (LM, factor 'EH x H', P = 0.039, Fig. 3d). In spring, 308 M. melolontha reduced the number of flower heads and total viable offspring per plant 309 independently of the other factors (LM, factor 'H', P < 0.001, Fig. 3a; respectively LM, factor 310 'H', P = 0.003, Fig. 3d). Evolutionary history and cytotype also affected plant fitness traits (Fig. 311 3). Triploid seeds from the summer flowering period germinated better, leading to a higher 312 number of offspring per plant (LM, factor 'PL', P =0.002, Fig. 3c; respectively LM, factor 'PL', P 313 < 0.001, Fig. 3d). During the spring flowering period, triploids produced more flower heads per 314 plant (LM, factor 'PL', P = 0.001, Fig. 3a), but fewer seeds per flower head (LM, factor 'PL', P 315 = 0.002, Fig. 3b), and showed no difference in germination rate and offspring per plant (Fig. 3c-316 d). Sympatric populations produced fewer flowers than allopatric populations during the spring 317 flowering period, and the number of viable offspring was lower for sympatric populations than 318 allopatric populations (LM, factor 'EH', P < 0.001, Fig. 3a; respectively LM, factor 'EH', P = 319 0.017, Fig. 3d).





#### 327 Root herbivory decreases vegetative growth

328 To determine whether reduced plant performance upon *M. melolontha* attack may 329 explain the changes in seed mass and subsequently in dispersal potential in sympatric 330 populations, we measured the impact of *M. melolontha* attack on shoot growth and biomass 331 and then correlated shoot biomass with seed mass. The length of the longest leaf, a proxy for 332 biomass accumulation (Huber et al. 2016a) varied through the season and was significantly 333 reduced by M. melolontha infestation (Fig. 4a). Shoot biomass was significantly reduced in 334 plants attacked by *M. melolontha* during both sampling periods (LM, factor 'Herbivory', P = 335 0.009 and LM, factor 'Herbivory', P = 0.047, Fig. 4b). Further, during the summer sampling 336 period, triploids produced more biomass than diploids (LM, factor 'Ploidy level', P < 0.001, Fig. 337 4b). However, none of these effects was different between sympatric and allopatric populations, 338 suggesting that the increase in seed dispersal capacity upon *M. melolontha* attack in sympatric 339 populations is not due to a general plant stress response.

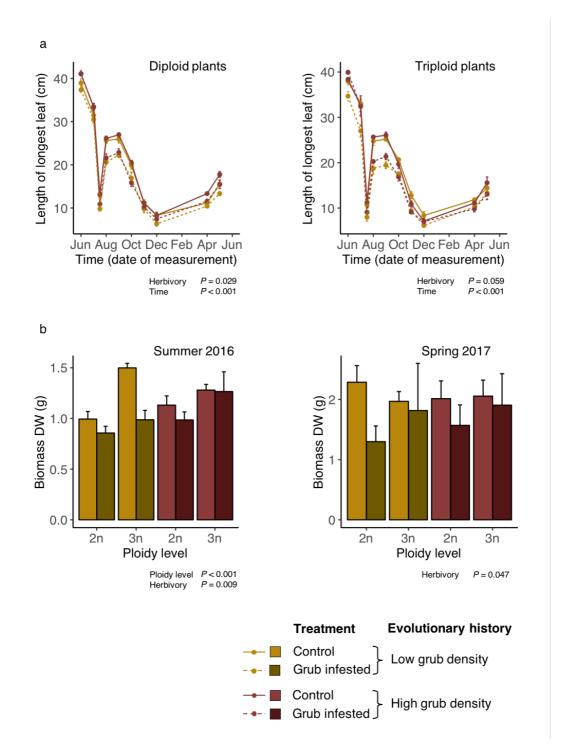
## 340 Correlations between vegetative growth, fitness and dispersal

341 Structural equation modelling (SEM) revealed that the sampling period ('Season') had 342 an influence on both vegetative and generative traits. In summer 2016, *T. officinale* plants 343 produced more seeds per head and seeds with longer pappus hair. In spring 2017, plants 344 produced more biomass, more flower heads per plant, heavier seeds with higher germination 345 rate, but shorter pappus hair (Fig. 5a).

346 Across all plants, vegetative biomass production explained seed mass and pappus hair 347 length (Fig. 5a). However, across populations and analysed for each sampling season 348 separately, shoot biomass was not correlated with average seed mass and hence dispersal 349 potential ( $R^2 = 0.009$ , P = 0.593 and  $R^2 = 0.080$ , P = 0.095, Fig. S3). Seed mass explained 350 pappus hair length and germination rate. (Fig. 5a). Pappus hair length was a negative predictor 351 of germination rate. The number of flowers per head was a negative predictor of seed mass. 352 while the number of seeds per head was a positive predictor of seed mass and pappus hair 353 length (Fig. 5a).

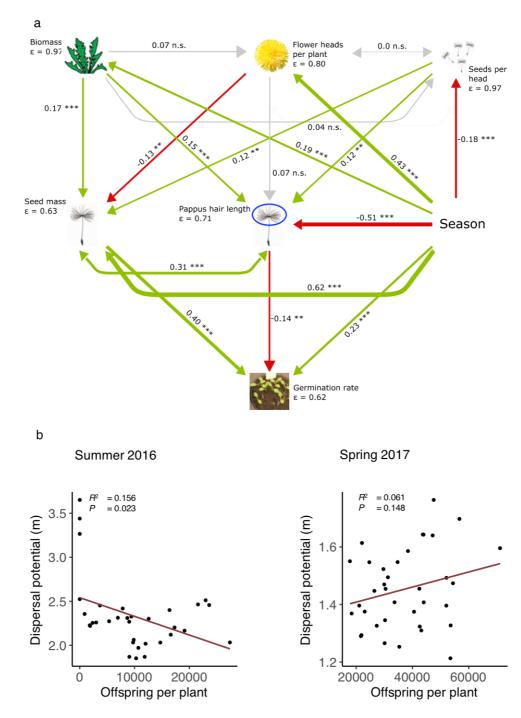
To further understand the relationship between dispersal potential (calculated from seed mass and pappus hair length) and total offspring per plant (calculated from flower heads, seeds per head and germination rate), we correlated these two parameters. Dispersal potential

- 357 and offspring per plant were negatively correlated for the summer flowering period, but not the
- 358 spring flowering period ( $R^2 = 0.156$ , P = 0.023 and  $R^2 = 0.061$ , P = 0.148, Fig. 5b). Together,
- 359 these results show that *T. officinale* dispersal and reproductive potential differ markedly
- 360 between the summer and spring flowering periods, and that there seems to be a tradeoff
- 361 between dispersal and reproduction during the summer flowering period, which is mostly driven
- 362 by seed mass, which increases germination, but decreases dispersal potential.



363

**Fig. 4 Rood herbivore attack reduces vegetative growth.** (a) Length of longest leaf in diploid and in triploid populations under infestation with *M. melolontha* (dashed lines) compared to plants without root herbivory (full lines). Vegetation was cut back at the end of July. (b) Above ground biomass decreased under root herbivory (hatched bars) compared to biomass of plants without root herbivory (plain bars). Mean values of population means (N = 3-6 populations) and standard errors (± SE) are shown. *P*-values from linear models are displayed (significant effects only). 2n: Diploid populations. 3n: Triploid populations.





373 Fig. 5 Dispersal and establishment are negatively correlated during the summer 374 flowering period (a) Structural equation model showing pathways between phenotypic plant 375 parameters. Positive links are indicated by green arrows and negative links are indicated by 376 red arrows. Grey arrows represent non-significant links. Red arrows coming from 'Season' 377 represent stronger effects of sampling period 'summer 2016', whereas green arrows coming 378 from 'Season' represent stronger effects of sampling period 'spring 2017'. Line thicknesses of 379 arrows correspond to the effect size given by their standardized path coefficients, which are 380 shown next to each path. Asterisks next to path coefficients indicate P-values (\*\*\*P < 0.001; 381 \*\*P < 0.01; \*P < 0.05; n.s.  $P \ge 0.05$ ). Residual variance ( $\epsilon$ ) is given for all response variables. 382 (b) Strength of correlation between dispersal potential and offspring per plant for seeds 383 collected in summer 2016 and in spring 2017. Linear regression lines of population means and 384 corresponding  $R^2$ - and P-values are shown. 385

## 386 Discussion

Plants can escape from unfavourable environments through seed dispersal. How seed dispersal capacity is affected by herbivory, however, is not well understood. Here, we show that *T. officinale* plants from sympatric, but not allopatric populations, increase their seed dispersal potential upon root herbivore attack in the field by reducing seed mass at the expense of the number of viable offspring. Below, we discuss the mechanisms and ecological implications of these findings.

393 Whether seed dispersal is an effective strategy to escape from hostile environments 394 depends on the distance that can be covered by the dispersing seeds relative to the size of the 395 unfavorable habitat. Attack by root-feeding M. melolontha larvae resulted in increased seed 396 dispersal potential in sympatric T. officinale plants during the summer flowering period. 397 Assuming an average wind speed of 2m/s, dispersal potential for grub-attacked sympatric 398 populations can be estimated to exceed dispersal of non-attacked populations by 41 cm. As 399 third instar larvae, *M. melolontha* can move up to 20 cm per day (Hasler 1986). An increased 400 seed dispersal distance of 41 cm may therefore not be sufficient to escape a patch with high 401 *M. melolontha* density. However, chances for long-distance dispersal induced by vertical wind 402 turbulence are also higher for seeds with lower falling velocity. Although long-distance dispersal 403 is relatively rare, producing seeds with lower fall rates does enhance the probability of T. 404 officinale offspring to be dispersed over hundreds of meters under favorable conditions, which 405 may allow the plant's offspring to escape from heavily root herbivore infested patches.

406 Theory predicts that increased seed dispersal is advantageous under locally stressful 407 conditions and may therefore be adaptive (Levin et al. 2003, Ronce 2007). To which extend the 408 herbivory-induced changes in seed traits observed here represents a form of adaptive plasticity 409 is not fully clear. Adaptive plasticity, defined as plasticity maintained through natural selection, 410 should lead to increased fitness (Miner et al. 2005). However, dispersal distance per se is not 411 a direct fitness measurement, as the benefit of dispersal depends on the environmental context 412 (van Kleunen & Fischer 2005; Teller et al. 2014). In this context, it is worth noting that we 413 observed an increase in seed dispersal potential during grub attack in sympatric T. officinale 414 populations, which have evolved in the presence of *M. melolontha* over the last decades, but 415 not in allopatric populations that have not been under white grub pressure. This result supports

the notion that seed dispersal traits are subject to natural selection (Cheptou et al. 2008; Ronce
2007; García-Verdugo et al. 2017; Saastamoinen et al. 2018) and indicates that root herbivores
may exert selection pressure on seed phenotypic plasticity.

419 In general, the seed dispersal capacity of a plant may increase under unfavorable 420 environmental conditions due to "stress flowering", leading to suboptimal seed development 421 and, thus, lighter seeds (Germain & Gilbert 2014, Begcy & Walia 2015). Our results suggest, 422 however, that physiological stress is unlikely to explain the increase in dispersal potential in 423 root-herbivore attacked dandelions. We found that M. melolontha attack reduces growth and 424 biomass accumulation of dandelion plants, but that these effects are similar in sympatric and 425 allopatric populations. Thus, we expect that the extent of stress that is imposed by root 426 herbivory does not differ between allopatric and sympatric populations. As only sympatric 427 populations adjust their seed phenotype upon root herbivore attack, we, therefore, propose that 428 this change represents a form of herbivore-induced plant response rather than a secondary 429 effect of physiological stress.

430 Seed mass influences dispersal potential and germination ability in opposite directions 431 and therefore represents a physiological trade-off. Many studies empirically support this 432 assumption (e.g. Soons & Heil 2002; Jakobsson & Eriksson 2003). Our study confirms that 433 vertical falling velocity and hence distribution distance of T. officinale seeds as well as 434 germination rate are strongly linked to seed mass. The capacity to disperse seeds thus comes 435 with a reduction in total viable offspring. Thus, dandelions may trade establishment ability 436 against higher dispersal potential under root herbivore attack. The high cost of higher dispersal 437 likely contributed to the inducibility of this phenomenon. Interestingly we found a significant 438 negative relationship between dispersal distance and number of viable offspring for the first 439 sampling period in summer, when root herbivory had a significant effect on dispersal, but no 440 significant link in the second sampling period, where root herbivory did not change seed 441 phenotypes. Thus, the negative relationship between dispersal and seed mass seems to be 442 context dependent and more strongly expressed under environmental stress.

Herbivores exert selection pressure on plants, leading to the evolution of tolerance and defense strategies (Núñez-Farfán et al. 2007). Earlier work demonstrated that sympatric dandelion populations that grow in the presence of *M. melolontha* produce higher amounts of

446 repellent root sesquiterpene lactones, which may benefit them by becoming less attractive to 447 M. melolontha (Huber et al. 2016b). The experimental setup in this study controlled for these 448 potential effects by confining root herbivores to individual dandelion plants. Our results suggest 449 that, when M. melolontha larvae do not have a choice between different plants, there is no 450 direct benefit to producing the repellent chemicals. On the contrary, sympatric populations 451 suffered from the same biomass reduction and did not produce more flowers or seeds under 452 root herbivore attack than allopatric populations. In combination with our earlier studies (Huber 453 et al. 2016a, Huber et al. 2016b, Bont et al. 2017), these results suggest that high herbivore 454 pressure selects for repellent root chemicals and higher dispersal potential, but not stronger 455 resistance or tolerance in dandelion. These results emphasize the need to evaluate the 456 evolution of plant defense strategies in a community context (Agrawal et al. 2012, Poelman & 457 Kessler 2016) and to include seed and dispersal traits into these analyses (Erb 2018).

Through feeding on roots, belowground insects can severely impair plant performance. From an evolutionary point of view, presence of root-feeders may therefore act as selective force on plant traits involved in seed distribution and establishment. In our study, exposure to *M. melolontha* resulted in reduced flower production, but it also resulted in the production of better dispersible seeds - depending on the sampling period and the evolutionary history of *T. officinale.* Thus, our findings suggest an important, but context-dependent influence of root herbivory on plant reproduction and dispersal traits.

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