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1 **RESEARCH IN CONTEXT**

2 From rhizoids to roots? Experimental evidence of mutualism between

3 liverworts and ascomycete fungi.

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1 Abstract

The rhizoids of leafy liverworts (Jungermanniales, Marchantiophyta) are commonly colonized by the ascomycete fungus Pezoloma ericae. These associations are hypothesized to be functionally analogous to the ericoid mycorrhizas (ErM) formed by P. ericae with the roots of Ericaceae plants in terms of bi-directional phosphorus-for-carbon exchange; however, this remains unproven. Here, we test whether associations between the leafy liverwort Cephalozia bicuspidata and P. ericae are mutualistic.

9 We measured movement of phosphorus and carbon between C. bicuspidata 10 and P. ericae using ³³P-orthophosphate and ¹⁴CO₂ isotope tracers in monoxenic 11 cultures. We also measured leafy liverwort growth, with and without P. ericae. We 12 present the first demonstration of nutritionally mutualistic symbiosis between a non-13 vascular plant and an ErM-forming fungus, showing transfer of fungal-acquired P to 14 the liverwort and of liverwort-fixed C to the fungus alongside increased growth in 15 fungus-colonized liverworts.

16 Thus, this ascomycete-liverwort symbiosis can now be described as 17 mycorrhiza-like, providing further insights into ericoid mycorrhizal evolution and 18 adding Ascomycota fungi to mycorrhizal fungal groups engaging in mutualisms with 19 plants across the land plant phylogeny. As P. ericae also colonizes the rhizoids of 20 Schistochilaceae liverworts, which originated in the Triassic and are sister to all other 21 jungermannialean liverworts associated with fungi, our findings point toward an early 22 origin of ascomycete-liverwort symbioses, possibly predating their evolution in the 23 Ericales by some 150 MY.

24

Key words: Cephalozia bicuspidata; ericoid mycorrhizal fungi; liverwort; Pezoloma
 ericae; symbiosis; carbon-for-nutrient exchange; mycorrhizas; mutualism

3

4 Introduction

5 Mycorrhizas are intimate symbioses formed between plant roots and soil fungi 6 that are prevalent across the globe in more than 80% of extant land plants (Smith and 7 Read, 2008). Through mycorrhizal associations, many plants engage in bi-directional 8 exchange of photosynthesis-derived plant carbon and fungal-acquired nutrients, 9 scavenged from sources beyond the root depletion zone or from soil pores too small 10 for roots to access (Smith and Read, 2008; Leake and Read, 2016). The role of 11 mycorrhizas in supplying extant land plants with nutrients, together with evidence of 12 mycorrhiza-like associations in Rhynie Chert plant fossils (Remy et al., 1994) has led 13 to the hypothesis that mycorrhizal fungi likely facilitated the evolution of land plants 14 >470 Mya (Pirozynski et al., 1975; Bidartondo et al., 2011; Leake, 2015). 15 Additionally, recent studies have shown that the genes required for mycorrhization 16 are conserved across all land plant lineages (Wang et al., 2010), including the earliest 17 diverging clade of liverworts – Haplomitriopsida (Fig. 1a, after Crandall-Stotler et 18 al., 2009). The later-derived leafy liverworts (Jungermanniidae) have not been 19 incorporated in such analyses and a critical caveat of molecular studies is that the 20 presence of genes does not necessarily imply functional significance. Indeed, for 21 many groups of extant plants that form mycorrhizas and mycorrhiza-like associations 22 where roots are absent, knowledge regarding the physiological function of the 23 symbiosis has been severely limited (Read et al., 2000, Field et al., 2015a). To date, 24 amongst fungus-associated early-branching land plants (i.e. liverworts, hornworts and 25 lycophytes) nutritional mutualisms have only been demonstrated in a handful of early

1 diverging thalloid and Haplomitriopsida liverworts (Field et al., 2012; 2015a; 2016),

2 (Fig. 1a), all which form symbioses with Mucoromycota fungi (Glomeromycotina

3 and/or Mucoromycotina) (Field et al. 2015b).

As far as can be established from published surveys and incidental 4 5 illustrations in floras (Pocock et al., 1984; Pocock & Duckett, 1985; Duckett et al.,1991, Ligrone et al. 2007) rhizoidal ascomycete associations occur in far more 6 7 liverwort species than those containing basidiomycetes, glomeromycetes and 8 mucoromycetes (Pressel et al., 2010). Following the accepted species names in the 9 Plant List (2013), we estimate that up to 1,000 late-diverging leafy liverworts are 10 likely to have ascomycete fungal symbionts, i.e. around 20% of liverwort species 11 worldwide compared to less than 100 with basidiomycetes, glomeromycetes and 12 mucoromycetes (see Table S2 for detailed breakdown). Reinforcing their symbiotic 13 rather than opportunistic status, the ascomycetes 1) induce swelling, branching and 14 septation of the rhizoids; and, 2) are ubiquitous rather than sporadic, in species where 15 they occur. In addition, transmission electron micrographs show healthy hyphae in 16 healthy host cells (Duckett et al., 1991; Pressel et al., 2010).

17 There are several families in the leafy liverworts (Jungermanniideae), 18 including Schistochilaceae, Lepidoziaceae, Calypogeiaceae, Cephaloziaceae and 19 Cephaloziellaceae, which consistently associate with Ascomycota fungi (Fig. 1a) 20 (Pressel et al., 2010). These fungal symbionts include Pezoloma ericae (D.J. Read) 21 Baral (syn. Rhizoscyphus ericae (D.J. Read) W.Y. Zhuang and Korf, 2004; 22 Hymenoscyphus ericae (D.J. Read) Korf and Kernan, 1983; and Pezizella ericae (D.J. 23 Read, 1974) (Duckett and Read, 1995; Read et al., 2000; Pressel et al., 2010)). 24 Notably, Pezoloma ericae is known to form ericoid mycorrhizas (ErM) with the roots 25 of Ericaceae plants and has previously been shown to provide nutrients to their

vascular plant hosts in exchange for fixed carbon (Read et al., 2003; Smith and Read,
 2008).

3 Ericaceous habitats are typically low in plant-available soil nutrients, 4 including N and/or P (Stribley and Read, 1974; Mitchell and Read, 1981; Leake et al. 5 1990; Bolan, 1991; Myers and Leake, 1996). The vascular plants inhabiting these 6 habitats, such as Calluna, Erica, Rhododendron and Vaccinium, grow together with 7 non-vascular plants, including the widespread leafy liverworts in the Cephaloziaceae 8 (Chambers et al., 1999; Upson et al., 2007) with which they 'share' fungal symbionts 9 (Duckett and Read, 1995; Read et al., 2000). It was recently shown that that this 10 shared mycobiont can bring benefits in terms of establishment and survival to 11 ericaceous plants (Kowal et al., 2015), and that the P. ericae-colonized liverwort may 12 serve as a source of fungal inoculum for vascular plants. It is possible that this effect 13 is driven by fungal-enhanced nutrition in liverworts and then in the vascular plant 14 species sharing the fungal symbiont, analogous to the nutritional role of arbuscular 15 mycorrhizal fungi (Glomeromycotina fungi) that associate with some thalloid 16 liverworts (Field et al., 2012). It may be that P. ericae associates in leafy liverworts 17 play a similar role to the Glomeromycotina or Mucoromycotina fungal partners of 18 thalloid liverworts by supplementing plant phosphorus (P) assimilation (Field et al., 19 2016).

Here, we aim to address the fundamental question of whether the ascomycete fungus P. ericae forms mycorrhiza-like associations with leafy liverworts equivalent to those formed by Glomeromycotina or Mucoromycotina fungi and thalloid liverworts. We traced the movement of P from P. ericae fungal hyphae to Cephalozia bicuspidata liverworts and the movement of carbon (C) from liverworts to the fungi using isotope tracers. We determined liverwort growth responses to colonization by

P. ericae fungi by measuring the size and mass of liverworts, grown both with and
 without P. ericae fungal symbionts.

3

4 Materials and Methods

5 We collected Cephalozia bicuspidata from Thursley Common, Surrey, in 6 autumn 2012 (OS grid reference SU900416). Mature sporophytes were harvested, 7 surface-sterilized and spores were cultured axenically (Duckett and Read, 1995) on 1.5% PhytagelTM (Sigma-Aldrich; ICP elemental analysis of Phytagel provided by 8 9 Sigma-Aldrich: 0.85% Ca, 0.35% Mg; 1.70% K, 0.15% P and 0.45% Na). No 10 additional nutrients were added to the culture medium. Pezoloma ericae was isolated 11 from the same liverwort collection and resynthesized with axenically-grown C. 12 bicuspidata using published methods (Kowal et al., 2015), thus satisfying Koch's 13 postulates. Molecular identification of the fungal isolate as P. ericae was carried out 14 previously (Kowal et al., 2015). Nomenclature for plants and fungi follows Hill et al. 15 (2008) and www.speciesfungorum.org, respectively. 16 17 Fungus-to-plant phosphorus transfer 18 We grew C. bicuspidata and P. ericae together in one compartment 19 ('liverwort and fungus') of 9 cm split-plate microcosms (Fig. S1) filled on both sides 20 of the divide ('B' in Fig. S1a) with 1.5% sterile Phytagel. Fragments of P. ericae 21 isolate (approximately 7 mm²) were inserted beneath the surface of the Phytagel and 22 an axenically-grown leafy liverwort stem (two per microcosm, c. 2 cm apart) was 23 gently pressed onto the surface, directly above the fungus fragment. Following

24 establishment of liverwort-fungal symbiosis, which is confined to the rhizoids (Fig.

1	1b), and growth of extraradical fungal hyphae (eleven weeks after planting), we	
2	introduced 0.1 MBq $H_3^{33}PO_4$ (i.e. 0.03 µg ^{33}P , specific activity 111 GBq mmol ⁻¹ ;	
3	Hartmann Analytics, Braunschweig, DE) into a well within the medium in the	
4	contiguous compartment of the plate. Each ³³ P-labelled well was then filled with	
5	1.5% sterile Phytagel. The barrier dividing the microcosm prevented the liverwort	
6	from encroaching into the compartment containing ³³ P while fungal hyphae were able	
7	to grow over the barrier and colonize the medium in both compartments (see Fig. S1).	
8	We prepared a total of 16 microcosms with an additional fungus-free (control)	
9	microcosm to measure non-fungal mediated diffusion of ³³ P into liverwort tissue. We	
10	also tested the effectiveness of the barriers in eight undivided microcosms containing	
11	C. bicuspidata without fungus with ³³ P-labelled wells placed the same distance from	
12	plants as in divided microcosms. All microcosms were sealed with Parafilm 'M'	
13	(Sigma), and placed in a controlled environment chamber (BDR16, Conviron,	
14	Winnipeg, MB, Canada). The temperature regime was typical of late-spring/summer	
15	for southeastern England with light intensity reflecting that at ground level beneath	
16	canopy vegetation, similar to that experienced by the liverworts in their natural	
17	environment (irradiance of 50 μmol m $^{-2}$ s $^{-1}$, 12 h : 12 h, light : dark, 16° C : 14° C	
18	day : night, 80% RH and 440 ppm [CO ₂]). After eight weeks, we removed the	
19	liverworts from the microcosms, and freeze-dried all plant tissues and growth	
20	medium containing fungal hyphae.	
21	Between 10-30 mg of plant tissue and growth medium for each microcosm	
22	was digested in 1 mL concentrated H ₂ SO ₄ for two hours and then heated to 365 °C for	

was digested in 1 mL concentrated H₂SO₄ for two hours and then heated to 365 °C for
15 minutes. After cooling, 100µL of hydrogen peroxide was added to each sample
before reheating to 365 °C for two minutes, resulting in a clear solution. Samples
were diluted up to 10 mL with distilled water before 2 mL of the diluted digest

solution was mixed with 10 mL of the liquid scintillant Emulsify Safe (Perkin Elmer).
Activity of the samples was determined using liquid scintillation counting (Packard
TriCarb 3100, Isotech, Chesterfield, UK). The amount of ³³P transferred to the plants
by the fungus in each microcosm was calculated using equations from Cameron et al.
(2007). This figure was then adjusted for passive movement of ³³P from the substrate
via diffusion by subtracting the mean amount of ³³P measured in liverworts harvested
from fungus-free control microcosms.

8

9 Plant-to-fungus carbon transfer

10 We filled nine cube-shaped vessels (Sigma Magenta GA-7-3, 77 mm \times 11 77 mm \times 97 mm) with Phytagel (1.5%) to a depth of 30 mm to prepare three replicate 12 microcosms with fungus only and six "complete" microcosms with liverwort and 13 fungus. A fragment of P. ericae isolate (ca. 7 mm²) was inserted beneath the surface 14 of the Phytagel for the fungus-only and complete microcosms. For the complete 15 microcosms, an axenically-grown leafy liverwort stem was gently pressed onto the 16 surface of the medium, directly above the fungus fragment. We maintained the plants 17 in the same controlled environment chamber and conditions as above. After six 18 months of growth, we labeled the microcosms with ¹⁴CO₂, separating the growth 19 medium from the chamber headspace with polythene (pre-cut with 25 mm diameter holes for the liverworts) in order to minimize direct diffusion of ¹⁴CO₂ into the 20 21 substrate. The polythene was sealed with anhydrous lanolin where it met the 22 substrate and along the edges of the Magenta vessels (Fig. S2b). We generated 0.5 MBq of ${}^{14}CO_2$ gas by adding 6.8 μ L ${}^{14}C$ -sodium bicarbonate to tubes in each 23 24 microcosm before introducing 500 µL 25% lactic acid. We allowed plants to fix 25 ¹⁴CO₂ for five hours in the middle of the day before introducing two Eppendorf tubes

1	containing 1 mL 2 M KOH into each microcosm for 30 min to trap any remaining
2	14 CO ₂ from the headspace of each microcosm. We then harvested, separated and
3	freeze-dried the liverworts and Phytagel containing fungal hyphae (see Methods S1)
4	before weighing, homogenizing and determining ¹⁴ C content by sample oxidation
5	(Packard Sample Oxidiser) and liquid scintillation counting (Packard Tri-Carb 3100).
6	We calculated total carbon (12 C plus 14 C) fixed by the plant and transferred to
7	the fungus as a function of the total volume and CO_2 content of the vessel headspace
8	and the proportion of supplied ${}^{14}\text{CO}_2$ label fixed by the plants, using equations from
9	Cameron <i>et al.</i> (2008). The specific activity of the source was 2.04 TBq Mol $^{-1}$.
10	To account for carbon movement through passive diffusion, and to assess the
11	effectiveness of the polythene/lanolin barrier, total plant-fixed fungal carbon in the
12	complete microcosms was determined by subtracting the mean carbon measured in
13	the fungus from the fungus-only control microcosms from each of the complete
14	microcosms.
15	
16	Liverwort growth
17	We measured C. bicuspidata growth both with and without fungal inoculation
18	looking first at changes in leafy liverwort surface area, and second at liverwort
19	biomass. In the first experiment, we introduced small fragments (c. 7 mm ²) of P.
20	ericae to 9 cm plates containing 1.5% Phytagel with axenically-grown leafy
21	liverworts (c. one year old). After fungal colonization of the liverwort rhizoids was
22	confirmed microscopically (Fig. 1b), we re-plated colonized liverworts and fungus-
23	free control liverworts individually using the same medium. Surface area
24	measurements of each liverwort stem (including leaves) were made before and after

1	six weeks using images taken with a Nikon Coolpix S10 digital camera and analyzed
2	using ImageJ software (Rasband, 1997-2012) (Fig. S3). For the second measurement
3	of the effect of P. ericae on C. bicuspidata growth, we freeze-dried and weighed the
4	liverworts from the complete microcosms used for the ¹⁴ C transfer experiment
5	(above) after they were harvested at six months and compared the mean to that of
6	plants from liverwort-only microcosms ($n = 6$). This allowed us to test for
7	differences in mean plant biomass in liverwort growth with or without fungus over a
8	longer time period than the surface area growth experiment described above.
9	Liverwort stems were weighed before initial planting to ensure similarity in mass
10	between sample groups.
11	
12	Statistics
13	After analyzing the data for normal distribution and homogeneity of variance,
14	we applied the appropriate statistical tests, i.e. parametric or non-parametric, using
15	GraphPad Prism (version 6.0h). The Mann-Whitney U test was used both for the dry
16	mass experiment and for testing barrier effect in the ³³ P experiment, owing to the low
17	number of datapoints; Student's t-test was used for other datasets.
18	
19	Results and discussion
20	Leafy liverwort-Pezoloma ericae symbiosis is mutualistic and mycorrhiza-like
21	In our radio-labeled liverwort microcosms, 1.786 ng g $^{-1}$ of 33 P was
22	assimilated from P. ericae into C. bicuspidata tissues (Fig. 2a, Table S1). There was
23	a significant difference in the liverwort ³³ P content and concentration between plant-
24	only microcosms with barriers and in plant-only microcosms without barriers (Mann-
	10

1 Whitney U = 0; $n_1=2$, $n_2=13$; P = 0.019 two-tailed (data not shown), further 2 demonstrating the efficacy of barriers in microcosms for preventing direct plant 3 access to ³³P-labeled wells. While we cannot exclude the possibility that the P 4 content of the Phytagel substrate used in our microcosms may have been directly 5 assimilated by the liverworts within the systems and therefore reduced plant demand 6 for fungal-acquired ³³P, our data unequivocally show movement of ³³P from the 7 fungus to the leafy liverwort.

8 The leafy liverworts transferred 0.019 ng (0.22 ng g^{-1}) of plant-fixed carbon 9 to their fungal partners (Fig. 2b); equivalent to 0.27% of the total amount of carbon 10 fixed during the labelling period (Table S1). By demonstrating unequivocal 11 exchange of fungal-acquired phosphorus and plant-fixed carbon between symbionts, 12 our results provide the first experimental evidence that associations between non-13 vascular plants and ascomycete fungal symbionts are nutritionally mutualistic. This 14 confirms previous hypotheses based on cytological evidence (van der Heijden et al. 15 2015) and culturing experiments (Duckett and Read, 1991; Upson et al., 2007; Kowal 16 et al. 2015). Additionally, liverworts resynthesized with P. ericae grew significantly 17 larger than liverworts grown under identical conditions without the fungus (P < 0.0118 for both, Fig. 3a,b). Together with our evidence of carbon-for-nutrient exchange 19 between symbionts, it is clear that the symbiosis between non-vascular leafy 20 liverworts and their fungal partners is mutualistic and mycorrhiza-like. This lays the 21 foundation to investigate further functional differences between liverwort-ascomycete 22 partnerships and previously documented exchanges between P. ericae and vascular 23 plants (Pearson and Read, 1973; Read and Stribley, 1973; Upson et al. 2008, Kowal 24 et al., 2015).

1	Our experiments aimed to uncover whether there is any movement of plant-
2	fixed carbon and fungal-acquired P between C. bicuspidata liverworts and P. ericae.
3	The evidence of ³³ P transfer from fungus to plant in our experimental microcosm
4	(Fig. 2a) strongly supports a nutritional role for P. ericae that is particularly
5	significant given that phosphorus can be one of the most limiting nutrients in the
6	ericaceous habitats where C. bicuspidata grows. It is possible that the enhanced
7	growth we observed in liverworts with fungal symbionts compared to those without
8	(Fig. 3) was a result of fungal remineralization of organic compounds leached from
9	the liverwort, rather than a direct result of increased uptake of fungal-acquired P.
10	However, the provision of ³³ P by the fungus, as shown here, supports the hypothesis
11	that the fungus plays a role in leafy liverwort P nutrition in natural environments.
12	
13	Evolutionary context
14	Our demonstration that the association between a leafy liverwort and the

15 ericoid mycorrhizal fungus P. ericae can be reciprocally beneficial provides 16 additional clues about the evolution of ErM, and lends further weight to the idea that 17 mutualisms between leafy liverworts and ascomycete fungi pre-dated their formation 18 in the Ericales, estimated at less than 100 MYA (Brundrett, 2004). Following their 19 discovery that ascomycetes belonging to the P. ericae group colonize the rhizoids of 20 Schistochilaceae liverworts, thought to have originated in the Triassic (Heinrichs et 21 al., 2007) as the sister group to all other fungus-containing lineages in the 22 Jungermanniales, Pressel et al. (2008) suggested that this association arose more than 23 250 MYA. Nonetheless, divergence time estimates for ascomycete lineages remain a 24 matter of debate particularly as the presumed recent origin of Leotiomycetes, the 25 clade containing P. ericae, is in the Cretaceous (James et al., 2006; Prieto and Wedin,

2013). Until ever-improving phylogenetic methods integrating fossil evidence
 (Beimforde et al., 2014) provide more accurate date estimates, recent origins for the
 association with multiple instances of host shifting between leafy liverworts and
 Ericaceae plants, remain a plausible alternative scenario (Selosse, 2005; Pressel et al.,
 2008).

6

7 Observations on previously reported liverwort-fungal symbioses

8 Although not directly comparable, given considerable differences in the 9 physiology of the liverwort hosts and in experimental design (i.e. axenic and edaphic 10 conditions), our finding of mycorrhizal-like associations between ascomycete fungi 11 and leafy liverworts now invites further comparisons with more ancient lineages of 12 liverworts, i.e. Haplomitriopsida (Treubia and Haplomitrium) and complex thalloid 13 liverworts (Neohodgsonia, Allisonia and Marchantia) and fungi - Mucoromycotina 14 and/or Glomeromycotina (Field et al. 2015a and 2016) (see Table S1; Fig. 1a). When 15 fungal-acquired ³³P uptake is normalized to biomass (Fig. 2a; Table S1), it appears C. bicuspidata gains significantly less ³³P from its fungal partner than Haplomitrium, 16 17 Treubia (Mucoromycotina only), Allisonia and Neohodgsonia (Mucoromycotina and 18 Glomeromycotina), but a roughly similar amount to Marchantia (Glomeromycotina 19 only). Cephalozia bicuspidata carbon allocation (in terms of absolute amount, Fig. 20 2b) to its symbiotic fungus also points to a relatively low carbon demand by P. ericae 21 on its liverwort host, especially when compared to other liverwort lineages (0.27% of 22 plant-fixed carbon transferred to the symbiotic fungus vs. 2.2% - 14.2% - see Table 23 S1). This may be partially influenced by the facultative biotrophic nature of P. ericae 24 allowing it to gain at least some organic carbon from dead organic matter, and the 25 extent of colonisation (i.e. restricted to rhizoids vs. extensive through thallus). It is

1 possible that the fungus in our experiments was able to derive some C directly from 2 the Phytagel substrate (which contains glucose and trace levels of nutrients; see 3 materials and methods). Regardless of the phosphorus-for-carbon measurements 4 presented in Table S1, it is evident that the relative "cost" of maintaining mycorrhiza-5 like symbioses varies between liverwort-fungal symbioses, with ErM-like 6 associations potentially requiring less liverwort investment in terms of photosynthate 7 allocation, and the liverwort-Mucoromycotina partnership requiring the most (Field et 8 al., 2016).

Having established that there is a mycorrhizal-like nutritional exchange
between the leafy liverwort C. bicuspidata and its fungal symbiont P. ericae,
investigations are now needed which include fungus-to-plant N transfer and more
natural experimental conditions to provide a robust platform to investigate functional
variation in plant-fungal symbioses across evolutionary lineages and ecological
gradients.

15

16 Future directions

17 The diversity of fungal symbionts across the land plant phylogeny is 18 becoming increasingly apparent; however, there is still a relative dearth of 19 information regarding the function of plant-fungal symbioses in many clades, 20 particularly the bryophytes. The largest remaining functional knowledge gaps now 21 are the basidiomycete symbioses in thalloid Aneuraceae and leafy Scapaniaceae and 22 Arnelliaceae liverworts (Bidartondo and Duckett, 2010). In pteridophytes only two 23 pioneering studies to date have demonstrated the reciprocal exchange of plant-fixed C 24 for fungal-acquired N and P between green sporophytes of Osmunda regalis and

Ophioglossum vulgatum and their Glomeromycotina symbionts (Field et al., 2012,
 2015c; Pressel et al., 2016).

3	Although we show transfer of C from C. bicuspidata to fungi in vitro, in	
4	habitats where C. bicuspidata grows, often in deep shade under a canopy of	
5	ericaceous plants, the fungal symbiont of C. bicuspidata may be getting a large	
6	proportion of its organic C from surrounding vascular plants via a shared fungal	
7	network, as demonstrated previously between Betula and the mycoheterotrophic	
8	liverwort Cryptothallus (Aneura) mirabilis via a shared basidiomycete fungus (Read	
9	et al., 2000; Bidartondo et al., 2003). How this affects ecosystem carbon and nutrien	
10	budgets in terms of storage and cycling remains to be uncovered.	
11	In conclusion, our findings provide a novel and important example of	
12	mycorrhizal functioning of an additional fungal lineage with non-vascular plants.	
13	Thus, our demonstration of symbiotic ErM-like functioning of P. ericae with a leafy	
14	liverwort, now adds the Ascomycota to the list of fungal groups engaging in	
15	mutualistic carbon-for-nutrient exchange across the land plant phylogeny. It has been	
16	proposed that C. bicuspidata liverworts with ErM fungal symbionts can be used to	
17	restore threatened heathlands by promoting establishment of native ericaceous plants	
18	(Kowal et al., 2015) that are limited by ErM inoculum availability (Diaz et al. 2006).	
19	Our results suggest that at least one of the mechanisms underpinning the potential	
20	role of C. bicuspidata in habitat restoration strategies is likely to be nutritional. The	
21	potential ecological applications of pioneer non-vascular plants as mycorrhizal	
22	reservoirs and vectors, particularly in habitat restoration, are only just becoming	
23	apparent. Evidence of a mutualism between a widespread non-vascular plant and	
24	ascomycete fungi that form mycorrhizas with dominant vascular plants should now	
25	encourage further ecological and physiological experiments.	

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Figures

Fig. 1 (a) Liverwort phylogram showing diversity in liverwort-fungal associations (after Crandall-Stotler et al., 2009) based on extensive sampling of nearly all the fungus-containing liverwort genera worldwide (Ligrone et al. 2007). (b) Cephalozia bicuspidata rhizoid resynthesized with Pezoloma ericae fungus (arrowed) in axenic culture.

Fig. 2 (a) Concentration of fungal-acquired 33 P in Cephalozia bicuspidata tissue (ng g⁻¹). (b) Total plant-derived carbon present in extraradical fungal mycelium after a five-hour labeling period (ng). Error bars denote 1 s.e. in both panels.

Fig. 3 (a) Change in surface area of leafy liverwort after six weeks growth with and without Pezoloma ericae fungal symbiont (n = 16, 22), P < 0.01^{**} (using t-test). (b) Total biomass of liverworts (dry weight; g) after six months of growth with and without P. ericae (Mann-Whitney U = 2; P = 0.0087^{**} ; n_{1,2} = 6 two-tailed). Error bars denote 1 s.e. in both panels.

Supplementary Figures and Data

Fig. S1 Diagrams and photographs of ³³P transfer experiment showing split plate experimental design set up before (a,b) and after (c,d) the fungus crossed the barrier. L – liverwort; H – hyphae; W – well and B – barrier.
Fig. S2 Diagram showing microcosm design for carbon transfer experiments.
(a) Cephalozia bicuspidata and Pezoloma ericae growing on 1.5% Phytagel (b) ¹⁴CO₂ released into chamber headspace, fixed by plant then transferred into

fungal hyphae (blue). Dotted black line shows placement of polythene with

lanolin seal in yellow. Polythene barrier minimizes ¹⁴CO₂ diffusion into Phytagel.

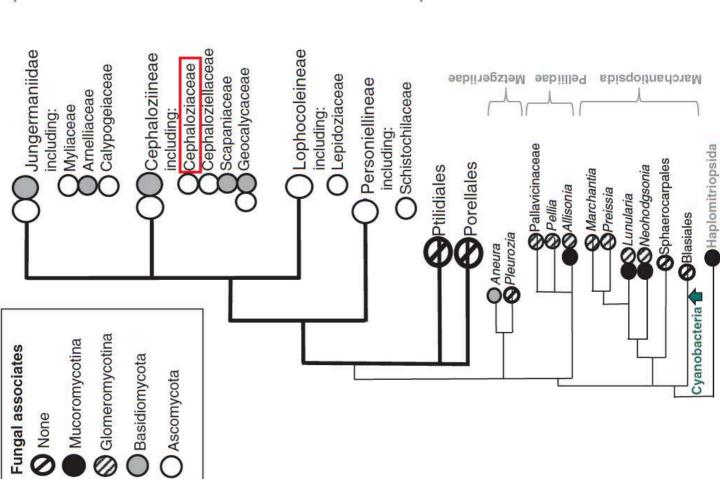
Fig. S3 Example of digitized pictures used to measure perimeter area of

liverworts. (a) Control without fungus (b) Inoculated with fungus (arrow).

Methods S1 Methods for harvesting plants and fungus for ³³P acid digestion, liquid scintillation and nutrient budgeting.

Table S1 Summary of carbon-for-nutrient exchange between *Cephalozia bicuspidata* and *Pezoloma ericae*, alongside data from previous studies of earlydiverging liverwort lineages and their fungal partners (below dotted line) (Field *et al.*, 2015a).

Table S2 Estimates of likely species numbers of liverworts worldwide with fungal symbionts.





Jungermaniidae

Jungermanniales

