

Table S1. Summary of evaluated morphological and physiological response variables for data analysis.

Model	Response variable group	Response variable	Error distribution (Power lambda link function) ¹
II	Morphology (biomass)	Roots (<2mm)	Gaussian (1)
		LMF	Gaussian (1)
		FRMF	Gaussian (1)
		R/S ratio	Gamma (0.182)
		FR/LB ratio	Gaussian (1)
III	Morphology (size)	Height (H)	Gaussian (1)
		Diameter (D)	Gaussian (1)
		H/D ratio	Gaussian (1)
IV	Physiology	PN	Gaussian (1)
		GS	Gamma (-0.101)
		Ci	Gaussian (1)
		ETR	Gamma (0.020)
		E	Gaussian (1)
		Pn/Gs ratio	Gaussian (1)
		Pn/E ratio	Gaussian (1)
		Ψ	Gamma (-0.062)

¹Power lambda link function [$g(\cdot) = \lambda^{\frac{1}{\lambda}} (\cdot - \lambda)^{-1}$] is the lambda (, numeric value inside the brackets) used for the monotonic transformations.

LMF: leaf mass fraction; FRMF: fine-root mass fraction; R/S ratio: root/shoot ratio; FR/LB ratio: fine-root-to-leaf biomass fraction; Height: height of the oak seedling; Diameter: basal diameter of the oak seedling; H/D ratio: height diameter ratio; Pn net photosynthesis; Gs: stomatal conductance; Ci: internal CO₂ concentration; ETR: electron transport rate; E: transpiration; Ψ: leaf water potential.

Table S2. Summary of Kaplan–Meier non-parametric models fitted to analyse the main factors—soil treatment and *P.c* infection—affecting the survival of oak seedlings.

Soil treatment	<i>P.c</i> infection	Time	N. risk	N. event	Survival	SE	Lower 95% CI	Upper 95% CI
Control	<i>P.c</i>	140	39	1	0.974	0.0253	0.926	1
		146	38	1	0.949	0.0353	0.882	1
		154	34	2	0.893	0.0507	0.799	0.998
		160	32	2	0.837	0.061	0.726	0.966
		162	25	2	0.77	0.0722	0.641	0.925
		166	23	2	0.703	0.0799	0.563	0.879
		168	18	3	0.586	0.0908	0.432	0.794
		174	6	2	0.391	0.128	0.206	0.743
0% BIO	<i>P.c</i>	56	52	1	0.981	0.019	0.944	1
		76	51	2	0.942	0.0323	0.881	1
		83	49	1	0.923	0.037	0.853	0.998
		89	48	3	0.865	0.0473	0.777	0.963
		105	45	2	0.827	0.0525	0.73	0.936
		112	43	3	0.769	0.0584	0.663	0.893
		119	40	3	0.712	0.0628	0.598	0.846
		133	37	3	0.654	0.066	0.537	0.797
		140	32	2	0.613	0.0679	0.493	0.762
		147	27	2	0.568	0.07	0.446	0.723
		155	24	1	0.544	0.071	0.421	0.703
		162	20	1	0.517	0.0725	0.393	0.68
		166	19	1	0.49	0.0736	0.365	0.657
		174	7	1	0.42	0.0904	0.275	0.64
Control	<i>P.c</i>	104	54	1	0.981	0.0183	0.946	1
		127	53	1	0.963	0.0257	0.914	1
		140	52	2	0.926	0.0356	0.859	0.998
		146	50	1	0.907	0.0394	0.833	0.988
		147	47	1	0.888	0.0431	0.808	0.977
		160	41	3	0.823	0.0538	0.724	0.936
		162	32	4	0.72	0.0673	0.6	0.865
		168	24	4	0.6	0.0784	0.465	0.775
12.5% BIO	<i>P.c</i>	70	56	1	0.982	0.0177	0.948	1
		74	55	1	0.964	0.0248	0.917	1
		83	54	2	0.929	0.0344	0.864	0.999
		89	52	1	0.911	0.0381	0.839	0.989
		90	51	1	0.893	0.0413	0.815	0.978
		92	50	2	0.857	0.0468	0.77	0.954
		96	48	1	0.839	0.0491	0.748	0.941
		97	47	2	0.804	0.0531	0.706	0.915
		103	45	1	0.786	0.0548	0.685	0.901
		105	44	2	0.75	0.0579	0.645	0.872
		109	42	1	0.732	0.0592	0.625	0.858
		111	41	4	0.661	0.0633	0.548	0.797
		119	37	3	0.607	0.0653	0.492	0.75
		125	34	1	0.589	0.0657	0.474	0.733
		127	33	1	0.571	0.0661	0.455	0.717
		133	32	2	0.536	0.0666	0.42	0.684
		147	28	2	0.497	0.0672	0.382	0.648

	154	22	1	0.475	0.0678	0.359	0.628
	168	15	3	0.38	0.0731	0.26	0.554
	105	54	1	0.981	0.0183	0.946	1
	123	50	1	0.962	0.0265	0.911	1
	127	47	1	0.941	0.0329	0.879	1
	139	46	1	0.921	0.038	0.849	0.999
	140	44	1	0.9	0.0425	0.82	0.987
Control	146	43	1	0.879	0.0464	0.793	0.975
	147	38	2	0.833	0.0543	0.733	0.946
	154	33	3	0.757	0.0646	0.641	0.895
	162	22	3	0.654	0.0786	0.517	0.828
	168	17	3	0.538	0.0886	0.39	0.743
	174	8	2	0.404	0.1059	0.242	0.675
	62	57	1	0.982	0.0174	0.949	1
	78	56	1	0.965	0.0244	0.918	1
	82	55	1	0.947	0.0296	0.891	1
25% BIO	83	54	1	0.93	0.0338	0.866	0.999
	89	53	1	0.912	0.0375	0.842	0.989
	90	52	1	0.895	0.0406	0.819	0.978
	103	51	1	0.877	0.0435	0.796	0.967
	104	50	1	0.86	0.046	0.774	0.955
	105	49	1	0.842	0.0483	0.753	0.942
P.c	106	48	1	0.825	0.0504	0.732	0.929
	111	47	1	0.807	0.0523	0.711	0.916
	112	46	1	0.789	0.054	0.69	0.903
	119	45	2	0.754	0.057	0.651	0.875
	127	43	2	0.719	0.0595	0.612	0.846
	140	39	1	0.701	0.0608	0.591	0.831
	154	32	1	0.679	0.0627	0.567	0.814
	160	31	1	0.657	0.0644	0.542	0.796
	161	26	1	0.632	0.0667	0.514	0.777
	168	22	4	0.517	0.0753	0.388	0.688

Bio 0% = seedlings that received no soil treatment, Bio 12.5% = seedlings that received 12.5% of Biohumim and Bio 25% = seedlings that received 25% of Biohumim. Control = seedlings were not infected by *P. cinnamomi* and P.c = seedlings were not infected by *P. cinnamomi*.

Table S3. Results of model selection of the generalised linear mixed models fitted to analyse the main factors—soil treatment (S) and biotic stress (B)—affecting different morphological response variables.

Response variable	S	B	S × B	AICc	deltaAICc
				25.1	0.00
	+			29.4	4.28
Root (<2mm) biomass		+		31.1	5.98
	+	+		35.4	10.28
	+	+	+	38.8	13.67
<i>Importance:</i>	0.05	0.11	<0.01		
	+	+	+	223.0	0.00
		+		223.5	0.51
Leaf mass fraction	+	+		224.8	1.8
				225.8	2.76
		+		227.2	4.16
<i>Importance:</i>	0.85	0.60	0.39		
				198.6	0.00
	+			199.5	0.93
Fine roots mass fraction		+		201.7	3.11
	+	+		202.6	4.01
	+	+	+	205.5	6.93
<i>Importance:</i>	0.19	0.40	0.02		
	+	+		82.3	0.00
		+		83.5	1.17
Root/Shoot ratio	+	+	+	83.7	1.35
				87.7	5.37
		+		87.8	5.48
<i>Importance:</i>	0.72	0.94	0.23		
				-3.9	0.00
	+			0.6	4.55
Fine roots/Leaf Biomass ratio		+		6.2	10.12
	+	+		10.7	14.61
	+	+	+	15.3	19.23
<i>Importance:</i>	<0.01	0.09	<0.01		
		+		367.5	0.00
	+	+		369.5	2.00
Height (H)				370.7	3.17
	+	+	+	371.3	3.79
		+		372.7	5.17
<i>Importance:</i>	0.84	0.33	0.08		
				180.3	0.00
	+			182.3	1.92
Basal diameter (D)		+		185.4	5.02
	+	+		187.3	6.93
	+	+	+	188.0	7.63
<i>Importance:</i>	0.09	0.29	0.02		
		+		191.3	0.00
	+	+	+	191.4	0.14
H/D ratio	+	+		193.6	2.28
				194.3	2.97
		+		196.2	4.93

<i>Importance:</i>	0.88	0.52	0.36
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Importance of predictor variable in the model averaging.

Maximal model = T × B + 1 | Mother tree.

Table S4. Results of model selection of the generalised linear mixed models fitted to analyse the main factors—soil treatment (S) and biotic stress (B)—affecting different physiological response variables.

Response variable	S	B	S × B	AICc	deltaAICc
Pn				328.7	0.00
	+			329.4	0.74
		+		331.0	2.29
	+	+		331.7	2.99
	+	+	+	333.2	4.54
<i>Importance:</i>		0.28	0.44	0.04	
Gs				-125.8	0.00
	+			-124.1	1.63
		+		-124.0	1.80
	+	+		-122.2	3.55
	+	+	+	-120.2	5.62
<i>Importance:</i>		0.31	0.32	0.03	
Ci				572.6	0.00
	+			574.2	1.60
		+		576.2	3.69
	+			577.9	5.30
	+	+	+	579.9	7.38
<i>Importance:</i>		0.15	0.69	0.01	
ETR				500.2	0.00
	+			502.1	1.90
		+		504.5	4.28
	+	+		506.3	6.12
	+		+	508.3	8.09
<i>Importance:</i>		0.96	0.33	0.07	
E				165.7	0.00
	+			167.7	2.05
		+		170.0	4.28
	+	+		172.1	6.39
	+	+	+	172.3	6.61
<i>Importance:</i>		0.13	0.28	0.02	
Pn/Gs				515.1	0.00
	+			516.2	1.03
		+		518.6	3.47
	+			519.7	4.53
	+	+	+	522.4	7.29
<i>Importance:</i>		0.16	0.63	0.01	
Pn/E				178.8	0.00
	+			180.9	2.07
		+		183.6	4.80
	+	+		185.7	6.86
	+	+	+	187.8	9.01
<i>Importance:</i>		0.09	0.27	<0.01	
Ψ				393.0	0.00
	+			395.1	2.13
	+			396.0	2.95

+	399.9	6.90
	402.5	9.44
<i>Importance:</i> 0.85 0.97 0.21		

Importance of predictor variable in the model averaging.

Maximal model = $T \times B + 1 |$ Mother tree.

Pn net photosynthesis; Gs: stomatal conductance; Ci: internal CO₂ concentration; ETR: electron transport rate; E: transpiration; Ψ: leaf water potential.