

Supplementary Material

Responses of fine-root biomass and production to drying depend on wetness and site nutrient regime in boreal forested peatland

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2 Supplementary Tables

Supplementary Table S1. FRB and FRP means, standard errors (SE), and numbers of samples (N). Due to the hierarchical structure of the data, SEs were estimated with linear mixed models.

Site	FRB, mean	FRB, SE	FRB, N	FRP, mean	FRP, SE	FRP, N	FRP/FRB
DP	191.8	34.4	9	95.5	28.7	14	0.50
DPdr	179.9	27.7	9	92.7	13.3	16	0.52
TP	250.3	41.2	9	118.4	25.0	16	0.47
TPdr	120.6	17.6	9	119.9	11.2	18	0.99
VS	205.7	49.2	9	42.3	12.6	14	0.20
VSdr	267.3	26.8	9	70.0	14.5	16	0.26
HS	155.5	31.6	9	53.6	10.8	9	0.34
HSdr	342.2	104.2	9	30.6	7.3	17	0.09

Supplementary Table S2. Results of the t-tests.

S2a) The mean differences between the undrained and drained pairs of the same site type for FRB (diameter ≤ 0.5 mm), with independent samples t-tests. Results with $p < 0.05$ are marked with *. ANOVA between sites for FRB showed a significant site effect ($F(7, 64) = 3.655$, $p = 0.002$).

FRB	T-test						
	N	Mean	SD	SE	df	t	p
DP	9	191.8	78.7	26.2			
DPdr	9	179.9	63.3	21.1	16	0.35	0.729
TP	9	250.3	123.5	41.2			
TPdr	9	120.6	39.8	13.3	16	3.00	0.008*
VS	9	205.7	131.2	43.7			
VSdr	9	267.3	80.5	26.8	16	-1.20	0.248
HS	9	155.5	94.7	31.6			
HSdr	9	342.2	193.0	64.3	16	-2.61	0.019*

S2c) The mean differences between the undrained and drained pairs of the same site type for FRP with independent samples t-tests. No p-values < 0.05 were found. ANOVA between sites for FRP showed a significant site effect ($F(7, 112) = 6.324$, $p = 0.001$).

FRP						T-test		
	N	Mean	SD	SE		df	t	p
DP	14	95.45	82.26	21.98				
DPdr	16	92.67	42.83	10.71		28	0.12	0.907
TP	16	118.38	66.49	16.62				
TPdr	18	119.94	47.57	11.21		32	-0.08	0.937
VS	14	42.28	47.02	12.57				
VSdr	16	69.98	58.04	14.51		28	-1.42	0.166
HS	9	53.58	32.42	10.81				
HSdr	17	30.64	30.07	7.29		24	1.80	0.084

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Response variable	Fixed effects estimates		Subplot level:								Plot level:		Random effects estimates		
			Pine basal area, m ²	Spruce basal area, m ²	Birch basal area, m ²	Alder basal area, m ²	Graminoid cov. %	Forbs cov. %	WT 2016&2017 July and August mean, cm	Soil temp. 5 cm, cum. mean 2016&2017	C/N ratio	K mg g ⁻¹	site	plot nested within site	residual
FRB, Forbs	Parameter	5.15					1.04						parameter is redundant	parameter is redundant	1053.42
	SE	5.06					0.42								190.74
	p	0.31					0.02								0.00
FRB, Shrubs	Parameter	-184.72								0.09	1.81		44.25	parameter not in the model	384.43
	SE	66.49								0.04	0.51		59.97		79.38
	p	0.01								0.04	0.01		0.46		0.00
FRB, Pine	Parameter	-66.91	3.70								3.05		parameter is redundant	497.92	1581.54
	SE	24.41	0.79								0.88			360.55	345.12
	p	0.01	0.00								0.00			0.17	0.00
FRB, Spruce	Parameter	-18.99			7.35	20.96							16740.07	673.72	3392.87
	SE	51.19			2.75	3.76							10696.11	792.00	740.38
	p	0.72			0.02	0.00							0.12	0.40	0.00
FRB, Birch	Parameter	-22.28		0.90	3.57				-1.25				18.03	74.01	256.77
	SE	10.35		0.39	0.73				0.51				60.75	74.01	60.52
	p	0.07		0.05	0.00				0.05				0.77	0.32	0.00
FRB, Alder	Parameter	-45.76						-2.54				207.28	383.38	parameter not in the model	276.24
	SE	20.12						0.41				58.96	263.83		52.70
	p	0.07						0.00				0.02	0.15		0.00

Response variable	Fixed effects estimates		Subplot level:								Plot level:		Random effects estimates		
			Pine basal area, m ²	Spruce basal area, m ²	Birch basal area, m ²	Alder basal area, m ²	Graminoid cov. %	Forbs cov. %	WT 2016&2017 July and August mean, cm	Soil temp. 5 cm, cum. mean 2016&2017	C/N ratio	K mg g ⁻¹	site	plot nested within site	residual
FRP, all roots	Intercept	48.70	2.94										295.04	323.00	2524.17
	Parameter	SE	13.25	0.92									362.63	304.12	385.43
	p	0.01	0.01										0.42	0.29	0.00
FRP, Conifers (FTIR)	Parameter	21.51	1.33										parameter is redundant	31.59	721.31
	SE	4.57	0.32										63.01	117.16	
	p	0.00	0.00										0.62	0.00	
FRP, Shrubs and Birch (FTIR)	Parameter	10.29	1.75										63.42	parameter not in the model	340.20
	SE	5.01	0.34										58.97		51.85
	p	0.08	0.00										0.28		0.00
FRP, Graminoids (FTIR)	Parameter	0.73					1.73						18.10	25.72	296.48
	SE	3.26					0.25						29.88	32.00	47.65
	p	0.83					0.00						0.55	0.42	0.00
FRP, Forbs (FTIR)	Parameter	2.88					0.44						1.76	3.25	66.65
	SE	1.32					0.10						5.15	6.21	10.71
	p	0.07					0.00						0.73	0.60	0.00