



*Supplement of*

## **Covariation of redox potential profiles and the water table level at peatland sites representing different drainage regimes: implications for ecological modelling**

**Markku Koskinen et al.**

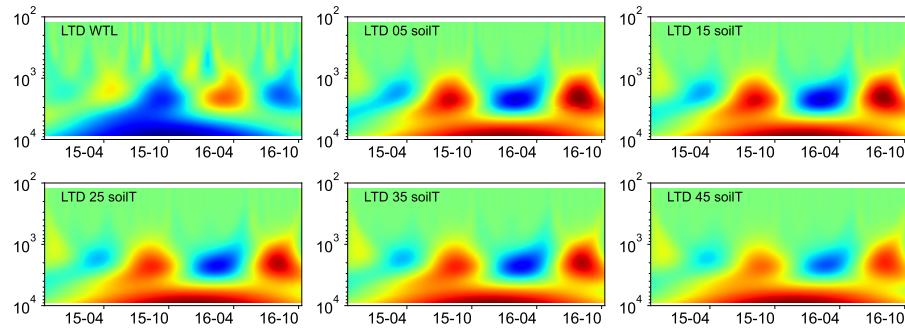
*Correspondence to:* Markku Koskinen (markku.koskinen@helsinki.fi)

The copyright of individual parts of the supplement might differ from the article licence.

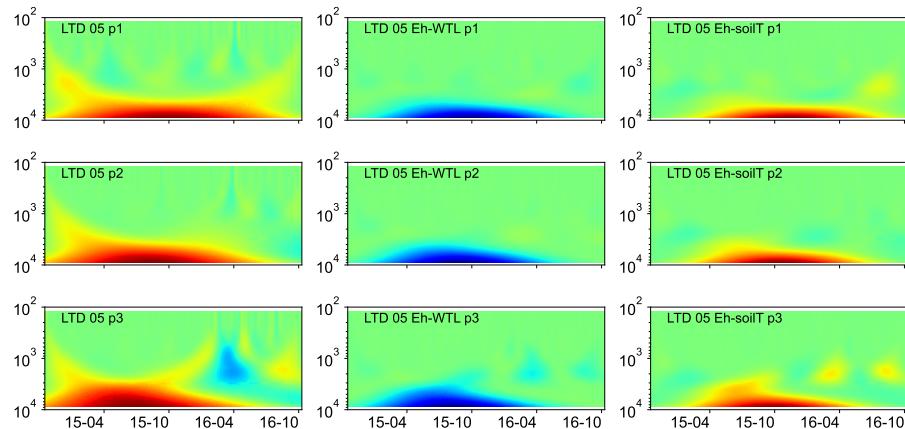
## S1. CWT figures

In all figures, the y-axis represents wavelet period in hours; notice that the axis is logarithmic and inverted (short periods, i.e. higher frequency is up). X-axis represents the date, formatted as the last two digits of year–month. Colour indicates wavelet coherence; dark red indicates strong synphase coherence, and dark blue indicates strong anti-phase coherence.

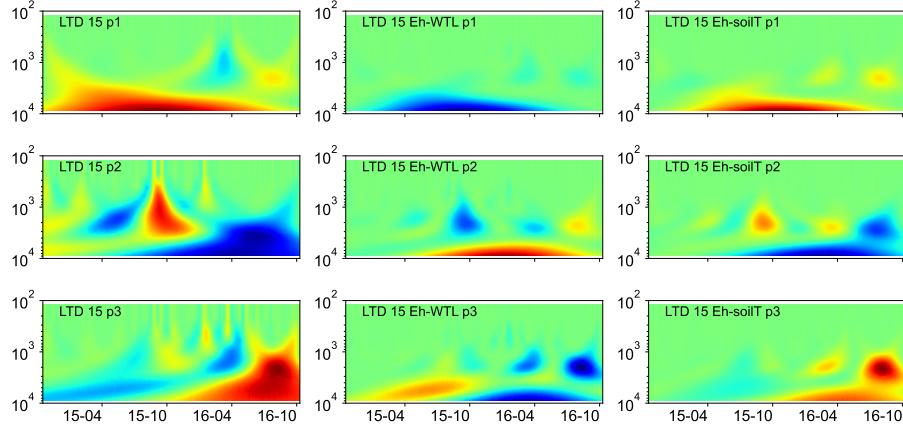
535 LTD



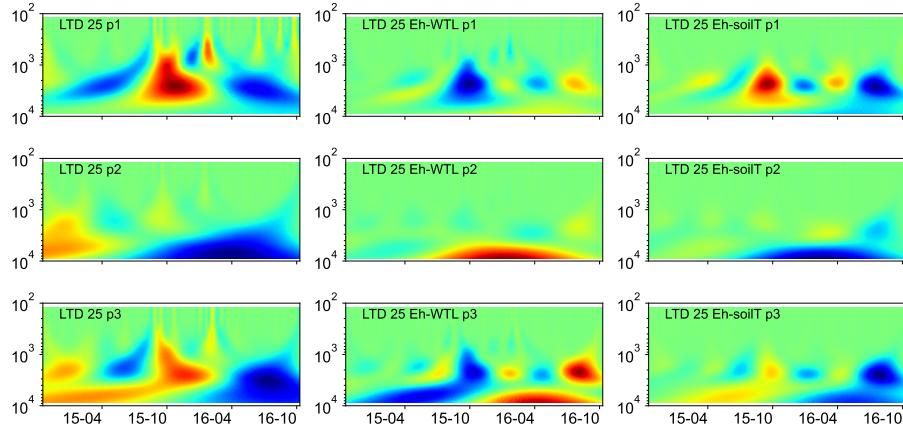
**Figure S1.** Left to right, top to bottom: Mexican hat wavelet coherence with 1) WTL data from the LTD plot; 2) soil temperature (soilT) at 5 cm depth; 3) soilT at 15 cm depth, 4) soilT at 25 cm depth; 5) soilT at 35 cm depth; 6) soilT at 45 cm depth.



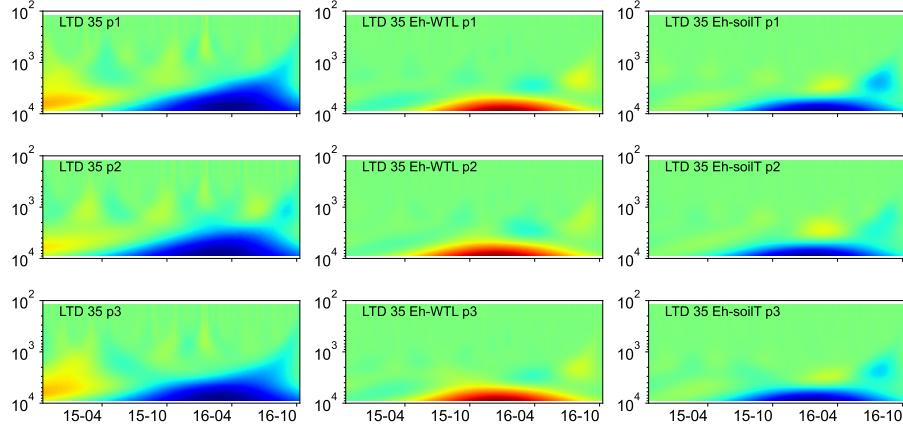
**Figure S2.** Left: Wavelet coherence between Mexican hat wave and  $E_h$  at LTD plot, 5 cm depth; centre: interaction between  $E_h$  and WTL wavelets; right: interaction between  $E_h$  and soil temperature wavelets at LTD plot, 5 cm depth. three probes plotted separately.



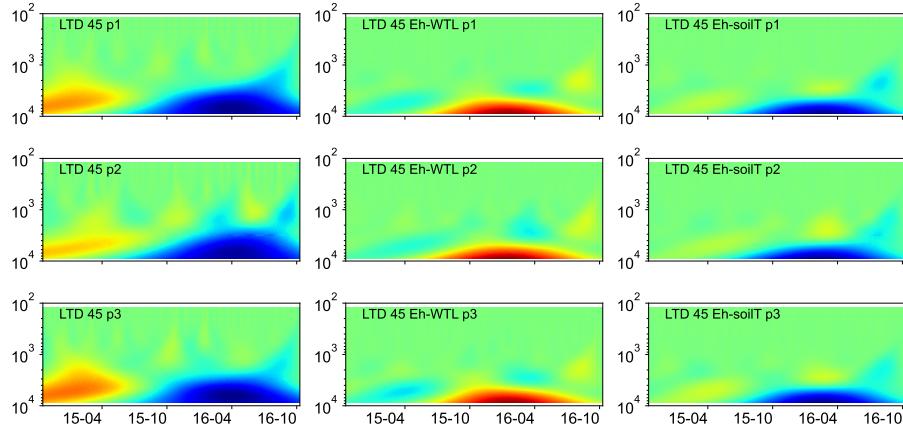
**Figure S3.** Left: Wavelet coherence between Mexican hat wave and  $E_h$  at LTD plot, 15 cm depth; centre: interaction between  $E_h$  and WTL wavelets; right: interaction between  $E_h$  and soil temperature wavelets at LTD plot, 15 cm depth. three probes plotted separately.



**Figure S4.** Left: Wavelet coherence between Mexican hat wave and  $E_h$  at LTD plot, 25 cm depth; centre: interaction between  $E_h$  and WTL wavelets; right: interaction between  $E_h$  and soil temperature wavelets at LTD plot, 25 cm depth. three probes plotted separately.

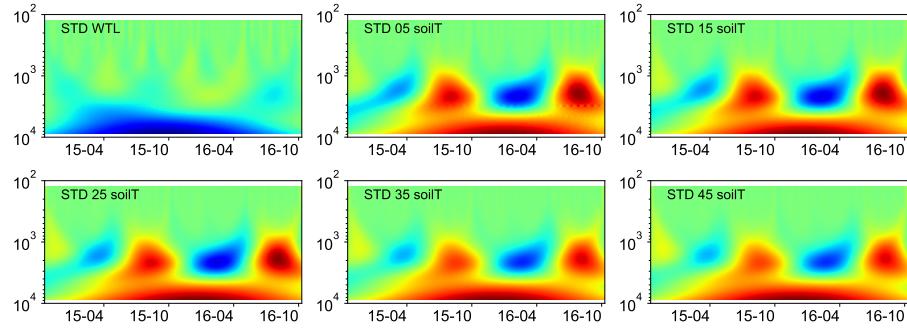


**Figure S5.** Left: Wavelet coherence between Mexican hat wave and  $E_h$  at LTD plot, 35 cm depth; centre: interaction between  $E_h$  and WTL wavelets; right: interaction between  $E_h$  and soil temperature wavelets at LTD plot, 35 cm depth. three probes plotted separately.

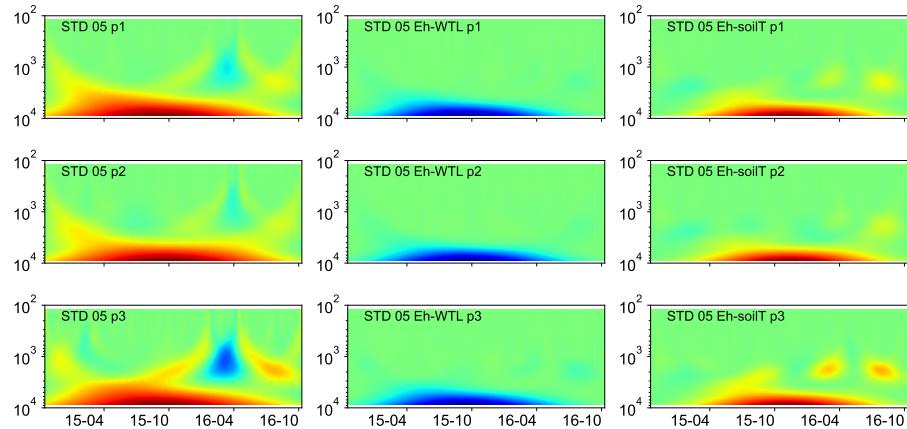


**Figure S6.** Left: Wavelet coherence between Mexican hat wave and  $E_h$  at LTD plot, 45 cm depth; centre: interaction between  $E_h$  and WTL wavelets; right: interaction between  $E_h$  and soil temperature wavelets at LTD plot, 45 cm depth. three probes plotted separately.

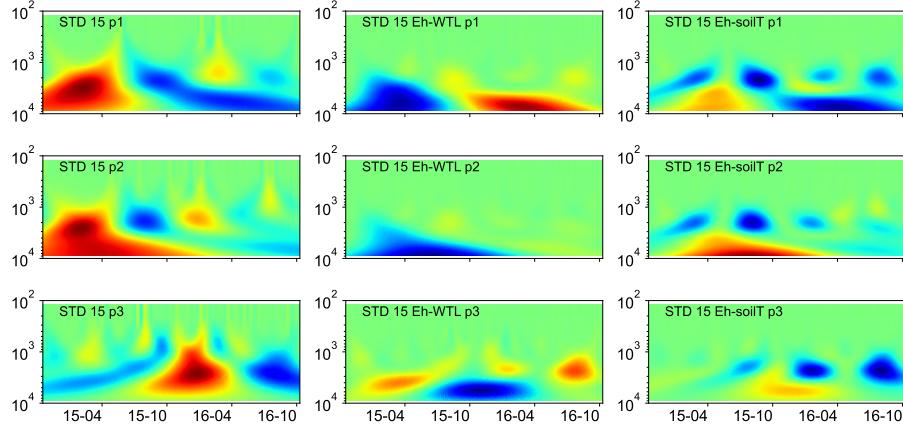
## STD



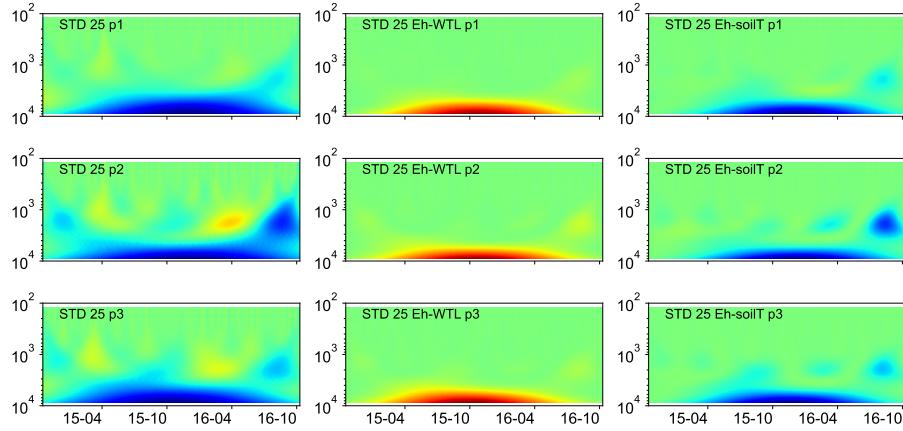
**Figure S7.** Left to right, top to bottom: Mexican hat wavelet coherence with 1) WTL data from the STD plot; 2) soil temperature (soilT) at 5 cm depth; 3) soilT at 15 cm depth, 4) soilT at 25 cm depth; 5) soilT at 35 cm depth; 6) soilT at 45 cm depth.



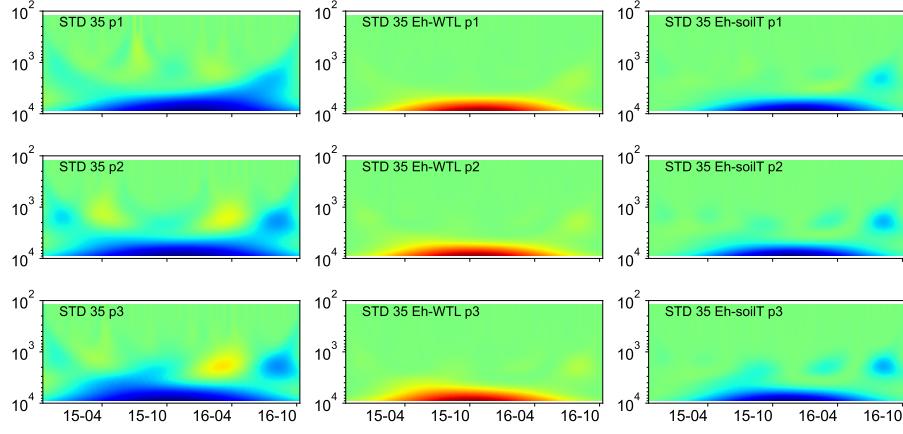
**Figure S8.** Left: Wavelet coherence between Mexican hat wave and  $E_h$  at the STD plot (5 cm depth); centre: interaction between  $E_h$  and WTL wavelets; right: interaction between  $E_h$  and soil temperature wavelets at the STD plot (5 cm depth). Three probes plotted separately.



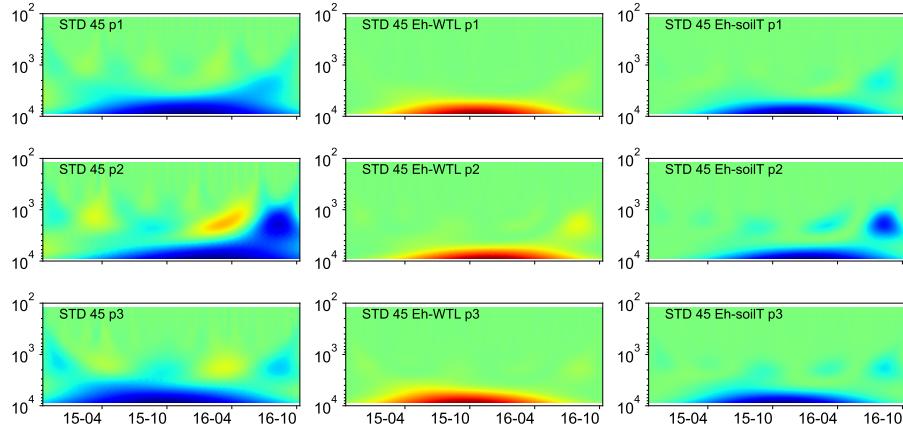
**Figure S9.** Left: Wavelet coherence between Mexican hat wave and  $E_h$  at STD plot, 15 cm depth; centre: interaction between  $E_h$  and WTL wavelets; right: interaction between  $E_h$  and soil temperature wavelets at STD plot, 15 cm depth. three probes plotted separately.



**Figure S10.** Left: Wavelet coherence between Mexican hat wave and  $E_h$  at the STD plot (25 cm depth); centre: interaction between  $E_h$  and WTL wavelets; right: interaction between  $E_h$  and soil temperature wavelets at the STD plot (25 cm depth). Three probes plotted separately.

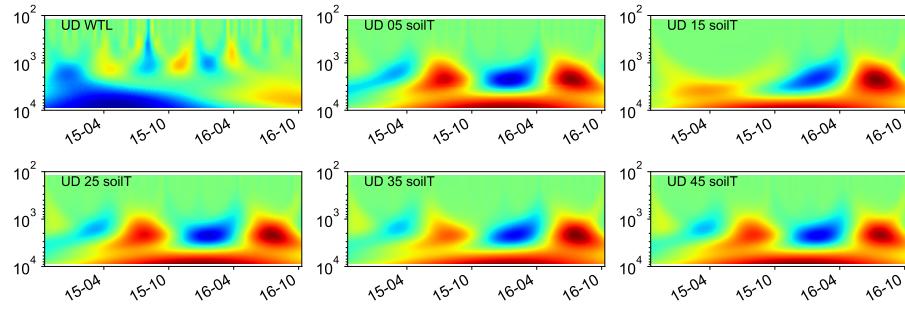


**Figure S11.** Left: Wavelet coherence between Mexican hat wave and  $E_h$  at STD plot, 35 cm depth; centre: interaction between  $E_h$  and WTL wavelets; right: interaction between  $E_h$  and soil temperature wavelets at STD plot, 35 cm depth. three probes plotted separately.

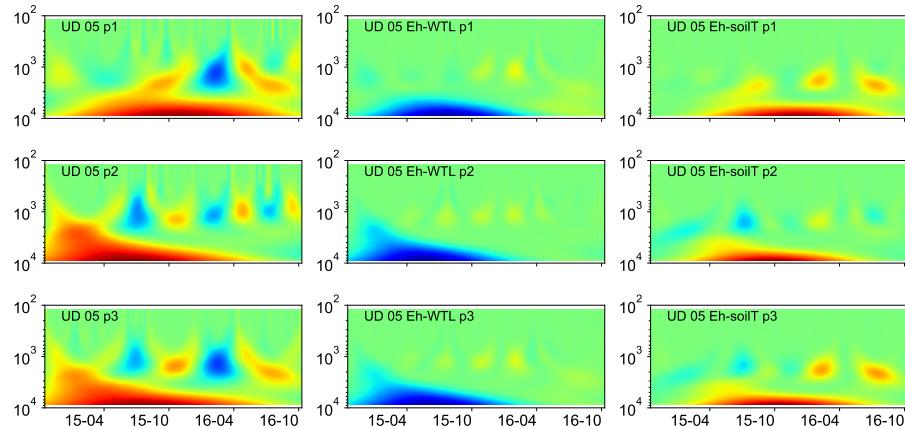


**Figure S12.** Left: Wavelet coherence between Mexican hat wave and  $E_h$  at the STD plot (45 cm depth); centre: interaction between  $E_h$  and WTL wavelets; right: interaction between  $E_h$  and soil temperature wavelets at the STD plot (45 cm depth). Three probes plotted separately.

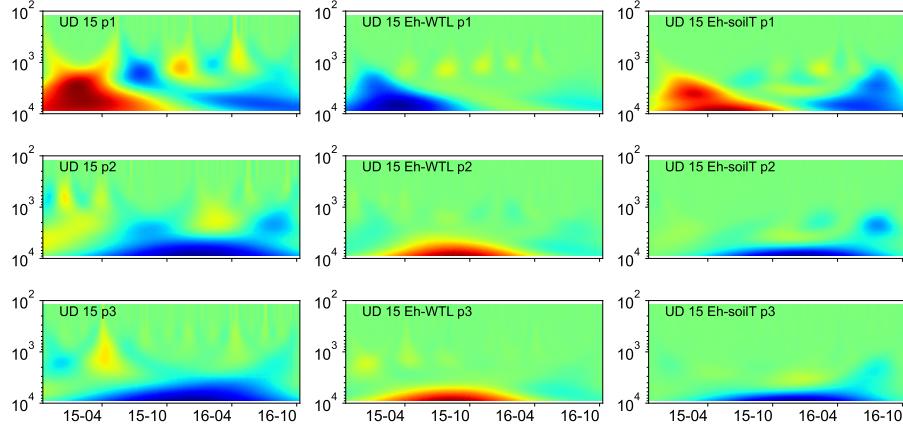
## UD



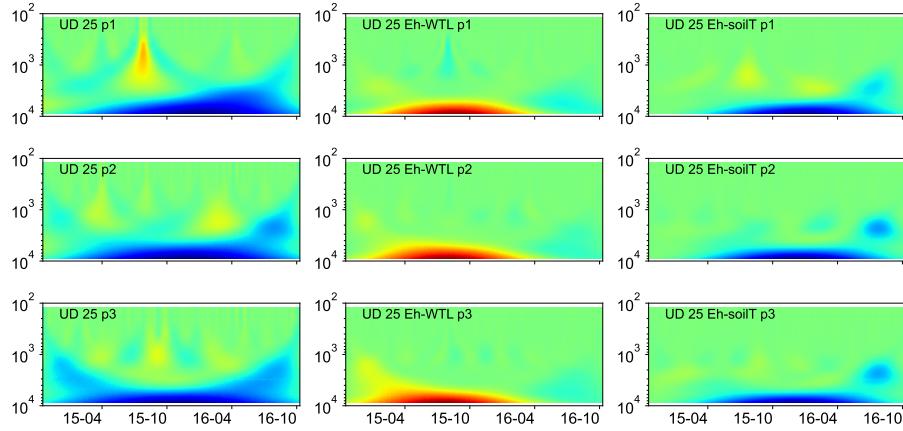
**Figure S13.** Left to right, top to bottom: Mexican hat wavelet coherence with 1) WTL data from the UD plot; 2) soil temperature (soilT) at 5 cm depth; 3) soilT at 15 cm depth, 4) soilT at 25 cm depth; 5) soilT at 35 cm depth; 6) soilT at 45 cm depth.



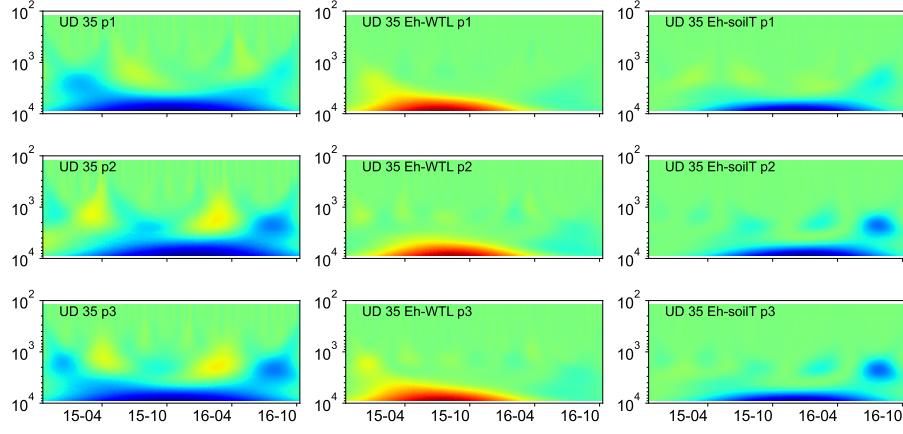
**Figure S14.** Left: Wavelet coherence between Mexican hat wave and  $E_h$  at the UD plot (5 cm depth); centre: interaction between  $E_h$  and WTL wavelets; right: interaction between  $E_h$  and soil temperature wavelets at the UD plot (5 cm depth). Three probes plotted separately.



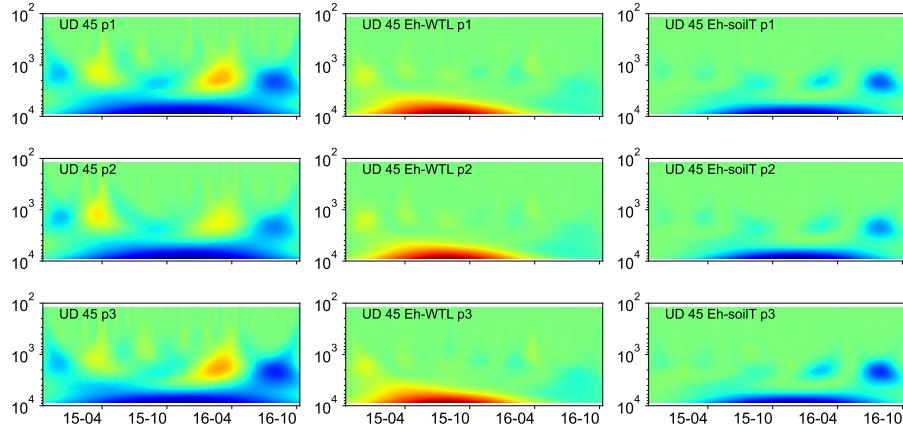
**Figure S15.** Left: Wavelet coherence between Mexican hat wave and  $E_h$  at UD plot, 15 cm depth; centre: interaction between  $E_h$  and WTL wavelets; right: interaction between  $E_h$  and soil temperature wavelets at UD plot, 15 cm depth. three probes plotted separately.



**Figure S16.** Left: Wavelet coherence between Mexican hat wave and  $E_h$  at the UD plot (25 cm depth); centre: interaction between  $E_h$  and WTL wavelets; right: interaction between  $E_h$  and soil temperature wavelets at the UD plot (25 cm depth). Three probes plotted separately.



**Figure S17.** Left: Wavelet coherence between Mexican hat wave and  $E_h$  at UD plot, 35 cm depth; centre: interaction between  $E_h$  and WTL wavelets; right: interaction between  $E_h$  and soil temperature wavelets at UD plot, 35 cm depth. three probes plotted separately.



**Figure S18.** Left: Wavelet coherence between Mexican hat wave and  $E_h$  at the UD plot (45 cm depth); centre: interaction between  $E_h$  and WTL wavelets; right: interaction between  $E_h$  and soil temperature wavelets at the UD plot (45 cm depth). Three probes plotted separately.

## S2. Enzyme activity tables

Enzyme activity presented as  $\mu\text{g}$  L-tyrosine for Protease and  $\text{NH}_4\text{-N}$  for Urease and Arg.Deaminase,  $\text{h}^{-1}$  g $^{-1}$  d.w. soil,  
540 respectively.

**Table S1.** Enzyme activity at the three mesotrophic study sites during Autumn 2014 (n=9; mean values  $\pm$  standard error (SE); data subjected to Tukey HSD tests and Box-Cox transformation; different letters express statistically significant results, where lowercase letters (a) refer to mean comparisons of homogeneous groups vertically, and uppercase letters (A) refer to horizontally between surfaces.

Treatment	Long-term Drainage (LTD)	Short-term Drainage (STD)	Undrained (UD)
Depth, cm	Protease	Protease	Protease
0-10	219.4 $\pm$ 11.4aA	428.0 $\pm$ 17.8aB	1519.2 $\pm$ 22.9aC
10-20	128.1 $\pm$ 14.4bA	410.5 $\pm$ 40.9abB	590.7 $\pm$ 12.1bC
20-30	125.7 $\pm$ 19.6bA	347.8 $\pm$ 17.5bB	358.8 $\pm$ 9.9cB
30-40	217.4 $\pm$ 23.9aA	220.6 $\pm$ 29.4cA	324.7 $\pm$ 14.0cB
40-50	202.5 $\pm$ 20.7aA	198.5 $\pm$ 26.4cA	314.5 $\pm$ 27.7cB
Depth, cm	Urease	Urease	Urease
0-10	127.2 $\pm$ 14.6aA	193.0 $\pm$ 17.2aB	1119.6 $\pm$ 10.1aC
10-20	96.6 $\pm$ 8.2aA	74.5 $\pm$ 11.3bA	339.9 $\pm$ 68.4bB
20-30	37.9 $\pm$ 3.1bA	49.0 $\pm$ 7.1bcA	66.4 $\pm$ 19.1cA
30-40	27.2 $\pm$ 4.0bcA	39.0 $\pm$ 3.5cA	71.4 $\pm$ 11.3cB
40-50	19.7 $\pm$ 6.0cA	41.4 $\pm$ 6.4cA	35.1 $\pm$ 8.1cA
Depth, cm	Arginine deaminase	Arginine deaminase	Arginine deaminase
0-10	33.5 $\pm$ 5.1aA	46.6 $\pm$ 8.3aA	31.8 $\pm$ 2.6aA
10-20	30.5 $\pm$ 2.9aA	50.8 $\pm$ 4.7aB	27.4 $\pm$ 1.7aA
20-30	42.8 $\pm$ 6.9aAB	58.1 $\pm$ 5.1aB	31.8 $\pm$ 1.1aC
30-40	32.6 $\pm$ 2.4aA	59.9 $\pm$ 1.7aB	56.5 $\pm$ 3.5bB
40-50	50.2 $\pm$ 5.7aA	108.6 $\pm$ 11.4bC	72.5 $\pm$ 5.9bB

**Table S2.** Enzyme activity at the three mesotrophic study sites during Spring 2015 (n=9; mean values  $\pm$  standard error (SE); data subjected to Tukey HSD tests and Box-Cox transformation; different letters express statistically significant results, where lowercase letters (a) refer to mean comparisons of homogeneous groups vertically, and uppercase letters (A) refer to horizontally between surfaces.

Treatment	Long-term Drainage (LTD)	Short-term Drainage (STD)	Undrained (UD)
Depth	Protease	Protease	Protease
0-10	282.8 $\pm$ 7.2aA	498.4 $\pm$ 21.0aB	1065.7 $\pm$ 34.5aC
10-20	164.6 $\pm$ 10.2bA	377.1 $\pm$ 17.8bB	700.6 $\pm$ 24.2bC
20-30	62.7 $\pm$ 6.4cA	281.0 $\pm$ 20.6cB	336.8 $\pm$ 12.2cB
30-40	113.9 $\pm$ 7.9dA	191.1 $\pm$ 10.2dB	298.4 $\pm$ 15.9cdC
40-50	213.6 $\pm$ 8.4eA	206.8 $\pm$ 21.4dA	247.1 $\pm$ 16.2dA
Depth	Urease	Urease	Urease
0-10	353.8 $\pm$ 25.8aA	446.8 $\pm$ 14.9aB	1366.6 $\pm$ 25.9aC
10-20	66.2 $\pm$ 9.7bA	38.6 $\pm$ 7.4cA	180.7 $\pm$ 8.1bB
20-30	27.9 $\pm$ 7.6cdA	53.5 $\pm$ 6.0bB	143.1 $\pm$ 9.6cC
30-40	39.4 $\pm$ 2.9bcA	17.6 $\pm$ 3.7cB	53.7 $\pm$ 5.3dA
40-50	17.4 $\pm$ 3.1dA	47.9 $\pm$ 3.2cB	53.5 $\pm$ 3.7dB
Depth	Arginine deaminase	Arginine deaminase	Arginine deaminase
0-10	11.0 $\pm$ 0.6aA	12.2 $\pm$ 1.2abA	7.4 $\pm$ 0.3aB
10-20	4.2 $\pm$ 0.3cA	6.2 $\pm$ 0.2cB	4.4 $\pm$ 0.5cA
20-30	3.2 $\pm$ 0.6cA	7.1 $\pm$ 0.5cB	8.4 $\pm$ 0.4aB
30-40	9.0 $\pm$ 0.8abA	10.0 $\pm$ 0.4bAB	11.1 $\pm$ 0.4bB
40-50	7.0 $\pm$ 0.5bA	13.1 $\pm$ 0.4aC	10.9 $\pm$ 0.2bB

**Table S3.** Enzyme activity at the three mesotrophic study sites during Summer 2015 (n=9; mean values ± standard error (SE); data subjected to Tukey HSD tests and Box-Cox transformation; different letters express statistically significant results, where lowercase letters (a) refer to mean comparisons of homogeneous groups vertically, and uppercase letters (A) refer to horizontally between surfaces.

Treatment	Long-term Drainage (LTD)	Short-term Drainage (STD)	Undrained (UD)
Depth	Protease	Protease	Protease
0-10	296.7±13.1aA	700.1±20.1aB	1605.6±34.9aC
10-20	255.3±3.6bA	362.3±14.9bB	786.5±30.0bC
20-30	115.1±5.2cA	350.0±11.4bB	308.3±14.6cB
30-40	133.8±5.7cA	271.1±11.3cB	360.9±17.8cC
40-50	133.9±5.0cA	256.2±11.0cC	214.95±10.3dB
Depth	Urease	Urease	Urease
0-10	431.0±13.7aB	162.5±7.7aA	1124.3±13.8aC
10-20	90.6±5.6bB	40.9±3.5bA	147.7±8.3bC
20-30	71.5±4.9bB	40.8±4.5bA	53.2±2.8cA
30-40	25.3±2.7cB	48.8±2.5bA	47.9±2.4cA
40-50	6.9±1.2dA	49.7±2.3bC	35.3±2.9dB
Depth	Arginine deaminase	Arginine deaminase	Arginine deaminase
0-10	12.8±0.8aA	8.8±1.0aB	7.5±0.4aB
10-20	6.0±0.8bA	9.4±0.6aB	8.6±0.6abB
20-30	7.6±0.3cA	9.1±0.3aB	8.4±0.3abAB
30-40	7.3±0.3cB	10.8±0.6aA	10.3±0.1cA
40-50	7.6±0.7cA	13.9±0.4bB	9.3±0.5bcA

**Table S4.** Correlations of  $E_h$  and enzyme activities within profiles, between depths. Year 2014 – Autumn, Mesotrophic fen. Treatment LTD and STD. Value for  $E_h$  is the mean of two weeks preceding the sampling. Correlations are significant at  $p < 0.05$ , N=5. Statistically significant correlations are designated \*\*.

Variable-Treatment						
$E_h$ -LTD	1					
Protease-LTD	-0.5999	1.0000				
Urease-LTD	0.7478	-0.0198	1.0000			
Arg. deam.-LTD	-0.4915	0.0133	-0.6312	1.0000		
$E_h$ -STD	0.4543	0.4167	0.8642	-0.4329	1.0000	
Protease-STD	<b>0.9645**</b>	-0.4144	<b>0.8956**</b>	-0.5579	0.6454	1.0000
Urease-STD	0.5318	0.3229	<b>0.8883**</b>	-0.4012	<b>0.9933**</b>	0.7087
Arg. deam.-STD	-0.7859	0.2471	-0.6702	0.8563	-0.4869	-0.7738
						-0.4998

**Table S5.** Correlations of  $E_h$  and enzyme activities within profiles, between depths. Year 2015 – Spring, Mesotrophic fen. Treatment LTD, STD and UD.  $E_h$  is the mean of two weeks preceding the sampling. Correlations are significant at  $p < 0.05$ , N=5. Statistically significant correlations are designated \*\*.

Variable-Treatment						
$E_h$ -LTD	1					
Protease-LTD	0.875	1				
Urease-LTD	<b>0.9178**</b>	0.7456	1			
Arg. deam.-LTD	0.5342	0.6857	0.6831	1		
$E_h$ -STD	<b>0.9482**</b>	0.7124	0.878	0.4201	1	
Protease-STD	<b>0.9102**</b>	0.5976	0.8679	0.2638	<b>0.9587**</b>	1
Urease-STD	<b>0.8961**</b>	0.7527	<b>0.9855**</b>	0.6694	0.8115	0.8296
Arg. deam.-STD	0.3318	0.686	0.3718	0.7731	0.0561	-0.0465
$E_h$ -UD	<b>0.9794**</b>	0.7774	<b>0.9449**</b>	0.5228	<b>0.9322**</b>	<b>0.9529**</b>
Protease-UD	<b>0.9425**</b>	0.6774	0.9125**	0.4261	<b>0.9924**</b>	<b>0.9773**</b>
Urease-UD	<b>0.9098**</b>	0.7316	<b>0.9955**</b>	0.6465	0.8552	0.8694
Arg. deam.-UD	-0.5038	-0.1413	-0.3124	0.3562	-0.685	-0.734

**Table S6.** Correlations of  $E_h$  and enzyme activities within profiles, between depths. Year 2015 – Summer, Mesotrophic fen. Treatment LTD, STD and UD. Value for  $E_h$  is the mean of two weeks preceding the sampling. Correlations are significant at  $p < 0.05$ , N=5. Statistically significant correlations are designated \*\*.

Variable-Treatment					
$E_h$ -LTD	1				
Protease-LTD	<b>0.9360**</b>	1			
Urease-LTD	<b>0.9499**</b>	0.8062	1		
Arg. deam.-LTD	0.8056	0.5478	<b>0.9141**</b>	1	
 <b>5</b>					
$E_h$ -STD	<b>0.9705**</b>	0.8264	<b>0.9879**</b>	<b>0.9223**</b>	1
Protease-STD	<b>0.9431**</b>	0.8079	<b>0.9973***</b>	<b>0.8926***</b>	<b>0.9767**</b>
Urease-STD	<b>0.9105**</b>	0.7128	<b>0.9629***</b>	<b>0.9750***</b>	<b>0.9807**</b>
Arg. deam.-STD	-0.4707	-0.5141	-0.5736	-0.3016	-0.4545
				-0.6145	-0.3646
$E_h$ -UD	<b>0.8984**</b>	0.6922	<b>0.9777***</b>	<b>0.9769***</b>	<b>0.9670***</b>
Protease-UD	<b>0.9866**</b>	<b>0.9292**</b>	<b>0.9636***</b>	<b>0.7979</b>	<b>0.9638***</b>
Urease-UD	<b>0.9552**</b>	0.7976	<b>0.9913***</b>	<b>0.9393***</b>	<b>0.9978***</b>
Arg. deam.-UD	-0.747	-0.6761	-0.7848	-0.6128	-0.7375
				-0.8224	-0.6433
					-0.5906
					-0.7322
					-0.7385
					-0.7233