

APPENDIX S2: SUPPLEMENTARY RESULTS

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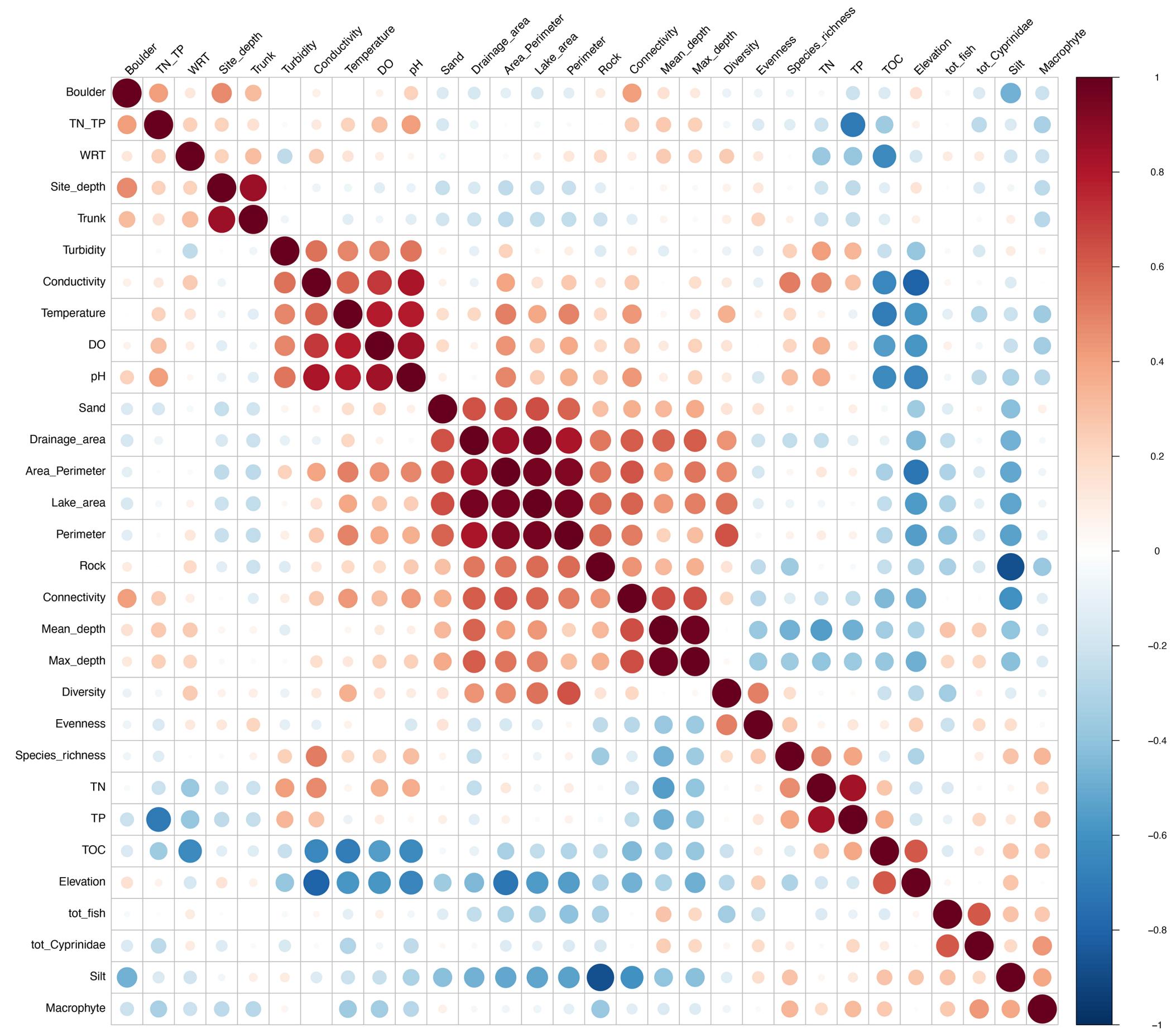


TABLE S2. Mean fish species length in the landscape across the different sampling methods.

Method	Mean (cm)	sd (cm)	N
Combined	6	3	4331
Minnow trap	6	3	1904
Seine net	5	2	2427

TABLE S3. Mean species length within each lake sampled across the different sampling method. Lake Tracy was omitted because of only one sampled fish.

Lake	Combined methods			Minnow trap			Seine net		
	Mean (cm)	sd (cm)	N	Mean (cm)	sd (cm)	N	Mean (cm)	sd (cm)	N
Achigan	4	2	73	18	NA	1	4	2	72
Beaver	4	1	1778	4	1	1004	4	1	774
Coeur	8	3	193	5	3	13	8	3	180
Cornu	11	8	27	7	10	13	14	4	14
Corriveau	6	2	672	7	2	283	5	1	389
Croche	10	4	281	12	3	191	6	1	90
Cromwell	7	3	182	8	4	44	6	2	138
Echo	6	2	515	7	5	51	6	2	464
Fournelle	7	3	77	6	2	12	7	4	65
Montaubois	10	5	71	13	5	36	8	2	35
Morency	5	2	183	5	2	104	5	3	79
Pin rouge	9	3	57	8	4	31	10	2	26
St-Onge	10	3	56	12	2	16	10	2	40
Triton	6	3	165	6	2	104	5	4	61

TABLE S4. References of the black spot disease occurrence in the fish species sampled our study system. *Ameiurus nebulosus* and *Esox masquinongy* have no mention of black spot disease. This table is not an exhaustive review.

	Black spot trematode species	References
<i>Lepomis gibbosus</i>	<i>Uvulifer ambloplitis</i>	Krull, 1934
	<i>Apophallus brevis</i>	Cone & Anderson, 1977
	Unknown	Steedman, 1991
<i>Ambloplites rupestris</i>	<i>Uvulifer ambloplitis</i>	Berra & Au, 1978
	Unknown	Steedman, 1991
<i>Ameiurus nebulosus</i>	NA	No mention
<i>Esox masquinongy</i>	NA	No mention
<i>Luxilus cornutus</i>	Unknown	Steedman, 1991; Hoffman, 1956
<i>Chrosomus eos x Chrosomus neogaeus</i>	Unknown	Paradis & Chapleau, 1994
<i>Pimephales notatus</i>	<i>Uvulifer ambloplitis</i>	Berra & Au, 1978
	Unknown	McAllister et al., 2013; Steedman, 1991; Hockett & Mundahl, 1989
<i>Pimephales promelas</i>	<i>Crassiphiala bulboglossa</i>	Vaughan & Coble, 1975; Wisenden et al., 2012
	<i>Uvulifer ambloplitis</i>	Berra & Au, 1978
	Unknown	Steedman, 1991
<i>Rhinichthys atratulus</i>	<i>Uvulifer ambloplitis</i>	Berra & Au, 1978
	Unknown	Steedman, 1991
<i>Semotilus atromaculatus</i>	<i>Uvulifer ambloplitis</i>	Berra & Au, 1978
	Unknown	McAllister et al., 2013; Steedman, 1991; Happel, 1991
	<i>Crassiphiala bulboglossa</i>	Krause et al., 1999; Hoffman, 1956
<i>Catostomus commersoni</i>	<i>Uvulifer ambloplitis</i>	Berra & Au, 1978
	Unknown	Steedman, 1991
<i>Umbra limi</i>	<i>Crassiphiala bulboglossa</i>	Hoffman, 1956
<i>Fundulus diaphanus</i>	<i>Crassiphiala bulboglossa</i>	Krause et al., 1999
<i>Micropterus dolomieu</i>	<i>Uvulifer ambloplites</i>	Berra & Au, 1978; Hunter & Hunter, 1938
<i>Perca flavescens</i>	<i>Crassiphiala bulboglossa</i>	Hoffman, 1956
	Unknown	Vaughan & Coble, 1975

TABLE S5. Observed landscape-scale fish community prevalence estimated by each method. All values are given in percentage.

Method	Prevalence (%)
Combined	30
Minnow trap	19
Seine net	20
Transect	36

TABLE S6. Observed lake-scale fish community prevalence estimated by each method. All values are given in percentage. Lake Tracy was not included because only one fish was caught through all methods. NAs mean that the lake was not sampled with the corresponding method.

Method	Prevalence (%)													
	Achigan	Beaver	Coeur	Cornu	Corriveau	Croche	Cromwell	Echo	Fournelle	Montaubois	Morency	Pin rouge	St-Onge	Triton
Combined	6	1	65	48	12	12	34	26	67	67	61	71	0	0
Minnow trap	0	1	23	54	37	11	97	60	58	74	94	87	0	0
Seine net	12	1	65	93	8	33	31	21	83	60	65	85	0	0
Transect	3	NA	66	47	5	10	23	28	65	NA	58	67	NA	0

TABLE S7. Observed site-scale fish prevalence estimated by each method. Affiliated lake are given as columns. All values are given in percentage. Samples with no captures were omitted to alleviate the table. Lake Tracy was not included because only one fish was caught through all methods. Lakes Beaver, Tracy, Montaubois and St-Onge were not sampled with the transect method.

TABLE S8. Method comparison of landscape final resampled prevalence estimates (N =35). Kruskal-Wallis chi-squared = 1288, p-value = 0. A Benjamini-Hochberg correction is applied on the dunn test.

Comparison	Adjusted p-value
Combined- Minnow trap	0.000
Combined- Seine net	0.000
Minnow trap- Seine net	0.000
Combined- Transect	0.000
Minnow trap- Transect	0.000
Seine net- Transect	0.000

TABLE S9. Landscape observed and resampled prevalence estimated by each sampling method. All values are given in percentage. N stands for the number of according sampling effort.

Method	Observed	N ₅	N ₁₀	N ₁₅	N ₂₀	N ₂₅	N ₃₀	N ₃₅
Combined	29.62	37.14	32.65	31.65	31.40	30.92	30.29	31.14
Minnow trap	19.46	39.49	32.94	30.16	27.60	26.12	24.94	24.39
Seine net	20.45	32.26	26.54	24.21	23.56	22.26	22.30	21.58
Transect	35.55	37.37	35.45	36.12	35.75	35.77	35.69	35.60

TABLE S10. Summary statistics of 999 resampling analyses. All values are given in percentage. N stands for the number of sampling effort implemented in the random resampling loop. Values are grouped according to sampling method

N	mean	sd	se	t score	margin error	lower bound	upper bound
Combined							
1	30.47	40.14	1.27	1.96	2.49	27.98	32.96
2	39.94	37.96	1.20	1.96	2.36	37.59	42.30
3	41.38	33.79	1.07	1.96	2.10	39.28	43.48
4	38.61	31.03	0.98	1.96	1.93	36.69	40.54
5	37.02	27.41	0.87	1.96	1.70	35.32	38.72
6	35.87	25.79	0.82	1.96	1.60	34.26	37.47
7	34.73	24.38	0.77	1.96	1.51	33.22	36.25
8	36.04	24.05	0.76	1.96	1.49	34.55	37.53
9	35.21	22.90	0.72	1.96	1.42	33.79	36.63
10	34.40	22.35	0.71	1.96	1.39	33.02	35.79
11	34.28	21.42	0.68	1.96	1.33	32.95	35.61
12	33.48	21.09	0.67	1.96	1.31	32.17	34.79
13	32.71	20.60	0.65	1.96	1.28	31.43	33.99
14	31.79	19.51	0.62	1.96	1.21	30.58	33.00
15	32.11	19.01	0.60	1.96	1.18	30.93	33.29
16	31.31	18.29	0.58	1.96	1.14	30.17	32.44
17	32.27	18.17	0.57	1.96	1.13	31.14	33.40
18	30.76	17.32	0.55	1.96	1.08	29.69	31.84
19	31.90	17.27	0.55	1.96	1.07	30.83	32.97
20	31.78	17.27	0.55	1.96	1.07	30.71	32.85
21	32.04	16.35	0.52	1.96	1.02	31.03	33.06
22	31.08	16.09	0.51	1.96	1.00	30.08	32.08
23	30.78	15.50	0.49	1.96	0.96	29.82	31.75
24	31.42	15.95	0.50	1.96	0.99	30.43	32.41
25	31.30	15.57	0.49	1.96	0.97	30.33	32.27
26	30.56	15.47	0.49	1.96	0.96	29.60	31.52
27	31.09	14.47	0.46	1.96	0.90	30.19	31.99
28	30.80	14.29	0.45	1.96	0.89	29.91	31.69
29	31.39	14.11	0.45	1.96	0.88	30.51	32.27
30	30.38	13.97	0.44	1.96	0.87	29.52	31.25
31	30.64	13.41	0.42	1.96	0.83	29.81	31.48
32	30.56	13.05	0.41	1.96	0.81	29.75	31.37
33	31.11	13.28	0.42	1.96	0.82	30.29	31.94
34	30.74	12.95	0.41	1.96	0.80	29.94	31.54
35	30.52	12.76	0.40	1.96	0.79	29.73	31.31
Minnow trap							
1	24.02	38.92	1.23	1.96	2.42	21.60	26.43
2	35.96	40.38	1.28	1.96	2.51	33.45	38.46
3	37.44	38.04	1.20	1.96	2.36	35.08	39.80
4	39.31	35.00	1.11	1.96	2.17	37.14	41.49
5	37.63	32.92	1.04	1.96	2.04	35.59	39.68
6	37.84	30.73	0.97	1.96	1.91	35.93	39.75
7	36.25	28.93	0.92	1.96	1.80	34.45	38.05
8	34.87	27.36	0.87	1.96	1.70	33.18	36.57
9	34.01	25.85	0.82	1.96	1.61	32.40	35.61
10	33.46	24.34	0.77	1.96	1.51	31.95	34.97
11	32.86	23.42	0.74	1.96	1.45	31.40	34.31
12	31.09	22.41	0.71	1.96	1.39	29.70	32.48
13	31.50	22.77	0.72	1.96	1.41	30.09	32.91
14	30.80	21.79	0.69	1.96	1.35	29.45	32.15
15	30.01	20.64	0.65	1.96	1.28	28.72	31.29
16	29.73	20.33	0.64	1.96	1.26	28.47	30.99
17	29.36	19.47	0.62	1.96	1.21	28.15	30.57
18	28.21	19.04	0.60	1.96	1.18	27.03	29.40
19	29.03	19.62	0.62	1.96	1.22	27.82	30.25
20	27.84	17.97	0.57	1.96	1.12	26.72	28.95
21	27.29	17.49	0.55	1.96	1.09	26.20	28.38
22	26.83	17.76	0.56	1.96	1.10	25.73	27.93
23	26.77	17.39	0.55	1.96	1.08	25.69	27.85
24	25.78	16.27	0.51	1.96	1.01	24.77	26.79
25	26.06	16.34	0.52	1.96	1.01	25.04	27.07
26	25.64	15.53	0.49	1.96	0.96	24.67	26.60
27	26.87	16.65	0.53	1.96	1.03	25.83	27.90
28	24.75	15.04	0.48	1.96	0.93	23.82	25.68
29	25.81	15.75	0.50	1.96	0.98	24.84	26.79
30	25.05	15.13	0.48	1.96	0.94	24.11	25.99
31	24.77	14.93	0.47	1.96	0.93	23.84	25.69
32	24.57	14.06	0.44	1.96	0.87	23.70	25.45
33	24.82	14.57	0.46	1.96	0.90	23.91	25.72
34	23.96	13.58	0.43	1.96	0.84	23.12	24.81
35	24.13	13.83	0.44	1.96	0.86	23.27	24.99
Seine net							
1	41.25	39.83	1.26	1.96	2.47	38.77	43.72
2	39.02	31.60	1.00	1.96	1.96	37.06	40.98
3	36.55	27.57	0.87	1.96	1.71	34.83	38.26
4	34.56	24.35	0.77	1.96	1.51	33.05	36.07
5	33.04	23.20	0.73	1.96	1.44	31.60	34.48
6	31.18	21.76	0.69	1.96	1.35	29.82	32.53
7	30.41	19.99	0.63	1.96	1.24	29.17	31.66
8	28.12	18.23	0.58	1.96	1.13	26.99	29.25
9	27.85	17.34	0.55	1.96	1.08	26.77	28.93
10	27.09	16.17	0.51	1.96	1.00	26.08	28.09
11	26.62	16.00	0.51	1.96	0.99	25.62	27.61
12	25.29	15.18	0.48	1.96	0.94	24.35	26.23
13	25.66	14.67	0.46	1.96	0.91	24.75	26.57
14	25.65	13.63	0.43	1.96	0.85	24.80	26.50
15	24.13	13.07	0.41	1.96	0.81	23.32	24.95
16	25.15	13.09	0.41	1.96	0.81	24.34	25.97
17	24.00	11.75	0.37	1.96	0.73	23.27	24.73
18	23.55	11.36	0.36	1.96	0.71	22.85	24.26
19	24.12	11.76	0.37	1.96	0.73	23.39	24.85
20	23.47	10.65	0.34	1.96	0.66	22.81	24.13
21	22.62	10.56	0.33	1.96	0.66	21.97	23.28
22	23.84	10.62	0.34	1			

TABLE S11. Estimated parametric coefficients and approximate significance of smooth terms of the fine-scale prevalence community GAMMs. The deviance explained (D^2) is given for every model as a measure of the model fit.

	Term	Smooth terms				
		Parametric coefficient		Statistic	p-value	
		Estimate	Standard error			
Evenness ($D^2 = 69.58\%$)	Intercept	-0.892	0.471	-1.895 ²	0.070	NA
	s(Evenness)	NA	NA	2.921 ³	0.095	0.707
	s(Lake)	NA	NA	3.658 ³	0.001	7.629
Diversity index ($D^2 = 78.80\%$)	Intercept	-0.971	0.597	-1.626 ²	0.115	NA
	s(Diversity)	NA	NA	50.579 ³	0.008	0.993
	s(Lake)	NA	NA	6.101 ³	0.000	8.720
Species richness ($D^2 = 73.70\%$)	Intercept	-0.900	0.553	-1.629 ²	0.114	NA
	s(Lake)	NA	NA	5.094 ³	0.000	8.375
	s(Species_richness)	NA	NA	7.517 ³	0.105	0.671
Non-host abundance ($D^2 = 87.49\%$)	Intercept	-0.825	0.509	-1.621 ²	0.117	NA
	s(Lake)	NA	NA	7.098 ³	0.000	8.796
	s(tot_Cyprinidae)	NA	NA	179.239 ³	0.000	1.750
Fish abundance ($D^2 = 80.46\%$)	Intercept	-0.761	0.546	-1.394 ²	0.174	NA
	s(Lake)	NA	NA	6.282 ³	0.000	8.449
	s(tot_fish)	NA	NA	55.368 ³	0.002	1.435
Elevation ($D^2 = 69.64\%$)	Intercept	-0.942	0.487	-1.933 ²	0.063	NA
	s(Elevation)	NA	NA	0.000 ³	0.396	0.000
	s(Lake)	NA	NA	4.594 ³	0.000	8.065
Drainage area ($D^2 = 69.72\%$)	Intercept	-0.970	0.488	-1.987 ²	0.056	NA
	s(Drainage_area)	NA	NA	0.077 ³	0.324	0.164
	s(Lake)	NA	NA	4.453 ³	0.000	7.935
Water residence time ($D^2 = 69.64\%$)	Intercept	-0.942	0.487	-1.933 ²	0.063	NA
	s(Lake)	NA	NA	4.594 ³	0.000	8.065
	s(WRT)	NA	NA	0.000 ³	0.830	0.000
Distance to nearest lake ($D^2 = 69.64\%$)	Intercept	-0.942	0.487	-1.933 ²	0.063	NA
	s(Connectivity)	NA	NA	0.000 ³	0.575	0.000
	s(Lake)	NA	NA	4.594 ³	0.000	8.065
Lake maximum depth ($D^2 = 69.64\%$)	Intercept	-0.942	0.487	-1.933 ²	0.063	NA
	s(Lake)	NA	NA	4.594 ³	0.000	8.065
	s(Max_depth)	NA	NA	0.000 ³	0.792	0.000
Lake mean depth ($D^2 = 69.64\%$)	Intercept	-0.942	0.487	-1.933 ²	0.063	NA
	s(Lake)	NA	NA	4.594 ³	0.000	8.065
	s(Mean_depth)	NA	NA	0.000 ³	0.768	0.000
Perimeter ($D^2 = 69.64\%$)	Intercept	-0.942	0.487	-1.933 ²	0.063	NA
	s(Lake)	NA	NA	4.594 ³	0.000	8.065
	s(Perimeter)	NA	NA	0.000 ³	0.590	0.000
Surface area ($D^2 = 69.16\%$)	Intercept	-1.151	0.424	-2.714 ²	0.011	NA
	s(Lake_area)	NA	NA	16.644 ³	0.010	2.828
	s(Lake)	NA	NA	1.694 ³	0.003	4.259
Area:Perimeter ($D^2 = 67.80\%$)	Intercept	-1.311	0.356	-3.682 ²	0.001	NA
	s(Area_Perimeter)	NA	NA	17.886 ³	0.000	2.966
	s(Lake)	NA	NA	0.000 ³	0.078	0.000
Conductivity ($D^2 = 56.19\%$)	Intercept	-0.665	0.177	-3.766 ²	0.001	NA
	s(Conductivity)	NA	NA	9.975 ³	0.000	2.617
	s(Lake)	NA	NA	0.000 ³	0.271	0.000
DO ($D^2 = 75.17\%$)	Intercept	-0.885	0.497	-1.780 ²	0.086	NA
	s(DO)	NA	NA	64.497 ³	0.016	1.046
	s(Lake)	NA	NA	4.548 ³	0.000	7.781
pH ($D^2 = 70.12\%$)	Intercept	-0.718	0.357	-2.014 ²	0.053	NA
	s(Lake)	NA	NA	1.761 ³	0.009	6.079
	s(pH)	NA	NA	35.861 ³	0.019	1.399
Turbidity ($D^2 = 76.24\%$)	Intercept	-0.923	0.491	-1.880 ²	0.070	NA
	s(Lake)	NA	NA	5.025 ³	0.000	8.236
	s(Turbidity)	NA	NA	4.051 ³	0.052	1.214
Temperature ($D^2 = 79.55\%$)	Intercept	-0.720	0.469	-1.535 ²	0.136	NA
	s(Lake)	NA	NA	4.144 ³	0.000	7.755
	s(Temperature)	NA	NA	69.780 ³	0.001	1.524
Trunk ($D^2 = 69.64\%$)	Intercept	-0.942	0.487	-1.933 ²	0.063	NA
	s(Lake)	NA	NA	4.594 ³	0.000	8.065
	s(Trunk)	NA	NA	0.000 ³	0.358	0.000
Transect depth ($D^2 = 69.64\%$)	Intercept	-0.942	0.487	-1.933 ²	0.063	NA
	s(Lake)	NA	NA	4.594 ³	0.000	8.065
	s(Site_depth)	NA	NA	0.000 ³	0.709	0.000
Macrophyte cover ($D^2 = 84.15\%$)	Intercept	-0.870	0.457	-1.906 ²	0.067	NA
	s(Lake)	NA	NA	6.203 ³	0.000	8.449
	s(Macrophyte)	NA	NA	44.108 ³	0.000	1.564
Boulder ($D^2 = 69.64\%$)	Intercept	-0.942	0.487	-1.933 ²	0.063	NA
	s(Boulder)	NA	NA	0.000 ³	0.811	0.000
	s(Lake)	NA	NA	4.594 ³	0.000	8.065
Rock ($D^2 = 69.64\%$)	Intercept	-0.942	0.487	-1.933 ²	0.063	NA
	s(Lake)	NA	NA	4.594 ³	0.000	8.065
	s(Rock)	NA	NA	0.000 ³	0.813	0.000
Sand ($D^2 = 74.31\%$)	Intercept	-1.024	0.494	-2.070 ²	0.047	NA
	s(Lake)	NA	NA	5.624 ³	0.000	8.061
	s(Sand)	NA	NA	4.159 ³	0.046	0.919
Silt ($D^2 = 69.64\%$)	Intercept	-0.942	0.487	-1.933 ²	0.063	NA
	s(Lake)	NA	NA	4.594 ³	0.000	8.065
	s(Silt)	NA	NA	0.000 ³	0.675	0.000
TOC ($D^2 = 69.64\%$)	Intercept	-0.942	0.487	-1.933 ²	0.063	NA
	s(Lake)	NA	NA	4.593 ³	0.000	8.065
	s(TOC)	NA	NA	0.000 ³	0.344	0.000
TP ($D^2 = 73.31\%$)	Intercept	-0.916	0.488	-1.875 ²	0.071	NA
	s(Lake)	NA	NA	4.554 ³	0.000	8.131
	s(TP)	NA	NA	9.240 ³	0.053	0.838
TN ($D^2 = 70.50\%$)	Intercept	-0.941	0.494	-1.907 ²	0.066	NA
	s(Lake)	NA	NA	4.654 ³	0.000	8.064
	s(TN)	NA	NA	0.394 ³	0.244	0.241
TN:TP ($D^2 = 80.96\%$)	Intercept	-0.977	0.487	-2.007 ²	0.055	NA
	s(Lake)	NA	NA	6.226 ³	0.000	8.447
	s(TN_TP)	NA	NA	17.630 ³	0.185	2.408
Null ($D^2 = 69.64\%$)	Intercept	-0.942	0.487	-1.933 ²	0.063	NA
	s(Lake)	NA	NA	4.594 ³	0.000	8.065

¹ Effective degrees of freedom

<p

TABLE S12. Transect sites water and habitat characteristics. The results are grouped by lake.

	Sampling_ID	Temperature (°C)	Conductivity (µS/cm)	Dissolved oxygen (%)	Turbidity (NTU)	pH	TOC (mg/L)	TN (mg/L)	TP (mg/L)	Trunk	Silt (%)	Sand (%)	Rock (%)	Boulder (%)	Macrophyte (%)	Mean depth (cm)
Achigan	ACHI_T_01	25.2	56.6	8.37	0.06	7.25	4.75	232.49	4.3049	15	0.0	5.0	74.8	20.2	3.8	118.5
	ACHI_T_02	24.3	55.2	8.55	0.38	7.38	4.67	259.85	6.2188	13	4.0	1.5	83.5	11.0	1.4	79.0
	ACHI_T_03	24.5	55.7	8.66	0.17	7.56	4.75	299.75	5.8841	26	28.7	0.0	20.8	49.5	6.9	115.0
	ACHI_T_04	24.5	55.6	8.67	0.18	7.38	4.67	238.19	5.4528	9	0.0	0.0	67.0	33.0	0.2	147.0
	ACHI_T_05	23.6	56.1	8.78	0.24	6.83	4.78	242.75	6.6700	1	9.0	43.8	45.6	1.6	36.7	37.5
	ACHI_T_06	24.3	55.9	8.60	0.16	6.98	4.83	240.47	4.7597	12	0.0	11.5	72.8	15.7	2.2	66.0
Beaver	BEAV_T_01	23.0	26.0	8.87	1.78	6.60	9.15	487.85	18.4269	NA	NA	NA	NA	NA	NA	NA
	BEAV_T_02	22.7	26.0	8.82	1.88	6.77	9.17	499.25	19.4058	NA	NA	NA	NA	NA	NA	NA
Coeur	COEU_T_03	23.0	53.4	8.26	0.20	7.13	5.60	331.67	7.3317	9	33.9	0.0	20.5	45.6	9.0	80.0
	COEU_T_04	22.8	53.2	8.26	0.28	6.78	5.64	341.93	6.9554	79	57.0	0.0	15.9	27.1	10.6	108.0
	COEU_T_01	23.6	53.8	8.37	0.29	7.38	5.75	419.45	9.5510	80	43.0	0.0	27.5	29.5	2.3	187.5
	COEU_T_02	23.3	53.5	8.38	0.33	7.25	5.63	370.43	8.4576	16	36.5	0.0	29.8	33.7	22.0	62.5
Cornu	CORN_T_01	22.6	147.8	8.69	0.74	7.33	3.97	324.83	6.9125	380	62.0	0.0	1.0	37.0	0.0	370.0
	CORN_T_02	22.5	149.6	8.00	0.41	6.87	4.05	292.91	7.6265	45	100.0	0.0	0.0	0.0	25.0	39.5
	CORN_T_03	22.6	152.3	8.10	1.34	6.77	4.05	374.99	13.9793	11	97.4	0.0	0.0	2.6	75.5	44.5
	CORN_T_04	22.6	147.8	9.10	0.13	7.61	4.13	357.89	8.7511	18	18.0	0.0	65.5	16.5	14.9	45.0
Corriveau	CORR_T_01	20.3	19.1	8.06	0.49	5.76	8.42	400.07	12.4658	108	77.5	0.0	14.0	8.5	35.0	107.0
	CORR_T_02	21.0	19.1	8.27	0.39	6.10	8.25	394.37	14.9332	33	57.5	6.0	12.0	24.5	27.5	89.5
Croche	CROC_T_01	21.3	14.2	7.84	0.38	6.09	6.36	332.81	7.0791	52	100.0	0.0	0.0	0.0	39.5	73.5
	CROC_T_04	21.9	14.3	8.20	0.19	5.89	5.39	260.99	8.0571	171	97.5	0.0	1.0	1.5	10.1	97.0
	CROC_T_06	21.6	14.2	7.71	0.24	6.05	6.30	298.61	8.8069	73	50.5	0.0	46.1	2.9	3.3	94.0
Cromwell	CROM_T_02	19.7	19.2	6.66	0.91	5.72	7.87	381.83	11.5249	79	80.0	0.0	6.0	27.0	56.0	76.0
	CROM_T_03	19.6	19.6	4.56	0.56	5.76	7.57	341.93	11.8224	27	71.0	0.0	14.0	15.0	61.5	84.5
	CROM_T_04	19.8	19.6	5.45	0.49	5.89	7.81	361.31	10.7160	53	89.8	0.0	5.8	4.4	69.0	117.0
Echo	ECHO_T_05	24.6	166.3	10.18	1.56	8.25	5.67	520.91	13.1379	10	84.0	0.0	7.0	9.0	58.5	52.0
	ECHO_T_01	23.5	174.3	9.84	1.80	8.00	5.58	492.41	13.4672	45	87.0	0.0	5.8	7.2	53.5	50.5
	ECHO_T_02	25.0	171.0	10.07	3.50	8.17	5.75	511.79	15.8648	27	59.5	19.0	10.5	10.5	49.0	39.5
	ECHO_T_03	23.4	159.9	9.77	1.54	8.15	5.65	519.77	15.7477	21	0.0	13.0	85.0	2.0	1.1	41.3
	ECHO_T_04	23.3	159.6	9.71	2.12	7.96	5.72	593.87	19.3587	17	75.0	0.0	19.5	5.5	11.5	51.0
Fournelle	FOUR_T_01	24.9	62.5	8.44	0.76	6.68	5.41	311.15	9.0404	41	96.9	0.0	3.0	0.1	31.5	82.0
	FOUR_T_02	24.5	60.7	8.94	1.58	7.38	5.42	328.25	10.2989	26	95.4	0.0	1.2	3.2	9.6	125.5
	FOUR_T_03	24.6	60.7	8.78	1.33	7.61	5.37	308.87	7.4852	57	100.0	0.0	0.0	0.0	8.5	61.5
Montaubois	MONT_T_01	22.6	60.9	8.26	0.03	6.44	3.88	303.17	6.1110	NA	NA	NA	NA	NA	NA	NA
	MONT_T_02	23.2	61.5	8.01	0.08	6.52	3.67	239.33	5.5773	NA	NA	NA	NA	NA	NA	NA
	MONT_T_03	22.8	61.7	8.17	0.02	6.64	3.70	265.55	5.0967	NA	NA	NA	NA	NA	NA	NA
Morency	MORE_T_03	23.2	128.6	9.29	0.42	8.17	4.25	355.61	8.4191	83	24.5	0.0	35.5	40.5	13.5	129.0
	MORE_T_04	22.3	126.1	9.52	0.43	8.28	4.30	355.61	6.6601	31	53.0	3.0	16.5	27.5	57.3	58.0
	MORE_T_01	24.8	132.7	10.38	2.38	8.50	4.49	356.75	7.5123	15	88.5	0.0	3.6	7.8	8.5	30.5
	MORE_T_02	24.0	129.3	9.53	4.32	8.16	4.30	341.93	6.8925	94	42.8	0.7	16.5	41.0	5.9	168.0
Pin rouge	PINR_T_01	21.0	45.5	7.70	0.39	6.74	8.27	502.67	13.6476	5	100.0	0.0	0.0	0.0	61.0	32.5
	PINR_T_02	20.9	45.9	8.13	0.35	6.42	8.29	460.49	13.4307	37	69.5	9.0	4.5	17.0	23.3	56.5
	PINR_T_03	20.7	45.7	8.08	0.12	7.01	8.15	408.05	8.1835	20	98.8	0.7	0.5	0.0	9.2	26.5
St-Onge	STON_T_02	22.3	94.4	7.64	1.45	6.55	13.30	656.57	25.7782	NA	NA	NA	NA	NA	NA	NA
	STON_T_01	22.8	99.1	7.10	1.36	6.18	13.10	530.03	17.6344	NA	NA	NA	NA	NA	NA	NA
Tracy	TRAC_T_02	24.1	72.9	8.24	0.03	6.35	3.87	467.33	8.6231	NA	NA	NA	NA	NA	NA	NA
	TRAC_T_01	23.0	71.1	8.31	0.08	6.45	3.31	272.39	4.8220	NA	NA	NA	NA	NA	NA	NA
Triton	TRIT_T_01	20.9	14.3	7.52	1.19	5.99	7.09	386.39	9.5666	27	98.8	0.0	0.0	1.2	57.5	128.5
	TRIT_T_02	20.9	14.2	7.55	0.82	5.69	7.93	379.55	10.1204	166	91.5	0.0	4.5	4.0	41.0	121.0

TABLE S13. Host specificity of the black spot disease at landscape-scale according to the different sampling methods. NaN means that no fish were caught in the corresponding category. All values are in percentage.

Species	Method			
	Combined	Minnow trap	Seine net	Transect
<i>Ambloplites rupestris</i>	52	22	71	54
<i>Fundulus diaphanus</i>	0	NaN	0	NaN
<i>Micropterus dolomieu</i>	28	0	9	31
Unknown centrarchids	0	NaN	0	NaN
<i>Lepomis gibbosus</i>	58	71	70	55
<i>Perca flavescens</i>	37	53	24	59
<i>Pimephales promelas</i>	2	2	2	NaN
<i>Chrosomus spp.</i>	0	0	0	NaN
<i>Pimephales notatus</i>	2	NaN	2	NaN
Unknown cyprinids	0	NaN	0	0
<i>Semotilus atromaculatus</i>	4	3	18	NaN
<i>Luxilus cornutus</i>	0	0	0	NaN
<i>Catostomus commersonii</i>	0	NaN	0	NaN
<i>Umbra limi</i>	0	NaN	0	NaN
<i>Rhinichthys atratulus</i>	0	NaN	0	NaN