

Determinants of community structure of zooplankton in heavily polluted river ecosystems

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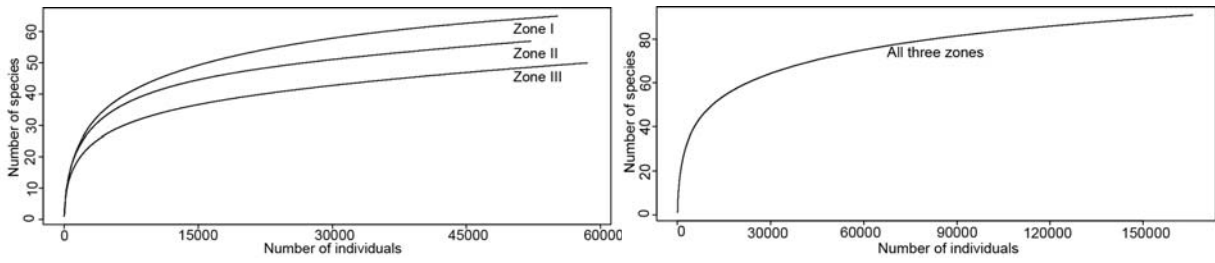
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Supplementary Figure S1 Rarefaction curves of counted individuals of rotifers. Left - the each zone, right - the combined set 94 samples.

Supplementary Table S1 Summary of principal component analyses on 13 biotic and abiotic factors in all 94 sampling sites. Factors in red were examined to be representing primary gradients of corresponding component. Distance = average geographical distance of a given sampling site from the other sampling sites, T = water temperature, SD = Secchi disk depth, TP = total phosphorus, SRP = soluble reactive phosphorus, TN = total nitrogen, NO₃-N = nitrate nitrogen, NH₄-N = ammonia nitrogen, Protozoa = the total abundance of all protozoa species, Crustacea = the total abundance of all crustacean species.

	Component				
	1	2	3	4	5
Longitude	-0.056	0.933	-0.109	-0.068	0.061
Latitude	0.107	0.862	0.296	0.21	0.047
Altitude	-0.233	-0.678	0.516	0.059	-0.097
Distance	-0.388	-0.575	-0.016	0.465	0.222
T	-0.02	-0.16	-0.832	-0.183	-0.11
SD	-0.142	-0.15	0.681	-0.166	-0.101
TP	0.918	0.135	-0.05	0.14	-0.066
SRP	0.912	0.126	-0.059	0.153	-0.089
TN	0.904	0.072	0.009	0.159	0.105
NO ₃ -N	0.805	-0.126	-0.11	-0.227	0.106
NH ₄ -N	0.249	0.02	0.025	0.839	-0.055
Protozoa	-0.007	0.072	-0.031	-0.01	0.921
Crustacea	-0.196	0.405	-0.301	0.298	-0.333

Supplementary Table S3 The average similarity of rotifer communities among sampling sites

within each zone based on the similarity percentage (SIMPER) analysis in the plain region of the

Haihe River Basin.

Zone I: Average similarity: 13.44					
Species	Average abundance	Average similarity	Similarity	Contribution (%)	Cumulation (%)
<i>Polyarthra trigla</i>	463.39	2.56	0.45	19.02	19.02
<i>Brachionus angularis</i>	92.68	1.79	0.42	13.31	32.34
<i>Brachionus urceus</i>	330	1.76	0.48	13.11	45.44
<i>Filinia maio</i>	157.5	1.75	0.37	13.04	58.48
<i>Pompholyx sulcata</i>	26.25	0.95	0.32	7.05	65.53
<i>Brachionus calyciflorus</i>	20.36	0.78	0.32	5.77	71.3
<i>Trichocerca</i> sp.	202.5	0.65	0.32	4.81	76.11
<i>Brachionus calyciflorus</i>	83.57	0.53	0.33	3.97	80.08
<i>Philodina</i> sp.	12.86	0.52	0.24	3.83	83.91
<i>Brachionus budapesllensis</i>	26.25	0.38	0.24	2.85	86.76
<i>Euchlanis</i> sp.	9.11	0.3	0.18	2.21	88.98
<i>Philodina megalotrocha</i>	35.36	0.3	0.12	2.21	91.18
Zone II: Average similarity: 19.22					
<i>Brachionus angularis</i>	192.32	4.58	0.78	23.81	23.81
<i>Polyarthra trigla</i>	445.18	3.63	0.58	18.87	42.67
<i>Filinia maio</i>	71.79	2.62	0.49	13.65	56.33
<i>Brachionus calyciflorus</i>	260.89	2.49	0.55	12.93	69.26
<i>Brachionus urceus</i>	73.93	1.93	0.4	10.02	79.28
<i>Keratella valga</i>	26.25	0.71	0.37	3.68	82.96
<i>Filinia longisela</i>	221.79	0.66	0.2	3.45	86.42
<i>Trichocerca pusilla</i>	56.79	0.64	0.28	3.31	89.73
<i>Trichocerca</i> sp.	71.25	0.56	0.26	2.9	92.63
Zone III: Average similarity: 24.49					
<i>Brachionus urceus</i>	131.05	5.83	0.81	23.8	23.8
<i>Brachionus angularis</i>	195.79	4.93	0.79	20.13	43.93
<i>Polyarthra trigla</i>	428.29	3.92	0.65	16.02	59.95
<i>Trichocerca</i> sp.	137.76	2.86	0.52	11.67	71.62
<i>Brachionus calyciflorus</i>	123.55	2.32	0.54	9.47	81.09
<i>Filinia maio</i>	112.5	1.29	0.35	5.28	86.37
<i>Brachionus budapesllensis</i>	24.87	0.65	0.29	2.67	89.03
<i>Keratella valga</i>	41.05	0.58	0.3	2.38	91.41