

Abundance of microorganisms and enzyme activity in the rapids-pool-benchland systems in natural Douliu River of China

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Abstract River structures and their ecosystem functions have been greatly destructed due to the human disturbance during the last decades. It is a pressing task to understand the complete structures and ecosystem functions of a natural river to restore the impaired river ecosystems. We found a basic structure of a natural river, i.e., the rapid-pool-benchland system, in the previous studies, so we selected the rapid-pool-benchland systems in the upper, middle and lower reaches of Duliu River, a natural river in China and tested the abundance of microorganisms and the enzyme activity in the sediments in these systems. Results indicated that the number of bacteria was far more than ammonifiers, actinomycetes, fungus and denitrifying bacteria in the sediments of all river reaches. In the upper reach, the microbial abundance in the sediments was more than the middle and lower reaches. In each of the rapid-pool-benchland system, the microbial abundance in the sediments of the pool was always highest, and that of the rapid was lowest. The catalase activity in the sediments of the benchland was higher than the rapid and pool but the phosphatase activity in the pool was higher than the rapid and benchland. The phosphatase and urease activities were significantly correlated to the abundance of the bacteria, ammonifiers, actinomycetes, fungus and denitrifying bacteria in the sediments. The rapid-pool-benchland system in a river obviously regulates the microbial abundance and the catalase and phosphatase activities in the sediments. However, the abundance of microorganisms only indicated a 0.434-0.836 level of significant correlations with the activity of four enzymes, and further influences biochemical functions.

Key word: Natural rivers, rapids, pools, benchlands, microorganisms, enzyme activity, sediments, Duliu River

INTRODUCTION

A river is a corridor connecting the terrestrial environment to the ocean realm and it plays an important role in the supply of water resources for sustenance of life systems, shipping and purification of polluted water (Wu, Chen, Zhang et al., 2008; Padmalal and Maya, 2014). Almost all the immemorial population assembly and the following ancient civilization origin on the coasts of rivers, and even to this day, many modern cities are also built along rivers (Wang, Wu, Zhang, et al., 2013). Arguably, a river is a footstone of the human civilization. However, human interventions, especially in developing countries, have imposed tremendous pressure on rivers, and consequently most of the rivers, especially the small rivers, have been altered to levels, often beyond their natural resilience capability (Dong, Sun, Peng, 2009; Padmalal and Maya, 2014). These interventions include many aspects. For example, river way is occupied due to the urbanization; river way is changed into a straight form leading to the disappearances of the meandering form; the natural riverbanks are replaced by concrete riverbanks for slope stability (Dong, Sun, Peng, 2009; Padmalal and Maya, 2014). As a result, the habitat heterogeneity of a river significantly declines, which results in the changes of biodiversity, self-purification capacity, landscape values and affinity

between humans and waterbody (Wang, Wu, Zhang, et al., 2013).

The habitat heterogeneity in rivers deals with many aspects, including a diversity of river forms, river structures, river materials, or river slopes. In river forms, scientists suggest that rivers generally present straight forms, bifurcation forms, middle-bar forms, wandering forms, braiding forms and so on (Niezgoda and Johnson, 2005; Yin, Li, Liu et al., 2012). These forms are products of the long-term interactions between water flow and a variety of environmental factors. The river structures can be reflected with a spatial variation of riverbed, waterflow, organisms, river banks along a transverse and a longitudinal direction of a river (Gregory, Benito, Downs, 2005). The river-bed materials consist of water, sediments, different sizes and forms of stones, organic matters, ions in water and organisms, and each of these materials in types and traits is also diverse (Asai, Takasaki, Muraoka et al., 2013). The river slope is variable along a longitudinal direction of a river, which is mainly related to landform and geological conditions. Clearly, a river is a complex system, but the system is becoming drab with a decline of the habitat heterogeneity.

In previous studies, we found a new river structure, the rapid-pool-benchland system in natural rivers by a great amount of field investigation (Fig 1 and Supplemental file 1) (Ma, Wang, Chen, 2014; Wu, Wang, Cheng, 2014; Chen, Ma, Wang, 2015; Wang, Wang, Liu, et al., 2015). The system continually occurs from the upper to lower reaches along a natural river, i.e., a rapid-pool-benchland system in the upper reach in a river directly links another rapid-pool-benchland system in the lower reach. In the system, there are also waterflow, organism, riverbed (sediment and pebbles) and river bank (dry land during most of time) that make up a complete river from a perspective of the different material compositions. However, the rapid-pool-benchland system explains a complete river in structure from a perspective of the habitat difference. In the rapid section in a rapid-pool-benchland system, there is a greater



Figure 1 A rapid-pool-benchland system in the upper reach of Yangtze River in China. The square symbol: pool; the triangular symbol: rapid; the circle symbol: benchland.

slope than pool and benchland, which generates a relatively great potential energy and consequently results in a rapid water flow. Moreover, there are bigger pebbles, stones or ratchel and few plants and sediments due to scouring effects. In addition, the rapid often

occurs in the middle section of a river (Fig 2). The pool in a rapid-pool-benchland system is generated by the long-term scouting effects of the water flow from the rapid section. In the pool, there are many fine-particle sediments, even sludge, disposed from the slow water flow, relatively rich plants and deeper water body than the rapid section. The pool generally occurs close to the river bank in a river (Fig 2). The benchland is often formed by transportation and sedimentation of floods and mainly consists of gravels and sediments. Some benchlands are directly river banks or links to the river bank. The most significant characteristic of the benchland is that there is alternation of wetting and drying because of a change of water levels during a year. As a result, many hydrophytes and xerophytes, and amphibious animals and insects are distributed in the benchland. In a natural river, there are significant relationships in areas and perimeters among the rapids, pools and benchlands, and the proportions of the areas or perimeters of the rapids to the pools or to the benchlands almost equal to a constant. Further studies indicate that the rapid-pool-benchland system is shaped by the long-term dynamical effects of the water flow and is a basic unit of the structure of a natural and healthy river. In such a river, there is a dynamic equilibrium among the rapids, pools and benchlands.

However, it remains unclear if abundance of microorganisms and the enzyme activity in the sediments of rapid, pool and benchland in a river will indicate different. The abundance of microorganisms and the enzyme activity in the sediments are important ecological functions in the rivers and referents of the river health. Microorganisms (bacteria, fungi, etc) process most of the organic waste from a variety of sources and they oxidize, reduce and precipitate mineral ions, thereby controlling the geochemical cycles in the rivers (Winiarski, Bedell, Delolme et al., 2006). The enzyme activity is an interesting parameter to measure the biological quality of soils or sediments because a number of studies have shown it to be one of the most sensitive parameters for evaluating the toxicity of certain pollutants in the soils or sediments (Garcia Hernandez, Costa, 1997; Tam, 1998; Ellis, Neish, Trett et al., 2001; Irha, Slet, Petersell, 2003; Benitez, Melgar, Nogales et al., 2004). Some recent studies have reported that enzymes in wetland systems play very important roles in matter transformation (Reboreda and Cacador, 2008). For example, there is a positive correlation between dehydrogenase activity and CO² emission, carbon cycle and biomass, and dehydrogenase activity is known to be sensitive to pollutants such as heavy metals (Garcia, Hernandez, Costa, 1994; Neto Ohannessian, Delolme et al., 2007). Ecologists have also found that there are significant positive correlations between the removal of total nitrogen and the number of soil microbes, and activities of soil enzymes in wetlands (Huang, Gao, Liu et al., 2012). In a lake, the differences in waterbody pollution can be well expressed by the number of microbes and activities of enzymes in sediments (Deng, Huang, Hu et al., 2009). The enzyme activity in river sediments is closely related to the anthropogenic nutrients and carbon loading input from the upper reaches of the river (Hill and Moffett, 2010; Millar, Payne, Ochs et al., 2015).

In the study, we selected three successively connecting rapid-pool-benchland systems respectively in the upper, middle and lower reaches of Duliu River, a typical natural river in Yungui plateau of China and collected the sediment samples in the rapids, pools and benchlands. Then, these sediment samples were delivered to laboratory to test the abundance of the microorganisms and the activity of several important enzymes. We assumed that the rapid-pool-benchland systems continually occurred to create habitat heterogeneity in the river, and consequently, which resulted in differences in the abundance of the microorganisms in the sediments of the rapid, pool and benchland systems. Correspondingly, the enzyme activity in the sediments of the rapid, pool and benchland systems also changed and further affected the biochemical functions. Therefore, we hope to answer the following questions: 1) if the rapid-pool-benchland system as a type of environmental heterogeneity

in a river directly influences the abundance of the microorganisms in the sediments, and further the enzyme activity? 2) If it does, to what degree can the relationships between the abundance of the microorganisms and the enzyme activity be explained by the rapid-pool-benchland system?

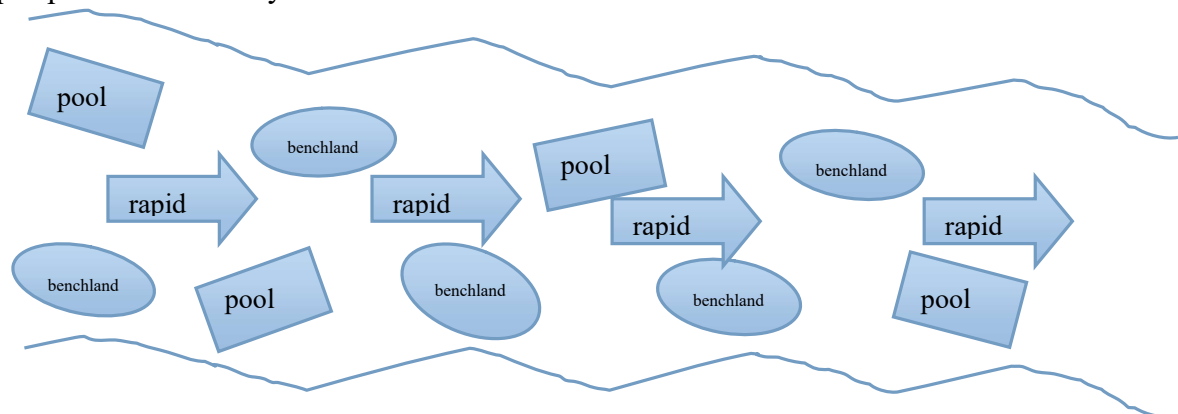


Fig.2 The typical model of the rapid-pool-benchland system

MATERIALS AND METHODS

Site description

Duliu River is a primary tributary of Pearl River, the third greatest river in China. Duliu River originates from Dushan county, flows through four counties in Guizhou Province, and finally joins Xi River, the upper reach of Pearl River. Total length of Duliu River is 310 km and the natural drop of the trunk stream is 84.5 m. The catchment area is 11,326 km². Annual mean rainfall ranges from 1350-1500 mm. The average annual flow is 145145 m³/s. The forest coverage is 74.1% in Duliu River watershed. There are no dams and artificial riverbanks along the river, so Duliu River is a natural river. We selected the upper Puan (26°04' N, 107°48' E), middle Sandu (25°58' N, 107°53' E) and lower Rongjiang (25°89' N, 108°50' E) river reaches, respectively as research objects. In each river reach, three rapid-pool-benchland systems successively connect with each other. From the three rapid-pool-benchland systems in each river reach to the upper or lower reaches, the rapid-pool-benchland systems connecting with each other are still seen. The parameters of the three river reaches are shown in Table 1.

Table 1 The parameters of the river reaches

River reaches	Longitudinal ratio	Bending coefficient	Rapid area (m ²)	Pool area (m ²)	Benchland area (m ²)
UR*	0.05±0.04	1.16±0.00	791.16±262.61	872.12±294	796.58±473.63
MR	0.02±0.01	1.05±0.00	1991.16±2262.41	1322.15±2041.96	1492.06±1686.82
LR	0.01±0.00	1.20±0.00	3128.52±714.4	3691.54±2994.8	12261.48±12307.98

*URR: the upper reach; MR: the middle reach; LRR: the lower reach.

Field sampling

In each set of the rapid-pool-benchland systems in the upper, middle or lower reach of Duliu River, we sampled the sediments in the rapid, pool and benchlands, respectively. Specifically, ten sampling sites were firstly even set up in the rapid, pool or benchland. Then, we used a small shovel with a long handle to collect about 300 g of the sediments at each site. The sediments from the ten sites were fully mixed and kept at 4 °C. The sediments in the rapid and pool were collected from the bottom of water. The total weight of the mixed samples of the sediments respectively collected from a rapid, pool or benchland was about

3kg. In the upper, middle and lower reaches, the number of the mixed samples was 27.

A test of abundance of microorganisms

Bacteria, fungus and actinomycetes: we adopted plate counting method to test the number of the bacteria, fungus and actinomycetes. We firstly made up the beef-protein mediums, Martin mediums and the No.1 medium of Gaos. After sterilization, these mediums were respectively sub-packed on three culture dishes that had been sterilized. Then, we weighed 10g of the fresh sediments and put them in a triangular flask with 100 ml of aqua sterilisata. Subsequently, the dilution of the soil solutions in the triangular flask was conducted in a sterile working chamber. The dilution proportions were respectively 10^{-6} , 10^{-7} and 10^{-8} for bacteria, and 10^{-4} , 10^{-5} and 10^{-6} for fungus and actinomycetes. We further used pipettes to transfer the diluted soil solutions to the culture dishes. The culture dishes were incubated at 28°C. Each of the dilution proportions was replicated three times.

Amonifying bacteria and denitrifying bacteria: we adopted a MPN (most-probable-number) method to test the number of the two types of bacteria. Firstly, the peptone agar mediums and the denitrifying bacteria mediums were respectively made up and were sub-packed into cuvettes. Each cuvette was filled with 5 ml, plugged using silica gel stopple and sterilized. In each of these filled cuvettes, a reversed Durham's fermentation tube was put to test the generation of gases. Then, we weighed 10g of the fresh sediments and put it in a triangular flask with 100 ml of aqua sterilisata. The triangular flask was oscillated for 30min to make suspension. Subsequently, the suspension was diluted into four levels of concentrations, i.e. the 10^{-4} , 10^{-5} , 10^{-6} and 10^{-7} times of the mother solutions. We used sterile pipettes to transfer these diluted solutions to the cuvettes filled with the mediums, and these cuvettes were incubated at 20-30°C. Each level of the concentrations was replicated three times. After three and five days, we checked the turbidity of the mediums for the amonifying bacteria. When the incubation continued till the seventh day, we used Nessler reagents to test if ammonia had been generated. For the denitrifying bacteria, we observed if bubbles had occurred in the cuvettes and if the solutions had become turbid after 14 days, and used Nessler reagents to test if ammonia had been generated. Based on the growth of the bacteria, we contrasted a MPN table to determine the number of the living bacteria per ml of the mother suspension.

Tests of the enzyme activity

Catalase: we adopted permanganate titration to test the catalase activity. Specifically, we weighed 5g of the fresh sediments and put them in a triangular flask. 25ml of H_2O_2 solutions and a little distilled water were decanted into the triangular flask, and then the triangular flask was continually oscillated for 30 minutes. The chemical reactions occurring in the triangular flask were terminated with addition of H_2SO_4 . The mixtures in the triangular flask were filtrated and the filtrates were titrated with $KMnO_4$. When the filtrates changed into red, the titration was stopped. The consumption volume (ml/g) of the $KMnO_4$ solution represented the catalase activity.

Alkaline phosphatase: the activity of alkaline phosphatase was tested with the phosphatase two sodium colorimetric method. We firstly weighed 5 g of the fresh sediments and put them in a triangular flask. Then, the toluene, disodium phenyl phosphate and borate buffer solution were decanted into the triangular flask. The triangular flask was continually incubated 24 h and then the mixtures in the triangular flask were filtrated. Potassium ferricyanide and 4-amino safety were used as chromogenic agents and were added to the filtrates. Finally, the filtrates were transferred to a volumetric flask, diluted to 750ml of constant volume for absorbance measurement.

Urease: the activity of the urease was tested by the phenol sodium hypochlorite

colorimetric method. We firstly weighed 5g of the fresh sediments and put them in a 50 ml volumetric flask. Then, we added 1ml of toluene to the volumetric flask, tightly plugged the volumetric flask and slightly waved it about 15 minutes. Succesively, 5ml of 10% solutions of urease and 10ml of citrate buffer solution (pH=7.6) were added to the volumetric flask and the soil solution in the volumetric flask must be fully mixed by waving. After the volumetric flask had been incubated at 37°C for 24 h, we used the distilled water at 38 °C to dilute the solution in the volumetric flask to constant volume. This moment, the toluene should be floated above the solution in the volumetric flask. After the solution in the volumetric flask was further waved to be evenly mixed, soliquoid was filtrated. Subsequently, 1ml of the filtrates was taken to put in a 50ml volumetric flask and 4 ml of sodium phenate solutions and 3 ml of chloros were added to the 50 ml volumetric flask. After coloration of the solutions in the volumetric flask, the solutions were distilled to constant volume. The absorbance of the solutions was tested after 1 h.

Dehydrogenase: we adopted (chloride three phenyl tetrazole) colorimetric method to test the activity of the dehydrogenase. We firstly weighed 5g of the sediments and put them in a triangular flask. Then, we added 2 ml of 1% triphenyltetrazolium chloride (TTC) solutions, glucose solutions, Tris-HCl buffer solutions, and Na₂S solutions to the triangular flask. The triangular flask was oscillated for 2h, and then the oxygen-free water and the sodium sulfite solutions were futher added to the triangulare flask. After the solutions in the triangular flask had been fully mixed, we took the solutions to test the absorbance of the solutions. The enzyme activity was indicated with the quantity of three phenyl formazen per unit soil and time.

RESULTS

Abundance of microorganics

The abundance in individual bacteria in the sediments collected from the rapids, pools or benchlands in these river reaches was highest among all types of microorganisms (Table 2). The abundance in individual amonifying bacteria, actinomycetes, fngus and denitrifying bacteria was respectively ranked second to fouth. One of the important causes why bacteria were more than actinomycetes and fngus was because the bacteria existed in the sediments with smaller individual than the actinomycetes and fngus. In natural ecosystems, there is generally a law, i.e., the individual size of organisms is negatively related to their invidual numbers. Secondly, the abuncance of these microorganisms in the pool in different reaches obviously exceeded the rapids and benchlands. In the pools, there was more nutrient accumulation in the sediments because of a slower water flow than in the rapids, which was banificial to the growth and reproduction of the microoganims. In the benchlands, floods often resulted in the alternate occurrence of waterflooding and dry environments and the delivary of the nutrients and consequently which was beneficial to the growth and reproduction of the aerobic and anaerobic microorganism. Greater loss of nutrients and disturbance of water flows in the rapids might refrain the growth and reproduction of the microorganisms than the pools and benchlands. Thus, the abundance of these microorganisms in the rapids presented relatively less than the pools and benchlands. Thirtly, the abudance of these microorganisms were higher in the upper reaches than the lower reaches because there were more COD input to the river in the upper reaches.

Enzyme activity

Catalase activity in the sediments averaged 3.638 ml/g in all rapids, pools and benchlands in the three river reaches in Dulu river and ranged from 1.74-5.465ml/g. The activity of the enzyme was relatively higher on average in the lower reaches than the upper and middle reaches ($P<0.01$, Fig 3A). However, the activity of the enzyme was lowest in the pools of all three river reaches and highest in the benchlands in the upper and middle reaches

($P < 0.01$) . In the rapids in the upper and middle reaches, the activity of the enzyme in the sediments presented moderate levels. These results occurred possibly because in the benchlands, there were high oxygen contents in the sediments under an alternative floodingwater conditions with a change of water levels in the river, which promoted the catalysis of the enzyme in decomposition of organic matter and correspondingly the activity of the enzyme. Similarly, the activity of the enzyme was also higher in the rapids than the pools because there was relatively richer dissolved oxygen in the rapids.

Table 2 The number of the microorganisms in the sediments of the rapid-pool-benchland system

River reaches	Systems	Bacteria ($\times 10^6 \cdot g^{-1}$)	Actinomycetes ($\times 10^4 \cdot g^{-1}$)	Fungus ($\times 10^4 \cdot g^{-1}$)	Amonifying bacteria ($\times 10^5 \cdot g^{-1}$)	Denitrifying bacteria ($\times 10^2 \cdot g^{-1}$)	Total ($\times 10^6 \cdot g^{-1}$)
Upper reaches	Pool	5.50±0.32	18.20±1.39	11.00±0.71	21.44±1.72	14.47±1.05	7.84±0.53
	Rapid	2.22±0.15	0.71±0.06	1.68±0.11	3.21±0.15	5.32±0.23	2.55±0.12
	Benchland	2.77±0.12	15.60±1.22	1.76±0.15	5.17±0.18	9.29±0.38	3.45±0.16
Middle reaches	Pool	2.14±0.17	10.01±1.07	4.60±0.21	5.21±0.20	4.31±0.17	2.68±0.13
	Rapid	0.32±0.02	0.61±0.03	1.08±0.05	0.33±0.01	1.22±0.01	0.66±0.02
	Benchland	0.96±0.08	8.55±0.57	1.12±0.04	1.46±0.08	1.79±0.01	1.20±0.04
Lower reaches	Pool	3.79±0.21	12.30±0.82	6.10±0.34	18.34±1.20	8.89±0.49	5.75±0.24
	Rapid	0.14±0.08	9.12±0.47	0.18±0.01	2.45±0.09	1.35±0.01	1.50±0.04
	Benchland	1.26±0.09	10.40±0.66	0.21±0.01	3.23±0.10	3.79±0.22	1.57±0.04

Phosphatase activity in the sediments averaged 3.066 ml/g in all rapids, pools and benchlands in the three river reaches and ranged from 0.357-8.889mg/g ml/g. The activity of the enzyme was relatively higher on average in the upper reaches than the middle and lower reaches ($P < 0.01$, Fig 3B), as just opposed to the catalase activity. However, the activity of the enzyme was lowest in the benchlands of all three river reaches and highest in the pools ($P < 0.01$) , as also opposed to the catalase activity. In the pools in the upper reaches, the activity of the enzyme was about 2 times higher than the rapids in the middle reaches and 3 times higher than the benchlands in the lower reaches. Moreover, the differences of the activity of the enzyme were also pretty obvious among the rapids, pools and benchlands in the upper, middle or lower reaches. Generally, the activity of the enzyme in the pools was 5 time higher than the benchlands. However, the difference between the pools and rapids was relatively small because of the habitat similarity. These interesting results might be caused by the relatively high pH in the waterbody in the upper reaches and pools.

Urease activity in the sediments averaged 1.245mg/g in all rapids, pools and benchlands in the three river reaches and ranged from 0.243-2.511mg/g. The activity of the enzyme did not present a regular change from the upper to lower reaches as the catalase and phosphatase activities (Fig 3C). Overall, the activity of the enzyme in the rapids, pools and benchlands in the lower reaches was higher than the upper and middle reaches. The differences of the activity of the enzyme were very significant among the rapids, pools and benchlands in the upper and lower reaches. However, there was not a significant difference in the activity of the enzyme among the rapids, pools and benchlands in the middle reaches ($P > 0.01$) . In the middle reaches, the benchlands got the highest activity of the enzyme, and the pools and rapids were ranked second and third, respectively, as was fully different from the results in the activity of the enzyme in the upper and lower reaches. The result in the middle reaches might had been affected by some stachastical factors.

The dehydrogenase activity in the sediments averaged 1.071 μ g/g·h in all rapids, pools

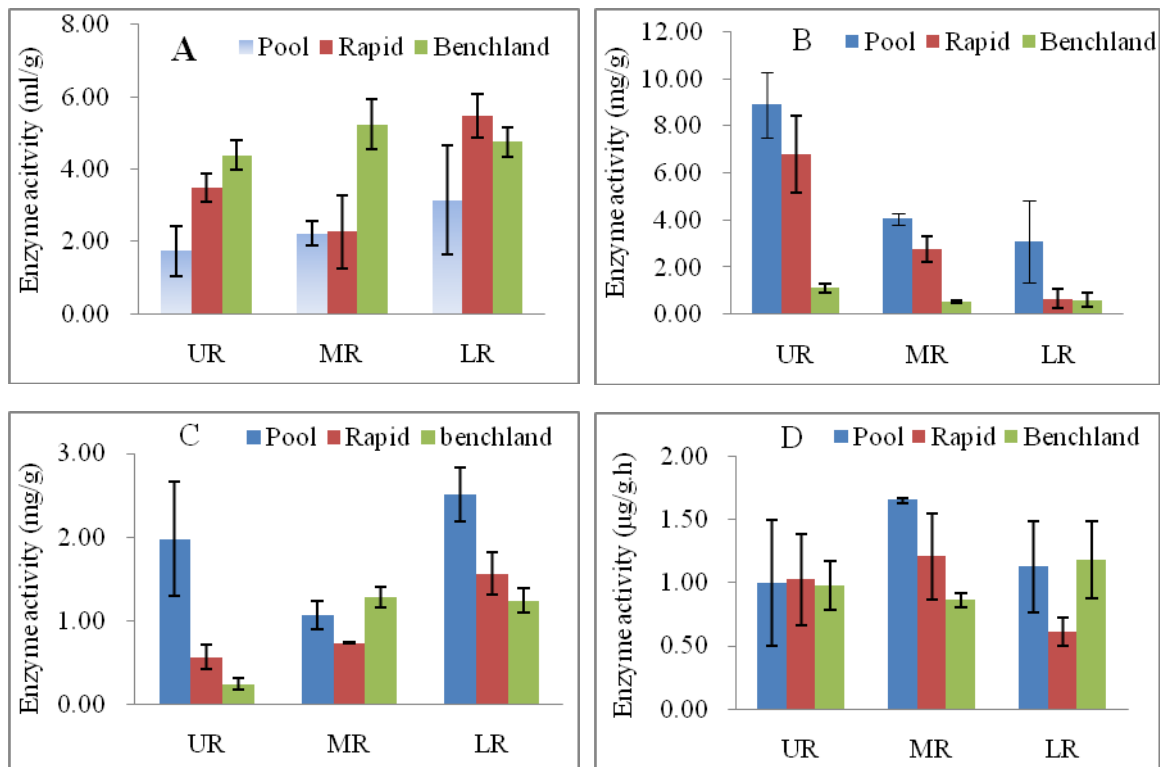


Figure 3 The enzyme activities in the sediments of the rapid-pool-benchland in the upper, middle and lower reaches in Dulu River. A: catalase; B: phosphatase; C: urease; D: dehydrogenase.

and benchlands in the three river reaches and ranged from 0.61-1.65 $\mu\text{g}/\text{g}\cdot\text{h}$, lower than the three enzymes as above. The activity of the enzyme did not also present a regular change from the upper to lower reaches as the catalase and phosphatase (Fig 3D). Moreover, the activity of the enzyme did not also indicated a significant difference among the lower, middle and lower reaches ($P>0.05$). In the upper reaches, the activity of the enzyme was almost fully identical in the rapids, pools and benchlands. The differences and regularity of the activity of the enzyme were very significant among the rapids, pools and benchlands in the middle reaches. The activity of the enzyme in the pools and benchlands was very similar but which was far greater than that in the rapids.

Relationships between the activity of the enzyme and the abundance of microorganisms

There were significant relationships between the abundance of all microorganisms and the activities of the urease and phosphatase in the sediments in the rapids, pools and benchlands in different river reaches (Table 3). However, statistical analysis only indicated a significant relationship of the catalase activity with two type of microorganisms, i.e., actinomycetes and bacteria among all tested microorganisms. Similarly, the dehydrogenase activity was only related to the bacteria and ammonifying bacteria as well. These high-significant relationships indicated that the enzymes and the corresponding microorganisms releasing the enzymes were important to the generation of the resultants catalyzed by these enzymes. Conversely, these low relationships indicated that the enzymes and the corresponding microorganisms releasing these enzymes did not play a key role in the formation of the resultants generated by the catalysis of these enzymes. The enzymes in the sediments come from a release of the cells of microorganisms and the abundant microorganisms in the sediments must produce a great amount of different enzymes, effectively catalyzing reactants to generate resultants, and consequently being tested high enzyme activity. However, the generation of some resultants is not fully dependent on the enzyme catalysis and other factors also affect the formation of the

resistants. For example, the hydrogen peroxide is catalyzed by the catalase to generate oxygen and water, but Mn and Fe also catalyze the hydrogen peroxide to generate the oxygen and water. Thus, no relationships between the enzyme activity and the microorganisms might have indicated that the effects of other factors perhaps exceed the catalysis effects of the enzymes from low abundant microorganisms.

Table 3 Correlation coefficients between the enzyme activity in the sediments and the abundance of microbes

Microorganisms	Catalase	Phosphatas	Urease	Dehydrogenase
Bacteria	0.424*	0.836**	0.651*	0.619*
Fugus	0.308	0.758*	0.498*	0.092
Actinomycetes	0.612*	0.712*	0.677*	0.128
Amonifying bacteria	0.232	0.653*	0.812**	0.372*
Denitrifying bacteria	0.077	0.689*	0.791*	0.108

Sign ** and * represent a significance at 0.95 and 0.99 confidence levels, respectively; N=27.

DISCUSSION

Previous much studies on abundance of microorganisms and activity of enzymes in the sediments mainly focused on the city riverway, wetlands and lakes under the pollution conditions to cope with the challenges of water pollution problems and less data has been collected from a natural river (Winiarski, Bedell, Delolme et al., 2006; Neto, Ohannessian, Delolme et al., 2007; Huang, Gao, Liu et al., 2012). In a natural river, the river structure is complete and consequently ecosystem functions of the riverway such as biodiversity, self-purification of waterbody, landscape values and affinity between humans and waterbody are also complete (Padmalal and Maya, 2014). The rapid-pool-benchland system is a basic unit in river structure in a natural river based on our previous studies (Ma, Wang, Chen, 2014; Wu, Wang, Cheng, 2014; Chen, Ma, Wang, 2015; Wang, Wang, Liu et al., 2015). Therefore, we selected the rapid-pool-benchland system as a representative of a complete structure in Dulu River and tested the abundance of microorganisms and activity of enzymes in the sediments to clarify the effects of the river structure on the microorganism patterns, and the relationships between the abundance of the microorganism and the biochemical functions. The final goal of the study is to clarify the important significance of the maintenance and restoration of a natural river structure.

In each of the rapid-pool-benchland systems, the abundance of the microorganisms (including bacteria, fungus, actinomycetes, amonifying bacteria and denitrifying bacteria) in the sediments of the pools in different reaches is significantly higher than the rapids and benchlands. Comparatively, the abundance of the microorganisms in the benchlands is also higher than the rapids. These results indicate that the habitat heterogeneity surely results in the differences of the abundance of the microorganisms and verify our assumption. However, the abundance of the bacteria, fungus, actinomycetes, amonifying bacteria and denitrifying bacteria in sediments respectively occupied 60%–99.07%, 0.012–0.172%, 0.28%–23.21%, 5.00%–31.89%, 0.009%–0.027% in total abundance of microorganisms regardless of the habitat difference in the rapid-pool-benchland systems and these proportions are similar to those in the artificial wetland (Du, Huang, Gao, et al., 2013) and in a calcareous purple paddy Soil (Gu YF, Yun X, Zhang XP, et al., 2008). This indicates that the habitat difference within the rapid-pool-benchland systems do not significantly affect the relative abundance among different groups of microorganism. The study results also indicated that the microorganisms was more abundant in the rapid-pool-benchland systems in the upper reaches, where there

were greater loads of organic pollution than the middle and lower reaches. This agrees with the previous results that some pollution matter in habitats can induce the growth of microorganism (Hill and Moffett, 2010; Millar, Payne, Ochs et al., 2015).

The enzyme activity in the sediments can reflect the efficiency in matter transformation, and the ability that organic pollution mater may be discomposed (Garcia et al. 1994; Neto Ohannessian, Delolme et al., 2007; Reboreda and Cacador, 2008). The enzyme activity is also one way of describing the general condition of soil microbial populations (Margesin, Zimmerbauer, Schinner, 2008). In the rapid-pool-benchland systems, the activity of catalase and phosphatase in the sediments presented significantly different among the rapids, pools and benchlands, indicating the habitat difference greatly induces variation of the activity of catalase and phosphatase. The catalase plays a great role in the decomposition of hydrogen peroxide and in the prevention of soil microorganism, plant roots and soils from harmfulness of hydrogen peroxide in sediments (Huang, Gao, Liu et al., 2012). The phosphatase catalyzes dephosphorylation of substrate in sediments to promote the decomposition of organic matter or produce the information moleculars to regulate biochemical functions. In the sediments of the rapids and pools, there is a higher activity of catalase and phosphatase, respectively (Fig.3), which suggests that the rapid-pool-benchland systems shape different environments for divergent biochemical functions in a river. Therefore, a habitat heterogeneity in a river structure, which can be denoted by a basic unit of a river, i.e., the rapid-pool-benchland systems, imposed different effects on ecosystem functions in a river. In addition, in the different river reaches in Dulu River, the activity of all enzymes was lower than that in the sediments in the lake (Liu, Xie, Li, 2013). This is mainly because the nutrients in the Dulu River were relatively poorer than those in the lake.

The biochemical functions in the sediments in different waterbodies are dependent on the catalysis of enzymes and the enzymes come from the secretion of microorganisms (Gardner and White, 2010). Thus, the abundance of microorganism should be positively correlated to the activity of enzymes. However, the analysis of the relationship suggests that there are positively and negatively linear relationships between the canonical variables of the microbes and the activity of catalase, urease and dehydrogenase in artificial wetlands possibly due to the interactions of these variables (Huang, Gao, Liu et al., 2012). In the study, the correlations between abundance of all types of microorganisms and the phosphatase and urease activities were very significant but dehydrogenase and catalase were only related to the abundance of two types of microorganisms, respectively. In the study of forest soils, although there is also a correlation between soil microorganisms and the enzyme activity, but a close correlation occurs between abundance of microorganism and physical and chemical property (Hu, Zhang, Gao et al., 2002). These results indicate that the enzyme activity in the sediments to affect the biochemical functions is decided by many factors including the abundance of microorganism, although the rapid-pool-benchland systems also regulate the abundance of microorganism and the enzyme activity in the sediments.

CONCLUSIONS

The enzyme activity is crucial to biochemical functions in the sediments in a river. The microorganism abundance partially affects the enzyme activity and further the biochemical functions. The rapid-pool-benchland systems, a basic unit of a river, reflect the habitat heterogeneity in a river, and directly regulate the abundance of microorganism and the enzyme activity in the sediments. The restoration of the river structures disturbed by humans towards the rapid-pool-benchland systems is beneficial to a complete function of the river ecosystem.

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