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2	<u>Sedimentary context controls the influence of ecosystem engineering by bioturbators on</u> <u>microbial processes in river sediments</u>
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By modifying the physical environment, ecosystem engineers can have inordinately large effects on surrounding communities and ecosystem functioning. However, the significance of engineering in ecosystems greatly depends on the physical characteristics of the engineered habitats. Mechanisms underlying such context-dependent impact of engineers remain poorly understood whereas they are crucial to establish general predictions concerning the contribution of engineers to ecosystem structure and function.

7 The present study aimed to decrypt such mechanisms by determining how the environmental 8 context modulates the effects of ecosystem engineers (bioturbators) on micro-organisms in 9 river sediments. To test the effects of environmental context on the role of bioturbators in 10 sediments, we used microcosms and recreated two sedimentary contexts at the laboratory by 11 adding a layer of either fine (low permeability) or coarse (high permeability) sand particles at 12 the top of a gravel-sand matrix. For each sediment context, we examined how the sediment 13 reworking activity of a bioturbating tubificid worm (Tubifex tubifex) generated changes of the physical (sediment structure and permeability) and abiotic environments (hydraulic discharge, 14 15 water chemistry) for micro-organisms. The biotic influences of the bioturbation process were 16 measured on microbial characteristics (abundances, activities) and leaf litter decomposition as 17 a major microbe-mediated ecological process.

18 Results showed that T. tubifex significantly increased hydraulic discharge (by about 6-fold) 19 and restore aerobic conditions in O₂-limited sedimentary habitats covered by fine sand (low permeability treatment). Consequently, worms stimulated microbial communities developed 20 21 on buried leaves and leaf litter breakdown increased (+30%). In contrast T. tubifex had a low influence on water exchanges and O₂ availability in highly permeable sediments. As 22 23 bioturbation did not modify the abiotic environment in sedimentary habitats covered by 24 coarse sand, T. tubifex did not influence the abundances and activities of microorganisms 25 developed on leaves.

1 Our study demonstrated that the significance of ecosystem engineering on the functioning of 2 aquatic ecosystems cannot be quantified without assessing the complex interactions between 3 bioturbation activities and sedimentary characteristics. We strongly suggest that context 4 dependency mainly modulates the effects of ecosystem engineers on other biota by 5 controlling the ability of engineers to modulate the availability of limiting factors for other 6 organisms.

7

8 Introduction

9 Habitat modification by engineer organisms has been recognized as a major ecological 10 process with large consequences for biodiversity and ecosystem functions in a broad range of 11 ecosystems (e.g. see Lavelle et al 1997, Crooks 2002, Mermillod-Blondin and Rosenberg 12 2006, Wright and Jones 2006, Wright et al. 2006, Badano and Marquet 2008, Gutiérrez et al. 13 in press). Ecosystem engineers can have inordinately large effects on communities by 14 modifying their surrounding physical environment (Jones et al. 1994, see Fig.1 for detailed 15 engineering sequence). The beaver (Castor canadensis) is a classical example of ecosystem 16 engineer. Its activities (essentially dam-building) can increase (i) the proportion of flooded soils (water and wetlands) in the landscape (Johnston and Naiman 1990) and (2) the retention 17 18 of sediment, organic material (Naiman et al. 1986) and nutrients (Naiman and Melillo 1984) 19 in the channel by decreasing water velocity, ultimately affecting the structure of animal and 20 plant communities in the landscape (Naiman et al. 1988, Hägglund and Sjöberg 1999, Wright 21 et al. 2002, Anderson and Rosemond 2007). Beside this emblematic example of beavers, it 22 exists a wide diversity of engineer organisms and then engineering mechanisms (Berke 2010) that significantly impact structure and functions. For example, the sediment reworking 23 24 activities of bioturbators can have marked effects on microbial communities developed on

1 sediments by affecting hydrological fluxes and biogeochemistry at water-sediment interface 2 (Mermillod-Blondin and Rosenberg, 2006; Mermillod-Blondin et al. 2004, Nogaro et al. 2009). Whatever the engineering mechanisms (e.g. through bioturbation, physical engineering 3 4 ...), a change in resource availability relative to the unmodified state may suffice to observe 5 positive or adverse engineering effects on some biotic variables (e.g. absolute and relative 6 abundance and richness of species in the surrounding communities) describing the structure of 7 communities (Bertness 1984a,b, Jones et al. 1997, Menge 2000). The theory of ecosystem 8 engineers also suggests that the highest effects of engineers on physical habitat produced the 9 highest changes of biotic variables (Jones et al. 1994). However, the influence of an 10 ecosystem engineering activity also depends on the environmental context in which it happens 11 (Crain and Bertness 2006, Wright and Jones 2006). Because ecosystem engineers affect 12 community through environmentally mediated interactions, a given engineering process 13 (similar density of a given engineer species) can have contrasted effects on some biotic 14 variables across environmental gradients and thus can appear as idiosyncratic (Jones et al. 15 2004, Moore 2006). However, most of studies dealing with ecosystem engineering only 16 examine the relationships between the engineer (e.g. number of individuals) and (i) the 17 physical characteristics of the environment or (ii) the other species, in a unique environmental 18 context. To establish general principles and predictions (for future changes) about the effects 19 of engineers on communities, there is a need to fully understand mechanisms underlying 20 context dependency and thus to examine the whole 'cause-effect relationships' sequence (see 21 Fig. 1) in contrasted environments.

In the present study, we aimed to tackle this context dependency in river sediments. The functioning of lotic ecosystems partly depends on the microbially-mediated biogeochemical processes (nutrient cycling, organic matter (hereafter OM) processing) realized in the hyporheic zone (hereafter HZ, sedimentary interface between surface water and groundwater)

1 (Grimm and Fisher 1984, Findlay 1995, Naegeli and Uehlinger 1997, Boulton et al. 1998, 2 Fellows et al. 2001). By controlling hydrological exchanges between the surface and the sediments, and thus chemical conditions (e.g. availability of dissolved oxygen DO) in the HZ, 3 4 sediment characteristics such as permeability appears as crucial factors controlling the 5 structure and activities of microbial communities developed on river sediments (e.g. Valett et 6 al. 1990, Brunke and Gonser 1997, Mermillod-Blondin and Rosenberg 2006). In this context, 7 bioturbators can have major influence on the HZ (and *a fortiori* on the whole-stream) 8 ecological functioning through modification of sediment structure and permeability (Nogaro 9 et al. 2009, Nogaro & Mermillod-Blondin 2009). However, the magnitude of bioturbation-10 driven change in hydraulic conductivity is expected to depend on physical hydraulic 11 conditions which are more or less constraining for micro-organisms living in HZ (Hakenkamp 12 and Palmer 2000, Mermillod-Blondin, in press). As physical hydraulic conditions are 13 essentially linked with sediment characteristics, the effects of bioturbators as ecosystem 14 engineers in river sediment are expected to be sedimentary-context dependent. By applying 15 the conceptual framework of ecosystem engineering in rivers, bioturbators are expected to 16 major influence on hyporheic ecological processes in sedimentary systems where: (1) they are 17 able to drastically modify habitat physical characteristics by actively reworking sediment, (2) these physical changes result in alteration of the hydraulic conductivity in the sedimentary 18 19 habitat and finally (3) the change in hydraulic conductivity modifies resource availability for 20 interstitial microorganisms involved in studied processes (Fig.1). In riverbeds covered by 21 excessive deposition of fine sedimentary particles, permeability, hydrological exchanges and 22 associated input of resources from the surface for interstitial organisms are reported to be low 23 (Beschta and Jackson 1979, Schächli 1992, Wood and Armitage 1997). In this context, 24 bioturbation would have more influence on water exchanges between surface and hyporheic 25 zone than in highly permeable sediments.

1 The aim of our study was therefore to examine how the environmental context (sediment characteristics) modulates the effects of an active bioturbator as ecosystem engineer in river 2 3 sediments. We tested the effects of the bioturbator Tubifex tubifex Müller (Oligochaeta, 4 Tubificidae) on the characteristics and activities of microbial communities in two sedimentary 5 habitats with contrasted textures (topped by coarse permeable sand *versus* topped by fine sand 6 with low permeability) by using slow filtration columns. For each habitat, we broke down and 7 examine the ecosystem-engineering process into detailed intermediate steps (see Fig.1) 8 including sediment reworking, physical (hydraulic conductivity) and abiotic (availability of 9 nutrients and electron acceptors used for OM mineralization) changes on the sedimentary 10 habitat. The ecosystem-engineering effects (biotic changes) were measured on the 11 characteristics (bacterial and fungal abundances) and activities (potential aerobic respiration, 12 potential denitrification, hydrolytic exoenzymatic activities) of the microbial community 13 developed on leaves buried in sediments. Decomposition of leaf litter (measurement of 14 breakdown rates) -a crucial microbially-mediated ecological process in river sediment- was 15 used as a final step in the engineering process. Bioturbators were expected to have the most 16 significant influence in sedimentary systems covered by fine particles, where restricted hydrological exchanges (low hydraulic conductivity and associated sharp decrease in O₂ and 17 following electron acceptors) are supposed to constrain microbial communities. 18

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20 Methods

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22 Experimental design

To address how sediment context modulates the effects of a bioturbator at the watersediment interface of rivers, we employed a factorial experimental approach in which the occurrence of *T. tubifex* and sediment texture were manipulated. Experiments were carried out in slow filtration columns (n = 16 experimental units, height = 35 cm and inside diameter = 10
cm; Mermillod-Blondin et al. 2005, Navel et al. 2010) filled with sediment, at constant
temperature (15 ± 0.5 °C) under a 12 h light / 12 h dark cycle.

4 These mesocosms were filled by successively adding gravel (2-4 mm diameter, 300 g) and 5 then sand (100-1000 μ m, 170 g), 8 times. We manipulated the surface sediment texture by 6 adding a 2 cm thick layer of either fine sand (90% of particles $< 150 \mu m$ diameter, low permeability: "fine sand treatment", n = 8 columns) or coarse sand (90% of particles > 300 7 8 μ m diameter, n = 8 columns) at the top of the gravel-sand base. The thickness of the top 9 sediment layer was in accordance with observations reported from riverbeds impacted by fine 10 sediment deposits (Wood & Armitage 1997). Analyses performed before the start of the 11 experiment indicated that the specific area and the amounts of total organic carbon (TOC), 12 nitrogen (TN) and phosphorus (TP) in the sediment were higher in the fine than in the coarse 13 sand (Table 1). All the sedimentary material was collected from the Rhône River, elutriated 14 and cleaned with deionised water to eliminate fauna and coarse particulate organic matter 15 (CPOM). Moreover, the whole-sediment layer was kept in the dark to suppress possible 16 photoautotrophic processes.

17 During sediment installation, a set of 35 leaf discs (diameter: 12 mm) of alder (Alnus glutinosa (L.) Gaertner), a common species along rivers characterized by fast leaf degradation 18 19 (Abelho 2001), was inserted between two circular sieves (3 mm mesh) at a depth of 9 cm 20 below the sediment surface in each column. Discs were cut avoiding central veins of leaves 21 collected from the riparian zone of the Rhône River during abscission (October 2008). Leaves 22 were conditioned in small-mesh bags immersed in a nearby river (located on the campus of 23 the University Claude Bernard Lyon 1, Lyon, France) for 10 days, i.e. a time sufficient to 24 allow microbial colonization (Suberkropp and Chauvet 1995). After installation of sediment and leaf litter, aerated artificial river water (96 mg.L⁻¹ NaHCO₃, 39.4 mg.L⁻¹ CaSO₄ · 2H₂O, 25

60 mg.L⁻¹ MgSO₄ · 7H₂O, 4 mg.L⁻¹ KCl, 19 mg.L⁻¹ Ca(NO₃)₂ · 4H₂O and 1.6 mg.L⁻¹
 (CH₃CO₂)₂CaH₂O; pH = 7.5; US EPA 1991) was added at a constant hydraulic head (ΔH = 3
 cm), to a depth of 10cm above the sediment surface of each mesocosm. Openings in each mesocosm allowed sampling water at different times during the experiment.

5 Seven days after sediment installation (T7) (time necessary to obtain a physico-chemical 6 stabilization of the system), we added a set of 100 individuals of T. tubifex to half of the 7 experimental units (n = 4 per treatment). The density of tubificid worms in the experimental units (around 12,800 individual.m⁻²) was in accordance with densities reported in field studies 8 (Fruget 1989, Martinet 1993). T. tubifex is a common deposit feeder that inhabits sandy and 9 10 muddy habitats, which can actively rework sedimentary particles (McCall and Fisher 1980) 11 and increase sediment permeability (Nogaro and Mermillod-Blondin 2009, Nogaro et al. 12 2009). The potential influence of T. tubifex on leaf litter degradation was expected to result 13 from the influence of T. tubifex as ecosystem engineers rather than a direct feeding on leaf 14 litter. To verify that T. tubifex do not feed on leaves, we conducted a preliminary experiment 15 using aerated aquatic microcosms in which 35 alder leaf discs were deposited at the surface of 16 a fine layer of sediment for 59 days. We measured that the occurrence of 100 individuals T. 17 tubifex did not significantly influence the leaf litter breakdown rate nor the microbial abundances and activities associated with leaf litter (unpublished data). 18

During the main experiment, hydraulic discharge rate was measured and water was sampled every 10 days at 4 depths to determine O_2 , NH_4^+ , NO_3^- , NO_2^- , PO_4^{3-} , SO_4^{2-} and dissolved organic carbon (DOC) concentrations, for all columns. At the end of the experiment, columns were dismantled and sediment was cut into slices to quantify sediment reworking and vertical distribution of invertebrates. Fungal biomass, total bacterial abundance, abundance of active eubacteria, potential aerobic and anaerobic activities and enzymatic activities involved in C and N cycles were determined on leaf discs, as described
 below. Leaf discs were then dried and weighed to quantify mass loss during the experiment.

3

4 Physico-chemical analyses

5 Every 10 days starting with day 6, a day before fauna addition (T6, T16, T26, T36, T46 and 6 T56), the outlet of each column was closed and water was shunted and sampled at +2 cm 7 above (H1) and -3 cm (H2), -8 cm (H3) and -13 cm (H4) below water-sediment interface 8 under similar hydraulic pressure conditions. An oxygen micro-sensor probe fitted in a glass 9 tube (OX 500, Unisense, Aarhus, Denmark) was used to determine O₂ concentration without contact with atmospheric oxygen during sampling. NH_4^+ , NO_3^- , NO_2^- , PO_4^{3-} and SO_4^{2-} 10 concentrations were determined following standard colorimetric methods (Grashoff et al. 11 12 1983) after filtration through Whatman GF/F filters (pore size: 0.7 µm; Millipore, Billerica, 13 MA, U.S.A.) using an automatic analyzer (Easychem Plus, Systea, Anagni, Italia). For DOC 14 measurements, water samples were filtered though Whatman HAWP filters (pore size: 0.45 15 µm; Millipore, Billerica, MA, U.S.A.) and acidified with 3 drops of HCl (35%). The DOC 16 concentration in water samples was measured with a total carbon analyzer (multi N/C 3100, 17 Analytik Jena, Jena, Germany) based on combustion at 900 °C after removal of DOC with 18 HCl and CO₂ stripping under O₂ flow.

19

20 Sediment reworking analyses

Particle redistribution induced by worms in the sedimentary matrix was estimated by the luminophore tracer technique (Gérino 1990). In each column, natural sediment particles (150-300 μ m) dyed with yellow luminescent paint were deposited uniformly at the top of the sedimentary matrix a few hours after the introduction of *T. tubifex* (at T7). During column dismantling (T59), the top 4 cm of sediment were cut into 0.5 cm thick slices, dried at 40 °C
(48 h) and homogenized before counting luminophores on 500 mg subsamples under U.V.
light (3 replicates per sampled slice). Vertical distribution of luminophores in the sediment
was obtained by expressing the density of particles (number.g⁻¹ dry sediment) obtained for
each slice as percentage of the total amount of luminophores obtained for the whole 4 cm top
sediment layer.

7

8 Vertical distribution of tubificid worms

9 After collecting subsamples on the top sediment for luminophore counting, sediment was 10 pooled into 5 cm thick sediment slices that were sieved (using a 500 μ m–diameter sieve) to 11 collect living tubificids. Individuals recovered in each slice were preserved in 96% ethanol 12 and counted under a dissecting microscope. For each column, the vertical distribution of 13 tubificid worms in the sediment was determined by reporting the abundance of worms in each 14 slice to the total amount of worms retrieved in the overall sedimentary column (results for 15 each slice were expressed as percent).

16

17 Microbial analyses

18 Fungal biomass

For each column, 5 leaf discs collected at the end of the experiment were stored at -80 °C and freeze-dried for 12 h before analysis. Fungal biomass was estimated with the ergosterol quantification method in which saponified products were obtained by methanol refluxing prior to saponification reaction using KOH/methanol (Gessner et al. 2003), following the protocol detailed in Navel et al. (2010). Ergosterol was isolated from saponified products by using Oasis HLB 3cc extracting columns (Waters Corporation, Milford, MA, U.S.A) and
elution with isopropanol. Mass of ergosterol in the sample was then calculated by using
HPLC system (HPLC 360/442, Kontron, Eching, Germany). Fungal biomass was estimated
from ergosterol amounts using a 182 conversion factor determined for aquatic hyphomycetes,
which are known to dominate fungal assemblages on decomposing litter (Gessner and
Chauvet 1993). Results were expressed in mg fungi.g⁻¹ dry mass of leaf litter.

7

8 Bacterial abundances

9 During column dismantling (at T59), 2 leaf discs were immediately collected and fixed in 4% paraformaldehyde in phosphate-buffered saline (PBS; 0.13M NaCl, 7mM NaHPO₄, 3mM 10 11 NaH₂PO₄; pH=7.2) for 10 h. Fixed samples were subsequently washed twice in PBS and were stored in ethanol and PBS (50:50) at 20 °C. After storage (2 weeks), leaf discs were 12 13 homogenized in 20 mL of 0.1% pyrophosphate in PBS using a sonicator with a 2 mm-14 diameter probe at 50 W for two periods of 60 s. All homogenized samples were finally 15 supplemented with the detergent NP-40 (Flucka, Buchs, Switzerland) to a final concentration 16 of 0.01 %. Aliquots (10 µL) of homogenized samples were spotted onto gelatine-coated slides 17 and were hybridized with Cy3-labelled oligonucleotide probe (mix of EUB 338, EUB 338 II 18 and EUB 338 III, eubacteria) and concomitantly stained with the DNA intercalating dye DAPI (200 ng.µL⁻¹, Sigma, Buchs, Switzerland) according to Navel et al. (2010). Numbers of 19 20 DAPI- and Cy3-bacteria were expressed as numbers of bacteria and numbers of active 21 eubacteria (hybridized with EUB 338, Karner & Fuhrmann 1997) per g dry leaf.

22

23 Microbial activities

All microbial activities were measured within the 24 h following columns dismantling, with
 leaf discs stored at 4 °C before analysis.

3 Enzymatic activities: β –glucosidase (EC: 3.2.1.21), β –xylosidase (EC: 3.2.1.37) and leucine aminopeptidase (EC: 3.4.11.1) activities were measured on 2 discs (2 times) by fluorimetry 4 5 using constant volume of substrate analogs: 4-methylumbelliferyl-ß-D-glucoside (MUF-glu; 6 750 µM, 2 mL), 4-methylumbelliferyl-xylosidase (MUF-xyl; 1000 µM, 2 mL) and L-Leucine-7 4-Methyl Coumarinyl-7-amideHCl (MCA-leu; 1000 µM, 2 mL), respectively. Incubation at 8 20 °C (40 min) was stopped by transferring into boiling water before centrifugation (5000 G; 9 4851 rpm, 3 min). Fluorimetry measurements were realised on a mix of supernatant (300 μ L) and buffer (30 µL, pH 10.4) using a microplate reader (SAFIRE, TECAN Group Ltd, 10 11 Switzerland) with excitation wavelength of 363 nm and emission wavelengths of 441 nm for 12 MUF-glu and MUF-xyl. Wavelengths were set at 343 nm (excitation) and 436 nm (emission) 13 for MCA-leu. Litter dry mass (drying at 70 °C for 48h) was determined at the end of analyses to express results as nmol of hydrolysed compound.h⁻¹.g⁻¹ dry leaf litter. For each sample, 14 values were corrected by the fluorimetric signal obtained with a formaldehyde-killed control 15 (measurements realised in similar conditions on 2 discs previously treated 30 min with a 39 % 16 formaldehyde solution). 17

Potential aerobic respiration and anaerobic denitrification activities were measured on leaf 18 19 discs following the slurry technique (Furutani et al. 1984). Leaf discs (n = 4 for respiration 20 and n = 6 for denitrification) were placed in 150 mL flasks supplemented with feeding solutions to optimize microbial activity. For the measurements of CO₂ production 21 22 (respiration), the incubation was conducted under aerobic conditions with 5 mL of a feeding solution of glucose (7.5 g.L⁻¹) and glutamic acid (7.3 g.L⁻¹). For the measurements of N₂O 23 production (denitrification), the incubation was under anaerobic conditions with a N₂ 24 atmosphere. The feeding solution was a mixture of 5 mL of a KNO₃ (2.2 g.L⁻¹), glucose (7.5 25

g.L⁻¹) and glutamic acid (7.3 g.L⁻¹) solution. Acetylene (10% v/v) was introduced in N₂
saturated atmosphere to stop N₂O-reductase activity. CO₂ and N₂O productions were
calculated from measurements of concentrations at 2 h and 6 h incubations by using gas
chromatography on a microcatharometer (M200 micro gas chromatograph, MTI Analytical
Instruments, Richmond, CA, U.S.A.). After the drying of leaf discs (70 °C for 48h), results
were expressed in µg of C or N.h⁻¹.g⁻¹ dry leaf litter.

7

8 Leaf litter degradation

9 For each column, the total dry mass of leaf litter after 59 days was calculated as the sum of 10 the dry masses of samples used in microbial analyses and that measured for the remaining leaf 11 material (common drying method: 70 °C for 48 h), with correction for the set of 5 discs that 12 were freeze-dried for fungal biomass assessment. Results were compared to the initial dry 13 mass determined on 5 additional sets of 35 alder discs (228.8 \pm 6.25 mg) at the start of the 14 experiment.

15

16 Data treatment

Repeated measures of hydraulic conductivity were analyzed using mixed model analysis of 17 18 variance with "sediment" ("coarse sand" vs. "fine sand"), "worms" ("with" vs. "without") and "time" as fixed factors, and experimental unit ("column") as random factor. Repeated 19 measures of vertical profiles in O_2 , DOC, NH_4^+ , NO_3^- , NO_2^- , SO_4^{2-} and PO_4^{3-} were analysed 20 21 similarly, with "depth" as additive fixed factor. Vertical distribution of tubificid worms was studied by using two-way analysis of variance (ANOVA) with "sediment", and "depth" as 22 23 main factor. Vertical distribution of luminophores was studied by using similar procedure 24 with "sediment", "depth" and "worms" as main factor. Influences of sediment permeability

and tubificid worms on data obtained on buried leaf litter (daily dry mass loss, fungal
biomass, total abundance of bacteria, abundance of active bacteria, % active bacteria,
enzymatic activities, potential aerobic respiration and potential denitrification) were examined
using two-way ANOVAs with "sediment" and "worms" as main factors. The method of
contrast was used to determine significant differences between treatments (Crawley 2002).

Hydraulic conductivity and microbial activities on leaves (glucosidase, 6 leucine aminopeptidase activity and potential denitrification activities) were log-transformed before 7 8 statistical analysis in order to fit the assumption of homoscedasticity. Abundances of 9 luminophores and worms retrieved at the end of experiment for each layer within a same 10 column were expressed as percent of the total abundance for the whole column, and were 11 arcsin-transformed before analyses. Statistical analyses were performed using JMP 8.0 (SAS 12 Institute, Cary, NC, U.S.A.), version 8.0.1. Significance for all statistical tests was accepted at 13 $\alpha < 0.05$.

14

15 Results

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17 Influence of sediment physical characteristics on sediment reworking activity.

The physical structure of the habitat influenced the vertical distribution of tubificid worms in sediment (Fig. 2; "sediment-by-depth interaction effect": $F_{(3,24)} = 83.44$, $P < 10^{-4}$). While the major part of individuals were retrieved in the top sediment layer when covered by fine sand (93% in the first 5 cm), most of worms were retrieved deeper in the sediment of columns covered with coarse sand (around 65% were found after 10 cm depth). The presence of worms in systems increased the transport of luminophores from the sediment surface to the sedimentary column (Fig. 2; "worms-by-depth interaction effect": $F_{(7.96)} = 29.78$, $P < 10^{-4}$). 1 This effect of worms on luminophore profiles was strongly influenced by the physical 2 characteristics of the sedimentary habitat ($F_{(7,96)} = 9.61$, $P < 10^{-4}$). The percentage of 3 luminophores buried at depth was less than 5% in the "coarse sand" treatment whereas it was 4 more than 20% in the "fine sand" treatment" (Fig. 2).

5

6 Influence of the sediment physical characteristics on hydraulic exchanges and microbial 7 processes involved in CPOM processing

Mean (\pm S.D.) hydraulic conductivity measured for the "fine sand" treatment was about 8-fold 8 lower than for columns topped with "coarse sand" (Fig. 3; 2.02 \pm 0.67 and 15.95 \pm 9 4.81 cm.h⁻¹, respectively). The decreases with depth of O_2 and NO_3^- concentrations (Fig. 4; 10 $F_{(3,276)} = 933.73$ and 134.90, respectively, $P < 10^{-4}$ for both) in the interstitial water were 11 higher in "fine sand" than in "coarse sand" treatment ("sediment-by-depth interaction effect": 12 $F_{(3,276)} = 200.85$ and 159.26 for O₂ and NO₃⁻ respectively, $P < 10^{-4}$ for both). This difference 13 14 was particularly marked in the top sediment layer (O₂ concentrations reduced by about 87% and 13% in "fine sand" and "coarse sand" treatments, respectively), and led to lower O₂ and 15 NO_3^{-1} concentrations in the sedimentary habitat covered by fine sand deposits ($F_{(1,12)} = 631.70$ 16 and 409.18 for O₂ and NO₃⁻ respectively, $P < 10^{-4}$ for both). Peaks of DOC, NH₄⁺, NO₂⁻ and 17 PO4³⁻ concentrations were only recorded in the "fine sand" treatment, leading to higher 18 concentrations of these solutes in "fine sand" than in "coarse sand" treatment (Fig. 4; $F_{(1,12)}$ = 19 93.02, 665.57, 193.27, 16.13, for DOC, NH_4^+ , NO_2^- and PO_4^{3-} respectively, $P < 10^{-4}$ for all). 20

Determinations of the dry mass of leaf litter retrieved at the end of the experiment (Fig. 5A) showed that the daily mass loss rate was 31% lower in the "fine sand" treatment than in the "coarse sand" treatment ($F_{(1,12)} = 14.39$, P = 0.043). In parallel, the total abundance of bacteria (Fig. 5B), the abundance of active bacteria (Fig. 5C), the fungal biomass (Fig. 5D), the

glucosidase activity (Fig. 5H) and the leucine aminopeptidase activity (Fig. 5I) were
significantly lower in "fine sand" than in "coarse sand" treatment (contrasts: comparisons
without *T. tubifex*: |t|₁₂ = 2.57, 2.87, 2.50, 5.18 and 7.14, respectively, *P* < 0.028 for all).
Potential aerobic respiration (Fig. 5E), potential denitrification (Fig. 5F) and xylosidase
activities measured on leaves were not significantly influenced by sedimentary conditions
(contrasts: |t|₁₂ = 1.43, 0.99 and 0.76, respectively, *P* > 0.176 for all).

7

8 Context-dependent influence of tubificid worms on hydrologic exchanges, biogeochemical 9 processes and CPOM processing

10 The influence of tubificid worms on hydraulic conductivity (Fig. 3) and water chemistry (Fig. 11 4) was dependent on the physical characteristics of the top sediment ("sediment-by-worms interaction effect": $F_{(1,12)} = 24.67, 48.41, 147.73, 13.76, 96.24$ and 4.04 for hydraulic 12 conductivity, O_2 , NO_3^- , SO_4^{2-} , NH_4^+ , and PO_4^{3-} concentrations, respectively, P < 0.044 for 13 all). While worms had low influence on hydraulic conductivity and concentrations of solutes 14 15 in "coarse sand" treatment, they increased by more than 6 fold the hydraulic conductivity in "fine sand" treatment. Consequently, the presence of tubificid worms increased O_2 and NO_3^- 16 concentrations and strongly reduced the peaks of solutes (DOC, NH_4^+ , NO_2^- and PO_4^{3-}) 17 18 released in the sedimentary columns with "fine sand" treatment.

Similarly, the influence of *T. tubifex* on microbial characteristics and associated processing of buried leaf litter was dependant on the physical structure of the sedimentary habitat (Fig. 4). We did not observe any influence of *T. tubifex* on microbial characteristics measured on leaves buried in "coarse sand" systems (Fig. 5; contrasts: $|t|_{12} = 0.24$, 0.83, 0.02, 1.32, 0.97 and 1.05 for total abundance of bacteria, abundance of active bacteria, fungal biomass, xylosidase, glucosidase and leucine aminopeptidase, respectively, P > 0.207 for all) except for potential aerobic respiratory activity ($|t|_{12} = 2.77$, P < 0.018). In contrast, *T. tubifex* had a positive influence on most microbial variables in "fine sand" treatment (contrasts: $|t|_{12} = 4.63$, 4.42, 3.05, 8.04 and 4.46 for total abundance of bacteria, abundance of active bacteria, xylosidase and leucine aminopeptidase respectively, P < 0.009 for all; and $|t|_{12} = 1.98$, P =0.071 for glucosidase activity) with the exception of potential denitrification ($|t|_{12} = 0.21$, P <0.835).

In parallel, *T. tubifex* increased the daily loss rate of leaf litter mass in the "fine sand"
treatment by about 30% (Fig. 4A; |t|₁₂ = 3.40, P < 0.006) where they counteracted the
negative influence of fine sediment deposition on leaf litter degradation (F_(1,12) = 5.61, P =
0.036). Such an effect was not observed in the "coarse sand" treatment (Fig. 5A, |t|₁₂ = 0.17, P
= 0.867).

- 11
- 12
- 13 Discussion
- 14

15 Contrasted biogeochemical processes induced by sediment characteristics.

16 Our study confirmed the expectation that the biogeochemical functioning of the hyporheic 17 zone is strongly influenced by the sediment context. The high hydraulic discharge rates in systems topped by a 2-cm thick layer of coarse sand (highly permeable systems) generated 18 aerobic conditions (O₂ concentration > 2 mg.L⁻¹) throughout the sedimentary matrix (to a 19 depth of 13 cm). As a consequence of the O_2 availability, NO_3^- and SO_4^{-2-} as less energetically 20 21 favourable electron acceptors for OM mineralization (Hedin et al. 1998)- were not consumed 22 for OM mineralization. In these conditions, we did not observe any significant production of solutes linked to anaerobic OM degradation in sediments (i.e. NH_4^+ , PO_4^{3-} and DOC) (Nogaro 23 24 et al. 2007). Hydraulic conductivity and discharge rates were 85% lower in sediment covered 25 by a 2-cm thick layer of fine sand, in comparison with systems topped by coarse sand. As a

consequence of the reduced hydraulic discharge rates, we observed sharp decreases in O₂ and 1 NO₃⁻ concentrations along depth in these systems. The rapid along-depth succession of 2 metabolic pathways was in accordance with predictable thermodynamic sequences (based on 3 free energy yields): O₂ being consumed first during OM mineralization in the oxic zone, 4 followed by the consumption of NO_3^- (denitrification), manganese and iron oxides, SO_4^{2-} and 5 6 carbon dioxide (Hedin et al. 1998, Baker et al. 2000, Kristensen 2000). Since O₂ was limiting in the first centimeters of sediments, most of the sedimentary matrix (and thus buried leaf 7 litter) was under anaerobic conditions, leading to the release of NH_4^+ , PO_4^{3-} and DOC. In such 8 O2-limited system, microbial abundances (total abundance of bacteria, abundance of active 9 10 bacteria and fungal biomass) and activities (glucosidase and leucine aminopeptidase 11 activities) were altered, and rates for microbial-mediated ecological processes occurring in the HZ were reduced (leaf litter decomposition was 30% lower than in not-stressed coarse-sand 12 13 treatment).

14 The contrasts in biogeochemical conditions between the two sedimentary contexts were 15 consistent with field studies showing that reduced hydrologic exchanges due to clogging 16 favoured the occurrence of anaerobic processes such as denitrification, sulphato-reduction and 17 methanogenesis (Dahm et al. 1987, Brunke and Gonser 1997, Boulton et al. 1998, Lefebvre et al. 2004). Our results are also in accordance with other studies showing that biogeochemical 18 19 conditions, in particular the availability of electron acceptors (mainly O_2 and NO_3^-), strongly 20 affect (i) the fungal colonization of leaves (Medeiros et al. 2009) and microbial enzymatic 21 activities such as cellulase and peptidase activities (Montuelle and Volat 1997) and (ii) OM 22 degradation rates (Chauvet 1988, Claret et al. 1998, Dahm et al. 1998, Lefebvre et al. 2005). It 23 is therefore clear that sediment contexts that lead to low hydrological exchanges and O_2 24 concentration in sediments limit the growth and activity of microbial communities and 25 ultimately the rates of microbial-mediated ecological process occurring in the HZ. Finally, we efficiently recreated the hydrological and biogeochemical functioning of 2 contrasted
 sediment contexts and highlighted the key role played by O₂ as resource for microorganisms
 developed in sedimentary habitats.

4

5 Modulation of bioturbator effects on biogeochemical processes by sediment context.

6 The present study confirmed our hypothesis that the effects of bioturbators on hydraulic7 conductivity and microbial-mediated processes depend on the sediment context.

8 As upward conveyors (feeding on sediment at depth and ejecting faecal pellets at the 9 sediment-water interface, Fisher et al. 1980, McCall and Fisher 1980), tubificid worms can 10 build networks of tubes and burrows that may extend as deep as 20 cm in sediments. 11 However, our results showed that both the bioturbation activity and the vertical distribution of 12 tubificid worms were modulated by the sedimentary characteristics. Using luminophore as 13 particle tracers, we noted that tubificid worms significantly reworked the top of the 14 sedimentary column with a fine sand layer whereas it was not the case with a coarse sand 15 layer. This contrast in bioturbation degree was linked to the vertical distribution of worms. 16 While worms used the whole sediment column in the systems topped with coarse sand, most 17 of tubificid worms were found in the top 0-5 cm in systems topped by a layer of fine sand. Fine sand probably acted as preferential feeding zone for T. tubifex (Juget 1979, Rodriguez et 18 19 al. 2001), which strongly influenced the vertical distribution of worms in the sedimentary 20 column. The different bioturbation activities exhibited by T. tubifex in the two sediment 21 contexts could explain their contrasting effects on hydraulic conductivity (see Fig.1 for the 22 successive effects of bioturbators). By producing galleries through the fine sand layer, T. 23 tubifex create water pathways that counteracted the adverse effect of fine sediment deposition 24 on water exchanges. The 6-fold increase in hydraulic conductivity due to T. tubifex in the

fine-sand treatment stimulated the exchanges of water and O₂ from surface to deep sediment 1 2 layers, restoring aerobic conditions in the sedimentary column (Fig. 1). Modification of aerobic-anaerobic conditions observed through the increase in O₂ and NO₃⁻ concentrations 3 was also associated with a lack of NH_4^+ , PO_4^{3-} and DOC accumulation in the sedimentary 4 5 habitat bioturbated by tubificid worms. The increase in electron acceptors (O_2 and NO_3) 6 availability for micro-organisms with T. tubifex has stimulated microbial communities 7 (abundances and activities) associated to the buried OM, leading to an increase by 30% of 8 OM breakdown rate in sedimentary systems covered by fine sand (Fig. 1). In contrast, 9 tubificid worms did not affect hydraulic conductivity and the subsequent chemical conditions 10 (availability of electron acceptors) in sediments topped by coarse sand. Consequently, 11 bioturbators did not influence microorganisms and microbial-mediated processes in 12 sediments.

13 Our study clearly demonstrated that the contribution of bioturbating invertebrates on 14 ecosystem processes was negatively correlated with the hydrologic exchanges occurring at the 15 water-sediment interface of the studied system, supporting conclusions from other studies 16 (Hakenkamp and Palmer 2000, Boulton et al. 2002). Bioturbators are able to strongly 17 influence water fluxes (through biological decolmation) in sedimentary habitats characterized 18 by low hydrologic exchanges (affected by the deposition of fine sediment particles), whereas 19 they only slightly modulate existing water fluxes in habitats with high hydrologic exchanges. 20 If the low influence of bioturbators on hydraulic conductivity in systems covered with coarse 21 sand could be linked to their low sediment reworking activity (Fig. 2), it could also resulted 22 from the reduced ability of bioturbation to increase hydrologic exchanges in a system that is 23 already highly permeable.

24

1 What underlies context dependency in the impact of engineers on other biota?

2 Most ecosystem engineering studies quantified the effects of engineers on some biotic 3 variables in a given habitat without taking into account the modulation of organism 4 engineering by environmental conditions (Crain & Bertness 2006). Our study clearly 5 demonstrates that the influences of bioturbators as engineers may vary across environmental 6 contexts. Few studies have already reported similar observations for various types of 7 ecosystems and various types of ecosystem engineers (e.g. Spooner and Vaughn 2006, 8 Nogaro et al 2009, de Moura Quierós A 2011) but they did not finely decrypt the complete 9 mechanisms by which ecosystem engineers and habitat characteristics interacted to shape 10 biological communities and/or ecosystem functions. More precisely, the environmental 11 context can influence the impacts of engineers on communities by modulating (i) the degree 12 with which its activity(ies) (or structure(s)) generates physical changes in the environment 13 and/or (ii) the degree with which physical changes can generate abiotic changes in the 14 environment (Fig. 1, see also Jones et al. 2010). As most of studies dealing with context 15 dependency examined only partially the engineering sequence, they could not provide 16 explanations about the mechanisms underlying context dependency. By examining the whole 17 engineering process as a detailed sequence of successive cause-effect relationships, our study allowed to conclude that the influence of bioturbators on microbial-mediated processes was 18 19 mainly mediated by the organism ability to change abiotic factors that are limiting for micro-20 organisms (i.e. O₂). This conclusion supports the general (and not tested before) idea that 21 ecosystem engineering is not an idiosyncratic process and that the magnitude of engineering 22 impacts on a biotic variable controlled by one or more limiting abiotic factors depends on the 23 degree to which the limiting abiotic factor(s) is/are modified relative to the unengineered state 24 (e.g., Gutiérrez et al. 2003, Gutiérrez and Jones 2006, Jones et al. 2010).

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- 25 Tables

Table 1. Specific area, total organic carbon (TOC), total nitrogen (TN) and total phosphorus
(TP) of the sediment used as top-sediment layer (mean ± SD, n= 4 for specific area, n=5 for
TOC, TN and TP).

		Specific area (cm ² g ⁻			
		1)	TOC (g kg ⁻¹)	TN (g kg⁻¹)	TP (mg kg ⁻¹)
	Coarse sand Fine sand	59.37 ± 0.34 1465.00 ± 33.17	0.97 ± 0.08 16.6 ± 1.6	0.17 ± 0.03 1.35 ± 0.06	3.76 ± 3.47 6.62 ± 1.36
5	i ine sanu	1403.00 ± 33.17	10.0 ± 1.0	1.55 ± 0.00	0.02 1 1.30
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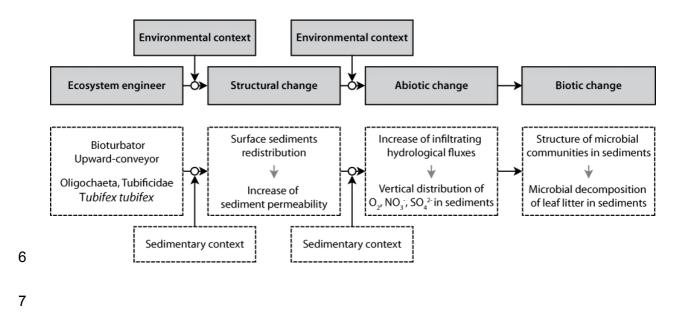
1	
2	Fig. 1. Successive steps for (i) a general case of ecosystem engineering sequence (in
3	grey) and (ii) the effects of <i>T. tubifex</i> on microorganisms in sedimentary habitats
4	(study case, in white). Note that feedbacks to the engineer and other relationships in the
5	ecosystem engineering can exist and are not represented here but can exist (for more
6	information, see Gutiérrez and Jones 2008, Jones et al. 2008).
7	
8	Fig. 2. Vertical distribution of <i>T. tubifex</i> in sediment and their associated effect on vertical
9	profiles of luminophores in sediment columns covered by fine sand or coarse sand.
10	
11	Fig. 3. Effect of <i>T. tubifex</i> on hydraulic conductivity measured every 10 days in sediment
12	columns covered by fine sand or coarse sand.
13	
14	Fig. 4. Effect of <i>T. tubifex</i> on depth profiles for O_2 , N-NO ₃ ⁻ , N-NH ₄ ⁺ , N-NO ₂ ⁻ , and DOC
15	concentrations determined at 4 depths (from H1: 2 cm above sediment interface, to H4: 13 cm
16	below sediment interface) after 36 days in sediment columns covered by fine sand or coarse
17	sand.
18	
19	Fig. 5. Effect of <i>T. tubifex</i> on leaf mass loss rate (A), and microbial characteristics measured
20	at the end of the experiment on leaves buried at 9 cm depth in sediment columns covered by

21 fine sand or coarse sand: total abundance of bacteria (B), abundance of active bacteria (C),

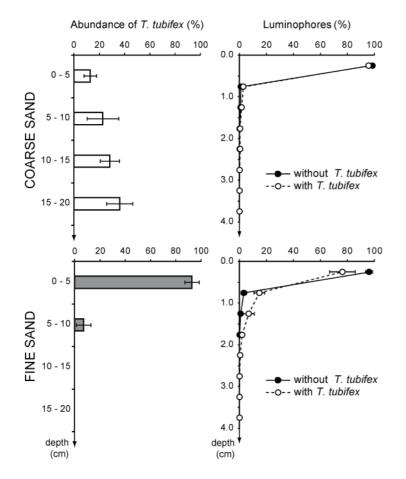
22 fungal biomass (D), potential respiratory activity (E), potential denitrification (F), xylosidase

23 (G), glucosidase (H), and leucine aminopeptidase activities (I).

- 1 2
- 3 Figures
- 4
- 5 Figure 1

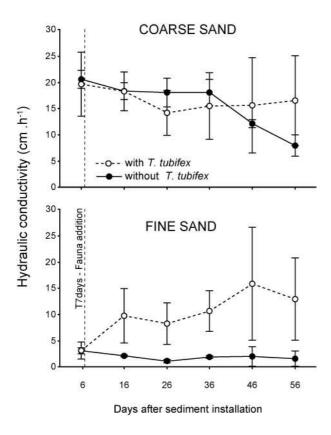


- 8
- 9 Figure 2

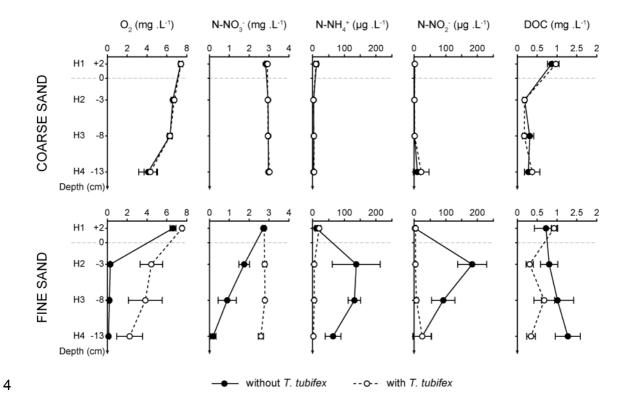








3 Figure 4





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