

The effect of sieve mesh size on the description of macroinvertebrate communities

Beatriz Barba, Aitor Larrañaga*, Ainhoa Otermin, Ana Basaguren and Jesús Pozo

Lab. Ecología de ríos, Dpto. Biología Vegetal y Ecología, Fac. Ciencia y Tecnología, Universidad del País Vasco/E. H. U., Apdo. 644. 48080 Bilbao.

* Corresponding author: aitor.larranagaa@ehu.es

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ABSTRACT

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Considerable time and effort is required to estimate the abundance and biomass of benthic macroinvertebrates, and often variable mesh size sieves are used to clean collected samples. We test whether the use of a mesh with a 1 mm pore size is adequate to obtain a valid description of a benthic macroinvertebrate community. Stream benthic surber samples were collected from 24 headwater streams. The densities, biotic indices and biological traits of macroinvertebrates retained in a 1 mm mesh ('> 1 mm' fraction) were compared to the same descriptors for the of macroinvertebrates retained in a 0.5 mm mesh sieve ('total'). We found that, if only the large fraction (>1 mm) is examined, the community descriptors are affected. Nevertheless, the observed changes were proportional and predictable for all of the variables describing invertebrate communities. Statistical differentiation of the tested metrics between sites was similar for both mesh sizes. Depending on the aim of the study (e.g., environmental impact assessments), the use of a 1 mm mesh sieve would be sufficient in describing macroinvertebrate communities.

Key words: Benthos, macroinvertebrates, processing, effort, mesh size.

RESUMEN

Efecto del tamaño de poro del tamiz en la descripción de las comunidades de macroinvertebrados

Para las estimas de abundancia y biomasa, el procesado de las muestras en laboratorio de invertebrados bentónicos requiere un tiempo y esfuerzo considerable y generalmente implica el uso de tamices de diferente luz de malla para lavar la muestra. Nuestro trabajo trata de comprobar si es suficiente para una descripción válida de la comunidad el uso de un tamiz de malla de 1 mm de poro en el procesado de muestras. Con tal propósito se recogieron muestras bentónicas de río en 24 tramos de cabecera y se compararon densidades, índices bióticos y rasgos biológicos de los invertebrados que eran retenidos en un tamiz de luz de malla de 1 mm (fracción '>1 mm') con los obtenidos usando una de 0.5 mm ('total'). Nuestro estudio revela que el análisis exclusivo de la malla gruesa afecta a los descriptores de la comunidad. Sin embargo, los cambios observados son proporcionales para todas las variables y se pueden predecir bien con ecuaciones lineales. Por otro lado, la diferenciación estadística entre estaciones es similar usando ambos tipos de malla, lo que en definitiva sugiere que, dependiendo del objetivo del estudio (por ejemplo, evaluación de impacto ambiental), el examen de la fauna retenida en un tamiz de 1 mm de poro puede ser suficiente para la descripción de las comunidades de macroinvertebrados.

Palabras clave: Bentos, macroinvertebrados, procesado, esfuerzo, tamaño del poro de malla.

INTRODUCTION

Benthic macroinvertebrates are the primary group of organisms used in stream water quality moni-

toring programs (Metcalf-Smith, 1996; Bonada *et al.*, 2006). Sorting benthic macroinvertebrates from sediments and organic matter, as well as counting and identifying them, takes consi-

derable time and effort (Ciborowski, 1991; Vlek *et al.*, 2006). While an exhaustive sorting of samples is the only way to assure a thorough description of the structure and composition of a macroinvertebrate community (Courtemanch, 1996; Cao *et al.*, 1998), techniques that optimize the cost-benefit for processing the samples have been developed (*e.g.* Barbour & Gerritsen, 1996; Vinson & Hawkins, 1996; Walsh, 1997; Metzeling & Miller, 2001; Vlek *et al.*, 2006).

The manner in which samples are collected and processed influences the description of the macroinvertebrate community being examined (Tanaka & Leite, 1998; Morin *et al.*, 2004; Boonsoong *et al.*, 2009); mesh size is a primary influencing factor (Battle *et al.*, 2007; Buss & Borges, 2008). A fine mesh gives a more precise estimate of the community but increases the effort needed for processing (Bartsch *et al.*, 1998). Thus a compromise between precision and effort must be made and can be achieved by using a larger mesh size.

Concerns about the imprecision of the estimates obtained using different mesh sizes would

be negligible if the variation of the community parameters was kept to a minimum, if the changes of these values were predictable or if statistical discrimination between sites was similar. Our objectives were to test the following questions: 1) whether mesh size (1 mm or 0.5 mm) affected macroinvertebrate community descriptors, such as density, taxonomic richness, biotic indices or other metrics based on biological traits; 2) whether there was a relationship between the metrics obtained by both meshes and 3) whether significant differences between sites remained consistent independent of the mesh size used.

METHODS

Study sites

Our study was conducted in 24 streams that drain over siliceous materials in tributaries of rivers in the Northern Iberian Peninsula (15 sites were in Agüera, 4 sites were in Barbadún, 3 sites we-

Table 1. Main characteristics of the study sites (mean \pm SE) during the three months before sampling ($N = 8$; DIN = dissolved inorganic nitrogen, SRP = soluble reactive phosphorus). *Características principales de las estaciones estudiadas (promedio \pm EE) en los tres meses previos a los muestreos ($n = 8$). DIN = nitrógeno inorgánico disuelto, SRP = fósforo reactivo soluble.*

Site	Stream order	Temp (°C)	pH	Alkalinity (meq/l)	Conductivity (μ S/cm)	Oxygen sat. (%)	DIN (μ g N/l)	SRP (μ g P/l)
1	2	9.7 \pm 0.6	7.1 \pm 0.1	0.48 \pm 0.04	106.0 \pm 4.6	96.8 \pm 0.6	804 \pm 58.1	13.2 \pm 3.7
2	1	10.1 \pm 0.6	7.3 \pm 0.1	0.80 \pm 0.09	140.7 \pm 8.5	97.7 \pm 1.8	1130 \pm 129.7	301.2 \pm 68.7
3	1	9.5 \pm 0.5	7.2 \pm 0.1	0.63 \pm 0.05	135.5 \pm 4.2	100.3 \pm 2.3	774 \pm 80.6	17.0 \pm 7.1
4	3	10.8 \pm 0.5	7.8 \pm 0.1	2.30 \pm 0.08	281.9 \pm 8.3	95.3 \pm 1.4	1217 \pm 46.4	37.2 \pm 8.8
5	1	10.4 \pm 0.3	7.4 \pm 0.1	1.18 \pm 0.07	183.0 \pm 6.0	95.4 \pm 0.4	262 \pm 61.8	14.5 \pm 7.9
6	1	10.8 \pm 0.4	7.3 \pm 0.1	1.15 \pm 0.05	191.3 \pm 4.9	95.8 \pm 1.6	515 \pm 42.8	15.7 \pm 2.6
7	1	10.9 \pm 0.5	7.3 \pm 0.1	0.91 \pm 0.03	133.55 \pm 7.9	97.6 \pm 0.5	538 \pm 48.3	5.0 \pm 2.1
8	1	11.4 \pm 0.5	7.2 \pm 0.1	0.97 \pm 0.04	143.4 \pm 5.82	80.8 \pm 0.9	818 \pm 61.8	13.7 \pm 3.2
9	2	9.9 \pm 0.4	7.1 \pm 0.1	0.41 \pm 0.02	101.4 \pm 3.1	100.2 \pm 1.9	558 \pm 48.4	3.7 \pm 1.6
10	3	10.7 \pm 0.3	7.4 \pm 0.1	0.72 \pm 0.04	134.7 \pm 5.8	98.6 \pm 0.7	894 \pm 58.2	19.7 \pm 6.5
11	2	9.8 \pm 0.6	7.2 \pm 0.1	0.50 \pm 0.04	84.5 \pm 7.6	105.4 \pm 4.7	505 \pm 51.4	9.4 \pm 2.4
12	2	10.0 \pm 0.6	7.2 \pm 0.1	0.46 \pm 0.04	82.8 \pm 6.9	105.3 \pm 3.4	480 \pm 40.7	8.6 \pm 5.6
13	1	10.9 \pm 0.6	6.4 \pm 0.2	0.16 \pm 0.02	67.6 \pm 3.4	100.0 \pm 1.5	1609 \pm 108.1	6.2 \pm 3.2
14	1	11.2 \pm 0.8	6.1 \pm 0.1	0.12 \pm 0.01	67.0 \pm 3.3	98.7 \pm 1.3	509 \pm 48.8	9.7 \pm 5.1
15	1	9.4 \pm 0.7	7.5 \pm 0.2	0.95 \pm 0.09	141.1 \pm 13.2	99.0 \pm 1.5	1169 \pm 48.6	3.4 \pm 2.1
16	1	11.4 \pm 1.2	6.6 \pm 0.1	0.44 \pm 0.18	96.2 \pm 19.2	98.5 \pm 1.7	609 \pm 71.6	8.5 \pm 4.8
17	1	9.2 \pm 0.7	7.0 \pm 0.1	0.27 \pm 0.02	93.3 \pm 4.4	98.1 \pm 1.3	962 \pm 53.9	6.6 \pm 2.3
18	1	10.3 \pm 0.7	7.1 \pm 0.1	0.75 \pm 0.08	214.2 \pm 17.9	99.8 \pm 1.7	828 \pm 123.7	7.53 \pm 3.5
19	1	10.6 \pm 0.8	6.8 \pm 0.1	0.23 \pm 0.02	64.8 \pm 4.8	100.0 \pm 1.6	317 \pm 33.8	4.4 \pm 1.89
20	1	10.7 \pm 0.9	7.1 \pm 0.1	0.43 \pm 0.04	119.8 \pm 14.8	101.2 \pm 1.2	753 \pm 94.5	8.1 \pm 2.4
21	2	11.8 \pm 0.4	6.5 \pm 0.1	1.88 \pm 0.12	82.8 \pm 2.6	99.1 \pm 0.6	1438 \pm 202.6	6.0 \pm 3.2
22	2	10.6 \pm 0.4	7.2 \pm 0.1	3.86 \pm 0.07	154.6 \pm 9.6	98.7 \pm 0.9	1591 \pm 121.6	102.1 \pm 14.9
23	1	12.6 \pm 0.4	7.8 \pm 0.1	1.84 \pm 0.25	491.9 \pm 7.2	96.3 \pm 0.9	1728 \pm 327.7	80.2 \pm 8.3
24	2	12.0 \pm 0.4	7.4 \pm 0.1	1.79 \pm 0.12	363.6 \pm 27.6	94.5 \pm 2.6	1751 \pm 197.2	53.7 \pm 11.4

re in Sámamo, and 2 sites were in Asón; see Table 1 for water physical-chemical characteristics). The climate is temperate, the annual precipitation rate is around 1600 mm, and the mean annual temperature is 14.3 °C. The primary land uses in this area are forestry, agriculture and stock breeding (Elosegi *et al.*, 2002).

Collecting and processing of benthic macroinvertebrate samples

In February and March 2003 (hereafter referred to as ‘sampling 1’) benthic samples were collected (Surber, 900 cm² area using a mesh with a 0.5 mm pore size) from sites 1 to 20 (five samples from randomly selected riffles) and were preserved in 70 % alcohol. Samples were sieved through a 1 mm mesh, which created two fractions: one captured in the 1 mm mesh sieve (referred to as the ‘>1 mm’ fraction) and one filtered through it. We refer to the sum of these two fractions as the ‘total sample’. Specimens were identified to genus (Plecoptera, Ephemeroptera and Crustacea), family, or class (Oligochaeta) following Tachet *et al.* (2002). In December 2003, a second benthic sample was taken (hereafter referred to as ‘sampling 2’) in sites 21 to 24 and samples were processed following the same methods outlined for ‘sampling 1’.

Data analysis

Indices

For ‘sampling 1’, we calculated eight indices for each data set (‘>1 mm fraction’ and ‘total sample’) and each site (one value per site was obtained from the lumped abundances of the five replica): taxonomic richness; Shannon diversity index (H) (Begon *et al.*, 1997); IBMWP (Iberian Biological Monitoring Working Party) (Alba-Tercedor & Pujante, 2000); IASPT (Iberian Average Score Per Taxon; calculated by dividing the IBMWP score by the number of families in each sample); EPT (Ephemeroptera, Plecoptera and Trichoptera) richness; percentage of shredders; percentage of other primary consumers and percentage of secondary consumers. While the me-

thodology for the IBMWP proposes the use of a kick net of 0.3 mm for sampling, we argue that a Surber net of 0.5 mm was sufficient in enabling us to calculate the IBMWP values; Torralba Burriel & Ocharan, (2007) found high similarities in their results obtained with the two types of nets, and we aim only to achieve internal comparability.

Biological and ecological traits

The analysis of biological and ecological traits focussed on shredder assemblages, a functional feeding group that reacts more clearly than others to environmental stressors in these highly organic matter dependant streams (Larrañaga, 2008; Larrañaga *et al.*, 2009). We analysed the following: maximum body size, life cycle length, number of reproduction cycles per year, dispersal rates, food, pH preference and trophic status preference. We ranked details about the traits as theoretical affinity scores for each modality within each trait at the genus level (Tachet *et al.*, 2002). Scores for each genus were multiplied by their density, and the resultant values for each modality for all genera were summed. Finally, relative scores for each modality were calculated for the shredder assemblage by dividing the total score in the modality by the total score for the corresponding trait.

Statistical analyses

We used three statistical approaches for our analysis. First, values of the indices obtained in the ‘>1 mm fraction’ were contrasted with those obtained in the ‘total sample’, using *t*-tests for paired data ($N = 20$). Second, regression analyses between the values calculated from the ‘>1 mm fraction’ and from the ‘total sample’ were performed for densities (n° m⁻²) of total invertebrates, feeding guilds and the most abundant taxa (mean relative density > 5 %), as well as for the relative scores of the trait modalities of shredders (untransformed data; $N = 99$; independent variable: ‘>1 mm fraction’). Third, we tested whether the between sites multiple comparisons remained unchanged with the ‘>1 mm fraction’ or the ‘total sample’ (hereafter referred to as interpretation analysis; data were log-transformed

Table 2. Comparison and regression analyses of the descriptors between the '>1 mm fraction' and 'total sample' ($N = 20$; $p < 0.001$ in all regressions). *Comparación y análisis de regresión de los descriptores entre '>1 mm' y 'total' (n = 20; p < 0,001 en todas las regresiones).*

Index	Comparison				Regression 'total' vs '> 1 mm'			
	'> 1 mm'	'total'	<i>t</i>	<i>p</i>	<i>b</i>	Cte	<i>R</i> ²	<i>F</i>
Taxonomic richness	27.2	32.4	10.46	< 0.001	0.860	9.047	0.72	46.9
Shannon diversity	2.4	2.3	1.48	ns	0.951	0.082	0.92	215.5
IBMWP	172.6	201.1	11.55	< 0.001	0.978	32.327	0.88	126.0
IASPT	6.3	6.2	3.25	0.004	0.665	1.964	0.65	33.3
EPT richness	16.5	19.1	1.97	ns	0.874	4.622	0.82	81.5
Shredder %	18.6	20.7	2.30	0.033	1.099	0.305	0.89	155.0
Other primary consumer %	75.5	74.4	1.03	ns	1.045	-4.437	0.88	135.8
Secondary consumer %	5.9	4.9	3.38	0.003	0.736	0.482	0.83	88.0

to approximate normality following Zar, 1999). We performed Tukey tests among the 20 sites (190 comparisons) for the '>1 mm fraction' and for the 'total sample' for the density and the trait related variables. We counted the number of pairwise comparisons for each variable that led to the same conclusion with both data sets (both sites being either statistically equal or statistically different); we considered these cases as 'equal' results. Alternatively, the number of comparisons that had significant differences between sites of the '>1 mm fraction' but not between sites of the 'total sample' were considered type I errors. Finally, we computed the number of comparisons between sites that were not significantly different considering the '>1 mm fraction' but different when considering the 'total sample' (type II errors).

'Sampling 2' was used to validate the relationships observed between the '>1 mm fraction'

and the 'total sample' obtained in 'Sampling 1'. Predictions of density in the 'total sample' for 'Sampling 2' were determined using regression equations constructed with the 'sampling 1' data set. The predicted values were thereafter compared with the 'total sample' observed values of 'sampling 2' by *t*-tests for paired data ($N = 20$). All statistical analyses were performed with SPSS 14.0 for Windows.

RESULTS

We found 67 taxa in the 'total sample', 61 of which also appeared in the '>1 mm fraction'. The only taxa that appeared exclusively in the 'total sample' (albeit in low densities) were Tabanidae (0.039 % of total abundance), *Asellus* (0.022 %), Molannidae (0.017 %), Psychomyi-

Table 3. Mean densities in the 'total sample' and the percentage of individuals lost when examining only the '>1 mm fraction'. The regression and interpretation analyses (see Methods) for the total invertebrates, the feeding guilds and the abundant taxa are included ($N = 99$; $p < 0.001$ in all regressions). *Densidades medias en el 'total' y porcentaje de individuos perdidos al estudiar sólo '>1 mm' junto con el análisis de regresión e interpretación (ver Methods) de la densidad total de invertebrados, de los grupos de alimentación y de los taxones abundantes (n = 99; p < 0,001 en todas las regresiones).*

Variable	Mean density	% lost in	Regression				Interpretation		
	(n° m ⁻²)	'>1 mm'	<i>b</i>	cte	<i>R</i> ²	<i>F</i>	Equal	Error I	Error II
Total invertebrates	4028.4	50.4	1.852	329.8	0.87	638.5	188	2	0
Shredders	731.9	56.7	1.894	131.9	0.78	354.1	182	8	0
<i>Echinogammarus</i>	411.1	55.8	1.748	90.8	0.78	346.7	174	8	8
Other primary consumers	3140.9	49.5	1.892	139.9	0.88	693.5	178	9	3
<i>Baetis</i>	678.9	55.1	1.727	146.5	0.87	648.8	186	4	0
Elmidae	254.6	59.2	2.078	36.6	0.91	1022.3	179	2	9
Oligochaeta	324.3	44.0	1.459	47.3	0.83	219.9	178	1	11
Simuliidae	354.9	35.2	1.258	62.9	0.95	1694.0	181	5	4
Chironomidae	1000.3	55.4	1.696	202.4	0.85	559.1	174	4	12
Secondary consumers	155.5	39.5	1.276	35.5	0.64	172.0	180	0	10

Table 4. Regression and interpretation analysis (see Methods) of the different modalities of the biological and ecological traits for the shredder assemblages ($N = 99$; $p < 0.001$ in all analyses). *Análisis de regresión e interpretación (ver Methods) de las diferentes modalidades de los rasgos biológicos y ecológicos del grupo de los fragmentadores* ($n = 99$; $p < 0,001$ en todos los casos).

Trait	Modality	Regression				Interpretation		
		<i>b</i>	Cte	R^2	<i>F</i>	Equal	Error I	Error II
Maximum body size	< 2.5 mm	0.930	1.48	0.50	96.8	168	15	7
	2.5-5 mm	0.881	3.89	0.53	111.4	164	18	8
	5-10 mm	0.875	3.69	0.91	934.2	168	9	13
	10-20 mm	0.760	1.79	0.85	563.3	168	3	19
	20-40 mm	0.883	2.91	0.93	1253.3	165	8	17
Life cycle length	≤ year	0.818	5.60	0.90	843.1	166	6	18
	> 1 year	0.818	12.63	0.90	843.1	169	5	16
Reproduction cycles per year	< 1	0.854	1.47	0.79	375.6	171	0	19
	1	0.862	7.88	0.91	1006.5	168	4	18
	> 1	0.892	3.26	0.93	1324.1	176	2	12
Dispersion	Aquatic passive	0.887	4.73	0.91	951.5	172	12	6
	Aquatic active	0.821	4.88	0.84	500.2	168	13	9
	Aerial passive	0.521	0.03	0.81	406.1	186	3	1
	Aerial active	0.913	3.66	0.78	340.3	171	9	10
Food	Fine sedim. + microorg.	0.709	0.00	0.76	309.3	190	0	0
	FPOM	0.819	2.43	0.83	476.3	167	3	20
	CPOM	0.502	15.17	0.32	44.8	169	3	18
	Microphytes	0.963	2.20	0.70	227.9	169	3	18
	Macrophytes	0.698	2.95	0.62	155.0	154	2	34
	Dead animals > 1 mm	0.736	2.99	0.77	332.3	175	5	10
	Macroinvertebrates	0.910	0.67	0.91	931.1	166	10	14
pH preference	< 4	0.841	1.26	0.84	513.8	181	1	8
	4-4.5	0.850	1.40	0.88	728.6	177	6	7
	4.5-5	0.587	7.16	0.51	99.8	178	7	5
	5-5.5	0.819	3.66	0.87	652.3	171	4	15
	5.5-6	0.871	2.60	0.88	724.3	173	10	7
	> 6	0.865	2.99	0.90	884.5	173	2	15
Trophic status preference	Oligotrophic	0.630	19.7	0.60	147.4	165	15	10
	Mesotrophic	0.680	14.7	0.67	200.5	152	18	20
	Eutrophic	0.767	0.69	0.55	120.2	172	18	0

dae (0.017 %), Psephenidae (0.005 %) and Valvatidae (0.005 %). Mesh size did not significantly affect the values obtained for the Shannon diversity index or for the EPT richness index. Taxa richness and the IBMWP index, however, were significantly higher and the IASPT was significantly lower for the ‘total sample’ than for the ‘> 1 mm fraction’ (Table 2). Considering IBMWP quality classes, 5 of the 20 sites (sites 4, 8, 9, 13 and 14) changed quality classes with mesh size.; they were classified as ‘non-contaminated’ or ‘non-significantly altered’ (IBMWP: 100-150) with values from the ‘>1 mm fraction’ and they were changed to the clean wa-

ters category (IBMWP > 150) when the ‘total sample’ was examined. Relative contribution of shredders was significantly lower in the ‘>1 mm fraction’ than in the ‘total sample’; for secondary consumers relative contribution was higher in the ‘>1 mm fraction’ and other primary consumers did not differ between fractions (Table 2). Regression analyses between the ‘>1 mm fraction’ and the ‘total sample’ were significant for all the indices analysed (Table 2).

When only the ‘>1 mm fraction’ was processed, 50.4 % of the 35 893 individuals collected in ‘sampling 1’ were lost (Table 3). Among the most abundant taxa, Elmidae suf-

Table 5. Comparison of the observed and estimated densities (mean \pm SE) for the total invertebrates, the feeding guilds and the abundant taxa for the ‘sampling 2’ sites ($N = 20$). *Comparación de las densidades observadas y esperadas del total de invertebrados, de los grupos de alimentación y de los taxones abundantes para las estaciones del segundo muestreo (‘sampling 2’) (n = 20).*

Variable	Average density (n ^o m ⁻² \pm SE)		Comparison	
	Observed	Estimated	<i>t</i>	<i>p</i>
Total invertebrates	6792.6 \pm 1428.8	6685.4 \pm 1523.7	0.77	ns
Shredders	632.6 \pm 151.9	636.6 \pm 151.6	1.82	ns
<i>Echinogammarus</i>	100.7 \pm 32.5	57.5 \pm 27.7	1.32	ns
Other primary consumers	5957.8 \pm 1419.0	5921.0 \pm 1456.6	1.29	ns
<i>Baetis</i>	211.1 \pm 36.9	116.0 \pm 18.0	2.29	0.03
Elmidae	278.5 \pm 52.5	325.0 \pm 71.3	1.51	ns
Oligochaeta	507.4 \pm 85.8	734.0 \pm 115.8	2.79	0.01
Simuliidae	75.5 \pm 12.6	74.6 \pm 11.2	0.12	ns
Chironomidae	950.4 \pm 117.2	882.7 \pm 115.0	0.89	ns
Secondary consumers	202.2 \pm 28.1	127.8 \pm 20.6	4.22	< 0.001

ferred the highest losses (59.2 %) and Simuliidae suffered the fewest losses (35.2 %) when examining just the ‘>1 mm fraction’ (Table 3). Regression analyses between the abundance of the feeding groups and the most abundant taxa in the ‘>1 mm fraction’ and the ‘total sample’ were significant ($p < 0.001$, Table 3). Abundance of Simuliidae, Elmidae and ‘other primary consumers’ had the best fit to the data, whereas ‘secondary consumers’ abundance had the worst fit (Table 3). Among the 190 possible comparisons (of abundances) between the 20 sites, the ‘equal’ cases (those that reach the same statistical conclusion studying both fractions) ranged from 174 (*Echinogammarus*) to 188, depending on the taxon or group analysed (average = 180; 94.7 % of the possible pairwise comparisons; Table 3). Type I errors (differences in the ‘>1 mm fraction’ and not in the ‘total sample’) ranged from 0 to 9 cases, whereas type II errors ranged from 0 to 12 cases (Table 3).

All linear regressions for the relative affinity scores of biological traits between the ‘>1 mm fraction’ and the ‘total sample’ were significant ($p < 0.001$; R^2 range: 0.32-0.91; Table 4). Of the 190 Tukey tests performed, between 152 and 181 cases (depending on the trait and the modality analysed) found ‘equal’ results with both mesh sizes (average = 169.9; 89.4 % of cases; Table 4). Type I errors ranged from 0 to 18 while type II errors ranged from 0 to 34 (Table 4). Validation of the model created with ‘sampling 1’

data by means of the comparison of the observed and the expected values for the ‘sampling 2’ was considered positive; 7 of the 10 variables examined resulted in no significant differences (Table 5). Only densities of *Baetis* and ‘secondary consumers’ were underestimated using the ‘sampling 1’ regression models; the abundance of Oligochaeta was overestimated by the regression model (Table 5).

DISCUSSION

The analysis of a fraction of the sample, here shown using a fraction trapped in a mesh pore size of 1 mm, can affect descriptors of macroinvertebrate communities (Tanaka & Leite, 1998; Morin *et al.*, 2004; Rodrigues *et al.*, 2007). In our results, only the Shannon diversity index, the EPT richness and the percentage of primary consumers resulted in similar findings for the two methods we used. We observed that a few taxa (5.2 on average) were only found in the ‘total sample’ we analysed. The absence of taxa in our study cannot be explained by the body length, however, as maximal body lengths for these taxa range from 2.5 to 40 mm (Tachet *et al.*, 2002) and individuals could have been retained in the 1 mm mesh, even at the early stages of their development (see Morin *et al.*, 2004). Nevertheless, the loss of taxa is quite relevant when we consider descriptors such as taxonomic richness and in-

dices, such as the IBMWP and the IASPT; the addition of new taxa substantially changes the final result. Concerning the IBMWP index, 25 % of the sites we analysed were classified in a higher quality class when examining the 'total sample'. Our findings suggest that caution should be taken when comparing studies that use different mesh sizes to measure macroinvertebrate community composition.

For the variables we analysed, spatial-temporal comparisons within a given study could be achieved for invertebrates trapped in larger mesh sizes. In our study, changes in the descriptors from the '>1 mm fraction' mesh to the 'total sample' were predictable by linear regression in all cases analysed. Additionally, these linear relationships were relatively constant for most of the taxa or groups we considered, at least for the temporal and spatial range that comprised our samples. Observed statistical discrimination among sites was also very similar considering both fractions. In the Tukey tests, the mean agreement between the between-site significant differences between the '>1 mm fraction' and the 'total sample' was nearly 95 %. As other studies have concluded (James *et al.*, 1995; Morin *et al.*, 2004; Gruenert *et al.*, 2007) a reliable spatial discrimination between sites can be obtained using a >1 mm size fraction.

There are other reasons why a coarse mesh might be used. First, while we have measured density to portray the importance of invertebrates, biomass can also be measured. A more precise estimation can be achieved when trying to solve ecological questions within an energetic perspective. The mesh size used produces a lower bias in biomass, as smaller individuals are the lightest, and their removal from the replicate causes little variation in the final mass estimates (Gage *et al.*, 2002; Morin *et al.*, 2004). Second, the traditional importance given to the structural descriptions of the community (Wright *et al.*, 2000) is giving way to the functional perspective (see Young *et al.*, 2008) that may use structural descriptors as complementary. Within this new approach, estimates based on coarser mesh sizes may be useful and may save time in studies of stream metabolism (Acuña *et al.*, 2005) and of leaf litter processing (Pozo *et al.*, 1998) data.

Third, in a highly complex system, such as river benthos where the number of replicates taken per site from the substrate is highly limited by the effort required to process them, the use of a coarser mesh could allow for the examination of a higher number of replicates (James *et al.*, 1995, Morin *et al.*, 2004). This measure could increase the precision, the degrees of freedom and the statistical power of stream macroinvertebrate community analyses (Bartsch *et al.*, 1998; Vlek *et al.*, 2006).

Values of structural descriptors for benthic macroinvertebrates change with mesh size but are predictable as they maintain proportionality among subsamples obtained by different mesh sizes. This linear relationship between the '>1 mm fraction' and the 'total sample' makes internal comparisons possible; we had accurate conclusions on average in 95 % of the cases we examined. The spatial and temporal extent to which this rule applies should be tested in more distanced areas.

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