COMPARATIVE DIVERSITY OF MOLLUSCAN FAUNA OF DIFFERENT AQUATIC HABITAT TYPES: THE LIWIEC RIVER CATCHMENT (EAST POLAND)

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ABSTRACT: Investigations of aquatic mollusc assemblages were conducted within a semi-natural catchment of a medium-sized lowland river Liwiec in five habitat types: the main Liwiec River channel, its secondary channels and six tributaries, as well as in natural ponds and man-made ditches within the river floodplains. To assess the contribution of each habitat type to the diversity, I used the species richness, diversity, rarity (species rarity index, SRI) and abundance, as well as the composition and structure of mollusc assemblages and compared them among the habitat types. The spatial pattern of mollusc diversity was analysed based on hierarchical partitioning. Over five years, 54 mollusc species were found, including three listed on the IUCN Red List or in Annexes II and IV of the EU Habitats Directive (Unio crassus Philipsson, Sphaerium rivicola (Lamarck) and Anisus vorticulus (Troschel)). The mollusc metrics in most cases did not differ significantly among the habitat types. All the habitats contributed relatively evenly to the mollusc diversity. The diversity at the site level was generated mainly by alpha component, whereas at the landscape scale habitat heterogeneity (beta component, i.e. $\beta$) was very important. In order to maintain mollusc diversity, conservation efforts ought to focus specifically on the most heterogeneous fragments of the Liwiec River catchment.

KEY WORDS: aquatic molluscs, threatened species, catchment, drainage ditches, floodplain, river

INTRODUCTION

The river catchment includes various types of lotic and lentic aquatic habitats, which form a complex system. However, macroinvertebrates of different habitat types have not been equally surveyed. Studies on riverine macroinvertebrate diversity focused primarily on main river channels, nonetheless, some of them did include floodplains and diverse aquatic habitats occurring there (e.g. GODREAU et al. 1999, TOCKNER et al. 1999, WARD et al. 1999). Studies conducted in streams were relatively numerous (e.g. LAKE 2000, HEINO et al. 2005, CLARKE et al. 2008 and references therein). Although small water bodies, including natural and man-made ponds and ditches, are very numerous their macroinvertebrate fauna remains under-investigated (e.g. PAINTER 1999, WILLIAMS et al. 2004, BIGGS et al. 2005, GALLARDO et al. 2008, 2009, 2014, VERDONSCHOT et al. 2011). Studies comparing macroinvertebrate assemblages in different water body types are relatively few (e.g. VERDONSCHOT 1990, WILLIAMS et al. 2004, BIGGS et al. 2007, 2017, DAVIES et al. 2008, VERDONSCHOT et al. 2011), however they still reveal that small water bodies contribute significantly to aquatic biodiversity across landscapes. Ditches (being man-made water bodies) appear important for regional stream biodiversity (SIMON & TRAVIS 2011) and display a considerable contribution to total aquatic biodiversity of the floodplain (ARMITAGE et al. 2003).

Aquatic molluscs are an important component of macroinvertebrate fauna in many riverine and riparian habitats (e.g. JURKIEWICZ-KARNKOWSKA 2015 and references therein). Species-rich malacocoenoses are found in different aquatic habitat types within river-floodplain systems (e.g. PIECHOCKI 1969, OBRLIJK et
Molluscs of the main Liwiec River channel were studied by Korycińska (2002) and Jurkiewicz-Karnkowska (2016). Comparative study of aquatic mollusc diversity within a short (about 10 km long) fragment of the Liwiec River and its floodplain included a stretch of the main channel, a secondary channel and a few remnants of the former river channel (Jurkiewicz-Karnkowska 2015). So far, ditches have not been included in the investigations within the Liwiec River floodplain.

The biodiversity characteristics of different water body types within catchments and their contribution to total diversity are important both for sustainable catchment management and for nature conservation. Such information is scanty, particularly for semi-natural catchments. The understanding of the biodiversity’s spatial pattern, including the contribution of alpha and beta diversity to the total diversity and determining which spatial scales most influence the diversity to the greatest extent, are crucial to conservation planning.

The main objectives of the present study were to assess the contribution of different aquatic habitat types to mollusc diversity at the landscape scale as exemplified by a semi-natural medium-sized lowland river catchment, as well as to compare conservation values of mollusc assemblages harboured by individual habitat types. The comparison of species richness, diversity, rarity and abundance, as well as composition and structure of mollusc assemblages among five habitat types: the main Liwiec River channel (ML), its secondary channels (SL) and six tributaries (T), natural ponds (P) and man-made ditches (D) within river floodplains was performed. Special attention was paid to the biodiversity and conservation value of ditches, representing the only man-made habitat type within the study area. Diversity partitioning within the individual habitat types and the entire study area was applied to better understand the spatial pattern of mollusc diversity and the contribution of alpha and beta diversity to the total diversity at different spatial scales.

MATERIAL AND METHODS

STUDY AREA

The study area comprised of the Liwiec River and its valley, as well as six of its tributaries (Stara Rzeka, Helena, Muchawka, Kostrzyń, Miedzanka and Osownica) and their valleys. Liwiec is the biggest left-bank tributary of the Bug River. It is a medium-sized river (ca. 142 km long), with a catchment area of 2,780 km². The mean long-term discharge (SSQ) recorded at a water gauge station in the lower course (17 km of the river course counting from the mouth) was 10.2 m³ s⁻¹ (Czarnecka 2005). The source area of the river is located at 160 m a.s.l. (52°36′24″N, 21°33′34″E) and the mouth at 85 m a.s.l. (52°05′39″N, 22°37′39″E), the mean river gradient is 0.52‰.

The valley is 120 km² in area, its width not exceeding 2 km, except at two short sections where it widens to over 5 km (within the upper and lower river sections). It has retained its natural character. The land is extensively used, mainly as meadows and pastures, with forests occupying a relatively small area.
The Liwiec River valley is covered by the Natura 2000 network (2 Special Areas of Conservation (SACs): PLB140002 and PLH140032) and partially by the Siedlce-Węgrów Landscape Protection Area. The river channel has been regulated in its upper section, the middle and lower sections have preserved a relatively natural character, however small hydro-technical constructions (weirs, culverts) are present nearly all along the river except at the mouth section, which has remained exceptionally natural. The river is fed by 10 tributaries, among which Muchawka, Kostrzyń and Osownica are the largest. Substantial fragments of the tributaries have been regulated, but their valleys have retained a relatively natural character. A considerable part of the Kostrzyń River valley is covered by the Natura 2000 network (three SACs:PLB140009 and PLH140036 and a fragment of PLH140032), two nature reserves and partially by the Mińsk Landscape Protection Area. The marshy valley of the Muchawka River within the boundaries of the city of Siedlce is a protected area (“Dolina Muchawki”).

The study included 139 sites (Fig. 1): 41 located in the main channel of the Liwiec River (ML1–L41), 11 in its secondary channels (SL1–SL11), 32 within its tributaries (T): Stara Rzeka (S1–S6), Helenka (H1, H2), Muchawka (Mu1–Mu7), Kostrzyń (K1–K5), Miedzanka (M1–M6) and Osownica (O1–O6), 34 in floodplain natural ponds, i.e. lentic water bodies of fluvial origin (P1–P34) and 21 in ditches (i.e. man-made channels) located within river floodplains (D1–D21). At each sampling site within the rivers and other water bodies the width was measured directly in the field with metre tape, the depth was assessed with calibrated pole. Current velocity was measured using a float and stopwatch. The width at the ML sites ranged from <5 to >15 m, in SL and T it ranged from 1 to 10 m, in P from 1 to >15 m and

Fig. 1. Study area and location of the investigated sites; ML – main Liwiec River channel, T – tributaries, SL – secondary channels of Liwiec, P – ponds, D – ditches; grey rectangle – location of the study area on the map of Poland
in D from <1 to 5m. The depth ranged from <0.2 to >1 m, with the lowest values in D, as well as in some of T and P sites. Current velocity at ML and T sites ranged from <0.1 to >0.25 m/s (depending on the slope), whereas the other sites were mostly stagnant and only a few of them were characterised by a slight water movement. Bottom sediments in the rivers were sandy or sandy-muddy, often with the admixture of detritus or gravel (depending on current velocity). In the other habitat types bottom sediments were generally muddy or sandy-muddy, mostly with a big admixture of detritus. ML and T sites were characterised by sparse or moderate abundance of macrophytes or lack of vegetation. Within the other habitat types macrophyte abundance was mostly moderate or high (especially in P). Canopy was lacking or it was poor, so most of the sites were open and others were relatively slightly shaded.

MOLLUSC SAMPLING

Molluscs were sampled during summer (July–mid-September) 2012–2016, as well as in late spring (mid-May–the beginning of June) in 2013, 2015 and 2016 using a hand net with a working side of 25 cm, mesh size of 0.5 mm and handle length of 2 m.

Individual habitats were investigated during 1–2 sampling events and every time usually 1–2 samples were taken, each of ca. 1 m² of the bottom area, covering all visually detected microhabitats. Samples from rivers were collected in the current and near the banks (never deeper than 1.5 m), ponds were investigated within the zone from the margin to the depth of 1.5 m, secondary channels of the Liwic River and ditches were surveyed mostly in the whole cross sections. In total, 270 samples were collected. Samples were washed directly in the field using a sieve of 0.5 mm mesh size and they were preserved in laboratory with 75% ethyl alcohol (except for the majority of Unionidae, which were identified in the field and returned to the water). In the laboratory the molluscs were sorted, counted and identified using the keys of PIECHOCKI (1979) and PIECHOCKI & DYDUCH-FALNIOWSKA (1993). Species names were updated according to PIECHOCKI & WAWRZYNIAK-WYDROWSKA (2016).

DATA ANALYSIS

Sampling bias was tested with a species accumulation curve and the abundance-based non-parametric estimators Chao1, Chao2 and ACE (COLWELL 2004). To compare species richness of the five habitat types, randomization was performed for equal numbers of sites treated as samples in each of them (n=11; the number of samples was constrained by the minimum sample size present in SL).

Dominance patterns of molluscs were estimated as the percentage of individual species within the total mollusc abundance (GÖRNY & GRÜM 1981). The frequency of individual species occurrence (i.e. the ratio of sites occupied, %F) within each of the habitat types was calculated (GÖRNY & GRÜM 1981).

Species were regarded as regionally uncommon when the frequency of their occurrence was lower than 5%. This value was accepted as a halfway between diverse values proposed in the literature (e.g. VERDONSCHOT et al. 2011, HEEGAARD et al. 2013).

The Shannon index H' (i.e. entropy) and Shannon true diversity (expH') (JOST 2006), where H' = -\sum(p_i \ln p_i)) were calculated based on mollusc abundance data. All these calculations were carried out with the EstimateS, v.9.0 software (COLWELL 2004).

Comparison of species rarity within and between the habitat types was conducted using a species rarity index (SRI), conceptually based on the Species Quality Score developed by FOSTER et al. (1989). To calculate SRI all species present were given a rarity category: common species (score 1), local species (i.e. confined to limited areas or widespread, but represented by few individuals, score 2), nationally scarce (based on PIECHOCKI & WAWRZYNIAK-WYDROWSKA 2016, score 4), included in Annexes II or IV of the EU Habitats Directive or legally protected in Poland (8; Council Directive 1992, Dz. U. 2016), included on the IUCN Red List of Threatened Species (IUCN 2018) – near threatened or vulnerable (16), endangered (32). Subsequently, the scores of all species at each site were summed up to obtain the Species Rarity Score, which in turn was divided by the number of species recorded at the site to give the SRI.

Following Whittaker’s terminology (WHITTAKER 1972), total species diversity in a set of communities may be partitioned into additive components within and among communities: \( y = \beta + \alpha \) (LANDE 1996, VEECH et al. 2002). In contrast to the classical, multiplicative approach (WHITTAKER 1972) additive partition is more consistent and can be applied to multiple spatial scales. According to JOST (2007) when the sizes of samples or assemblages are unequal, only Shannon diversity measures (diversity indices of order one) can be decomposed into meaningful independent alpha and beta components. Species richness (diversity index of order 0) weights each assemblage equally, regardless of its true weight, and for this reason it is not a satisfactory measure, particularly when weights are important. Additive partitioning of Shannon entropy (H') was used to study the hierarchical partitioning of diversity, because H' is the only diversity index which can be additively partitioned into independent alpha and beta components. The relative contribution of the within and among sample (\( \alpha_i \) and \( \beta_i \)), among site (\( \beta_s \)) and among habitat (\( \beta_h \)) diversity to total diversity (\( y \)) of the study
area was assessed. To enable proper interpretation of the results of diversity partitioning the obtained Shannon alpha, beta and gamma entropies were converted to true diversities (their number equivalents, i.e. exponentials). The results of multiplicative decomposition of true Shannon diversity and species richness were compared with the results of additive partitioning of Shannon entropy. MacArthur’s homogeneity measure (MacArthur 1965), $\exp(H_{\alpha})/\exp(H_{\gamma})$, was calculated to show the proportion of landscape diversity contained in the average sample.

Beta diversity was used as the measure of dissimilarity among sites to avoid strong bias in Jaccard’s index, which may result from small sizes of mollusc assemblages in some cases.

RESULTS

COMPOSITION AND STRUCTURE OF MOLLUSC ASSEMBLAGES

In total, 54 mollusc species were recorded during the study (Table 1). The highest number of species (43) was found in the main channel of the Liwiec River (ML), followed by the tributaries (T, 41 species) and ditches (D, 37 species); the lowest species richness was recorded in secondary channels of the Liwiec River (SL) and ponds (P) – 35 species in each.

Table 1. Frequency distribution (%F) of mollusc species in five habitat types within the Liwiec River catchment: main river (ML), tributaries (T), secondary channels (SL), floodplain ponds (P) and ditches (D); rarity categories of species are given in parentheses; * – alien species

| Species                  | ML | T  | SL | P  | D  |
|--------------------------|----|----|----|----|----|
| Gastropoda               |    |    |    |    |    |
| Viviparidae              |    |    |    |    |    |
| Viviparidae              |    |    |    |    |    |
| Bithynia tentaculata     | (1)| 3  | 2  | 16 | 20 |
| Valvatidae               | (1)| 1  | 0  | 0  | 0  |
| Bithynia tentaculata     | (1)| 31 | 48 | 72 | 34 |
| Valvatidae               | (1)| 7  | 2  | 24 | 16 |
| V. macrostoma Mörch, 1864| (2)| 2  | 0  | 4  | 2  |
| V. piscinalis O. F. Müller, 1774 | (1)| 12 | 8  | 16 | 4  |
| V. piscinalis O. F. Müller, 1774 | (1)| 12 | 8  | 16 | 4  |
| V. piscinalis O. F. Müller, 1774 | (1)| 12 | 8  | 16 | 4  |
| Lyneidae                 |    |    |    |    |    |
| Acroloxus lacustris      | (1)| 1  | 2  | 0  | 0  |
| Stagnicola corvus        | (1)| 0  | 1  | 0  | 0  |
| Stagnicola corvus        | (1)| 0  | 1  | 0  | 0  |
| Stagnicola corvus        | (1)| 0  | 1  | 0  | 0  |
| Stagnicola corvus        | (1)| 0  | 1  | 0  | 0  |
| Physidae                 |    |    |    |    |    |
| Aplexa hyporum (Linnaeus, 1758) | (1)| 1  | 2  | 0  | 16 |
| Physella acuta Draparnaud, 1805* | (2)| 1  | 2  | 0  | 0  |
| Physa fontinalis (Linnaeus, 1758) | (1)| 5  | 5  | 25 | 36 |
Table 1. continued

| Species                      | ML | T | SL | P | D |
|------------------------------|----|---|----|---|---|
| **Gastropoda**               |    |   |    |   |   |
| **Planorbidae**              |    |   |    |   |   |
| *Ancylus fluviatilis*        | (1)| 0 | 2  | 0 | 0 |
| O. F. Müller, 1774           |    |   |    |   |   |
| *Anisus calcifornis*         | (2)| 3 | 2  | 0 | 14|
| (Sandberger, 1875)           |    |   |    |   |   |
| *A. leucostoma*              | (1)| 5 | 0  | 4 | 10|
| (Millet, 1813)               |    |   |    |   |   |
| *A. spirorbis*               | (1)| 1 | 0  | 0 | 4 |
| Linnaeus, 1758               |    |   |    |   |   |
| *A. vortex*                  | (1)| 6 | 17 | 56| 54|
| Linnaeus, 1758               |    |   |    |   |   |
| *A. vorticulus*              | (8)| 0 | 0  | 24| 20|
| Troschel, 1834               |    |   |    |   |   |
| *Bathyomphalus contortus*    | (1)| 0 | 13 | 20| 26|
| Linnaeus, 1758               |    |   |    |   |   |
| *Gyraulus albus*             | (1)| 9 | 19 | 12| 4 |
| O. F. Müller, 1774           |    |   |    |   |   |
| *G. crista*                  | (1)| 3 | 0  | 4 | 2 |
| Linnaeus, 1758               |    |   |    |   |   |
| *G. laevis*                  | (4)| 0 | 0  | 0 | 2 |
| Alder, 1838                  |    |   |    |   |   |
| *G. rossmaessleri*           | (2)| 5 | 4  | 0 | 6 |
| (Auerswald, 1852)            |    |   |    |   |   |
| *Hippus complanatus*         | (1)| 1 | 4  | 8 | 10|
| Linnaeus, 1758               |    |   |    |   |   |
| *Planorbarius corneus*       | (1)| 5 | 17 | 64| 64|
| Linnaeus, 1758               |    |   |    |   |   |
| **Sphaeriidae**              |    |   |    |   |   |
| *Planorbis planorbis*        | (1)| 5 | 6  | 4 | 14|
| Linnaeus, 1758               |    |   |    |   |   |
| **Bivalvia**                 |    |   |    |   |   |
| **Unionidae**                |    |   |    |   |   |
| *Anodonta anatina*           | (1)| 11| 0  | 4 | 0 |
| Linnaeus, 1758               |    |   |    |   |   |
| *A. cygnea*                  | (2)| 1 | 4  | 0 | 0 |
| Linnaeus, 1758               |    |   |    |   |   |
| *Unio crassus*               | (32)| 7 | 0  | 0 | 0 |
| Philipsson, 1788             |    |   |    |   |   |
| **Unio pictorum**            | (1)| 19| 8  | 12| 0 |
| Linnaeus, 1758               |    |   |    |   |   |
| **Unio tumidus**             | (1)| 15| 12 | 4 | 0 |
| Philipsson, 1788             |    |   |    |   |   |
| **Sphaerium corneum**        | (1)| 53| 60 | 52| 24|
| Linnaeus, 1758               |    |   |    |   |   |
| **S. rivicola**              | (16)| 7 | 0  | 12| 0 |
| (Lamarck, 1818)              |    |   |    |   |   |
| Number of species            |    | 43| 41 | 35| 35|
| Total                        |    | 54| 54 | 54| 54|

Table 2. Comparison of the numbers of species found within five habitat types and the entire study area with the expected species richness calculated with non-parametric abundance-based estimators ACE, Chao1 and Chao2; the values in parentheses show percent of estimated richness comprised by a number of species found.

| Habitat type          | Number of species found | ACE       | Chao1     | Chao2     |
|-----------------------|-------------------------|-----------|-----------|-----------|
| The Liwiec River (ML) | 43                      | 46.13 (93.2%) | 46.00 (93.5%) | 50.02 (85.7%) |
| Tributaries (T)       | 41                      | 46.48 (88.2%) | 42.67 (96.1%) | 48.61 (84.3%) |
| Secondary channels (SL)| 35                     | 36.83 (95.0%) | 35.00 (100%)  | 35.73 (98.0%) |
| Ponds (P)             | 35                      | 35.5 (98.6%)  | 35.00 (100%)  | 35.16 (99.5%) |
| Ditches (D)           | 37                      | 38.46 (96.2%) | 38.00 (97.4%) | 42.33 (87.4%) |
| **Total**             | **54**                  | **54.75 (98.6%)** | **54.25 (99.5%)** | **54.14 (99.7%)** |
The accumulation curve for the entire study area approached an asymptote. Although accumulation curves for the habitat types have not achieved asymptotes, the species lists could be considered complete in most of the cases, due to them containing more than 90% of the expected numbers of species calculated with the non-parametric estimators ACE, Chao1 and Chao2, or almost complete in four cases, where they contained 84–88% of the expected numbers of species (Table 2).

The comparison of species richness among the five habitat types, based on equal numbers of sites (n=11) revealed only small differences, with the highest number of species in the secondary channels and ponds and the lowest in the ditches (Fig. 2).

From among all molluscs recorded, 21 species (38.9%) occurred in all habitat types and 8 (14.8%) were unique (i.e. found only in one habitat type): two, four and two species were found exclusively in ML, T and D, respectively.

Cluster analysis based on relative abundance of mollusc species in each habitat type revealed a relatively high similarity of the malacoфаunas (Fig. 3). The highest similarity (i.e. the lowest distance) was found between ML and T, somewhat smaller between P and D. The malacoфаuna of SL showed a greater similarity to those in ML and T as compared to P and D.

The dominance pattern varied among the habitat types (Fig. 4). The malacoсоенoses of ML and T sites were dominated by small bivalves – *Sphaerium corneum* and representatives of the genus *Pisidium*. *Bithynia tentaculata* was the only gastropod species with a considerable proportion in the dominance structure within the rivers. The mollusc assemblages of SL and D were characterised by a considerable percentage (i.e. over 5% of the total abundance for individual species) of both gastropods and small bivalves. The malacoсоенoses of P sites were dominated by pulmo-
nate gastropods of the family Planorbidae. In SL and D sites all species contributed more than 2% to the total mollusc abundance. In the other habitat types, the percentage of species with a low contribution to the total abundance (<2%) increased in the following order: P (11.4%), ML (31.7%) and T (34.4%).

None of the species was common (i.e. F>50%) at the scale of the total study area and very few such species occurred within individual habitat types: S. corneum in ML, three bivalves (S. corneum, P. nitidum and P. subtruncatum) in T, B. tentaculata, P. corneus, A. vortex and S. corneum in SL, two gastropods (P. corneus and A. vortex) in P and four gastropods (B. tentaculata, A. vortex, P. planorbis and P. corneus) in D (Table 1). Regionally uncommon species (i.e. with frequencies lower than 5%) were numerous, as they comprised 44.4% of the total number of species within the entire study area. They also formed a large fraction of malacoofaunas within individual habitat types: 34.9%, 29.3%, 20.0%, 25.7% and 27.0% in ML, T, SL, P and D, respectively.

Within the study area six nationally rare (rarity category 4) and threatened species were found (over 11% of all mollusc species recorded). They occurred in all habitat types: three in ML, two in T, SL and P, and one in D (Table 1). Three species deserve a special mention: Unio crassus, Sphaerium rivicola and Anisus vorticulus. A. vorticulus is included in Annex II of the EU Habitats Directive (Council Directive 1992) and is legally protected in Poland (Dz. U. 2016). U. crassus and S. rivicola are included on the IUCN Red List of Threatened Species (IUCN 2018; EN and VU, respectively); furthermore, the first is listed in the EU Habitats Directive (Council Directive 1992, Annexes II and IV) and is legally protected in Poland (Dz. U. 2016). A. vorticulus was found at ten sites (8 in P and 2 in SL), U. crassus at five sites within the lower Liwiec River section, S. rivicola at eight sites (4 in ML and 4 in SL).

Several other species, which are not legally protected in Poland and their conservation status in Europe is LC (Least Concern), also deserve some comments, among others Valvata macrostoma, Anisus calciformis, Gyraulus laevis, Pisidium crassum, P. moitessierianum, P. hibernicum and P. pulchellum. V. macrostoma and G. laevis exhibit a decreasing population trend, they are relatively rare in Poland and other European countries, in some of them they are legally protected (Piechocki & Wawrzyniak-Wydzowska 2016, IUCN 2018). Within the study area V. macrostoma was found at seven sites and G. laevis only at one. A. calciformis is relatively common in some regions of Poland, but its population trend in Europe is unknown because of the scarcity of data (Piechocki & Wawrzyniak-Wydzowska 2016, IUCN 2018). It was found at 15 sites within the study area. All three gastropods are threatened with habitat loss and modification (channel regulation, drainage, pollution). Pisidium moitessierianum is not threatened through its range in Europe, but it is locally declining. Its general population trend is unknown (Piechocki & Wawrzyniak-Wydzowska 2016, IUCN 2018). Within the study area it was found only at 3 riverine sites. P. pulchellum is known from rather few localities in Poland, in Europe it is widely distributed, but locally declining. The general population trend is unknown (Piechocki & Wawrzyniak-Wydzowska 2016, IUCN 2018). It was collected from six sites within the study area. P. crassum and P. hibernicum are relatively rare in Poland, they are not included in the IUCN Red List of Threatened Species (IUCN 2018), even as DD (Data Deficient). The above mentioned pill-clams are threatened with habitat modification – pollution, changes to flow regime, dredging (Piechocki & Wawrzyniak-Wydzowska 2016, IUCN 2018). Within the study area they were found at three and six sites, respectively.

Table 3. Mean values (±SD) and ranges (in parentheses) of total species richness, mean species richness per site, Shannon index (H’), true diversity (exp(H’)), abundance and species rarity index (SRI) of molluscs in the main Liwiec River channel (ML), its tributaries (T), secondary channels of Liwiec (SL), ponds (P) and ditches (D)

|        | ML     | T      | SL     | P      | D      |
|--------|--------|--------|--------|--------|--------|
| H’     | 1.59±0.49 | 1.42±0.52 | 1.86±0.77 | 1.46±0.42 | 1.44±0.49 |
| exp(H’)| 5.50±2.60 | 4.74±2.61 | 9.12±10.81 | 4.68±1.82 | 4.68±2.09 |
| Mean   | 25.1±31.1 | 76.2±81.3 | 80.4±66.9 | 140.0±257.6 | 116.9±136.6 |
| Mean   | 1.91±2.20 | 1.16±0.20 | 1.66±0.91 | 1.24±0.29 | 1.11±0.13 |
|        | (1–11.33) | (1–1.60) | (1–3.50) | (1–2.0) | (1–1.37) |
**COMPARISON OF SPECIES RICHNESS, DIVERSITY, ABUNDANCE AND RARITY AMONG HABITAT TYPES**

The total number of species at individual sites varied from 1 to 21, but the mean values were fairly similar in all habitat types (Table 3, P>0.05). The highest proportion of sites with total species richness ≥10 was recorded within SL (45.5% of all sites within this habitat type), followed by T (34.4%), ML (29.3%), P (26.5%) and D (23.8%).

The mean number of species per site (i.e. alpha diversity) at individual sites ranged from 1 to 18 and there were no considerable differences among habitat types except the nearly significantly lower value in ML than in P (P=0.0539, Table 3).

The values of Shannon entropy (H') and true diversity (exp(H')) for individual sites varied widely (0–3.70 and 1–40.45 for H' and exp(H'), respectively). The mean H' and exp(H') values in individual habitat types ranged from 1.42 to 1.86 and 4.68 to 9.12, respectively (Table 3), however the differences among the habitat types were mostly not significant except the higher exp(H') value in SL as compared to T, P and D (P=0.0444, P=0.0400 and P=0.0397 for the comparisons with T, P and D, respectively).

The mollusc abundance at individual sites ranged from 1 to 1,400 ind./m², with the highest values (>100 ind./m²) recorded in 11 ponds and seven ditches (mainly due to abundant occurrence of a few desiccation-resistant species), as well as at two ML sites, nine T sites and four SL sites. The mean abundance of molluscs within ML sites was distinctly lower when compared to the other habitat types (Table 3, P=0.0046, P=0.0409, P=0.0000 and P=0.0003 for the comparisons with T, SL, P and D, respectively). Its values were fairly similar among SL, P and D and slightly lower in T as compared to P and D.

The species rarity index (SRI) at individual sites varied widely (1–11.33, Table 3), but the mean SRI values did not differ significantly among the habitat types (P>0.05). The highest SRI value was found in ML (11.33), the maximum value in SL was 3.5, whereas in T, P and D the SRI values were always below two. The proportion of sites with SRI values exceeding one increased as follows: ML (41.5%), T (53.1%), D (57.1%), P (61.8%), SL (63.6%).

**PARTITIONING OF MOLLUSC DIVERSITY ACROSS SPATIAL SCALES**

Additive partitioning of Shannon entropy (H') proved that the total entropy of 3.11 found within the entire study area can be divided into a mean within habitat type entropy (i.e. $\alpha_1$) of 2.60 and between habitat type entropy (i.e. $\beta_1$) of 0.51 (Fig. 5). The within habitat type entropy consisted of mean within site entropy (i.e. $\alpha_2$) of 1.55 and between site entropy (i.e. $\beta_2$) of 1.05. The within site entropy can be divided into mean within sample entropy (i.e. $\alpha_3$) of 1.17 and between sample entropy (i.e. $\beta_3$) of 0.38. The total entropy in each of the five habitat types can be divided into additive components in a similar way (Fig. 5). Based on additive partitioning the percentages of total landscape diversity attributed to site-specific diversity ($\alpha_i+\beta_i$), habitat heterogeneity ($\beta_i$) and habitat variability ($\beta_i$) were 49.84% (i.e. 37.62% +12.22%), 33.76% and 16.40%, respectively. This indicates a relatively high proportion of alpha diversity at the site level and the highest beta diversity at the medium scale (i.e. habitat heterogeneity, $\beta_i$). This regularity was also visible in all habitat types. The results of additive partitioning of Shannon entropy were generally consistent with the results of multiplicative decomposition of Shannon true diversity and species richness (Table 4).

**Table 4. Results of multiplicative decomposition of species richness and Shannon true diversity (exp(H')) in the distinguished habitats (ML, T, SL, P, D) and the entire study area (Total)**

|        | $\alpha_1$ | $\beta_1$ | $\beta_2$ | $\beta_3$ | $\gamma$ |
|--------|------------|-----------|-----------|-----------|---------|
| **Species richness** |            |           |           |           |         |
| ML     | 4.42       | 1.76      | 5.53      | 43        |         |
| T      | 6.00       | 1.35      | 5.10      | 41        |         |
| SL     | 7.40       | 1.41      | 3.35      | 35        |         |
| P      | 6.90       | 1.13      | 4.50      | 35        |         |
| D      | 6.90       | 1.19      | 4.50      | 37        |         |
| Total  | 5.80       | 1.40      | 4.60      | 1.40      | 54      |
| **exp(H')** |            |           |           |           |         |
| ML     | 3.10       | 1.77      | 2.81      | 15.46     |         |
| T      | 3.74       | 1.27      | 2.18      | 10.35     |         |
| SL     | 4.92       | 1.85      | 1.63      | 14.85     |         |
| P      | 4.10       | 1.14      | 2.92      | 13.67     |         |
| D      | 4.04       | 1.16      | 2.80      | 13.09     |         |
| Total  | 3.71       | 1.43      | 2.55      | 1.67      | 22.47   |
The proportions of diversity components varied among the habitat types. The highest proportion of site-specific diversity was found in SL. The highest within sample diversity (\(\alpha\)) was found in SL and the lowest in ML, whereas the highest between sample diversity (\(\beta\)) was recorded in ML and the lowest in P. The highest between site diversity (habitat heterogeneity, \(\beta\)) was noted in P and the lowest in SL.

DISCUSSION

During the study, rich and diverse malacoфаuna was found within a semi-natural Liwiec River catchment. In terms of landscape species richness (i.e., \(\gamma\)-diversity) lotic habitats – the Liwiec River and its tributaries – collectively supported a slightly richer malacoфаuna than the ponds, secondary channels of the Liwiec River and ditches together (50 and 44 species, respectively). However, species richness and Shannon diversity measures (\(H^'\) and \(\exp(H^')\)) were relatively equally distributed among the five habitat types.

The high mollusc gamma diversity in the Liwiec River and its tributaries could result from the high beta diversity, despite the connectivity within the river channels. This might be caused by the influence of stochastic events on the mollusc assemblages, such as the interaction of dispersal with local competition (and/or predation). The considerable disturbance of the river bed (with mostly sandy bottom sediments), low productivity within the river channel and stochastic extinction of less adapted species may also be important (Jurkiewicz-Karnkowska 2016 and references therein).

The total mollusc species richness found in the Liwiec River and its tributaries was similar to that recorded by Bôdis et al. (2016) from the Danube, Tisza and their tributaries and higher than in the Wkra River and its tributaries in Poland (Lewin 2014) or the Mediterranean Guadiana River (Perez-Quintero 2011). It was also similar or higher, compared to the values reported from other lowland and upland medium-sized rivers in Poland (Piechocki 1981), as well as some Lithuanian and Czech rivers of a similar nature (Pliuraitë & Kesminas 2004, Beran 2013).

The comparison of species richness for equal numbers of sites in every habitat type revealed that the secondary channels of the Liwiec River were the most species-rich habitat within the study area (but only slightly richer than ponds). This is consistent with the results of Bôdis et al. (2016) reporting the highest mollusc species richness and Shannon index (\(H^'\)) in side channels of the Danube, as well as with the earlier study within a short section of the Liwiec River and its valley (Jurkiewicz-Karnkowska 2015) and may result from both favourable environmental conditions (low water flow or even lentic conditions and abundant macrophytes) and habitat heterogeneity.

The proportion of regional diversity contained in the average sample reflected by MacArthur’s homogeneity measure (\(M=\exp(H^\gamma)/\exp(H^\alpha)\)) was relatively low both for the entire study area (\(M=0.165\)) and for individual habitat types (0.201, 0.361, 0.331, 0.300 and 0.309 for ML, T, SL, P and D, respectively).

Ponds and ditches have been reported to make an important contribution to aquatic biodiversity (e.g., Painter 1999, Armitage et al. 2003, Biggs et al. 2007, Davies et al. 2008, Gallardo et al. 2009, Simon & Travis 2011). Ponds are commonly recognised as a very heterogeneous habitat type which results in a high variability of their invertebrate assemblages, including molluscs (e.g., Scheffer et al. 2006, Davies et al. 2008). The malacoфаuna found in the investigated ponds was poorer than that of the Liwiec River and its tributaries. Similar results were obtained by Lewin (2014), who found more mollusc species in the Wkra River than in its oxbow lakes. However, when equal numbers of sites were compared, ponds investigated within the valleys of the Liwiec River and its tributaries supported more species than the main Liwiec channel, its tributaries and ditches, which is consistent with the results of Williams et al. (2004) from agricultural landscape in Southern England. The total number of mollusc species found in the studied ponds was lower than in the young permanent floodplain water bodies of the Bug River and higher than in the considerably desiccating and temporary ones (Jurkiewicz-Karnkowska 2009). It was also higher than in oxbow lakes of the Polish rivers Grabia and Wkra (Piechocki 1969, Lewin 2014) and 12 oxbow lakes of the Odra River (Piechocki & Szlauer-Łukaszewska 2013). The number of gastropod species was similar to the value reported from the water bodies of an active floodplain of the upper Rhine (Obdlik & Fuchs 1991).

The number of species collected in ditches was similar to the value reported from Wicken Fen, UK (Painter 1999) and higher than in ditches within agricultural areas in the Netherlands (Verdonschot et al. 2011). This resulted most likely from the location within river floodplains and the permanent presence of water in almost all surveyed ditch sites. According to the literature, species-rich ditches typically occur in floodplain and coastal environments or low-lying fen landscapes (e.g., Painter 1999, Armitage et al. 2003, Biggs et al. 2007).
Ditches, being the only habitat type of anthropogenic origin included in the present study, supported species-rich and abundant malacocoenoses with the number of species similar to that in the ponds and secondary channels of the Liwiec River (slightly smaller compared to equal numbers of sites, \( n=11 \)). It was the only habitat type where *Gyrulaus laevis*, rare in Poland, was found, as well as *Musculium lacustre*. This bivalve, like several other species recorded from the ditches (*Valvata macrostoma*, *Ladislavella terebra*, *Aplexa hypnorum*, *Anisus calculiformis*, *Gyrulaus rossoeckleri*), is not legally protected in Poland, but those species are threatened with extinction due to the disappearance of suitable habitats, resulting from land drainage. The importance of ditches, as a habitat supporting uncommon gastropods, has been previously reported, for example by Watson & Ormerod (2004).

Similarity among the malacoфаunas of the five habitat types was relatively high, indicating connectivity (hydrological or via biotic vectors, e.g. birds, insects, amphibians) among them. Some authors reported a large proportion of freshwater species in a range of water body types, which reflected the occurrence of a network among different kinds of freshwater habitats (e.g. Williams et al. 2004, Biggs et al. 2017, Bodis et al. 2016). The relatively low habitat variability (\( \beta \)), revealed by hierarchical partitioning of diversity, was consistent with the relatively high similarity among malacoфаunas of different habitat types.

In conclusion, within the semi-natural river-floodplain systems belonging to the Liwiec River catchment, the mollusc diversity was relatively equally distributed among the habitat types, only the mean species richness was lower in the main Liwiec River channel compared to the ponds and exp(H') was higher in the secondary channels of the Liwiec than in its tributaries, ponds and ditches. However, the mollusc diversity was not distributed evenly across the landscape. The diversity at the site level was generated mainly by alpha component, whereas at the landscape scale habitat heterogeneity (beta component) was very important. The significance of habitat variability (\( \beta \)) was distinctly smaller. The aquatic malacoфаuna of the Liwiec River catchment was rich and almost free of alien species (save for a few individuals of *Physella acuta* only, which were found at two riverine sites). Conservation value of mollusc assemblages of the five habitat types did not differ significantly. Nationally rare and threatened species were present in all of them. Although the highest SRI value was recorded in the main Liwiec River channel, a higher proportion of sites with SRI>1 was found in the secondary channels of the Liwiec, ponds and ditches. All the habitat types deserve attention in order to maintain diversity at the landscape scale. The care for their good ecological condition, particularly protection against pollution and eutrophication, is vastly important to maintain alpha diversity at individual sites. In the case of semi-natural habitats (main Liwiec River channel, its side channels and tributaries, small floodplain water bodies) preservation of hydrological processes structuring the habitat heterogeneity would be important for maintaining beta diversity at a medium scale (\( \beta \)). This seems at least partially achievable, due to the fact that a substantial part of the Liwiec River valley and some parts of the valleys of its tributaries are encompassed by different forms of areal protection (the Natura 2000 network, landscape protection areas) and the entire area is extensively used, mainly as meadows and pastures. Maintaining a high biodiversity of ditches requires their management in rotation, which ensures the presence of a range of ditch ages (e.g. Clarke 2015). For this reason, such a way of management should be executed within the Liwiec River catchment. Conservation efforts should focus mainly on the most heterogeneous fragments of the Liwiec River catchment to support mollusc diversity at the regional scale.

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