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Update on selected topics in acanthocephalan parasites research
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Summary

The respectable community of parasitologists aimed at the broad-spectral research of acanthocephalan parasites met at the 9th Acanthocephalan Workshop. The workshop took place in the beautiful surroundings of the High Tatras, Slovakia in the Congress Centre Academia, Stará Lesná near Tatranská Lomnica on September 9 – 13th. This special event was hosted by the Slovak Society for Parasitology, the Institute of Parasitology of the Slovak Academy of Sciences, Košice, Slovakia, and the Czech University of Life Sciences Prague, Czech Republic. It consisted of nearly three dozen lectures presented by distinguished acanthocephalan specialists who came from 13 countries and five continents. Vibrant discussions and creating new plans for future collaborations were accompanied by local mountain touring that offered the venue richly endowed with nature, deep forests and beautiful mountains. The contributions were addressed to resolve current systematic, taxonomic, biological, behavioural, ecological, and related topics. Presented results showed the most recent progressive developments comparable with all the other parasitic worm groups. The 10th Acanthocephalan Workshop will be hosted by Dr. Marie-Jeanne Perrot-Minnot, Université de Bourgogne Franche-Comté, Dijon, Bourgogne, France, in 2022.

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SYSTEMATICS AND TAXONOMY
Variability in the Acanthocephala
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Unique and unusual features in the many species of acanthocephalans described by Amin from fish, amphibians, reptiles, birds, and mammals, in various parts of the world including South America, Vietnam, Japan, the United States, the Middle East, and North

and East Africa, are described. The presentation is in five parts. (1) An introductory section dealing with the classification, general morphology, ecology, and life cycles of the Acanthocephala. (2) Unusual anatomical features of taxonomic or of questionable taxonomic importance addressing variations in the proboscis, proboscis hooks, male and female reproductive organs, and lemnisci. Newly described structures including (a) Para-receptacle structure (PRS) and hoods in certain species as well as a new order of Acanthocephala from Vietnamese birds, are also featured. (3) Structural and functional relationships explaining the relationship between the metamorphosis of the giant nuclei in Eoacanthocephala and worm reproductive cycle. (4) Host-parasite relationships elucidating the relationships between worm anatomy and biology during worm growth. (5) Curiosities in reviews and revisions highlighting taxonomically based zoo-geographical patterns and trends

in the genera *Neoechinorhynchus*, *Polymorphus*, and *Pallisentis*. A comprehensive treatment of the acanthocephalans of South America and those marine forms off the Eastern United States is also included here. A look at the September, 2013 classification scheme of the Acanthocephala is included covering 4 classes, 26 families, 157 genera, and 1298 species are included

Acanthocephalans in the Helminthological collection of the Institute of Parasitology (IPCAS), České Budějovice, Czech Republic

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The Institute of Parasitology of the Biology Centre of the Czech Academy of Sciences (IPCAS) is a governmental research institution established in Prague in 1962 and relocated to České Budějovice (Budweis) in South Bohemia in 1985. It is the principal institution in the Czech Republic devoted exclusively to basic parasitological research including helminthology. Thanks to the effort of František Moravec, world expert in the systematics of fish nematodes and former curator, the Institute hosts one of the largest helminthological collections in Central Europe. The collection is extraordinarily rich mainly in fish helminths such as nematodes and monogeneans. However, collection of acanthocephalans is also worth mentioning as it includes a total of 107 species (92 identified to the species level) of 7 orders and 18 families; as many as 80 species are from fishes. Type specimens of 16 species (holotypes of 8 taxa) are also deposited in the collection; 14 of them are parasites of teleosts, one of amphibians and one of mammals. In two cases, the new species were type species of newly erected genera, namely *Pseudogorgorhynchus* Moravec, Wolter et Körting, 2000 and *Sharpilosentis* Lisitsyna, Scholz et Kuchta, 2015, both from fishes. Specialists in acanthocephalan taxonomy are much welcome to request specimens deposited in the IPCAS. Deposition of type and voucher material of any acanthocephalans in the future would be much appreciated.

An integrated taxonomic approach to understanding species diversity of fish-parasitizing *Neoechinorhynchus* species

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Neoechinorhynchus Stiles and Hassall, 1905 is a highly diverse and widespread genus of acanthocephalan that parasitizes fishes and turtles on six continents. The literature base for the genus grew rapidly throughout the 20th century, mostly as a result of original descriptive work recognizing the diversity across the world. There have been, however, relatively few contributions regarding North American species of *Neoechinorhynchus* since the turn of the 21st century, and a particular lack of use of modern molecular tools. Several recent studies used DNA sequence data to understand species relationships and uncover cryptic species of *Neoechinorhynchus* in Mexico, but there is still no molecular phylogenetic analysis focused on species of *Neoechinorhynchus* from the USA or Canada. Here we present the phylogenetic results of the first such study, based on nuclear DNA from the internal transcribed spacer (ITS) and large ribosomal unit (LSU) regions, including molecular data from multiple specimens of each of 17 total species (two from turtles and 15 from fishes) from the United States and Canada. We also present morphological evidence supporting the boundaries of several of these species. In addition to giving the first look at relationships between these species, this work has uncovered three previously uncharacterized species from white suckers of Central New York State. Plus, we have found that a species of another neoechinorhynchid genus, *Octospinifer* van Cleave, 1919 nests among species of *Neoechinorhynchus*, lending further support to the recently recognized paraphyly of this large and somewhat enigmatic genus.

***Pomphorhynchus* spp. in the Rhine system – recent situation and future perspectives**

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Over the last few years contrasting results regarding the distribution of two previously synonymised acanthocephalan species, *Pomphorhynchus laevis* and *Pomphorhynchus tereticollis*, have been published for the Rhine river system. Shortly after its redescription, Emde et al. (2012) stated a high invasive potential for *P. tereticollis*, due to its occurrence in the invasive round goby. Recently, a contradictory study by Hohenadler et al. (2018) showed a gradual replacement of *P. tereticollis* by *P. laevis* in eel. However, a distinction between the two congeners remains difficult, since both studies used either a morphological or molecular approach for the differentiation and furthermore, they only took account of unsuitable hosts. Hence, fish of different taxa from the Upper Rhine near Karlsruhe were examined for acanthocephalan infestation to clarify the recent distribution of the congeners. To facilitate distinguishing between the two species, randomly selected *Pomphorhynchus* worms were verified molecularly by

sequencing the COI-locus. Proboscis hooks of the same specimens were counted and measured, allowing the analysis of hook patterns with the Proboscis Profiler (Wayland, 2010). Significant differences between the proboscis armature of *P. laevis* and *P. tereticollis* allowed a sufficient species separation. Principal Component Analysis of the hook profiles revealed a correlation between molecular and morphometric traits, thus yielding a potential new option for morphological determination.

Examination showed high infection rates with mature *P. laevis* in common barbel and chub, while *P. tereticollis* was only found at low abundance in chub and brown trout. For the first time co-infections of both species are recorded in an individual host. Unsuitable hosts, such as catfish, eel and round goby, were exclusively infected with immature *P. laevis*. In light of the recent findings the distribution of *Pomphorhynchus* spp. in the Rhine River and tributaries is discussed, as well as the influence of invasive Ponto-Caspian fish and amphipod species as acanthocephalan hosts have been re-evaluated.

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Never-ending story of *Pomphorhynchus* spp. – back to the type material?

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Pomphorhynchus laevis (Zoega in Müller, 1779) was considered to be the predominant species in European fishes for a long time, characterized by high morphological and physiological intraspecific variability (e.g. Kennedy, 1984). During the first decade of the 21st century, previously synonymised *Pomphorhynchus tereticollis* (Rudolphi, 1809) was resurrected (Bombarová et al., 2007; Špakulová et al., 2011): taking into account the ICZN rules, the Ru-

dolphi's type material was compared with newly collected worms from the type locality (Baltic coast) and the type fish host (*Platichthys flesus*), and the worm morphology was combined with new comparative genetic characteristics (gene sequencing, karyotype). The existence of these species is apparent but their distinction is difficult. The same applies to other congeners described so far in Europe using only morphological tools. The latest extensive study of *P. laevis* and *P. tereticollis* phylogeography throughout Western Palaearctic (Perrot-Minnot et al., 2018) showed that *P. tereticollis* occurs within Western and Central parts of Europe showing weak geographical and genetic structuring, while *P. laevis* sensu lato is representing by five lineages partially matching several major biogeographical regions. That means we have a big task ahead of us – i.e. to harmonize so far existing European *Pomphorhynchus* species with the recently described lineages, re-describe their morphology, and to re-establish species spectrum within the whole Europe including its Peri-Mediterranean and Ponto-Caspian parts. The best way may probably be to use newly acquired acanthocephalans from type localities and type fish hosts (in the absence of ethanol-fixed material in museum collections). Further, re-descriptions on basis of integrative approaches should include morphology, gene sequencing and maybe also chromosome analysis. A promising beginning may be *Pomphorhynchus* spp. from Sava River, the İşıklı Lake, the Volga River etc.

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Integrative taxonomy of acanthocephalans from Brazilian mammals

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Since the early twentieth century, species of acanthocephalans have been reported in mammals from different geographic regions in Brazil (Vieira et al. 2008). However, there is still a lack of adequate, complete taxonomic, phylogenetic and ecological information regarding these species. As part of a PhD project carried out with the support of the Oswaldo Cruz Foundation (FIOCRUZ), Rio de Janeiro, Brazil, we aimed to identify acanthocephalan species collected from wild mammals in Brazil and investigate their phylogenetic relationships. Specimens of acanthocephalans were identified based on morphological and morphometric features using light and scanning electron microscopy and DNA analyses. We identified *Gigantorhynchus echinodiscus* (Gigantorhynchidae) in a giant anteater, *Myrmecophaga tridactyla*, from the Cerrado in the state of São Paulo. Although the species has been previously reported in this host, genetic data were absent. Our study provides details on the structures as a bipartite cylindrical proboscis with a crown of 18 hooks and numerous small hooks, pseudo-segmented body, long leminisci, large testes and ovoid eggs. Such features agree with the genus *Gigantorhynchus* characteristics (Yamaguti, 1963). Our genetic study included new sequences of partial 28S rRNA gene. Phylogenetic analyses were inferred using maximum-likelihood in PhyML 3.0 and Bayesian analysis in MrBayes version 3.2.6, this last using the CIPRES platform. Results showed that *G. echinodiscus* is closely related to *Mediorhynchus* sp., both forming a well-supported monophyletic group (scores=0.92/0.95). The family Gigantorhynchidae includes two genera, *Gigantorhynchus* and *Mediorhynchus*, both possessing bipartite proboscis with large hooks and small hooks, pseudo-segmented body, long or short and slender leminisci, large testes, eight compact cement glands and ovoid eggs. Genetic data were congruent with morphological studies, classifying both species within the family Gigantorhynchidae (Amin, 2013). This study extends our knowledge

about acanthocephalans from Brazilian mammals, emphasizes the importance of integrative taxonomic studies to better understand their taxonomy and evolution.

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Redescription of *Acanthogyryus (Acanthosentis) maroccanus* (Dollfus, 1951) (Acanthocephala: Quadrigyridae) from the Algerian barb *Luciobarbus callensis* (Cypriniformes: Cyprinidae) from Algeria

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The fresh water acanthocephalan *Acanthogyryus (Acanthosentis) maroccanus* Dollfus, 1951 was insufficiently described in *Luciobarbus setivimensis* (Cypriniformes: Cyprinidae) in Morocco. Although this acanthocephalan has been reported from cyprinid fishes in northern Africa, its detailed morphological description has never been provided. In this study, we present novel morphological data including scanning electron microscopy micrographs (SEM), based on specimens found in the Algerian barb *Luciobarbus callensis* from the Oued Charef Dam Lake, Algeria. The most typical characteristics of *A. maroccanus* are: (i) short proboscis armed with 3 incomplete circles of 6 rooted hooks each; (ii) proboscis hooks of different lengths; (iii) trunk shorter than 8.5 mm; (iv) trunk with 17–19 circles of spines, covering almost completely the anterior part of the trunk; (v) 3–4 ventral and 6–8 dorsal giant hypodermal nuclei; and (vi) male reproductive structures (sperm duct and Saeftigen's pouch) extending into bursa. Our observation also revealed the presence of the parareceptacle structure and vaginal sleeve in females of *A. maroccanus* from Algeria. Additionally, we obtained partial sequences of the large ribosomal subunit (28S) and mitochondrial cytochrome c oxidase 1 (*cox1*)

genes for this species; representing the first molecular data for the species-rich genus *Acanthogyrus* Thapar, 1927.

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Acanthocephalans from California sea lions (*Zalophus californianus*): New records and evidence for synonymy of *Corynosoma obtuscens* with *Corynosoma australe*

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California sea lions (CSLs) (*Zalophus californianus*) are one of the most abundant and recognized pinniped species in the North Pacific. To increase the limited knowledge addressing acanthocephalans parasitizing CSLs, 33 animals including pups, juvenile and adult sea lions were examined at The Marine Mammal Center (TMMC), California, USA, in 2012, 2015 and 2016. Totally, 2,268 specimens of acanthocephalans representing 5 species from three genera *Andracantha* (*A. phalacrocoracis* and *Andracantha* sp.), *Corynosoma* (*C. strumosum*, *C. obtuscens*) and *Profillicollis* (*P. altmani*) were found. The total prevalence was 73 % and intensity of infection varied from 1 to 1,225 (average = 94.5; median = 19). *P. altmani* and *A. phalacrocoracis*, predominantly parasites of fish-eating birds, were registered in CSLs for the first time and only in juvenile CSLs. Significant differences were observed in the species diversity in different age groups of CSLs. Detailed morphological and molecular examinations of *C. obtuscens* collected from CSLs gave us solid evidences for a synonymy of this species with *Corynosoma australe*. Comparison of the metrical and quantitative characters of 51 specimens of *C. obtuscens* (24 males, 27 females) from CSLs from California, USA and published data on *C. australe* from different hosts and regions, including type material of

the both species (Johnston, 1937; Smales, 1986; Hernández-Orts et al., 2017) did not reveal any significant differences between these species. The level of the genetic divergence in the *cox1* sequences from *C. obtuscens* from CSLs and *C. australe* from others marine mammals and penguins reaching 1.4 – 1.6 % can be considered as an intraspecific variability. Thus, according to the rule of priority the name *C. obtuscens* should be recognized as the junior synonym of *C. australe*.

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Acanthocephalans from common carp (*Cyprinus carpio* L.) from the Yangtze River basin, China

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Within the years 2009–2018, a total of 1,687 fish specimens belonging to 36 species of 11 families were collected and examined for intestinal parasites throughout the Yangtze River basin. The present work resumes acanthocephalan parasites from the family Illiosentidae Golvan, 1960 collected from the intestine of common carp (*Cyprinus carpio* L.) from the Tai Hu Lake in Wuxi, Jiangsu Province, China. A total of 17 market-size common carps (total length of 30 – 50 cm) were purchased at the fish market or from local fishermen and examined. In total, 21 specimens of acanthocephalans were found in 6 carps (prevalence of 35 %). Worms were washed in tap water, relaxed in refrigerator and then fixed in 4 % formalin solution under slight pressure. Several specimens were fixed in 96 % molecular-grade ethanol for molecular analyses. For final morphological examination, temporary microscopic preparations in Berlese's medium were made; line drawings were made using a drawing attachment on a Zeiss Axio Imager M1 compound

microscope equipped with DIC optics and a digital imaging system. All the studied specimens were assigned to the Illiosentidae based on the family-specific morphology of the worms and the presence of 8 cement glands in males. However, specimens recently found in the common carp differed from all the existing genera of the family Illiosentidae by the i) structure of the reproductive system of females, i.e. vagina lacking a muscular sphincter; ii) presence of the terminally pointed protruding tail end in the form of a dome with a muscular base; iii) proboscis receptacle having a goblet-shaped widening in the anterior part of worms of both sexes, absent in representatives of other genera. In addition, recently found specimens are the only representatives of the family Illiosentidae hitherto found in freshwater ecosystems; all other species of the family were referred to parasitize marine fish. Therefore, a new acanthocephalan genus and species are to be proposed from the carp *C. carpio* from the Yangtze River basin, China.

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DIVERSITY AND GEOGRAPHICAL DISTRIBUTION

Acanthocephala from sharks and rays (Elasmobranchii)

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Acanthocephalan parasites, although relatively common in bony fish, are not usually found in cartilaginous fish (sharks and rays) and parasite-host switching between bony and cartilaginous fishes has not been reported. On occasion infections of elasmobranchs have been recorded, but usually the infection has been light, the worms not mature, or a single worm only recovered. In such instances the infections have either been considered accidental, or, in some circumstances, the result of an infected definitive host being consumed by the shark or ray. At least 14 species of acanthocephalan have been so recorded. A further 4 species have been described only from elasmobranchs (Weaver & Smales, 2014). Dissection of elasmobranchs from Australian waters is continuing and here we present preliminary results. Firstly, the probable accidental infection of *Sphyrna mokarran*, the great hammerhead, with *Corynosoma cetaceum* whose usual hosts are dolphins. Secondly, an as yet undescribed species from the wobbegong, *Sutorectus tentaculatus*. In this case the worms, males and females up to 50 mm long, were mature and the females were gravid. The species cannot be easily placed in any family and the morphologically similar species have not been recorded from any other fish found in Australian waters. This represents the fifth example of a species known only from elasmobranchs and indicates that in some cir-

cumstances elasmobranchs may act as suitable definitive hosts for acanthocephalan parasites.

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Acanthocephalans in northern fur seals (*Callorhinus ursinus*) from St. Paul Island, Alaska: results of four-year observations

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Several species of acanthocephalans (Acanthocephala: Polymorphidae) are known to parasitize in northern fur seals (NFSs) (*Callorhinus ursinus*). Acanthocephalans from the genera *Corynosoma* and *Bolbosoma* were found in NFSs in different parts of their range, including the NFSs from the Pribilof Islands (St. Paul and St. George Islands), where about 50 % of the NFS world population have breeding rookeries. The aim of the present study was to examine a species diversity of acanthocephalans parasitizing the NFSs at different rookeries of St. Paul Island, Alaska. During 2011–2014, gastrointestinal tracts of 756 humanely harvested NFS subadult males were examined during the annual Aleut subsistence harvest at five rookeries: Lukanin (165), Polovina (164), Gorbach (125), Zapadnyj (182) and Morzhovyi (120). All acanthocephalans (1,170 specimens) were collected manually, fixed in 70 % ethanol and identified by the morphology after their clarification in the Berlese medium. Totally, 8 species of acanthocephalans from two genera *Corynosoma* and *Bolbosoma* were found in NFSs with a total prevalence of 43.4 %. *Corynosoma strumosum* was the most prevalent species (20.9 %); the prevalence of the other species was lower: *C. similis* – 13.5 %, *C. semerme* – 14.6 %, *C. villosum* – 12.8 %, *C. validum* – 1.9 %, *C. alaskensis* – 0.1 %, *C. cameroni* – 0.7 %, *B. nipponicum* – 1.6 %. The intensity of infection of acanthocephalans in NFSs was low – from 1 to 31 (average = 3.6±4.3; median = 2) parasites per seal. Differences in the species composition and prevalence were observed in NFS subpopulations from the five rookeries. The highest biodiversity of acanthocephalans was observed in subpopulations from Morzhovyi rookery (8 species); the lowest – in Lukanin (6 species). Comparison of our results with data previously collected in Alaska and other parts of the North Pacific revealed a decrease of the NFS infection with acanthocephalans during last four decades.

Diversity of Acanthocephala in some teleost fishes from Kalaat El Andalous, Tunisia

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In Tunisia, the first record of Acanthocephala was reported in 2015 from the Bizerte lagoon. In order to improve knowledge about this group, we extended our investigations to other localities such as lagoon and sea of Kalaat El Andalous (Northern Tunisia). During two years (April 2015 to May 2017), a total of 671 fishes belonging to four families, Gobiidae (*Gobius niger*, *G. paganellus* and *G. geniporus*), Scorpaenidae (*Scorpaena porcus*), Sparidae (*Lithognathus mormyrus*) and Belonidae (*Belone belone* and *B. gracilis*) were examined. Acanthocephalans were collected from the digestive tract of the fish and identified on the basis of the morphological features. The present study allowed us to identify three families of Acanthocephala including Illiosentidae Golvan, 1960: *Telosentis exiguus* von Linstow, 1901, Arhythmacanthidae Yamaguti, 1935: *Acanthocephaloides propinquus* Dujardin, 1845 and *A. irregularis* Amin, Oğuz, Heckmann, Tepe & Kvach, 2011, and Echinorhynchidae Cobbold, 1879: *Acanthocephalus lucii* Müller, 1776. The solution of parasitic infra-communities showed that they are species-poor; 75.90 % of fishes harboured only a single parasite species, whilst 23.09 % of them were infected with two species of Acanthocephala. *Telosentis exiguus* and *A. propinquus* were found along the entire digestive tract, whereas *A. irregularis* and *A. lucii* were restricted to its posterior part. The overall dynamics of the parasitism showed that *T. exiguus* is the most frequent species with the highest percentage of the prevalence (23.58 %) in *G. niger*. *Acanthocephaloides irregularis* and *A. lucii* were reported only in *S. porcus* and *L. mormyrus*, respectively. *Acanthocephalus lucii* was recorded in *L. mormyrus* from the Mediterranean Sea for the first time. The data of the present study were compared with results available from other localities and countries.

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Acanthocephalans of Carangidae and Mullidae from Northern Tunisia: Ecology and morphology.

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The acanthocephalan fauna of fish is poorly known in Tunisian waters and published data are restricted to those of Gargouri et al. (2015). In order to fill this gap, our survey is focused on acanthocephalans of commercially important fish in Bizerte Lagoon (Northern Tunisia). A total of 470 fish specimens belonging to Carangidae (*Trachurus trachurus*, *Caranx rhonchus*, *Trachinotus ovatus*) and Mullidae (*Mullus surmuletus*) were collected and examined for thorny headed worms. Five acanthocephalan species belonging to four different families were recovered, namely two species of Arhythmacanthidae Yamaguti, 1935: *Acanthocephaloides* sp. Meyer, 1932 and *Breizacanthus irenae* Golvan, 1969, and one of Illiosentidae Golvan, 1960: *Telosentis exiguus* (von Linstow, 1901), Polymorphidae Meyer, 1931: *Southwellina hispida* Van Cleave, 1925 and Pomphorhynchidae Yamaguti, 1939: *Longicollum pagrosomi* Yamaguti, 1935, respectively. The species showed flexibility in the spatial distribution within the host intestine being present in different parts of the gut, except for the cystacanth of *S. hispida*, which often occupied the mesentery of the anterior intestine. The latter species was recorded in *T. trachurus* from the Mediterranean Sea for the first time with a prevalence of 2.5 %. A new host record is similarly noted for *Acanthocephaloides* sp. (P = 1.7 %) recovered from *C. rhonchus*. This carangid also harboured the pomphorhynchid *L. pagrosomi* with a very low prevalence (1.11 %). *Breizacanthus irenae*, with the relatively important prevalence (P = 25 %), was collected from *M. surmuletus*. The most widespread acanthocephalan species (generalist) was *T. exiguus* (P = 35 % in *T. trachurus* and P = 27.8 % in *C. rhonchus*). The light and scanning electron microscopy showed that *T. exiguus* exhibited variations in the trunk size and number of proboscis hooks, compared with other previously reported morphological observations (Dezfuli & Sbrenna, 1990, Kvach & Sasal, 2010.). These preliminary data enhance the understanding the plasticity of acanthocephalans and promote further taxonomical research within this parasite group in the Mediterranean area.

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Acanthocephalans of the genus *Neoechinorhynchus* of the Baikal rift zone

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Neoechinorhynchus Stiles et Hassal, 1905 represents one of the largest genera within Acanthocephala. The latest revision of the genus (Amin, 2002) provided information on 88 species split into two subgenera. Only two species of this genus, namely *N. rutili* and *N. crassus*, were listed in the Key to the parasites of the freshwater fish fauna of the USSR (1987). However, more recent study on *Neoechinorhynchus* in northeastern Russia (Atrashkevich et al., 2016) showed that the genus is more species-rich there. We examined acanthocephalans of this genus in various cyprinid fishes from the water-bodies of the Baikal basin and found the fragmented presence of a single species *N. rutili*. Its occurrence was restricted to the mouths of the rivers Selenga and Upper Angara and adjacent shallow water zones, and it was absent in the open Baikal as well as the upper and middle streams of the abovementioned rivers. The prevalence of *N. rutili* differed in various fish hosts, being high in *Leuciscus idus* (P = 18.4 %), lower in *L. leuciscus baicalensis* (P = 1.4 %) and low in the *Rutilus rutilus* (P = 0.6 %). In general, the acanthocephalans of the Baikal Lake, including *N. rutili*, are limnophilic species and we suppose that the favorable conditions could be associated with a high water velocity in the rivers of the Baikal area and the drift of infected intermediate hosts into the lake. Another congeneric species, *N. tumidus*, was found far from the Baikal in the isolated Lake Baunt (northern Transbaikalia, the River Lena Basin) in *Lota lota* (70.3 %), *Coregonus lavaretus* (75.3 %) and *L. leuciscus baicalensis* (26.3 %).

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LIFE CYCLE AND ECOLOGY

Divergent selective pressures on male and female acanthocephalans: morphological and ecological consequences

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Acanthocephalans are dioecious parasites with a polygynous mating system. Both experimental and observational data drawn from several species suggest that males have a shorter life-span and intensely compete for the access to non-mated females, with larger body sizes likely rendering greater reproductive pay-offs. Regardless of whether females are choosy (this is not known), they live longer than males to produce the offspring in suitable microhabitats where nutritional resources are abundant. Accordingly, adult males and females should develop different strategies to face divergent selective pressures. In this presentation, we explore this question in two key aspects, i.e., the investment in holdfast structures, and microhabitat selection, using species of *Corynosoma* as a model. Firstly, we examined the size of body spines in cystacanths and adults of *C. australe* and *C. cetaceum*. We found that spines grow in both species, but only in females, which also have significantly larger spines than males. This sexual dimorphism does not result from a pure allometry and conform to the hypothesis that females need to withstand the extreme flow conditions prevailing in marine mammals for longer. Secondly, we compared the intestinal distribution of non-mated and mated females, and males, of *C. australe* and *C. strumosum* in two pinniped species. We found that both species share a strikingly similar pattern: the distribution of non-mated females is significantly more anterior than that of mated females, which concentrate in intestinal regions where a nutrient absorption occurs. The distribution of males, however, is significantly more anterior than that of females. Apparently, males, but no females, experience a trade-off between growing in suitable microhabitats and contacting non-mated females that migrate towards these microhabitats. A comparison of body sizes between sexes in *C. strumosum* along the intestine conforms to this interpretation.

Cryptic species and unexpected intermediate host specificity in the acanthocephalan *Polymorphus minutus*

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Polymorphus minutus is a bird-infecting acanthocephalan and is the most commonly reported species of this genus in central Europe. Previous data suggests that *P. minutus* comprises of different lineages or even cryptic species exploiting different intermediate hosts. Our aim was to test the genetic diversity of *Polymorphus* c.f. *minutus* depending on locality and intermediate host species, to examine if *P. minutus* can be considered a cryptic species. We sampled amphipod intermediate hosts infected with *Polymorphus* cf. *minutus* cystacanths originating from 27 sites in Germany and France. Parasites and hosts were identified using integrated datasets (COI and/or morphology for hosts and COI + ITS1-5.8S-ITS2 and morphology for parasites). Mitochondrial and nuclear genetic data strongly supported the existence of three cryptic *Polymorphus* cf. *minutus* (type 1-3) species. The morphological analyses indicated minor differences in hook dimensions between the types, while no difference was observed in the analysis with "Proboscis Profiler". The three types revealed a high degree of intermediate host specificity: *Polymorphus* type 1 was only encountered in *Gammarus fossarum* type B, *Polymorphus* type 2 in *Echinogammarus* sp. and *Echinogammarus berilloni*, and *Polymorphus* type 3 in *Gammarus pulex* and *Gammarus roeselii*. These results point to a so far neglected cryptic diversity of the genus *Polymorphus* in Central Europe. Furthermore, *Polymorphus* type 2 is most likely a non-native parasite in Germany that co-invaded with *E. berilloni* from the Mediterranean area. Potentially, type 3 originated from South-East Europe and migrated to Germany by *G. roeselii*, where it was captured in *G. pulex* as intermediate host. Therefore, our findings can be seen in the context of ecological globalization in terms of the anthropogenic displacement of intermediate hosts and its impact on the dispersal of parasite species. Moreover, our data clearly point to a taxonomic revision of *Polymorphus* c.f. *minutus*.

Massive infections of *Neoechinorhynchus buttnerae* in an Amazonian fish, *Colossoma macropomum*, with notes on the parasites biology

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For over a decade, an acanthocephalan *Neoechinorhynchus buttnerae* has caused production losses for *Colossoma macropomum* producers in Brazil. In some cases, the prevalence of the parasite reaches up to one hundred percent being accompanied with high infections causing a morphological damage of the intestine and compromise the growth of the fishes. The life cycle of this parasite was recently described under experimental conditions using intermediate hosts fed with the eggs of adult specimens of *N. buttnerae* collected from *C. macropomum*. Every stage of the development was photographed, measured, drawn and described. In total, ten larval developmental cycles were studied. This information might help future studies in hopes to control the massive infections in the most produced native fish of Brazil.

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Host-switch of *Corynosoma australe* (Polymorphidae) from mammal seals to bird penguins: exploiting an ancestral resource?

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Corynosoma australe is common acanthocephalan specific to otariid pinnipeds in temperate and sub-Antarctic waters of the Southern Hemisphere. This acanthocephalan is trophically-transmitted and is regularly exposed to hosts and non-hosts through complex food web interactions. In fact, immature specimens of *C. australe* have been reported in higher trophic levels hosts (e.g. sharks, birds and cetaceans) in which apparently, they cannot grow or reproduce. In this study we investigate an apparent host-switching event of *C. australe*, since gravid worms were collected from the intestine of Magellanic penguins *Spheniscus magellanicus* from Brazil. A morphological study showed that the acanthocephalans from penguins belong to *C. australe*. Partial fragments of the 28S rRNA and mitochondrial *cox1* genes were amplified from isolates from penguins and two pinniped species (i.e. South American sea lion *Otaria flavescens* and South American fur seal *Arctocephalus australis*) to confirm this identification. Infection parameters clearly differed between penguins and the two pinniped species, which were significantly lower in the former. The sex ratio of *C. australe* also differed between penguins and pinnipeds; in penguins it was strongly biased against males, while in pinnipeds it was close to 1:1. Interestingly, females of *C. australe* from the South American sea lion were significantly smaller than those from penguins and the South American fur seal, but fecundity was lower and more variable in females collected from penguins. At first glance, the occurrence of reproductive individuals of *C. australe* in Magellanic penguins could be interpreted as an adaptive coloniza-

tion of a novel avian host through favourable mutations. However, it could also be considered, perhaps more likely, as an example of ecological fitting through the use of a plesiomorphic (host) resource, since the ancestors of *Corynosoma* infected aquatic birds.

BIOLOGY, ACANTHOCEPHALANS IN POLLUTED ENVIRONMENTS

Characterization of nutritional relationships between acanthocephalans and their hosts as compared to other host-parasite associations using stable isotope analyses

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Stable isotope analysis of carbon and nitrogen can deliver insights into trophic interactions between organisms. While many studies on free-living organisms are available, the number of those focusing on trophic interactions between hosts and their associated parasites still remains scarce. Information about some taxa such as acanthocephalans or monogeneans is either very limited or even completely missing. Additionally, available data revealed different and occasionally contrasting patterns, which is most likely depending on the parasite's taxonomic position and its degree of development. Among others, we have determined $\delta^{13}\text{C}$ and $\delta^{15}\text{N}$, in different acanthocephalan-host systems considering larval as well as adult stages of aquatic and terrestrial hosts. Herewith, we are able to evaluate the trophic position of acanthocephalans with respect to their hosts and to analyse a potential trophic level shift associated to the transmission from intermediate to definitive hosts.

In the context of a meta-analysis we have added this information to already existing data for approximately 100 host-parasite associations considering many taxonomic groups of parasites. This data compilation allowed us to describe patterns of nutritional relationships between hosts and their parasites. Taxa such as cestodes, trematodes and acanthocephalans as well as larval nematodes were mostly depleted in ^{15}N . Parasitic crustaceans (Copepoda, Isopoda, Cirripedia) and gastropods showed usually similar or lower isotope values, whereas parasitic insects, arachnids, fish as well as adult nematodes were enriched in ^{15}N (on average 4.6 ‰). In the taxa Cestoda, Trematoda and Acanthocephala processed nutrients provided by the host are assimilated via the body surface. Taxa such as Nematoda, Crustacea, Insecta, Arachnida and others, actively feed on tissues or fluids of their host organisms. Whereas the latter behave similar to micropredators, the nutrient absorbing parasite taxa appear to be commensal-like organisms

(acanthocephalans, cestodes) with respect to their nutritional strategy.

Combined effects of parasitism and anthropogenic stressors in the freshwater amphipod *Gammarus fossarum*: Impacts on multiple traits

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The interest in acanthocephalan fish parasites as model organisms in ecotoxicology increased ever since pioneer studies more than 20 years ago. More recently, the need for assessing multiple stressors has been emphasized, as well as the interest in combining behavioural and physiological responses with life-history traits. We address here whether the infection of the freshwater amphipod *Gammarus fossarum* with *Pomphorhynchus tereticollis* interacts with chemical pollutant and temperature. Using a fully crossed factorial design, we investigated whether infection at the population level (naïve populations without parasite historical record vs non-naïve populations with historical records) and at the individual level (infected and uninfected organisms) changes the effect of the carbamate insecticide (methomyl) and the temperature. For this, we assessed the expression of various behavioural and physiological traits (general defence systems), and food intake and survival as components of functional response and fitness respectively. Infection with *P. tereticollis* has pervasive effects on most of the traits measured. At the population level, uninfected gammarids from non-naïve populations exhibited decreased AChE and antioxidant potential, and increased Glutathion-S-transferase activity, lipid peroxidation and Pro-phenoloxidase activity. At the individual level, infection induced decreased feeding rate, increased AChE and GST activity, and decreased anti-oxidant potential, lipid peroxidation, and phenoloxidase activity. By contrast, methomyl alone had limited effect, restricted to decreased AChE activity (its target) and feeding rate. We evidenced very few interactions between stressors; however an additive effect of population infection history and methomyl exposure was evidenced on AChE activity and feeding rate. Temperature mainly modulates the effects of other factors – methomyl, population, and infection- therefore making the predictions about the outcome of global changes in temperature complex. Given these pervasive effects of population parasite history and individual infection, parasitism should be taken into account when running ecotoxicological monitoring, to gain realism in their conclusions when addressing ecological risk.

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Mechanistically modelling metal accumulation in fish-parasite systems: Ongoing attempts

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Fish are affected by both, exposure to metals and infection with parasites. Each of these stressors might have effects on the response of fish to the other. Some efforts have recently been done in developing kinetic models for a prediction of the metal accumulation in fish-parasite systems. Our previous model allows the investigation of relationships between its accumulation in the fish and acanthocephalans, but does not include the mechanisms how metals are accumulated in parasites. Physiologically-based pharmacokinetic (PBPK) model has been used for simulating the organ-specific accumulation of chemicals. However, the capacity of this model for mimicking fish-parasite systems has not been investigated. We aimed to integrate mechanisms for the simulation of metal accumulation in the system of the chub *Squalius cephalus* and acanthocephalan *Pomphorhynchus tereticollis* by developing of PBPK model, in which acanthocephalans were considered a compartment, similar to the blood, storages (muscle, skin, bone, and carcass), gill, kidney, liver, and intestine. The metal accumulation in the system was modelled as a function of internal (i.e. exchange between different compartments) and external (i.e. exchange with water) factors. The flow from the blood to the other compartments depends on diffusive exchange and fraction of metals dissolved in the blood plasma. The flow to the blood is a function of diffusive exchange and blood-tissue partitioning. The model was calibrated by MATLAB-based AMIGO Toolbox. As a detoxifying organ, the liver was found to accumulate Ag at the highest level. Initial results from the model calibration for Ag show a good

potential of the model for simulation the accumulation in the muscle, gill, and liver. However, further attempts are being examined to improve the efficiency of the model in simulation of Agdelayed accumulation in the intestine as well as in acanthocephalans.

Comparison of the metal accumulation capacity of acanthocephalans from fish and mammalian definitive hosts

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Recent research on the metal accumulation potential of parasites provided strong evidence that various taxa can be used as sentinels for metal pollution. Among the taxa of parasites tested so far with respect to their metal accumulation capacity acanthocephalans have been found to accumulate metals to a much higher degree than the host and the host's surrounding environment. However, until now, most studies focused on limnetic host-parasite associations, whereas only few papers considered marine acanthocephalan-host systems. Moreover, most studies were performed on fish as definitive hosts, information about the accumulation potential of acanthocephalans from (marine) mammals is scarce. Here, we summarize data of several metal accumulation studies that were performed on marine acanthocephalans (belonging to the genera *Neoechinorhynchus* and *Corynosoma*) collected from their definitive fish (different species of mullets, Mugelidae) and mammalian hosts (common seal, *Phoca vitulina*). Moreover, based on a well-studied limnetic acanthocephalan species (*Pomphorhynchus laevis* from its cyprinid host) we also address the usefulness of acanthocephalans as sentinel organisms for rare earth elements (REE), which are – although being present only in low environmental concentrations – mainly introduced into ecosystems due to anthropogenic activities. The results confirmed that acanthocephalans from fish accumulated various elements to a higher extend than the host itself. This trend was found for both essential (e.g. Cu, Mn, Zn) and non-essential metals (e.g. Cd, Pb) as well as for some REE, especially when compared to host's muscle tissue. Acanthocephalans from the mammalian host did not show such pronounced accumulation patterns and the concen-

trations of most elements were lower or comparable to those found for kidney or liver tissue. Interestingly, several elements such as Ag, Cd, Mn, Ni and V were accumulated to a higher degree in the parasite compared to muscle, intestine and stomach tissue of the host. The differences in accumulation patterns will be discussed with respect to differences in pollution levels of the host's habitats and also to differences in uptake mechanisms and metabolism of metals in mammalian and fish hosts.

Insights from chub – acanthocephalan system in polluted environments: Towards host decontamination?

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Few studies have investigated the accumulation and metabolisation of organic micropollutants in host-parasite systems despite their ubiquity, chronic release and potentially higher toxicity of their metabolites. The fate of organic pollutants in host-parasite systems was examined in a freshwater fish, the chub *Squalius cephalus*, naturally infected by acanthocephalan *Pomphorhynchus laevis*. Infected (N = 74) and uninfected fish (N = 55) were caught by electrofishing operations from the Marne River and its tributaries in September 2016. The occurrence and levels of several persistent micropollutant families (polychlorinated biphenyls (PCBs), organochlorine pesticides (OCPs), polybrominated diphenyl-ethers (PBDEs)) and metabolizable ones (polycyclic aromatic hydrocarbons (PAHs), phthalates, pyrethroid pesticides), as well as their metabolites, were characterized in parasites and various tissues of the fish (muscle, liver and stomach content). Infected chub were significantly less contaminated by OCPs ($P \leq 0.01$) but had higher levels of pyrethroid pesticides ($P \leq 0.05$) than uninfected fish. PAHs and phthalates reached higher levels in the acanthocephalans than in chub tissues ($[C_{[P]laevis} / C_{[chub\ tissue]}] > 1$). The highest concentrations of PAHs were detected in parasites (Median [Min ; Max], in ng.g⁻¹ of dry weight (dw), 1922 [875.64 ; 9779.30]), followed by the stomach content (591.80 [321.28 ; 1714.17]), liver (92.29 [0.34 ; 637.39]) and muscle (6.74 [0.89 ; 147.76]), with levels up to 5,000 times higher in parasites than in the host muscle. Our results suggest (1) a potential transfer of organic pollutants from the host to parasites, (2) a lower ability

for parasites to metabolize these compounds and/or (3) a higher sensitivity of parasites to the pollution compared to their fish host. Experimental validations will be conducted to confirm this host decontamination by acanthocephalans and the potential benefits for chubs, through the analysis of suitable biomarkers for a host health assessment (oxidative stress, immunity, body condition).

Application of acanthocephalans as bioindicators of metal exposure in the karst Krka River influenced by industrial and municipal wastewaters

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Naturally present trace metal concentrations in rivers are low, even extremely low in karst rivers, which make them very sensitive to the anthropogenic influence. The Krka River is a typical karst river in the Republic of Croatia whose lower part was proclaimed a National Park due to its exceptional natural beauty. The aim of the present study was to evaluate metal exposure in the Krka River few kilometres upstream of the park border, in the area threatened by industrial and municipal wastewaters, using bioindicator organisms: fish brown trout (*Salmo trutta* Linnaeus, 1758), gammarids (*Gammarus fossarum* Koch, 1836) and fish intestinal parasites, acanthocephalans (*Dentitruncus truttae* Sinzar, 1955). During the last few decades, acanthocephalans were recognized as potential indicators of metal exposure in the aquatic environment due to their effective metal accumulation, which is order of magnitude higher, compared to commonly used bioindicator organisms (Sures et al., 2017). Concentrations of 16 microelements and 4 macroelements were measured in digested fish intestine, gammarids and acanthocephalans from the Krka River source as a reference site and downstream with the wastewater impact. Prior to metal measurement by high resolution inductively coupled plasma-mass spectrometer (HR ICP-MS), samples were acid digested in a drying oven at 85°C for 3.5 hours using concentrated HNO₃ and H₂O₂. The prevalence of acanthocephalans in fish from the reference site was 100 % and from the polluted area 80 %, while the mean intensity of infection per fish was around 30 in both locations. More effective accumulation of most metals was found in acanthocephalans compared to fish and gammarids in both locations, especially of toxic Cd, Pb, Sr and Tl, in which average levels were 4-39 times higher in acanthocephalans. Metal concentrations were mostly higher in organisms from the wastewater impacted area, confirming anthropogenic influence and useful contribution of acanthocephalans as sensitive bioindicators of bioavailable metal levels.

References:

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Accumulation of heavy metals in fishes and its parasites from the Ružín water reservoir, biomonitoring study

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In this study, the actual content of 10 heavy metals was analysed in different organs of European perch (*Perca fluviatilis* L.) as well as its parasites (*Acanthocephalus lucii*, *Proteocephalus percae*) from the Ružín water reservoir. Heavy metals from reservoir sediments accumulated mostly in the kidneys (almost 30-times more than in fish muscles). Toxic lead was mostly concentrated in fish bones. In the muscles of perch, the only mercury exceeded the maximum permissible level (1.05 mg.kg⁻¹; limit 0.5 mg.kg⁻¹). Concerning toxic element accumulation, acanthocephalans, in comparison to the tapeworms and fish organs, absorbed them most intensively. The study was supported by VEGA project No. 2/0125/17.