Aus dem Fachbereich Wildtierkrankheiten des Leibniz Instituts für Zoo- und Wildtierforschung, Forschungsverbund Berlin e.V. eingereicht beim Fachbereich Veterinärmedizin der Freien Universität Berlin

PARASITES AND IMMUNITY IN ANTHROPOGENICALLY IMPACTED WATERFOWL POPULATIONS

Inaugural-Dissertation
zur Erlangung des Grades eines
Doktors der Veterinärmedizin
an der
Freien Universität Berlin

vorgelegt von **Hanna Prüter**Tierärztin aus Lüneburg

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GENERAL INTRODUCTION

Central European freshwater habitats, which provide breeding, roosting and molting habitats for different waterfowl species, are highly anthropogenically impacted ecosystems. Among others, human impact on freshwater habitats and its biodiversity include eutrophication, restructuring, environmental pollution and introduction of nonnative species (Meybeck 2003; Framing Committee of the Global Water System Project 2004; Strayer 2010; Vörösmarty et al. 2010). Particularly species biodiversity and quality of freshwater habitats are threatened by environmental pollutants originating from industry, agriculture and hunting. Additionally, native species biodiversity, community structures and ecosystem functioning may be threatened by the introduction of non-native species (Dudgeon et al. 2006; Carpenter et al. 2011).

Both environmental pollution and vertebrate invasion can interact with health indices of native populations such as parasite biodiversity, exposure to pollutants, parasite transmission and immunity. Environmental habitat pollution can result in acute or chronic intoxication of vertebrates. Such intoxication can result in a decrease of vertebrate immune competences, which subsequently facilitates disease outbreaks and parasite transmission (Franson 1986; Vallverdú-Coll et al. 2015b, a, 2016). Additionally, biodiversity loss may increase the transmission rate and risk of infectious diseases (Keesing et al. 2010). Moreover, parasite transmission originating from invasive species can be of high risk for the health of native species (Callaway and Ridenour 2004; Crowl et al. 2008; Pyšek and Richardson 2010). Thus, studying the epidemiological roles of native and invasive waterfowl species is needed to broaden the understanding of anthropogenic impact on health indices of specific species and species communities in freshwater habitats.

Out of the complex interplay of anthropogenic impact on freshwater ecosystems and health indices of freshwater species, parasite indices and immunity of two waterfowl species of the family Anatidae (waterfowl: ducks, geese and swans) were investigated. These species were chosen as they are characteristic species of freshwater habitats. Thus, an interplay between anthropogenic impacted fresh water habitat conditions and health indices such as parasite-host interactions, prevalence and abundance of parasites/infectious agents, immunity and disease transmission of Anatids (Figure 1) can be expected.

Out of the family Anatidae, the mallard (*Anas platyrhynchos*) as the most common native breeding duck and the most heavily hunted game bird in Germany was selected as study species (Gedeon et al. 2015; Deutscher Jagdverband e.V. 2018). Mallards have been shown to be affected by environmental pollution, predominantly caused by lead leading to both acute and chronic lead intoxication (Birkhead and Perrins 1985; Guitart et al. 1994; Mateo et al. 1997, 1998; Kelly and Kelly 2004; Degernes et al. 2006; Mateo 2009; Binkowski et al. 2013; Ferreyra et al. 2014, 2015). Moreover, mallards occur in all types of freshwater habitats including those of recreational use and are final hosts for bird schistosomes, whose cercaria might cause swimmer's itch in humans (Cort 1928).

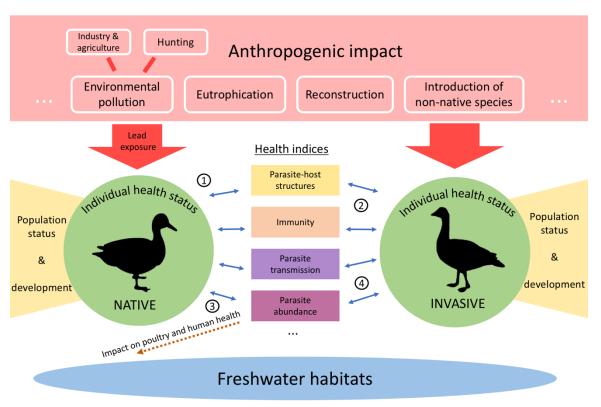


Figure 1: Schematic illustration of the interplay between anthropogenic impact and health indices of the two study species in freshwater habitats. Approaches of the thesis are highlighted with numbers of the different chapters (1-4).

Additionally, several species of the family Anatidae were introduced to non-native regions and successfully established as invasive populations (Duncan et al. 2003). Several health-related hypotheses have been formulated to characterize the interaction of health-indices of non-native species with the probability to become invasive for anthropogenically introduced species (Blossey and Notzold 1995; Torchin et al. 2003; Callaway and Ridenour 2004; Lee and Klasing 2004). The Egyptian goose (*Alopochen aegyptiacus*) was chosen as second study species as one of the most successful invasive

goose species in Europe (Gedeon et al. 2015) to investigate potential health related mechanisms behind the successful invasion of this species as well as its epidemiological role in the infection cycles of pathogens, which could be of relevance for native species, livestock and humans.

This thesis aims to highlight the impact of anthropogenic actions on health indices of waterfowl species from freshwater habitats. It evaluates the impact of environmental lead pollution and the introduction of non-native species on health indices of the study species in freshwater habitats. This research has the potential to broaden the general understanding of the complex interplay between humans and wildlife and consequently could be of relevance for nature conservation.

Thesis outline

The main body of this thesis consists of four chapters addressing different subjects related to health indices of both study species, which are either impacted by anthropogenic changes or may impact human and poultry health (Figure 1).

To study the **impact of anthropogenic changes to parasite and host community structures** in freshwater ecosystems, two main approaches were selected. First, to address the long-term effect of environmental lead pollution on parasite biodiversity in waterfowl, the chronic effect of lead burden on intestinal helminthes species richness and infection intensity in mallards were investigated (chapter 1). Second, health-related mechanisms, which might function as underlying mechanisms in the successful invasion of the Egyptian goose, were studied to test and potentially improve existing hypotheses in the field of animal invasion using the Egyptian goose as a model species. Therefore, parasite prevalence and variance of immune marker of Egyptian geese from Germany were compared to those of native Egyptian geese from Namibia to investigate potential differences in the light of the successful invasion of Europe by this species (chapter 2).

To study potential threats for native species, poultry and humans, originating from waterfowl, two additional approaches were selected. First, to gain knowledge on the prevalence and abundance of potentially zoonotic helminthic infections, neural bird schistosome infections in mallards, whose cercaria might cause swimmer's itch in humans, were studied. Thus, prevalence and abundance of neural bird schistosomes were investigated to study which species of neural bird schistosomes occur at which prevalence in mallards from German fresh waters (chapter 3). Second, parasites and selected

infectious agents of Egyptian geese in Germany were investigated to gain knowledge on their epidemiological role in transmission cycles of diseases relevant for native species and poultry. Additionally, the prevalence of these selected infectious agents was compared to published data from native geese species, to discuss them in the frame of contrasting ecology of the different geese species (chapter 2 and 4).

Key concepts

Key concepts, which are relevant to the studies presented in chapter 1-4, will be introduced in the following.

Life history theory and resource tradeoffs

Environmental factors, available resources, predators and parasites are important factors determining population growth rate and success of both native and invasive populations (Shea and Chesson 2002). Generally, resources are limited and must be allocated between different life history traits or stages, such as individual growth, dispersion, reproduction and immunity (Sheldon and Verhulst 1996; van der Most et al. 2011). Parasites, which compete for resources with the host, are energetically costly as they cause tissue damage or deprive the host from resources and induce immune reaction. Hosts with a high parasite burden will shift energetic resources into immunity. Thus, parasitic infection may decrease energetic investment of the host into other life history traits and therefore impact hosts population dynamics (Schmid-Hempel 2011).

Moreover, Atkinson, Thomas, and Hunter (2008) state that parasitism and its impact on the hosts should generally be considered in the context it occurs, including the impact of anthropogenic changes to the environment. According to the authors, wild birds are equipped with a suite of evolved traits, which enable them to successfully survive, reproduce and stay healthy under given environmental conditions. However, anthropogenic changes, which affect wild bird populations, may lead to an increased need for other trades, which might not have been developed and therefore are not available. Following this argumentation, Schlaepfer, Runge, and Sherman (2002) state that in changed habitat conditions (e.g caused by anthropogenic disturbance, altered environments) adaptations to the natural environment can function as "evolutionary

traps", meaning that naturally evolved trades may no longer be optimal but rather unfavorable for the hosts.

In summary, anthropogenic changes to the environment can be energetically costly for the host. Particularly, environmental pollutants can induce tissue damage and thus resources must be used for reparation directly. Moreover, pollutants may indirectly affect vertebrate hosts by altering parasite community structures and immune defenses of the host or directly affect parasite health (Rocke and Samuel 1991; Grasman and Scanlon 1995; Grasman 2002; Sures 2004; Sures et al. 2017). In addition, species invasion can impact resource tradeoffs as it may result in relatively lower parasitic burden (Enemy release, see below) for the invasive species. Consequently, resources can be allocated from immunity into reproduction and growth. Allocating resources from immunity to other physiological processes, such as reproduction, has been suggested to increase invasion success (Lee and Klasing 2004). Thus, studying parasite-host interactions in the frame of environmental pollution and species invasion allows to investigate resource tradeoffs induced by anthropogenic changes and threats to biological systems, which have been shown to be high in freshwater habitats (Vörösmarty et al. 2010).

Environmental lead pollution and its impact on host-parasite interactions (in birds)

Environmental pollution originating from industry, agriculture and hunting build up in freshwater habitats (Carpenter et al. 2011). Generally, high stability and persistence of pollutants can lead to severe contamination of both sediment and water bodies. Contamination entering food chains can be of risk for livestock and human health. Moreover, wildlife health including the health of plants, invertebrates and vertebrates, habitat quality and vertebrate biodiversity are threatened by environmental pollutants (Acevedo-Whitehouse and Duffus 2009).

One of the most prominent pollutants can be found among heavy metals. Especially, lead is a highly toxic heavy metal that is known to be toxic to living organisms including humans and animals, primary affecting the central nervous system, renal and hepatic function as well as the hematopoietic system. Lead is non-biodegradable and cannot be physiologically excreted from the body. In vertebrates, lead gets mainly stored in bones, which serve as long-time accumulation matrix (Beyer and Meador 2011; Flora et al. 2012).

Beside the negative impact of lead on human health also wildlife health is threatened by lead accumulating in the environment (Arnemo et al. 2016). Lead pollution of freshwater habitats originates from different anthropogenic sources such as mining, fishing using lead fishing weights and hunting. Lead based shot gun pellets used for waterfowl hunting form the main source of environmental lead pollution (Birkhead and Perrins 1985; Sears 1988; Scheuhammer and Norris 1995; Mateo et al. 1997, 1998; Meharg et al. 2002; Gómez-Ramírez et al. 2011; Newth et al. 2013; Ferreyra et al. 2014).

Environmental lead pollution leading to intoxications may impact host-parasite interactions by either directly affecting the parasites or the immune system and resource balance of its vertebrate hosts. Depending on the lead dosage, acute (lethal) or chronic (sublethal) symptoms can manifest in vertebrate hosts. In vertebrates, acute intoxication may result in severe clinical symptoms that include central nervous, renal and hepatic dysfunction directly leading to death (Sears 1988; Degernes et al. 2006; Flora et al. 2012; Newth et al. 2013), and thus not directly affecting host-parasite interactions. In contrast, sublethal dosages or chronic intake lead to sublethal effects of lead (Martinez-Haro et al. 2011; Vallverdú-Coll et al. 2016) with important fitness consequences for the vertebrate host. For instance decreased reproductive success and immunosuppression were found in vertebrates affected by chronic lead intoxication (Franson 1986; Trust et al. 1990; Rocke and Samuel 1991; Grasman and Scanlon 1995; Youssef et al. 1996; Singh et al. 2003; Eeva et al. 2005; Pikula et al. 2010; Vallverdú-Coll et al. 2015a, 2016).

Parasites can benefit from weakened immune function leading to an increase of parasite load in vertebrates (Franson 1986). However, parasites might also be directly poisoned by lead in the intestines of the hosts. Intestinal parasites of ducks are often exposed to lead as oral intake of hunting pellets is the main route of lead intoxication in these birds (Mateo et al. 1997; Ferreyra et al. 2014). Moreover, parasite related immune defenses might be enhanced by lead intoxication (Degen et al. 2005; Eeva et al. 2005; Gao et al. 2007; Cizauskas et al. 2014; Vallverdú-Coll et al. 2015b; Bertellotti et al. 2016). Additionally, intestinal helminths were found to accumulate lead in higher dosages than their vertebrate hosts. Thus, intestinal parasites may function as lead sink and reduce lead levels in vertebrate hosts (Sures et al. 2017).

In summary, the impact of anthropogenic induced environmental lead pollution on host-parasite interactions is highly complex. Here, this topic is addressed by studying chronic lead intoxication in mallards and its association with intestinal helminth parasite species richness and infection intensity (**chapter 1**).

Invasive vertebrate species and their health-related aspects relevant during invasion

Invasive species are major threats to native species biodiversity and health (Lövei 1997; McGeoch et al. 2010). With increasing globalisation leading to global biodiversity loss and climate change resulting in habitat changes, studying the mechanisms that promote species invasion is important for risk assessment, conservation and public health (Pyšek and Richardson 2010; Chown et al. 2015). Underlying mechanisms can be anthropogenically induced (such as for instance: number of introduced individuals) or arise from the traits of the introduced species, including its physiology, behaviour or pathogens.

Several health-related hypotheses have been formulated to study and understand species invasion and its impact on native species. The "Novel Weapon" hypothesis states that neozootic species bring novel pathogens to the novel range which harm native species (Callaway and Ridenour 2004). Beside this direct transmission of parasites from invasive to native species, parasites, infectious agents and host immune function of invaders have been considered to play important mechanistic roles during the process of vertebrate invasion. For instance the loss of parasites during invasion ("Enemy Release Hypothesis") and the ability to shift energetic resources away from immunity into reproduction and growth ("Evolution of Increased Competitive Ability") enable nonnative species to successfully invade new areas (Blossey and Notzold 1995; Torchin et al. 2003; Lee and Klasing 2004).

However, support for the above-mentioned hypotheses are non-consistent among studies of invasive vertebrates, which might be caused by the complexity of the vertebrate immune system as well as parasite-host interactions. Thus, studying parasites, pathogens and immunity of native and invasive Egyptian goose populations can provide new insights into the mechanisms of anthropogenic induced vertebrate invasion. In **chapter 2**, the Egyptian goose was used as a model species for vertebrate invasion and it was aimed to study the above mentioned major hypotheses in the field of invasion biology by investigating immunity and parasites of both a native and an invasive population. Moreover, it was aimed to combine existing hypotheses with the role of newly acquired parasites in the invasive range. Furthermore, in **chapter 4**, antibody seroprevalence against selected viruses and prevalence of selected bacteria in the invasive population of

Egyptian geese were studied to evaluate the role of this species in the transmission and maintenance of pathogens relevant for wild birds and poultry.

Avian helminth infections affecting humans-bird schistosomes as an example

Zoonotic diseases originating from waterfowl can be of risk for human health. Predominantly viral and bacterial infectious agents have been demonstrated to be transmitted from waterfowl to humans sometimes inducing severe clinical symptoms (Peiris et al. 2007; Benskin et al. 2009). One of the most intensively studied viral disease of this kind is the avian influenza virus, where waterfowl species function as reservoir hosts. These viruses can also be transmitted to humans and can result in severe symptoms in patients and occasionally death (Peiris et al. 2007).

Also avian helminth parasites can impact human health. Especially trematodes from the Schistosomatidae family, including bird schistosomes can cause diseases in humans (Kolárová et al. 2010; Soldánová et al. 2013). Human schistosomes of the genus *Schistosoma* (e.g. *Schistosoma mansoni, Schistosoma haematobium*) are agents of schistosomiasis, affecting humans in the subtropics and tropics. However, also in Europe, bird schistosomes of the genus *Trichobilharzia*, which cause trichobilharziosis in various bird species – especially species of the family Anatidae – with pathogenesis comparable to human schistosomiasis (Wojcinski et al. 1987), can affect human health. Cercaria of bird schistosomes may induce allergic dermatitis in humans ("swimmers itch"), which occurs after repeated cercaria penetration of the skin of sensitized individuals (Cort 1928). Humans engaged in recreational water activities (e.g. swimming, etc.) are most likely to be exposed to these cercariae (Chamot et al. 1998). "Swimmers itch" occurs worldwide with the exception of Antarctica (Horák et al. 2002) and is regarded as a reemerging disease in Europe (Horák and Kolářová 2011; Soldánová et al. 2013).

Due to their aquatic habitat, waterfowl of the family Anatidae are most likely the main reservoir species of bird schistosomes. However, only a few studies have examined the prevalence of neural bird schistosomes in waterfowl from Europe (Rudolfová et al. 2002, 2007; Jouet et al. 2008) and so far, *T. regenti* was the only neural bird schistosome species described in European birds. In **chapter 3** neural bird schistosome species prevalence in mallards from German freshwaters was studied.

Study species

Among vertebrates, water birds of the family Anatidae (ducks, geese and swans) are characteristic species of freshwater habitats, which have been shown to be affected by habitat pollution caused by factors such as lead intoxication (Birkhead and Perrins 1985; Guitart et al. 1994; Mateo et al. 1997, 1998; Kelly and Kelly 2004; Degernes et al. 2006; Mateo 2009; Binkowski et al. 2013; Ferreyra et al. 2014, 2015). Additionally, birds of the family Anatidae form one of the six major groups of invasive birds and were mainly introduced for hunting and exhibition purpose (Duncan et al. 2003).

Out of the family Anatidae, the mallard (*Anas platyrhynchos*) as the most common native breeding duck species and the Egyptian goose (*Alopochen aegyptiacus*) as one of the most successful invasive goose species in Europe (Gedeon et al. 2015) were chosen as study species. In Germany, both species share breeding, roosting and molting habitats. Additionally, both can be considered as synanthropic species, which often occur in anthropogenic impacted habitats in Germany.

Mallard (Anas platyrhynchos)

The mallard is the most common breeding duck species in Europe (Gedeon et al. 2015) and almost all races of the domestic duck are descended from the mallard (Herre and Röhrs 1990). Wild mallards are highly flexible in breeding habitat choice, which includes almost all freshwater water habitat types in Germany including highly anthropogenic impacted environments. The long-term population trend of breeding mallards in Germany is considered to be stable although it fluctuated in recent decades (Gedeon et al. 2015). European mallards are partial migratory. Recoveries of ringed individuals show that the autumn and winter population is composed of both German breeding birds and animals from mainly North-east and Eastern Europe (Bairlein, Dierschke, and Dierschke 2014).



Figure 2: Male (back) and female (front) mallard (Picture: Susanne Auls)

Mallard populations are impacted by hunting and its indirect effects of environmental lead pollution. The mallard is the most heavily hunted game duck in Germany (Deutscher Jagdverband e.V. 2018) and a general intake of lead from the environment in mallards leading to chronic lead intoxication is likely. Intake from the environment can occur via contamination of the sediments with old ammunition and other anthropogenic lead pollution of aquatic habitats. Furthermore, the mallard is a particularly suitable model species to study lead effects as the species can be immunosuppressed by chronic lead intake (Vallverdú-Coll et al. 2016). Moreover, mallards harbor a variety of parasitic helminth species belonging to all major helminth classes including bird schistosomes (Boch and Schneidawind 1988).

Egyptian goose (Alopochen aegyptiacus)

European neozootic Egyptian geese escaped from captivity in the late 19th century and became established as breeding population since the 1970s. Over the last decades, Egyptian geese in central Europe show rapid population growth (Wahl et al. 2011). In Germany, the population increased more than 30-fold over the last 15 years from 250-300 breeding pairs in 1999 to at least 8,000 pairs in 2013 (Arnold et al. 2013; Gedeon et al. 2015). Thus, among birds, the Egyptian goose is one of the most successful neozootic species to Europe (Bauer and Woog 2008). Moreover, in 2017 the Egyptian goose

became listed as invasive species in Germany according to "EU Regulation No 1143/2014 on the prevention and management of the introduction and spread of invasive alien species". This EU regulation "sets out rules to prevent, minimize and mitigate the adverse impact on biodiversity of the introduction and spread ... of invasive alien species" (Article 1) including the need for a scientific risk assessment (Nehring and Skowronek 2017).



Figure 3: Male (right) and female (left) Egyptian goose (Picture Manuela Merling de Chapa)

European Egyptian geese occur in a variety of freshwater habitats (Dietzen et al. 2015), often come into close contact with humans and frequently use anthropogenic structures (e.g. buildings, bridges, roofs) for nesting. Additionally, they often share their habitats with mallards, which they usually dominate by aggressive behavior (personal observations). Especially the Rhine and Mosel area (50.4°N, 7.6°E), which was chosen for field work, is a favored habitat in Germany. Here, Egyptian geese often use tutored lawns (e.g. turfs, parks) for grazing, where they are frequently fed with bred by humans. Those birds have rather low fear of humans and lawns are often riddled with their feces (personal observations). Thus, if Egyptian geese carry zoonotic infectious agents, pathogen transmission to humans is quite possible.

In their native range in Africa, Egyptian geese are hosts of parasite species from all major parasitic classes (trematodes, cestodes, nematodes, acanthocephalans) (Alexander and McLaughlin 1997). Additionally, they have been reported to carry avian

influenza A virus and avian avulavirus 3 (Shihmanter et al. 1998; Pfitzer et al. 2000; Thompson et al. 2008). Regarding avian influenza A virus, they were described as excellent sentinels having the highest total risk score compared to 15 other anatid species (regarding: range, abundance, mobility, shared roosts, mixed flocks, foraging and anthropogenic association) (Cumming et al. 2008; Abolnik et al. 2010). However, the epidemiological role of Egyptian geese in their invasive range in Europe is largely unknown.

Compared to its native range, the Egyptian goose faces fewer predators in novel European habitats (Gyimesi and Lensink 2010). Therefore, its population size can be expected to continuously increase in the next decades, raising the importance of gaining knowledge on the underlying mechanisms of their invasion success, their health status and on infectious agents and relevant pathogens they may carry.

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CHAPTER 1

Chronic lead intoxication decreases intestinal helminth species richness and infection intensity in mallards (*Anas platyrhynchos*)

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CHAPTER 2

The immunological plasticity hypothesis explains contradictory immunity pattern in invasive species

In preparation for submission

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Abstract

Several related hypotheses have been formulated to explain and understand the success of invasive species. Often, immunity and parasites have been linked to invasion success. While most of the related hypotheses are supported by empirical studies, there are contradictory results for some species. In many cases, hypotheses have been tailored for certain simplified scenarios. Here, we propose a new hypothesis (the Immunological Plasticity Hypothesis) which can explain contradictory findings and help to understand the role of parasites and immunity during vertebrate invasion. This hypothesis states that immune plasticity enables invaders to reduce the overall physiological cost of immunity while maintaining the ability to efficiently defend against novel parasites. Comparing a broad spectrum of immune effectors and parasite prevalences in an invasive and a native Egyptian goose population we find support for our hypothesis and discuss the implications for studying immunity and parasite infection during invasion processes.

Introduction

Invasive species are major threats to global biodiversity (Lövei 1997; McGeoch et al. 2010). Moreover, they may affect animal and public health by playing important epidemiological roles in spreading and maintaining several micro- and macro-parasites (from here on termed 'parasites') (Strauss, White, and Boots 2012). With increasing globalisation and biodiversity loss, studying the underlying mechanisms which promote species to become invasive, is essential for risk assessment, species conservation efforts and public health (Pyšek and Richardson 2010).

Although, the introduction of exotic, neozootic species to novel habitats is common, only few become true invaders. After introduction, neozootic species need to become established and spread to successfully invade a new region (Duncan, Blackburn, and Sol 2003). Environmental factors, available resources and natural enemies (i.e. predators and parasites) are important factors determining the population growth rate and invasion success of such species (Shea and Chesson 2002). Resources are typically limited and must be allocated between different life history traits or stages, such as individual growth, dispersion, reproduction and immunity (Sheldon and Verhulst 1996; van der Most et al. 2011). Allocating resources from immunity to other physiological processes, such as reproduction, has been suggested to increase invasion success (Lee and Klasing 2004).

Allocating resources away from immunity is especially beneficial when parasite burden is low. According to the "Enemy Release Hypothesis" (ERH) invading hosts show reduced parasite burdens by escaping the parasites in their native range when these are absent in the new range (Torchin et al. 2003). The ERH has support from studies on vertebrates including different bird species (Shwartz et al. 2009; Lima et al. 2010; Marzal et al. 2011; Valente et al. 2014; Clark et al. 2015; Ellis et al. 2017; Keogh et al. 2017; Diagne et al. 2016; Torchin et al. 2003). Generally, both parasite species richness and prevalence have been found to be lower in invasive than in native populations because specialist enemies are absent in the new region and host switching by specialist parasites of native species is rare (Keane and Crawley 2002; Torchin et al. 2003).

Based on the ERH, the "Evolution of Increased Competitive Ability Hypothesis" (EICA) states that invasive plants that escape from parasites can reduce the investment in defence mechanisms and thereby allocate more energetic resources to reproduction and dispersal (Blossey and Notzold 1995). Lee and Klasing (2004) refined this hypothesis (revised-EICA) focusing on the vertebrate immune system. They suggested that successful vertebrate invaders are more likely to decrease investment into energetically costly immune defences and compensate with immunity that incurs less energetic and pathological costs. Thus, down regulated inflammatory responses which are costly might lead to an increase in less costly responses (e.g. antibody-mediated immunity) (Lee and Klasing 2004).

In contrast to the revised-EICA, Cornet et al. (2016) suggest that newly acquired local parasites might induce an equivalent immune response in invasive and native populations. The authors show that only a subset of studies on invasive vertebrates support the revised-EICA, whereas others could not find evidence supporting the revised-EICA hypothesis. In a common garden experiment, Keogh et al. (2017) showed that invasive shore crabs (*Hemigrapsus sanguineus*) are less infected with rhizocephalan parasites than native crabs but at the same time more susceptible to experimental infection with these parasites. This indicates that reducing immune investment might increase the risk of getting infected by new parasites for invasive species. Also, Brown and Shine (2014) suggest that trade-offs between different immune effectors are more important for invasive species than the general downregulation of costly traits. Moreover, Møller and Cassey (2004) propose that invasive bird species with strong immune response towards novel parasites are potentially more likely to become successful invaders. Thus, several studies indicate that the revised-EICA might underestimate the impact of novel parasites on invasive populations.

The inconsistencies are addressed here by a new hypothesis that combines (1) the existing concepts underlying the enemy release and reduced immune investment in invasive populations with (2) the ability to defend against novel parasites in a new range. Our hypothesis posits that plasticity of immunity as opposed to generally reduced investment in immune function provides invading species with the necessary flexibility to colonize novel environments. Plasticity of immunity could enable invaders to balance reduced immune investment against their original parasites with increased defence against novel parasites. Thus, individuals which do not face novel parasites will reduce their immune investment, whereas individuals who face novel parasites will be able to increase the appropriate immune defense. Accordingly, we name this hypothesis "Immunological Plasticity Hypothesis" (IPH).

Depending on the immune effector, its main role in defence against parasites, and the stage of the invasion process we propose three different scenarios (Figure 1 b-d). If the enemy release effect is dominant (e.g. early stages of invasion; invasion front), immune investment would be down-regulated as suggested by the revised-EICA hypothesis (Figure 1b). In this case, low parasite prevalence and reduced overall costs of immunity can be expected. After introduction, individuals of invasive populations would increasingly face new parasites. Invasive animals must provide effective defences against these novel parasites. Here, low prevalence for some and higher prevalence for other parasite species (more heterogeneous than in the early stages) on a population level are likely. Thus, some individuals must be able to defend against the new parasites requiring higher immune investment than native individuals (Møller and Cassey 2004) whereas others may still invest less in immunity compared to their native relatives. Plasticity of immunity would be most prominent during this time and accordingly we would expect an increased variance in immune measures (Figure 1c). In later phases, prevalence of parasites might increase further and thus, defence against new parasites must increase. Immune effector levels would increase due to increasing need to defend against new parasites (Figure 1d).

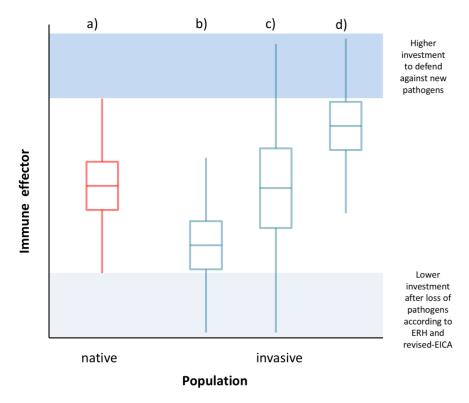


Figure 1: Predictive changes of the means and variances of immune effectors between native (a) and invasive (b-d) populations; (b) decrease in mean according to the "Enemy Release Hypothesis" (ERH) and "revised- Evolution of Increased Competitive Ability Hypothesis" (revised-EICA), (c) increase in variance if the effect of enemy release allows decreasing investment into immunity but defence against new parasites must increase as new parasites affect the hosts, (d) the effect of increased investment to defend against new parasites is stronger than the enemy release effect.

Based on the IPH, the prediction was that the differences in mean and variance between immune effectors of native and invasive Egyptian geese would depend on the specific pattern of loss and gain of parasites. If only parasite loss is observed, we predict a decrease in the means of high cost and a potential increase in low cost immune effectors as shown in Fig. 1b. If the invasive population is infected with new parasites, an increase of variance in immune effectors as in Fig 1c would be predicted. After having successfully established (later phase of invasion), mean values of specific immune effectors needed to defend against those new parasites can be expected to stay high (Fig 1d). To assess potentially congruent patterns, parasite communities and mean and variance of immune effectors of a native and an invasive population of the Egyptian goose (*Alopochen aegyptiacus*) were investigated. The Egyptian goose is one of the most successful invasive bird species in Europe, whose population continues to increase (Bauer and Woog 2008; Gyimesi and Lensink 2010) and which has been listed as an invasive species in Germany according to "EU Regulation No 1143/2014 on the prevention and management of

the introduction and spread of invasive alien species" and the BfN script 47 in 2017 (Nehring and Skowronek 2017).

Material and Methods

Sampling:

Parasite prevalence and immunity of Egyptian geese from a native population in Namibia were investigated and compared to those of a currently spreading invasive population of the same species in Germany. In both regions, geese were sampled during ringing procedures (live trapped) or dissected after hunting (necropsy).

Live trapping

Twenty-one Egyptian geese (9 male, 12 female) were live trapped in Namibia (22.35°S, 17.05°E) (native range) in February 2016. Additionally, data from a subset of 110 adult Egyptian geese (65 male, 45 female) investigated by Prüter et al. (2018a) were included in this comparative study. German geese were sampled in the Rhine and Mosel areas (50.4°N, 7.6°E) (invasive range) in 2015 (n=78) and 2016 (n=32) in different months (supplementary data Table S1). Sex and reproductive status were recorded. Reproductive status was defined as breeding (e.g. guiding gosling, showing territorial behavior with a partner, having an egg-laying active cloaca) or non-breeding (e.g. not fulfilling criteria of breeding and/or being part of a nonfamily-flock). All Namibian birds were non-breeding individuals. Blood was drawn from the vena metatarsalia plantaris superficialis using needles with a diameter of 0.06 mm for males and 0.04 mm for females. A fresh blood smear was prepared at capture and air dried. Blood samples kept at cool were centrifuged and sera were frozen in liquid nitrogen within eight hours after blood draw. Pharyngeal swabs were collected using sterile cotton swabs. Sera, blood cell pellets and pharyngeal swabs were kept frozen at -80°C till further analysis. Sampling in Germany was authorized by the Landesuntersuchungsamt Rheinland-Pfalz (G 15-20-005) and Landesamt für Natur, Umwelt und Verbraucherschutz Nordrhein-Westfalen (LANUV) (84-08.04.2015.A266). Permission to collect samples in Namibia was granted to GM and HK by the Ministry of Environment and Tourism (MET). Permission to export sample material from Namibia was granted by a MET export permit (No. 107513), and samples were transported to Germany in compliance with the Nagoya Protocol on Access to Genetic Resources.

Necropsy

Additionally to live trapping, twenty-six free ranging Egyptian geese (17 male, 9 female) hunted during the autumn/winter season 2014/2015 and 2015/2016 in the North and West of

Germany and twenty-seven Egyptian geese (11 male, 16 female), which were shot in February 2016 during regular pest control in Central Namibia were dissected. One of twenty-seven was live trapped and sampled before death and is thus included in both groups (live trapped and necropsy). Geese from Germany were kept frozen at -20°C after hunting until further analysis. Namibian geese were dissected immediately *post mortem*. During necropsy, ectoparasites, intestinal helminthes and nasal leeches were collected. Additionally, pharyngeal swabs were taken for molecular analyses.

Determination of parasites:

Both macro-parasites (ectoparasites, nasal leeches (Euhirundidae), intestinal helminthes) and selected micro-parasites (blood parasites (Haematozoa), bacteria, viruses) of Egyptian geese from the two populations were determined. Hereafter we use the term "parasites" combining macro- and micro-parasites and only explicitly distinguish between the type of parasites when differences can be expected and/or occur.

During necropsy, wing and breast feathers were macroscopically checked for the presence of ectoparasites. The upper beak was cut open and macroscopically investigated for the presence of nasal leeches. Intestinal helminthes were extracted from the intestine of the birds and were determined to the family level based on morphology. Additionally, blood smears of all live-trapped animals were investigated for the presence of blood parasites during immunological cell counts (Doster and Goater 1997).

To compare with bacterial prevalence of adult German Egyptian geese (included in Prüter et al. 2018a) (red shaded in Table 2)), the Namibian birds were screened for DNA from *Mycoplasma* spp. and *Riemerella (R.) anatipestifer* using conventional 16S rRNA-based PCR assays as described by Prüter et al. (2018a). To verify the specificity of the *Mycoplasma* PCR assay, products with a clear band were further investigated by sequence analysis, again following the procedure described by Prüter et al. (2018a). Only samples with a clear sequencing result were assessed as positive.

Serological methods were used to investigate the seroprevalence of antibodies (Ab) against Influenza A virus (IAV), *Avian avulavirus 1* (AAvV-1) and *West Nile virus* (WNV). For the detection of Ab against IAV, a commercial competitive enzyme linked immunosorbent assay (ELISA) was used following the manufacturer instructions (ID.vet, Grabels, France, Influenza A Antibody competition, FLUACA ver 0917DE).

A commercial competitive ELISA for detection of Ab against AAvV-1 (former Avian paramyxovirus 1; syn. Newcastle disease virus) was used and evaluated according to the

manufacturer protocol (ID.vet, Grabels, France, Newcastle Disease Competition, NDVC ver 0913 DE). Additionally, sera were tested applying a commercial competitive ELISA for Ab against Flaviviridae including WNV following the manufacture protocol (ID.vet, West Nile Competition, WNC ver 1014-1P DE).

Immunological assays:

Due to the complexity of the immune system, several eco-immunological tests were used to quantify both the cellular and humoral parts of the acquired and innate immune responses of Egyptian geese (Demas et al. 2011). Most of the methods are not species specific and have been used in a wide variety of free-living avian species, including different waterfowl (Matson et al. 2006; Giraudeau et al. 2010; Bourgeon et al. 2010). We quantified the levels of different humoral (natural antibodies, complement, lysozyme and haptoglobin) and cellular (monoctyes, heterophils, eosinophils and basophils) effectors of innate immunity. For adaptive immunity we measured the total immunoglobulin Y (IgY) concentration and the number of lymphocytes (Matson et al. 2006). Sample sizes (n) for each assay were dependent on the total amount of serum available from each individual and therefore differ between the tests (Table 1).

Table 1: Total sample sizes (n), sample sizes grouped by sex $(n \, \partial, n \, \varphi)$ and year of sampling of blood and serum samples from Namibian (native) and German (invasive) Egyptian geese (*Alopochen aegyptiacus*) for each immunological effector grouped by the costs of immunity (low costs vs. high cost according to (Klasing 2004; Lee and Klasing 2004))

Immunological effectors		Invasive 2015 (Germany)				Invasive 2016 (Germany)		Native 2016 (Namibia)
Low cost	n	$n \circlearrowleft, n \subsetneq$	n	$n \triangleleft, n \supsetneq$	n	$n \mathcal{O}, n \mathcal{O}$		
IgY	74	44♂, 30♀	26	16♂, 10♀	21	9♂, 12♀		
Lysozyme	76	43♂, 33♀	30	18♂, 12♀	20	9♂, 11♀		
Natural antibodies, complement	75	43♂, 32♀	24	16♂,8♀	21	9♂, 12♀		
High cost	n	Sex ratio	n	Sex ratio	n	Sex ratio		
Granulocytes (basophil, eosinophil, heterophil), Total leucocytes, Lymphocytes, Monocytes	77	45♂, 32♀	31	19♂, 12♀	21	9♂, 12♀		
Haptoglobin	72	42♂, 30♀	23	15♂, 8♀	21	9♂, 12♀		

Immunoglobulin Y

Total IgY, the avian equivalent to mammalian IgG, was assessed using a sensitive ELISA with commercial anti-chicken antibodies (Martinez et al. 2003; Bourgeon et al. 2010). 96-well high-binding ELISA plates (82.1581.200, Sarstedt) were coated with 100 µl of diluted serum sample (2 samples per bird 1:16000 diluted in carbonate–bicarbonate buffer) and incubated first for

1 h at 37°C and then overnight at 4°C. After incubation, the plates were washed with a 200 μ l solution of phosphate buffer saline and PBS–Tween, before 100 μ l of a solution of 1% gelatine in PBS–Tween was added. Plates were then incubated at 37°C for 1 h, washed with PBS–Tween and 100 μ l of polyclonal rabbit anti-chicken IgY conjugated with peroxidase (A-9046, Sigma) at 1:250 (v/v) was added. Following 2 h incubation at 37°C, the plates were washed again with PBS–Tween three times. After washing, 100 μ l of revealing solution [peroxide diluted 1:1000 in ABTS (2,20-azino-bis- (3-ethylbenzthiazoline-6-sulphonic acid))] was added, and the plates were incubated for 1 h at 37°C. The final absorbance was measured at 405 nm using a photometric microplate reader (μ Quant Microplate Spectrophotometer, Biotek) and subsequently defined as total serum IgY levels (Bourgeon and Raclot 2006).

Lysozyme

To measure lysozyme concentration in serum, we used the lysoplate assay (Giraudeau et al. 2010): 25 μl serum were inoculated in the test holes of a 1% Noble agar gel (A5431, Sigma) containing 50 mg/100 ml lyophilized *Micrococcus lysodeikticus* (M3770, Sigma), a bacteria which is particularly sensitive to lysozyme concentration. Crystalline hen egg white lysozyme (L6876, Sigma) (concentration: 1, 1.25, 2.5, 5, 6.25, 10, 12.5, 20 and 25 μg/ml) was used to prepare a standard curve for each plate. Plates were incubated at room temperature (25-27°C) for 20 h. During this period, as a result of bacterial lysis, a clear zone developed in the area of the gel surrounding the sample inoculation site. The diameters of the cleared zones are proportional to the log of the lysozyme concentration. This area was measured three times digitally using the software ImageJ (version 1.48, http://imagej.nih.gov/ij/) and the mean was converted to a semi-logarithmic plot into hen egg lysozyme equivalents (HEL equivalents, expressed in μg/mL) according to the standard curve (Rowe et al. 2013).

Haemolysis-haemagglutination assay

The levels of the natural antibodies and complement were assessed by using a haemolysis—haemagglutination assay as described by (Matson, Ricklefs, and Klasing 2005) adjusted to the limited volume of serum. After pipetting 15 µl of serum into the first two columns of a U-shaped 96-well microtitre plate, 15 µl sterile PBS were added to columns 2-12. The content of the second column wells was serially diluted (1:2) until the 11th column, resulting in a dilution series for each sample from 1/1 to 1/1024. The last column of the plate was used as negative controls, containing PBS only. We then added 15 µl of 1% rabbit red blood cells (supplied as 50% whole blood, 50% Alsever's solution, Envigo) suspension to all wells and incubated at 37°C for 90 min. After incubation, in order to increase the visualisation of agglutination, the plates were tilted at a 45° angle at room temperature. Agglutination and lysis, which reflect the

activity of the natural antibodies and the interaction between these antibodies and complement (Matson, Ricklefs, and Klasing 2005; Pap et al. 2010), was recorded after 20 and 90 min, respectively. Haemagglutination is characterised by the appearance of clumped red blood cells, as a result of antibodies binding multiple antigens, while during haemolysis, the red blood cells are destroyed. Titres of the natural antibodies and complement were given as the log2 of the reciprocal of the highest dilution of serum showing positive haemagglutination or lysis, respectively (Matson, Ricklefs, and Klasing 2005; Pap et al. 2015).

White blood cell counts

To count leucocytes, blood smears were prepared, air-dried and stained using Giemsa- and May-Grünwald staining. Smears were examined at 1,000×magnification with oil immersion and the relative number of different types of leucocytes was assessed by counting 100 leucocytes. The number of white blood cells of different types was expressed per 10⁴ erythrocytes (Pap et al. 2015).

Haptoglobin

We measured haptoglobin concentrations with a commercial kit (TP801, Tri-Delta Diagnostics, Inc.) following the instructions of the manufacturer. Haptoglobin concentrations (mg/ml) in undiluted serum samples were calculated according to the standard curve on each plate (Matson et al. 2006).

Statistical analyses:

Parasite prevalence

To investigate potential differences in the prevalence of parasites between native and invasive Egyptian geese, we used Fisher's exact tests.

Immunity

The means and variances of the different immune effectors were compared between the invasive and native Egyptian geese populations. To this end, we used linear mixed-effects models (LMMs). These models typically assume that the response variable is a function of a linear combination of some predictor variables, random effects, and a normally and independently distributed error. Importantly, the variance of this error (the so-called residual variance) is usually assumed to be constant. Thus, the estimated effects of the predictors only describe changes in the mean of the response variable, but not to the variance around that mean. Here, models were used in which the variance was allowed to be a linear function of some predictors. Thus, we were able to estimate simultaneously the effect of predictors upon the

variation in the mean of the response variable and also in the residual variation around that mean.

Different immune effectors were used as response variables (Table 1). As predictors for the mean sex (male vs. female), reproductive status (breeding vs. non-breeding) and invasion status (native vs. invasive) were included as fixed effects and month of sampling as a random effect. As predictors for the variance, we included invasion status (native vs. invasive) as the single predictor, which allowed us to test our prediction that the variance in immune effectors is higher among invasive compared to native individuals.

Some of the immune effectors were transformed (see tables supplementary data S2, S3, S4) to ensure normality of residuals. For haptoglobin we were not able to perform a transformation that ensured normality, because of the high proportion of values below the detection threshold. To account for this, we performed a general linear mixed model (GLMM) with a binominal error distribution and with a binary response variable (haptoglobin being either above or below the detection threshold). Thus, for haptoglobin we were only able to test for a change in mean but not for a change in variance. Total leucocytes were analysed followed by a separate analysis of differential white blood cell counts.

The LMMs and GLMMs were implemented using the R package *glmmTMB* version 0.2.0 (Brooks et al. 2017). Potential collinearity of predictors was tested by calculating variance inflation factors using the R package *car* version 2.1-6 (Fox and Weisberg 2011). All statistical analyses were performed using R version 3.3.2 (R Core Team 2016).

Results

Parasite prevalence:

Blood parasites and all groups of macro-parasites (ectoparasites, nasal leeches, intestinal cestodes and trematodes), except intestinal nematodes, were found at lower prevalence in the invasive than in the native population of Egyptian geese. However, none of the observed trends reached statistical significance (Table 2). DNA of the bacterium *R. anatipestifer*, which has a prevalence of 67.0% in adult invasive Egyptian geese from Germany (Prüter et al. 2018a) was not detected in the pharynx of native geese from Namibia. This difference in the prevalence was highly significant for *R. anatipestifer* (Table 2). *Mycoplasma* DNA., which was not detected in the German geese (Prüter et al. 2018a), was detected in the pharynx of two individuals from Namibia. However, the results did not reach statistical significance (Table 2). Seroprevalence of selected viruses (IAV, AAvV and WNV) were all lower in geese from the

invasive population but this effect was only statistically significant for antibodies against IAV (Table 2).

Immunity:

Males showed significantly higher levels of total leucocytes, haptoglobin abundance and lymphocytes than females. Total leucocytes and lymphocytes were significantly higher in birds that were in breeding status than in non-breeding individuals. No statistically significant effects on the other immune effectors were associated with sex or reproductive status (supplementary data Table S2, S3 and S4).

Differences in the mean immune function measures:

Of the four assays measuring 'low cost' immune effectors, only mean haemolysis concentration was significantly higher in the invasive population (Figure 2). The mean of total leucocytes ('high cost' immune effector) was not significantly different between the two study populations (Figure 2). Detailed analyses of the mean of differential white blood cells demonstrated significantly higher mean levels of heterophils and lymphocytes in the invasive population, whereas mean eosinophil concentration was significantly lower in this group. No statistically significant differences in mean monocytes and mean basophil numbers between the two groups were observed. In contrast, haptoglobin concentration (a 'high cost' immune effector) was significantly higher in abundance in the invasive (65%) than in the native (7.3%) population (Figure 3).

Differences in variance in immune function measures:

Variance of all 'low cost' immune effectors was significantly higher in the invasive population (Figure 2). The variance of total leucocytes did not significantly differ between the two groups (Figure 2). The variance in the concentration of eosionophils, heterophils and monocytes was significantly higher in the invasive geese. Variance in basophils and lymphocyte counts did not significantly differ between the groups (Figure 4).

Table 2: Results of the parasite screening and serology of Egyptian geese from Namibia and Germany (Prüter et al. 2018a). Total sample sizes (n), number of infected and non-infected individuals (Ratio) and prevalences (%) of macro parasites, bacteria and seroprevalences against selected viruses in the native Namibian and invasive German population of Egyptian geese (*Alopochen aegyptiacus*); Trend: \downarrow higher prevalence in the native than in the invasive population; \uparrow higher prevalence in the invasive than in the native population; Outcome of Fisher's exact test comparing prevalences of the two groups (p-value).

			Namibia Germany				Fisher test		
Parasitology:	Group	n	Ratio	Prevalence	n	Ratio	Prevalence	Trend	p
	Ectoparasites	27	11/16	40.74	26	3/23	11.54	\downarrow	0.08
	Euhirundidae	27	1/26	3.85	26	0/26	0	\downarrow	1
	Intestinal helminthes	27	6/21	22.2	26	4/22	15.38	\downarrow	0.74
	Cestoda	27	2/25	7.4	26	1/25	3.8	\downarrow	1
	Nematoda	27	0/27	0	26	1/25	3.8	↑	1
	Trematoda	27	4/23	14.8	26	1/25	3.8	\downarrow	0.36
	Heamatozoa	21	1/20	4.76	110	0/110	0	\downarrow	0.17
Bacteria:	Target genes	n	Ratio	Prevalence	n	Ratio	Prevalence	Trend	p
	Riemerella anatipestifer 16S rRNA gene	47	0/47	0	94	63/31	67.02	↑	< 0.001
	Mycoplasma spp. 16S rRNA gene	47	2/45	4.44	94	0/94	0	\downarrow	0.12
Serology:	Antigen	n	Ratio	Prevalence	n	Ratio	Prevalence	Trend	p
	IAV	21	9/12	42.86	105	9/96	8.57	↓	0.003
	AAvV-1	20	2/18	10	102	4/98	3.92	\downarrow	0.27
	WNV	13	1/12	7.69	56	0/56	0	\downarrow	0.2

Abbreviations: IAV= Influenza A virus; AAvV-1= Avian avulavirus 1; WNV= West Nile virus

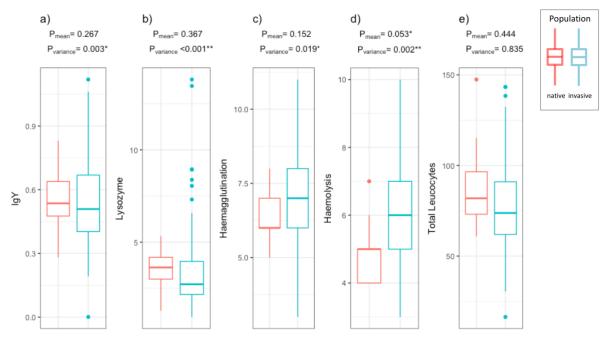


Figure 2: Differences in distributions of low cost (a-d) and high cost (e) immune measures between native and invasive Egyptian geese are shown; red= native; blue=invasive; P = p-values for the effects of population (native vs. invasive) on the mean (P_{mean}) and variance ($P_{variance}$) of the respective immune measure.

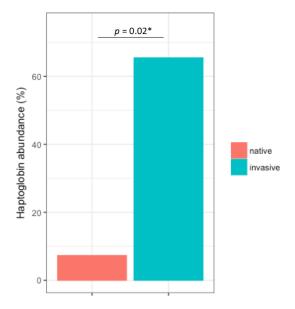


Figure 3: A barplot for the differences in haptoglobin abundance between native and invasive Egyptian geese is shown; red=native, blue=invasive; *p*-value for the effects of population (native vs. invasive) on the mean haptoglobin from the binomial distribution model. (It was not possible to fit a Gaussion model for haptoglobin concentration. Thus, the assessment of the difference in variance was not possible).

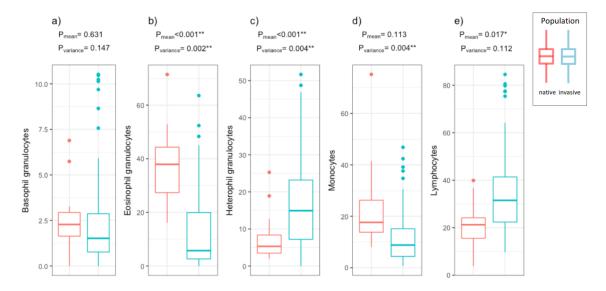


Figure 4: Differences in distributions of the differential white blood cells (a-e) between native and invasive Egyptian geese are shown; red= native; blue=invasive; P = p-values for the effects of population (native vs. invasive) on the mean (P_{mean}) and variance ($P_{variance}$) of the respective immune measure.

Discussion

Our results support the immune plasticity hypothesis (IPH). All but one of the studied parasite's prevalence or seroprevalence decreased when compared to native and invasive populations of Egyptian geese. Only the bacterium *R. anatipestifer* showed a significant increase in the invasive population, suggesting that this population faces novel pathogens during invasion. Thus, we predicted an increased variance of immune effectors, especially those of relevance for the defence against this novel pathogen, in the invasive compared to the native population (Figure 1c). As expected, the variance for six out of ten immune effectors in the invasive population of Egyptian geese was higher than in the native population. Moreover, five out of eleven immune effectors had higher mean values in the invasive population when compared to their native conspecifics. Most of these immune effectors are considered energetically costly (Lee and Klasing 2004), which is inconsistent with the predictions of the revised-EICA hypothesis. Still, the variance results are consistent with the predictions arising from the IPH demonstrating that existing hypotheses insufficiently covered the complex interplay of loss and gain of parasites in invasive populations.

Most studies of invasive vertebrate hosts and their pathogens focus on helminths, against which hosts respond with a Th2 response and eosinophils are one of the main immune

effectors (Al-Sabi et al. 2013; Romeo et al. 2014; Valente et al. 2014; Diagne et al. 2016). Thus, previous tests of the revised-EICA related predictions, which focused mainly on macroparasites, might be biased by the pathogens and immune effectors investigated. Here, low mean eosinophil granulocytes in invasive geese indicate a lower impact of macro-parasites on the immune system than in the native population (consistent with the ERH). However, variance of eosinophil granulocytes was significantly higher in the invasive than in the native population. The evidence suggests that Egyptian geese did not only lose parasites (ERH) but gained and are affected by novel ones. Thus, combining the results of mean and variance of immune marker has the potential to detect combined effects in immunological studies (IPH).

The R. anatipestifer prevalence, which was significantly higher in the invasive geese in Germany, indicated that invasive Egyptian geese encounter different pathogens in their new environments (Prüter et al. 2018a). R. anatipestifer is a bacterium of relevance for domestic ducks and geese, leading to severe clinical symptoms (Hubálek 2004; Hinz et al. 1998). The lack of clinical symptoms in the invasive Egyptian geese infected with R. anatipestifer may indicate that they are more tolerant than expected (Raberg, Graham, and Read 2009; Cornet et al. 2016) or that waterfowl may carry R. anatipestifer as a commensal bacterium (Ryll et al. 2001; Prüter et al. 2018a). Immune defences effective against micro-parasites, especially bacteria (heterophils, lymphocytes, haemolysis) were found to be higher on average in the invasive than in the native goose populations. Haptoglobin, which is an energetically costly acute phase protein with bacteriostatic function (Matson et al. 2006), was significantly more abundant in the invasive than in the native population of Egyptian geese, inconsistent with the EICA-related predictions. However, a significantly higher mean value of lymphocytes without evidence for higher variance in the invasive population indicate, that bacteria might be particularly important pathogens affecting this invasive population. Thus, immune defences against bacteria needs to be maintained in the invasive population. This is consistent with the predictions arising from the IPH (Fig. 1d).

The complexity of the immune system and the high specificity of different immune traits towards specific parasites may be more important than previously acknowledged (Morand et al. 2015). Furthermore, White and Perkins (2012) suggested that the higher the plasticity of the immune system of individuals of a founder population, the more likely the species becomes a successful invader (pre-selection). Additionally, Ghalambor et al. (2007) hypothesized that the period of persistence of invasive species (period after introduction and before rapid population growth starts) might be dependent on phenotypic plasticity. Thus, changes in immune

investment and immune plasticity in the different stages of invasion are likely (for the different phases of invasion see Duncan et al. 2003).

The Egyptian goose (*Alopochen aegyptiacus*) is a successful invasive bird species having and continuing to spread throughout Western Europe since the mid-20th century (Gyimesi and Lensink 2010). The impact of newly acquired pathogens on this population is most likely still ongoing and potentially increasing. This might explain why plasticity in immunity is prominent. The higher bactericidal investment but less plasticity in bacterial related immunity, might indicate that population growth rates can be expected to slow in the future, as immune investment increases. After the invasion has progressed further or completed, the variance of all effectors could decrease (Figure 1d). Thus, our findings indicate that amongindividual variation in immunity is on average higher in invasive compared to native Egyptian geese but that these effects depend on the parasites and therfore differ among immune effectors.

Compared to mallards (*Anas platyrhynchos*), a native species to Germany's avifauna, which shares the same habitat with the invasive Egyptian goose population, both native and invasive Egyptian geese from this study reveal lower parasite prevalence and load (Prüter et al. 2018b). Low parasite prevalence in the invasive population might indicate that the Egyptian goose is not a suitable host for parasites native to German waterfowl. However, this comparably low macro-parasite prevalence and richness in both populations might suggest that the Egyptian goose is highly immunocompetent and thus a prime water bird invasive species (Morand et al. 2015).

Contradictory effects of invasion on immune function have been also reported in amphibians. Immune defences are weaker in invasive cane toads (*Rhinella marina*) that move further distances indicating a trade-off between dispersal and immune investment (Brown and Shine 2014). Cane toads on the invasion-front were found to have higher bactericidal and phagocytic activity than in more established populations consistent with our observations in Egyptian geese (Brown et al. 2015). However, the overall immune investment in cane toads at the invasion front was lower compared to established populations (Llewellyn et al. 2012). In contrast, Cuban treefrogs (*Osteopilus septentrionalis*) at the invasion front were found to have a reduced bactericidal ability compared to frogs from more established areas (Goetz et al. 2018). These contradictory results from amphibians may reflect a complex mechanistic difference in immune investment of invading species. We suggest that studying plasticity of the immune system during range expansion of theses amphibians might provide further insights into the drivers of invasion of those species and might help to explain contradictory findings. Based on the IPH we expect that immune plasticity benefits individuals of invasive populations (1) in the

early stages of invasion by reducing investment in the immune system and (2) in later stages of invasion by providing effective defence against novel pathogens.

Invasive populations can have higher or lower genetic diversity than their native conspecifics (Roman and Darling 2007; Edelaar et al. 2015). Plasticity in immunity might either compensate for potentially lower genotypic variation (phenotypic plasticity) of invasive populations or for higher genetic diversity. Thus, the higher plasticity in immunity in the invasive population of this study may be caused by high phenotypic plasticity or potentially be an indicator of increased genetic variation. Studies of the genotypic diversity of the immune system (e.g. MHC and other immune genes) (Acevedo-Whitehouse and Cunningham 2006) of Egyptian geese from both populations could help to clarify the underlying mechanisms of the higher variance in immune effectors between the two populations in this study.

Intraspecific comparisons ('bio geographic' approach), (Roy and Lawson Handley 2012; Cornet et al. 2016) are essential for studying the IPH as interspecific comparisons of immune function are less robust (Matson et al. 2006). Common garden experiments combined with immune challenge or infection experiments would help to clarify the susceptibility to infection during range expansion of invasive species. Moreover, we recommend reanalysing data from studies on the revised-EICA by applying our approach of simultaneously investigating changes in mean – and in the variance – of immune markers. This approach could identify drivers of successful invasion and potentially predict population trends for neozootic species.

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Statement of authorship

GÁC, HP, KM and MF designed the study; HP, ST, JM, HK, ADG, KM, DL, GÁC coordinated different parts of the study; HP, ST and NB collected the data in Germany; HP, GM, JM, RP and GÁC collected the data in Namibia; HP, GvSH, KM, DL, GÁC performed laboratory analysis; HP and MF analysed the data; HP wrote the manuscript with the contributions of all the co-authors.

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Supplementary data

Table S1:

Month of compling	Status	Origin	non-b	reeding	breeding		
Month of sampling	Status	Origin	male	female	male	female	
April	invasive	Germany	1	0	7	4	
June	invasive	Germany	1	1	17	12	
July	invasive	Germany	3	0	11	11	
October	invasive	Germany	2	3	2	2	
November	invasive	Germany	1	0	0	0	
		2016					
February	native	Namibia	9	12	0	0	
June	invasive	Germany	1	0	12	9	
July	invasive	Germany	2	0	5	3	

Table S2: Summary output of the GLMMs to test for differences in the mean (in the conditional model) and variance (in the dispersion model) of

the respective "low cost" immune effector compared between native and invasive Egyptian geese

the respective low	cost immune effector con	iparcu octi	veen native a	ina mvas	sive Egyptian	geese				
	Effect on t	he mean				Effect on the variance				
Response	Predictor (fixed effects)	Estimate	Std. Error	Z	p		Estimate	Std. Error	Z	p
	Intercept	0.75001	0.02141	35.04	<2e-16 ***	Intercept	-3.8156	0.1412	-27.025	< 2e-16 ***
IgY	Status	0.04835	0.04357	1.11	0.2671	status	-1.0285	0.3457	-2.975	0.00293 **
181	Sex	-0.04258	0.02446	-1.74	0.0817.					
	reproduction	-0.04270	0.04169	-1.02	0.3057					
	Intercept	1.11349	0.07420	15.006	<2e-16 ***	Intercept	-1.1904	0.1368	-8.70	< 2e-16 ***
log(Lysozyme)	Status	0.14721	0.16312	0.903	0.367	status	-1.2431	0.3462	-3.59	0.00033 ***
log(Lysozyme)	Sex	-0.07812	0.08457	-0.924	0.356					
	reproduction	0.01005	0.15863	0.063	0.949					
	Intercept	7.2142	0.1927	37.44	<2e-16 ***	Intercept	0.5771	0.1414	4.080	4.49e-05 ***
Haemagglutination	Status	-0.5815	0.4062	-1.43	0.1523	status	-0.9183	0.3396	-2.704	0.00684 **
Tracinaggiumation	Sex	-0.3684	0.2204	-1.67	0.0947 .					
	reproduction	-0.2368	0.3857	-0.61	0.5393					
	Intercept	6.16166	0.17711	34.79	<2e-16 ***	Intercept	0.3872	0.1415	2.736	0.00622 **
Haemolysis	Status	-0.72516	0.37448	-1.94	0.0528 .	status	-0.7976	0.3407	-2.341	0.01922 *
Tracillorysis	Sex	-0.03806	0.20527	-0.19	0.8529					
	reproduction	-0.42019	0.35082	-1.20	0.2310					

Effect on the variance

0.13976

44.35 <2e-16 ***

-0.21 0.835

Estimate Std. Error

-0.07186 0.34425

Intercept 6.19793

status

Table S3Summary output of the GLMMs to test for differences in the mean and variance of the respective "high cost" immune effector compared

between native and invasive Egyptian geese

reproduction

Effect on the mean										
Response	Predictor (fixed effects)	Estimate	Std. Error	Z	p					
	Intercept	85.331	5.928	14.396	<2e-16 ***					
Total leucocytes	status	-9.760	12.762	-0.765	0.4444					
	sex	-11.555	3.993	-2.894	0.0038 **					
	reproduction	16.122	7.078	2.278	0.0227 *					
	Intercept	1.7129	0.4198	4.080	4.5e-05 ***					
log(hantaglahin hinamial)	status	-2.0296	0.8368	-2.425	0.0153 *					
log(haptoglobin binomial)	sex	-0.9708	0.4681	-2.074	0.0381 *					

0.2173

0.7166

0.303 | 0.7617

6	2
U	J

Table S4: Summary output of the GLMMs to test for differences in the mean and variance of the differential white blood cells compared between

native and invasive Egyptian geese

native and myasiv	Effect on the mean							Effect on the variance				
Response	Predictor (fixed effects)	Estimate	Std. Error	z	p		Estimate	Std. Error	Z	p		
	Intercept	1.42532	0.10786	13.214	<2e-16 ***	Intercept	-0.5348	0.1348	-3.966	7.3e-05 ***		
basophil	status	0.11548	0.24031	0.481	0.631	status	-0.4880	0.3368	-1.449	0.147		
granulocytes	sex	-0.19872	0.13009	-1.528	0.127							
	reproduction	-0.03004	0.21373	-0.141	0.888							
	Intercept	3.03294	0.24407	12.426	< 2e-16 ***	Intercept	1.1671	0.1350	8.645	< 2e-16 ***		
eosinophil	status	2.76064	0.52452	5.263	1.42e-07 ***	status	-1.0498	0.3394	-3.093	0.00198 **		
granulocytes	sex	-0.08153	0.28233	-0.289	0.773							
	reproduction	0.17518	0.50016	0.350	0.726							
	Intercept	3.9632	0.1946	20.362	< 2e-16 ***	Intercept	0.7067	0.1349	5.241	1.6e-07 ***		
sqrt(heterophil	status	-1.5726	0.4205	-3.740	0.000184 ***	status	-0.9608	0.3371	-2.850	0.00437 **		
granulocytes)	sex	-0.2066	0.2263	-0.913	0.361127							
	reproduction	0.2738	0.3974	0.689	0.490832							
	Intercept	39.444	4.819	8.184	2.74e-16 ***	Intercept	5.1786	0.1397	37.07	<2e-16 ***		
lymphoxytog	status	-25.537	10.672	-2.393	0.01672 *	status	-0.5684	0.3575	-1.59	0.112		
lymphocytes	sex	-8.089	2.351	-3.441	0.00058 ***							
	reproduction	11.680	4.135	2.825	0.00473 **							
	Intercept	2.16666	0.20471	10.584	<2e-16 ***	Intercept	-0.3884	0.1391	-2.792	0.00523 **		
log(monogytog)	status	0.68564	0.43262	1.585	0.113	status	-1.0521	0.3607	-2.917	0.00353 **		
log(monocytes)	sex	-0.04229	0.13565	-0.312	0.755							
	reproduction	0.12671	0.24652	0.514	0.607							

Table S5: Methods of the parasite screening and serology of Egyptian geese from Namibia and Germany.

Parasitology:	Group	Transmission mode	Method	Material
	Ectoparasites	direct	morphology	carcasses
	Euhirundidae	direct	morphology	carcasses
	Intestinal helminthes		morphology	carcasses
	Cestoda	vector	morphology	carcasses
	Nematoda	direct	morphology	carcasses
	Trematoda	vector	morphology	carcasses
	Heamatozoa	vector	morphology	blood smear
Bacteria:	Target genes	Transmission mode	Method	Material
	Riemerella anatipestifer 16S rRNA gene	direct	PCR	pharyngeal swab
	Mycoplasma spp. 16S rRNA gene	direct	PCR	pharyngeal swab
Serology:	Antigen	Transmission mode	Method	Material
	IAV	direct	ELISA	serum
	AAvV-1	direct	ELISA	serum
	WNV	vector	ELISA	serum

Abbreviations: IAV= Influenza A virus; AAvV-1= Avian avulavirus 1; WNV= West Nile virus; ELISA= Enzyme linked immunosorbent assay; PCR= Polymerase chain reaction

CHAPTER 3

Having bird schistosomes in mind the first detection of *Bilharziella polonica* (Kowalewski 1895) in the bird neural system

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CHAPTER 4

Sane and sound -A serologic and molecular survey for selected infectious agents in neozootic Egyptian geese (Alopochen aegyptiacus) in Germany

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GENERAL DISCUSSION

This thesis focuses on two anthropogenic impacted waterfowl species of freshwater habitats, the mallard, as the most common native duck species and the Egyptian goose, which is currently invading Europe. The anthropogenic driven changes to freshwater habitats/ecosystems, which were emphasized were environmental lead pollution (**chapter 1**) and the introduction of the Egyptian goose to Europe (**chapter 2 and 4**). Additionally, neural bird schistosomes in mallards from German freshwaters, whose cercaria might cause swimmer's itch, were investigated (**chapter 3**).

Impact of anthropogenic changes to parasite and host community structures

Using the example of intestinal helminth parasites in mallards, it could be shown that anthropogenic induced environmental lead pollution resulting in long term accumulation of lead in mallards, is related to significant loss of intestinal helminth species richness and infection intensity in mallards (**chapter 1**). Thus, studying intestinal helminths as sentinels for lead pollution in mallards, clearly highlights that lead intoxication not only directly affects waterfowl health (as it has been shown by others before) (Sears 1988; Degernes et al. 2006; Newth et al. 2013; Martinez-Haro, Green, and Mateo 2011; Vallverdú-Coll et al. 2016) but freshwater ecosystem species communities in a more complex manner than previously acknowledged.

Parasite-host systems are coevolved and balanced systems in which the parasites take away energetic resources from the host without running the risk to kill the host. External influences, which throw this parasite-host system out of balance, might lead to severe consequences for the vertebrate host by weakening its immune system and/or affecting its parasite lifecycles (Schmid-Hempel 2011). Here, it is shown that environmental lead pollution negatively impacts biodiversity indices of intestinal helminth parasites of wild mallards.

Moreover, intestinal helminth parasites may accumulate higher lead dosages than the vertebrate host and therefore function as lead sink (Sures et al. 2017). Hence, intestinal helminthes might protect their hosts from higher lead dosages. Results of **chapter 1** indicate that also in mallards, intestinal helminths might function as lead sink. Still, lower infection intensities in ducks showing higher lead levels in bones indicate

susceptibility towards lead also in those parasite species. Consequently, intestinal helminthes, which accumulate lead are presumably threatened by excessive chronic lead intake.

Summarizing, the results of **chapter 1** underline the importance of studying parasite-host interactions in the frame of anthropogenic changes to the environment. Beside the health of vertebrate hosts, parasite biodiversity is a subject of conservation concern (Gómez and Nichols 2013). Vice versa, imbalanced parasite-host interactions might bare health risks for vertebrate hosts. Hence, it can be recommended to generally lower anthropogenic induced environmental pollution to preserve healthy species communities in German freshwater habitats.

Moreover and beside the indirect impact of environmental lead on parasite species richness and infection intensity, this thesis shows that mallards from German fresh waters show comparably high chronic lead burden (Ferreyra et al. 2014, 2015). This finding indicates that lead pollution in German freshwater habitats is potentially still high although the use of lead based ammunition for hunting waterfowl is prohibited in most of the German federal states (JWMG, n.d.; BayJG 1978; LJG-NRW 1994; NJagdG 2001; BbgJagdDV 2004; LJG 2010). Still, due to their partly migratory behavior, lead intoxication in mallards could also be caused by high lead burden of freshwater habitats in other European countries. Nevertheless, European studies on lead burden of waterfowl indicate that the compliance of hunters to use lead-free ammunition only increased after enforcement and vigilance of rangers was intensified (Mateo et al., 2013). Since the regulation on the use of lead-free ammunition to hunt waterfowl is not consistent within Germany, a standardization of law and regular controls could help to lower lead burden in freshwater habitats and consequently reduce the risk of species biodiversity losses.

By studying parasites and immunity of native and invasive Egyptian goose a new hypothesis ("Immunological plasticity hypothesis") was developed, which integrates the role of newly acquired pathogens into existing hypotheses on species invasions. In support of the novel hypothesis the variance of immune marker is significantly higher in the invasive than in the native population of Egyptian goose. Thus, the invasive Egyptian goose population in Germany seems to be both able to reduce investment into immunity if possible (at low parasite burden) and defend against newly acquired parasites when necessary. This flexibility in immune investment might be the fundament of the successful invasion of the Egyptian goose to Europe and is potentially a general mechanism helping non-native species to become invasive.

Potential threats for native species, poultry and humans

Generally, birds can function as vectors for infectious diseases which can affect livestock and are potentially of zoonotic risk. For instance, cercaria of bird schistosomes, including *T. regenti*, whose final hosts are water birds, may cause cercarial dermatitis (swimmer's itch) in humans, which is regarded as a re-emerging disease (Horák and Kolářová 2011; Soldánová et al. 2013). Particularly climate change was shown to increase the risk of re-emerging diseases. Climate change driven changes in behavior traits of migratory birds, shifts in seasonal or temperature dependent processes of intermediate hosts (snails) and the frequency of transmission and intensity of infection, are potential risk factors regarding the dispersal of bird schistosomes to new regions and therefore increases the availability in wetland habitats (Horák and Kolárová, 2011).

In chapter 3, a 21% prevalence of Trichobilharzia regenti, a neural bird schistosome, was found in native mallards from German freshwaters. This result is consistent with previous studies in water birds, which reported prevalences of T. regenti ranging from 24 to 74.5% in different definitive bird hosts (Kolářová, Skirnisson, and Horák 1999; Rudolfová, Sitko, and Horák 2002; Jouet et al. 2008). Moreover, Bilharziella polonica was detected either between the brain membranes (meninges) and the brain, in the spinal cord or in the intestine of 12% of the mallards, which is the first description of this species from the birds neural system. The presence of B. polonica in the birds neural system indicates that this species has a broader spectrum of target organs in the definitive host than previously described and that the spectrum of neural bird schistosome species is broader than previously acknowledged. Thus, for B. polonica, parasite-host interaction is of higher complexity than previously known, which should be further investigated for different definitive host species as well as for potential accidental hosts. Monitoring bird schistosomes in vertebrate hosts in the light of anthropogenic changes to ecosystems (such as climate change, pollution, species invasion) should be considered to increase knowledge on this particular parasite-host system in the view of the fact that swimmer's itch is re-emerging in Europe.

In **chapter 4**, it was shown that Egyptian geese are frequent carriers of *Riemerella* anatipestifer and furthermore provided serological evidence of exposure to selected viral pathogens of relevance for native birds and poultry. This finding indicates that invasive Egyptian goose are involved in native pathogen transmission cycles and therefore might

potentially spillover and spillback pathogens to native species and poultry. Additionally, in **chapter 4**, seroprevalence of viral pathogens were compared between Egyptian geese and published results from studies on other goose species (Kruckenberg et al. 2011; Kistler et al. 2012; Brown et al. 2010; Hlinak et al. 1998; Bradshaw and Trainer 1966; Bönner et al. 2003). Reported differences between different goose species are likely driven by season or ecological differences between the species. Thus, future research of pathogen transmission cycles among wild birds and their epidemiological roles should include both the role of native and invasive species in the maintenance and spread of pathogens. This will help to assess potential changes to health indices of native species (e.g. parasite-host interactions, parasite/pathogen abundance and disease transmission) caused or promoted by invasive species.

Outlook

Freshwater species biodiversity is threatened by human activities and anthropogenic impact on a global scale (Vörösmarty et al. 2010). Waterfowl health and population dynamics have been found to be negatively impacted by climate change, industrialization of agriculture, increasing recreational use of water habitats, invasion of species, intensification of ship traffic, aridification of wetlands and hunting (Wahl et al., 2011). Increasing knowledge on waterfowl health, their epidemiological roles, threats and parasites can help to assess those threats in terms of biodiversity loss in freshwater habitats. Particularly the results of **chapter 1**, which highlight the negative impact of lead pollution on parasite species richness in freshwater ecosystems, are applicable to implement and evaluate strategies of international wetland protection. Thus, reducing lead pollution in freshwater ecosystems could help to protect species biodiversity in German freshwater habitats.

Invasive species can significantly harm native species and biodiversity (Lövei 1997; McGeoch et al. 2010). Especially freshwater ecosystems are vulnerable and threatened by the impact of invasive species (Strayer 2010). Thus, monitoring of non-native species and their potential to become invasive is of great importance for biodiversity conservation especially in freshwater ecosystems. Moreover, invasive species may affect animal and public health by playing important epidemiological roles in spreading and maintaining several micro- and macro-parasites (Strauss, White, and Boots 2012). Thus, studying health related aspects in the frame of species invasion may help to

reduce negative impact of invasive species on native biodiversity. Predictions arising from the "Immunological Plasticity Hypothesis", proposed in **chapter 2**, should be tested in studies on invasive vertebrates to investigate the predictive power of variability in immune markers for population trends of non-native species including the probability to become invasive.

In Europe, based on §5 Regulation (EU) No 1143/2014, a risk assessment of invasive species conducted by each European Union member state is mandatory. According to the formulation of Article 5, this risk assessment should be based on available scientific knowledge. Regarding this European legislation, the results from **chapter 4**, which indicate that German Egyptian geese are involved in transmission cycles of infectious agents relevant for wild birds and poultry, can be a basis for the evaluation of its health impact on native species (Article 5 f). Moreover, the new hypothesis proposed in **chapter 2** may help to create projections of likely future distributions of invasive species (§5 e) in general by studying variance of immune markers in the different phases of invasion.

In conclusion, this thesis highlights the importance to encounter the complex interplay between health indices of waterfowl and anthropogenic impacted environmental factors such as pollution and species invasion to studies on habitat biodiversity and wildlife health. Moreover, it provides one of the very few studies on infectious agents and immunity of invasive birds and raises a new hypothesis in the field of vertebrate invasion, which will help to study health related drivers of invasion processes.

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ÜBER DIE ROLLE VON PARASITEN UND DES IMMUNSYSTEMS IN ANTHROPOGEN BEEINFLUSSTEN WASSERVOGELPOPULATIONEN

ZUSAMMENFASSUNG

Der menschliche Einfluss auf Süßwasserökosysteme ist immens. Die vorliegende Arbeit befasst sich mit den Wechselwirkungen zwischen menschlichen Einflüssen auf Süßwasserökosysteme und Parasiten sowie die Immunsysteme der dort lebenden Wasservögel. Hierfür wurden Parasitenindizes und Immunparameter von Stockenten als die häufigste heimische Entenart und der Nilgans als eine der häufigsten invasiven Gänsearten in Deutschland untersucht.

In Kapitel 1 wurde untersucht, welchen Einfluss chronische Bleibelastung bei Stockenten auf deren Parasitenbiodiversität hat. Dazu wurden Stockenten hinsichtlich chronischer Bleibelastung sowie Artenvielfalt und Infektionsintensität von Magendarmhelminthen untersucht. Sowohl Artenvielfalt als auch Infektionsintensität waren signifikant niedriger in Enten mit höherer chronischer Bleibelastung. Dieses Ergebnis deutet darauf hin, dass nicht nur die Ente als Wirt, sondern auch die Parasiten selbst durch Blei negativ beeinflusst werden. Dies zeigt, dass eine menschlich verursachte Bleibelastung der Umwelt, welche zu Langzeitbleibelastungen bei Wasservögeln führt, einen signifikanten Biodiversitätsverlust von Magendarmhelminthen zur Folge haben kann

In **Kapitel 2** wurden Parasitenprävalenzen und Immunparameter von einer invasiven und einer heimischen Nilganspopulation verglichen. Hierbei konnte gezeigt werden, dass die Varianzen der Immunparameter der invasiven Population signifikant höher sind als die der heimischen Population. Zudem wird eine neue Hypothese ("Immunological Plasticity Hypothesis") beschrieben, die bereits bestehende Hypothesen zum Einfluss von Parasiten und Immunparameter auf Mechanismen der Invasionsbiologie verbindet und um den Einfluss neuerlich erworbener Parasiten erweitert. Diese neue Hypothese kann helfen, gesundheitsbezogene Antriebe von Invasionsprozessen besser zu verstehen.

In **Kapitel 3** wurden Stockenten hinsichtlich des Auftretens und der Prävalenz neuronaler Vogelschistosomen untersucht, deren Zerkarien beim Menschen Badedermatitis auslösen können. Mit einer Prävalenz von 21% war *Trichobilharzia*

regenti die häufigste Vogelschistosomenart in den Stockenten. Neben *T. regenti* konnte *Bilharziella polonica* aus verschiedenen Bereichen des zentralen Nervensystems von 12% der Stockenten extrahiert werden. Diese Entdeckung ist die erste Beschreibung *B. polonicas* im Nervensystem von Vögeln und lässt die Schlussfolgerung zu, dass das Spektrum neuronaler Vogelschistosomen größer ist als bisher angenommen. Beide Ergebnisse unterstreichen die Wichtigkeit für ein Monitoring von Vogelschistosomen in Wirbeltierwirten.

In **Kapitel 4** konnte gezeigt werden, dass Nilgänse in Deutschland Träger von *Riemerella anatipestifer* sind. Zusätzlich deutet das Auftreten von Antikörpern gegen Pathogene, die von Bedeutung für heimische Vögel und Hausgeflügel sind (Influenzavirus A, *Aviäres Avulavirus 1*, Aviadenoviren, *Enten-Atadenovirus A* (syn.: egg drop syndrome 1976 virus), auf einen Kontakt mit diesen Erregern hin. Diese Ergebnisse zeigen, dass invasive Nilgänse in Deutschland in Zyklen heimischer Krankheitserreger eingebunden sind und diese Art möglicherweise als Überträger der untersuchten Erreger fungieren kann.

Mit dieser Dissertation wird deutlich, dass das komplexe Zusammenspiel zwischen menschlichen Einflüssen auf die Umwelt und Gesundheitsaspekten von Wasservögeln in Studien zur Artenvielfalt in Süßwasserökosystemen einbezogen werden sollten. Zusätzlich wurde eine neue Hypothese vorgestellt, anhand derer zukünftige Studien die gesundheitsbezogenen Mechanismen der Invasion von Wirbeltieren besser beurteilen können und die helfen kann, das Feld der Invasionsbiologie generell weiter zu entwickeln.

PARASITES AND IMMUNITY IN ANTHROPOGENICALLY IMPACTED WATERFOWL POPULATIONS

SUMMARY

This thesis aimed to investigate interactions between human driven ecosystem changes and waterfowl species in the frame of immunity and parasitic infections. This interplay was investigated studying parasite indices and immunity of the mallards and Egyptian geese in the context of lead pollution in mallards, helminthic parasites of zoonotic potential in mallards, the role of immunity and parasites in the invasion process of Egyptian goose and infectious agents of Egyptian goose which could be of relevance for native species, livestock and humans.

In **chapter 1**, it was studied how lead pollution affects parasite diversity by investigating intestinal helminth species richness and infection intensity in mallards exposed to environmental lead. Parasite species richness and infection intensity was found to be significantly lower in birds with higher chronic lead levels suggesting both host and parasites respond to lead exposure. Thus, it was shown that anthropogenic induced environmental lead pollution resulting in long term intoxication of mallards, is related to significant biodiversity loss in intestinal helminth species communities of mallards.

In **chapter 2**, parasite prevalence and immunity of an invasive and a native Egyptian goose populations were compared, showing that the variance of immune marker is significantly higher in the invasive than in the native population. Moreover, a new hypothesis ("Immunological Plasticity Hypothesis") was raised in the field of vertebrate invasion, which aims to combine existing hypothesis with the impact of newly acquired pathogens on invasive species. The new hypothesis will help to study health related drivers of invasion processes.

In **chapter 3**, prevalence of neural bird schistosomes, whose cercaria might cause swimmer's itch in humans, in mallards was investigated. Showing a prevalence of 21%, *Trichobilharzia regenti* was the most prevalent neural bird schistosome. Beside *T. regenti*, *Bilharziella polonica* was detected either between the brain membranes (meninges) and the brain, in the spinal cord or in the intestine of 12% of the mallards, which is the first description of this species from the birds neural system. This finding

indicates that the spectrum of neural bird schistosome species is broader than previously acknowledged. Both findings highlight the importance of monitoring this parasitic group in vertebrate hosts.

Chapter 4 shows that Egyptian geese are frequent carriers of *Riemerella* anatipestifer and furthermore provides serological evidence of exposure to selected viral pathogens of relevance for native birds and poultry. Thus, invasive Egyptian geese are involved in native pathogen transmission cycles and therefore might potentially spillover and spillback pathogens to native species and poultry.

In conclusion, this thesis highlights the importance to encounter the complex interplay between health indices of waterfowl and anthropogenic impacted environmental factors such as pollution and species invasion to studies on species biodiversity in freshwater ecosystems. Moreover, it provides one of the very few studies on infectious agents and immunity of invasive birds and proposes a new hypothesis in the field of vertebrate invasion, which will help to study health related drivers of animal invasion.

PUBLIKATIONSVERZEICHNIS

Scientific publications:

Prüter H, Sitko J, Krone O (2016): Having bird schistosomes in mind- The first detection of *Bilharziella polonica* (Kowalewski 1895) in the bird neural system. *Parasitology Research*, doi: 10.1007/s00436-016-5359-9

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Scientific presentations:

2015, Talk, Poster: Impact of invasive bird species on ecosystems in Europe and its human drivers, Endoparasites and immunity of native and invasive waterfowl in Germany, 6th European Wildlife Disease Association student Workshop, Verier-du-lac, France

2015, Poster: Endoparasites and immunity of native and invasive waterfowl in Germany, 10th International Conference on Behaviour, Physiology and Genetics of Wildlife, Berlin, Germany

2015, Talk: Endoparasitenfauna und Immunfunktion bei heimischen und invasiven Wasservögeln im Vergleich, Deutsche Ornithologen-Gesellschaft (DO-G) 148. annual meeting – 3rd price of young scientist talk, Konstanz, Germany

2015, Invited speaker: Egyptian goose (*Alopochen aegyptiacus*) – a new European citizen, Scientific society Windhoek, Namibia

2015, Talk: Abundance of nasal bird schistosomes in Mallards from German wetlands, Tagung der DVG-Fachgruppe "Parasitologie und parasitäre Krankheiten", Berlin, Germany

2016, Talk, poster: Immune function of the Egyptian goose (*Alopochen aegyptiacus*) – a highly successful neozootic waterbird species in Europe, Serological survey of selected viral pathogens in Egyptian geese (*Alopochen aegyptiacus*) from Germany, 12th Conference of the European Wildlife Disease Association (EWDA), Berlin, Germany

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SELBSTSTÄNDIGKEITSERKLÄRUNG

Hiermit bestätige ich, dass ich die vorliegende Arbeit selbständig angefertigt habe. Ich versichere, dass ich ausschließlich die angegebenen Quellen und Hilfen in Anspruch genommen habe.

Berlin, den 25.06.2019

Hanna Prüter

