1 Large female northern pike (Esox lucius) do not connect spawning areas across a lagoon network in 2 the southern Baltic Sea Lukyanova Olga<sup>1,2</sup>, Arlinghaus Robert<sup>1,4</sup> & Dhellemmes Félicie<sup>1,3,5</sup> 3 4 <sup>1</sup>Department of Fish Biology, Fisheries and Aquaculture, Leibniz Institute for Freshwater Ecology and 5 *Inland Fisheries, Berlin, Germany* 6 <sup>2</sup>AZTI, Marine Research, Basque Research and Technology Alliance (BRTA), Pasaia, Spain 7 <sup>3</sup>Excellence Cluster "Science of Intelligence", Berlin, Germany 8 <sup>4</sup>Division of Integrative Fisheries Management, Faculty of Life Sciences and Integrative Research 9 Institute on Transformations of Human-Environment Systems, Humboldt Universität zu Berlin, Berlin, 10 Germany 11 <sup>5</sup>Center for Adaptive Rationality, Max Planck Institute for Human Development, Berlin, Germany 12 13 **Corresponding author:** 14 Félicie Dhellemmes 15 Science of Intelligence, Marchstraße 23, 10587 Berlin, Germany felicie.dhellemmes@gmail.com 16 17 18 **Funding:** 19 European Maritime and Fisheries Fund and the State of Mecklenburg-Vorpommern (grant numbers MV-I.18-LM-004 and B730117000069; BODDENHECHT) 20 21 22

#### Abstract

Large individuals may serve as keystone connectors, a role recently demonstrated in Atlantic cod (*Gadus morhua*). To examine whether this pattern extends to other coastal fish species, we analysed capture-mark-recapture data for 666 individuals out of 4597 tagged coastal northern pike (*Esox lucius*) and acoustic tracking data from 318 individuals in the southern Baltic Sea, with total lengths of the individuals ranging from 28 cm to 126 cm. Neither mark-recapture nor telemetry data revealed a relationship between individual body length and sex, distance between capture and recapture, connectivity, maximum horizontal displacement, and among-year spawning site fidelity. Instead, connectivity and movement ranges were correlated between years and repeatable, suggesting consistent inter-individual variation unrelated to body length. These findings suggest that large pike do not serve as keystone connectors, likely due to their reproductive biology as total spawners. In total spawners, spatial bet hedging might be realised through mechanisms other than the use of variable spawning sites by individuals within a season, including ecotype evolution of different spawning phenotypes and variable spawning times or locations across years.

# Keywords

mark-recapture, acoustic telemetry, BOFFFF, site connectivity, spawning

#### Introduction

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The relevance of exceptionally large fish in the conservation of fish populations has gained substantial attention over the past two decades, particularly following the paper by Berkeley et al., (2004) on the positive effect of maternal age on larvae survival in rockfish (Sebastes melanops) (Birkeland & Dayton, 2005). This research interest has culminated in the catchy concept of BOFFFF big old fat fecund female fish – and the supposed relevance of large and old fish for population dynamics and stock recovery (Hixon et al., 2014). Major syntheses have outlined the various mechanisms by which large fish contribute to offspring production, recruitment, and population stability (e.g., Barneche et al., 2018; Hixon et al., 2014; Hsieh et al., 2010a; Kopf et al., 2024; Marshall et al., 2021). Consequently, stock assessment methods and harvest management strategies are increasingly incorporating measures to conserve large and old fish (Froese, 2004; Griffiths et al., 2024), with growing interest in approaches that protect both small, immature individuals and highly fecund large fish, e.g., through harvest slots (Ahrens et al., 2020; Gwinn et al., 2015) or marine protected areas (Marshall et al., 2021). The discussion on the relevance of large fish in population dynamics and conservation often centres on two key size-dependent maternal effects: (1) the influence of female size on egg and offspring phenotype and quality (e.g., Berkeley et al., 2004), and (2) size-dependent increases in relative fecundity per female mass, characterized by hyperallometric relationships between body mass and egg number (e.g., Barneche et al., 2018; Hixon et al., 2014a). However, the size-dependent maternal effects on egg and offspring quality regularly reported in laboratory trials (e.g., Berkeley et al., 2004; Hixon et al., 2014) may not manifest in the wild and may appear only in studies where environmental influences are controlled (Marshall et al., 2010). Further, population-level fish recruitment is under strong density-dependent feedback in fishes, potentially diminishing the relevance of spawning stock composition and the importance of size-dependent effects for population dynamics and fisheries yield (Ahrens et al., 2020; Cooper et al., 2013). Similarly, the fecundity advantage of large fish may not be generally impactful for the population as a whole, as large fish are typically in low abundance

(Andersen et al., 2019), unless the hyperallometry of the mass-fecundity relationship is exceptionally strong (Ahrens et al., 2020; Marshall et al., 2021). Given that exceptionally large fish are usually rare in most stocks relative to younger and hence smaller spawners, and the degree of hyperallometry in mass-fecundity varies across species and populations (Barneche et al., 2018), the conservation value of protecting large fish similarly varies among fisheries (Marshall et al., 2021). Ultimately, the optimal size limit, or more generally, optimal size-selectivity in fisheries exploitation, will also depend on management objectives and hence social values (Ahrens et al., 2020). Beyond size-dependent impacts on reproductive traits (reviewed by Green, 2008; Hixon et al., 2014; Marshall et al., 2008), body length influences numerous other biological processes that have received less attention from conservation and fisheries management perspectives. Moreover, female size is often given more attention than male size in most exploited stocks, based on the assumption that sperm is rarely limiting (Hixon et al., 2014). Yet, body size in both sexes plays a role in sexual selection, mate choice, and decisions related to spawning and foraging locations (Hixon et al., 2014; Uusi-Heikkilä, 2020). Size variation also affects vulnerability to environmental stressors, such as heat waves or anoxic conditions, and influences dominance hierarchies and trophic interactions in complex ways (Ahti et al., 2020; Roos & Persson, 2013). Importantly, both inter- and intraspecific movement rates and dispersal often increase with body size in fishes (Minns, 1995; Tamburello et al., 2015), and body length reduces natural mortality (Lorenzen, 2022) and increases fecundity of female fish in most species (Barneche et al., 2018). Accordingly, large body size is a key trait under selection in various migratory (e.g., Burns & Bloom, 2020) and non-migratory fishes (e.g., Monk et al., 2021). In Pacific lamprey (Entosphenus tridentatus) and herring (Clupea harengus), for instance, larger fish migrate greater distances (Hess et al., 2014; Slotte, 1999). Similar patterns have been reported for Atlantic salmon (Salmo salar, Jonsson et al., 1991), brown trout (Salmo trutta, L'Abée-Lund, 1991), and American shad (Alosa sapidissima, Glebe & Leggett, 1981). An inverse relationship between body length and migration distance upstream, however, was reported in Pacific salmon (Crossin et al., 2004). Possible reasons may include differences in life-history (i.e. semelparity v. iteroparity), and

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selection acting differentially on salmonid species that travel only modest distances upriver and ones that must conserve energy for the downriver return to the ocean (Crossin et al., 2004). In another example of a typically non-migratory species, larger northern pike (Esox lucius) exhibited greater movement, larger home ranges, and larger reproductive output in a small lake than smaller, less mobile individuals (Monk et al., 2021). A positive size-movement relationships could be caused by greater swimming speeds at larger body sizes (Ohlberger et al., 2006), reduced cost of swimming (Alexander, 2003), improved condition (Bernatchez & Dodson, 1987), or accumulated experience in locating optimal spawning or foraging sites (Reebs, 2001; Rose, 1993; Webster, 2017). As large fish also face lower natural predation risk (Lorenzen, 2022), they may be less constrained to move and thereby act as keystone connectors, linking distant spawning habitats and thereby facilitating gene flow (Olsen et al., 2023). In social fishes, larger individual fish may also exhibit leadership in locating food patches (Reebs, 2001). Despite its relevance, the movement aspect of size variation and the potential impacts of size-selective mortality typical for fisheries on habitat connectivity remain underexplored, even though these factors may contribute to understanding why size-truncated stocks exhibit greater population fluctuations than those with more natural age and size structure (Anderson et al., 2008; Hsieh et al., 2010). Independent of the ongoing discussion about the conservation value of protecting large fish (most recently reviewed in Kopf et al., 2024), the socio-economic value of large-sized fishes is often substantial. This is especially true for recreational fisheries, where angler satisfaction tends to scale exponentially with increasing capture rates of large fish (Arlinghaus, 2024; Arlinghaus et al., 2014; Beardmore et al., 2015; Birdsong et al., 2022). Therefore, many recreational anglers voluntarily release large trophy fish (Arlinghaus et al., 2007), making their conservation feasible in fisheries where post-release mortality is low (Ahrens et al., 2020). In a recent study, Olsen et al. (2023) used acoustic telemetry in coastal sites of Norway to demonstrate that large female Atlantic cod (Gadus morhua) act as key spawning site connectors. This

is an important finding because connectivity among spatially separated spawning sites might provide

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a major buffering mechanism contributing to population resilience. The authors argue that removing these keystone individuals through fisheries could isolate meta-populations, increasing the risk of local extinctions. The work by Olsen et al. (2023) in batch spawning cod motivated us to investigate whether similar size-dependent connectivity exists in a coastal population of northern pike in the southern Baltic Sea, an extreme brackish environment for this freshwater species (Arlinghaus et al., 2023a; Rittweg et al., 2024). Northern pike is a phytophilic, large-bodied piscivorous freshwater fish (Craig, 1996; Skov & Nilsson, 2018), heavily exploited in both commercial and recreational fisheries in a positively size-selective manner (Arlinghaus et al., 2018). The species regularly occurs in coastal areas across the Baltic Sea where salinities average below 10 PSU (Practical Salinity Unit; Jacobsen & Engström-Öst, 2018). Pike are mesothermal and annual single-spawners, i.e. species that reproduce only once during the breeding season each year (also known as total spawners), with timing varying by latitude between February and May (Raat, 1988). During spawning, individual females release eggs with groups of a few males over a period of a maximum of few days depending on temperature fluctuations (Clark, 1950; Raat, 1988; Svardson, 1949). Males seem to preferentially spawn with larger, more fecund females (Fabricius & Gustafson, 1958), and larger males were found to sire a greater number of offspring in a natural lake (Pagel, 2009). In the southern Baltic, northern pike has diversified into ecotypes ranging from brackish residents to anadromous and freshwater subpopulations (Rittweg et al., 2024). Previous telemetry and markrecapture studies in the Baltic Sea have shown that pike is largely sedentary with limited home ranges (Dhellemmes et al., 2023a; Flink et al., 2023; Jacobsen et al., 2017; Karås & Lehtonen, 1993), forming a spatially structured meta-population across a network of brackish lagoons (Lukyanova et al., 2024) and interconnected rivers (Nordahl et al., 2019; Tibblin et al., 2016). Population connectivity is generally low outside the spawning season but increases significantly prior to and during spring spawning, when pike activity and space use increase (Cook & Bergersen, 1988;

Dhellemmes et al., 2023a; Diana, 1980; Flink et al., 2023; Lukyanova et al., 2024; Raat, 1988).

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Increases in space use during spawning are especially pronounced in coastal sites where multiple ecotypes, such as brackish spawners and anadromous fish (Müller, 1986; Sunde et al., 2022; Tibblin et al., 2016), co-exist and spawning migrations are regularly observed (Flink et al., 2023; Lukyanova et al., 2024). Anadromous fish that forage in coastal sites but return to freshwater streams for spawning rely on these migrations to spawn successfully (Tibblin et al., 2015). Yet also fully coastal ecotypes have been reported to engage in spawning migrations towards enclosed, sheltered bays that are used for spawning (Flink et al., 2023; Jacobsen et al., 2017; Lukyanova et al., 2024). Therefore, movement is a key contributor to successful spawning in coastal pike, and like the case of Atlantic cod (Olsen et al., 2023), larger pike might serve as key connectors among different areas and lagoons. However, it remains unclear whether and to what extent the largest individuals act as key connectors among spawning sites, which are located in tributaries, nearshore areas, and bays within the lagoons (Flink et al., 2023; Roser et al., 2023). Most importantly, as the reproductive biology of northern pike as total spawner strongly differs from Atlantic cod as batch spawner, spawning site connectivity as a function of length might emerge more across years than within a spawning season. In pike, as in many other fish species (Minns, 1995), space use is positively correlated with body size (Monk, 2019; Monk et al., 2021; Rosten et al., 2016). However, such positive relationship between body length and activity or home range size is not universally observed in pike (Dhellemmes et al., 2023a; Jepsen et al., 2001; Kobler et al., 2008; Koed et al., 2006). Discrepancies among studies may stem from differences in the size gradients studied, local environmental conditions, local ecosystem extension or population-specific characteristics. Additionally, past fishing pressure may play a role, as larger, faster-growing, and more active pike are selectively harvested by passive fishing gear, such as gill nets or by recreational angling (Carlson et al., 2007; Edeline et al., 2007; Monk et al., 2021). Across generations, such selective harvesting could favour smaller, less active phenotypes, potentially eroding the positive body length-movement relationship over evolutionary time. Also, the timing and duration of sampling and behavioural observation may influence findings, as pike are generally sedentary ambush predators for much of the year (Diana, 1980), with elevated activity

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primarily occurring before and during the spring spawning season (Lukyanova *et al.*, 2024; Raat, 1988). Therefore, size-related behavioural differences may be more pronounced prior to and during the spawning season and less evident in other seasons. No study so far has examined whether body length, in either males or females, enhances connectivity among spawning sites or whether larger individuals indeed act as key connectors, as observed in Atlantic cod (Olsen *et al.*, 2023). Collecting such data requires studying a large sample of differently sized fish in widespread coastal areas or very large lakes where long-range movements can be detected.

We equipped a coastal lagoon area of 1200 km² with an acoustic telemetry system, offering a suitable setting to examine the size and sex-dependency of spawning site connectivity in pike and assessed whether body length plays a key role in connecting different areas during the spring spawning period and outside spawning in a coastal pike population inhabiting a vast, interconnected lagoon ecosystem in the southern Baltic Sea across multiple years (for a review of the study area, see Arlinghaus *et al.*, 2023a). Following Olsen *et al.* (2023), we hypothesized that the largest individuals of both males and females would act as keystone connectors, linking spawning and foraging sites over large distances, resulting in positive relationships of body length and network connectivity both during and outside spring spawning time and especially between years.

#### **Materials and Methods**

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## Study area and telemetry array

Our study was conducted in the southwestern Baltic Sea, in the lagoons and freshwater tributaries bordering the islands of Rügen, Hiddensee, Fischland-Darß-Zingst, and Usedom in northeastern Germany (Figure 1). Like the rest of the Baltic Sea, the interconnected lagoons of the study area are brackish but exhibit substantial inter-lagoon variation in average salinity, ranging from 2 (oligohaline) to 9 PSU (mesohaline) along a northeast-to-southwest gradient, with the most isolated lagoons being the least saline (Arlinghaus et al., 2023a; Figure 1). In March 2020, we deployed an array of 140 acoustic telemetry receivers (VR2Tx, Frequency: 69 kHz, MAP-113, Innovasea Systems Inc. DE, U.S.A) across the study area covering 1200 km<sup>2</sup> of the total 1600 km² lagoon area on German territory (Dhellemmes et al., 2023a, Figure 1). The telemetry array was developed to gather data on area connectivity over a broad spatial scale rather than detect finescale movements. The array comprised the most important documented or suspected spawning sites of pike (Roser et al., 2023), both within the lagoons and in major inflowing streams and rivers (Figure 1). Receivers were retrieved, downloaded, and redeployed with a fresh battery in the winters of 2021, 2022, and 2023, in partnership with the Institut für Fisch und Umwelt (FIUM), Rostock, Germany. Over the course of the study, 13 receivers were lost, and 6 were relocated to enhance coverage in areas of interest. More details on receiver deployment are provided in Dhellemmes et al. (2023a).

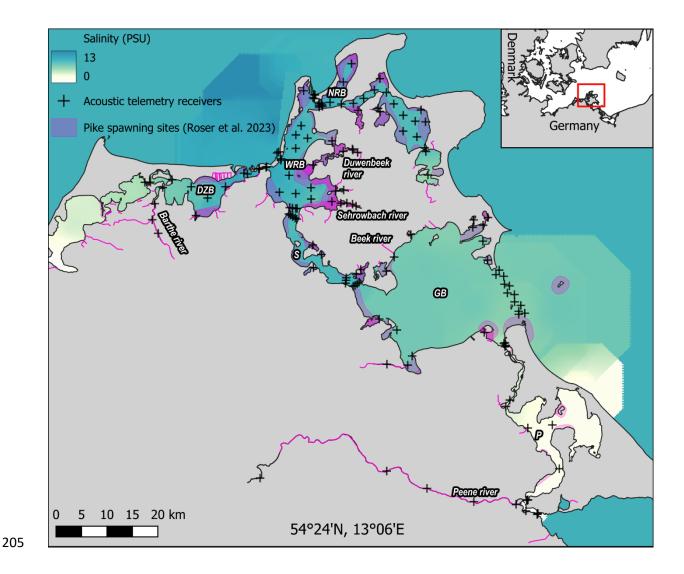
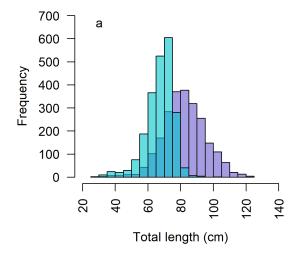


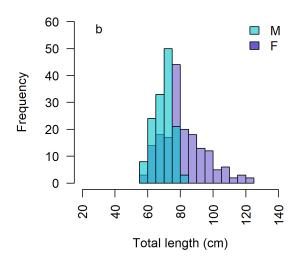
Figure 1. Map of the study area, including receiver locations, the average salinity gradient and the pike's spawning sites. Pike spawning site data are taken from Roser et al. (2023), where spawning sites are aggregated from different sources, which may agree on the location of a spawning site, leading to an overlap in the polygons and a brighter pink colour on the map. Italicised black and white labels indicate the different areas. DZB: Darß-Zingst Bodden; WRB: Western Rügen Bodden; NRB: Northern Rügen Bodden; S: Strelasund; GB: Greifswalder Bodden; P: Peenestrom.

## Fish captures and tagging

The research strategy involved tagging pike across as much of the size range of the species as possible (minimum total length for external ID tag = 28.0 cm; minimum total length for internal acoustic tag = 55.7 cm) and throughout all lagoons and most larger freshwater streams to observe year-round behaviour, with a particular focus on the period before and during the spring spawning

season and across a minimum of two spawning seasons. Between January 2020 and January 2023, we captured 5836 individual pike in collaboration with local fishers, angling guides, and through our own sampling. Sampling was non-random across the study sites, instead reflecting preferred fishing grounds by cooperating fishers and specific lagoons aligned with other project objectives (Arlinghaus et al., 2023b). Capture methods included rod and reel fishing, fyke nets, gillnets (in cool and cold water where fish can be released alive after capture), and electrofishing in the freshwater tributaries. Upon capture, pike were externally sex determined (cloacal shape or spilling of gametes upon gently pressing the body cavity; females = 2664, males = 2509, unknown = 663; Casselman, 1974), measured to the nearest millimetre (total length in cm; mean  $\pm$  SD = 73.6  $\pm$  16.1, min = 12.6, max = 126.2; Figure 2, a) and weighed to the nearest gram (mean  $\pm$  standard deviation (SD) = 2983  $\pm$  1979 g, min = 11.3, max = 17000). The health of all captured pike was assessed visually (i.e., colour, liveliness); healthy individuals that had not yet been tagged and measured more than 28 cm received external ID tags (N = 4597; females = 2319, males = 2183, unknown = 95; mean total length =  $75.2 \pm$ 13.6 SD, min = 28.0, max= 121.0; Figure 2; Floy T-bar anchor, Floy Tag & Mfg. Inc., NE, U.S.A.). Each tag indicated the web address and phone number of the recapture database (www.boddenhechtforschung.de), through which anglers and fishers could report their catch (date, ID, length, gear, and location, i.e., a lagoon name) to enter a raffle for fishing-related prizes. Overall, 17 % of the fish were captured and tagged by the research team and 83 % by cooperating guides and fishers.





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Figure 2. Frequency distribution of total length (mm) in our sample. a) All pike captured for the study and b) pike equipped with acoustic tags. Males (M) and females (F) are represented in blue and purple respectively.

Between February and December 2020, we selected 317 pike (males = 139, females = 177, unknown

= 1) to receive internal acoustic transmitters (N = 120, MM-R-16 50 HP, approx. 6-year battery life, dry weight = 35 g, in-water weight = 18.9 g; N = 196, MM-R-16 33 HP, approx. 3.5-year battery life, dry weight = 26.7 g, in-water weight = 13.6 g, random pulse rate: 60–180 s, Frequency = 69 kHz, MAP-113, Lotek Wireless Inc., ON, Canada), with 300 individuals tagged in the beginning of spawning season (February-March). Selection criteria included body weight (ensuring that in-water tag weight was always below 2 % of the pike's body mass; Jepsen *et al.*, 2005), visual assessment of their health (lethargic or otherwise damaged pike were excluded), and capture location (Table 1). Pike selected for telemetry were spread widely across the study area in capture locations both in lagoons and freshwater tributaries. They averaged 76.4 cm in total length (± 12.4 SD; min = 55.7, max = 121.0; Figure 2, b) and 3761 g in weight (± 2111 SD; min = 1388, max = 15000). To further motivate reports of captures of telemetry pike in the database, these pike received a white external ID tag, indicating a € 100 reward for the first report of each individual via our website or phone number. The acoustic receivers remained operational until the end of February 2023, allowing for the generation of up to three years of data per individual (Dhellemmes *et al.*, 2023a). Upon download, the data was filtered for false detections using *ATfiltR* (Dhellemmes *et al.*, 2023a). Upon download, the data was filtered

<u>Table 1. Capture locations, sex, and total length (in cm) of pike selected for transmitter implantation</u>
in 2020. Capture locations can be seen in Figure 1.

	Female (Mean ± SD; min; max)	Male (Mean ± SD; min; max)	Unknown (Mean ± SD; min; max)
Barthe river	6 (83.1 ± 17.3; 63.6; 115.2)	5 (62.1 ± 1.7; 61.2; 65.1)	0

Beek river	0	2 (67.4 ± 6.0; 63.1; 71.6)	0
Duwenbeek river	1 (77.3)	6 (68.8 ± 4.9; 60.1; 73.0)	0
Darß-Zingster Bodden (DZB)	31 (88.5 ± 10.4; 69.6; 110.5)	9 (72.9 ± 2.9; 68.9; 77.4)	1 (84.1)
Greifswalder Bodden (GB)	17 (70.6 ± 12.9; 58.9; 106.5)	12 (68.3 ± 4.6; 59.1; 74.8)	0
North Rügen Bodden (NRB)	27 (81.0 ± 8.5; 65.6; 96.6)	9 (69.7 ± 6.0; 60.5; 76.2)	0
Peenestrom (P)	26 (81.9 ± 11.7; 67.5; 110.0)	12 (66.7 ± 6.1; 58.6; 79.1)	0
Peene river	14 (73.4 ± 14.1; 56.4; 106.4)	11 (64.3 ± 5.9; 56.3; 73.5)	0
Strelasund (S)	27 (87.9 ± 14.3; 64.0; 121.0)	24 (73.5 ± 4.4; 61.6; 81.5)	0
Sehrowbach river	5 (83.1 ± 3.5; 79.0; 87.0)	8 (67.6 ± 4.7; 61.5; 76.1)	0
Western Rügen Bodden (WRB)	22 (78.3 ± 16.4; 55.7; 120.6)	42 (71.1 ± 5.3; 58.2; 82.4)	1 (74.1)

# Recapture distance

In the event of a recapture of a tagged pike (from our own sampling or via reports in our participatory mark-recapture database by fishers, guides and anglers), we calculated the in-water distance between each pike's initial capture and recapture locations. This included recaptures of both externally tagged and telemetry-tagged pike. The calculations were done using the *gdistance* package (Etten, 2017), which involved creating a transition layer from a shapefile of the land masses around our study area and then calculating the shortest path through this layer. This was done

independently of the capture and recapture dates, encompassing the entire study period. The average time between capture and recapture was 250 days (± 229 days).

## Connectivity and maximum horizontal displacement based on biotelemetry

We used the telemetry data to calculate connectivity and maximum horizontal displacement (i.e., maximum in-water distance between all detections of an individual pike, MHD) for February, March, April, and May to focus on the prime spawning season for pike. For comparison, we also calculated connectivity and MHD for each pike outside of the spawning period. MHD was computed in water using the same technique as described above for recapture distance. To assess connectivity, we constructed movement networks as unipartite undirected networks in the *igraph* package (Csàrdi *et al.*, 2025; Csàrdi & Nepusz, 2006), with nodes representing the acoustic receiver locations, and edges reflecting subsequent detections of individuals moving between these locations. Following the method in Olsen et al. (2023), each fish was assigned a connectivity score based on the number of unique edges detected within the time window of interest.

#### Among-year spawning site fidelity

To assess whether larger pike are more likely to change spawning sites among years, we estimated site fidelity in two ways. First, we used a binary variable indicating whether a pike was detected at the exact same location during spawning time across years (yes = 1, no = 0). Second, we calculated the Sørensen index (Sørensen, 1948), a metric commonly used in community ecology, which describes the overlap between a set of data points (here, areas) and can be used to assess site fidelity (Lenormand *et al.*, 2021). Sørensen values of 1 indicate identical area use and values <1 indicate increasing dissimilarity. Both metrics were calculated at two spatial scales: a broad area scale and a finer receiver scale (Figure 1).

#### Statistical analysis

We primarily focused on the size-dependency of movement metrics and connectivity, with a positive size-dependency supporting the assumption that the largest individuals act as key connectors between spawning sites and lagoon areas. We conducted all our analyses in R version 4.3.1 (R Core

Team, 2021). We compared metrics within and outside of the spawning season using t-tests. Distance between capture and recapture, MHD, and connectivity score (both for the spawning season and the rest of the year) were each fitted as response variables, with sex and total length as fixed effects in an interaction. For distance between capture and recapture, we added the time (in days) between each capture as a fixed effect. The capture area of each pike was added as a random effect to account for differences in receiver coverage and pike captures between the areas. We also added individual ID as a random intercept to MHD and connectivity models to account for repeated measures of the same individuals across years. Connectivity models were fitted using Poisson distribution, while MHD and recapture distance models were fitted using Gaussian distribution. To assess whether pike behaved in similar ways across years, we assessed the consistency and repeatability of pike behaviour by estimating Pearson's correlations between metrics in one year and the next, and adjusted repeatability (i.e., controlling for the effects of area, total length and sex) for each model using the rptR package (Stoffel et al., 2017) with 1000 bootstraps. To further our analysis of among-year variation in space use, we modelled site fidelity (binary: 0 or 1, at both area and receiver levels) and the Sørensen index (continuous: 0 to 1, at both area and receiver levels) as functions of total length interacting with sex while accounting for area and pike ID as random effects. Site fidelity was modelled using a Bernoulli distribution, while the Sørensen index was modelled using a beta distribution. All of our models were constructed in brms (Bürkner, 2017), using 4 chains of 120000 iterations each, with a thinning interval of 100 and a burn-in of 20000. We used leave-one-out cross-validations based on expected log pointwise predictive density (ELPD) based on the loo criterion in the loo package (Magnusson et al., 2019; Vehtari et al., 2017) to test whether or not the interaction between sex and total length should be left in the models. If the difference in ELPD between the models was >4, we estimated their predictive performance to differ and used the model with the highest ELPD for the analysis. If the difference in ELPD was <4, models were considered similar in their predictive performance, and we used the simplest model in our analysis. We assessed model fit by visually

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inspecting the posterior chains and considered fit to be satisfactory if no patterns could be observed. We interpreted effects as meaningful when their 95% credible intervals excluded zero. Pike of unknown sex (n = 14 for mark-recapture, and n = 2 for telemetry) were removed from the analysis.

## Visual assessment of spawning site use

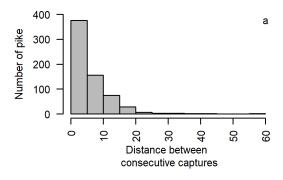
For all telemetry pike with an MHD greater than 10 km during the spawning season, we conducted an additional visual inspection of movement patterns. This included examining spatial maps of their displacements and plots showing the distance to their first detection during the spawning period (see Figure 6 for an example). We focused on the most mobile fish, i.e., individuals with broad-scale movements (MHD > 10 km), as the spatial resolution of our telemetry array was insufficient to detect finer-scale movement patterns and because Baltic pike have been observed to undertake spawning-related movements of this magnitude (Dhellemmes *et al.*, 2023a; Karås & Lehtonen, 1993). Within each year, we assessed whether individuals may have visited multiple spawning sites by identifying distinct peaks in the distance-from-first-detection plots (see Figure 6). For individuals with data spanning multiple years, we also examined whether they returned to the same areas across years. This analysis did not aim to rigorously test hypotheses regarding multi-site spawning, but rather to provide additional descriptive observations that could inform and guide future studies seeking to investigate these behaviors in greater depth.

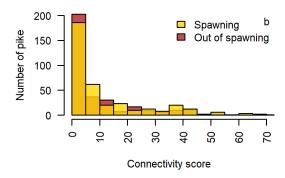
# Results

# Mark-recapture

Out of our 4597 tagged fish, 666 individuals were recaptured at least once (with 51 pikes having multiple recaptures). We were able to calculate the distance between capture and recapture and had total length and sex data for 546 of these. The minimum recorded distance was 0 km, and the maximum was 58.6 km, with a mean of 5 km ( $\pm$ 6.2 SD) and a median of 2.6 km (Figure 3, a). Leave-one-out cross-validations suggested that the sex by total length interaction resulted in a poorer model fit (difference in expected log pointwise predictive density (ELPD) = -0.6, standard error

(SE) = 0.5). We found males and females to be similar in their intercept (i.e., the average movement distance among captures), and neither time between captures nor total length had an effect on the distance between captures (Table 2, Figure 4).





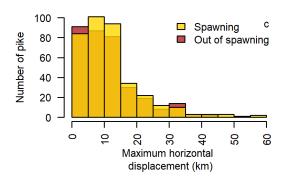


Figure 3. Frequency distribution of the response variables. (a) Distance between capture and recapture of individual pike. (b) Connectivity score during spawning season (February to May) and out of spawning. (c) Maximum horizontal displacement during spawning season and out of spawning.

<u>Table 2. Model output for the distance between capture and recapture.</u> Significant slopes and significant differences in intercept are highlighted in bold. Two samples are presented: the full sample, and a sample excluding the two most mobile fish.

		Estimate	Estimated error	Credible interval	Upper 95% Credible interval
ıle	Intercept (taken as Sex:Female)	0.5	2.39	-4.22	5.28
Full sample	Sex:Male	0.06	0.7	-1.29	1.42
	Total Length (cm)	0.05	0.03	-0.01	0.11
	Time between captures (days)	0.00	0.00	-0.00	0.00

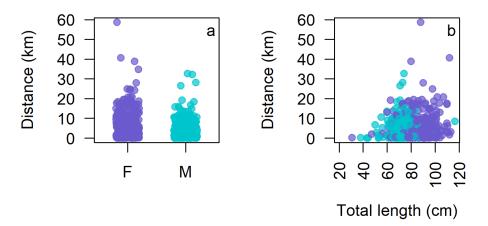


Figure 4. Distance between consecutive captures as a function of sex (a) and total length (b).

#### Connectivity and maximum horizontal displacement based on biotelemetry

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358 Out of 318 tagged fish, 292 generated data for an average duration of 439 days (time between first 359 and last detection, min = 0, max = 1065). After removing pike of unknown sex and those detected on 360 only a single receiver, the final sample size was N = 242 (females = 133, males = 109). Of those, N = 361 128 individuals produced data in two distinct years, and N = 48 in three distinct years. In total, we 362 analysed over 1.6 million (1649010) detections. 363 Connectivity (taken as the number of network edges per pike during a given period) was on average 364 12.56 (± 15.67 SD, min = 1, max = 87) during the spawning period (February, March, April, May) and 365 9.45 (± 12.55 SD, min = 1, max = 63) outside of spawning (Figure 3, b). MHD was on average 11.74 km 366  $(\pm 9.61 \text{ SD, min} = 0.14, \text{max} = 57.61)$  during spawning and 10.85 km  $(\pm 8.58 \text{ SD, min} = 0.14, \text{max} =$ 367 44.80) outside of the spawning period (Figure 3, c). T-tests indicated that connectivity was higher 368 during spawning (t = 2.92, p = 0.003), but MHD was not (t = 1.28, p = 0.19). 369 The interaction between body length and sex decreased model fit in the case of connectivity during 370 spawning and was therefore excluded (connectivity spawning ELPD = -11.4, SE = 5.6). In all other 371 cases, the inclusion of the interaction did not improve model fit, which also led to the exclusion of 372 the interaction term (connectivity out of spawning ELPD = -1.1, SE =4.1; MHD spawning ELPD = -0.1, 373 SE = 1.5, MHD out of spawning ELPD = -0.2, SE = 1.0). Higher connectivity and MHD during the 374 spawning time and the rest of the year were not explained by body size (Table 3, Figure 5, b, e, h, k), 375 rejecting the hypotheses of a positive size-dependency of spawning site connectivity. Males had 376 higher MHD out of the spawning period (Table 3, Figure 5, j), but we detected no other effect of sex 377 on movement metrics (Figure 5, a, d, g). 378 Connectivity, both within and outside of spawning season was well correlated between years (within 379 spawning Pearson's  $r_{(143)} = 0.52$ , p < 0.0001; out of spawning  $r_{(134)} = 0.7$ , p < 0.0001; Figure 5, c, f) but 380 had low repeatability at the individual pike level (within-spawning intra class correlation = 0.06 [0, 381 0.19], p=0.01; out of spawning intra class correlation = 0.19 [0.08, 0.37], p < 0.0001). MHD within and

outside of spawning time was also highly correlated among years (during spawning Pearson's  $r_{(143)}$  =

0.44, p <0.0001; out of spawning Pearson's  $r_{(134)}$  = 0.43, p <0.0001; Figure 5, i, l) and was strongly repeatable at the individual pike level (intra class correlation = 0.35 [0.2, 0.48], p < 0.0001; out of spawning intra class correlation = 0.26 [0.1, 0.39], p < 0.001).

<u>Table 3. Model output for the connectivity and maximum horizontal displacement based on acoustic</u>

<u>telemetry.</u> Significant slopes and significant differences in intercept are highlighted in bold.

		Estimate	Estimated	Lower 95%	Upper 95%
			error	Confidence	Confidence
				interval	interval
Connectivity	Intercept	1.15	0.57	0.06	2.25
(spawning)	(taken as				
	Sex:Female)				
	Sex:Male	-0.08	0.15	-0.38	0.22
	Total Length	0.00	0.01	-0.01	0.01
Connectivity	Intercept	1.29	0.69	-0.02	2.64
(out of	Sex:Male	-0.10	0.18	-0.45	0.25
spawning)	Total Length	0.00	0.01	-0.01	0.02
MHD	Intercept	11.47	4.71	2.17	20.60
(spawning)	Sex:Male	1.28	1.40	-1.45	3.96
	Total Length	-0.02	0.05	-0.12	0.09
MHD (out of	Intercept	10.80	4.51	1.74	19.57
spawning)	Sex:Male	2.61	1.25	0.2	5.02
	Total Length	-0.02	0.05	-0.13	0.08

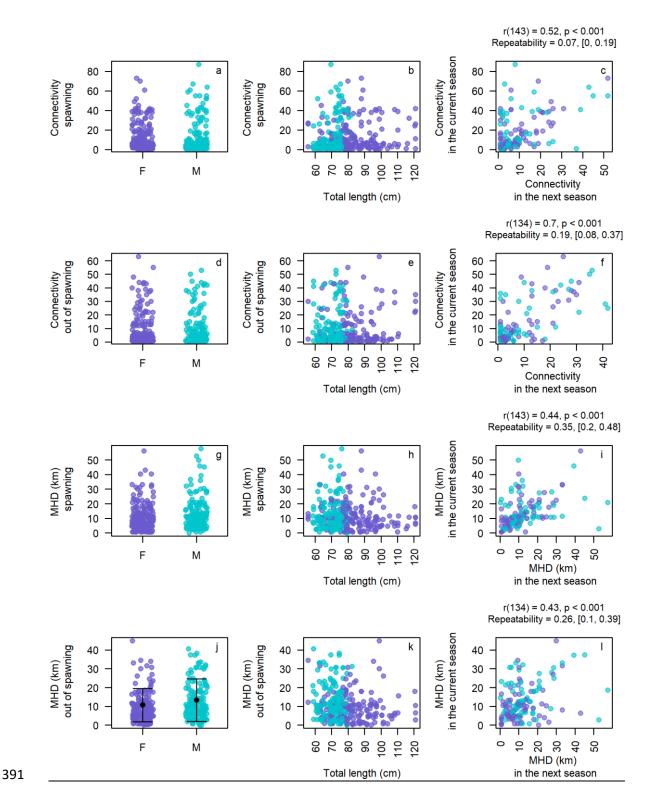


Figure 5. Scatter plot of the raw data for our response variables during the spawning period (February to May, panels a, b, c, q, h, i) and for the rest of the year (d, e, f, j, k, l) as a function of sex and total length and from one year as a function of the next. Connectivity during spawning and for the rest of the year is not explained by sex (a, d) or size (b, e), but individuals are consistent across years (c, f), although not repeatable during spawning. MHD during spawning is not explained by sex (d) or size

(e), but males have higher MHD out of spawning (j). Individuals are consistent and repeatable in MHD across years (i, l). Estimates and credible intervals are shown on panel j.

## Among-year spawning site fidelity

In all four among-year analyses, we dropped the sex by total length interaction as it did not improve model fit (site fidelity between area ELPD = -0.8, SE =0.3; site fidelity between receivers ELPD = -1.1, SE =0.6; Sørensen index between areas ELPD = -0.8, SE =0.5; and Sørensen index between receivers ELPD = -0.7, SE =1.1). Across all models, we found no effect of either total length or sex on among-year spawning site fidelity (Table 4). This indicates that although some individuals may change spawning sites across years, such differences were not explained by body size or sex.

<u>Table 4. Model output for the indices of site connectivity among-year as a function of body size and sex.</u> Significant slopes and significant differences in intercept are highlighted in bold.

		Estimate	Estimated	Lower 95%	Upper 95%
			error	Confidence	Confidence
				interval	interval
Site fidelity	Intercept	-0.53	1.77	-3.96	3.05
(area level)	(taken as				
	Sex:Female)				
	Sex:Male	0.07	0.49	-0.87	1.06
	Total Length	0.02	0.02	-0.02	0.06
Site fidelity	Intercept	1.27	2.28	-3.20	5.75
(receiver	Sex:Male	-0.25	0.58	-1.43	0.87
level)	Total Length	-0.04	0.03	-0.10	0.02
Sørensen	Intercept	1.52	0.68	0.19	2.83
index (area	Sex:Male	-0.09	0.19	-0.45	0.28
level)	Total Length	0.00	0.01	-0.01	0.02

Sørensen	Intercept	1.25	0.79	-0.31	2.80
index	Sex:Male	-0.21	0.23	-0.66	0.26
(receiver	Total Length	-0.01	0.01	-0.03	0.01
level)					

# Visual assessment of spawning site use

across different years (for example see Figure 6).

Out of the 93 individuals with an MHD greater than 10 km during the spawning season, 83% (N = 77, females = 42, males = 33) exhibited pronounced displacement behaviour in at least one year, i.e., they travelled to areas distinct from their non-spawning residency areas, as indicated by clear peaks in distance-from-first-detection plots (see Figure 6; Supplementary file 2, section 2 B). Among these, 13% (N = 12, females = 5, males = 7; Supplementary file 2, section 2 A) visited two different spawning sites in at least one year.

Of the fish with multi-year data (N = 62), 84% (N = 52) were recorded visiting a location during spawning season distinct from where they resided the rest of the year, indicating migratory behaviour. Among these migrants, 65% (N = 34; 20 females, 14 males; Supplementary file 2, section 1B) returned to the same spawning area for at least two years, with 7% of migrants (N = 4; 2 females, 2 males; Supplementary file 2, section 1A) recorded repeatedly using two distinct spawning sites in

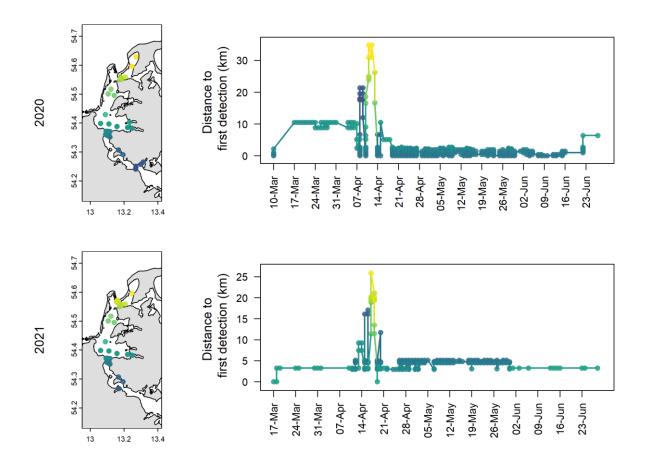


Figure 6. Map of detections (left) and distance from the first detection for pike BH-90099, an 88 cm female, in 2020 and 2021. In both years, the pike makes a brief visit to the south (blue receivers), followed immediately by a visit to the north (yellow receivers). This movement pattern may indicate visits to different spawning sites, although actual spawning behavior cannot be confirmed with our technology. In 2020, the pike travelled both slightly farther south and farther north compared to 2021, suggesting possible use of different areas between years.

#### Discussion

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In disagreement with the study hypothesis, neither mark-recapture nor telemetry data indicated higher connectivity or broader movement ranges in larger coastal northern pike of either sex during and out of the spawning season, except for males exhibiting higher maximum horizontal displacements (MHD) out of spawning. Site fidelity and repeated area use among years were also independent of size and sex in all analyses. These findings suggest that larger individuals of the total spawner pike do not serve as key spawning site connectors, unlike in the batch spawning cod (Olsen et al., 2023). However, at the individual level and unrelated to body length, we detected that both connectivity and spawning-season movement ranges were highly correlated between years and repeatable, as was connectivity outside the spawning season. This behavioral consistency suggests strong inter-individual variation in adult pike (i.e., the presence of behavioral types or personalities), aligning with prior research on pike in our study area (Dhellemmes et al., 2023a) and in other study systems and life stages in pike (Kobler et al., 2009; Laskowski et al., 2016; McGhee et al., 2013; Monk et al., 2021; Pasquet et al., 2016). Our additional visual assessment of the 93 most active pike (MHD > 10 km; 29% of all tagged fish) also pointed to the consistency in behaviours of these individuals, e.g., return to the same spawning area across the years. Some (13%) of these wide-roaming individuals visited multiple spawning sites within a single season, with no apparent sex- or size-based differences. Agreeing with our results, previous studies have found that behavioral consistency in pike was unrelated to body length (Nyqvist et al., 2012). Our results suggest that larger pike are unlikely to be key connectors among spawning (in the spawning time) or foraging areas (outside spawning time), contrary to what was found in the Atlantic cod in Norwegian fjords (Olsen et al., 2023). The disagreement between the two studies may stem from differences in reproductive biology and behaviour in Atlantic cod and coastal northern pike. Atlantic cod is a batch spawner, which exhibits a lek-based reproductive system with pelagic eggs (Nordeide & Folstad, 2000) and where females arrive at spawning territories previously established by larger, dominant males (Brawn, 1961; Dean et al., 2014). Cod are determinate batch spawners,

typically engaging in size-dependent release of small egg batches over multiple weeks to months across several sites (Kjesbu, 1989; Roney et al., 2018). Cod also show strong patterns of spawning site fidelity, as they are presumed to return to high-quality sites year after year (Skjæraasen et al., 2011). In contrast, while strong spawning and natal site fidelity have also been reported for northern pike (Miller et al., 2001; Tibblin et al., 2015), this species is characterised as total spawners, with females, often surrounded by multiple males who arrive earlier to spawning sites, releasing small batches of adhesive eggs onto underwater vegetation typically within a single day over several hours (Billard, 1996; Fabricius & Gustafson, 1958; Lindroth, 1946; Svardson, 1949). In rare exceptions, individual females release eggs over up to three days, but not involving long-range changes in spawning locations (Clark, 1950). There are reports that older and longer females predominate in the first days of the reproductive periods on spawning sites, and younger females seem to more frequent at the end of the spawning period (Sukhanova, 1979; Wright & Shoesmith, 1988). Male pike have more extended movements during spawning and occupy slightly larger territories than females, exhibiting intrasexual aggression in fights over access to females (Billard, 1996; Fabricius & Gustafson, 1958; Vostradovsky, 1983). In marine coastal areas, pike typically spawn in sheltered, vegetated bays, or in freshwater tributaries at traditional spawning grounds (Flink et al., 2023; Roser et al., 2023; Tibblin et al., 2016). Because pike eggs and larvae are adhesive to underwater vegetation, recruitment is localized (Billard, 1996) as there is no long-range movements of pelagic eggs, unlike in cod. The lek mating characteristic of Atlantic cod, combined with strong intrasexual selection (particularly among males), pelagic eggs, spatial bet-hedging (i.e., releasing eggs by large fish in different spawning sites, Olsen et al., 2023), and highly biased reproductive success in the wild (Hutchings et al., 1999; Roney et al., 2018; Rowe et al., 2008) may render the species more prone to show size-dependency in longrange movements so that the larger, more fecund or competitively superior female cod have an advantage in migrating to optimal spawning sites and can use multiple locations as a bet-hedging strategy to distribute egg production across space and time. This is facilitated by their ability to release pelagic eggs over multiple weeks (Kjesbu, 1989; Roney et al., 2018), allowing them to adapt to temporal environmental variability (Rogers et al., 2017).

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Such spatial bed-hedging within a spawning season by linking multiple spawning sites through movement is apparently not possible in the total spawning pike who mostly releases eggs within a single day, reducing any possibility to reap fitness benefit through long-range movements across multiple spawning sites. However, spatial bet hedging in total spawners such as pike may occur through the choice of variable spawning sites or variable timing of reproduction (Tibblin et al., 2016) across different spawning seasons. However, we also found no evidence for such size-dependent behaviour in our study system, agreeing with previous reports that revealed a high degree of natal site fidelity and high consistency in both lentic and coastal pike across years (Miller et al., 2001; Tibblin et al., 2015, 2016). Therefore, we believe spatial bet hedging at the individual level is far less pronounced in total spawners such as coastal pike compared to batch spawners such as cod, and instead, other mechanisms of spatial bet hedging through individual variation, such as ecotype evolution of brackish residency over anadromy and development of a long spawning season lasting up to two months (Arlinghaus et al., 2023a; Pagel et al., 2015), may provide population resilience in total spawning fish species. That said, the exploratory visual inspection of pike's movement patterns of the most widely roaming individuals suggested that some individuals (13%), both females and males, may visit multiple spawning sites within a single season, with several fish (4%) doing so repeatedly across the years. This could indicate that a small fraction of pike might engage in some form of spatial bed-hedging within a season, where individuals travel substantial distances to visit different spawning sites (Figure 6), but importantly, this pattern was unrelated to body length. Using oviduct transmitters and other tagging technologies could help to gain more precise estimates of

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spawning time and location for bony and cartilaginous fish (Gardner & Höök, 2024; Sulikowski & Hammerschlag, 2023), including pike (Pierce, 2004).



Figure 7. A ca. 95cm pike captured in a ditch during our sampling effort (Photo: P. Roser).

In the study area, coastal pike have evolved three reproductively isolated ecotypes that express four behavioural phenotypes: brackish residents, a cross-habitat ecotype favouring low-salinity areas, an anadromous ecotype that forages in brackish water but migrates into streams to spawn and freshwater residents that live and spawn in tributaries to the lagoons (Rittweg *et al.*, 2024; Roser *et al.*, 2023). The evolution of ecotype diversity may represent a pike's own spatial bet-hedging strategy, but one that does not rely on large pike acting as spawning site connectors and instead relies on population differentiation into ecotypes with different migratory tendencies. That said, depending on local conditions, while migrations are adaptive for the anadromous ecotype, large body size may be disadvantageous in small, shallow streams, potentially leading to selection against large body size in this ecotype (Figure 7). Previous studies have indeed failed to record very large and old anadromous individuals migrating into shallow streams draining into the lagoons (Rittweg *et al.*, 2024; Roser *et al.*, 2023). Furthermore, the lack of size-dependent relationships in the long-range migration distance and connectivity in the coastal pike (Dhellemmes *et al.*, 2023a) suggests that these ecotypes increase activity and space use during spawning time, but do so primarily through

localised movements within specific lagoons (Lukyanova et al., 2024). This aligns with previous findings in the study area, which also reported no strong relationship between space use and body length (Dhellemmes et al., 2023a), contrasting with other studies in pike (e.g., Rosten et al., 2016). However, size-dependent space use in pike has not been consistently reported in freshwater systems (e.g., Jepsen et al., 2001; Kobler et al., 2008; Koed et al., 2006a). While the lack of size-dependency in empirical data in past research could be related to low contrast in the size range or limitations in the studied ecosystem dimension (e.g., small lake), these factors are unlikely to apply in our research, given the large size range (50–126 cm) of our tagged pike and extensive study area (ca. 1,200 km<sup>2</sup>). We speculate that the lack of size-dependency both in overall space use (Dhellemmes et al., 2023a) and spawning site connectivity (this study) in the coastal pike population of Rügen, Germany, is related to the specific reproductive biology of pike as total spawner as discussed above. In addition, the result may in part stem from behaviour-selective harvesting and corresponding evolutionary adaptations. Far-roaming larger phenotypes and their underlying genotypes may have been systematically removed in decades of intensive harvesting in the study area (Arlinghaus et al., 2023a; van Gemert et al., 2022; Roser et al., 2024), altering the population's phenotypic and genetic composition, and eliminating individuals with a tendency to move a lot before and during spawning. The study area has been intensively harvested by small-scale commercial and recreational fisheries for more than a century (Arlinghaus et al., 2023a), which is sufficient time for the evolutionary impacts of harvesting to manifest in pike (Matsumura et al., 2011). Both gill nets (Carlson et al., 2007; Edeline et al., 2007) and recreational angling gear are known to target not only larger individuals but also those with an elevated space use (Monk et al., 2021), which often have higher fitness in the wild (Monk et al., 2021). In the study region, the so-called pre-spawn gill net fishery actively targets pike in spawning aggregations in bays (Arlinghaus et al., 2023a), exploiting their increased activity levels and migration into sheltered spawning bays (Flink et al., 2023). It is conceivable that the largest individuals and those with the longest migration distance have been selectively removed from the

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population, erasing the potential for size-dependent spawning-related movements to be revealed in current time.

Male pike were found to display higher MHD than females outside of the spawning season, when controlling for body size, in agreement with Jepsen *et al.* (2001) in one of two study lakes, but disagreeing with Koed *et al.*, (2006) in a river population. The difference between the sexes, did not apply, however, to the spawning season, suggesting that both males and females travel similar long-range distances to reach their spawning grounds. Earlier work had revealed that males appear more active during the spawning season than females (Lucas, 1992), but this study only looked at localized movements and not long-range movements. The observation that males may use more space than females is in agreement with past studies conducted in our study area (Dhellemmes *et al.*, 2023a). Movement differences between the sexes may be due to the cost of gamete production being lower for males than females (Jonsson *et al.*, 1997), resulting in males having more residual energy to dedicate to movements. They may also be driven by intersexual competition for resources, with males and females specializing in different foraging strategies, or due to sex dimorphism in size, which could mean that smaller males are under greater risk of cannibalistic or other types of predation, motivating greater displacements in the search for safe refuges (Haugen *et al.*, 2006, 2007; Li & Kokko, 2021).

Several limitations should be noted. First, we inferred the connectivity among spawning sites by analysing movement metrics during the key spawning period known for pike, without observing exactly where individual pike spawned. However, our receiver array broadly covered the most important known spawning sites (Figure 1), making our study likely robust for assessing long-range movement patterns. Second, given our array design, we were unable to track fine-scale behaviours during spawning site selection, and size-dependent micro-level site selection and associated differences in activity (e.g., Lucas, 1992) may still occur on local scales. The telemetry system we used was not designed to detect fine-scale movement resolution, which is why we cannot conclusively answer this question. Future studies focusing on specific lagoons and using biologgers and micro

transmitters inserted into the oviduct, capable of direct spawning site detection, will be necessary to determine whether pike of different lengths choose particular spawning sites and connect them through localised movements within specific lagoons.

To conclude, we found no evidence that larger pike of either sex serve as key connectors of different spawning and foraging sites within and across spawning seasons in different years. The lack of size-dependent connectivity suggests that spatial bet-hedging within a spawning season through connecting spawning sites maybe biologically impossible in the total spawner pike, and choosing of different spatially separated spawning sites across years may not confer sufficient fitness advantages for pike for evolution to happen. That said, we found indications that some individuals might still exhibit spatial bet-hedging behaviour during a spawning season by choosing multiple possible spawning sites independently of sex or size, although further research is needed to confirm that the pike indeed release multiple egg batches in different sites and not only visit multiple sites to choose the one to ultimately release their eggs. Our findings do not diminish the importance of conserving large fish, given their high social value (e.g., high angler preference for catching large pike; Koemle et al., 2022) and fecundity-related benefits of large-sized pike females (Ahrens et al., 2020). However, the conservation value of large pike cannot be motivated by their key role as spawning site connectors, unlike the case of batch spawning Atlantic cod (Olsen et al., 2023).

# **Ethics statement**

The research was completed following German legislation for animal experimentation, approved by Landesamt für Landwirtschaft, Lebensmittelsicherheit und Fischerei MecklenburgVorpommern—

Veterinärdienste und Landwirtschaft—under grant no. 7221.3-1-052/19.

#### Data availability statement

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The data are available in the European Tracking Network repository: Dhellemmes F, Arlinghaus R (2021) Boddenhecht telemetry dataset. https://marineinfo.org/id/dataset/7859

#### **Acknowledgments**

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