Title Page 1

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- Title: An environmental resistance model to inform the biogeography of aquatic invasions in 4 complex stream networks
- 5
- **Running title:** Environmental resistance to invasions 6
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- 27

28 Abstract

- Aim Freshwater invasions are a global conservation issue. Emerging tools for biogeographic 29
- analyses can provide critical information for their effective management and monitoring. Here, 30
- we propose a method to assess the distribution of environmental resistance of stream ecosystems 31
- to biological invasions by coupling multi-stage habitat potential models for non-native species. 32
- 33 Location Andean Patagonia (Chile and Argentina).
- 34 Taxa North American beaver (Castor canadensis), Chinook salmon (Oncorhynchus
- tshawytscha), and coho salmon (O. kisutch). 35
- **Methods** Environmental resistance to invasive species was mapped throughout a large region of 36
- 37 Patagonia by stacking multi-stage habitat relationships for each target species and assessing the
- complementation between critical habitats at multiple scales. We generated an environmental 38
- model of stream networks derived from high-resolution topographic and climatic data 39
- representing 15,406 drainage basins (> 1 km²) covering an area of 369,791 km². We quantified 40
- the intrinsic potential of stream reaches (100 m and 1,000 m) to sustain high-quality habitats and 41
- assessed habitat complementation (i.e., abundance and proximity) at the sub-basin scale as a 42
- proxy for environmental resistance. 43

44 **Results** Our model revealed high heterogeneity in the distribution of environmental resistance to

- invasions throughout the study region, providing case-specific insights for the research and
- 46 management of invaders.
- 47 Main conclusions Environmental resistance modelling is a novel method to study the
- 48 biogeography of riverine invasions. Our approach is compatible with additional sources of
- 49 information about species and the environment, and shows versatility to diverse invasion
- scenarios and data sources. This method can be useful in prioritising research and management
- of incipient and spreading invasions, especially for large and data-poor regions.

Keywords: aquatic, complementarity, hydrogeomorphology, invasibility, lotic, non-native, risk,
 resistance, riverscapes, suitability

54 55

1. Introduction

56 Biological invasions are a primary driver of global ecological change (Lockwood et al., 2013)

- and a source of increasing attention in biogeographic studies (Hastings et al., 2005; Acevedo et
 al., 2016). Spatially explicit habitat modelling can provide valuable information for risk
- assessment and spatial prioritisation of monitoring and management of invasions (Jeschke &
- 60 Strayer, 2008; Jiménez-Valverde et al., 2011). However, the apparent idiosyncrasies and lack of
- final strayer, 2008, Jinenez-Varverde et al., 2011). However, the apparent thosyncrastes and fack
 predictability of invasions in freshwater ecosystems (Moyle & Light, 1996; García-Berthou,
- 62 2007) and the high structural complexity of stream networks (Fausch et al., 2002; Wiens, 2002)
- 63 present additional challenges to modelling the spatial distribution of aquatic invaders. The
- 64 management and control of introduced species depend on introduction intensity (propagule
- 65 pressure), the invasive traits of introduced species (e.g., vagility, fertility, phenotypic plasticity;
- 66 Kolar & Lodge, 2001), and the compatibility of their life-history requirements with critical
- attributes of the receiving ecosystem (e.g., suitability, connectivity). Therefore, habitat modelling
 applied to freshwater invasions warrants the consideration of unique ecological relationships
- between introduced species and the spatial structure of freshwater ecosystems.

In the absence of empirical information about presence/absence, mapping the habitat 70 suitability for invaders in stream networks can be approached using mechanistic knowledge of 71 the interplay between landscapes, populations, and behaviour as seen in native ranges (Poff, 72 73 1997; White et al., 2014). Conceptually, riverscapes represent freshwaters running through landscapes (Ward, 1998; Fausch et al., 2002; Wiens, 2002) as highly confined, linear, and 74 hierarchical stream networks (Ward, 1998; Montgomery, 1999; Fausch et al., 2002; Benda et al., 75 2004; Humphries et al., 2014). The biophysical heterogeneity (Ward, 1989) of riverscapes is 76 77 shaped by multiple longitudinal (i.e., upstream-downstream), lateral (i.e., channel-floodplain), vertical (i.e., surface-hyporheic), and temporal dimensions. Yet, the distribution and dispersal of 78 79 freshwater species are generally shaped by the longitudinal dimension of riverscapes (Huet, 1959; Vannote et al., 1980; Penaluna et al., 2021). Predictive models able to assess these critical 80 components of the longitudinal distribution of stream habitats can be used to evaluate the 81 propensity of receiving riverscapes to support novel species. 82

The habitat intrinsic potential model is a mechanistic approach to characterise the structural capacity of individual stream sections to provide high-quality habitats for a target species (Burnett et al., 2007). In native ranges, this approach has been used to identify historical (Agrawal et al., 2005) and degraded (Burnett et al., 2007) habitats and to monitor habitats for juvenile animals (Matter et al., 2018), species relocations (Dittbrenner et al., 2018), and population abundances (Flitcroft et al., 2012; Petro et al., 2018; Romey, 2018). In addition, this approach has been expanded to represent both multiple spatial scales (Firman et al., 2011) and
species' life stages (Flitcroft et al., 2012). The use of the intrinsic potential approach in invasion
biology is promising, for example, in predicting possible interactions between native and
introduced species (Jalbert et al., 2021; Manning et al., 2022). Yet, the lack of standard protocols
to assess the spatial distribution of ecologically-relevant hydrogeomorphic processes, as well as
the inclusion of multiple invaders' life stages, have hindered the use of intrinsic potential models
to answer questions about the biogeography of aquatic invaders.

96 Species relate to their ecosystems at multiple spatial-temporal scales, and species-specific relationships between invaders and their landscapes are essential to develop meaningful invader-97 habitat models (Brown et al., 2008; Vicente et al., 2019). Every species requires specific 98 99 environmental conditions and resources, and those requirements can present predictable ontogenic changes. For instance, species often require multiple habitat types for the completion 100 of their life cycles in stream networks (e.g., Schlosser, 1995a), a process known as "habitat 101 complementation" (Dunning et al., 1992). The availability of complementary habitats can be 102 viewed as an environmental filter to invasions (Moyle & Light, 1996). Therefore, a 103 comprehensive understanding of the relationships between species life-histories and relevant 104 landscape attributes is critical for assessing a region's vulnerability to the colonisation and 105 expansion of introduced species. Under this paradigm, the traditional intrinsic potential approach 106 using single life-stage characterisations of habitat suitability provides an incomplete 107 108 representation for species with complex life histories. Thus, habitat models incorporating habitat complementation requirements can enhance intrinsic potential models to map and quantify 109 environmental resistance to invasions of species with complex life histories. 110

Here, we develop and implement a method for mapping environmental resistance to 111 invasions based on habitat intrinsic potential and habitat complementation models at different 112 spatial scales. Our models include a clear description of standard steps and procedures that will 113 ensure reproducibility. We provide a proof-of-concept and partially validate our model using 114 three species introduced in freshwater environments of southern South America, including the 115 North American beaver (Castor canadensis Kuhl, 1820), Chinook salmon (Oncorhynchus 116 tshawytscha Walbaum, 1792), and coho salmon (O. kisutch Walbaum, 1792). Our approach is 117 transferable across species and ecosystems and aims to advance the understanding of spatial 118 patterns of these and other aquatic invasions by prioritising surveillance efforts for regions at 119 high risk, but too large to be sampled comprehensively. 120

121 122

2. Materials and Methods

We mapped the distribution of environmental resistance to the invasion of non-native species following five sequential steps (**Fig. 1**). In Step 1, we assembled multi-stage models of specieshabitat relationships. In Step 2, we characterised habitats at stream reach and basin scales using a hydro-geomorphic model. In Step 3, we classified the intrinsic potential for each life stage for each reach. In Step 4, we assessed habitat complementation at the sub-basin scale. In Step 5, we mapped the distribution of environmental resistance to species invasion.



- 131 Figure 1. Workflow diagram of our spatially-explicit modelling framework to map the
- distribution of environmental resistance to the invasion of non-native species in stream networks
- showing 5 key steps.
- 134

135 **2.1 Study area**

136 Our study area encompassed the Andean Patagonian region, including all Pacific basins and the

- archipelago of Tierra del Fuego (Fig. 2). This region is characterised by complex mountainous
 terrain and high heterogeneity in geophysical and bioclimatic conditions (Amigo & Ramírez,
- 139 1998; Olson et al., 2001). The region's physiography has been shaped by the Andes Mountains,
- featuring active volcanos, fjords, glacial fields, islands (>40,000), and diverse valley types
- 141 (Börgel, 1965). The extensive latitudinal range of our study region (36-55 °S) includes multiple
- 142 climates and hydrological conditions (Abell, 2002). Most human-related influences are
- 143 concentrated in the north (38-41°S), whereas pristine protected areas predominate throughout the 144 south (41-55°S).
- 145

146 **2.2 Target species used to illustrate our modelling procedure**

- 147 North American beaver and Pacific salmon (Chinook and coho salmon in this study) are
- ecological engineers and keystone species in their native ranges. Thus, the potential
- 149 consequences of their introductions in novel environments will likely result in far-reaching
- 150 impacts on receiving systems. For instance, beavers can alter ecosystems across multiple
- organisation levels, from the geomorphology of channels to water chemistry to the entire
 composition of aquatic and terrestrial fauna (Naiman et al., 1988; Anderson et al., 2020; Brazier
- composition of aquatic and terrestrial fauna (Naiman et al., 1988; Anderson et al., 2020; Brazier
 et al., 2021). Similarly, Pacific salmon contribute marine-derived nutrients to terrestrial and
- freshwater ecosystems sustaining many species and communities (Naiman et al., 2002). In
- addition, these focal species could interact symbiotically in invaded regions. Beaver ponds can
- improve local habitat conditions (Alexander, 1998), growth (Arismendi et al., 2020), abundance
- 157 (Virbickas et al., 2015), spatial distribution (Leidholt-Bruner et al., 1992; Mitchell & Cunjak,
- 158 2007), and overall population productivity of salmonids (Pollock et al., 2004). In contrast, beaver
- dams can also influence connectivity for dispersal and spawning migrations (Lokteff et al.,
- 160 2013). Therefore, environmental models generated jointly for these invasive species could have
- 161 synergistic advantages for studying their biogeography.
- Pacific salmon (Basulto, 2003) and the North American beaver (Skewes et al., 2006)
 have been introduced in South America (Chile and Argentina) during the last century. They have
 had different levels of invasion success, ecological impacts, and societal values. See Appendix
 S1 in Supporting Information for a summary of the history of introduction, invasion success,
- and effects of these species in the study region.
- 167
- 168



Figure 2. Map of the study area in Andean Patagonia (Chile and Argentina, 36.8 – 55.7 °S)
encompassing a total area of 369,791 km², and the generalised geographic ranges of our three
non-native target species. For North American beaver and Chinook salmon, coloured lines
represent reportedly invaded ranges, and for non-native coho salmon, coloured areas represent
the distribution of potential propagule sources from aquaculture production during the last ten
years (2010-2019). Black stars indicate the locations of potentially naturalised coho salmon

detections (Górski et al., 2017; Chalde et al., 2019). Map projection: South America Albers

177 Equal-Area Conic.

178

179 2.3 Step 1: species-habitat relationships

180 The development of mechanistic species-habitat relationships requires the identification of variables determining the quality of physical habitats using remote sensing data and Geographic 181 Information Systems (GIS; Lunetta et al., 1997). The habitat intrinsic potential model requires 182 183 the delineation and characterisation of stream networks based on high-resolution elevation data (< 30-m, Fig. 1) and uses simple geohydrological variables (e.g., discharge, gradient, valley 184 confinement) that can predict suitable environments (e.g., flow velocity, substrate size, hyporheic 185 exchange, pool-riffle ratio). This procedure allows for the customisation of species-habitat 186 relationships for different life stages that can be transferred to any region of interest. 187

We revised all intrinsic potential models published for our three target species to develop 188 transferable species-habitat relationships for applications related to introductions to novel regions 189 outside their native ranges. For the North American beaver, these models have been successfully 190 implemented to map the potential of streams to support dam-building habitats in Oregon and 191 Washington, USA (Suzuki & McComb, 1998; Dittbrenner et al., 2018; Petro et al., 2018). For 192 Chinook (Agrawal et al., 2005; Busch et al., 2013; Bidlack et al., 2014; Matter et al., 2018) and 193 coho salmon (Burnett et al., 2007; Romey, 2018), intrinsic potential models have been described 194 for different populations, phenotypes, and life stages. The differences in species-habitat 195 196 relationships among these studies make it difficult to choose a single species-habitat model for

- applications outside their native ranges, especially considering the mixed genetics of Pacific
- salmon introductions in Patagonia (Neira et al., 2014; Correa & Moran, 2017; Gomez-Uchida,
- Cañas-Rojas, et al., 2018). Instead, we combined species-habitat relationships from native rangesto incorporate the variability of tolerable and optimal conditions described for different
- 201 phenotypes of the same target species. Using this approach, we combined trapezoid-shaped
- piece-wise linear functions (i.e., trapezoidal functions) by enveloping the whole area under the
- 203 curve described for the relationship between each species and environmental variable (i.e.,
- envelope assembly; **Fig. 1, Step 1**). This simple aggregation technique acknowledges the high
- degree of uncertainty in predicting patterns that might occur in the receiving ecosystems. In
- addition, it gives value to species plasticity traits (i.e., related to environmental requirements),
- which have been highly associated with invasion success (Alcaraz et al., 2005; Olden et al.,
 2006; Strayer, 2010; Arismendi et al., 2014). The resulting species-habitat functions were
- 200, Strayer, 2010, Anshendr et al., 2014). The resulting species-habitat functions were 209 interpreted as representations of all known tolerable (trapezoid's feet) and optimal (trapezoid's
- shoulders) environmental ranges described for each species and life stage. More details of this
- 211 procedure are provided in **Appendix S2**.
- 212

213 2.4 Step 2: hydrogeomorphic model

There are no publicly available datasets of stream networks in South America to support the

creation of habitat intrinsic potential models. Thus, we used publicly available topographic and

- climatic datasets to delineate and characterise the stream networks of our study region (**Fig. 1**, **Star 2**). For this number we developed a systemiced processing to allow in ArecUS Proc 2.8
- Step 2). For this purpose, we developed a customised processing toolbox in ArcGIS Pro 2.8
 (Esri, 2022) through 'arcpy' (code available at https://doi.org/10.5061/dryad.3tx95x6jf). We
- followed similar procedures described by Miller (2003) for hydrogeomorphic watershed analysis
 based on Digital Elevation (DEM) and precipitation models (*sensu* Benda et al., 2007; Clarke et
- 221 al., 2008).

Based on a DEM of 12.5-meter resolution (downscaled from 30-m SRTM DEM by the 222 Alaska Satellite Facility; https://asf.alaska.edu) and a mean annual precipitation model (30 arc-223 seconds; Fick & Hijmans, 2017), we generated a spatially explicit model of the stream networks 224 of Andean Patagonia (36.8 – 55.7 °S). This environmental model covered an area of 369,791 225 km^2 comprised of 15,406 river basins (> 1 km²). The digital drainage network represented 226 671.971 linear km of streams > 0.01 m³s⁻¹. Stream reaches were characterised at two spatial 227 resolutions (i.e., 100 m and 1,000 m line sections) based on the most frequently applied variables 228 229 for our target species in the scientific literature (see Appendix S2), including 1) mean annual discharge, 2) channel gradient, 3) channel width, 4) valley width, 5) valley confinement, and 6) 230

- *anadromous connectivity.* For an expanded description of environmental modelling steps, see
 Appendix S3.
- 232 Appe 233

234 2.5. Step 3: habitat potential model

Using the species-habitat relationships for each of our three target species (**Fig. 3**), we scored reach-scale geophysical attributes ranging from 0 (low) to 1 (high) based on their potential to

sustain habitats at any given time. Scores were combined using geometric means to calculate a

stream reach intrinsic potential index (**Fig. 1, Step 3**). This averaging method enabled the

combination of partially compensatory environmental variables in determining the presence of

- suitable habitats. For instance, optimal discharge and valley conditions can compensate for sub-
- optimal channel gradients. However, if another variable's score approximates zero (i.e., limiting

factor), its weight is higher in the final intrinsic potential index (more details in Appendix S3, section 2.5).

244 245

2.6. Step 4: habitat complementation model

Pacific salmon require different lotic habitats to complete life cycles (Flitcroft et al., 2019). Thus, 246 an approach that includes habitat complementation (Schlosser, 1995b) is warranted to better 247 address ecological questions in riverscapes (Fausch et al., 2002). Here, we evaluated the 248 complementation of freshwater habitats for Pacific salmon by estimating the intra-basin 249 abundance and proximity of potential spawning and rearing habitats. For North American 250 beaver, given habitat intrinsic potential models were only available for a single life stage (i.e., 251 252 dam building), we assessed habitat complementation between high-potential aquatic and terrestrial environments (i.e., forest cover) as shown in high-resolution land cover data (10-253 meter, Karra et al., 2021). The spatial density of high-potential habitats (linear km/sub-basin area 254 for beavers, linear km/accessible linear km for salmon) for different life stages were 255 compounded in a Habitat Complementation Index (HCI). The HCI consisted of four discrete 256 classes (Fig. 1, Step 4), namely no habitat complementation (HCI = 1; missing potential habitats 257 258 for one or both stages), low habitat complementation (HCI = 2; potential habitats are available, but in low densities, overall), medium habitat complementation (HCI = 3; potential habitats are 259 available in moderate densities, overall), and high habitat complementation (HCI = 4; high 260

- density of potential habitats, overall). More details about the calculation of HCI for each species
 are provided in Appendix S3, section 2.6.
- 263

264 2.7 Step 5: Environmental resistance model

We adapted this habitat complementation model to map the distribution of environmental resistance to species naturalisation and invasion (**Fig 1, Step 5**). We estimated environmental resistance as the inverse of the HCI, resulting in an Environmental Resistance Index (ERI) with

- four discrete classes. Areas providing high habitat complementation were classified as having
- low environmental resistance to invasion (ERI = 1). In contrast, areas with no habitat
- 270 complementation where classified as having very high resistance to invasion (ERI = 4).
- 271

272 **2.8 Model applications**

273 2.8.3 North American beaver

We mapped the distribution of environmental resistance to the invasion of beavers in the Tierra

del Fuego archipelago (**Fig. 4c**) and compared it with current beaver distribution data (**Fig. 4a**;

Huertas Herrera et al., 2020) and priority management zones (**Fig. 4b**; Parkes et al., 2008).

- 277 Priority areas for the management of invasive beavers were designed by the Chilean Agriculture
- and Cattle Service (Servicio Agrícola y Ganadero, SAG) to prevent the expansion of the species
- to the mainland (Parkes et al., 2008). We identified high priority areas that were still uninvaded

280 by beavers but presenting low resistance to beaver invasion. This application provides explicit

information for the spatial narrowing of surveillance and prevention efforts of beavercolonization.

We also tested the predictive potential of environmental resistance modelling using an invasive beaver dam occurrence dataset for Tierra del Fuego (Huertas Herrera et al., 2020). This dataset consisted of 206,203 locations of active and abandoned beaver dams digitised from highresolution satellite imagery. We fitted logistic linear regressions to test the sensitivity and specificity of our beaver habitat model to predict the presence-absence of beaver dams at subbasin and stream reach (1 km scales, using HCI indexes and available linear km of potential

habitats as explanatory variables. Additionally, we assessed the performance of our

environmental resistance model for identifying degrees of invasion (based on sub-basin dam

- densities) using Mann-Whitney U tests for contiguous classes (Fig. 4c).
- 292 293

294 2.8.1 Chinook salmon

We mapped and quantified the abundance and proximity of potential habitats for the rearing and 295 spawning life stages of Chinook salmon. We identified areas with the lowest environmental 296 resistance to a continued invasion of Chinook salmon, and areas with decreased habitat potential 297 298 and connectivity, i.e. high resistance. In addition, we illustrated our approach to answering management questions at smaller spatial extents (i.e., basin level). Using the case of the Toltén 299 River basin, we mapped the distribution of environmental resistance in a system that supports a 300 301 self-sustaining, naturalised Chinook salmon population. The Toltén River basin supports emergent recreational and artisanal fisheries of growing interest for public agencies and scientists 302 (Cid-Aguayo et al., 2021). We combined our environmental resistance assessment with current 303 304 land and water uses to illustrate model applications for different management scenarios affecting the biogeography of this salmon fishery. 305 306

307 *2.8.2 Coho salmon*

We combined our environmental resistance assessment with current information about coho 308 salmon propagule pressure represented by salmon farming production in marine net-pens 309 between 2010 and 2019. We summarised historical data of massive salmon escapes and total 310 aquaculture production (a surrogate for leakage escapes) at aquaculture management zones from 311 Los Lagos and Aysén districts (41-47°S), where most coho salmon aquaculture exists (Soto et 312 al., 2022). The overlap between areas with low environmental resistance and high propagule 313 pressure can be interpreted as areas at risk of non-native coho salmon naturalisation. In addition, 314 we overlapped our environmental resistance assessment with the most recent reports of likely 315 naturalised salmon (Górski et al., 2017; Chalde et al., 2019) to illustrate the proximity of 316 freshwater habitats that could support these self-sustaining populations. These maps can assist 317 decision-makers in the spatial allocation of surveillance and pre-emptive management efforts. 318

319

320 3. Results

321 **3.1** Species-habitat relationships for our three target species

In our review of the intrinsic potential modelling literature, we identified 17 original studies modelling the relationships between reach-scale geomorphic variables and the habitats of our three target species (**Fig. 3**; Table S1). All available studies focused on native ranges. Studies showed differences among species, populations, and life stages, as well as in their selection of geophysical variables and the shape of habitat functions (more details in **Appendix S2**).



329 Figure 3. Species-habitat suitability relationships obtained from a literature review to classify geophysical conditions based on their potential to provide habitats for our three target species. 330 The upper panel shows suitability curves for classifying dam-building habitats for North 331 332 American beaver (a). The lower panels show Chinook (b) and coho salmon (c) rearing (green line) and spawning (purple line) suitability curves. Rearing suitability curves for both species 333 shared discharge, gradient, and valley constraint variables, while Chinook salmon spawning 334 models predominantly used channel width instead of discharge. Our habitat model included two 335 suitability curves for each salmon species representing the swimming capacity of adult migrants 336 through stream gradients at two different scales: 100-m (grey line) and 1,000-m stream reaches 337 (black line). 338

- 339
- 340

341 **3.2** Distribution of environmental resistance for target species and illustrative applications

342 *3.2.1 North American beaver*

- 343 We focused on the archipelago of Tierra del Fuego and adjacent areas where beavers have not
- 344 yet colonised. In this region, we delineated 3,647 basins draining directly to the ocean and
- subdivided basins > 50 km^2 into units averaging 120 km^2 . Our model identified 32,163 stream
- km with high geophysical potential to sustain dam-building beaver habitats, representing 41.5%
- of the modelled stream network. At the sub-basin level (**Fig. 4c**), our model returned 18.9%
- 348 $(11,645 \text{ km}^2)$ of the area having low resistance (ERI = 1) to the invasion of beavers, 27.5%
- 349 (16,966 km²) showed moderate resistance (ERI = 2), 16% (9,851 km²) showed high resistance
- (ERI = 3), and 37.6% (23,235 km²) showed a very high resistance (ERI = 4). Areas of low environmental resistance were concentrated in south-eastern and central-western parts of the
- environmental resistance were concentrated in south-eastern and central-western parts of the main island and the Brunswick Peninsula. The comparison of environmental resistance with
- 352 main island and the branswick remission of environmental resistance with 353 priority management zones (by SAG; Fig. 4b) showed high overlap of low environmental
- resistance with top priority areas for the containment of beaver invasion (i.e., removal zones),
- 355 suggesting an increased risk of expansion into the continent. Further, logistic regressions showed
- satisfactory performance in identifying beaver occurrences at the sub-basin (Area Under the
- 357 Curve of the Receiver Operating Characteristic, AUC = 0.82, 64% sensitivity, 82% specificity)
- and reach scales (AUC = 0.86, 83% sensitivity, 73% specificity). Lastly, Mann-Whitney U tests
- showed a high discrimination rate in the degree of invasion (**Fig. 4a**) among environmental
- resistance classes (*p*-values < 0.000001; **Fig. 4d**), proving helpful predicting the distribution of
- 361 beaver invasions at the sub-basin scale.



Figure 4. Model results for North American beaver, showing (a) map of environmental resistance at the sub-basin level. Plot (b) shows priority management zones for invasive beaver by the Chilean Agriculture and Cattle Service (Servicio Agrícola y Ganadero, SAG), here extending the "sustained control" zone into Argentina. (c) shows sub-basin beaver dam densities based on the occurrence dataset by Huertas Herrera (2020). The boxplot in (d) compares dam densities at each environmental resistance class (without outliers) using the same colours as in

- 369 (c), with red depicting low resistance, yellow medium resistance, and orange showing low
- resistance sub-basins. **** denotes high significance levels (p-values < 0.0001) in class
- 371 discrimination obtained from Mann Whitney U tests. Map projection: South America Albers
- 372 Equal-Area Conic.
- 373
- 374 3.2.2 Pacific salmon
- We mapped the distribution of environmental resistance to Pacific salmon (Chinook and coho
- salmon) invasions for the entire study area (Web Map available at <u>https://arcg.is/1WuqDH</u>). No
- areas were excluded based on our air temperature exclusion criterion (see Appendix S3, section
- **2.3**), whereas approximately 8.1% (54,437 stream km) were excluded due to exceedingly high
- 379 glacial influence.
- 380
- 381 *3.2.2.1 Chinook salmon*

- 382 The anadromous connectivity analysis for Chinook salmon in Andean Patagonia showed high
- access potential (reaches with anadromous connectivity scores > 0.75) of 167,736 stream km.
- 384 The intrinsic potential classification showed that 31,540 km had a high potential to sustain
- Chinook salmon rearing habitats, and 45,434 km had a high potential for spawning habitats.
- Most sub-basins (46.1%, 165,739 km²) had a very high resistance (ERI = 4) to Chinook salmon
- invasion (mostly systems that were not accessible to anadromous salmon). Sub-basins with high
- resistance to invasion (ERI = 3) covered 25.4% (91,317 km²) of the study area, while those with medium environmental resistance (ERI = 2) corresponded to 18.9% (67,942 km²), and those with
- medium environmental resistance (ERI = 2) corresponded to 18.9% ($67,942 \text{ km}^2$), and those with low resistance (ERI = 1) comprised only 9.6% ($34,684 \text{ km}^2$). Results for large basins (>1,000
- 391 km^2) are summarized in Fig. 5a.
- We mapped the interaction of potential habitats for Chinook salmon with different land
- use (i.e., forest, agriculture) and a hypothetical scenario of hydropower development in the
- Toltén River Basin (**Fig. 5b**). There were 2,071 km of high potential habitat for Chinook salmon in this basin representing low resistance to its invasion, from which 15% overlapped with
- in this basin representing low resistance to its invasion, from which 15% overlapped with
 agricultural lands and 16.4% with forest cover. In the hypothetical scenario of hydropower
- agricultural lands and 16.4% with forest cover. In the hypothetical scenario of hydropowerdevelopment, our model showed access obstruction to almost half of potential spawning habitats
- 398 (51.4%, 1,065 km) and 38.3% (794 km) of potential rearing habitats (**Fig. 5b**). This hydropower
- senario would block Chinook salmon access to 38% (129 km) of high potential streams co-
- 400 occurring with forest cover.



Figure 5. Chinook salmon potential habitats and environmental resistance. Panel (a) shows the
 distribution of environmental resistance in large basins (> 1,000 km²) throughout the study area.
 Very high resistance class sub-basins are not displayed since most are not accessible to

anadromous salmon). This was the dominant class for most basins. Panel (b) zooms into the

406 Toltén River basin (8,398 km²), displaying rivers coloured based on their potential for spawning

and rearing. Terrestrial pixels represent optimal (trees, green) and suboptimal (crops, yellow)

408land covers for salmon. Blocked environments by the hypothetical hydropower development

scenario appear as hatched areas. Map projection: South America Albers Equal-Area Conic.

- 410
- 411 *3.2.2.2 Coho salmon*

412 There were 153,222 stream km accessible to coho salmon in the study area (**Fig. 6a**). These

environments included 58,414 stream km with a high potential to sustain coho salmon rearing

and 62,356 km with a high potential for spawning. Sub-basins with very high resistance (ERI =

4) to coho salmon invasion represented 43.5% (156,641 km²) of the study area, whereas high
resistance sub-basins corresponded to 6.3% (22,506 km²). Medium resistance (ERI = 2) sub-

basins covered 21.3% (76,610 km²), and low resistance sub-basins were 28.9% (103,925 km²) of the study area.

We mapped environmental resistance to the invasion of coho salmon in the Lakes district 419 of southern Chile, a region with high historical aquaculture production (therefore, high levels of 420 propagule pressure due to aquaculture escapes). We identified three areas presenting a high risk 421 to the establishment of coho salmon populations, including the Reloncaví fjord and bay, the 422 interior river basins of Chiloé, and the area of Melimoyu-Puyuhuapi (Fig. 6b). In these three 423 424 areas, coho salmon aquaculture production has surpassed 40 million tonnes during the last three decades and is associated to persistent fish escapes from aquaculture facilities (Soto et al., 2001; 425 Sepúlveda et al., 2013; Gomez-Uchida, Sepúlveda, et al., 2018). The proximity of salmon farms 426 to low-resistance freshwater systems for coho salmon breeding and recruitment suggests a higher 427 428 likelihood for their success in establishing naturalised populations.

429 We also mapped potential habitats for coho salmon in basins near reported naturalised salmon in northern Magallanes (Górski et al., 2017; Fig. 6c) and the Beagle Channel (Chalde et 430 al., 2019; Maldonado-Márquez et al., 2020); Fig. 6d). From these reports, we evaluated 431 environmental resistance for ten basins, including four basins where coho salmon were not 432 detected (i.e., Maravilla River, Santa María River, Cambaceres River, Moat River). In basins 433 with coho salmon detections, our model showed that 56% of the area consisted of sub-basins 434 with low environmental resistance. In basins with no detections, only 35% of their area had low 435 resistance. Basins with the highest percentage of low environmental resistance areas included the 436 437 Coicopihue (100%), Puma (100%), Santa María (100%), and Varela (75%) river basins.



Figure 6. Coho salmon habitat and environmental resistance. Panel (a) shows the three areaswhere we provide illustrative applications of our approach for coho salmon. Panel (b) shows the

- 441 distribution of environmental resistance relative to coho salmon aquaculture production between
- the years 2010-2019 (Soto et al., 2022) and three labelled areas with the highest risk of invasion,
- 443 (c) the distribution of potential stream habitats for rearing and spawning near areas where
- 444 naturalised coho salmon have been captured (Górski et al., 2017), and (d) areas with detection of

445 coho salmon using environmental DNA (Chalde et al., 2019) and live captures (Maldonado446 Márquez et al., 2020). Map projection: South America Albers Equal-Area Conic.

447 448

4. Discussion

Region-wide surveillance and monitoring of invasive species are logistically unfeasible and costprohibitive, making spatial prioritisation tools extremely useful for increasing efficiency in the allocation of resources. We present a mechanistic habitat model to map environmental resistance to riverine invasions based on the availability and connectivity of potential habitats for multiple life stages of target species. We illustrate the benefits of this method with applications to three case studies of non-native species with varying environmental requirements, propagation history, invasion success, and perceived impacts on receiving ecosystems.

Our approach has advantages for invasion modelling due to its effective leveraging of 456 publicly available, high-resolution satellite data, and the simplicity of its interpretation. By 457 employing relatively static geomorphic predictors and long-term averaged climatic models, the 458 habitat intrinsic potential approach quantifies the geophysical capacity of freshwater habitats to 459 sustain certain biotic conditions at any given time or management scenario (Burnett et al., 2007). 460 Invasion risk assessments based on this modelling approach require fewer updates over time 461 while still allowing the integration of temporally dynamic data (e.g., propagule pressure, land 462 cover, future climatic scenarios) as complementary determinants of the expression of that habitat 463 464 potential for different target species.

The described "geophysical envelope" model assembly method can leverage disparate 465 information sources to generate transferable species-habitat models that are robust to the 466 uncertainties of niche expression in invaded regions. Species-habitat data about optimal and 467 tolerable environmental ranges can be collected from literature reviews, expert surveys, and field 468 studies for large spatial-temporal extents and multiple populations. By using the most extreme 469 470 bounds (e.g., 90th percentile) of environmental tolerance and optimality reported for a species, this simple assembly method can accommodate the broad range of uncertainty surrounding the 471 expression of ecological niches at invaded regions (Bates & Bertelsmeier, 2021). This 472 environmental resistance model is more robust to changes in habitat selection by invasive species 473 than model assemblies using central tendency metrics (e.g., averaged models), which can fail to 474 comprehensively represent the broad range of usable geophysical conditions by an invader. 475

By stacking multi-stage intrinsic potential models, our habitat complementation model 476 477 integrates critical attributes of landscape complexity to illustrate the probability of the establishment of invasive populations. Habitat modelling in stream ecosystems requires the 478 understanding of complex life-history traits, such as the requirement for different critical habitats 479 for different life stages (Solomon, 1973; Flitcroft et al., 2012, 2014). Thus, multi-stage habitat 480 models offer a more meaningful quantification of the landscape's potential to sustain populations 481 with complex life histories. This approach can also help identify life-stages requiring more 482 483 restricted environmental ranges, facilitating the identification of environments where monitoring and control efforts would be more effective. 484

The ability of this habitat model to integrate diverse sources of information has utility for case-specific applications in invasion risk assessments. We exemplify this by using land-cover data to identify environments with low geophysical resistance to invasion. In addition, the location and intensity of introduction events likely play an important role in the success and distribution of invasions (e.g., Arismendi et al., 2009). Habitat models compatible with propagule pressure data improve our ability to map invasion risk for species introduced within large spatial extents. Additional biotic variables (i.e., food, potential competitors, presence of
other invaders, presence/absence of predators) could also be integrated through dynamic models
representing invasion risk in varying scenarios (Buchadas et al., 2017; Soto et al., 2022);
however, this is at the expense of increased complexity of model development and interpretation.
This methodology can also be extended to study the risk of negative interactions between native
and invasive species by assessing the overlap between potential habitats among multiple species
(Jalbert et al., 2021; Manning et al., 2022).

498 We illustrate the advantages of our environmental resistance model to inform decisionmaking and resource allocation with applications to three priority species invading a large region 499 of Patagonia. For North American beaver, the application demonstrates its utility in identifying 500 501 overlaps between areas with low environmental resistance and priority management zones (i.e., removal and active control zones) that were previously designed by local agencies, suggesting a 502 high risk of expansion onto the continent where invasion control could become logistically 503 504 unfeasible. The study region shows high abundance and connectivity of potential habitats for invasive Chinook salmon, although with marked patterns of regional heterogeneity that could 505 inform the management and monitoring of the species. Our findings also illustrate that northern 506 507 areas, where there is a concentration of Chinook salmon stakeholders (e.g., Toltén River basin) have the lowest environmental resistance to salmon invasion. However, this region presents 508 higher rates of land degradation (e.g., agriculture, forestry, hydropower development) as 509 510 compared to more pristine, though less abundant, low-resistance environments in southern basins. Our model's application to coho salmon exemplifies its potential to inform the research 511 of invasive species with long-range dispersal capacity, for which risk areas can encompass vast 512 spatial extents. We identify large areas (> 10,000 km²) with high coho salmon invasion risk, 513

- 514 where low resistance systems co-occur with high propagule pressure from net-pen aquaculture
- escapes (e.g., Reloncaví, Interior Chiloé, and Melimoyu-Puyuhyapi), and where naturalised coho
- salmon have been recently reported (e.g., Górski et al., 2017; Chalde et al., 2019; Maldonado-
- 517 Márquez et al., 2020). These results have an immediately useful management application by518 aiding in the planning and prioritisation of field monitoring.
- In conclusion, we contribute to advancing research in the biogeography of invasions with 519 an environmental resistance model for riverine landscapes. Our habitat intrinsic potential 520 approach is readily transferable to other widely introduced species as intrinsic potential models 521 exist for many of them (e.g., rainbow trout, Burnett et al., 2007; northern pike, Jalbert et al., 522 523 2021; brook trout, Manning et al., 2022). Complex and unique relationships between species and their environments can be assessed using patterns of habitat complementation as critical 524 determinants of the sustainability of invasive populations. Applications of this flexible and 525 transferable method may have disproportionately high value when applied to invasions occurring 526 527 in data- and resource-poor regions by aiding in the spatial allocation of efforts to monitor the
- 528 occurrence, success, and impacts of invasive species on receiving ecosystems.
- 529

530 Data Availability

- 531 The data supporting the findings of this study are publicly available through Dryad
- 532 (<u>https://doi.org/10.5061/dryad.3tx95x6jf</u>) and the source code through Zenodo
- 533 (<u>https://doi.org/10.5281/zenodo.6618378</u>).

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- 789
- 790 **Conflict of Interest**

- 791 The authors declare no conflicts of interest.
- 792

793 Biosketch

- **J. Andres Olivos** is interested in the landscape ecology of aquatic biodiversity and gives
- rephasis on the effects of climate and land-use changes, hydrological disturbances, and
- biological invasions, as the most pressing issues for sustainability and conservation. This article
- is a coproduction with a multi-disciplinary research team working on the management and
- conservation of fish and wildlife at broad spatiotemporal scales.
- 799

800 Authors contributions

J.A.O. led the conceptualization, data collection, geospatial modelling, analysis, elaboration of

figures, and writing of the manuscript. **I. A.** and **B. P.** provided critical guidance in the

- 803 conceptualization and writing of the manuscript. **R. F.** and **G. G.:** provided extensive feedback
- and guidance. A. H. H., and J. F.: provided comprehensive datasets for the development and
- validation of habitat models. All authors reviewed, edited, and approved the submission of the
- 806 manuscript.