RESEARCH PAPER

Spatial pattern of freshwater habitats and their prioritization using additive partitions of beta diversity of inhabitant piscine assemblages in the Terai–Dooars ecoregion of Eastern Himalayas

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Abstract

Underlying spatial and habitat attributes of a river network are crucial to comprehend the bio-spatial arrangements within it, the study of which sufers from a paucity of information. Despite several reports on various piscine assemblages, no study contributes to understanding the characteristic attributes of the freshwater habitats of the sub-Himalayan Terai–Dooars ecoregion. Therefore, this study aims to uncover such underlying features through a precise understanding of the spatial profle of freshwater habitats and additive partitions of piscine beta diversity. A signifcant spatial association is found in the upper stretches of most of these torrential freshwater reaches confned to the eastward of the River Teesta basin to the tributaries of River Jaldhaka. Such a pattern is aligned with a higher local contribution to beta diversity (LCBD) values. The spatial map of LCBD indicates that the mid-altitude (100 > elevation > 2000 m) region contains unique or rare species assemblages. This fact is further confrmed by the spatial aggregation of characteristically adapted hill stream fsh species with higher species contribution to beta diversity (SCBD) values. The results are further explained by relevant climatic, topographic, nutrients (sediments), and habitat attributes of which climate, topographic, substrate, and land cover features are the most contributory factors. Such variables are subjected to severe modulation following increasing anthropogenic pressure and changing climatic conditions, leading to the jeopardy of these freshwater habitats. Therefore, prime importance should be accorded to the ecological restoration value of these spatially structured torrential freshwater habitats for conservation and monitoring in the coming days.

Keywords Fish · Spatial Structures · Beta Diversity · LCBD · SCBD · Terai–Dooars · Eastern Himalayas

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Introduction

Spatial arrangements of biodiversity varying upon the different spatial scales need to be understood for efficient conservation assessment and planning (Ferrier et al. [2007;](#page-13-0) Margules and Pressey [2000\)](#page-14-0). The assemblage of both native and exotic species within a given landscape is contingent on the degree of spatial heterogeneity (Davies et al. [2005\)](#page-12-0). However, the spatial process directly corresponds to the species sorting following the dynamics of environmental gradients, attributed by various factors at a given spatial extent (Cottenie [2005](#page-12-1); Heino et al. [2015a](#page-13-1); Jackson et al. [2001](#page-13-2)). Therefore, understanding the active spatial flters for a species at varied habitats is crucial, resulting from dispersal limitation, neutral dynamics, and spatially structured environmental attributes (Leprieur et al. [2009;](#page-14-1) Sharma et al. [2011\)](#page-14-2). As in-feld sampling over a large space is not convenient with time and effort, it persuades the employment of choosing

surrogates for the species under study (Ferrier et al. [2007](#page-13-0)). Such surrogates include habitat types, stream sections, landscapes often stemmed from remote sensing imagery, abiotic environmental classes, climate, and terrains (Ferrier et al. [2007](#page-13-0)).

Beta diversity accounts for the change in species composition between certain places representing the diferentiation component of diversity and assemblage (McKnight et al. [2007](#page-14-3)). Being central to a wide array of ecological and evolutionary understanding (McKnight et al. [2007\)](#page-14-3), beta diversity has long been used on many aquatic organisms to describe crucial turnover of species with the response to specifc environmental characteristics resulting from temporal and spatial dissimilarities (Angeler [2013;](#page-12-2) Melchior et al. [2017](#page-14-4); Viana et al. [2016\)](#page-14-5). Legendre and De Cáceres [\(2013\)](#page-13-3) have developed another two components to emphasize the contribution of habitat towards the existing pattern of dissimilarities in biodiversity (Vilmi et al. [2017](#page-15-0); Yao et al. [2020](#page-15-1)). These components are described as a local contribution to beta diversity (LCBD), indicating the uniqueness of the site and its degradation as well as the species contribution to beta diversity (SCBD), indicating unique and rare assemblages of species with conservation values (Legendre and De Cáceres [2013;](#page-13-3) Sor et al. [2018;](#page-14-6) Vilmi et al. [2017\)](#page-15-0).

The freshwater ecosystem depicts an epitome of researches concerning distinct species assemblage, spatial heterogeneity, and insuperable barriers (Leprieur et al. [2009\)](#page-14-1). The freshwater fsh communities are regulated by niche diferences and dispersal limitations following ecological changes limited at a connected stretch of waters (Chase and Leibold [2003](#page-12-3); Heino et al. [2009;](#page-13-4) Hubbell [2001](#page-13-5)). Climate, energy, and habitat diversity work in close association, resulting in a diverse piscine assemblage globally (Guégan et al. [1998](#page-13-6); Leprieur et al. [2011\)](#page-14-7). Therefore, dismantling spatial features arising at a particular site and relevant local environmental, catchment, and climatic factors is critical (Heino et al. [2007](#page-13-7); Perez Rocha et al. [2018\)](#page-14-8) in shaping freshwater fsh biodiversity. The lotic habitats of freshwater fshes are under serious threat. They are subjected to climate change, channel modifcation, fragmentation, fow alteration, and degradation (Bhatt et al. [2016;](#page-12-4) Goswami et al. [2012a,](#page-13-8) [b](#page-13-9)).

Such threats, coupled with the observed trend of anthropogenic infuences, pose more considerable infuence over the Himalayas (Singh [2015\)](#page-14-9). The freshwater network primarily prevalent in the sub-Himalayan Terai–Dooars (TED) ecoregion has been prioritized for conservation based on the richness of fsh species with higher conservation values, endemism, and vulnerabilities (Bhatt et al. [2016](#page-12-4)). The major rivers of the TED ecoregion in Northern Bengal, India, are River Teesta, Jaldhaka, Torsha, and Mahananda (Jayaram and Singh [1977](#page-13-10)). However, understanding the spatial distribution and underlining species sorting in such exclusive freshwater habitats is overlooked due to much research efort into the identifcation of fsh species, reporting exclusive alpha diversity at various local scales (Bhowmik et al. [2016](#page-12-5); Dey et al. [2015a](#page-12-6) ,[b](#page-12-7)[, c](#page-12-8), [d](#page-12-9)[, 2019\)](#page-12-10), eco-physiological studies in captivity (Dey et al. [2015a](#page-12-6); [b,](#page-12-7) [c,](#page-12-8) [d](#page-12-9)), preserving germplasm and reporting genetic variabilities (Dey et al. [2015a](#page-12-6); [b](#page-12-7), [c,](#page-12-8) [d](#page-12-9); Kundu et al. [2019\)](#page-13-11). A frm understanding of spatial arrangements of fsh communities would reveal the complex interaction of multiple factors, which shapes their habitats in a more varied or nested manner considering their adaptations, invasions, dispersal limitations, and mass efects (Leprieur et al. [2009;](#page-14-1) Leroy et al. [2019;](#page-14-10) Planque et al. [2011](#page-14-11); Shurin et al. [2009;](#page-14-12) Wiersma and Urban [2005\)](#page-15-2). Such a pattern gives direct insight into the response of biological communities to climate and environment, which often serves as a basis of sound management of their commercial or conservation interest (Leprieur et al. [2009;](#page-14-1) Planque et al. [2011](#page-14-11)). Therefore, the lack of such foundation stumbles conservation planning and monitoring over this freshwater habitat, becoming more vulnerable following changing geo-climatic conditions (Akhter et al. [2019;](#page-11-0) Barman and Das [2014](#page-12-11); Goswami et al. [2019\)](#page-13-12) and increasing anthropogenic pressure (Karmakar [2011](#page-13-13); Naha et al. [2019](#page-14-13); Singh [2015\)](#page-14-9).

Therefore, this study reveals the pattern and reasons behind the unique fsh assemblage in the sub-Himalayan TED ecoregion freshwater network. We delineated the signifcant spatial variables explaining the spatial structure of fish assemblages using spatial distance among the sites, identifying if the scales have a signifcant association. Then, we conducted a decomposition of the spatial model at relevant scales into submodels, aiming to reveal species–environment relationships. Next, using beta dissimilarities, we analyzed the individual contribution of fsh species and their habitat through partitioning the total beta diversity while explaining the latter by relevant environmental variables. The results were compared to identify the freshwater habitats and their correspondence in unique piscine assemblage, subjected to ecological conservation and management considering the threats of dynamic climatic and anthropogenic events in the sub-Himalayan TED ecoregion. This study signifes the frst attempt to construct such a profle from this region to explore ecological sensitivity in species conservation values and eco-restorations.

Materials and methods

Study area

The sub-Himalayan TED ecoregion has moist and dense riverine forests along the foothills of the snow-capped Kanchenjunga range along the Eastern Himalayas (Barman and Das [2014](#page-12-11); Kandel et al. [2016](#page-13-14)). Northern Bengal (NB) comprises the areas within West Bengal, India, confned to

the north of the river Ganges (Barman and Das [2014](#page-12-11)). Innumerable streams are draining these alluvial foodplains of the TED ecoregion in NB (Barman and Das [2014;](#page-12-11) Paul et al. [2009;](#page-14-14) Rudra [2018\)](#page-14-15). The freshwater reaches of this region are more dynamic due to continuous deposition in the channel, increasing the height of the riverbeds (Chakraborty and Datta [2013](#page-12-12)). The river channels are experiencing frequent shifts following anabranching and changing river courses (Akhter et al. [2019;](#page-11-0) Goswami et al. [2019\)](#page-13-12) Sub-tropical monsoon climate causes excessive precipitation in the sub-Himalayan regions leading to the higher flow in these torrential courses. In the summer months, they are replenished by snow-melt waters (Akhter et al. [2019;](#page-11-0) Bhatt et al. [2012](#page-12-13); Panja et al. [2020](#page-14-16); Rudra [2018\)](#page-14-15). This study has been conducted on a vast drainage network of Teesta–Neora–Jaldhaka rivers, including watersheds of River Teesta, River Chel, River Neora, River Dharala, River Murti, and River Jaldhaka, draining through the TED ecoregion in NB, India (Fig. [1](#page-2-0)). Along the banks of these rivers, the TED ecoregion has several reserve forests and national parks, such as Singalila National Park, Mahananda Wildlife Sanctuary, Neora

Valley National Park, Gorumara National Park, and Chapramari Wildlife Sanctuary, which indicate the importance of this ecoregion (Bhattacharya [2019\)](#page-12-14). A stream network of these freshwater reaches was derived using digital elevation layers (MERIT-HYDRO DEM) (Yamazaki et al. [2019](#page-15-3)) in the Arc GIS platform (V.10.1).

Fish sampling

A total of 31 sampling sites were selected based on a pilot study upon these freshwater reaches. Later, a tri-seasonal fish sampling, i.e., pre-monsoon, monsoon, and post-monsoon, was conducted during 2016–2019. A 90 m reach, at each location was sampled using the electro-fishing method by electro-fsher (300 V, 3–4A, DC) followed by gill nets, cast nets, and dragnets, respectively. The dimensions of the nets used in samplings were constant (obtained from the same source) for all the areas surveyed. Fishes were identifed following existing literature (Barman and Das [2014](#page-12-11); Jayaram and Singh [1977;](#page-13-10) Menon [1999](#page-14-17); Shaw [1938](#page-14-18); Talwar and Jhingran [1991](#page-14-19)). The removal method of estimation was

Fig. 1 Study area comprising a freshwater network of sub-Himalayan Terai–Dooars ecoregion confned into Northern Bengal of Eastern Himalayas

applied in three consecutive eforts (Bohlin et al. [1989\)](#page-12-15). The captured fsh specimens were counted, and a single representative was preserved in a 10% formalin solution.

Environmental data

Seven bioclimatic variables, four topographic, two substrates, and two land-cover attributes had been considered under the environmental profle. The ecological success, reproductive behaviors, and physiology of freshwater fshes are signifcantly driven by variability in temperature and precipitation (Barbarossa et al. [2021](#page-12-16); Ficke et al. [2007\)](#page-13-15). Despite being coarse-scale modulators, previous studies (Domisch et al. [2011](#page-12-17), [2015](#page-12-18), [2013](#page-12-19), [2011](#page-14-7); Durance and Ormerod [2007](#page-13-16); Leprieur et al. [2009;](#page-14-1) Oberdorff et al. [1999](#page-14-20); Reyjol et al. [2007\)](#page-14-21) found the signifcant contribution of annual mean temperature, the max temperature of the warmest month, min temperature of the coldest month, annual precipitation, precipitation of wettest month, precipitation of driest month and evatransportation in shaping the spatial assemblage of freshwater fshes. Furthermore, the topographical characteristics of streams, i.e., elevation, slope, stream order, and terrain position index, directly correspond to local ecological attributes of freshwater habitats viz. water temperature, dissolved oxygen, and fow regimes (Austin [2007;](#page-12-20) Domisch et al. [2011](#page-12-17), [2013](#page-12-19); Kuemmerlen et al. [2014\)](#page-13-17). On the other note, stream substrate characteristics (upland & valley bottom characteristics, soil sediment) and landcover attributes (normalized diference vegetation index and land cover) control the productivity, pH, turbidity, and nutrient dynamics of water which are responsible for various species-specifc adaptations (Brooks et al. [2005;](#page-12-21) Efenberger et al. [2006](#page-13-18); Fausch et al. [2002;](#page-13-19) Kozel and Hubert [1989\)](#page-13-20). Therefore, these ffteen variables (Table [1\)](#page-3-0) are pertinent to understand freshwater species distribution. All these variables were obtained in 30 arc-second resolution and sampled for the selected 31 sampling sites. The respective sources of the environmental data (Haynes et al. [2018;](#page-13-21) Huntington et al. [2017;](#page-13-22) Karger and Zimmermann [2019;](#page-13-23) Pelletier et al. [2016](#page-14-22); Trabucco and Zomer [2019](#page-14-23); Yamazaki et al. [2019\)](#page-15-3) are listed in Table [1.](#page-3-0) Slope, stream order (Strahler), and topographic position index were calculated using the DEM raster (Yamazaki et al. [2019](#page-15-3)) in the QGIS platform (QGIS version 3.10.0-A Coru $\tilde{A} \pm a$) (<https://www.qgis.org/en/site/>). Before fitting into analytical models, these variables were resampled, standardized, and stacked for the study region and sampled against the 31 sites to conduct a Pearson's correlation among them. Predictors with high collinearity (Pearson's $r \ge 0.8$) (Domisch et al. [2011](#page-12-17); Thuiller et al. [2014\)](#page-14-24) were discarded (See Supporting Information: Fig. S1).

Data analysis

Identifcation of the spatial structures

The spatial arrangements of habitats signifcantly structure the fish community composition considering space, scale, and connections within a landscape (Jackson et al. [2001](#page-13-2); Legendre and Legendre [2012;](#page-13-24) López‐Delgado et al. [2019](#page-14-25)).

Table 1 List of environmental variables used as explanatory variables for the fsh assemblage patterns in the freshwater river system of Terai*–* Dooars ecoregion, Eastern Himalayas

Category	Variables	Abbreviation	Source
Bio-Climatic	Annual mean temperature	B1	Climatologies at high resolution for the earth's land surface areas (https://chelsa-climate.org/) (Karger and Zimmermann 2019)
	Max temperature of the warmest month	B ₅	
	Min temperature of the coldest month	B6	
	Annual precipitation	B12	
	Precipitation of wettest month	B13	
	Precipitation of driest month	B14	
	Eva-transportation	EV	Potential evapotranspiration climate Database version 2 (https://cgiarcsi.community/) (Trabucco and Zomer 2019)
Topography	Elevation	EL	MERIT Hydro: global hydrography data sets (http://hydro.iis.utokyo.ac.jp/yamadai/MERIT_Hydro/) (Yamazaki et al. 2019)
	Slope	SL	Calculated using EL in QGIS 3.16.0
	Stream order	SO	
	Terrain position index	TPI	
Substrate	Upland & valley bottom characteristics	UV	Oak ridge national laboratory distributed active archive center (ORNL) DAAC) (https://daac.ornl.gov/) (Pelletier et al. 2016)
	Soil sediment	SS	
Land Cover	Normalized difference vegetation index	NDVI	Climate Engine (http://climateengine.org/) (Huntington et al. 2017)
	Landcover	LC	IPUMS-TERA (https://terra.ipums.org/) (Haynes et al. 2018)

This widely recognized concept would reveal the role of space in habitat heterogeneity and spatial aspects of biotic and abiotic conditions (Hanski [2001;](#page-13-25) Jackson et al. [2001](#page-13-2)). The framework of Moran's eigenvectors map (MEM) has gained popularity as a recent family of spatial analysis techniques (Ali et al. [2010](#page-11-1); Jackson et al. [2001](#page-13-2); Legendre and Legendre [2012;](#page-13-24) Perez Rocha et al. [2018](#page-14-8)). This machinery aims to model the correlation structure present at each scale, linking with the spatial heterogeneity of environmental factors (Ali et al. [2010](#page-11-1)). Based on this framework, a distancebased Moran's eigenvectors map (dbMEM) (Dray et al. [2006\)](#page-12-22) would identify the characteristic spatial scales through a spatial fltering technique (Blanchet et al. [2008](#page-12-23)) to defne a set of spatial proxy variables and their selection to explain the spatial structure of the response variable under study, i.e., fsh species assemblage. The dbMEMs corresponding to smaller eigenvalues usually represents very fne-scale spatial patterns, where spatial autocorrelation is presumed low. On the contrary, dbMEMs corresponding to larger eigenvalues signify coarse spatial variability scales, often selected to defne prominent spatial structures. Furthermore, dbMEMs corresponding with positive eigenvalues (Positive Moran's I) depicts a positive spatial association, which is more crucial to consider than a negative spatial association (Ali et al. [2010;](#page-11-1) Grifth and Peres-Neto [2006](#page-13-26); Legendre and Legendre [2012](#page-13-24)).

Initially, the spatial coordinates of the sites and a Hellinger transformed presence–absence data set was subjected to defne dbMEMs (Blanchet et al. [2008](#page-12-23); Dray et al. [2006](#page-12-22); Legendre and Legendre [2012](#page-13-24)) and their spatial association with the fish species assemblage of the freshwater habitats. The data was checked for linear trends and detrended if present. The dbMEM eigenvectors were computed, and those with positive association (positive Moran's I) were retained. Then the dbMEMs were tested in a redundancy analysis (RDA) with species data for signifcance and run a forward selection with double-stopping criteria (adjusted R square of the RDA and α level of rejection) (Blanchet et al. [2008](#page-12-23)) to select signifcant dbMEMs. Now a new RDA was run with the signifcant dbMEMs to tests the signifcance of the axes. Based on the results, maps were drawn for each signifcant axes identifying spatial structures among the sites. (Borcard and Legendre [2002](#page-12-24); Borcard et al. [1992;](#page-12-25) Dray et al. [2012;](#page-12-26) Legendre and Legendre [2012\)](#page-13-24). This analysis was performed in the R platform with the *quickMEM* function from package *adespatial*. The total variation in the fsh species assemblage was further partitioned using the spatial (dbMEMs) and environmental variables (Perez Rocha et al. [2018](#page-14-8); Vilmi et al. [2017\)](#page-15-0) to assess their relative and cumulative roles through partial RDA testing.

The site scores of each signifcant RDA axes were further explained by the environmental variables (Borcard and Legendre [2002](#page-12-24); Borcard et al. [1992;](#page-12-25) Legendre and Legendre [2012\)](#page-13-24) to identify the signifcant association of environmental variables resulting in positive-broad scale spatial structures. This assessment was achieved by applying boosted regression trees (BRT) models (Elith et al. [2008\)](#page-13-27). The boosted regression tree (BRT) is much exploited as an excellent modeling tool for the predictive purpose of ecological researches (Elith et al. [2008\)](#page-13-27). However, BRT modeling with smaller data set would face a slight penalty for using larger trees which are usually overcome using low learning rates and smaller decision trees (Elith et al. [2008](#page-13-27)). BRT is advantageous in accommodating diferent predictors, with no dependencies on response data transformation, outlier removal, and handling complex non-linear relationships while considering interactions between the predictors to reduce predictive errors (Carslaw and Taylor [2009](#page-12-27); Elith et al. [2008](#page-13-27); Jafari et al. [2014](#page-13-28); Panja et al. [2020](#page-14-16)). Considering such an advantage, previous studies have appropriately used this machine learning model using a small data set (Jafari et al. [2014](#page-13-28); Panja et al. [2020](#page-14-16), [2021b](#page-14-26), [a;](#page-14-27) Zhang and Ling [2018\)](#page-15-4). A combination of learning rates (high to low) was tried to achieve the minimum 1000 trees initially based on the tenfold cross-validation method. Since the present study dealt with a comparatively smaller data set, tree complexity was set to 3 (Elith et al. [2008\)](#page-13-27). A pseudo determinant factor, $D²$ was calculated for the fitted model accounting for their credibility (Nieto and Mélin [2017\)](#page-14-28). The environmental variables contributing highest to the model are identifed.

Beta diversity measures

Legendre and De Cáceres ([2013\)](#page-13-3) emphasized the advantages of estimating beta diversity (BD_{Total}) as the total variation of the community matrix (Z) . They showed that Z could be linked with the beta diversity assessments computed from the dissimilarity matrix of community composition while establishing the closer relevance of other beta diversity measures (Anderson et al. [2006](#page-12-28); Ricotta and Marignani [2007;](#page-14-29) Whittaker [1972\)](#page-15-5). Based on such principle, BD_{Total} could be disintegrated into two attributes accounting for contributions of single sites to overall beta diversity and individual species to overall beta diversity (Heino and Grönroos [2017;](#page-13-29) Legendre and De Cáceres [2013](#page-13-3)). The former is referred to as local contributions to beta diversity (LCBD) which comparatively indicates the ecological uniqueness of the study sites. The latter attribute is defned as species contributions to beta diversity (SCBD) which denotes the degree of variation of individual species across the study region (Legendre and De Cáceres [2013\)](#page-13-3). The LCBD values identify sites with more (or less) contribution than the mean to beta diversity, which comparatively indicates the uniqueness of species composition at a particular site. LCBD directly corresponds to the number of rare species across geographical space, while were negatively related to the occurrences of common species, therefore, accounting for dispersal limitation as well as the local environment and community characteristics (Legendre and De Cáceres [2013](#page-13-3); Vilmi et al. [2017](#page-15-0); Yao et al. [2020\)](#page-15-1). On the other note, SCBD indices are not the same as indicator species for a given group of sites. Instead, it directly corresponds to signifcant variations imposed by each species across the study area (Cáceres and Legendre [2009](#page-12-29); Dufrêne and Legendre [1997;](#page-13-30) Legendre and De Cáceres [2013](#page-13-3)).

Z could be obtained by calculating a matrix of squared deviations along the column means based on computing community dissimilarity matrices from the transformed species presence–absence data. Then the total sum of squares (SS_{Total}) is obtained when summing all the squared values, which forms the initial basis of BD _{Total}.

 $BD_{\text{Total}} = \frac{SS_{\text{Total}}}{(n-1)}$,n represent the number of sites (Legendre and De Cáceres [2013\)](#page-13-3).

Computation of SS_{Total} is advantageous to directly assess the contributions of individual species and individual sampling units to the overall beta diversity (Heino and Grönroos [2017](#page-13-29); Legendre and De Cáceres [2013](#page-13-3)). Therefore, SCBD and LCBD indices were developed on total sum squares of species composition, thereby calculating the proportion of jth species and the ith sampling unit for SCBD and LCBD viz. $SCBD_j = SS_j/SS_{Total}$ and $LCD_i = SS_i / SS_{Total}$ (Legendre and De Cáceres [2013\)](#page-13-3). Sites with higher LCBD values are identified. The fish species with higher SCBD values than the average value of all are identifed.

BRT was ftted similarly with the LCBD values against the predictor environmental data set. The signifcant variables were identifed, contributing highest to explain the LCBD attributes of each site and compared with the spatial model. LCBD site scores were further assessed through a correlation approach with the signifcant RDA axes about surrogacy in conserving and monitoring these freshwater ecosystems. Based on the best trees of the suited model, the LCBD was predicted over the restacked spatial map

of selected environmental variables to construct a spatial LCBD profle for this study area.

All the statistical analyses have been performed in the R platform (Team R [2015](#page-14-30); Team RC [2013\)](#page-14-31) using packages, namely, *raster, maptools, rgeos, dismo,* and *gbm*. The spatial maps are created using the Arc GIS platform (version 10.1).

Results

Initially, a total of 175 fsh species were recorded from the sampling. However, discarding exotic species (See Supporting Information: Table S1) from the study (Bhowmik et al. [2016](#page-12-5); Sarkar and Pal [2018](#page-14-32)), a total of 170 indigenous fsh species with 11,560 individuals has been recorded from the sampling along 31 sites (See Supporting Information: Table S2 and Fig. S2). River Jaldhaka has the highest species richness with 158 fsh species. In comparison, the lowest has been observed in River Neora with 22 fish species.

Partitioning spatial component of ecological variation

A total of nine spatial variables, i.e., dbMEM eigenvec-tors (Table [2\)](#page-5-0), have been produced with R^2 of the global $model = 0.484$. Among them, four dbMEM eigenvectors have been forward selected (Table [2](#page-5-0)) at $p < 0.001$. In the fnal RDA model, two canonical axes are signifcant, i.e., RDA 1 (Axes1) at *p*<0.001 and RDA 2 (Axes 2) at *p*<0.05 (Table [2;](#page-5-0) Fig. [2](#page-6-0)), explaining the underlying spatial association of sites resulting into species sorting in these selected freshwater habitats. It appears that Axes 1 (Fig. [2\)](#page-6-0) indicates substantial positive spatial infuence on the species assemblages around the upper stretches of most of these freshwater reaches confned to the TED ecoregion. However, Axes 2 (Fig. [2\)](#page-6-0) indicates a secondary spatial pattern occurring specifcally in River Teesta and its tributaries around the mid altitudinal zone.

In variation partitioning, the selected spatial variables significantly influence the fish species variation in the partial RDA model at $p < 0.001$ (See Supporting

Table 2 Details of analysis uncovering the broadscale spatial pattern of fish assemblage in the freshwater river system of Terai*–*Dooars ecoregion, Eastern Himalayas

¶*dbMEM* Distance-based Moran's eigenvector, *MEM* Moran's eigenvector, *RDA* Redundancy analysis

Fig. 2 Signifcant canonical axis of redundancy analysis: Axes 1 and 2 of four forward selected dbMEM eigenvectors in Distance-based Moran's Eigenvector Maps (dbMEM) analysis explaining the spatial association of freshwater habitats regarding diferential fsh assem-

blages in the freshwater network of sub-Himalayan Terai–Dooars ecoregion. (Darker and larger the circle indicates the larger and positive eigenvalues depicting stronger positive spatial association)

Information: Table S3). The purely spatial components have explained 22%, while purely environmental factors have explained the variation of 11%. However, they both cumulatively address 2% of the total constrained variation (Fig. 3).

Additive components of beta dissimilarity

The higher LCBD values are projected in the upper stretches of River Chel, Neora, Murti, Jaldhaka, and mid altitudinal reaches of River Teesta (Fig. [4](#page-7-1)a). RDA Axes 1 depicting substantial spatial patterns in the sites are highly correlated (Pearson's $r > 0.8$) with the respective LCBD values (Fig. [4b](#page-7-1)). The higher SCBD values are observed for 54 fish species. The top ten fish species with significant contribution are *Neolissochilus hexastichus, Neolissochilus hexagonolepis*, *Garra lamta, Tor tor, Badis badis, Devario aequipinnatus, Garra annandalei, Crossocheilus latius, Amblyceps mangois, and Barilius vagra* (Fig. [5\)](#page-8-0).

Boosted regression models

A Pearson's correlation test has identifed the highly correlated predictors among ffteen predictor environmental variables. As the highly correlated variables would increase the uncertainty in the model and decrease robustness, they were removed from the predictor set (See Supporting Information: Fig. S1). Therefore, eight variables have been retained, which are annual mean temperature (B1), annual precipitation (B12), precipitation of driest month (B14), slope (SL), stream order (SO), land cover (LC), upland & valley bottom characteristics (UV) and soil sediment (SS).

B14, SO, UV, LC, B1, B12, and SL have a more substantial infuence in the spatial model (Table [3](#page-8-1)), explaining Axes 1 with BRT (See Supporting Information: Fig. S3). However, B12, UV, B14, SO, LC & B1 are relevant (Table [3\)](#page-8-1) to explain Axes 2 in the BRT model (See Supporting Information: Fig. S3). Therefore, B14, B12, B1, SO, UV, and LC are cumulatively responsible for the underlying spatial profle in the freshwater habitats of differential fish species assemblage. On the other note, LCBD values of these freshwater

Fig. 3 Variation partitioning of environmental and spatial variables explaining piscine assemblages in the freshwater network of sub-Himalayan Terai–Dooars ecoregion

Fig. 4 a Local contribution to beta diversity (LCBD) values of freshwater habitats in the freshwater network of sub-Himalayan Terai– Dooars ecoregion (Larger the circle indicates the higher values). **b** Correlation between the LCBD values and signifcant RDA axes (A1:

Axes1 & A2: Axes 2) of the spatial model depicting the spatial association of freshwater habitats in the sub-Himalayan Terai–Dooars ecoregion

Fig. 5 List of 54 fsh species with higher species contribution to beta diversity (SCBD) values in the freshwater network of sub-Himalayan Terai–Dooars ecoregion. (For species names, see Supporting Information: Table S2)

Table 3 Modelling output with boosted regression trees using environmental predictors of fsh assemblage in the freshwater river system of Terai*–*Dooars ecoregion, Eastern Himalayas

Sl. No.	Feature modeled with boosted regression trees	Cross-valida- tion correla- tion	Pseudo determi- nant coefficient (D^2)	Variables with a higher relative influence (>5)
	RDA Axes 1	$0.13 + 0.03$	0.86	Precipitation of the driest month, Stream order, Upland & valley bottom, Landcover, Annual mean temperature, Annual precipitation & Slope
2.	RDA Axes 2	$0.13 + 0.04$	0.85	Annual precipitation, Upland & valley bottom, Precipitation of the driest month, Stream order, Landcover & Annual mean temperature
3.	LCBD.	$0.05 + 0.24$		Precipitation of the driest month, Stream order & Annual mean tempera- ture

RDA Redundancy analysis, *LCBD* Local contribution to β diversity

habitats are best explained by B14, SO $\&$ B1 (Table [3](#page-8-1)) in the BRT model (See Supporting Information: Fig. S3).

The projected spatial map of LCBD (Fig. [6](#page-9-0)) following the best trees projects higher values in the upstream reaches of River Jaldhaka, Murti, Chel, Neora. Comparatively, lower LCBD values are dispersed around the plains of Terai and Dooars, confned into the district of Jalpaiguri and Cooch Bihar. Moderate LCBD values are predicted within mid altitudinal stretches of River Teesta.

Discussion

In this study, a prominent spatial association determining the fish assemblages among the water reaches is observed. Such spatial features contribute at large to the diferential fish species assemblage in these freshwater habitats. At the same time, they are substantially associated with climatic (precipitation of driest month, annual precipitation, annual mean temperature), topographical (stream order), substrate (upland & valley bottom characteristics), and land cover (landcover) characteristics of these freshwater reaches of TED ecoregion of NB. On the other note, the LCBD profle

is aligned with axes 1 of signifcant spatial association of these freshwater habitats. However, such profle is best explained by climatic (precipitation of driest month, annual mean temperature) and topographical (stream order) attributes. Furthermore, higher SCBD values emphasized 54 fsh species which are diferentially distributed among these freshwater reaches of the TED ecoregion of NB.

The freshwater habitats of this study belong to diferent freshwater river reaches that are not necessarily connected (Fig. [1\)](#page-2-0). Our result suggests a spatial metacommunity structure among the freshwater habitats presumed to result in these broad-scale spatial associations (Grönroos et al. [2013](#page-13-31); Heino et al. [2015a](#page-13-1), [b;](#page-13-32) Thompson and Townsend [2006](#page-14-33)). However, the more delicate spatial association is not achieved as they are hard to discern (Ali et al. [2010](#page-11-1); Legendre and Legendre [2012](#page-13-24)) and might be associated with local mass efect dynamics (Borcard and Legendre [2002;](#page-12-24) Heino et al. [2015a,](#page-13-1) [b](#page-13-32)). Such an approach is apposite considering the identification of broad-scale spatial structures and coarse-scale resolution of relevant environmental variables explaining them. Broad-scale spatial variables tend to be associated with dispersal limitation (Heino et al. [2015a,](#page-13-1) [b;](#page-13-32) Heino et al. [2015a](#page-13-1), [b](#page-13-32); López‐Delgado et al. [2019](#page-14-25)) as the watercourse

Fig. 6 Projection of spatial map of local contribution to beta diversity (LCBD) of freshwater habitats in the freshwater network of sub-Himalayan Terai–Dooars ecoregion

distances are more stringent for stream organisms (Altermatt [2013](#page-12-30)). Therefore, the community dynamics of each habitat rely signifcantly upon spatial autocorrelation (SA) (Shurin et al. [2009](#page-14-12)) than changes in rates of movements.

The results indicate a substantial role of climate, topography, landscape, and substrate features behind the prominent positive spatial association of the habitats. Such fndings are relevant as a signifcant geomorphological recess has been observed in the east of River Teesta to River Jaldhaka–Diana channel, further segmented by Chel–Mal, Mal–Murti, Jaldhaka–Gathia interfuves (Goswami et al. [2019\)](#page-13-12). This region is predominantly manifested by a transitional zone between the Eastern Himalayan mountains and the upper Gangetic plains (Chakraborty and Datta [2013\)](#page-12-12), where a marked diference exists in annual temperature, precipitation, and climate extremities (Das [2020](#page-12-31); Panja et al. [2021b](#page-14-26), [a](#page-14-27); Rudra [2018](#page-14-15); Sam and Chakma [2019](#page-14-34)). The substrate composition of freshwater channels is characterized by the piedmont fans, channel deposition, frequent shifting of courses (Chakraborty and Datta [2013](#page-12-12); Goswami et al. [2019](#page-13-12)) which cumulatively

modulated by topography. Such characteristics might have resulted in the recurrence of high and low flow regimes and a pronounced temperature gradient in these freshwater habitats (Bandyopadhyay et al. [2014](#page-12-32); Chakraborty and Datta [2013](#page-12-12); Goswami et al. [2012a,](#page-13-8) [b](#page-13-9); Guha et al. [2007](#page-13-33); Panja et al. [2020](#page-14-16)). The spatial structure in the mid altitudinal freshwater habitats of River Teesta and Chel might be associated with the stream frequency, braiding, and drainage density (Akhter et al. [2019](#page-11-0); Dhali et al. [2020](#page-12-33)) which has previously exhibited a positive spatial pattern autocorrelation (Akhter et al. [2019\)](#page-11-0). Goswami et al. ([2019\)](#page-13-12) explained that two extrinsic factors, i.e., tectonism and climate coupled with extreme anthropogenic events (Dhali et al. [2020](#page-12-33)), have been causing land cover changes and interfuve characteristics of these rivers, a signifcant modulator of the observed spatial structures of our results. Such variables are presumed to create the spatial association and segregation among these freshwater reaches by modulating fow regimes, pH, dissolved oxygen, turbidity, and the ecological integrity of the river channel and aquatic habitats (Akhter et al. [2019](#page-11-0); Comiti et al. [2011](#page-12-34); Dhali et al. [2020\)](#page-12-33), leading to more spatially structured piscine assemblages. Such fnding is further strengthened by the higher infuence of spatial variables than environmental factors in variation partitioning (Fig. [3](#page-7-0)) (de Campos et al. [2019](#page-12-35); Erős et al. [2012](#page-13-34); Leonidas et al. [2020](#page-14-35)). However, the unexplained remnant variation is comparatively large and could be often described for several local factors, biotic interactions, and their lack of fdelity to include in the study (López‐Delgado et al. [2019;](#page-14-25) Perez Rocha et al. [2018\)](#page-14-8).

The degree of dispersal limitation (Cottenie [2005](#page-12-1); Villéger et al. [2013\)](#page-14-36) is arduous to fathom for each fish species; instead, the focus has been given to its additive partitions of beta dissimilarities, i.e., LCBD and SCBD (Heino [2011](#page-13-35); Leibold et al. [2004](#page-13-36)). Previously LCBD indices were accounted to assess the uniqueness of the aquatic habitats of a fsh community (Legendre and De Cáceres [2013\)](#page-13-3). The higher LCBD values indicate unusual species composition and species-poor sites requiring ecological restoration (Legendre and De Cáceres [2013;](#page-13-3) Panja et al. [2021a;](#page-14-27) Vilmi et al. [2017](#page-15-0)), which is refected for the freshwater habitats of the transitional zone between the Eastern Himalayan mountains and the upper Gangetic plains. Such a pattern is aligned with the spatial association of the sites. This alignment is also supported by the positive correlation between signifcant spatial Axes 1 and LCBD values. Therefore, the LCBD analysis captures the broad-scale spatial association of these freshwater habitats and infers factual information about these freshwater habitats. The LCBD attribute is explained by precipitation of driest month, annual mean temperature, and stream order. The facts support these fndings that most Eastern Himalayan foothill rivers become feeble during the post-monsoon season due to low discharge (Ayaz et al. [2018](#page-12-36); Dhali et al. [2020;](#page-12-33) Rudra [2018](#page-14-15)). These torrential freshwater reaches harbor strong temperature gradients as they are replenished by snow-melt waters and originated under dense canopies at higher elevations (Akhter et al. [2019](#page-11-0); Barman and Das [2014](#page-12-11); Bhatt et al. [2016,](#page-12-4) [2012;](#page-12-13) Das [2020;](#page-12-31) Panja et al. [2020;](#page-14-16) Rudra [2018](#page-14-15)). Following the requirements colder, higher flow modulated oxygen-rich habitats of some characteristically adapted fsh species; the assemblages might become spatially segregated (Jackson et al. [2001](#page-13-2)), leading to unique assemblage structures. On the other note, higher order streams are prevalent in the torrential upland rivers, viz. Chel, Neora, Mal, and Murti, while lower order streams are predominant in large-scale rivers, such as River Jaldhaka and Teesta (Akhter et al. [2019;](#page-11-0) Dhali et al. [2020](#page-12-33); Goswami et al. [2019](#page-13-12)). The lower stream orders are usually negatively associated with species richness (Beecher et al. [1988;](#page-12-37) Platts [1979](#page-14-37)). Therefore, stream order has a signifcant relationship with LCBD, negatively correlated species richness (Legendre and De Cáceres [2013](#page-13-3)). However, these variables are similar to the variables explaining spatial association except for substrate and land cover attributes. Such diference is

presumed to be raised due to the more direct role of substrate and land cover in shaping these freshwater habitats (Akhter et al. [2019;](#page-11-0) Ali et al. [2010](#page-11-1); Biswas and Paul [2020](#page-12-38); Dhali et al. [2020](#page-12-33); Goswami et al. [2019\)](#page-13-12) in comparison to the diferential distribution of fsh species (Chakrabarty and Homechaudhuri [2015;](#page-12-39) Panja et al[.2021b,](#page-14-26) [2020\)](#page-14-16).

It is apparent that these habitat attributes have facilitated the enriched assemblage of rare and characteristically adapted fsh species from genus *Neolissocheilus, Garra, Tor, Devario, Barilius, Psilorhynchus, Crossochelius,* and *Glyptothorax* of the EH (Barman and Das [2014](#page-12-11); Chakrabarty and Homechaudhuri [2015;](#page-12-39) Goswami et al. [2012a](#page-13-8); Panja et al. [2021a](#page-14-27), [b\)](#page-14-26). The SCBD analysis emphasized these genera, which have driven the higher contribution towards the beta dissimilarity (Fig. [5](#page-8-0)), resulting in higher LCBD of their habitats (Heino and Grönroos [2017;](#page-13-29) Legendre and De Cáceres [2013\)](#page-13-3). SCBD values also associate with general species characteristics (for example, species niche & degree of occupancy) and adaptive traits of a species (Heino and Grönroos [2017](#page-13-29); Legendre and De Cáceres [2013\)](#page-13-3). Considering the top contributors in SCBD analysis, *Neolissocheilus hexastichus* and *N. hexagonolepis* are migratory Mahseer fish species that prefer riffles and pools, characterized by the higher water current and substrate coarseness (Arunachalam [2010;](#page-12-40) Froese and Pauly [2011](#page-13-37); IUCN [2020\)](#page-13-38). They are also experiencing threats of extinction, being categorized as near threatened in IUCN red list (IUCN [2020\)](#page-13-38). Although least concerned (IUCN [2020\)](#page-13-38), *Garra lamta* and *G. annandalei* dwell in swift and clear torrential hill streams and exhibit a high degree of adaptation against rocky substratum (Froese and Pauly [2011;](#page-13-37) Nagar et al. [2012\)](#page-14-38). Unlike the other Mahseer species, *Tor tor* inhabits rapid streams with rocky substrate exhibiting upstream spawning migration into more oxygen-rich cascades, riffles, deep pools, and reservoirs (Froese and Pauly [2011](#page-13-37); Menon [1999\)](#page-14-17). *Amblyceps mangois*, *Devario aequipinnatus*, *Barilius vagra,* and *Crossocheilus latius* also inhabit hill streams and prefer mid-hill clear waters with coarser bedrock pebbles, gravel, and stones (Froese and Pauly [2011;](#page-13-37) Menon [1999](#page-14-17); Talwar and Jhingran [1991](#page-14-19)). In contrast, *Badis badis* dwells in tropical freshwaters with a moderate temperature and lower pH. Therefore, the choice of freshwater habitats characterized by a similar range of variables in these fsh species might have led to spatial aggregation and unique or rare species composition in the upper reaches of River Chel, Neora, Murti, Jaldhaka, and upper-west stretches of River Teesta. Such inference is refected in both the spatial and LCBD models, which are fairly explained by characteristic climate, topography, substrate, and land cover attributes. However, the predictive model with LCBD is overftted, indicating reduced fdelity and the need for higher resolution studies in his feld (Elith et al. [2008;](#page-13-27) Nieto and Mélin [2017\)](#page-14-28). Previous studies (Milardi et al. [2018](#page-14-39), [2019](#page-14-40)) accorded a higher native species richness in upland sites with significant SCBD than exotic fish species. Besides, exotic fshes usually override the critical environmental drivers (relevant to native) as they uniquely rely upon geography and human-mediated dispersal limitations (Gavioli et al. [2019;](#page-13-39) Leprieur et al. [2009\)](#page-14-1). However, evidence regarding the impact of exotic fshes on functional diversity, predation, and trophic overlap with the native fsh lack from this region; therefore, the present inferences are solely based on native fsh species and presumed to be less moderated by exotics considering the spatial scale of the study (Davies et al. [2005](#page-12-0); Gavioli et al. [2019](#page-13-39); Milardi et al. [2019](#page-14-40)).

The torrential freshwater reaches of these vast TED ecoregions are experiencing a frequent change in river courses with increasing habitation and altered land use pattern (Chakraborty and Datta [2013;](#page-12-12) Dhali et al. [2020](#page-12-33); Naha et al. [2019](#page-14-13)). A severe trend of deforestation due to natural and anthropogenic hazards regarding replacement, settlements, mining, pebble displacements, and cultivation is leading to increased siltation and vulnerability of these river beds (Akhter et al. [2019](#page-11-0); Chakraborty and Datta [2013](#page-12-12); Dhali et al. [2020](#page-12-33); Goswami et al. [2012a](#page-13-8), [b;](#page-13-9) Naha et al. [2019](#page-14-13); Panja et al. [2020](#page-14-16); Rudra [2018](#page-14-15)). Climate change would aggravate such perils more disastrously, leading to severe degradation of this spatially structured freshwater habitat with unique or rare piscine assemblage (Barman and Das [2014](#page-12-11); Goswami et al. [2012a,](#page-13-8) [b](#page-13-9); Panja et al. [2021b\)](#page-14-26). Due to their spatial association and dispersal limitation, a wide range of fsh species will be experiencing immense threats, while their freshwater habitat will be in jeopardy in the future (Barman and Das [2014;](#page-12-11) Bhatt et al. [2016](#page-12-4), [2012;](#page-12-13) Bhattacharya [2019](#page-12-14); Chakraborty and Datta [2013;](#page-12-12) Goswami et al. [2012a](#page-13-8); Naha et al. [2019](#page-14-13); Panja et al. [2020](#page-14-16); Rudra [2018\)](#page-14-15).

Conclusion

This study reveals the underlying control of freshwater habitats resulting in freshwater fsh species sorting as a frst of its kind information in the TED ecoregion of the EH. The decomposition of spatial models and local contribution has led to identifying the spatial range of freshwater habitats with unique assemblage and higher eco-restoration values. Such habitats must be prioritized for monitoring and conservation assessments following constant alteration in tectonic, climate, and anthropogenic events. The inferences also raise concern for managing and restoring these characteristic freshwater habitats that share signifcant association and more nested assemblage of unique and rare fsh species. Therefore, using spatial decomposition and additive beta partitions would be benefcial to demarcate the habitats and identify the characteristically adapted species for conservation and prioritization. A future application of such analytics through a multi-taxa approach across the expansive landscape would reveal important information on the freshwater habitats of ecologically sensitive ecoregion of EH.

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Authors' contributions SP has formulated, generated, identifed the samples, analyzed the data, and prepared the manuscript primarily. AP has generated the data, participated in the analysis, and drafting the manuscript secondarily. MC has generated data and identifed the samples. SH has supervised the study and edited the manuscript before fnal submission.

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Data availability The raw data is not being submitted presently at this moment. It cannot be reproduced before the publication of the manuscript. However, it may be shared in the review/revision stage for better analytical clarity during review/ revision. All the necessary data for peer review has been submitted with the manuscript and supporting information.

Declarations

Conflict of interest The authors are declaring here no confict of interest.

Ethical approval This study has been conducted by following the ethical guidelines endorsed by the University of Calcutta, University Grant Commission, and Govt. of India. No vertebrate animals have been sampled, which are already forbidden to be captured from the wild. No surveys and sampling procedures were extended to the protected areas and the water bodies within. The authors are now declaring the fulfllment of all ethical commitments subjected to this research work.

Informed consent All the authors have given their full consent for the publication of this manuscript in this journal.

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