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How the Freshwater Biodiversity Information System (FBIS) is supporting national freshwater fish conservation decisions in South Africa

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In South Africa, anthropogenic pressures such as water over-abstraction, invasive species impacts, land-use change, pollution, and climate change have caused widespread deterioration of the health of river ecosystems. This comes at great cost to both people and biodiversity, with freshwater fishes ranked as the country's most threatened species group. Effective conservation and management of South Africa's freshwater ecosystems requires access to reliable and comprehensive biodiversity data. Despite the existence of a wealth of freshwater biodiversity data, access to these data has been limited. The Freshwater Biodiversity Information System (FBIS) was built to address this knowledge gap by developing an intuitive, accessible and reliable platform for freshwater biodiversity data in South Africa. The FBIS hosts high quality, high accuracy biodiversity data that are freely available to a wide range of stakeholders, including researchers, conservation practitioners and policymakers. We describe how the system is being used to provide freshwater fish data to a national conservation decision-support tool-The Department of Forestry, Fisheries, and the Environment (DFFE) National Environmental Screening Tool (NEST). The NEST uses empirical and modelled biodiversity data to guide Environmental Impact Assessment Practitioners in conducting environmental assessments of proposed developments. Occurrence records for 34 threatened freshwater fishes occurring in South Africa were extracted from the FBIS and verified by taxon specialists, resulting in 6 660 records being used to generate modelled and empirical national

distribution (or sensitivity) layers. This represents the first inclusion of freshwater biodiversity data in the NEST, and future iterations of the tool will incorporate additional freshwater taxa. This case study demonstrates how the FBIS fills a pivotal role in the data-to-decision pipeline through supporting data-driven conservation and management decisions at a national level.

KEYWORDS

freshwater fish, South Africa, FBIS, screening tool, conservation, SDM (species distribution model), threatened species, decision-making

Introduction

Freshwater ecosystems are among the most biodiverse habitats on earth, covering less than 1% of the planet's total surface area, but accounting for nearly 25% of all vertebrates and more than 50% of all fishes (Hughes et al., 2021; Fricke et al., 2022). However, despite providing essential ecosystem services, protection of freshwater habitats and their associated biodiversity remain a low priority for policymakers when developing protected areas and legislation (Hughes et al., 2021). Recent global studies (Abell et al., 2007; Adams et al., 2015; Hughes et al., 2021; Williams-Subiza and Epele, 2021) have highlighted significant gaps in protected area networks for freshwater systems, and freshwater biodiversity is declining twice as fast as in marine and terrestrial ecosystems (Grooten and Almond, 2018), with nearly a third of freshwater fishes now threatened with extinction (WWF, 2020; IUCN Red List, 2020; Hughes et al., 2021).

In South Africa, freshwater fishes are recognised as the country's most threatened species group (Skowno et al., 2019). Of the 106 formally described native fish species, 27 are threatened with extinction (classified as Vulnerable, Endangered or Critically Endangered according to the IUCN), with at least eight reported to be in decline over the last decade (Chakona et al., 2022). Ongoing taxonomic revisions indicate that several endemic taxa are genetically distinct and have narrower distribution ranges than previously thought (Chakona et al., 2022). As such, some threatened endemic 'species suites' are likely to be split taxonomically, resulting in individual species being more vulnerable to extinction and raising the total number of threatened taxa in the country (Chakona et al., 2022). The key anthropogenic pressures impacting freshwater fishes in South Africa are land-use change (O'Brien et al., 2019; Chakona et al., 2020a), pollution (Wepener et al., 2011; Horak et al., 2021), excessive abstraction of water (Dallas and Rivers-Moore, 2014; Cerrilla et al., 2022; Evans et al., 2022), spread of invasive species (Ellender and Weyl, 2014; Weyl et al., 2020; Zengeya et al., 2020; Cerrilla et al., 2022) and climate change (Dallas and Rivers-Moore, 2014; Ziervogel et al., 2014). Consequently, there is an urgent need to develop current, data-driven conservation plans and policies to guide effective management and protection of freshwater habitats in South Africa, to safeguard their unique biodiversity and to sustain their essential ecosystem services.

South Africa's protected area network (Republic of South Africa, 2004a; Republic of South Africa, 2004b; Chakona et al., 2022), covers less than 10% of South Africa's total land area (Skelton et al., 1995; Russell, 2011; Skowno et al., 2019). Whilst protected areas generally provide some level of protection for freshwater fishes, few protect entire catchments (Skelton et al., 1995; Acreman et al., 2019; Jordaan

et al., 2020), which is problematic given the linear nature of river ecosystems (Jordaan et al., 2020). Of the country's rivers, only 18% are regarded as Well Protected and 12% as Moderately Protected, with the remainder of the being classified as Not or Poorly Protected (Department of Environmental Affairs, 2016). Even rivers considered as Well or Moderately Protected are often not in a pristine or healthy condition. For example, Kleynhans (2000) and Nel et al. (2009) found that almost half of the river systems falling within protected areas in South Africa are already degraded upstream as a result of human activities (Kleynhans, 2000; Nel et al., 2009). Russell (2011) found that of the 19 national parks managed by South African National Parks, only 13 included habitat for freshwater fishes, with protection often being an unintended byproduct of the targeted protection of threatened terrestrial plants and mammals. Moreover, protection of freshwater fishes within protected areas is compromised by human-linked impacts such as climate change, invasive species and habitat degradation impacts further upstream (Impson et al., 2002; Abell et al., 2007), with 84% freshwater fish regarded as under-protected (Jordaan et al., 2020). As such, South Africa's current network of protected area network does not adequately protect freshwater fishes (Nel et al., 2004; Abell et al., 2007; Jordaan et al., 2020).

Having recognised the limitations of the current protected areas network, expansion of these areas in South Africa is supported through the country's commitment to the Convention on Biological Diversity (CBD) in The National Protected Area Expansion Strategy (NPAES, 2016; Republic of South Africa, 2010; Department of Environmental Affairs, 2016). In conjunction with the NPAES, the Government of South Africa also published amendments (24 (5) (a) and 24 (5) (h)) to the National Environmental Management Act, 1998 (Act 107 of 1998; Republic of South Africa, 1998), stipulating that future developments will need to be guided by an objective Environmental Impact Assessments (EIA) process that takes into consideration presence or absence of threatened taxa (South African National Biodiversity Institute (SANBI), 2021). As such, the Department of Forestry, Fisheries, and the Environment (DFFE) developed a National Environmental Screening Tool (screening.environment.gov.za; SANBI, 2021)-hereafter, the 'NEST'. The NEST is a national web-enabled application that allows applicants seeking environmental authorisation for development to screen their proposed development site for any environmental sensitivities, for example, the presence of threatened species (DFFE, 2021). The NEST uses empirical and modelled biodiversity data (packaged as 'sensitivity layers") to guide EIAs of proposed developments-a process that has in the past been criticized for not being sufficiently transparent or robust (i.e., datadriven). A brief description of the NEST and an explanation of the species distribution layers are outlined below.

The NEST consists of theme-specific spatial datasets, which have been assigned various sensitivity levels (SANBI, 2021), allowing for pre-screening of the proposed development footprint. The NEST assesses the likelihood that a proposed development will have a negative impact on either the environment or any threatened species that may occur in the same area. One of the main components of the NEST is the Plant and Animal Sensitivity Layers, which uses a fourtiered sensitivity rating system to identify and classify habitat where threatened species occur (SANBI, 2021). These layers are briefly described below (SANBI, 2021).

- Very high layer-Habitat for highly range-restricted threatened taxa that has an extent of occurrences of less than or equal to 10 km². For each taxon, critical habitat is manually mapped at a fine scale by taxon experts. Combined data for all taxa are combined into a single spatial layer: Very high sensitivity layer.
- **High layer**-The current distribution of threatened taxa are included in the high sensitivity level by developing spatial polygons around known recent occurrence records (defined by the NEST as records collected since the year 2002 with reasonably high spatial accuracy). Combined data for all taxa are combined into a single spatial layer: High sensitivity layer.
- Medium layer–Species distribution models (SDMs), which use species occurrence records combined with multiple environmental variables to quantify and predict areas of suitable habitat, were used to model suitable habitat areas where threatened species are expected to occur. Combined data for all taxa are combined into a single spatial layer: Medium sensitivity layer.
- Low sensitivity layer-Areas where no threatened taxa are currently known or expected to occur.

Whilst the NEST and associated Species Assessment Guidelines (SANBI, 2021) do currently include aquatic habitat sensitivity layers, they do not currently include any freshwater species-specific layers (SANBI, 2021). Developing sensitivity layers for freshwater taxa requires access to reliable and comprehensive freshwater biodiversity data. However, despite a wealth of current and historic biodiversity data existing for South Africa's freshwater ecosystems, access to these data has been limited by the lack of a dedicated and resourced freshwater information system (Dallas et al., 2022).

Several databases have been developed in South Africa in the past for collating and preserving freshwater biodiversity data at both national and provincial levels (Dallas et al., 2022). However, these databases are generally difficult to access, use different data formats and standards, and are not always maintained due to limited resources and funds (Dallas et al., 2022)—a problem also experienced in both terrestrial and marine ecosystems. Some South African organisations (such as the South African Institute for Aquatic Biodiversity and the Albany Museum) publish their freshwater biodiversity data to the Global Biodiversity Information Facility (GBIF), but the data currently available on GBIF under-represent what is actually available for South African freshwater fish, and require substantial time and effort to clean, format and analyse. Whilst these platforms all contribute valuable data, none of them adequately meet the needs for national freshwater conservation and decision-making in a South Africa (Dallas et al., 2022). However, such information is critical for informing national and international biodiversity assessments, measuring the impact of anthropogenic activities (such as climate change), and enhancing our ability to make informed policy, management, and conservation decisions at both provincial and national levels (Dallas et al., 2022). As such, access to comprehensive and reliable freshwater biodiversity data is imperative and will help to safeguard critical freshwater biodiversity from anthropogenic threats, and will allow freshwater resources in South Africa to be sustainably used and managed.

The Freshwater Biodiversity Information System (FBIS; freshwaterbiodiversity. org), an open-access, online platform for freshwater biodiversity data in South Africa launched in 2017, was developed by the Freshwater Research Centre (FRC) in partnership with SANBI and Kartoza to bridge this data gap by improving access to comprehensive and reliable freshwater biodiversity data. The FBIS is a powerful, open-access system for hosting, standardising, analysing and serving freshwater biodiversity data for South Africa. As such, the FBIS functions as a repository for South African freshwater data, and has been populated with data from key available sources including data mobilised through manual literature searches, and data pulled in via links with existing online platforms such as GBIF (Dallas et al., 2022). The FBIS represents the first comprehensive, accessible, national-level resource for freshwater biodiversity data records in South Africa, and thus provides an opportunity for national-level freshwater biodiversity data to be utilised by researchers, policymakers, and conservation practitioners in real time (Dallas et al., 2022).

Here we describe how the FBIS was recently used to provide freshwater fish data for informing a national-level conservation decision-support tool (the NEST), and we evaluate the role of the FBIS in the data-to-decision pipeline. Specifically, we examine how data extracted from the FBIS were used to develop national freshwater fish sensitivity layers-core components of the NEST-thereby allowing for the first inclusion of freshwater species spatial coverage in the tool, and the potential for improved freshwater biodiversity conservation at a national scale. This case-study demonstrates the importance of collecting and collating comprehensive, high quality biodiversity data sets, and being able to synthesize these data and make them accessible for analysis and uptake and use in conservation planning and decision making at a national scale. We also highlight the importance of expert consultation and multi-disciplinary stakeholder engagement during various stages of this process.

Methods

Data source

The development of the NEST sensitivity layers for freshwater fish was a collaborative effort that included

TABLE 1 List of threatened (Vulnerable, Endangered and Critically Endangered) South African freshwater fish taxa included in the NEST analysis. Conservation status and the number of occurrence records sourced from the Freshwater Biodiversity Information System (FBIS) pre and post 2002 are shown. Taxa #1–27 are formally-describes species, while taxa #28–34 represent recognised genetically-distinct lineages. Threat categories are: VU = Vulnerable; EN = Endangered; CR = Critically Endangered.

#	Scientific name	Common name	IUCN Conservation Status	SANBI Conservation Status	Number of Records Pre 2002	Number of Records Post 2002
1	Austroglanis barnardi	Barnard's rock catfish	EN	EN	66	34
2	Chetia brevis	Orange-fringed largemouth	EN	EN	17	27
3	Chiloglanis bifurcus	Incomati suckermouth	CR	CR	81	45
4	Chiloglanis emarginatus	Pongolo suckermouth	VU	VU	26	27
5	Ctenopoma multispine	Many-spined climbing perch	LC	VU	21	3
6	Enteromius gurneyi	Redtail barb	VU	VU	73	14
7	Enteromius treurensis	Treur River barb	CR	CR	20	10
8	Labeo rubromaculatus	Tugela labeo	VU	VU	29	16
9	Labeo seeberi	Clanwilliam sandfish	EN	EN	42	230
10	Marcusenius caudisquamatus	Natal bulldog	EN	EN	1	8
11	Oreochromis mossambicus	Mozambique tilapia	VU	VU	1,581	1,292
12	Pseudobarbus afer	Eastern Cape redfin	EN	EN	93	95
13	Pseudobarbus asper	Smallscale redfin	VU	VU	294	54
14	Pseudobarbus burchelli	Barrydale redfin	CR	CR	18	32
15	Pseudobarbus burgi	Berg River redfin	EN	EN	84	109
16	Pseudobarbus capensis	Berg-Breede River whitefish	EN	EN	45	24
17	Pseudobarbus erubescens	Twee River redfin	CR	CR	122	20
18	Pseudobarbus phlegethon	Fiery redfin	EN	EN	118	72
19	Pseudobarbus quathlambae	Maloti minnow	EN	EN	94	59
20	Pseudobarbus senticeps	Krom River redfin	CR	CR	31	4
21	Pseudobarbus skeltoni	Giant redfin	EN	EN	2	36
22	Pseudobarbus swartzi	Gamtoos redfin	VU	VU	130	35
23	Pseudobarbus trevelyani	Border barb	EN	EN	23	10
24	Pseudobarbus verloreni	Verlorenvlei redfin	EN	EN	29	24
25	Sandelia bainsii	Eastern Cape rocky	EN	EN	83	71
26	Serranochromis meridianus	Lowveld largemouth	EN	EN	75	10
27	Silhouettea sibayi	Sibayi Goby	EN	EN	10	2
28	<i>Kneria</i> sp. nov. south africa		EN	EN	29	19
29	<i>Marcusenius</i> sp. nov. kosi		EN	EN	0	4

(Continued on following page)

TABLE 1 (*Continued*) List of threatened (Vulnerable, Endangered and Critically Endangered) South African freshwater fish taxa included in the NEST analysis. Conservation status and the number of occurrence records sourced from the Freshwater Biodiversity Information System (FBIS) pre and post 2002 are shown. Taxa #1–27 are formally-describes species, while taxa #28–34 represent recognised genetically-distinct lineages. Threat categories are: VU = Vulnerable; EN = Endangered; CR = Critically Endangered.

#	Scientific name	Common name	IUCN Conservation Status	SANBI Conservation Status	Number of Records Pre 2002	Number of Records Post 2002
30	<i>Pseudobarbus</i> sp. nov. breede		N/A	VU	216	324
31	<i>Pseudobarbus</i> sp. nov. doring		CR	CR	26	19
32	<i>Pseudobarbus</i> sp. nov. heuningnes		EN	EN	7	48
33	<i>Pseudobarbus</i> sp. nov. keiskamma		EN	EN	38	9
34	<i>Pseudobarbus</i> sp. nov. keurbooms		N/A	EN	19	16

contributions from several organisations: FRC, SANBI, NRF-South African Institute for Aquatic Biodiversity (NRF-SAIAB), CapeNature, Endangered Wildlife Trust (EWT), DFFE, Mpumalanga Tourism and Parks Agency (MTPA), Free State Department: Economic, Small Business Development, Tourism and Environmental Affairs, and University of Cape Town (UCT), as well as multiple other individuals and organisations via SANBI's National Freshwater Fish Observation Group (see Supplementary S3 and S4). Occurrence records for 34 threatened (Vulnerable, Endangered and Critically Endangered classified as per the IUCN) freshwater fishes occurring in South Africa (Table 1) were downloaded on 14 September 2022 from the FBIS online database (FBIS, 2022). These 34 fish taxa included 27 formally described species and seven lineages, yet to be formally described (Chakona et al., 2022). Formally described species have been assessed both globally (on the IUCN Red List) and nationally (Red List of South African Species; SANBI, 2022), whereas undescribed lineages have only been assessed nationally (SANBI, 2022). Only primary, secondary, and catadromous threatened freshwater fishes were included, while marine peripheral/sporadic and primary marine fishes were excluded.

Data cleaning and validation

A series of online workshops were held with freshwater fish taxon specialists from across South Africa during 2021 to verify the quality and comprehensiveness of the data downloaded from the FBIS (see Supplementary Table S1). Relevant experts were identified for each threatened fish taxon, based on prior involvement in species Red-Listing assessments, and consulted. Occurrence records for each fish taxon from the FBIS were thoroughly scrutinized, erroneous records were removed, and missing data sets identified and added. Resultant occurrence data sets were used to generate updated distribution maps for each taxon, and these were sent to all specialists for approval prior to further analysis.

Data analysis

All data cleaning, processing and analyses were conducted using R Software (R Core Team, 2020; Version 3.5.0). Data visualizations and final maps were produced using R Software (R Core Team, 2020; Version 3.5.0) and ArcGIS Pro (Esri Inc, 2022). Cleaned and validated occurrence records were used to produce different sensitivity layers, for inclusion into the NEST. The protocols for developing the different sensitivity layers followed those described by DFFE (2021). A brief description of how each sensitivity layer was developed is outlined below. Final data and spatial layers were presented to taxon specialists for review before being combined and submitted for inclusion in the NEST.

Very high sensitivity layers

The 'very high' sensitivity category only applied to freshwater fishes that were assessed as Critically Endangered. Given that freshwater fish occur in linear river systems, the criteria used to develop the Very high sensitivity layer were adapted as follows: *All historic (pre-2002) and current (post-2002) occurrence records of freshwater fish taxa that are categorised as Critically Endangered*. As such, all known, valid, historic occurrence records were used to build the very high sensitivity layer. Occurrences were intersected with the FEPA Sub-Quaternary Catchment layer (Nel, 2011; SANBI, 2011) to create catchment-specific occurrence polygons, indicating the presence of a Critically Endangered freshwater fish in that catchment.

High sensitivity layers

The high sensitivity category only applied to freshwater fishes that were assessed as Vulnerable or Endangered. The 'high' sensitivity layer is comprised of all valid, post-2002 occurrence records (SANBI, 2021). Occurrence records for each freshwater fish (i.e., the assumed, current distribution of the species) were plotted in R (R Core Team, 2020; Version 3.5.0). Occurrence data



were then intersected with the River Freshwater Ecosystem Priority Areas (FEPA) Sub-Quaternary Catchment layer (Nel, 2011; SANBI, 2011) to create catchment-specific occurrence polygons, indicating the recent presence of a Vulnerable or Endangered freshwater fish in that catchment.

Medium sensitivity layers

Taxa that qualify for inclusion in the 'medium' sensitivity layer were categorised as either Vulnerable or Endangered. For these taxa, species distribution models (SDMs) were used to generate predictive geographic ranges. All valid occurrence records for each freshwater fish were used to independently develop a unique, accurate SDM for each taxon using a Bayesian additive regression trees (BART) algorithm *via* functions from the *embarcadero* R package (Carlson, 2020). A comprehensive suite of environmental and hydrological variables was used to generate these SDMs (see Supplementary Table S2). Modelled distributions were converted to a binary output (presence/absence) for each species using a threshold that maximizes the true skill statistic from the species SDM.

Results

Initially, 6 068 records were downloaded from the FBIS (Figure 1; FBIS, 2022). Data cleaning, guided by expert consultation, resulted in the deletion of 354 erroneous records

(Figure 1). An additional 946 records provided by the experts originating from private, unpublished datasets owned by relevant experts (Figure 1) were uploaded to the FBIS. The final cleaned dataset was also tagged in FBIS as "DFFE Screening Tool 2022", which will ensure that expert input and feedback will not be lost and will help streamline this process in the future.

In total, 6 660 occurrence records for the 34 threatened (Vulnerable = 8; Endangered = 20; Critically Endangered = 6) freshwater fishes were used to develop the NEST layers (Table 1; Figure 1). Of the 34 threatened freshwater fishes included in the NEST, 14 had fewer than 50 records in South Africa (Table 1). Three threatened taxa, *Marcusenius caudisquamatus* (n = 9), *Marcusenius* sp. nov. kosi (n = 4) and *Silhouettea sibayi* (n = 12) were found to have less than 20 records. Only two taxa, *Oreochromis mossambicus* (n = 2 873) and *Pseudobarbus* sp. nov. breede (n = 540), had more than 500 records (Table 1).

Modelled taxon distribution maps (contributing to the medium sensitivity layer) and current taxon distribution maps (high sensitivity layer) were produced for all taxa classified as Vulnerable and Endangered, except for *S. sibayi*, which had no records post-2002. For Critically Endangered taxa, occurrence data were used to develop a single distribution map for each of the seven taxa (making up the very high sensitivity layer). Example outputs are presented for two fish species, namely, *Labeo seeberi* (Endangered; Figure 2) and *Pseudobarbus erubescens* (Critically Endangered; Figure 2), illustrating finalised occurrence data (Figures 2A, E), SDMs derived from these data (Figures 2B, F), and how these were used to develop the medium (Figure 2C), high (Figure 2D) and very high (Figure 2G) sensitivity layers for the DFFE Screening Tool.



FIGURE 2

Labeo seeberi occurrence records (A), species distribution model (B), medium sensitivity area (C) and high sensitivity area (D) located in the Olifants-Doring primary catchment area of South Africa. *Pseudobarbus erubescens* occurrence records (E), species distribution model (F), and very high sensitivity area (G) located in tertiary catchment E21.



The final composite (combined across all taxa) national sensitivity layers (medium, high, and very high) for all threatened freshwater fishes are presented in Figure 3. The medium sensitivity layer for freshwater fish spans 49 385 km of river and covers a total catchment area of 251 264 km², whilst the high sensitivity layer spans 15 162 km of river and covers a catchment area of 117 412 km², with the very high sensitivity layer providing much-needed protection for Critically Endangered freshwater taxa spanning 1,024 km of river and covering a total catchment area of 5 992 km².

Discussion

We present the first inclusion of freshwater species coverage in a key national conservation decision-support tool—the DFFE EIA NEST—and describe how data from the FBIS were used as the basis for threatened fish sensitivity layers (core elements of the tool) through a multi-stakeholder, collaborative approach (see Supplementary S3 and S4). Sensitivity layers were successfully developed for 34 threatened freshwater fishes occurring in South Africa and combined to produce national-level sensitivity layers for inclusion into the tool. Given that the majority of South Africa's rivers are either poorly protected or not protected (Department of Environmental Affairs, 2016), the updated coverage of freshwater fishes in the NEST should provide much-needed protection for the country's threatened freshwater fishes by preventing or minimising further destruction of critical freshwater habitats due to new developments.

Importantly, all new applications for development under the EIA regulations will be compelled to make use of the tool before authorisations are granted. This will identify and protect sensitive catchment areas that in the past would have been overlooked due to a lack of access to freshwater biodiversity data. This will no doubt support future conservation efforts, especially in locations where threatened taxa occur outside of formal, protected areas (Impson et al., 2002; Nel et al., 2004; Abell et al., 2007; Russell, 2011; Jordaan et al., 2020). In addition to the conservation and policy benefits, the

distribution maps produced for the medium, high and very high sensitivity layers can also help guide expansion of protected areas and stewardship initiatives and can be used in future national biodiversity assessments and IUCN Red List assessments.

The FBIS supported the development of freshwater fish sensitivity layers through providing quick and easy access to reliable and comprehensive freshwater fish occurrence records for South Africa. The relatively large number of records provided by the FBIS is especially noteworthy given that most distribution modelling and mapping of threatened taxa are heavily hampered by limited access to occurrence records (Stockwell and Peterson, 2002; Wisz et al., 2008). A substantial percentage of these data originating from published literature and private data repositories would not have been available had it not been for a platform like the FBIS. Whilst databases like GBIF and FishBase do provide opportunities for the scientific community to share and access freshwater fish data, there are not any concerted efforts to drive the data collation process (especially within freshwater systems in South Africa). In this regard, the FBIS has a proven advantage over global biodiversity databases, through leveraging personal relationships and thorough ongoing stakeholder engagement to ensure that both the quantity and quality of records uploaded to the FBIS are maximised. Furthermore, the inclusion of various taxon specialists, government organisations and conservation authorities (see Supplementary S1) via multiple online workshops ensured that the final occurrence dataset was cleaned efficiently and included only verified, high-certainty records. This culture of cooperation and collaboration was one of the cornerstones of success in developing the freshwater fish sensitivity layers for the NEST and should be adopted by all future bioinformatics projects that seek to impact conservation management and planning on a national scale.

Understandably, we did encounter some limitations whilst developing the NEST sensitivity layers. Firstly, there were 14 taxa with fewer than 50 records in South Africa, with three taxa having less than 20 records each (Table 1). Data scarcity increased the difficulty of producing accurate SDMs (Stockwell and Peterson, 2002; Wisz et al., 2008) potentially reducing the accuracy of the sensitivity layers for these taxa. This highlights the urgent need to increase threatened species monitoring efforts at both provincial and national levels (Chakona et al., 2022). We recommend, based on their low numbers of recent records, that the following taxa be considered as a very high priority for baseline data collection and monitoring: Chetia brevis, Ctenopoma multispine, Enteromius treurensis, Kneria sp. nov. south africa, Labeo rubromaculatus, Marcusenius caudisquamatus, Marcusenius sp. nov. kosi, Pseudobarbus senticeps, Pseudobarbus skeltoni, Pseudobarbus sp. nov. doring, Pseudobarbus sp. nov. keiskamma, Pseudobarbus sp. nov. keurbooms, Amatolacypris trevelyani, and Silhouettea sibayi. In this regard, a standardised sampling protocol for freshwater fishes is currently under development in South Africa.

Secondly, the NEST sensitivity layers are only restricted to the modelled and current distribution of threatened taxa. Several studies have highlighted the importance of ensuring that catchment-level impacts (Skelton et al., 1995; Kleynhans, 2000; Nel et al., 2004; Abell et al., 2007), specifically upstream-impacts of anthropogenic activities, also be considered when developing protected areas for freshwater systems. The current iteration of the NEST does not fully account for this, providing protection for threatened taxa at the sub-

quaternary catchment level only. Future iterations of the tool should consider ways to provide upstream protection at a larger primary catchment scale. Lastly, the overall success in terms of providing onthe-ground protection to South Africa's threatened freshwater fishes will be dependent on local government authorities correctly and competently interpreting the outputs of the NEST and making responsible land-use change decisions by preventing developments where the risk to threatened taxa is deemed to be too high.

The freshwater fish component of the NEST provides a critical first step towards adequately incorporating threatened freshwater taxa into the EIA and development application process. However, there are additional steps that could improve upon and update the tool, thereby supporting more effective freshwater conservation in the future. Firstly, there is an urgent need to update and resolve the taxonomy of South Africa's freshwater fishes, as there are a number of distinct genetic lineages that await formal description (Chakona et al., 2015, 2020b; Martin and Chakona, 2019; Bronaugh et al., 2020; Kambikambi et al., 2021; Mazungula and Chakona, 2021; Ramoejane et al., 2021). Once the taxonomy of South Africa's freshwater fishes has been revised, the FBIS and NEST can be updated accordingly. Secondly, the NEST process described here can now be replicated for other species groups such as freshwater invertebrates. Although anurans are already included in the NEST (SANBI, 2021), there is also the potential to use the FBIS and the methodology developed here to improve upon these sensitivity layers. Lastly, distribution mapping used for threatened species should be repeated for non-native freshwater species (specifically invasives) occurring in South Africa-the primary threat to native freshwater fishes in the country (Weyl et al., 2020; Zengeya and Wilson, 2020; Chakona et al., 2022). Worryingly, the distributions of many of the country's non-native fish species are not well known, and a lack of spatial data is holding back reliable system-wide invasive species assessments (De Moor, 1996; Ellender and Weyl, 2014; Zengeya and Wilson, 2020). Although mapping non-native species distributions falls beyond the scope of the NEST, this could provide insights into non-native species impacts on native taxa, and assist with identifying key areas for alien species management interventions (Weyl et al., 2020). Actioning these next steps will amplify the impact that the NEST may have on preventing further population declines in South Africa's freshwater fish fauna.

This case study demonstrates how the FBIS has been used successfully to provide spatial data on threatened freshwater fishes to inform a national-level conservation decision-support tool in South Africa. Key to the success of this project was investing substantial time and effort into manually identifying, collating and cataloguing historic biodiversity data into a standardised, digital format. This generated comprehensive, highquality biodiversity data sets that, through the FBIS, were then made accessible for analysis and uptake into national conservation planning and decision-making. Collaboration, networking and stakeholder engagement from the outset encouraged data-sharing and facilitated inter-disciplinary skill-sharing (e.g., modelling, mapping and data management skills)-two critical elements of the projects' success - and we recommend this approach to similar bioinformatics efforts elsewhere, particularly in developing countries.

Data availability statement

The datasets presented in this study can be found in online repositories. The names of the repository/repositories and accession number(s) can be found in the article/Supplementary Material. Alternatively, the dataset can be accessed via the Freshwater Biodiversity Information System (freshwaterbiodiversity.org).

Author contributions

MK is the primary author and was involved in all aspects of publishing this manuscript. JS, CG, and HD contributed to the conception and design of the project as well as assisting with writing the first draft of the manuscript. DH contributed to conception and design of the work as well performing statistical analysis and assistance with writing the first draft of the manuscript. JP, DV, DI, AC, DR, NJ, BP, MJ, RB, FR, TZ, AH, and NR-M all assisted with data cleaning and provided conceptual design and advice throughout the project. All authors contributed to manuscript revision, read, and approved the submitted version.

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Conflict of interest

The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

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Supplementary material

The Supplementary Material for this article can be found online at: https://www.frontiersin.org/articles/10.3389/fenvs.2023.1122223/ full#supplementary-material

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