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Recognizing knowledge gaps in biodiversity for the preservation of endangered flora

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Abstract

Biodiversity, the variety of life on Earth, is of paramount importance for ecological stability and human well-being. Among the most vulnerable components of biodiversity is endangered flora, encompassing a wide range of plant species facing the threat of extinction. Preserving these endangered plant species is critical for maintaining ecosystem balance and ensuring sustainable natural resources for future generations. To achieve effective conservation, identifying and addressing knowledge gaps in biodiversity research becomes imperative. This study presents a comprehensive review of the current state of knowledge concerning endangered flora preservation. It explores the following key areas: Threat Assessment and Monitoring: Understanding the threats faced by endangered plant species is the first step towards their conservation. Research efforts should prioritize the assessment of threats such as habitat loss, climate change impacts, invasive species, and over-exploitation. Regular monitoring is essential to gather data on population dynamics, geographic distribution, and State of the environment influences protection tactics. Taxonomy and Species Identification: Accurate taxonomy and species identification are fundamental for designing targeted conservation measures. In many cases, taxonomic uncertainties hinder conservation efforts, leading to mismanagement and neglect of certain plant species. Advanced genetic techniques and the integration of traditional ecological knowledge can aid in resolving taxonomic challenges. Climate Change Adaptation: Climate change poses significant challenges to endangered flora preservation.

Keywords: Biodiversity, Endangered flora, Geographic distribution, Climate change.

1. Introduction

In the tropics, the existence of life upon planet or human knowledge of presence is dispersed widely. These last several generations has seen a major rise in attention amongst scientists and conservationists in attempts to uncover biodiversity knowledge gaps. Their

awareness has been stoked by the mounting evidence showing that communities are vanishing as a result of habitat destruction, over exploitation of environmental materials, garbage, the spread of alien species, and changed climatic patterns (León–Cortés *et al.*, 2023). Underwater ecosystems, these have most of the greatest densities of species on the world, make up less than 1% of the total area. These ecosystems also support more than 50% of all fish species and over 25% of all vertebrates. But though supplying crucial ecosystem services, politicians continue to place little emphasis on safeguarding freshwater habitats and the biodiversity that goes along with them when creating conserved spaces and laws (Sinthumule, 2023). Utilization of botanical types occurs everywhere due to the strong consumer for pharmacological, visual, and decorative goods worldwide. Developing botanical material extraction has set its sights on Sub-Saharan since it is home to a significant supply of rare and precious plants. Recognized to produce vital oils employed in the fragrance industry, sandalwood is a versatile, resistant to drought tree. Due to the considerable loss in the primary producers of rosewood fuel, it became a potentially profitable wildlife (Lannuzet *et al.*, 2022). Indigenous peoples have utilized the cultural knowledge of their local ecosystem for millennia to not only function but also to maintain the cultural distinctiveness. Although classic understanding was around for countless centuries, the occidental professional establishment has only recently come to recognize it as a useful resource of ecosystem understanding (Isbell, 2023). The absence of information regarding biological richness and transportation, also described as the Nonlinear gaps, is particularly pronounced in extremely diversified places even while data was essential for protection management. Further then greater prediction computer techniques have been created, released, and used on ever-larger biological databases to combat various problems (Kor and Diazgranados, 2023).

Buechley *et al.*, (2019) present World wildlife or well both the biological ecology are during risk from invasive alien plants (IAPs). Since the players and data are numerous and dispersed, it is regarded as a single of the factors that influences wildlife and changes the ecological functions and economical circumstances of the community, demanding cooperation in the administration of IAPs. Since 80% of property was domestically held, Collaborative Adaptive Stewardship (CASt) is essential for IAPs preservation. The goal was to compile and compile data from hundreds of specialists in wildlife from across the globe who jointly research all significant taxed and regions in clean water, terrestrial, and aquatic environments. Buechley *et al.*, (2019) provide we created a poll to find areas of worldwide agreement, address information gaps for overlooked topics and locations, and check for notable variations in estimations and viewpoints among associations of experts. Where applicable, we contrasted surveyed findings to data from other sources. It might be difficult for protected areas (PA) in biodiversity centres to maintain several nationally uncommon and vulnerable botanical varieties on a cost at periods. Stephenson *et al.*, (2022) provide choice of which species to monitor might be made easier for PA managers by prioritizing species based on their regional danger of extinction. On the species' presence and danger of destruction in the PA, there still is frequently barely any documentation provided. As a result, PA managers frequently utilize the regional or international Red List to determine which species are most important at the PA layer. Designated places are required on a worldwide scale to preserve biodiversity and cultural property. Pirie *et al.*, (2022) present Despite being vital to plants, historical collection sites are not recognized or acknowledged under the world property cover, even though these are crucial to the ecology of regulated places. In order to establish the argument that historical examples and related in presence plants communities are bio-cultural legacy funds, we set out to explain the management and historical worth of botanical collection exhibits and to make the connections of human

beings, scientific society, and landscape obvious. Kulak *et al.*, (2022) present Birds provide essential ecosystem roles, were especially vulnerable to death, and are frequently utilized as key type and weather monitors. However, there is no worldwide structure in place to organize rescue efforts and study on them. They create a special studies and preservation purpose score (RCPI) to identify global study and protection goals, identify the factors that threaten elimination for the initial time, and concentrate expert activities on raptors. Shan *et al.*, (2023) provide Many customers, particularly governments, businesses, and societal cultures, have access to the knowledge required should take wise choices about nature, compromising attempts and continue preserve, rebound from, and manage the planet. In this essay, we discuss the need of enhancing wildlife observation, assess the issues at hand, and provide potential solutions. Due to the vast information spaces, it is imperative to expand environmental surveillance, especially in low developing countries with substantial biological diversity. Achieng *et al.*, (2023) provide Ecological DNA studies have proven themselves as a reliable method for identifying ecological populations and are very promising in assisting programs to analyze, conserve, and control wildlife a territory with outstanding ecology, which is endangered by industrial stresses as well as a fundamental shortage of study facilities which restricts environmental analysis and supervisors. Optimizing over protection requires prioritizing ecological efforts, particularly while the ecological situation increases. Mandrak *et al.*, (2019) present sought help cover up knowledge gaps out species. In order to quantify the worldwide distributions of 88 species of Aloe, we produced a collection of herb garden discount coupons and frequency records. It then utilized this collection to determine the main risks to Aloe in this area and predict the danger of endangerment. Moire *et al.*, (2023) present during the past few generations, the territory and number of big carnivorous animals had significantly decreased. Whereas precise community estimates and history tracking are necessary for protection strategy and the supervision of vulnerable organisms, we have proof that wildlife tracking can't be uniformly dispersed or taking place when it is least required. Fleischmann, (2019) provide a typical approach for managing environmental incursions places minimal attention on the community efforts and instead relies heavily on state-run, extensive controlled measures. Never the less, helping should enable identify, eradicate, and confine invasive alien plant species (IAPS). To increase the effectiveness of unwanted alien organism administration, it is crucial to comprehend the level of voluntary engagement and their motives.

2. Materials and methods

Endangered flora

Employing the Registry of Animals to cross-check 123 Wildlife and Preservation fragments ecological endemism, we created one repository of the indigenous plants of contains information on 175 relationships, 1061 groups, 10,965 species, and 762,655 shipping records for bacteria and the team of p athletes.. Central Plants Checklist and the Angiosperm of this method team were used as biological authority to establish individual classification, which was then evaluated against geographically frequency information of national Botany Database. Data of variations with subtypes was combined together form one genus. The samples are georeferenced in to quarter degrees squares (QDS), or around 25 km by 25 km, to approach the geographical accuracy of previous data. The National Vegetation Map project, the Custodians of Endangered Wildflowers initiative, the ecological.

Atlas Product (n = 245,407), the ACOCKS databases (n = 66,068), and lesser programs (n = 138,957) were used as sources for most current and accurate plants appearance information. By placing reallocation information from protologues, developments, gathering latitude-longitude data using Internet the planet, and flowers, some species (n = 1054) that

were left out of these records was personally retrieved.

Genetic data

They used a store in R version 3.5.2 to query Gen Bank, a vast visible records with DNA for roughly 260,000 officially explained organisms, for papers for each of indigenous organisms, coming back an amount of DNA sequences accessible to everyone organisms, in order to evaluate taxonomic partiality in DNA order.

Species Distribution Modeling

The globe Claims collection was used to source 19 raster-based bioclimatic variables with a geographical with a 10 arc-minute accuracy, almost matching the accuracy for their information about organism's presence. As ecological indicators, those bioclimatic variables were subsequently combined with The packages gbm, lmtree, boot, dismo, and randomForest were used in R version 3.5.2 to build species distribution models (SDMs) using generalized straight theories, randomly jungles, and gradient boost devices. Some common modeling techniques forecast the appropriateness of an organism' environment using presence-absence records. Because the database doesn't contain any actual delays, pseudo-absences were created using underlying information tied to national frontiers. The model was tested using 25% of the occurrence documents, and the strategy was trained using 75% of the experience datasets. To avoid models under-fitting, repeated data was eliminated. By merging the multiple models results adjusted by the median of Area under the Curve (AUC), which assigns greater value to the method that provides a superior calculation, composite results are created. AUC values below 0.5 were assigned a score of 0% for forecasts. The limit that optimizes the total of the real positivity rate and actual-negative rate was applied to convert averaged forecasts into quantitative presence-absence images. While they discovered the total accuracy of spatial estimations using SDMs dropped as the number of frequency areas rose, we only fitting SDMs to animals having at least 5 appearances (n = 8295). ArcGIS 10.5, in accordance with Goldsmith et al., was used to determine the creatures range over individuals having three or four frequency locations (n = 691). The QDS in that a wildlife had just a few appearances (n = 1961) was regarded as the organism' region.

Climate suitability

Applying our general libraries in R, raster data from each SDM conclusion was overlaid to identify areas with wide thermal range for a variety of flora. Following the removal of spatial elements for each composite output into the equalized measurement of areavaries from km of 25 to km of 9 25 latticesamong10.6 in ArcGIS, the aggregate results were then compared to the correctness of the fundamental biological characteristics in the repository. Excellent regard tissues convey regions via indicate weather identifying that the behind a significant section of the supply of the showed animals, although the fact that we do not believe animals are living in this environment specifically because other space elements ranging within smaller geographical levels, organisms relations, and previous danger might affect each species' real-life rewards. Since the anticipated richness for coastal grid cells with 50% of land would probably underestimate their actual richness, these cells were not included. We utilized a hierarchical going to calculate the quantity of variety from appearance data and the overall amount of appearance reports into every tissue of the hexagonal matrix used below in order to produce maps of reported biodiversity and population efficiency.

Spatial and taxonomic gaps

They measured the sampling percentage, where we referred as being the proportion of species found from presence reports compared to expected diversity from SDMs, in order to identify potential regional gaps in classification collecting activity. The percentage of frequency data compared to predicted diversity from SDMs or what we call samples density, was computed the measure in collecting effort. Thus, it created the distribution relates in verified population locations. They superimposed a map of major highways on the sample densities map "road effect," which refers to propensity in collecting stronger at areas were widely accessed. In order to pinpoint prospective regions for biological selection, we are additionally assessed the proportion of biological diversity with genetic information compared to forecasted diversity from SDMs. In order to determine if dangerous organisms was most probable to possess genetic knowledge, we categorized the species possessing sequences of information according to IUCN Red List danger level. Finally, we looked at how much the gaps in ecological records around one axis matched the gaps in information around other directions spanning area and groupings.

3. Results**The spatial distribution of biodiversity data gaps**

Mounting estimates using SDMs and collecting existing biodiversity from frequency reports allowed us to map the regional characteristics of actual and predicted indigenous diversity onto 1790 grid cells. We demonstrate that 4% of the nation's indigenous organisms have not being tested, despite the fact that our SDMs reveal that every cell in the grid encloses distance having a temperature which is contained in a seasonal region length of at least 69. SDMs and reported records both have a comparable general layout in terms of relative diversity, although the latter's entire grids ecosystems, the predicted established biodiversity from SDMs is in good agreement with present knowledge. The Tropical Cape Biome, also known as the Coral Cape Ecological Hotspot, is predicted to have a peak indigenous quantity of 4908 species per grid cell. The grid cells with the greatest predicted unique complexity are covered by the communities of Indian Oceans Coastal Area. Flora is the ecological with the greatest mean predicted total wealth. This flora is located in the Western Floristic Country, biological hotspots with a predicted high of 5,303 varieties for grid cell unique complexity.

Distribution of biodiversity data gaps

This repository contains species from 175 families and 1061 genera, with a wide range in the organization of biological data on a taxonomy and chronological level. The top 10 families in terms of the number of native species. Sixty-one percent of the database's indigenous plants belong to these 10 groupings. One isolated individual only represents 40 groups. 61% of the database's indigenous organisms are represented by these 10 groupings. A single native species covers forty different family groups. The Proteaceae, Asteraceae, and Fabaceae families have the most unique species 9 location occurrence records. The top 10 families by sample make up 69% of all the database's records of occurrences. In our study, a single record represents three identical groups. As would be predicted if all species had an equal likelihood of getting tested, In overall, better species-rich groups had being analyzed beyond thoroughly than less species-rich groups. The sample rigor varies quite noticeably between relationships, though. For instance, the Anemiaceae family contains 508 entries despite having just one endemic species; nonetheless, the many groups that are substantially underrepresented may be of additional protection significance.

Diversity of threaded

In terms of the distribution of endangered by family, the IHR's plants known as Fab (84 threatened species) and Cyperaceae (species at 65) and Poaceae (species at 36) families were the most numerous (**Figure 1**). The endangered groups were also divided into five life forms: climber (species at 15), bryophyte (species at 05), pteridophyte (species at 15), and plant (species at 286; **Figure 2**). According to their woodland coverage and spatial location, the vulnerable species are distributed significantly unequally among the states (**Table 1**).

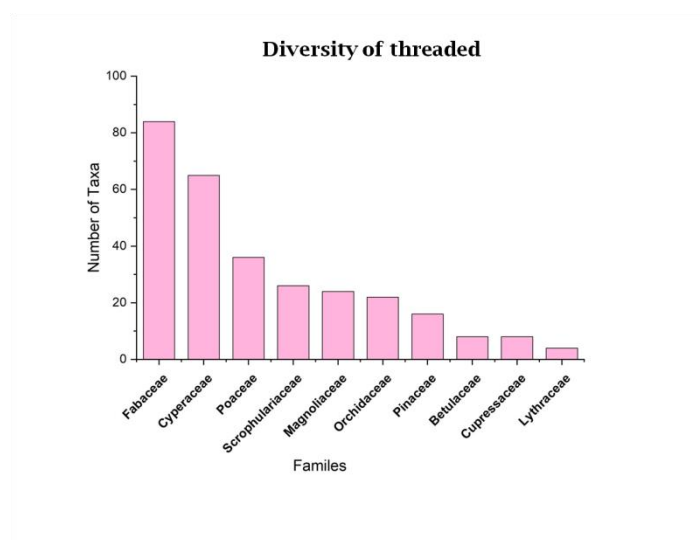


Figure 2: Dominant families of threatened of IHR

Table 1: Outcomes of IHR

Families	Number of Taxa
Fabaceae	84
Cyperaceae	65
Poaceae	36
Scrophulariaceae	26
Magnoliaceae	24
Orchidaceae	22
Pinaceae	16
Betulaceae	8
Cupressaceae	8
Lythraceae	4

The states in Eastern Himalayan (SK, MN, ML, TR, MZ, NL, &AR) and two component countries (AS hills and WB hills) are home to more threatened animals than the states of Western Himalayan (UK, JK, and HP). The material on these Eastern Himalayan republics, however, has received less research and is still mostly unknown in terms of wildlife. SK (203 species), HP (species at 190), and JK (species at 189) represent the IHR sectors with the greatest threatened groups (see **Figure 2 and table 2**).

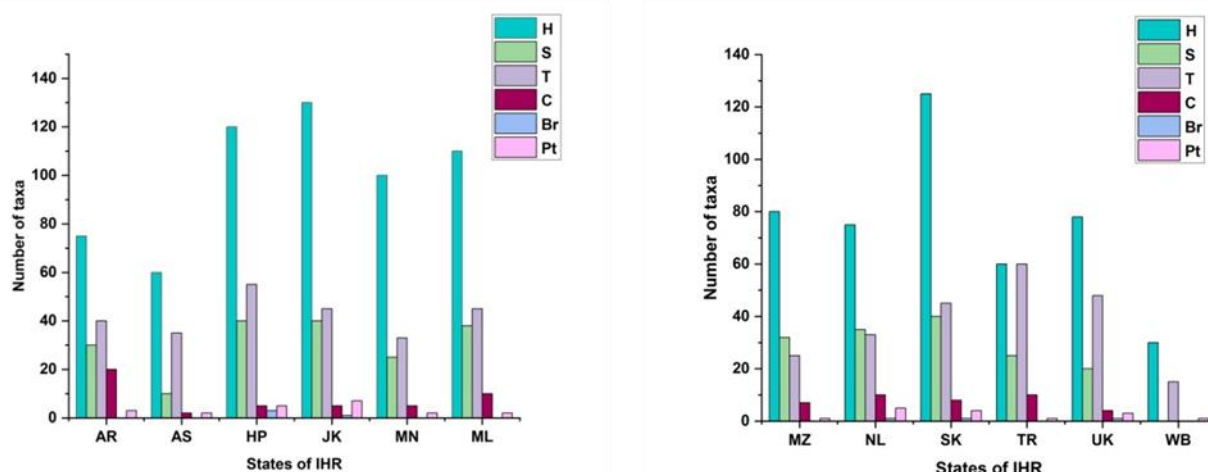


Figure 2: Species at risk within the IHR: H represents herbs, S represents shrubs, T represents trees, C represents climbing enthusiasts, Br represents bryophytes, and Pt represents pteridophytes

Table 2: Life of threatened taxa

States of IHR	Number of taxa					
	H	S	T	C	Br	Pt
MZ	80	32	25	7	0	1
NL	75	35	33	10	1	5
SK	125	40	45	8	1	4
TR	60	25	60	10	0	1
UK	78	20	48	4	1	3
WB	30	0	15	0	0	1
AR	75	30	40	20	0	3
AS	60	10	35	2	0	2
HP	120	40	55	5	3	5
JK	130	40	45	5	1	7
MN	100	25	33	5	0	2
ML	110	38	45	10	0	2

Population trends of threatened taxa

Responding to the variety of ecological and social disruptions, the quantity of plants components has changed through the past few generations. Analyzing the patterns of the endangered plant communities in IHR, it was shown that 38% of the species had a stable tendency of populace, 12% had a falling trend, and just 2% had a rising trend (Figure 3 and table 3). According to IUCN statistics, just 12% of the species is now on the decline, and 35% of the vulnerable plant species are not identified since the location is inaccessible or there isn't enough published research.

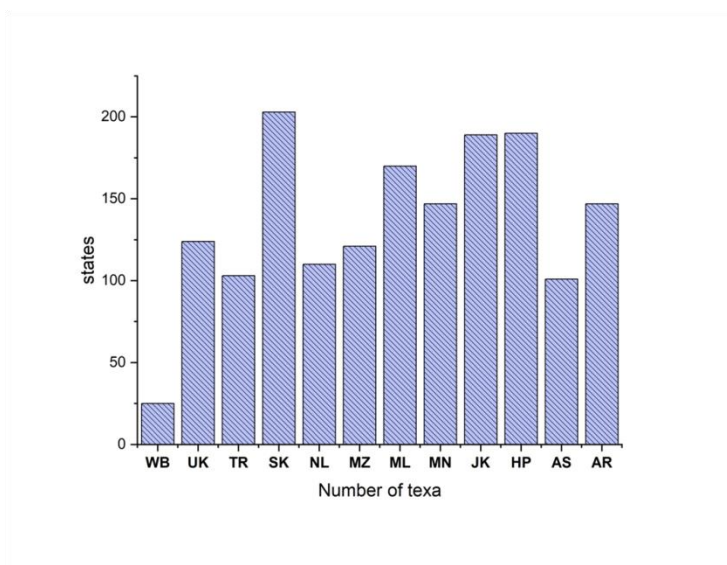


Figure 3:State wise distribution (IHR)

Table 3:Numerical outcomes of distribution state (IHR)

Number of texa	states
UK	124
WB	25
SK	203
TR	103
MZ	121
NL	110
MN	147
ML	170
HP	190
JK	189
AR	147
AS	101

4. Discussion

It is discovered about the biological connections and interdependence that threatened plants have inside their natural surroundings. Sustainability encompasses not solely distinct species as well as the complicated network of interspecies connections. This is difficult to create strategies for preservation that fit the requirements of threatened plants holistically if their connections are not fully understood. To close those spaces and guarantee such vital flora varieties' sustained existence, an integrated strategy integrating cutting-edge conserving tactics, grassroots involvement, and research is analyzed in detail.

5. Conclusion

Protecting threatened plant species is critical to preserving the natural environment and providing supplies required for humanity to survive. Filling up data shortages in studies on biodiversity is enhancing preservation tactics. Putting Threat Evaluation and Tracking are essential to comprehending and reducing the effects of excessive use, invasions, disappearing habitats, and worldwide warming. Precise taxonomic and species recognition are essential to creating conservation approaches that are both successful and integrate conventional environmental information. They also highlight the necessity of cutting-edge

genetics tools. The report also emphasizes how urgently environmentally friendly adaptation techniques are needed the regard to environmental conditions that are shifting. Through the establishment of focused efforts to conserve and the closure of these information spaces, they may work toward the preservation of threatened plant species, which will strengthen habitats and ensure an environmentally sound legacy throughout the future.

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