



Succinct insight of the Ecological Indices for plant species

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Abstract : Introduction: It is reasonably multifaceted to enumerate biodiversity, because of the colossal amount of indices predictable for this purpose. Ecological indices intend to epitomize the general attributes of communities that consent to compare different regions, taxa, and trophic levels as well. These indices are of simple significance for environmental monitoring and conservation. There is no accord about, for the indices for their appropriateness and information shared.

Objective: Several common ecological/diversity indices in an array from simple to complex statistical analyses have been discussed in order to determine which indices are better apposite for definite analyses. The common diversity indices such as Species richness (S), Shannon's diversity (H'), Simpson's diversity ($D1$), Simpson's dominance ($D2$), Simpson's evenness (E), and Berger-Parker dominance (BP) have been discussed in this insight. The trait based measures of diversity allows having an insight whether or not they perform similarly to the superior studied species diversity. Path analysis can be employed for the determination whether compound indices detected additional liaisons between diversities of diverse organisms and traits than basic indices.

Conclusion: Thus, this exhibited that common diversity indices emerge interchangeable in simple analyses. But when taking into account for the complex interactions, the preference of index can intensely amend the construal of results. Data mining for the identification of the index, producing the most momentous results should be circumvented. But at the same time allowing for analyses using multiple indices provides superior insight into the interactions in a system.

Keywords: Ecological indices, Common diversity indices, Plant species, Path analysis, path models.

Introduction

Biodiversity symbolizes the multiplicity and heterogeneity of either organisms or traits at all hierarchical levels of life, ranging from molecules to ecosystems. In general, the focal point is on species diversity, but genetic and chemical diversity, are also imperative and edifying. Even after making a decision on which form of diversity to measure, the predicament that prevails is of quantifying biodiversity as there is no distinct index that sufficiently summarizes the concept [1].

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Richness (S) refers to the numeral of species or attributes present. It is the effortless metric which is worn to signify diversity [2], and thus it lingers as the general widespread applied index [3]

Intuitively, species or trait abundance is furthermore significant for diversity, and the comparative abundance of species can be integrated into the indices embodying diversity. The commonest of these indices was anticipated by Berger and Parker, and has a methodical relationship with the geometric string of the species abundance model [4], and testifies the relative abundance of solitary the most profuse species in the population [5].

There have been copious endeavours to generate compound indices that coalesce measures of richness and abundance. Principal among these are the Shannon's diversity (H') and Simpson's diversity (D_1) indices, which diverge in their theoretical basis and elucidation [6]. H' has its groundwork in information theory and symbolizes the ambiguity about the distinctiveness of an anonymous entity.

In a vastly varied and uniformly disseminated system, an anonymous entity could fit in to any species, leading to a soaring ambiguity in prophecy of its characteristics. In a reduced amount of assorted systems subjugated by one or a few species, it is easier to envisage the individuality of an anonymous entity and there is not as much of vagueness in the system [7]. This metric is widespread in the ecological narrative, regardless of its abstract perception [8]. D_1 is harmonized of Simpson's innovative index and symbolizes the prospect that two arbitrarily chosen entity belonging to different species [9]. D_2 is closely interrelated to D_1 , being the converse of Simpson's inventive index [10]. Together of these alterations dole out to make the index amplified as diversity instinctively increases. Although both are worn, D_2 is more widespread [11].

Objective

The undertaking of such swot has commonly been to gain a better understanding of the human – environment relationship and the aspect affecting it and to stumble on the superior ways to portray plant knowledge patterns. Incidence based species richness, species accumulation curve and similarity measures have also been utilized to evaluate and envisage species richness, to appraise sampling effort and match up to the resemblance of species account for traditional ethnomedicinal data [12].

Ecological Indices

Twenty two different ecological indices can be categorized into;

- ❖ Richness Index (Inclusive of Margalef and Menhinick index)
- ❖ Diversity Index (Inclusive of Shannon and Waver Index, Simpson index Berger-Parker dominance index, McIntosh D, Fisher's Alpha and Hill diversity index),
- ❖ Evenness Indexes (Pielou J or E1, Buzas and Gibson or E2, Heip or E3, Hill or E4 and E5)
- ❖ Species Accumulation Curve
- ❖ Rarefaction
- ❖ Richness Estimators (Chao and Lee 1, abundance based coverage estimator, Incidence based coverage estimator, I^{st} and II^{nd} order jackknife. Hill's diversity numbers to compare proportions of rare, intermediate and common species [13].

Biological community is the set of individuals of all the species of an area which is inclusive of all the organisms present in the area. Extent of species diversity is one of the most imperative tools in the categorization of the communities. To each sample, community, or conversion between two communities, a value is associated that is referred to Diversity Index [14]. The study of diversity is done at three levels:

- ❖ α – **Diversity**: It is intra-community diversity. It can be measured in terms of species richness and evenness.
- ❖ β – **Diversity**: It is inter-community diversity and connotes the change in species along a gradient of habitats.
- ❖ γ – **Diversity**: This is the highest level of diversity measurement. It is measured at the pooled communities.

There are quite a few restraints associated with the extent of diversity:

- ❖ Diversity of a habitat varies with day time and the season due to incursion and retreat of species.
- ❖ Sampling variations and veiled species causes variations in diversity indices.

- ❖ Diverse sampling methods and diversity measures yields altered diversity indices for the same area.
- ❖ Most of the mathematical postulations in the formulation of diversity measures may not embrace exact [15].

Discussion

Hill's numbers endows with a technique to illustrate the liaison involving diversity indices and the standards of N_1 (Shannon – Wiener, base e), N_2 (Reciprocal of Simpson's index, $1/D$) and N'' (Reciprocal of Berger – Parker index), corresponds with the evaluation of profuse, very profuse, and most profuse species in a sample, respectively. The value of N'' can be construed as quantify of the familiar species, $N_1 - N''$ can be elucidated as assess of the number of intermediate species, and $N_0 - N_1$ matches to a measure of rare ones [16].

Michaelis – Menten means estimators were the paramount estimator because the curve loomed a horizontal asymptote. E_1 , also called the Shannon J_0 or Pielou's J , is possibly the most numerous evenness index used; but is sturdily exaggerated by species richness, and the totting up of rare species (or singletons) can significantly change the value of E_1 . Therefore it is recommended for the use of E_1 and E_2 (Buzas and Gibson E). A broad-spectrum predicament with all measures of evenness, however, is that they presuppose the totality of number of species that could probably be sampled, is known [17].

The **Evenness Ratios** are observed or taken in to consideration as the observed species numbers ought to be always being less than true species richness, with the probable exclusion of E_4 and E_5 . E_4 and E_5 remain comparatively constant with sampling disparities and hence tend to be self-determining of sample size. This is for the reason that, E_4 and E_5 are calculated as ratios, where S is in both the numerator and the denominator; thus in actual fact cancelling the brunt of the number of species in the sample. However, E_4 and E_5 are not completely unaffected by the outsized number of singletons which are found in undersized samples, inclusive of the samples collected in the preliminary stages of research at a site before an satisfactory sample size is mounted up [18].

Rarefaction provides a scheme of assessment linking different communities, whereby each community is 'rarefied' and back to an equal number of sampled specimens. It enables evaluating sampling efforts. Interpolation and sample-based rarefaction eliminates the requirement for re-sampling methods and thus authorizes an undeviating statistical assessment of the species richness between sampled sets. Rarefaction curve can be utilized to evaluate sampling effort and investigated disparities in plant use par category of uses, for example, age and gender within unlike communities. Thus, it can be advocated that such methods have the probable to be broadened to species richness between sites, as well as for the approximation of the number of species with an absolute census of the plants used or trade being possible. Though rarefaction can be constructive, it is very susceptible to the underlying pattern of species abundance, this implies that with much lower species evenness will often result in lower estimates of species diversity than those with even abundances, despite of species diversities in veracity are equal [18].

Species Accumulation Curves (SAC) are known to augment the assessment of Ethnomedicinal studies and thus creates prospect for rational squabble that provides advancements in scientific and practical knowledge. This curve can also be worn as a way of estimate for species richness. This is done commonly with fitting function such as the asymptotic Michaelis – Menten algorithm or Non – Asymptotic estimators such as log – linear model. This technique has been utilized to stumble on out the association between village studies and plants utilized in recoveries. SAC also provides an indication about a decline in the number of fresh species mentioned per interview. Further interviews are likely to draw out additional less prominent species. SAC can be utilized to study the traditional use of genotype and thus a comparison can be done it with nonparametric estimators of species richness. Thus, this curve has been utilized to assess the expected richness of used and known plants by the number of plant species [19].

The **Jackknife** is valuable because it is identified to decrease preconception and for estimation of species richness. Another vital feature of the Jackknife estimator of species richness is that the estimator is based on the incidence or dearth of a species in a given plot rather than on the profusion of the species [20].

Conclusion

In Ethnobotany there are many structures and outlines for collection of data which comprises of field audits, factiontreatise and visit to herbalist. Each and every informant or an event may have aptitude to engender a huge statistics which in due course fabricates a maze of rhetorical information. With appreciation of this entanglement, now Ethnobotany has tailored scores of statistical advances that can solve the data matrix into a logical conclusion. This study shows that there are many quantitative approaches have been utilizing all over the world, out of which some of the techniques are well adapted by many researchers, while some techniques are region specific or rather objective specific. Two specific branches namely Anthropology and Ecology significantly chipped in to build "*Quantitative Ethnomedicine*". In the current review, surplus diversity indices were compiled for the first time which provided a vital database for progressive researchers. This technical paper also compiled the advanced information of assorted ecological methods, currently utilized in Quantitative Ethnomedicine.

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Conflict of Interest

Nil

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