Biodiversity of reefs: inferring from sparse data

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Abstract. Data on occurrences of a particular organism from publications and museum specimens can be used to infer occurrence of members of that species in places where sampling has not been done and at times in the past and future. Programs to make such inferences are based on knowledge of the habitat correlates of the species and determining where else in the world those habitat parameters occur. Precision of such an inference depends on 1) accurate and precise knowledge of the species’ habitat requirements, 2) detailed spatially-explicit environmental data, and 3) comprehensive taxonomic and nomenclatural information. Such inferential tools can be important in understanding biogeographic consequences of climate change, in predicting where invasive species might persist, and in recognizing invasive species.

Key words: biogeographic modeling, habitat requirements, taxonomy.

The problem
Species range maps like those in many field guides (eg that to anemonefishes and their hosts by Fautin and Allen 1992, and that to marine life of southern Africa by Branch et al. 2005), are abstractions of the places where members of the species of concern are known to occur. Although the open Indian Ocean between Sumatra and Sri Lanka is depicted by Fautin and Allen (1992) as being within the range of the sea anemone Heteractis aurora (Fig. 1), one could not actually find the animal there because, as Fautin and Allen (1992) point out, these animals are confined to shallow water. Thus they occur along land, in strips so narrow they are not easily indicated on small-scale maps – so the range is represented as covering the entire sea bounded by known or inferred occurrences at the extremes. Local guides portray a smaller portion of the globe, and so can indicate ranges in strips along the coast, as Branch et al. (2005) do, but the strips are continuous, and it is likely that members of most species occur in only some of the places indicated, occurrences dictated by the presence of appropriate habitat. Such abstractions over-represent the range of a species.

Maps indicating actual occurrences are typical in technical publications, such as Wallace (1999), and in the growing number of biodiversity websites such as that to sea anemones and relatives (Fautin 2008: eg Fig. 2). With few exceptions, it is likely that members of a species occur in more than the places indicated on the maps, which are where they have been collected. These concrete depictions under-represent the range of a species.

A solution
As has been known for many years (Allee et al. 1949), to a first approximation, an organism’s range can be inferred from abstracting its habitat characteristics based on places the organism is known to occur, then assuming the organism occurs where its habitat does. Computerized methods make the four-step process of developing such range maps theoretically straightforward. 1) Occurrence records of a species and 2) maps depicting the distribution of relevant environmental parameters are overlain. 3) An algorithm abstracts the values of the environmental parameters that coincide with occurrence records, then 4) maps the occurrence of the environmental parameters within the values known to be compatible with the existence of the organism.

The distribution of these values of environmental parameters is a first-cut inference of the occurrence of habitat suitable for the species and thus the potential range of the species. Modeled environmental data for the past and future allow inferences to be made about historical biogeography and how ranges may shift with climate change.

Many models have been developed to carry out such computations, differing in assumptions and underlying algorithms. Some are referred to as “niche models” but few actually consider biological features other than by proxy with habitat. Sixteen such modeling programs were tested against one another
Some considerations
Presumably, precision of the environmental data and knowledge of a species’ habitat requirements correlate directly with reliability of a range inference. Accuracy of the inference of a species’ range should also increase with improved knowledge of the known distribution of the species. One way to maximize information of a species’ occurrence is to include all records for the species regardless of the scientific name that was used for it.

Although ideally a single scientific name refers to a single species, many species have been given more than one name (such names are synonyms; International Commission of Zoological Nomenclature 1999). The sea anemone species for which the valid name is *Heteractis aurora* (Quoy and Gaimard, 1833) has been referred to in publication by 11 names (Fautin 2008): it was described as a new species three times, has been placed in nine genera, and has been misidentified.

The points on distribution maps in “Hexacorallians of the World” (Fautin 2008), which depict occurrence records from the published literature, are color-coded by the name used for the species. Because synonyms are a matter of taxonomy – they depend on a
scientist’s opinion, and scientists may legitimately differ on whether two names (or more) refer to a single species – this convention allows a user to distinguish among the records by name, disregarding those that the user considers do not refer to the species in question. In fact, such a depiction can be informative in deciding whether a particular synonymy is justified.

Figure 3 is the KGS Mapper output of suitable habitat for *Heteractis aurora* based on 75 occurrence records and the environmental parameters mean depth, mean and minimum surface seawater temperature, and maximum and minimum salinity. These parameters were selected because they are relevant to the biology of these anemones. The animals are zooxanthellate, so occur only in shallow water, and are confined to warm, fully saline seas. They are also known only from the Indo-Pacific. Within that natural range, the model output depicts the animal’s potential range; this can be used to infer occurrence of members of the species in places where sampling has not been done, which can be useful, for example, in planning fieldwork, or inferring areas into which the animals might expand as climate changes. Outside that natural range (in the Caribbean, for example), the model output can be used to infer where the species might unnaturally occur, as, for example, where it might invade and persist. On the website “Hexacorallians of the World,” the darker the reddish color, the more likely the habitat is to be suitable (Guinotte et al. 2006).

Figure 4 depicts the habitat suitable for *Heteractis aurora* based on the same environmental parameters as for Figure 3, but uses only those occurrence records that referred to the species by one of its synonyms, *Radianthus koseirensis*. Presumably both the smaller number of records and the restriction of them to the western portion of the species’ range leads to the conclusion that the species is likely to be
far less widespread than is seen in Figure 3. When rendered in black and white, the difference in model output is easiest to perceive in Southeast Asia, where the shaded area of inferred occurrence is more widespread and darker in Figure 3 than Figure 4. Plotting occurrences using a species name rather than a species concept (by including synonyms) can be not only incomplete but may be misleading. That is because more than one species may have been given the same scientific name (such names are termed homonyms; International Commission of Zoological Nomenclature 1999). Homonyms are not a matter of opinion, but are objectively verifiable, and thus fall within the province of nomenclature. The anemone species to which the name *Heteractis aurora* properly belongs occurs strictly in tropical waters of the Indo-Pacific. That species was first placed in the genus *Actinia* (to which nearly all sea anemones were initially assigned). What is clearly a different species in Britain was also given the name *Actinia aurora*. Applying a model to infer distribution from records that include occurrences of the British species would obviously give a meaningless result.

**Conclusions**

Use of modeling programs to infer the range of a species from known occurrences and features of the environment benefits from

- accurate and precise knowledge of the species’ habitat and natural history (to select environmental parameters relevant to the organism’s life that control its distribution)
- knowledge of biogeography (where the organism would be expected to occur, so invasions can be distinguished from occurrences in places where the species might be expected but has not previously been recorded)
- consideration of taxonomy and nomenclature (what other names have been applied to it – and which applications of a name do not refer to the species in question)

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