

1 For the virtual issue, "Red Sea and Western Indian Ocean Biogeography"

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3 **Red Sea and Western Indian Ocean Biogeography**

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24 **EXECUTIVE SUMMARY**

25 The Red Sea and Western Indian Ocean (WIO) represent iconic and strategic locations in our
26 tropical oceans, and yet few studies address the origins, maintenance, and distribution of marine
27 fauna of these locations. This virtual issue of the *Journal of Biogeography* (JBI) serves to
28 illustrate the biogeographical advances in these habitats at the periphery of the vast Indo-Pacific
29 region with an editorial introduction, 14 contributions from an international workshop, and a set
30 of 24 papers previously published in JBI. The selection includes a number of influential research
31 studies that present novel data, those that apply methods to synthesize general patterns, and some
32 that utilize newer genetic methods to infer patterns of the evolutionary history of coral reef
33 ecosystems within this region. Several key themes emerge from this collection of papers:

- 34 • There appears to be no single, repeated, or uniform explanation for the region's evolutionary
35 history; the evolutionary path for each species appeared to be unique. It does not appear that
36 there was a single, all-encompassing vicariance event that shaped the evolution of regional
37 species.

- 38 • Many cryptic species have been revealed by molecular work, both in groups with ambiguous
39 taxonomy and within the best-characterised groups. This approach clarifies patterns of
40 regional endemism.

- 41 • Many of the previously defined or postulated biogeographical regions mirror genetic
42 boundaries, but there are important exceptions for some species.

- 43 • There is substantial potential for regional species to serve as model study systems to address
44 topics such as the impacts of climate change on biogeography; understanding the connections
45 among generation of biodiversity, evolution, and the environment; and elucidating

46 mechanisms that maintain biodiversity and/or promote speciation in the 'marginal' seas of the
47 WIO.

48 **EDITORIAL**

49 “A moment later I was overboard. Now, in the bright midday light, the coral looked even more
50 brilliant and colourful than the night before. Directly below me was a peach-coloured bush;
51 from Klunzinger’s writings I immediately recognized it as a cup star coral. There were violet-
52 hued bunches of pistil coral, and between them I spotted a vermilion *coenopsammia*. There were
53 luxuriant growths of dozens of species of star coral in the shape of brightly coloured balloons
54 and blisters; snow-white thistle coral filled the hollows and staghorn coral extended their
55 delicate rose-coloured tips. And brilliant coral fish were flitting about these fairy-tale gardens
56 like butterflies.”

57 – Hans Hass, recounting his first dives in the Red Sea, from *Challenging the Deep* (1973)

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59 Hans Hass made his first pioneering dives within the Red Sea in 1949. He relied on works
60 from the great naturalists dating back to the previous century, but still found great excitement in
61 observing these fauna in their natural habitat. Since the time of Hass’ explorations, much has
62 changed. While traditional natural history expeditions still play an important role in scientific
63 discovery, the modern era of biogeography now seeks mechanisms (or processes) to explain the
64 observed patterns on a level that few in Klunzinger’s or Hass’ time could have imagined.
65 Establishing the distribution of a species is now followed by questions of genetic similarity to
66 conspecifics throughout their range. In some cases, the focus has shifted to understanding the
67 spatial and temporal variability of a “hologenome” as a factor in local adaptation, recognizing
68 that while a species may appear to be genetically homogeneous, the multitude of
69 microorganisms, symbionts, and viruses associated with each individual may be highly variable
70 in space and time (Zilber-Rosenberg & Rosenberg, 2008). And yet there remain some areas of

71 the world where physical acquisition of samples and deposition of these samples into natural
72 history collections still play a particularly important role in elucidating biogeographical patterns
73 (Rocha *et al.*, 2014). The Red Sea and Western Indian Ocean (WIO), for example, form a marine
74 region bordered by many countries in which the socio-economic and political realities have
75 limited access for many scientists, thus hindering a modern perspective of general biogeography
76 in the region. While local expertise and knowledge undoubtedly exist here, the accessibility of
77 this information to the international scientific community is sometimes restricted (Berumen *et*
78 *al.*, 2013; Loya *et al.*, 2014). This region is particularly interesting because it includes multiple
79 biogeographical provinces (Spalding *et al.*, 2007; Briggs & Bowen, 2012), each with distinct
80 faunal communities and suites of endemic species (e.g., DiBattista *et al.*, 2016a), and these
81 provinces have numerous intersections within the region (e.g. DiBattista *et al.*, 2015a). While
82 many consider the WIO region to be “marginal” or “peripheral” habitat, it may still play an
83 important role in global biogeography (e.g., as an exporter of biodiversity) (e.g. Bowen *et al.*,
84 2013).

85 Herein we introduce a special virtual issue in the Journal of Biogeography (JBI) titled
86 "Red Sea and Western Indian Ocean Biogeography". The overall aim of the present virtual issue
87 is to highlight advances in our understanding of marine biogeography in this fascinating region.
88 The virtual issue arises largely as the result of a workshop convened by the King Abdullah
89 University of Science and Technology (KAUST) in Saudi Arabia to bring together expertise on
90 regional biogeography in early 2014. As a leading journal in this field, JBI was a logical place to
91 house the proceedings of this workshop. The virtual issue highlights previously published works
92 in JBI and provided the impetus for new papers to find a common repository. It currently
93 includes 24 previously published papers and 14 papers specifically submitted to the thematic

94 issue and it creates the opportunity to add additional papers to the collection as they are
95 published.

96 The Red Sea and WIO are reported to host endemism and biodiversity hotspots. Two review
97 papers capture the past and present patterns of endemism surrounding the Arabian Peninsula.
98 DiBattista *et al.* (2016a) summarize the patterns of endemism for several broad taxonomic
99 groups, finding consistently high levels of endemism within the Red Sea and variable patterns in
100 adjacent regions. Turning to the question of how these patterns of endemism arose, DiBattista *et*
101 *al.* (2016b) explore competing hypotheses but conclude that there is no single explanation.
102 Focusing on the corals of the Indian Ocean, Obura (2016) reaches a similar conclusion; multiple
103 origins of diversity are likely responsible for the current biogeographical patterns. McClanahan
104 (2015) offers a perspective on applications of such biogeographical patterns in connection with
105 management efforts to conserve reef fisheries; based on data from seven southwestern Indian
106 Ocean countries, he proposes a “portfolio approach” that spreads risk by managing a variety of
107 habitats and locations, but with a particular focus on protecting biomass in high-compliance
108 areas.

109 Several studies in this virtual issue focus on wide-ranging phylogeographical patterns,
110 placing the region in a much broader context. Ahti *et al.* (2016) examine cosmopolitan wrasses,
111 finding that Indian Ocean and Pacific Ocean sister species may be better described as subspecies.
112 Waldrop *et al.* (2016) highlight the evolution of range-restricted butterflyfish species (Red Sea
113 and NW Indian Ocean) and isolated populations (Hawai’i), concluding that genetic divergence is
114 high among these peripheral biogeographical provinces. On the other hand, in a wide-ranging
115 butterflyfish species, DiBattista *et al.* (2015b) find that genetic differentiation among studied
116 populations was insufficient to justify a species- or subspecies-level distinction of a previously

117 proposed Red Sea subspecies, despite different colour morphs across its range (*i.e.*, the two
118 colour morphs do not sort into reciprocally monophyletic groups of mtDNA haplotypes).
119 Fernandez-Silva *et al.* (2016) examine a goatfish species and find that phylogeographical breaks
120 are congruent with existing boundaries of biogeographical provinces, and that these peripheral
121 biogeographical provinces may represent evolutionary incubators for reef fishes (*sensu* Bowen *et al.*
122 *al.*, 2013). Broadening the taxonomic scope, Iacchei *et al.* (2016) find similar concordance of
123 phylogeographical and biogeographical barriers in the spiny lobster. Higher resolution studies
124 such as Priest *et al.* (2016) explore fine-scale genetic and life history patterns in a grouper around
125 the Arabian Peninsula, suggesting that cryptic speciation exists as a result of biogeographical
126 barriers within the Arabian region. Van der Ven *et al.* (2015) explore population genetics of an
127 *Acropora* coral species along the Kenyan and Tanzanian coasts, concluding that high
128 connectivity was most likely explained by local oceanographic features rather than by
129 geographical distance. Within the Red Sea, Reimer *et al.* (2016) examine the type (*i.e.*, clade) of
130 symbiotic algae residing in a zoantharian host and find that the local thermal regime is the most
131 likely driver of their distribution. As invasive species become even hotter topics in various parts
132 of the world, the role of the Red Sea fauna in the Lessepsian invasion of the Mediterranean is
133 examined by Bariche *et al.* (2015), who use DNA barcoding to identify numerous cases of
134 invasive fishes. Bariche *et al.* (2015) revealed some possible cases of previously unrecognized
135 invasions as well as invasions by cryptic species. The authors found some taxa with unresolved
136 taxonomic status, highlighting the need for updated and vouchered genetic libraries for the
137 region.

138 Our selection of papers from the JBI archives highlights seminal work on the origin of
139 tropical marine species (Briggs, 2000) and a realignment of biogeographical provinces (Briggs &

140 Bowen, 2012, 2013). We include studies that apply phylogenetic or phylogeographical methods
141 in order to advance our understanding of drivers of speciation and subsequent dispersal in reef
142 fish (Cowman & Bellwood, 2013; DiBattista *et al.*, 2013; Gaither & Rocha, 2013; Hodge *et al.*,
143 2014; Ludt & Rocha, 2015; Hodge & Bellwood, 2016), non-coral invertebrates (Reid *et al.*,
144 2006; Hoareau *et al.*, 2013), or corals (Pinzón *et al.*, 2013) and their symbionts (LaJeunesse *et al.*,
145 2010). Collectively, these studies show that there are not "one size fits all" generalities for
146 patterns of speciation and genetic connectivity within the WIO. Rather, each taxon may have its
147 own unique evolutionary history (even congeners may have different histories) with
148 correspondingly unique mechanisms. We additionally include studies focused on an important
149 transition zone along the South African coastline (Bustamante & Branch, 1996; Primo &
150 Vazquez, 2004; Teske *et al.*, 2014), studies teasing apart key processes affecting both ancient
151 (Fernandes *et al.*, 2006) and more recent migration (Roberts *et al.*, 1992; Golani, 1993; Kiflawi
152 *et al.*, 2006) within and out of the Red Sea, studies focused on the isolating effects of the Arabian
153 Sea upwelling (Kemp, 1998; Schils & Coppejans, 2003), a study using holothuroids as a model
154 to understand zoographical patterns in the Western Indian Ocean (Samyn & Tallon, 2005), and
155 studies that remind us we must not now nor ever forget to consider the ecology of a system when
156 defining biogeographical boundaries (Choat, 2006).

157 Our sincere gratitude goes to the Editor-in-Chief (Peter Linder, who handled the transition in
158 the role from Rob Whittaker); the handling editors for individual papers in the virtual issue; and
159 numerous authors and reviewers, whom all created and shaped the content of the final
160 contribution. Input from Dr Michelle Gaither and three anonymous reviewers improved this
161 introductory note. The virtual issue represents a major step forward in our ability to consider the

162 “big picture” supporting the origins and maintenance of marine fauna in the Red Sea and WIO,
163 thus following in the footsteps of the likes of Klunzinger and other regional explorers.

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165 *“I suspected that the Red Sea still had a lot to reveal to us, a lot which had remained hidden*
166 *from me on my first visit. I was to be proved right.”*

167 – Hans Hass, from *Challenging the Deep* (1973)

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