



NIMBioS

National Institute for Mathematical
and Biological Synthesis



NIMBioS Interdisciplinary Seminar

3:30 p.m.*, **Monday, September 28, 2015**

**Join us for refreshments at 3 p.m.*

Dr. Stuart Pimm

**Doris Duke Professor of Conservation
Nicholas School of the Environment, Duke Univ.**

“The laws of biodiversity and what we do not understand about them”

The practice of conservation biodiversity starts with understanding which species and which places are priorities in preventing species decline and extinction. A minority are obvious and familiar: large bodied vertebrates, especially predators, are hunted or persecuted over their large geographical ranges. The large majority of threatened vertebrates have small geographical ranges leading them to be vulnerable to habitat destruction. Armed with where these species live, we can make strategic choices of which places are world priorities — the “biodiversity hotspots” — and downscale to tactical decisions on exactly where and how to protect them. There’s an obvious problem. We are protecting biodiversity based on 0.1% of species. Taxonomists are not going to name, let alone map, the remaining 99.9% anytime soon. This motivates a search for general patterns of biodiversity — I call them “laws” — and an understanding of them that might help address whether they are likely to apply equally to known and unknown taxa. Wallace promulgated the most famous law of biodiversity: “Every species has come into existence coincident both in space and time with a pre-existing closely allied species.” A simple enough pair of observations, yet ones with profound implications. There are others. The sizes of species ranges have consistent log-normal distributions. Widespread species are locally common, whereas local species are generally rare. Species densities vary greatly from place to place. Species with small geographical ranges are geographically concentrated and generally not in the places where the largest numbers of species live. There is the familiar species-area relationship and there is a distinctive pattern of species loss where habitats are destroyed. To be credible, the theories to explain these laws need to be mechanistically credible, testable against new data, and make hitherto unexpected predictions. Most theories fail these criteria, suggesting considerable effort is needed to understand biodiversity patterns.

Location: Hallam Auditorium, Rm 206, NIMBioS, Claxton Education Bldg, 1122 Volunteer Blvd.

The seminar will be live streamed. Visit <http://www.nimbios.org/videos/livestream>.

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The National Institute for Mathematical and Biological Synthesis (NIMBioS) brings together researchers from around the world to collaborate across disciplinary boundaries to investigate solutions to basic and applied problems in the life sciences. NIMBioS is sponsored by the National Science Foundation with additional support from The University of Tennessee, Knoxville

