

A Mini Review on Evolutionary Biology

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Abstract

Not everyone agrees that evolutionary biology has a place in medical education. Deans of medical schools and other educators frequently request proof that understanding evolutionary biology will increase the effectiveness of medical specialists. Citing concrete examples is a straightforward response. For instance, doctors need to be aware of how antibiotic resistance has developed, how pathogen phylogenies can be tracked, how selection has shaped the mechanisms that control protective responses like pain and fever, and the intricate relationships between evolution, the environment, and aging-related diseases. However, focusing only on such obvious applications undersells the value of evolutionary biology in medicine. Although it doesn't directly apply to daily life, basic science education in medicine is important for understanding the body and disease. "The principles that underlie biological complexity, genetic diversity, interactions of systems within the body, human development, and influence of the environment guide our understanding of human health, and the diagnosis and treatment of human disease," according to the AAMC-HHMI report's overarching principle number 2. Calculus, physics, and chemistry proficiency are required not because these subjects are frequently used in clinics but rather because doctors who are proficient in them will have a better understanding of the human body and will be able to make better medical decisions.

Keywords: Biology • Evolutionary • Genetic diversity

Introduction

The evolutionary synthesis (ES) often known as the modern synthesis or the ES, as Huxley termed it, gave rise to the current conceptual framework of evolutionary theory. Any discussion of extending or altering present theory will benefit from some knowledge of the ES's past and how the field has developed since. My understanding of the history of biology, and specifically the history of evolutionary biology, is that it has generally been gradual rather than paradigm-shifting, building on earlier discoveries [1]. For instance, the "rule" of independent assortment had to be changed to take linkage into account immediately after Mendel's "laws" were discovered and canonised in the first decades of the twentieth century. From being a discrete "factor," the "gene" evolved into a trinity of recon, muton, and cistron. Phylogeny, field investigations of natural selection, evolutionary genomics, and a plethora of genetic phenomena that were unimaginable in the 1940s or even the 1960s have all been steadily incorporated into theory, informing (and occasionally foretelling) it. Transposable elements, exon shuffling, chimeric genes, gene duplication and gene families, whole-genome duplication, de novo genes, gene regulatory networks, intragenomic conflict, kin selection, multilevel selection, phenotypic plasticity, maternal effects, morphological integration, evolvability, coevolution, and more are all recognised and studied in modern evolutionary biology. Some of these phenomena and ideas were unknown or only dimly understood a few decades ago [2].

Description

It is crucial to understand that "mutation" in population genetics theory refers

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to any new alteration of the genetic code that is stably passed down through generations. Although the fundamental principles of population genetics were not altered by the revelation of the molecular basis of inheritance following the ES, our knowledge of the evolutionary process and its history was substantially expanded. The fundamental hypothesis, for instance, leaves open the question of what constitutes mutation-gene duplication, a single base pair alteration, the insertion of a transposable element into a regulatory sequence, or the doubling of the entire genome, for instance. New types of mutations, such transposable elements, have been incorporated into the population genetics framework as they are identified [3].

Conclusion

Natural selection was and still is frequently believed to have its roots in the natural environment, although the creators of the ES were well aware that selection had much broader roots. Wright emphasised epistasis for fitness, in which prevalent alleles at one locus affect the selective value of alleles at another locus. Fisher described the evolution by selection of sex ratio, selfing, and outcrossing and he provided a genetic interpretation of Darwin's idea of sexual selection. "Internal selection"; mutations can affect how physiological and developmental processes work, which in turn affects viability and reproduction. Molecular, behavioural, ecological, or other types of data are required for a causal explanation of any occurrence of selection, but demonstrating the fundamental process of biological evolution occurs at the population level rather than the individual level and involves changes in the frequency of heritable variations within populations from generation to generation rather than the individual organism. While Rensch, Simpson, and Mayr, organism-focused evolutionary biologists, had a more comprehensive conception of evolution, including phenotypic evolution, speciation, and differential proliferation of clades, while acknowledging that phenotypic evolution and speciation occur by changes in allele frequencies, Dobzhansky defined evolution as a change in allele frequencies. Although they acknowledged allele frequencies as the fundamental, generation by generation mechanism of change, Rensch and Simpson did not discuss allele frequencies when they wrote about "Evolution above the species level" and "Tempo and mode in evolution," respectively [4,5].

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