



Special Issue

Biological Statistical Mechanics

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Message from the Guest Editors

Dear Colleagues,

Any natural entity can be imagined as a system made up of interacting elements. This allows for the development of phenomenological “laws” shared by network-like systems only dependent on their wiring architecture. We can compare this situation with the success of classical thermodynamics, even if the founding fathers of this science were erroneously convinced that heat was a fluid. The difference with classical thermodynamics is that in the case of biological systems we cannot rely on macro-parameters like volume or pressure, but we must seriously consider the particular correlation structure of the system at hand. This is why focusing on state transitions (e.g., differentiation, development, onset of diseases, ecosystem de-stabilization) in which we expect abrupt changes of the system correlation structure is probably the most fruitful direction to establish a “biological statistical mechanics”. This Special Issue is devoted to the collection of statistical mechanics-inspired approaches to biological systems at any scale of definition from cell biology to ecology and epidemiology. The issue is of crucial importance given the evident failure of strictly deterministic molecular biology approaches to predicting system-level properties of biological entities.

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