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Introducing a handbook for gene regulatory network research using evolutionary computation, with applications for computer scientists, computational and system biologists. This book is a step-by-step guideline for research in gene regulatory networks (GRN) using evolutionary computation (EC).

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Using evolutionary algorithms to study the evolution of gene regulatory networks controlling biological development Alexander Spirov Computer Science and CEWIT, SUNY Stony Brook, Stony Brook, NY, USA; and the Sechenov Institute of Evolutionary Physiology and Biochemistry, St. Petersburg, Russia

Using evolutionary algorithms to study the evolution of ...

The evolution of high throughput technologies that measure gene expression levels has created a data base for inferring GRNs (a process also known as reverse engineering of GRNs). However, the nature of these data has made this process very difficult.

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