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| Collana | Studies in Computational Intelligence, , 1860-9503 ; ; 1109 |
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| Soggetti | Computational intelligence Computer simulation Neural networks (Computer science) Biological models Computational Intelligence Computer Modelling Mathematical Models of Cognitive Processes and Neural Networks Biological Models |
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| Nota di bibliografia | Includes bibliographical references. |
| Nota di contenuto | 1. Exponential time-differencing method for the solution of diffusive HIV model -- 2. An effective technique for solving a model describing biological species living together -- 3. Neuro-Swarming Integrated Heuristic with an Interior-Point Scheme for Novel design of Lane-Emden Nonlinear Prediction Singular Differential Model -- 4. A fitted operator method for a system of delay model of tumor cells dynamics within their micro-environment -- 5. A Mathematical Model to Study Regulatory Properties and Dynamical Behaviour of Glycolytic Pathway using Bifurcation Analysis -- 6. On solutions of fractional biological models using reproducing kernel Hilbert space method -- 7. An operational matrix based method to find the solution of the fractional tumour immune vitamins model -- 8. Analysis of a fractional stage-structured model with Crowley-Martin type functional response by Lagrange polynomial based method -- 9. Qualitative Theory and Approximate Solution to Typhoid Fever Model Subject to Non Singular Kernel Type Derivative -- 10. Study of the SIRI epidemic model |

described by the Caputo derivative -- 11. Unlocking Biological systems through mathematical Modelling -- 12. Implementation of vaccination in epidemic model for COVID-19.

Sommario/riassunto

This book discusses computational methods related to biological models using mathematical tools and techniques. The book chapters concentrate on numerical and analytical techniques that provide a global solution for biological models while keeping long-term benefits in mind. The solutions are useful in closely understanding biological models, and the results will be very useful for mathematicians, engineers, doctors, scientists and researchers working on real-life biological models. This book provides significant and current knowledge of biological models related to real-life applications. The book covers both methods and applications.