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Computational network biology: Data, models, and applications. Author links open overlay panel Chuang Liu a Yifang Ma b Jing Zhao c Ruth Nussinov d e Yi-Cheng Zhang i a Feixiong Cheng f g h Zi-Ke Zhang j a. Show more. ... studying the biological systems from network biology has attracted much attention in recent years, and networks have long ...

### Computational network biology: Data, models, and ...

An ecological network is a representation of the biotic interactions in an ecosystem, in which species are connected by pairwise interactions. These interactions can be trophic or symbiotic. Ecological networks are used to describe and compare the structures of real ecosystems, while network models are used to investigate the effects of network structure on properties such as ecosystem stability.

### **Ecological network - Wikipedia**

1.022 - Introduction to Network Models. Amir Ajorlou. Laboratory for Information and Decision Systems Institute for Data, Systems, and Society Massachusetts Institute of Technology. Lecture 20. Lecture 20. Introduction to Network Models 1 /

## 1.022 - Introduction to Network Models

These data then need to be integrated into various network models or multiscale models. Models are a fundamental step in the scientific discovery. In this article, we described different types of models that have been used in biology for knowledge discovery and predictions. However, building a good model is a hard task.

## Mathematical modeling of biological systems | Briefings in ...

population coding; neural network model; Much information that the brain processes and stores is temporal in nature. Therefore, understanding the processing of time in the brain is of fundamental importance in neuroscience  $(1 \downarrow \downarrow -4)$ . To predict and maximize future rewards in this everchanging world, animals must be able to discover the temporal structure of stimuli and then flexibly anticipate or act correctly at the right time.

## Understanding the computation of time using neural network ...

logical model must have a clear interpretation such as a contact rate or a duration of infection. Models can be used to assess many quantitative conjectures. For example, one could check a conjecture that AIDS incidence would decrease if90% of the sexually active heterosexual population started using condoms consistently.

# Three Basic Epidemiological Models

Population models are used to determine maximum harvest for agriculturists, to understand the dynamics of biological invasions, and for environmental conservation. Population models are also used to understand the spread of parasites, viruses, and disease. Another way populations models are useful are when species become endangered.

#### **Population model - Wikipedia**

1. Introduction Part I. Single Neuron Models: 2. Detailed neuron models 3. Two-dimensional neuron models 4. Formal spiking neuron models 5. Noise in spiking neuron models Part II. Population Models: 6. Population equations 7. Signal transmission and neuronal coding 8. Oscillations and synchrony 9. Spatially structured networks Part III. Models of Synaptic Plasticity: 10.

# [PDF] Spiking Neuron Models: Single Neurons, Populations ...

A synthesis of contemporary analytical and modeling approaches in population ecology The book provides an overview of the key analytical approaches that are currently used in demographic, genetic, and spatial analyses in population ecology. The chapters present current problems, introduce advances in analytical methods and models, and demonstrate the applications of quantitative methods to ...

#### Population Ecology in Practice | Wiley

We consider the recently introduced edge-based compartmental models (EBCM) for the spread of susceptible-infected-recovered (SIR) diseases in networks. These models differ from standard infectious disease models by focusing on the status of a random partner in the population, rather than a random individual.

# Incorporating Disease and Population Structure into Models ...

In this work, we focus on two types of spiking network models: clustered networks and classic balanced (non-clustered) networks. To study single-trial population activity, we used factor analysis (FA), a linear dimensional- ity reduction method [2, 4, 20–22]. A key feature of FA is that it

partitions spike count

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