



Knowledge Discovery in Proteomics (Chapman & Hall/ Crc Mathematical Biology and Medicine)

Igor Jurisica, Dennis Wigle

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Systems biology

In each area, the authors describe the challenges created by the type of data produced and present potential solutions to the problem of data mining within the domain. They take a systems approach, covering individual data and integrating its computational aspects, from data preprocessing, storage, and access to analysis, visualization, and interpretation.

With clear exposition, practical examples, and rich illustrations, this book presents an outstanding overview of this emerging field, and builds the background needed for the fruitful exchange of ideas between computational and biological scientists.

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