



Bioinformatics: Problem Solving Paradigms

Volker Sperschneider

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What is this book good for? Imagine you are a computer scientist working in the bioinformatics area. Probably you will be a member of a highly interdisciplinary team consisting of biologists, chemists, mathematicians, computer scientists ranging from programmers to algorithm engineers, and eventually people from various further fields. A major problem within such interdisciplinary teams is always to find some common language, and, for each member of some discipline, to have profound knowledge of what are the notions, basic concepts and goals of the other participating disciplines, as well as of what they can contribute to the solution of their own problems. This does, of course, not mean that a computer scientist should do the job of the biologist. Nevertheless, a computer scientist should be able to understand what a biologist deals with. On the other hand, the biologist should not do the computer scientist's job, but should know what computer science and algorithm engineering might contribute to the solution of her/his problems, and also how problems should be stated in order for the computer scientist to understand them. This book primarily aims to show the potential that algorithm engineering offers for the solution of core bioinformatics problems.

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