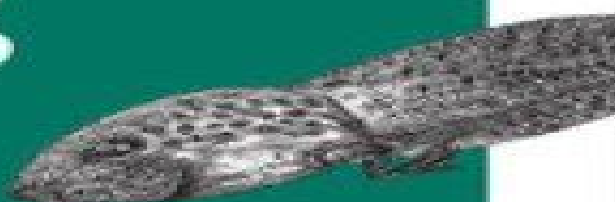


*An Introduction to Perl for Biologists*



# Beginning Perl for Bioinformatics

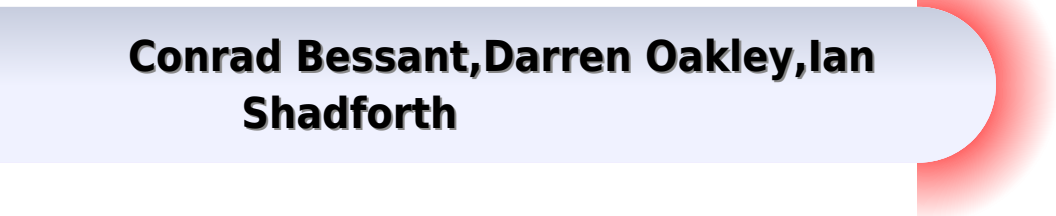


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# Beginning Perl For Bioinformatics

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**Mastering Perl for Bioinformatics** James Tisdall, 2003-09-25 Historically programming hasn t been considered a critical skill for biologists But now with access to vast amounts of biological data contained in public databases programming skills are increasingly in strong demand in biology research and development Perl with its highly developed capacities in string handling text processing networking and rapid prototyping has emerged as the programming language of choice for biological data analysis Mastering Perl for Bioinformatics covers the core Perl language and many of its module extensions presenting them in the context of biological data and problems of pressing interest to the biological community This book along with Beginning Perl for Bioinformatics forms a basic course in Perl programming This second volume finishes the basic Perl tutorial material references complex data structures object oriented programming use of modules all presented in a biological context and presents some advanced topics of considerable interest in bioinformatics The range of topics covered in Mastering Perl for Bioinformatics prepares the reader for enduring and emerging developments in critical areas of bioinformatics programming such as Gene finding String alignment Methods of data storage and retrieval SML and databases Modeling of networks graphs and Petri nets Graphics Tk Parallelization Interfacing with other programming languages Statistics PDL Protein structure determination Biological models of computation DNA Computers Biologists and computer scientists who have conquered the basics of Perl and are ready to move even further in their mastery of this versatile language will appreciate the author s well balanced approach to applying Perl s analytical abilities to the field of bioinformatics Full of practical examples and real world biological problem solving this book is a must for any reader wanting to move beyond beginner level Perl in bioinformatics      *Perl for Bio Informatics I* David Scott, 2013-01-07 Go from scripts to functions to programs Learn bio programming with the Perl development paradigm Learn to write tested documented object oriented code the Perl way      [Handbook of Hidden Markov Models in Bioinformatics](#) Martin Gollery, 2008-06-12 Demonstrating that many useful resources such as databases can benefit most bioinformatics projects the Handbook of

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*XML for Bioinformatics* Ethan Cerami,2006-06-02 Introduction The goal of this book is to introduce XML to a bioinformatics audience It does so by introducing the fundamentals of XML Document Type Definitions DTDs XML Namespaces XML Schema and XML parsing and illustrating these concepts with specific bioinformatics case studies The book does not assume any previous knowledge of XML and is geared toward those who want a solid introduction to fundamental XML concepts The book is divided into nine

chapters Chapter 1 Introduction to XML for Bioinformatics This chapter provides an introduction to XML and describes the use of XML in biological data exchange A bird s eye view of our rst case study the Distributed Annotation System DAS is provided and we examine a sample DAS XML document The chapter concludes with a discussion of the pros and cons of using XML in bioinformatic applications Chapter 2 Fundamentals of XML and BSML This chapter introduces the fundamental concepts of XML and the Bioinformatic Sequence Markup Language BSML We explore the origins of XML de ne basic rules for XML document structure and introduce XML Na spaces We also explore several sample BSML documents and visualize these documents in the TM Rescentris Genomic Workspace Viewer

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