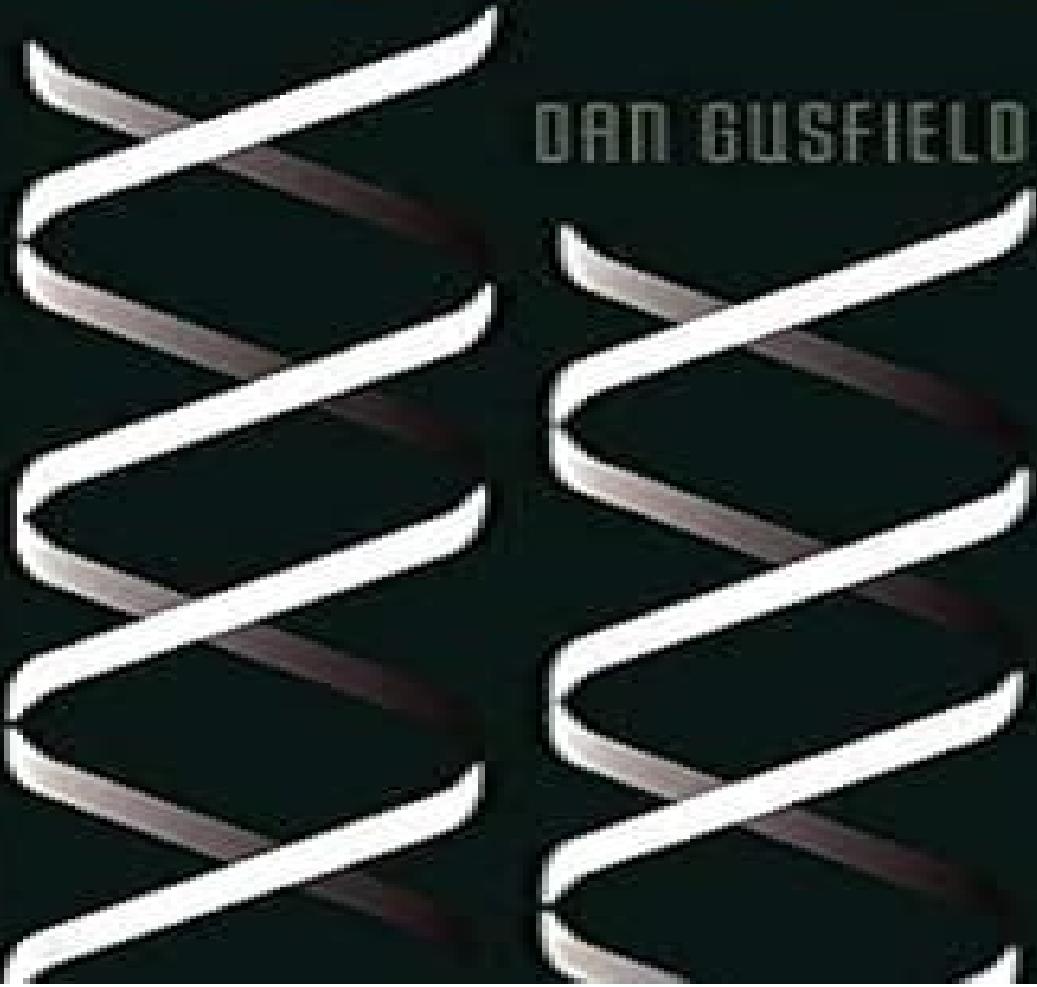




ALGORITHMS ON STRINGS, TREES, AND SEQUENCES

*Computer Science and
Computational Biology*

DAN GUSFIELD



Algorithms On Strings Trees And Sequences Computer Science And

Ming-Yang Kao

Algorithms On Strings Trees And Sequences Computer Science And:

Algorithms on Strings, Trees, and Sequences Dan Gusfield,1997-05-28 String algorithms are a traditional area of study in computer science In recent years their importance has grown dramatically with the huge increase of electronically stored text and of molecular sequence data DNA or protein sequences produced by various genome projects This book is a general text on computer algorithms for string processing In addition to pure computer science the book contains extensive discussions on biological problems that are cast as string problems and on methods developed to solve them It emphasises the fundamental ideas and techniques central to today s applications New approaches to this complex material simplify methods that up to now have been for the specialist alone With over 400 exercises to reinforce the material and develop additional topics the book is suitable as a text for graduate or advanced undergraduate students in computer science computational biology or bio informatics Its discussion of current algorithms and techniques also makes it a reference for professionals

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Theoretical Computer Science Mario Coppo,Elena Lodi,2005-09-28 This book constitutes the refereed proceedings of the 9th International Conference on Theoretical Computer Science ICTCS 2005 held at the Certosa di Pontignano Siena Italy in October 2005 The 29 revised full papers presented together with an invited paper and abstracts of 2 invited talks were carefully reviewed and selected from 83 submissions The papers address all current issues in theoretical computer science and focus especially on analysis and design of algorithms computability computational complexity cryptography formal languages and automata foundations of programming languages and program analysis natural computing paradigms quantum computing bioinformatics program specification and verification term rewriting theory of logical design and layout type theory security and symbolic and algebraic computation **Encyclopedia of Algorithms** Ming-Yang Kao,2008-08-06 One of Springer s renowned Major Reference Works this awesome achievement provides a comprehensive set of solutions to

important algorithmic problems for students and researchers interested in quickly locating useful information This first edition of the reference focuses on high impact solutions from the most recent decade while later editions will widen the scope of the work All entries have been written by experts while links to Internet sites that outline their research work are provided The entries have all been peer reviewed This defining reference is published both in print and on line

Algorithms and Theory of Computation Handbook, Volume 1 Mikhail J. Atallah,Marina Blanton,2009-11-20

Algorithms and Theory of Computation Handbook Second Edition General Concepts and Techniques provides an up to date compendium of fundamental computer science topics and techniques It also illustrates how the topics and techniques come together to deliver efficient solutions to important practical problems Along with updating and revising many Advances in Computers Marvin Zelkowitz,2009-05-11 This is volume 75 of Advances in Computers This series which began publication in 1960 is the oldest continuously published anthology that chronicles the ever changing information technology field In these volumes we publish from 5 to 7 chapters three times per year that cover the latest changes to the design development use and implications of computer technology on society today In this present volume we present five chapters describing new technology affecting users of such machines In this volume we continue a theme presented last year in volume 72 High Performance Computing In volume 72 we described several research projects being conducted in the United States on the development of a new generation of high performance supercomputers **Handbook of Computational Molecular Biology** Srinivas Aluru,2005-12-21 The enormous complexity of biological systems at the molecular level must be answered with powerful computational methods Computational biology is a young field but has seen rapid growth and advancement over the past few decades Surveying the progress made in this multidisciplinary field the Handbook of Computational Molecular Biology of Proceedings of the Seventeenth Annual ACM-SIAM Symposium on Discrete Algorithms SIAM Activity Group on Discrete Mathematics,Association for Computing Machinery,Society for Industrial and Applied Mathematics,2006-01-01 Symposium held in Miami Florida January 22 24 2006 This symposium is jointly sponsored by the ACM Special Interest Group on Algorithms and Computation Theory and the SIAM Activity Group on Discrete Mathematics Contents Preface Acknowledgments Session 1A Confronting Hardness Using a Hybrid Approach Virginia Vassilevska Ryan Williams and Shan Leung Maverick Woo A New Approach to Proving Upper Bounds for MAX 2 SAT Arist Kojevnikov and Alexander S Kulikov Measure and Conquer A Simple O 20 288n Independent Set Algorithm Fedor V Fomin Fabrizio Grandoni and Dieter Kratsch A Polynomial Algorithm to Find an Independent Set of Maximum Weight in a Fork Free Graph Vadim V Lozin and Martin Milanic The Knuth Yao Quadrangle Inequality Speedup is a Consequence of Total Monotonicity Wolfgang W Bein Mordecai J Golin Larry L Larmore and Yan Zhang Session 1B Local Versus Global Properties of Metric Spaces Sanjeev Arora L szl Lov sz Ilan Newman Yuval Rabani Yuri Rabinovich and Santosh Vempala Directed Metrics and Directed Graph Partitioning Problems Moses Charikar Konstantin Makarychev and Yury Makarychev Improved Embeddings of Graph Metrics

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