

Handapparat_AG-ABI, Sonderstandort Takustr. 9, Informatik (Präsenzbestand)

Autor (Name, Vorname)	Titel	Verlag: Ort, Jahr (Ausgabe/Auflage)	Signatur
Altmann / Dunker / Hunter / Murray / Klein (Eds.)	Pacific Symposium on Biocomputing 2006	(Proceedings) World Scientific: New Jersey et al., 2006 06	Mau 1 a / J 2.0
Austern, Matthew H.	Generic Programming and the STL	Addison-Wesley: Boston et al., 1999	Aus.M 1a / XD.3.2
Balzert, Heide	Lehrbuch der Objektmodellierung [+ CD-ROM]	Spektrum Akademischer Verlag: Heidelberg, Berlin, 1999	Balz.H 1c / XD.2.0
Bevington, Philip R. / Robinson, D. Keith	Data Reduction and Error Analysis	McGraw-Hill: New York, 2003 (3)	Bev.P 1 a / XG.3
Bishop, Christopher M.	Pattern Recognition and Machine Learning	Springer: New York, 2006	Bi.C 2a / XG.3
Blanchette, Jasmin / Summerfield, Mark	C++ GUI Programming with Qt 4 [+ CD-ROM]	Trolltech AS/Prentice Hall: Upper Saddle River et al., 2006	Bln 1 a / XD.3.2
Bono, Edward de	Six Thinking Hats	Penguin: London et al., 2000. (© 1985, 1999)	Bon.E 5c / A 3.0
Bosch, Karl	Elementare Einführung in die angewandte Statistik	vieweg studium: Braunschweig, Wiesbaden, 2000 (7)	Bos 4h / H 3.0
Carnegie, Dale	The Quick & Easy Way to Effective Speaking	Pocket Books: New York, 1977 (© 1962)	Crn 1 d / A 3.0
Chao, Kun-Mao / Zhang, Louxin	Sequence Comparison. Theory and Methods	Springer: London, 2009	Chao.K 1 a / XE.1
Cristianini, Nello / Hahn, Matthew W.	Introduction to Computational Genomics. A case studies approach	CUP: Cambridge et al., 2007	Cri.N 2 a / XJ.3
Cristianini, Nello / Shawe-Taylor, John	An Introduction to Support Vector Machines	CUP: Cambridge et al., 2003	Cri.N 1b / XF.1.1
Davies, Kevin	The \$ 1,000 Genome. The Revolution in DNA Sequencing and the New Era of Personalized Medicine	Free Press: New York/NY, 2010	Davi.K 1 a / XA.1
Deonier, Richard C. / Tavaré, Somon / Waterman, Michael S.	Computational Genome Analysis. An Introduction	Springer, 2005	Deo.R 1 a / XJ.3
Dümbgen, Lutz	Stochastik für Informatiker	Springer: Berlin et al., 2003	Dünn 2a / XG.3
Durbin, Richard / Eddy, Sean / Krogh, Anders / Mitchison, Graeme	Biological sequence analysis. Probabilistic models of proteins and nucleic acids	CUP: Cambridge et al., 2009 (13)	XL.0 1b/ XL.0
Eckel, Bruce	Thinking in C++	Prentice Hall: New Jersey, 2000 (2)	Ec.B 2a / XD.3.2
Eidhammer, Ingvar / Flikka, Kristian / Martens, Lennart / Mikalsen, Svein-Ole	Computational Methods for Mass Spectrometry Proteomics	John Wiley & Sons: Chichester, West Sussex, 2007	WC 2600 / E34
Eidhammer, Ingvar / Jonassen, Inge / Taylor, William R.	Protein Bioinformatics. An Algorithmic Approach to Sequence and Structure Analysis	John Wiley & Sons: Chichester, West Sussex, 2004	Eid.I 1a / XJ.3

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Evjen, Bill / Hanselmann, Scott / Muhammad, Farhan, Sivakumar, Srinivasa / Rader, Devin	Professional ASP.NET 2.0	Wiley: Indianapolis/IN, 2006	XD.3.2 80 a / XD.3.2
Ewens, Warren J. / Grant, Gregory R.	Statistical Methods in Bioinformatics. An Introduction	Springer: New York, Berlin, Heidelberg, 2001	Ewen 2b / H 13.0
Ewens, Warren J. / Grant, Gregory R.	Statistical Methods in Bioinformatics. An Introduction. 2nd Edition	Springer: New York, 2005 (2)	Ewen 2c / H 13.0
Feibelman, Peter J.	A Ph.D. Is Not Enough - VERLUST -	New York, 1993	Fei.P 1a / XK.m
Gogol-Döring, Andreas / Reinert, Knut	Biological Sequence Analysis Using the SeqAn C++ Library	CRC Press: London/UK, Boca Raton/USA, 2010	Gog.A 1 a / XE.1
Gonick, Larry / Criddle, Craig	The Cartoon Guide to Chemistry	HarperCollins: New York, 2005	Gon.L 2 a / XA.1
Gonick, Larry / Smith, Woollcott	The Cartoon Guide to Statistics	HarperCollins: New York, 2005 (© 1993)	Gon.L 1a / XG.3
Gonick, Larry / Wheelis, Mark	The Cartoon Guide to Genetics	HarperCollins: New York et al., 2005 (© 1983)	Gon.L 3 a / XA.1
Goossens, Michel / Mittelbach, Frank / Samerin, Alexander	Der Latex-Begleiter	Addison-Wesley, 2000 / Pearson Studium, 2002	Goo.M 2d / XI.7.2
Griffith, Arthur	GCC: The Complete Reference	McGraw-Hill/Osborne: New York et al., 2002	Grif.A 1a / XD.3.4
Hand, David / Mannila, Heikki / Smyth, Padhraic	Principles of Data Mining	MIT Press: Cambridge et al., 2001	Hand 3a / XH.2.8
Hastie, Trevor / Tibshirani, Robert / Friedman, Jerome	The Elements of Statistical Learning	Springer: New York et al., 2001	Has.T 2a / XG.3
Haubold, Bernhard / Wiehe, Thomas	Introduction to Computational Biology. An Evolutionary Approach	Birkhäuser: Basel, Boston, Berlin, 2006	Hau.B 1 a / XJ.3
Higham, Nicholas J.	Handbook of Writing for the Mathematical Sciences	Siam: Philadelphia, 1998 (2)	Higha 1b / A 1.0
Hunt, Andrew / Thomas, David	The Pragmatic Programmer	Addison-Wesley: Boston et al., 1999	Hunt.A 1 a / XD.2.3
Hütt, Marc-Thorsten / Dehnert, Manuel	Methoden der Bioinformatik	Springer: Berlin, Heidelberg, 2006	Hütt 1a / XJ.3
Jones, Neil C / Pevzner, Pavel	An Introduction to Bioinformatics Algorithms	The MIT Press: Cambridge/Mass, London/UK, 2004	Jone.N 1a / XF.2.0
Josuttis, Nicolai M.	Object-Oriented Programming in C++	John Wiley & Sons, Ltd: Chichester/UK, 2003	Jolsu 2 a / XD.3.2

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Josuttis, Nicolai M.	The C++ Standard Library. A Tutorial Reference	Addison-Wesley: Boston et al., 2002 (10)	Jos.N 1a / XD.3.2
Jurisica, Igor / Wigle, Dennis	Knowledge Discovery in Proteomics	CRC Press: London/UK, Florida/USA, 2006	Jur 1 a / XJ.3
Koegel, Timothy J.	The Exceptional Presenter	Greenleaf Book Group Press: Austin/Texas, 2007	Koeg 1 b / A 3.0
Kopka, Helmut	Latex. Band 1: Einführung	Addison-Wesley, 2000 / Pearson Studium, 2002 (3)	Ko.H 3.1 I / XI.7.2
Lengauer, Thomas (Ed.)	Bioinformatics - From Genomes to Therapies. Vol. 1: Molecular Sequences and Structures	Wiley-VCH: Weinheim, 2007	XJ.3 7.1a / XJ.3
Lengauer, Thomas (Ed.)	Bioinformatics - From Genomes to Therapies. Vol. 2: Molecular Interactions	Wiley-VCH: Weinheim, 2007	XJ.3 7.2a / XJ.3
Lengauer, Thomas (Ed.)	Bioinformatics - From Genomes to Therapies. Vol. 3: Molecular Function	Wiley-VCH: Weinheim, 2007	XJ.3 7.3a / XJ.3
Lippman, Stanley B.	Essential C++	Addison-Wesley: Reading, Massachusetts et al., 2000	Lipp 2a / XD.3.2
Liu, Jun S.	Monte Carlo Strategies in Scientific Computing	Springer: New York, 2004	Lin.J 1 b / H 11.2
Lund, Ole / Nielsen, Morten / Lundegaard, Claus / Kesmir, Can / Brunak, Soren	Immunological Bioinformatics	MIT Press: Cambridge et al., 2005	XJ.3 5a / XJ.3
Mäkinen, Veli / Belazzougui, Djamel / Cunial, Fabio / Tomescu, Alexandru I.	Genome-Scale Algorithm Design. Biological Sequence Analysis in the Era of High-Throughput Sequencing	CUP: Cambridge/UK, 2015	
Manly, Bryan F. J.	Multivariate Statistical Methods: A primer	Chapman & Hall/CRC: Boca Raton, 2005 (3)	Maul 3a / XG.3
Manly, Bryan F. J.	Randomization, Bootstrap and Monte Carlo Methods in Biology	Chapman & Hall/CRC: Boca Raton, 2007 (3)	Maul 2b / H 3.0
McGovern, James / Tyagi, Sameer / Stevens, Michael E. / Mathew, Sunil	Java Web Services Architecture [+ CD-ROM]	Morgan Kaufmann Publishers: San Francisco, 2003	XC.2.4 30a / XC.2.4
Meffert, Beate / Hochmuth, Olaf	Werkzeuge der Signalverarbeitung	Pearson Studium: München et al., 2004	Mef 1a / XI 5.4
Mehlhorn, Kurt / Sanders, Peter	Algorithms and Data Structures. The Basic Toolbox	Springer: Heidelberg, 2008	Meh 4a / XF.2.0
Meyers, Scott	Effective STL -VERLUST-	Addison-Wesley: Boston et al., 2001 (3)	Mey.S 2a / XC.3.2
Meyers, Scott	Effective STL	Addison-Wesley: Boston et al., 2001 (3)	Mey.S 2b / XD.3.2
Motwani, Rajeev / Raghavan, Prabhakar	Randomized Algorithms	CUP: New York et al., 2007 (9)	Mot.R 1 e / XF.2.2

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Myers, Glenford J. / Badgett, Tom / Thomas, Todd / Sandler, Corey	The Art of Software Testing	John Wiley & Sons: Hoboken/NJ, 2004 (2)	Myer 2c / XD.2.5
Navarro, Gonzalo	Compact Data Structures. A Practical Approach	CUP: New York/NY, 2016	ST 230 / N322
Ohlebusch, Enno	Bioinformatics Algorithms. Sequence Analysis, Genome Rearrangements, and Phylogenetic Reconstruction	E. Ohlebusch, 2013	ST 134 / O37
Parida, Laxmi	Pattern Discovery in Bioinformatics. Theory & Algorithms	Chapman & Hall/CRC: Boca Raton, 2008	Pari 1a / XI.5.0
Piantadosi, Steven	Clinical Trials. A Methodologic Perspective	John Wiley & Sons: Hoboken/NJ, 2005 (2)	Pia 1a / XG.3
Polanski, Andrzej / Kimmel, Marek	Bioinformatics	Springer: Berlin, Heidelberg, 2007	Pola.A 1 a / XJ.3
Postman, Neil / Weingartner, Charles	Teaching as a Subversive Activity	Delta Book, Dell Publishing Co., Inc.: New York/NY, 1969	Post.N 1 a / XA.m
Sagan, Carl	The Demon-Haunted World. Science as a Candle in the Dark	Ballantine Books: New York, 1997 (© 1996)	Sag.C 1 b / A 3.0
Schmidt, Bertil (Ed.)	Bioinformatics. High Performance Parallel Computer Architectures	Taylor & Francis: Boca Raton et al., 2011	XB.2.4 / 1a
Schölkopf, Bernhard / Smola, Alexander J.	Learning with Kernels	The MIT Press: Cambridge/Mass., London/UK, 2002	Schö.B 1a / XI.2.0
Semple, Charles / Steel, Mike	Phylogenetics	OUP: Oxford, New York, 2003	Se.C 1a / XJ.3
Siek, Jeremy G. / Lee, Lie-Quan / Lumsdaine, Andrew	The Boost Graph Library	Addison-Wesley: Boston et al., 2002	Sie.K 1a / XD.2.2
Smith, R. Martin	Understanding Mass Spectra. A Basic Approach	John Wiley & Sons: Hoboken/NJ, 2004 (2)	Smt.R 1 1 / XJ.3
Srivastava, Sudhir (Ed.)	Informatics in Proteomics	Taylor & Francis: Boca Raton et al., 2005	XJ.3 6a / XJ.3
Srivastava, Sudhir (Ed.)	Informatics in Proteomics	Taylor & Francis: Boca Raton et al., 2005	XJ.3 6b / XJ.3
Stephen, Graham A.	String Searching Algorithms	World Scientific: Singapore et al., 1994	Steph 1b / XF.2.2
Stroustrup, Bjarne	The C++ Programming Language	Addison-Wesley: Boston et al., 2001 (3)	St.B 1e / XD.3.2
Sung, Wing-Kin	Algorithms in Bioinformatics. A practical introduction	Taylor & Francis: Boca Raton et al., 2010	Winki 1 a / XF.2.0
Tramontano, Anna	The Ten Most Wanted Solutions in Protein Bioinformatics	Taylor & Francis: Boca Raton et al., 2005	Tram 1 a / XJ.3
Vandevoorde, David / Josuttis, Nicolai M.	C++ Templates. The Complete Guide	Addison-Wesley: Boston et al., 2003	Van.D 1a / XD.3.2

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Weiss, Mark Allen	Data Structures & Algorithm Analysis in C++	Addison-Wesley Longman, 1999 (2)	Weis.M 2a / XE.1
Wickham, Hadley	ggplot2. Elegant Graphics for Data Analysis	Springer Science+Business Media, 2009	Wic.H 1a / XG.3
Williams, David	Weighing the Odds, A Course in Probability and Statistics	CUP: Cambridge et al., 2001	Will.D 3 b / H 2.0
Zimmermann, Karl-Heinz	An Introduction to Protein Informatics	Kluwer Academic Publishers: Boston, Dordrecht, London, 2003	Zimm 1a / XJ.3
Zinsser, William	Writing to Learn	Harper: New York et al., 1989 (© 1988)	Zins.W 2 c / A 3.0
Zvelebil, Marketa / Baum, Jeremy O.	Understanding Bioinformatics	Garland Science, Taylor & Francis: New York/USA, Abingdon/UK, 2008	Zve 1e / XJ.3
Zvelebil, Marketa / Baum, Jeremy O.	Understanding Bioinformatics	Garland Science, Taylor & Francis: New York/USA, Abingdon/UK, 2008	Zve 1f / XJ.3