

1. Record Nr.	UNISA996465989803316
Titolo	Applications of Evolutionary Computation [[electronic resource] ] : EvoApplications 2012: EvoCOMNET, EvoCOMPLEX, EvoFIN, EvoGAMES, EvoHOT, EvoASP, EvoNUM, EvoPAR, EvoRISK, EvoSTIM, and EvoSTOC, Málaga, Spain, April 11-13, 2012, Proceedings // edited by Cecilia Di Chio, Alexandros Agapitos, Stefano Cagnoni, Carlos Cotta, Francisco Fernández de Vega, Gianni A. Di Caro, Rolf Drechsler, Anikó Ekárt, Anna I. Esparcia-Alcázar, Muddassar Farooq, W.B. Langdon, Juan-J. Merelo-Guervós, Mike Preuss, Hendrik Richter, Sara Silva, Anabela Simões, Giovanni Squillero, Ernesto Tarantino, Andrea Tettamanzi, Julian Togelius, Neil Urquhart, Sima Uyar, Georgios N. Yannakakis
Pubbl/distr/stampa	Berlin, Heidelberg : , : Springer Berlin Heidelberg : , : Imprint : Springer, , 2012
ISBN	3-642-29178-3
Edizione	[1st ed. 2012.]
Descrizione fisica	1 online resource (xxv, 550 pages) : illustrations
Collana	Theoretical Computer Science and General Issues, , 2512-2029 ; ; 7248
Disciplina	004.0151
Soggetti	Computer science Computer programming Computer networks Computer science—Mathematics Computer vision Artificial intelligence Theory of Computation Programming Techniques Computer Communication Networks Mathematical Applications in Computer Science Computer Vision Artificial Intelligence
Lingua di pubblicazione	Inglese
Formato	Materiale a stampa
Livello bibliografico	Monografia
Note generali	International conference proceedings. Editors: Cecilia Di Chio, Alexandros Agapitos, Stefano Cagnoni, Carlos Cotta, Francisco Fernández de Vega, Gianni A. Di Caro, Rolf Drechsler, Anikó Ekárt, Anna I. Esparcia-Alcázar, Muddassar Farooq, W.B. Langdon, Juan-J. Merelo-Guervós, Mike Preuss, Hendrik Richter, Sara

Silva, Anabela Simões, Giovanni Squillero, Ernesto Tarantino, Andrea Tettamanzi, Julian Togelius, Neil Urquhart, Sima Uyar, Georgios N. Yannakakis.

Nota di bibliografia

Includes bibliographical references and index.

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This book constitutes the refereed proceedings of the International Conference on the Applications of Evolutionary Computation, EvoApplications 2012, held in Málaga, Spain, in April 2012, colocated with the Evo\* 2012 events EuroGP, EvoCOP, EvoBIO, and EvoMUSART. The 54 revised full papers presented were carefully reviewed and selected from 90 submissions. EvoApplications 2012 consisted of the following 11 tracks: EvoCOMNET (nature-inspired techniques for telecommunication networks and other parallel and distributed systems), EvoCOMPLEX (algorithms and complex systems), EvoFIN (evolutionary and natural computation in finance and economics), EvoGAMES (bio-inspired algorithms in games), EvoHOT (bio-inspired heuristics for design automation), EvoIASP (evolutionary computation in image analysis and signal processing), EvoNUM (bio-inspired algorithms for continuous parameter optimization), EvoPAR (parallel implementation of evolutionary algorithms), EvoRISK (computational intelligence for risk management, security and defense applications), EvoSTIM (nature-inspired techniques in scheduling, planning, and timetabling), and EvoSTOC (evolutionary algorithms in stochastic and dynamic environments).

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2. Record Nr.	UNINA9910817935303321
Titolo	Addiction research methods // edited by Peter G. Miller, John Strang, Peter M. Miller
Pubbl/distr/stampa	Chichester, West Sussex, U.K. ; ; Ames, Iowa, : Wiley-Blackwell/Addiction Press, 2010
ISBN	9786612482427 9781282482425 1282482424 9781444318852 1444318853 9781444318869 1444318861
Descrizione fisica	1 online resource [402 pages]
Altri autori (Persone)	MillerPeter (Peter Graeme) StrangJohn MillerPeter M <1942-> (Peter Michael)
Disciplina	616.86/027
Soggetti	Substance abuse - Research - Methodology Psychiatry - Research - Methodology
Lingua di pubblicazione	Inglese
Formato	Materiale a stampa
Livello bibliografico	Monografia
Nota di bibliografia	Includes bibliographical references and index.
Sommario/riassunto	Addiction Research Methods' is a comprehensive handbook for health professionals, policy-makers and researchers working and training in the field of addiction. The book provides a clear, comprehensive and practical guide to research design, methods and analysis within the context of the field of alcohol and other drugs. The reader is introduced to fundamental principles and key issues; and is orientated to available sources of information and key literature. Written by a team of internationally acclaimed contributors, the book is divided into six major sections: Introduction; Research Design; Basic Toolbox; Biological Models; Specialist Methods; and Analytical Methods. Each chapter offers an introduction to the background and development of

the discipline in question, its key features and applications, how it compares to other methods/analyses and its advantages and limitations. FEATURES \* List of useful websites and assistive technology. \* Case study examples \* List of useful hermeneutics \* Recommended reading list \* Contains exercises to help the reader to develop their skills.

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