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Altri autori (Persone)	BrayR.A GibsonD.I
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Blair, 1979; 35 Family Notocotylidae Luhe, 1909; 36 Family Nudacotylidae Barker, 1916; 37 Family Opisthotrematidae Poche, 1926; 38 Family Rhabdiopoeidae Poche, 1926; 39 Superfamily Allocreadioidea Looss, 1902; 40 Family Allocreadiidae Looss, 1902
41 Family Batrachotrematidae Dollfus & Williams, 196642 Family Opecoelidae Ozaki, 1925; 43 Family Opistholebetidae Fukui, 1929; 44 Superfamily Lepocreadioidea Odhner, 1905; 45 Family Lepocreadiidae Odhner, 1905; 46 Family Acanthocolpidae Luhe, 1906; 47 Family Apocreadiidae Skrjabin, 1942; 48 Family Brachycladiidae Odhner, 1905; 49 Family Deropristidae Cable & Hunninen, 1942; 50 Family Enenteridae Yamaguti, 1958; 51 Family Gorgocephalidae Manter, 1966; 52 Family Gyliauchenidae Fukui, 1929; 53 Family Liliatrematidae Gubanov, 1953; 54 Family Megaperidae Manter, 1934; Bibliography; Index; A; B; C
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Sommario/riassunto

This is the second of three volumes of Keys to the Trematoda, a series on the systematics and identification of the Class Trematoda. The book presents the taxa in the Order Echinostomida and some of those in the Order Plagiorchiida, with keys for their identification at the superfamily, family, subfamily and generic levels. The keys are based on critical examination of specimens by subject experts, and generic diagnoses are accompanied by illustrations of important morphological characters. This volume includes seven echinostomidan superfamilies (the echinostomatoids, haploporoids, haplosplanc

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Collana	Theoretical Computer Science and General Issues, , 2512-2029 ; ; 8602
Disciplina	005.1
Soggetti	Algorithms Artificial intelligence Pattern recognition systems Computer networks Computer science Application software Artificial Intelligence Automated Pattern Recognition Computer Communication Networks Theory of Computation Computer and Information Systems Applications
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Nota di contenuto	Nature-inspired techniques for telecommunication networks and other parallel and distributed systems -- Evolutionary algorithms and complex systems -- Evolutionary computation in energy applications -- Evolutionary and natural computation in finance and economics -- Bio-inspired algorithms in games -- Evolutionary computation in image analysis, signal processing, and pattern recognition -- Nature-inspired techniques in industrial settings -- Bio-inspired algorithms for continuous parameter optimization -- Parallel implementation of evolutionary algorithms -- Computational intelligence for risk management, security and defence applications -- Evolutionary computation in robotics -- Evolutionary algorithms in stochastic and

dynamic environments -- EC and related techniques in bioinformatics and computational biology.

Sommario/riassunto

This book constitutes the thoroughly refereed post-conference proceedings of the International Conference on the Applications of Evolutionary Computation, EvoApplications 2014, held in Granada, Spain, in April 2014, colocated with the Evo* 2014 events EuroGP, EvoCOP, and EvoMUSART. The 79 revised full papers presented were carefully reviewed and selected from 128 submissions. EvoApplications 2014 consisted of the following 13 tracks: EvoCOMNET (nature-inspired techniques for telecommunication networks and other parallel and distributed systems), EvoCOMPLEX (evolutionary algorithms and complex systems), EvoENERGY (evolutionary computation in energy applications), EvoFIN (evolutionary and natural computation in finance and economics), EvoGAMES (bio-inspired algorithms in games), EvoIASP (evolutionary computation in image analysis, signal processing, and pattern recognition), EvoINDUSTRY (nature-inspired techniques in industrial settings), EvoNUM (bio-inspired algorithms for continuous parameter optimization), EvoPAR (parallel implementation of evolutionary algorithms), EvoRISK (computational intelligence for risk management, security and defence applications), EvoROBOT (evolutionary computation in robotics), EvoSTOC (evolutionary algorithms in stochastic and dynamic environments), and EvoBio (EC and related techniques in bioinformatics and computational biology).