

Melentiev Energy Systems Institute SB RAS
Russian Foundation for Basic Research
Laboratory of Algorithms and Technologies for Networks Analysis,
Higher School of Economics, Nizhny Novgorod

XVII Baikal International School-Seminar
METHODS OF OPTIMIZATION
AND THEIR APPLICATIONS

Abstracts

July 31 – August 6, 2017
Maksimikha, Buryatia

Irkutsk
2017

Numerical study of bioinspired methods for solving global optimization problems

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Keywords: global optimization, bioinspired optimization methods, test problem, meta-heuristic approach.

Bioinspired optimization methods performs models representing simplified analogues of some biological processes (population evolution, collective behavior of self-organizing agents, etc.). They refer to the so-called meta-heuristic approaches, which do not guarantee the global extrema finding, but often allows one to find a good approximation in a rather short time. In fact, meta-heuristics describe a set of rules for the search process implementation (mostly a stochastic process). It aims to find near-optimal solution by the target function. In recent practice, large-scale problems with an “expansive” target functions are widely solved with the use of hybrids and multimethod schemes based on both bioinspired meta-heuristics and deterministic, in particular, gradient algorithms. Thus it becomes important to have some information about different meta-heuristics effectiveness and any preliminary assessment of their capabilities [1, 2, 3]. The work deals with the construction of an empirical rating, we perform a numerical study of more than 20 different modern bioinspired methods. For results of the research to be representative, we have defined a unified set of tests (synthetic and applied problems, dimensions of 100 or more variables). For each problems we generated a set of 100 starting points, each studied method runs on these sets with the same limit of an objective function computations. The main result of the work is a simple statistics (average, standard deviation, minimum and maximum value of the function, real time) of each method work on 100 starts for the whole set of the tests. Every considered method possessing its own parameters was searched for their possible values (using fixed step on a uniform grid). Thus, the results obtained are close to optimal in the sense of the methods parameters sets.

Acknowledgments. This work is partly supported by the Russian Foundation for Basic Research (project N 17-07-00627).

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