

# Editorial: application of optimization algorithms in chemistry

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Editorial on the Research Topic

### [Application of Optimization Algorithms in Chemistry](#)

Molecular structure optimization, fitting potential energy functions to *ab initio* and experimental data, and spectral assignment are among the hardest optimization tasks in molecular sciences. These are fundamental problems in chemistry, but they can also be relevant in molecular physics and biochemistry. In past decades, several methodologies have been proposed to help in the above mentioned tasks, and some of them are already incorporated into computational tools, such as GMIN ( [Wales and Scheraga, 1999](#) ; [Wales, 2010](#) ), Gradient Embedded Genetic Algorithm or GEGA ( [Alexandrova and Boldyrev, 2005](#) ), OGOLEM ( [Hartke, 1993](#) ; [Dieterich and Hartke, 2017](#) ), Birmingham Cluster Genetic Algorithm or BCGA ( [Johnston, 2003](#) ; [Shayeghi et al., 2015](#) ), Evolutionary Algorithm for Molecular Clusters or EA\_MOL ( [Llanio-Trujillo et al., 2011](#) ; [Marques and Pereira, 2011](#) ), Global Reaction Route Mapping or GRRM ( [Ohno and Maeda, 2006](#) , [2019](#) ), Automated Mechanisms and Kinetics or AutoMeKin ( [Martínez-Núñez, 2015a](#) , [b](#) ; [Martínez-Núñez, 2020](#) ), and Genetic Algorithm fitting or GAFit ( [Rodríguez-Fernández et al., 2017](#) , [2020](#) ). Most of these computational programs are interfaced with well-known packages that perform electronic-structure calculations and, hence, allow for a direct assessment of the semi-empirical, density functional theory (DFT) or *ab initio* energy of the system during the optimization process. Another relevant methodology to explore low-energy landscapes is the parallel-tempering Monte Carlo technique, which has been also applied in the calculation of thermodynamic properties.

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Global geometry optimization studies are, now, being extended to systems of increasing complexity. In particular, global optimization algorithms have been applied to a great diversity of chemical systems, including atomic and molecular clusters as well as colloidal aggregates and biomolecules.

Nonetheless, optimization work needs, in general, a large number of computational resources and, hence, improvements in algorithms to relieve the burden. Major challenges are concerned with the treatment of systems with increasing size and incorporating higher levels of theory in the molecular model. Also, multi-component aggregates pose an important combinatorial problem and require novel optimization strategies. Although the use of state-of-the-art spectroscopic techniques to probe the structure of clusters has allowed for close collaborative work involving computational and experimental achievements, there is still room for greater improvement in this effort. In particular, comparisons between theoretical and experimental spectroscopic data will benefit from significant improvements in algorithms devoted for the spectral assignment.

Pursuing those purposes, we believe the collection of papers for the present Research Topic illustrates the broad scope of computational strategies for global optimization applications in chemistry. All contributions describe optimization strategies for a great diversity of chemical systems.

Basin-hopping (BH) is able to generate a coarse-grained mapping of a potential energy surface (PES) in terms of local minima, which can then be used to gain insights into molecular dynamics and thermodynamic properties as pointed out by [Zhou et al.](#) in their contribution. These authors also show

how unsupervised machine learning tools can be employed to enhance BH searches, which result in more efficient identification of local minima and transition states connecting them.

[Jana et al.](#) employ a particle swarm optimization (PSO) method to search for small  $C_n$  clusters. PSO is another useful algorithm for a stochastic search in multidimensional space. The method has proven efficient in hard optimization problems compared with traditional methods.

[Hernández-Rojas and Calvo](#) also employ BH method, this time to predict low-energy structures of adamantane clusters by using both coarse-grained and atomistic potential models. Although coarse-grained models are appealing for the complex clusters that are studied, the comparison with atomistic potentials shows that some relevant structural details are not captured by the former.

As for seeking conformational minima of flexible acyclic molecules, [Ferro-Costas and Fernández-Ramos](#) propose an algorithm that combines a systematic variation of torsion angles with a Monte Carlo search. This methodology has been applied to calculate multi-structural partition functions of several alcohols ranging from n-propanol to n-heptanol and was also tested with the amino acid L-serine.

[Panadés-Barrueta et al.](#) put forward a fully automated method to generate highly-accurate semiempirical potential energy surfaces. They use global optimization techniques and automated PES sampling algorithms to refine

specific reaction parameters of semi-empirical Hamiltonians, which can be subsequently employed in quantum dynamics studies.

In turn, [Wang et al.](#) carry out a microsolvation study of  $\text{Na}^+$  with water by applying a genetic algorithm combined with density functional theory to obtain low-energy structures of the clusters. Also, a new genetic algorithm is proposed by [Silva et al.](#) for the prediction of structures of nanoparticles. This work explores the efficacy of new evolutionary operators to treat Lennard-Jones and carbon clusters.

[Khatun et al.](#) develop a global optimizer which grows the cluster by adding atoms one by one. The method is tested by studying transition-metal clusters and binary and ternary nanoalloys of such elements.

[Cova and Pais](#) review and discuss deep learning strategies for optimizing the prediction of chemical patterns, which includes accelerated literature searches, analysis and prediction of physical and quantum chemical properties, transition states, chemical structures, chemical reactions, and also new catalysts and drug candidates.

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All authors listed have made a substantial, direct and intellectual contribution to the work, and approved it for publication.

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## Conflict of Interest

The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

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