Finding Many Stable Molecular Arrangements

Conformational Searching with Genetic Algorithms

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Outline

- 1. Background Information
- 2. The Genetic Algorithm
- 3. Conclusion

Background Information

Computational methods require knowledge of molecular structure

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We may need information about one or more low-energy conformations

 \Rightarrow Ok, let's find them all!

Possible Solutions

Many techniques are well established

method	description	implemented, e.g., in	
grid-based	based on grids of selected Cartesian or internal coordinates (e.g., grids of different torsional angle values of a molecule)	MOE ¹³	
rule/knowledge-based	use known (e.g., from experiments) structural preferences of compounds	ALFA, 14 CONFECT, 15 CORINA and ROTATE, 16,17 COSMOS, 18,19 OMEGA ²⁰	
population-based metaheuristic	improve candidate solutions in a guided search	Balloon, ²¹ Cyndi ²²	
distance geometry	based on a matrix with permitted distances between pairs of atoms	RDKit ²³	
basin-hopping ²⁴ / minima hopping ²⁵	based on moves across the PES combined with local relaxation	ASE, ²⁶ GMIN, ²⁷ TINKER SCAN ²⁸	
"Names of freely available programs are highlighted in boldface.			

Possible Solutions

Many techniques are well established None are perfect

method	description	implemented, e.g., in	
grid-based	based on grids of selected Cartesian or internal coordinates (e.g., grids of different torsional angle values of a molecule)	CAESAR, Open Babel, Confab, MacroModel, MoE ¹³	
rule/knowledge-based	use known (e.g., from experiments) structural preferences of compounds	ALFA, 14 CONFECT, 15 CORINA and ROTATE, 16,17 COSMOS, 18,19 OMEGA ²⁰	
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- · Parallel-Scalable

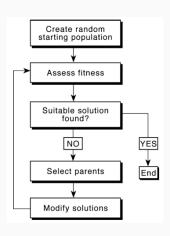
The Genetic Algorithm

Outline

- Inspired by biological evolution
- Evolve a population over generations
- · Survival of the fittest
- · Requirements:
 - Represent individuals as vector
 - · Fitness function

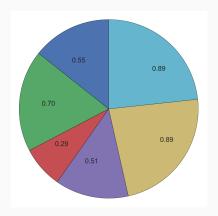
•
$$V = (x_1 y_1 z_1 x_2 y_2 z_2 ... x_N y_N z_N)$$

•
$$F = \frac{E_{max} - E}{E_{max} - E_{min}}$$



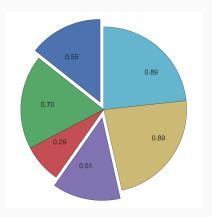
Selecting Parents

- Several methods are common
- Reinforce good characteristics
- · Still give losers a chance
- · 'Breed' pairs of winners

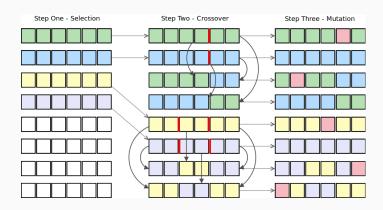


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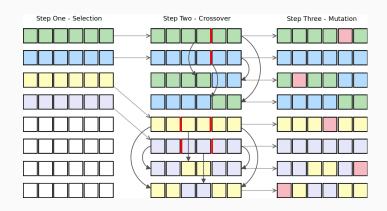
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The Next Generation



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Crossover distinguishes this from Monte Carlo

Conclusion



Backup slides

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