Review Article Open Access

# An Optimization Model of Molecular Voronoi Cells in Computational Chemistry

## Jiapu Zhang<sup>1,2\*</sup>

<sup>1</sup>Molecular Model Discovery Laboratory, Department of Chemistry and Biotechnology, Faculty of Science, Engineering and Technology, Swinburne University of Technology, Hawthorn Campus, Hawthorn, Victoria 3122, Australia

<sup>2</sup>Graduate School of Sciences, Information Technology and Engineering and Centre of Informatics and Applied Optimization, Faculty of Science, The Federation University Australia, Mount Helen Campus, Mount Helen, Ballarat, Victoria 3353, Australia

#### **Abstract**

In computational chemistry or crystallography, we always meet the problem that requires distributing N particles in one square unit with the minimal neighbor distance. Sometimes this problem is with special or complex constraints. This short article will build a molecular optimization model for the problem, and then will show one example of the application of this model

**Keywords:** Computational chemistry; Crystal molecular structure; Optimization model; Optimized Voronoi cells distribution

We consider the problem that requires distributing  $N \ (\ge 1)$  particles in one three - dimensional (3D)  $2a \times 2b \times 2c$  box/cell/unit with the minimal neighborhood distance. Let us define that  $d_{ij}$  is the direct-distance variable between particle i  $(1 \le i \le N)$  and particle j  $(1 \le i \le N, j \ne i)$ . Direct-distance means particles i and j have a direct interaction relationship, for example, in computational chemistry, the van der Waals (vdW) contact [1], or solvent accessible surface area (ASA) contact, etc. to each other. Denote  $(x_{i1}, x_{i2}, x_{i3})$  and  $(x_{j1}, x_{j2}, x_{j3})$  the coordinates of particles i and j, respectively. Then, for the convenience of practical computations [2,3], we can build an optimization model for the above problem:

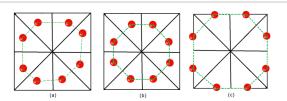
$$\min f(x) = \left(\sum_{i=1}^{N-1} \sum_{j=i+1}^{N} d_{ij}\right)^{2} \tag{1}$$

$$= \left(\sum_{i=1}^{N-1} \sum_{j=i+1}^{N} \left(x_{i1} - x_{j1}\right)^2 + \left(x_{i2} - x_{j2}\right)^2 + \left(x_{i3} - x_{j3}\right)^2\right)^2$$
 (2)

$$-a \le x_{i1}, x_{j1} \le a, -b \le x_{i2}, x_{j2} \le b, -c \le x_{i3}, x_{j3} \le c, i, j = 1, ..., N$$
(3)

This might be a problem of Voronoi diagram and the unit is called Voronoi cell. In computational chemistry, some crystals own special structures of the Voronoi cells; in such a case, we may add some additional constraints to Eq. (3). Clearly, the well-known Lennard-Jones Clusters problem [4] is one case of the above optimization problem Eqs. (1)  $\sim$  (3). Some computer-aided design models can be looked as the problem Eqs. (1)  $\sim$  (3) [5,6,7]. Any optimization algorithms can be used to solve Eqs. (1)  $\sim$  (3) but global optimization algorithms (e.g., in [8]) are more preferred to use.

Example 1: We give a 2D Voronoi cells example (Figure 1). 2D is



**Figure 1:** The optimization model to distribute 8 particles into 8 Voronoi cells of a square unit: (a) initial distribution given, (b) optimal (octagon) distribution inner the square, and (c) optimal (octagon) distribution onto the boundary of the square. The green dashed line denotes there is a direct contact relationship between the two particles they link (e.g., the two atoms have the vdW interactions).

a special case of 3D. We distribute 8 particles in one 2D square with the minimal neighborhood distance among them, with a constraint that each particle is only in one of the 8 Voronoi cells of the square. Figure 1(a) shows the initial solution that is given to the problem. Figure 1(b) and Figure 1(c) show the optimal (octagon) distribution of the 8 particles inner the square and onto the boundary of the square, respectively, after we solve the optimization problem Eqs.  $(1) \sim (3)$ .

Example 2: We give a 3D Voronoi cells example (movies in [9]). The Lennard-Jones clusters problem is clearly a 3D Voronoi cells problem. Cameron et al. presented 4 movies to illuminate how the atoms to be arranged and at last reach the minimal energy states [9].

# Acknowledgments

This research was supported by a Victorian Life Sciences Computation Initiative (VLSCI) grant numbered VR0063 on its Peak Computing Facility at the University of Melbourne, an initiative of the Victorian Government (Australia). The author thanks Dr. Li S. to describe a similar 2D Voronoi cells problem in solar energy cells schedule. Last but not least, the author appreciates all the referees for their comments and suggestions, which have significantly reproved this article.

## References

- OlechnoviÄ□K, Venclovas C (2014) Voronota: A fast and reliable tool for computing the vertices of the Voronoi diagram of atomic balls. J Comput Chem 35: 672-681
- Zhang J, Gao DY, Yearwood J (2011) A novel canonical dual computational approach for prion AGAAAAGA amyloid fibril molecular modeling. J Theor Biol 284: 149-157.
- Zhang JP, Hou YT, Wang YJ, Wang CY, Zhang XS, et al. (2011) The LBFGS quasi-Newtonian method for molecular modeling prion AGAAAAGA amyloid fib- rils. Nat Sci 4(12A) (Special Issue on Bioinformatics, Proteomics, Systems Biology and Their Impacts to Biomedicine): 1097–108.
- Zhang JP (2015) The hybrid idea of optimization methods applied to the energy minimization of (prion) protein structures focusing on the ß2–a2 loop. Biochem Pharmacol (Los Angel) accepted on 03-Jul-2015.

\*Corresponding author: Jiapu Zhang, Molecular Model Discovery Laboratory, Department of Chemistry and Biotechnology, Faculty of Science, Engineering and Technology, Swinburne University of Technology, Hawthorn Campus, Hawthorn, Victoria 3122, Australia, Tel: +61-3-9214-5596; +61-3-5327-6335; E-mail: jiapuzhang@swin.edu.au; j.zhang@federation.edu.au

Received July 12, 2015; Accepted August 06, 2015; Published August 13, 2015

**Citation:** Zhang J (2015) An Optimization Model of Molecular Voronoi Cells in Computational Chemistry. Biochem Pharmacol (Los Angel) 4: 179. doi:10.4172/2167-0501.1000179

Copyright: © 2015 Zhang J. This is an open-access article distributed under the terms of the Creative Commons Attribution License, which permits unrestricted use, distribution, and reproduction in any medium, provided the original author and source are credited.

- Ma DL, Chan DSH, Chung-Hang Leung (2011) Molecular docking for virtual screening of natural product databases. Chem. Sci 2011: 1656-1665.
- Chan DS, Yang H, Kwan MH, Cheng Z, Lee P, et al. (2011) Structure-based optimization of FDA-approved drug methylene blue as a c-myc G-quadruplex DNA stabilizer. Biochimie 93: 1055-1064.
- 7. Ma DL, Lai TS, Chan FY, Chung WH, Abagyan R, et al. (2008) Discovery
- of a drug-like G-quadruplex binding ligand by high-throughput docking. ChemMedChem  $3\colon 881\text{-}884.$
- 8. Zhang JP (2015) Molecular structures and structural dynamics of prion proteins and prions. Springer ISBN 978-94-017-7317-1.
- Lennard-Jones38 Clusters: http://www.cims.nyu.edu/~cameron/rareevents. html, Retrieved on 05 August 2015.