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Molecular networks by invariant shape coordinates and deformation indexes

Andrea Lombardi¹

¹ Dipartimento di Chimica, Biologia e Biotecnologie, Unibersità di Perugia, Perugia, Italy Corresponding author: <u>andrea.lombardi@unipg.it</u>

The classification of large molecules according to structural similarities is a relevant issue in biochemistry [1], even more in the upcoming big data era. The proper choice of parameters, containing invariant structural information, is a convenient way to induce mapping and grouping of structures, depending on predominant structural motifs, individual amino acid geometry or connectivity properties. Few-body hyperspherical coordinates provide the appropriate framework in which shape parameters and deformation indices [2-4] may be defined and applied to characterize complex structures-

Shape coordinates and invariant deformation indexes

The hyperspherical coordinates can be thought of as a generalization of the ordinary spherical coordinates, in the three-dimensional space, to spaces of higher dimensions. The radius is replaced by a "hyperradius", and the two *azimuthal* and *polar* spherical angles are replaced by a set of "hyperangles", by which the higher dimension Cartesian space can be parametrized.

The hyperspherical coordinates of a given system of $N \ge 2$ particles with masses m_1, \dots, m_N and Cartesian position vectors in the center of mass reference frame, $\mathbf{r}_1, \dots, \mathbf{r}_N$, can be derived by introducing a new set of mass *scaled* vectors, $\mathbf{q}_{\alpha} = (m_{\alpha}/M)^{1/2} \mathbf{r}_{\alpha}$ ($1 \le \alpha \le N$), where M is the total mass of the system, and building up a $3 \times N$ position matrix denoted by Z arranging column-wise the components of the \mathbf{q} vectors, as follows:

$$Z = \begin{pmatrix} \mathbf{q}_{1,1} & \dots & \mathbf{q}_{1,N} \\ \mathbf{q}_{2,1} & \dots & \mathbf{q}_{2,N} \\ \mathbf{q}_{3,1} & \dots & \mathbf{q}_{3,N} \end{pmatrix}$$

The next step is allowing to act rotations (left-multiplication) on the position matrix by an orthogonal matrix R^t , transpose of a matrix $R \in O(3)$. Then, alternative rotations can be performed in the so called *kinematic space* [5], by right-multiplication of Z by a $N \times N$ matrix $K \in O(N)$, Z' = ZK. In special cases, such as when the center of mass is the reference, the K matrices have a zero last column, and one only has to deal with a $3 \times (N - 1)$ reduced Z matrix. This lower dimensional matrix Z is called *reduced position matrix*. The *singular value decomposition* theorem [5] applied to the $3 \times n$ position matrix Z (where n = N or n = N - 1) leads to a product of three matrices:

$$Z = R \Xi K^t$$

where $R \in O(3)$ and $K \in O(n)$ are 3×3 and $3 \times n$ orthogonal matrices, respectively, as previously outlined. The new $3 \times n$ matrix Ξ contains all zero entries, with the possible exception of the diagonal elements, $\Xi_{11} = \xi_1$, $\Xi_{22} = \xi_2$, $\Xi_{33} = \xi_3$ for which the inequality

$$0 \le \xi_1 \le \xi_2 \le \xi_3$$

holds.

The values ξ_i , (i = 1, 2, 3) are known as the *singular values* of the matrix *Z* and are uniquely determined, while the factors *R* and *K* in Eq. 3 are not, therefore they are invariant under both ordinary and kinematic rotations [5]. They are directly connected to the moments of the



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inertia of the system and define the global shape of the system [5]. If $N \le 3$ and Z is the full $3 \times N$ position matrix, then the smallest singular value ξ_1 is necessarily zero. The hyperradius ρ is defined as follows:

$$\rho^2 = \xi_1^2 + \xi_2^2 + \xi_3^2$$

This property can lead to applications to molecular dynamics and to the systematic study of the minimum energy structures of *N*-particle aggregates and large molecules such as proteins. Besides the hyperradius, two *deformation indices* can be defined by combining the ζ 's, measuring deviations from the spherical top shape [5]:

$$\xi_{-} = \frac{\xi_1^2 - \xi_2^2}{\rho^2}$$

Which is zero for prolate top configurations, and

$$\xi_{+} = \frac{\xi_{2}^{2} - \xi_{3}^{2}}{\rho^{2}}$$

which is zero for oblate top configurations. By definition, when both indexes are zero, one has a spherical rotor. An additional remarkable property of the ξ 's is that in the case N = 4, is that ξ_3 can take negative values and the chirality sign (mirror images) of the system depends on its sign.

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References

[1] S. Konno, T. Namiki and. K. Ishimori, Quantitative description and classification of protein structures by a novel robust amino acid network: interaction selective network (ISN), Sci. Rep. 16654 (2019)

[2] A. Lombardi, AIP Conference Proceedings, Simmetry and deformations of clusters and biomolecules by invariant shape coordinates, 2343 (1), 020004 (2021)

[3] A. Lombardi, N. Faginas-Lago, N. and L. Pacifici, Protein Networks by Invariant Shape Coordinates and Deformation Indexes. In: Gervasi, O., Murgante, B., Misra, S., Rocha, A.M.A.C., Garau, C. (eds) Computational Science and Its Applications – ICCSA 2022 Workshops. ICCSA 2022. Lecture Notes in Computer Science, vol 13382, 348-359 (2022)

[4] V. Aquilanti, A. Lombardi, E. Yurtsever, Global view of classical clusters: the hyperspherical approach to structure and dynamics, Phys.Chem. Chem. Phys. 4 (20), 5040 (2000)

[5] M. B. Sevryuk, A. Lombardi, and V. Aquilanti, Hyperangular momenta and energy partitions in multidimensional many-particle classical mechanics: The invariance approach to cluster dynamics, Physical Review A 72, p. 033201 (2005)

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