



Mini Review

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Molecular Chirality as an Integral Biomarker of Lactation, Nutrition and Cognitive Development Express-Review

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Abstract

It becomes an axiom that all living organisms are composed predominantly of L-amino acids (L-AAs) and D-sugars. However, in many biomedical studies, there is a significant discrepancy between the amount of available information regarding D-AAs biology and a degree of pragmatic implementation. It is applying to the fields of nutrition and lactation. The mutual interaction of internal and environmental sources of D-AAs in an organism, including the food, microorganisms, is influenced by the spontaneous and enzymatic racemization. The interacting sources of D-AAs are recognized as an essential determinant in the embryonic/infant development, the lifestyle of an adult an organism, and aging. Progress in the analytical methods of chiral discrimination is promising to revolutionize the field of neuroscience, neurodegeneration, psychology, drug production, and the food industry, which should be reflected in the field of lactation. However, the attention to the D/L AAs balance in the area of breastfeeding is overlooked until the present. In short, our aim is predicting the importance of the serine racemase (SerRs) enzymes in human breast milk production and digestion.

Keywords: Lactation; Enzyme; Serine Racemase; Racemization; Post-Translational Modification; Protein Folding; Biochirality

Abbreviations: L/D-AAs: L/D-amino acids; SerR: Serine Racemase; DAAO: D-AA oxidase; D-Ala: D-Alanine; D-Asp: D-aspartic Acid; D-Arg: D-arginine; D-Glu: D-glutamic acid; D-Gln: D-glutamine; D-Leu: D-leucine; D-Lys: D-lysine; D-Met: D- Methionine; D-Ph: D-phenylalanine; D-Ser: D-serine; PTMs: Post-translational modifications; PhTs^{NE}: Non-Equilibrium Phase Transitions; SerRs: Serine Racemase; DAO, DAAO: D-Amino Acid Oxidase

Introduction

Biochirality

"D-amino acids, the enantiomeric counterparts of L-amino acids, were long considered to be non-functional or not even present in living organisms" [1]. "D-Amino acids are now emerging as potential biomarkers" [2]. "Accumulating evidence points to distinctive roles of D-AAs in non-ribosomal physiology" [3]. "Chiral recognition is the principle governing most biomolecular interactions" [4]. Molecular biochirality chirality (BioCh)* has its physical and chemical ground [4,5]. Our concern is the molecular BioCh in relation to the science of nutrition. Under the impact of new coming results, the intuitive impression of the absolute biological homo-chirality is replaced by the concept of prevalent chirality. Consequently, the descriptive terminology is shifted from "mysterious","attractive," and "surprising" to the simple

and objective "fundamental". It becomes an axiom that all living organisms are composed predominantly of L-amino acids (L-AAs) and D-sugars. The main features characterizing a new level of understanding are.

- The physiology of an organism is driven by the fine dynamical balance between the prevalent and inferior (L- and D-) isoforms [6].
- The primary sources of D-AAs in the organism are the enzymatic and spontaneous post-translational modifications (PTM). The significant part of D-AAs pool (from 10- to 20 %) in different organs of an organism (such as kidney, brain, and others) is independent on the racemase enzymes activity [7,8].
- The physical basis of PTMs of protein folding is the non-equilibrium phase transitions (PhTs^{NE}) [9].



The physical basic of the prevalent AAs chirality (as a deviation from an equilibrium racemic state (antiracemization), and the tendency to spontaneous racemization is grounded on the thermodynamics laws. The quantitative relation between the entropy and energy are revealing the Parity Violation Energy Difference (PVED) in the organic molecules [6].

d) An enormous number of macromolecular spatial transformations require the statistical interpretation of the thermodynamics of PTMs.

e) The transformation of protein's spatial symmetry is inevitably accompanied by the alterations of the corresponding thermodynamic state.

f) Spontaneous, nonenzymatic reactions in proteins (including racemization) are known to have the relevance to protein misfolding, aggregation, aging, neurodegeneration, and nutrition. Current evidence indicates that D-AAs are present in the high concentration in microorganisms and the lower level in plants, mammals and humans [7]. Such evolutionary consistency is associated with a variety of life-supporting cellular functions, including cell division, cell proliferation, and cell signaling. Involvement of D-AAs in neurotransmission underlies their role in the laterality of brain structure and functions [9,10]. These functions include learning, memory, spatial perception, and control of behavior. Currently, molecular chirality is widely recognized as a useful biomarker in the

epidemiology and pharmacology [11-13]. At the same time, it is just at the very initial phase in the field of nutrition [14-16].

Source of D-AAs in Human Organism

Post-Translational Modifications

There are several internal and environmental sources of D-AAs in an organism which are influenced by the mutual interaction. Among them, the most frequently considered are the food, microorganisms, and different forms of racemization [1,17]. The primary internal mechanism underlying the appearance of D-AAs containing peptides/proteins is attributed to the enzymatic and spontaneous post-translational racemization (Figure 1). Biogenesis (synthesis and degradation) of D-AAs containing peptides/proteins is performed by the catalytic activity of specific enzymes, including serine racemase (SerR) and D-AA oxidase (DAAO) [18]. According to many experimental evidence, D-Ser plays a distinct role in the biochirality of the food [19], the mammal's physiology and neurotransmission [20,21]. The reason for the specificity of D-Ser in the biochemistry of an organism remain to be studied.

In the organism, D-Ser is synthesized by SerR and degraded by DAAO. D-Ser level in mammals is highest in urine and serum [22]. The key regulator of intra-body chiral AAs level (including D-Ser) is urinary excretion by the kidney [23]. PTMs include both the intrinsic spontaneous and enzyme-driven racemization [1]. As the consequence, D-AAs are detected in all major areas, including kidney, liver, skin and the brain [23-28].

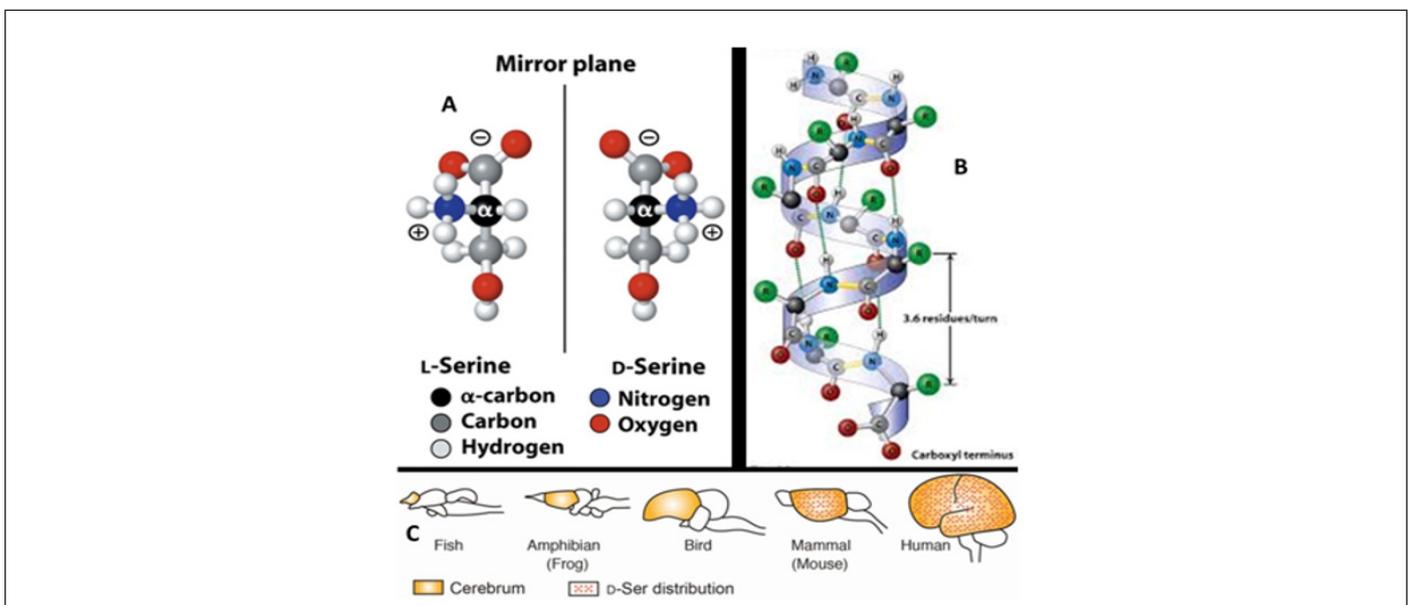


Figure 1: A, B, C

A. All living organisms are composed predominantly of L-amino acids (L-AAs). Example of the Serine (Ser) amino acid.

B. The L and D-isoforms of AAs are different in the way the chain of AAs is folding into the secondary and higher-order structures a result of the angle of the atomic bond. Example of alpha-helix. Adopted from {<https://socratic.org/questions/differences-between-alpha-helix-and-beta-pleated-sheet-structures-of-pro>}.

C. Distribution of D-Ser in the CNS of vertebrates. Typical vertebrate brain structures are shown with the cerebrum highlighted in orange. D-Ser (red dots) occurs exclusively in mammalian (including human) cerebrum. Adopted from {Adopted from [53-54]}.

Microbiota

The interactions between epithelial cells and luminal microbiota have a significant impact on host nutrition and health through the metabolism of dietary components [29]. Microbiota is a critical player in the age-dependent (from infant to adult) food consumption. Notable, that in mammals, many D-AAs are produced by microbiota (gut and intestinal), including D-Ala, D-Arg, D-Asp, D-Gln, D-Glu, D-Leu, D-Lys, D-Met, D-Phe, D-Ser, and D-Trp [30]. Many of them are known as racemization prone. According to recent evidence, about 30% of the human D-AAs are derived from microbial synthesis [31-33]. Microbial D-AAs are involved in communication with the host D-AAs [3]. The essential role of D-AAs recognition in the host-microbe interface is supported by many experimental facts including following that "D-phenylalanine and D-tryptophan regulate chemotaxis of neutrophils through a G-coupled protein receptors, D-Ser has a bacteriostatic role in the urinary tract, D-Ph, and D-Leu inhibit innate immunity through the sweet taste receptor in the upper airway, and D-Trp modulates immune tolerance in the lower airway" [32]. In this situation, it is not surprising that age-related alteration in the intestinal microbiota found to be associated with the brain amyloid-beta (A-beta) plaque deposition [34]. The molecular mechanism of such association could be the microbiota-induced expression of the enzymes (such as D-AAs racemases** and DAO). **Currently, two racemase enzymes have been found in mammals signaling pathways: Ser-Rs and Asp-Rs [17]. Of these two enzymes, only SerRs can be found in human tissue [1]. D-Ser and D-Asp are (so far) the only D-AAs in humans originating from intrinsic racemization [1]. Both enzymes are previously unrecognized mediators of microbe-host interplay, including microbiota-induced racemization.

Diet: Food Processing and Racemization

The presence of D-AAs in foods, including milk, beer, wine [35-38], vegetables, fruit and rice [19,38] is well known. Over the past decades, it has become clear that food processing techniques contribute to the racemization of L-AAs into D-AAs [1]. However, the studies devoted to the role of internal racemization in the digestion are practically absent.

Conclusion

Progress in the analytical methods of chiral discrimination is promising to revolutionize the field of neuroscience, neurodegeneration, psychology, drug production, food industry and lactation. A relevant example is biomarkers in kidney pathology, which, until recently, was a lack of effective biomarkers to the predictive prognosis of chronic kidney disease. The attention to peptide/protein chirality opens a new era of diagnostic and treatment [23]. However, in many biomedical studies, there is a significant discrepancy between the amount of available information regarding D-AAs biology and a degree of implementation. It is

applying to the field of lactation, due to the fact, that D-AAs are the essential players in the embryonic and infant development [40, 41]. Milk proteins of different animals have a high and wide range of the AAs concentrations, which are the subjects of racemization. Consequently, the significance of breastfeeding should be re-considered from the prism of biochirality. Indeed about 20 necessary and not essential AAs found in the breast milk and AAs estimation is a routine element of human milk protein quality control. It is well known that the heating of natural and artificial milk is accompanied by racemization [42]. It is also widely accepted that gut microbiota is an essential mediator of breast milk impact on the host body AAs metabolism [43, 44]. However, attention to the D/L AAs balance is overlooked until the present.

The rare exception is the several recently published review [1,45] which can be recommended as a useful stereo-chemistry-related introduction to nutritionists. Currently, only a little data exists on D-AAs needs in infants and children [46]. Therefore, the role of D-AAs in this field is only on the way to appreciate it. The new strategy of breastfeeding is focused on education to convince that both of two factors are essential: the most effective role of the immediate skin-to-skin contact of a mother with an infant, and the specificity of D-AA's contents. Nonetheless, the attention to the degree of AAs racemization remains to be low or absent. It is known that milk proteins are "predigested" by the diverse family proteolytic enzymes within the human mammary gland [47,48]. The importance of enzymes in human breast milk production and digestion by infants was both predicted and demonstrated [4,47-49]. These experimental facts are in sharp contrast with the situation regarding the enzymes of the post-translational racemization role in the breastfeeding. The implication of AAs racemases in the preparation of milk products is known for the centuries [50]. However, currently an exploration of the enzymes of human milk does not extend to the action of the racemase enzymes (widely expressed in an organism). Based in this discrepancy we are predicting the importance of the serine racemase (SerRs) enzymes family in human breast milk production and digestion. It is notable, that L-Serine (L-Ser) {non-essential amino acid}, is required to synthesize membrane lipids such as phosphatidylserine and sphingolipids [51]. Ser-L deficiency inevitably compromises cell membrane function [52-54] Esaki et al. Therefore, excessive racemization can be also harmful to all significant cellular functions, including the processes of milk production in an organism of mother and milk digestion by infants. Intake of proteins is the key nutritional factors in human diet. The balance of plant and animal proteins has a differential significance for aging stage from infant to ageing adults. At the early stage the developmental aspects are the primary concerns. From the third decade of life a gradual decline in muscle mass and alteration of muscle morphology gain the significance [55,56]. These results have an immediate link to the fundamental role of biochirality in general, prevalence of

L-amino acids, and protein aging in human physiology. Giving the fact that abnormal D-AAAs level is associated with the chronic and acute kidney diseases [57] and recent spark of interest to link the brain-gut-microbiome axis, [58, 59] it is logical to assume that the nutritional aspects of protein chirality are the essential determinants in the complex of issues: from the early development to the age-related decline of the perceptual and cognitive function [59].

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