



Structural and functional characterization of Latroductins I and II, low molecular weight peptidic components from *Latrodectus hesperus* venom

Esperanza Rivera-de-Torre¹, Javier Narbona¹, David Pantoja-Uceda², Gustavo Titau², Miguel A. Treviño², Belén Patiño³, Pilar Medina⁴, Javier Maraver-de-Paz¹, Sara García-Linares¹, José G. Gavilanes¹, Jessica E. Garb⁵, María Ángeles Jiménez², Álvaro Martínez-del-Pozo¹

¹Department of Biochemistry and Molecular Biology, Faculty of Chemistry, Complutense University, 28040 Madrid, Spain. ²Department of Biological Physical Chemistry, Institute of Physical Chemistry Rocasolano, CSIC, Serrano 119, 28006 Madrid, Spain ³Department of Microbiology III, Faculty of Biology, Complutense University of Madrid, 28040 Madrid, Spain ⁴Unidad de Protección de Cultivos, E.T.S.I. Agrónomos, Universidad Politécnica de Madrid, 28040 Madrid, Spain ⁵Department of Biological Sciences, University of Massachusetts Lowell, Lowell, MA, USA
email: esperanza.rivera.detorre@gmail.com

Latroductins (Ltds) are low molecular weight proteins of around 70 amino acids (6-8kDa), with an acidic isoelectric point and a high content of disulfide bonds that have been detected within black widow spider venom (*Latrodectus* spp). They have been isolated in small amounts from the venomous cocktail of black widow species but have not been characterized in deep detail. They usually appear as an omnipresent component in latrotoxin (LTXs) preparations. These LTXs are the most characteristic high molecular weight proteins (110-140kDa) within *Latrodectus* spp venom, whose toxic activity relies in the formation of pores through presynaptic neural membranes. However, Ltds natural biological function is not yet known. The scarce available data suggest that they are essential to increase the neurotoxic activity of LTXs by increasing their binding affinity for the membrane. The association between these two groups of proteins (Ltds and LTXs) has driven some authors to propose their mutual assembly into a *latrotoxin-macromolecular-complex*. In order to contribute to sort out the specific role of Ltds in black widow spider venom, we have cloned the highly expressed LtdI and II proteins from *L. hesperus* in a suitable vector for their production in the yeast *P. pastoris*. The proteins were successfully purified to homogeneity in milligram amounts through several chromatographic steps. They did not show neither antimicrobial nor antifungal activities. Their six cysteine residues were forming disulfide bonds, as supported by mass spectrometry data. Circular dichroism characterization was consistent with thermostable and fully folded globular peptides, showing high α -helical content. Production of Ltds labelled with ¹⁵N and ¹³C isotopes allowed using NMR to determine their three-dimensional structures with atomic resolution.

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