

NMR-Spectroscopic Screening of Crude Venom of Mesobuthus Cyprius

Alexandra Primikyri¹, Socrates Tsinoglou², Evroula Hapeshis³, Christos Petrou³, Yiannis Sarigiannis³

- ¹ Department of Chemistry, University of Ioannina, Ioannina, Greece
- ² MedVenom Ltd, Nicosia, Cyprus

³ Department of Life and Health Sciences, University of Nicosia, Nicosia, Cyprus

† Presented at the First Canadian Peptide and Protein Community Virtual Symposium, 27–28 May 2021; Available online: https://cppc2021.sciforum.net/

Published: 27 May 2021

Abstract: Mesobuthus cyprius, one of the two endemic scorpions in Cyprus, belongs in the family of Buthidae which is geographically distributed worldwide and is the largest of the scorpion families. Moreover, from a clinical perspective, Buthidae is the most important scorpion family as several members of this family are toxic to mammals and can be dangerous to humans. Even though Mesobuthus Cyprius was discovered in 2000 using molecular phylogenetics there are no other published data regarding the peptide and protein composition, the toxicity, or any other activity of the venom. It is impressive, that a broad variety of bioactive substances in scorpion venoms may be considered as a source for drug discovery and development. Direct NMR spectroscopic analysis of unpurified biological extracts is a powerful tool for the discovery of natural products especially in complex mixtures like venoms. It permits partial or complete structural characterization of its major components as well as of many minor components. Herein, we report the application of 1D and 2D NMR spectroscopy for the first time in the analysis of the Mesobuthus Cyprius venom and the identification of a wide range of biomolecules and peptides. Samples were dissolved in D2O and based on COSY, TOCSY, NOESY, HSQC and HMBC spectra, structures were proposed for the venom's major components as well as for many minor components. Final identification of individual compounds can be accomplished through synthesis of proposed structures and additional Liquid Chromatography -Mass Spectrometry Analysis.