

Structure of an Antimicrobial Peptide Buforin II

Gwan-Su Yi^{*}, Chan Bae Park[†], Sun Chang Kim[†] and Chaejoon Cheong^{*}

^{*}Magnetic Resonance Group, Korea Basic Science Institute, Taejon 305-333, and [†]Department of Biological Sciences, Korea Advanced Institute of Science and Technology, Taejon 305-701, Korea.

The structure of an antimicrobial peptide buforin II has been studied by ¹H NMR and CD spectroscopy. Buforin II is flexible and random structure in H₂O but the parts of buforin II become helical structure in trifluoroethanol (TFE)/H₂O (1:1, v/v) solution. From the restrained molecular dynamics calculation using NMR data, we obtained the possible conformation of buforin II in TFE/H₂O solution. The calculated structure contains an α -helix in Val-12 to Arg-20 segment and irregular helix in Pro-11 to Gly-7 segment. These helical region shows amphiphilicity. The possible conformation of this antimicrobial peptide in amphiphilic environment is discussed.