

DETERMINATION OF VALINE AND LEUCINE ISOMERS IN PEPTAIBOLS

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Peptaibols are peptid-like oligomers produced as extracellular secondary metabolites by *Trichoderma* species. Some peptaibols known as effective antibacterial and antifungal agents. They are amphiphilic, which allows them to form voltage-dependent ion channels in cell membranes causing leaks, which finally lead to the death of the cells. These compounds contain non-ribosomally synthesized, non-proteinogenic amino acids, like α -aminoisobutyric acid, ethyl-norvaline, isovaline, and hydroxyproline. Regarding these molecules, the N-terminus is acetylated, and the C-terminal is an amino-alcohol. Due to their diverse nature and bioactivities, the comprehensive elucidation of the structure of newly discovered peptaibols is important.

A lot of efforts have been made to analyse the amino acid sequence of peptaibols using mass spectrometry, however, the isobaric amino acids cannot be specified, thus the configuration of the compounds could not be determined. The hydrolysis of peptaibols followed by liquid chromatographic separation both problems could be solved-, e.g. isobaric amino acids could be separated, and due to chiral derivatisation D-, and L-isomers can be distinguished.

In this work, we focused on the achiral and chiral separation of the isobaric amino acids, valine, isovaline, leucine and isoleucine by HPLC-UV analysis. For these purposes, two HPLC methods were developed. For achiral separation the standard amino acid mixture was firstly derivatised with o-phthalaldehyde (OPA) and then 9-fluorenylmethoxycarbonyl chloride (Fmoc). For chiral derivatisation, Marfey's reagent was used to separate the D-, and L-amino acids. Finally, the developed methods were applied for hydrolysed pure peptaibol fractions.

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