

BIO 2023

Mass spectrometry

> Proteomics Studies: Characterization of Biotechnological Products

Identification

- Peptide mapping
- · N-terminal amino end sequence
- Molecular mass

Structure

- Mapping of disulfide bridges
- Primary sequence
- Post-translational modifications
- Mass protein identification
- Bioinformatic analysis of peptides and proteins
- Studies of Metabolomics of directed and non-directed plant extracts
- Chemical stability studies of active ingredients

Molecular and Microbiological Methods

- Cell culture assays
 - Cytotoxicity and cell viability assays. Induction of hypoxia in cell culture to assess drug response. Study of gene expression in cell culture
- Development and validation of methods for determining impurities by real-time PCR.

Development and validation of methods for determining elemental impurities by ICP-OES or ICP-MS

Raman spectroscopy (non-destructive technique)

- -Identification of raw materials
- -Determination of active substances
- -Polymorphic evaluations
- -Monitoring reactants, intermediates, and end product concentrations, determining pathways, kinetics, mechanisms, endpoints, and yields for various reaction types, such as Diels-Alder, Fischer esterification, Grignard, and hydrogenation.
- -Optimize the conditions of the process. Drug synthesis and crystallization. Identification of falsified medicines

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