Amino Acid Quantitation of Cell Culture Media Matrices via an Integrated CE-MS Analyzer

Overview

Optimization of cell culture media and process conditions are critical for achieving optimal cell growth, productivity, and product quality. Additionally, cell culture media contains many components that can complicate analysis via traditional analytical tools. The coupling of miniature mass spectrometry (MS) to a microchip capillary electrophoresis (CE) platform provides a simple solution for at-line amino acid quantitation.

Core Technology





Ion Trap Mass Spectrometer



Ion trap mass spectrometer operates at ≈ 1 Torr, eliminating need for turbomolecular pumps

Integrated At-Line Amino Acid Analyzer







- Integrated analyzer includes autosampler and all CE-MS components
- Consumable kit optimized for analysis of 200 samples
- Automated on-board algorithms compute concentrations, complete calibrations, and generate pass/fail reports



Ji Young L. Anderson¹, William Morgan-Evans², Bethany Kerr², Reagan Draper¹, Kenion H. Blakeman¹, Milla Neffling¹, Graziella Piras¹ ¹908 Devices Inc., Boston, MA; ²CPI, Darlington, UK



Feed Optimization





- Essential amino acid profiles in ambr250 bioreactors were measured across 12 days
- **Standard feeding:** feed medium was added every other day starting on day 3 Custom feeding: complete feed was supplemented with amino acids based on observed depletion patterns

Feed Strategy Refinement



- titer production
- Customized feeding strategy resulted in up to 25% increase in titer

Subsets of the depleted amino acids were fed to determine which were critical to

The pyramid-based strategy was beneficial for titer increase in several conditions





• Three different media types were spiked with 10x Trypsin EDTA solution • For each media type, a neat sample and a Trypsin EDTA spiked sample

- were prepared

- difference.

Literature References

• Using REBEL diluent, CHO media samples were diluted 100x, insect media samples were diluted 200x, and T cell media samples were diluted 25x • Quantitative results were obtained by analyzing the samples on a REBEL system with n=12 replicates (3 vials, 4 reps per vial)

• 90% of the results had <20% difference between the basal media and spiked media samples. Within the calibrated range, 94% had <15%

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