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changes that occur in specific subpopulations of cells. A protein that is highly expressed (up arrows) in the cells of interest might be detected at low abundance overall due to low expression (down arrows) in background cells.

Cell-selective proteomics for biological discovery

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Protein research is a

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continuous pipeline;
Discovery proteomics involves a small set of samples (typically up to 8) in which hypotheses are tested, and key proteins are identified and quantified. These proteins can then be verified and validated in greater numbers of samples to account for biological variation.

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proteomics is
complicated.

Proteomics emerged as
a result of two factors:
maturation of genomic
global sequencing
profiling and the
unprecedented
technological boom in

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mass spectrometry (MS) in the 1990s. The term “proteomics” was quickly accepted and the field expanded.

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Proteomics is the characterization of all proteins in a biological system including the

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protein spatial distribution and temporal dynamics, the protein response to environmental stimuli, and the differences in protein states between diseased and control biological systems.

Proteomics - an overview | ScienceDirect Topics

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