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Proteomics seeks to identify and to characterize all the proteins synthesized in a cell or a tissue. Based on this information, one can then try to understand how individual proteins or protein collectives function within an organism. The first half of the course focuses on current methodology used to analyze and identify proteins.

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Proteomics Workflows. The first step of proteomics is sample preparation. In this step, we are trying to extract protein from cells. In the second step, we use methods such as 2D electrophoresis to separate different proteins. Then we try to cut proteins into peptides since peptides are easier to detect.

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