

NEWS

Relative LC-MS/MS-based label-free protein quantification allows comparing strains growing conditions or different strains amongst themselves (see Guillaume and all.). By further data analysis, it is then easy to get a large view on the ongoing events in the organism.

Label-free quantitative proteomics of two *Bifidobacterium longum* strains. Guillaume and all. 2009.

<http://www.ncbi.nlm.nih.gov/pubmed/19328873>

High-resolution nano LC-MS/MS quantitative proteomics and CORAVALID™ data processing: The efficient tool

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