

“ We have published several papers utilizing IPA software. The software is currently a standard tool for proteomics or RNA-seq analysis.

— Maciej Lalowski, Adjunct Professor, University of Helsinki,
Faculty of Medicine, Meilahti Clinical Proteomics Core Facility

Source: Maciej Lalowski, Adjunct Professor, University of Helsinki,
Faculty of Medicine, Meilahti Clinical Proteomics Core Facility



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