

## Modeling the potential of kidney disease with an integrated organoid omics map





Proteome of kidney organoids evolves with duration in culture. a) Volcano plot of proteomic differential expression analysis (log<sub>2</sub> fold change of label-free quantification intensity comparing D29 with D21). Proteins are represented by dots and the black line illustrates the significance cut-off (two sided t test, FDR 25 proteins) were distinguished and are indicated with colored rectangles. These five clusters are shown in (d). d) GO enrichment analysis of unbiased clustering (clusters >25 proteins) of proteins during organoid differentiation (D21–D29) (Fisher's exact test, FDR



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