

Data Supplement

GPR19 coordinates multiple molecular aspects of stress responses associated with the aging process

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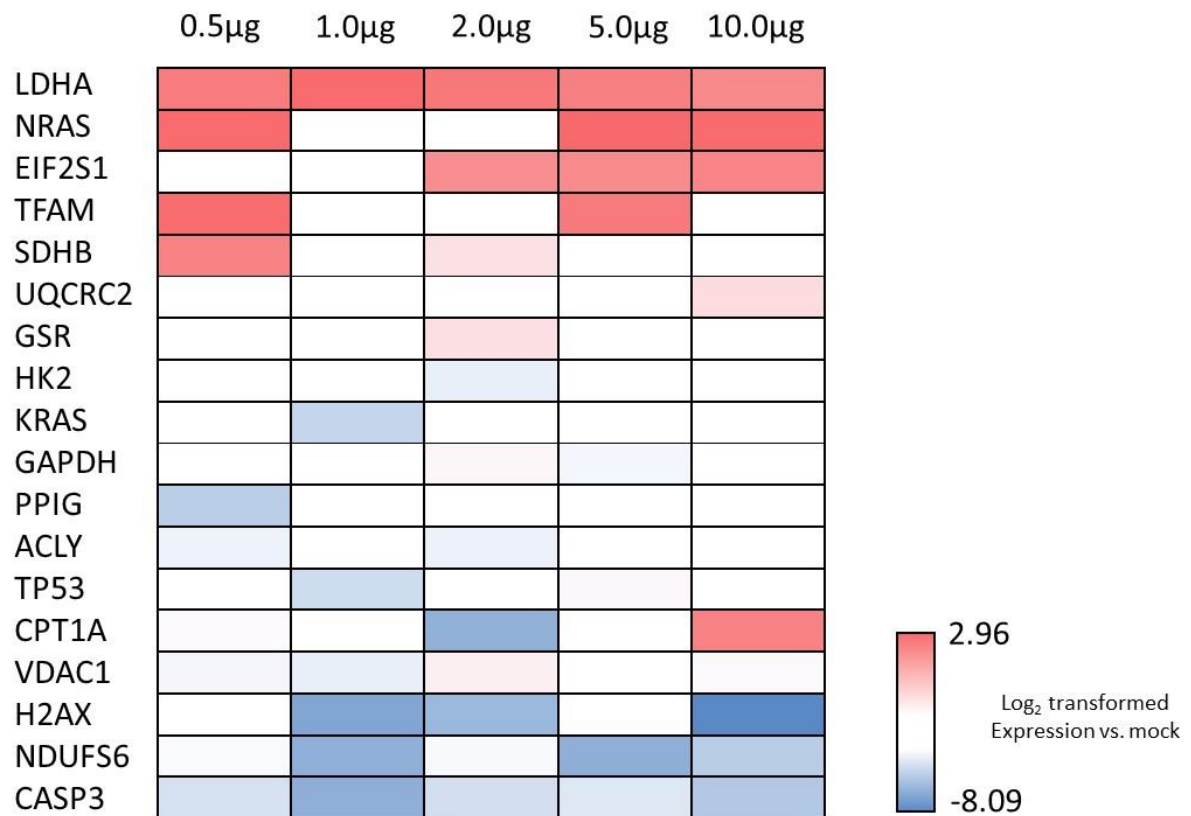


Figure S1. Expression heatmap for proteins common between metabolic pathways and Pujana ATM, BRCA1 and CHEK2 pathways. For each differentially expressed protein the expression value – compared to the expression found in mock transfected cells – is demonstrated with upregulated proteins denoted in red and downregulated proteins identified in blue.