

Supplementary Figure 2

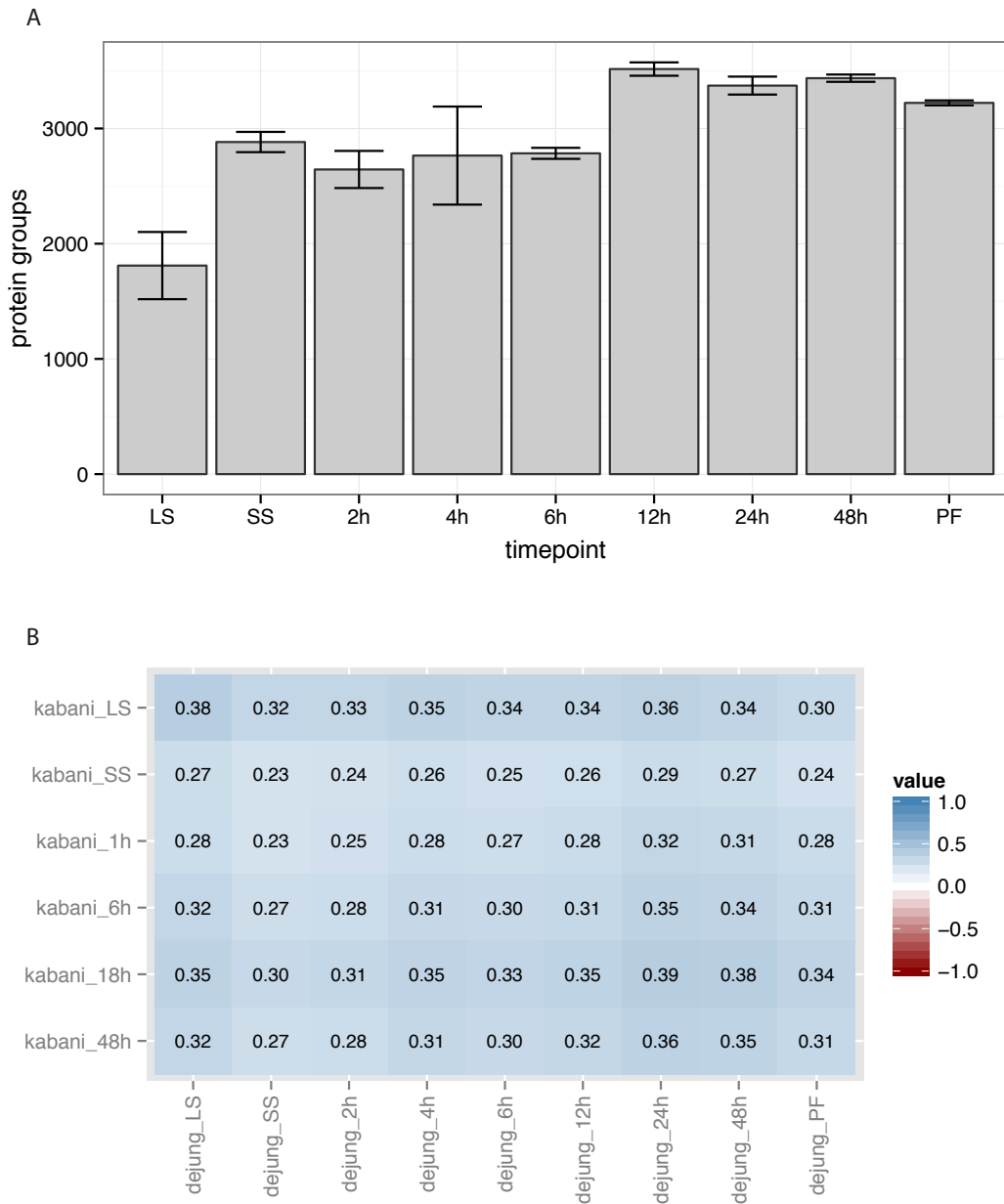


Fig S2: Proteome complexity and comparison of proteome expression profiles with the transcriptome during the differentiation process. (A) Protein groups count per timepoint. **(B)** The overall correlation ranges from $r=0.23$ to $r=0.39$. A global maximum is found when comparing the transcriptome 18 h post induction with the proteome 24 h post induction. The transcriptome and proteome of the LS stage shows the second highest correlation.