

Figure S8 – Individual 50S r-proteins H/M ratio changes in proteome during stationary phase (1/4)

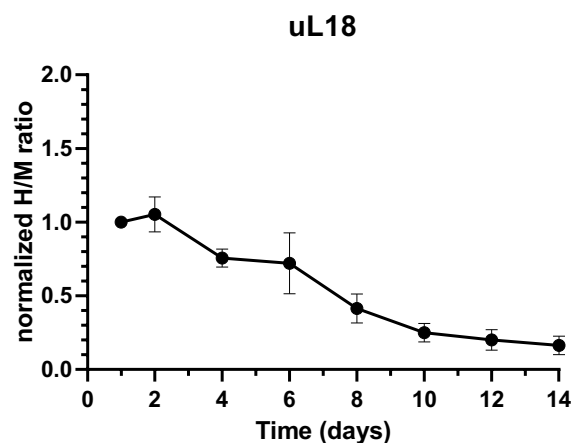
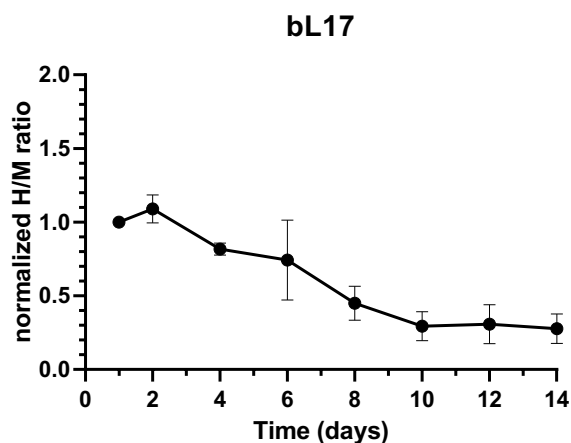
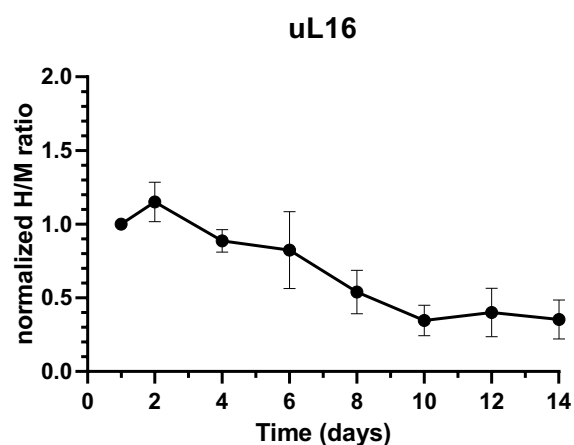
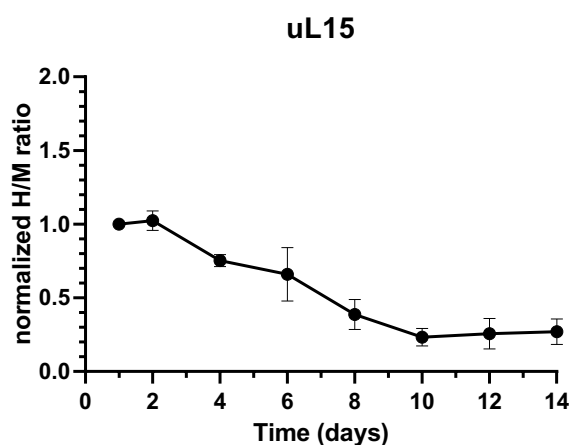
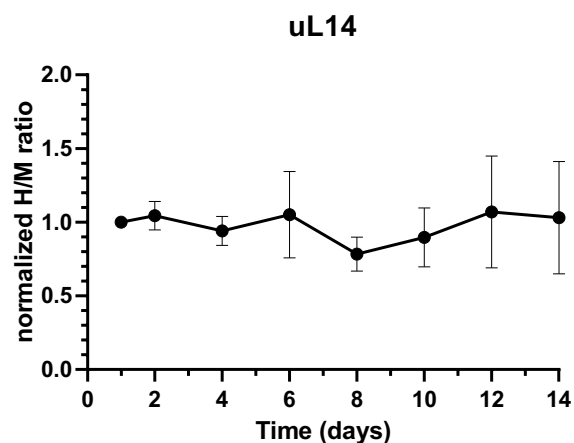
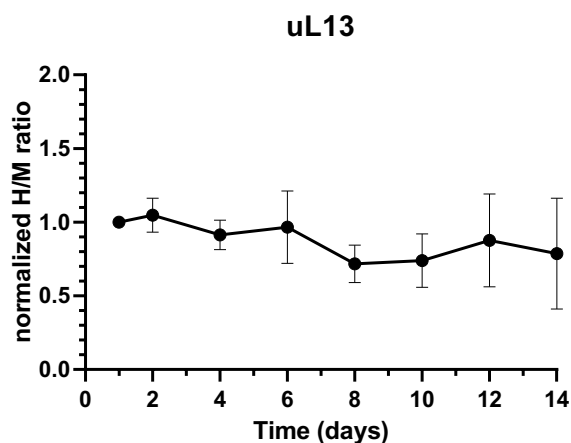
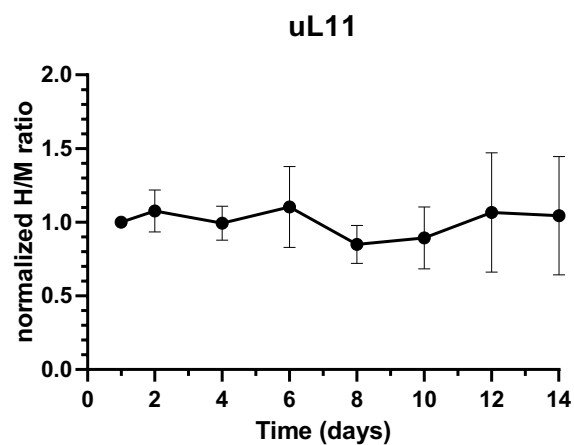
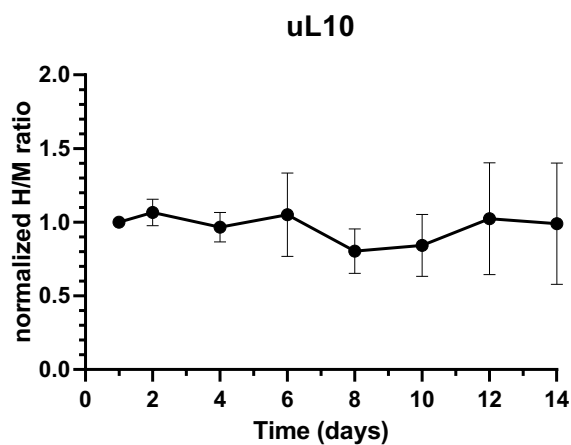


Figure S8 – Individual 50S r-proteins H/M ratio changes in proteome during stationary phase (2/4)

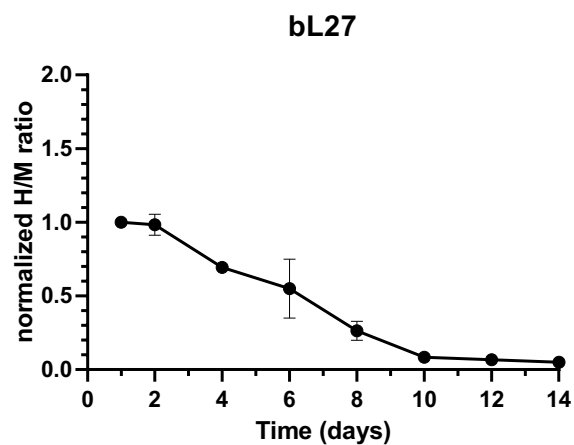
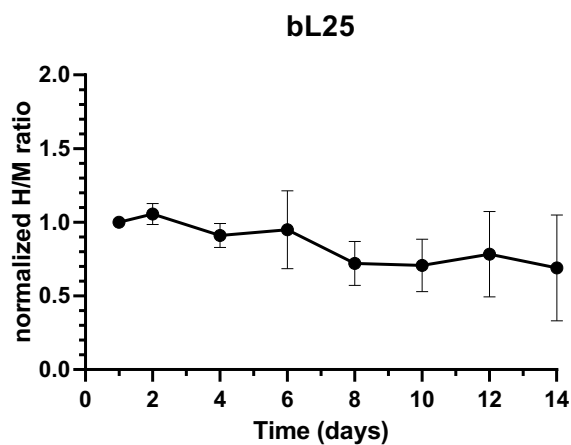
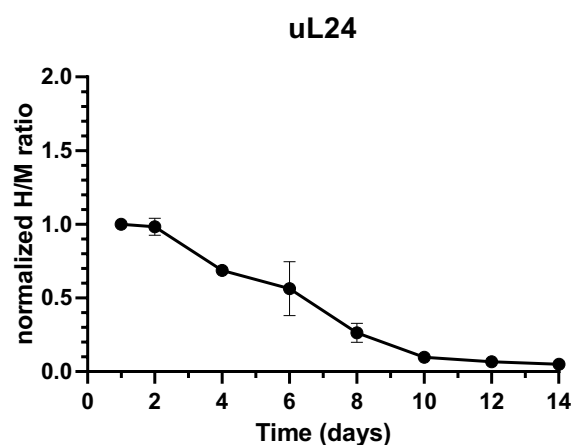
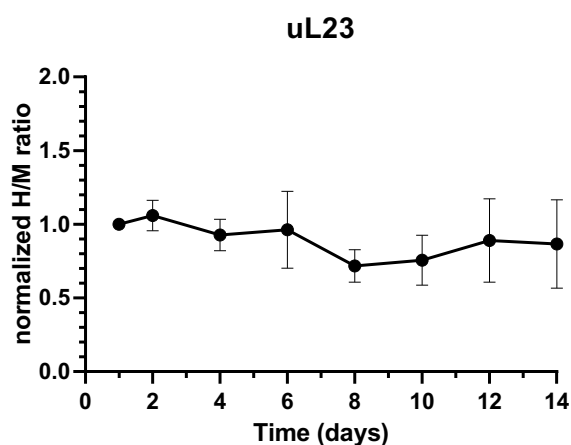
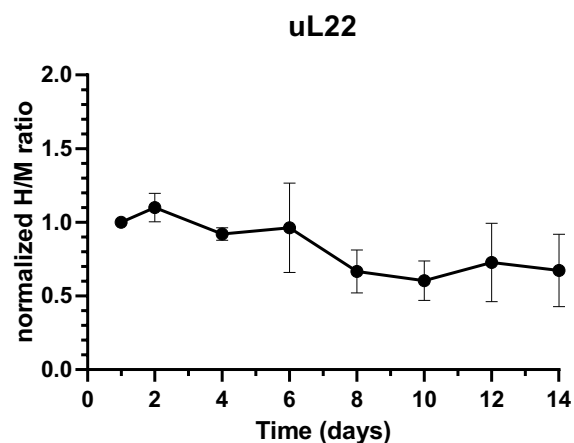
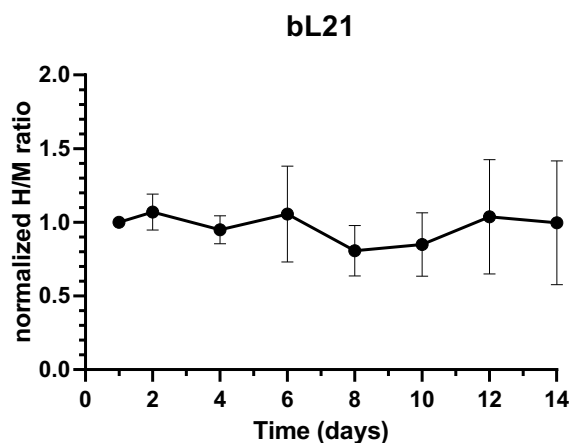
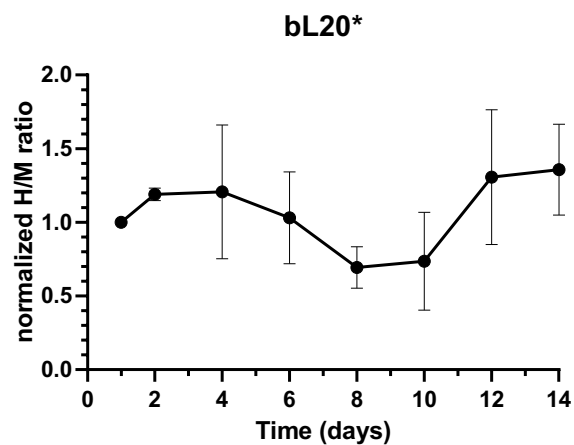
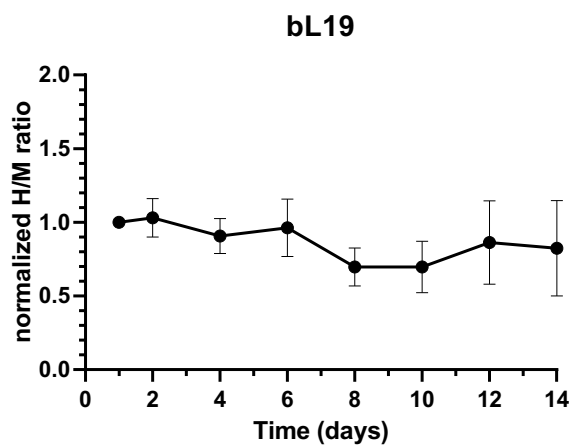


Figure S8 – Individual 50S r-proteins H/M ratio changes in proteome during stationary phase (3/4)

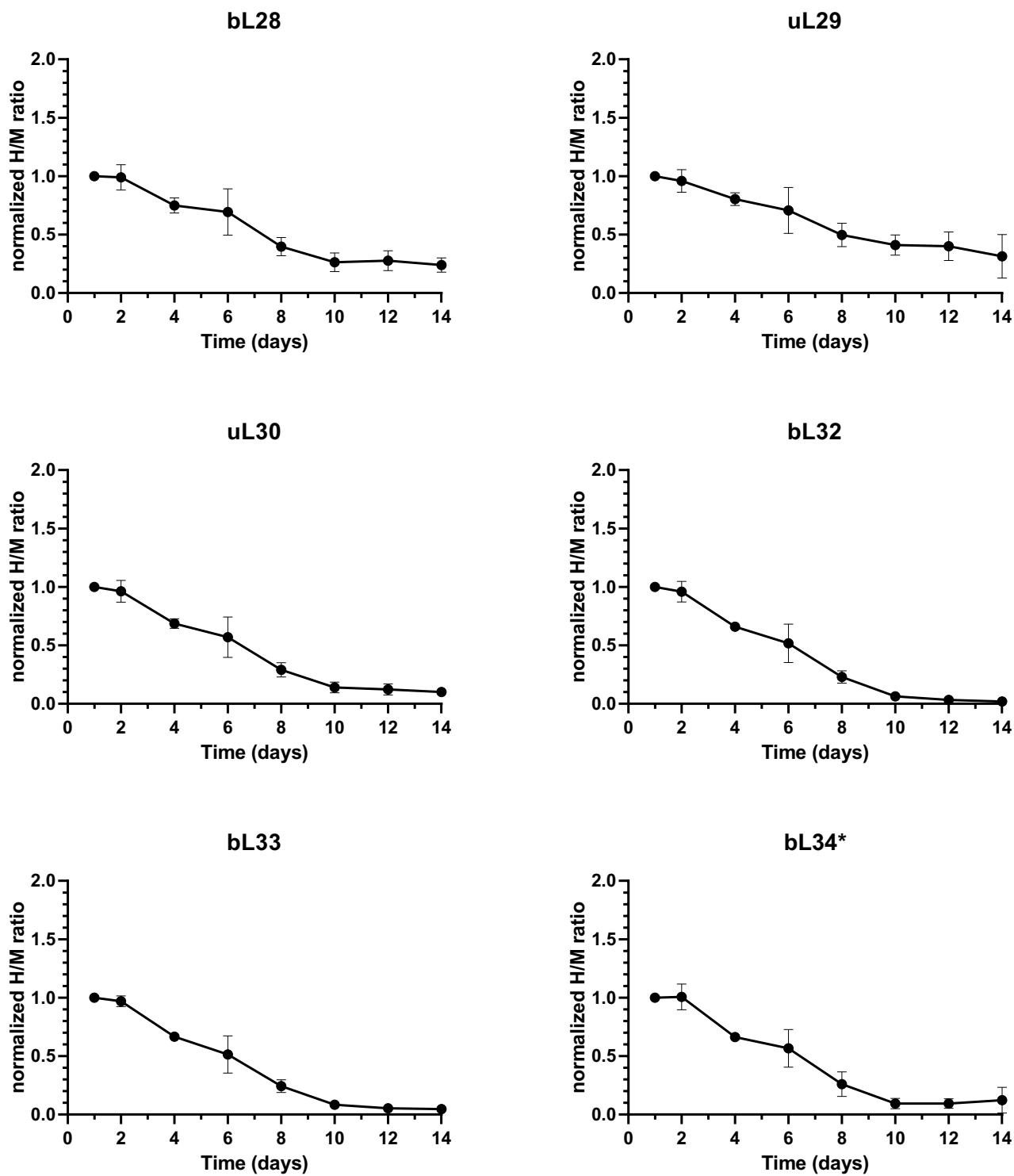


Figure S8 – Individual 50S r-proteins H/M ratio changes in proteome during stationary phase (4/4)

Figure S8 – Individual 50S r-proteins H/M ratio changes in proteome during stationary phase. This figure is related to Figure 5B in the main text. Cells were collected over the course of 14 days, lysed, and total protein from samples was mixed in a 1:1 ratio with total protein from the reference cells containing medium-heavy labeled arginine and lysine. Samples were analyzed using LC-MS/MS. Normalized H/M ratio describes r-proteins relative quantities against day 1. Values shown in the figure are the means of three independent biological experiments with standard deviation (n=3; mean \pm SD).