



Figure S4 - bL31 and bL36 paralogues in *E.coli* early stationary phase proteome. Samples were collected over the course of two days. Total protein from the cells was mixed with reference ribosomes containing a known quantity of bL31A and bL36A (A reference) or bL31B and bL36B (B reference) and analyzed using qMS. The relative quantity of r-proteins is shown as the L/M ratio (Y-axis) and is normalized to the average L/M ratio of 50S r-proteins in the sample. Values shown in the figure are the mean of two independent biological experiments with standard deviation (n=2; mean \pm SD).