

**Figure S1. Chaperone levels in different genotypes.**

Protein levels were determined by western blots for all genotypes relative to the wild type (1.0). Data are shown for two molecular chaperones (a) DnaK and (b) GroEL for each genotype (x-axis) before (white bars) and after (hatched bars) evolution in LB. Bars show standard errors. Neither chaperone was overexpressed after evolution (main text).

