

**Table S1.** Levels of aggregates, acetylated proteins and viable cells in *E. coli* strains with impaired acetylation/deacetylation pathways. The bacteria were grown in TB7 medium supplemented with 0.4% glucose or 25 mM acetate.

Strains	Aggregates (%)	Relative level of acetylation	Viable cells (%)
WT	5.0 ± 0.50	1.0 ± 0.09	65 ± 5
$\Delta ackA$	7.3 ± 0.78	2.1 ± 0.20	98 ± 2
$\Delta pta$	5.1 ± 0.51	0.4 ± 0.04	75 ± 6
$\Delta ackA\text{-}pta$	5.5 ± 0.57	0.2 ± 0.03	75 ± 7
$\Delta acs$	3.9 ± 0.37	0.6 ± 0.06	60 ± 5
$\Delta yfiQ$	4.1 ± 0.38	0.3 ± 0.02	67 ± 6
$\Delta cobB$	5.2 ± 0.54	2.7 ± 0.22	60 ± 6
$\Delta ackA\text{-}pta\text{-}acs$	2.9 ± 0.24	0.01 ± 0.002	55 ± 4
$\Delta ackA\text{-}pta\text{-}yfiQ$	3.2 ± 0.28	0.1 ± 0.01	55 ± 5
WT	7.0 ± 0.64	3.7 ± 0.04	60 ± 4
$\Delta ackA$	7.0 ± 0.55	3.7 ± 0.39	70 ± 6
$\Delta pta$	9.2 ± 0.53	4.4 ± 0.51	95 ± 4
$\Delta ackA\text{-}pta$	7.5 ± 0.70	2.9 ± 0.32	80 ± 4
$\Delta acs$	6.1 ± 0.80	3.4 ± 0.37	81 ± 6
$\Delta yfiQ$	6.4 ± 0.63	3.0 ± 0.32	60 ± 5
$\Delta cobB$	7.2 ± 0.60	3.9 ± 0.41	70 ± 6
$\Delta ackA\text{-}pta\text{-}acs$	5.1 ± 0.76	0.2 ± 0.02	65 ± 5
$\Delta ackA\text{-}pta\text{-}yfiQ$	4.9 ± 0.50	0.2 ± 0.03	65 ± 5