

Supplementary Table 1. Bacterial strains used in this study

Bacterial strain ⁱ	Relevant genotype	Source or reference
BG214	<i>trpC2 metB5 amyE sigB37 xin-1 attSPβ attICEBs</i>	Strain collection
BG703	+ Δ <i>ruvAB</i>	(1)
BG190	+ Δ <i>recA</i>	(2)
BG855 ^b	+ Δ <i>recU</i>	This work
BG651 ^b	+ Δ <i>recA</i> Δ <i>recU</i>	This work
BG1019	+ <i>recU56</i>	This work
BG1021	+ <i>recU71</i>	This work
BG1023	+ Δ <i>recA</i> <i>recU56</i>	This work
BG1025	+ Δ <i>recA</i> <i>recU71</i>	This work

^aAll strains are isogenic with BG214. ^bThe entire *recU* gene coding region was deleted.

1. Sanchez H, *et al.* (2005) The RuvAB branch migration translocase and RecU Holliday junction resolvase are required for double-stranded DNA break repair in *Bacillus subtilis*. *Genetics* 171: 873-883.
2. Ceglowski P, Luder G, Alonso JC (1990) Genetic analysis of *recE* activities in *Bacillus subtilis*. *Mol Gen Genet* 222: 441-445.