1 Supplemental figures



Fig. S1. Comparison of competence-regulated RFP fluorescence intensities.
Cells of strain carrying P_{ssbB}.RFP were grown in C+Y medium, and induced at OD600
0.05 with CSP (1 µg/ml) and analyzed after 1.5h. Average fluorescence intensities at
the single cell level measured by fluorescence microscopy using an mCherry filterset,
polychroic mirror QUAD2 (•) or a tritc filter set; polychroic mirror QUAD1 (•). Values
are in arbitrary units [A.U.]. Single cell fluorescence levels were determined using
MicrobeTracker (1).



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Fig. S2. Comparison of tagRFP, mCherry and mKate2 in a C-terminal protein 15 fusion to the histone-like protein HIpA. (A) Non-deconvolved, unprocessed 16 micrographs of merodiploid D39 strains expressing HIpA-TagRFP (KB1-65), HIpA-17 mKate2 (MK119) and HlpA-mCherry (MK218), respectively. Scale bar equals 2 µm. 18 The filter sets TRITC for HIpA-TagRFP or mCherry for HIpA-mKate2 and HIpA-19 mCherry were used. (B) Immunodetection of fusion proteins in whole cell extracts of 20 strains KB1-65 (hlpA, hlpA-tagRFP), MK119 (hlpA, hlpA-mKate2), MK218 (hlpA, 21 hlpA-mCherry) and KB1-64 (hlpA-mCherry) using anti-RFP antibodies. 22

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25 Supplementary references

Sliusarenko O, Heinritz J, Emonet T, Jacobs-Wagner C. 2011. High throughput, subpixel precision analysis of bacterial morphogenesis and
 intracellular spatio-temporal dynamics. Mol Microbiol 80: 612–627.