



Figure S1. Standard curves used for quantitative Western blot and qRT-PCR. Plotted standard curves for the quantitative Western blot analysis of Hag (A), CsrA (B), and FliW (C) with nanograms (ng) on the x-axis and signal on the y-axis. Each dot represents a replicate for wild type and signal intensity within the standard curve. Panel D) Plotted standard curve used for qRT-PCR with Ct value on the x-axis and the number of hag transcripts on the y-axis. Blue dots represent the standard curve generated when the in vitro transcribed hag transcript was mixed at a 1:1 ratio with RNA extracted from the hag deletion background (DS1677). Red dots represent the standard curve generated when the in vitro transcribed hag transcript was added to the lysed hag deletion background and re-extracted (see methods for more detail).