



Figure S1. Regulation of *Spy_0186* and *Spy_0195* by CsrR in GAS strains 854 and DLS003. Data represent fold-change in transcript abundance of *Spy_0186* and *Spy_0195* as assessed by qRT-PCR in strain DLS003ΔR relative to that in DLS003 (left) or in strain 854Δ*csrR* relative to that in 854 (right). Values are mean±SEM of duplicate samples of three biological replicates.