**Figure S1. Complementation analysis to further assess the SoxR-dependence of genes identified by RNA-Seq.** qRT-PCR was performed on RNA isolated from WT/pSET152, Δ*soxR*/pSET152, and a Δ*soxR* strain complemented with wild-type *soxR* (pSoxR), that were grown on R2YE plates for 3 days. The expression levels of all genes were standardized to the level of the constitutively expressed housekeeping sigma factor, *hrdB,* and normalized to expression in WT/pSET152.