

## Supplementary Data

Table S1

*Streptomyces coelicolor* strains used in the study

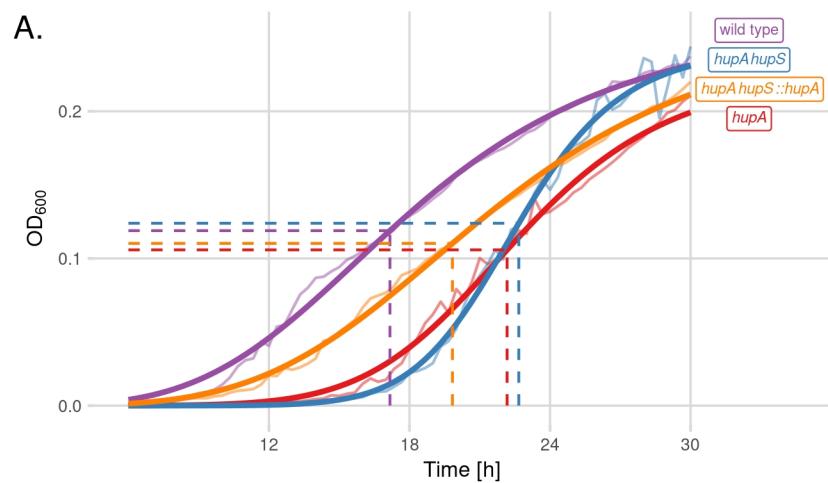
Strain	Relevant genotype	Sources or reference
M145	SCP1- SCP2-	Bentley et al., 2002
ASMK011	M145 $\Delta hupA::scar$	Strzałka et. al., 2022
K304	M145 $\Delta hupS::higro$	Salerno et. al., 2009
ASMK019	M145 $\Delta hupA::scar \Delta hupS::higro$	This study
ASMK019.2	M145 $\Delta hupA::scar \Delta hupS::higro$ pSET152 <hupa></hupa>	This study

Table S2

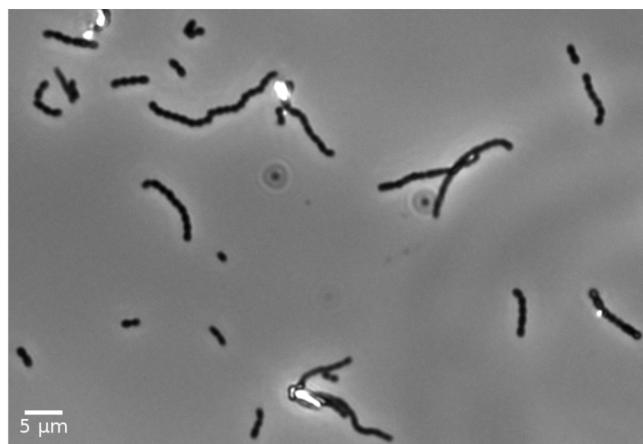
Oligonucleotides used in the study

Primer name	Sequence
sigma_FW	GGCTGGAGGCAAGGTTCGTG
sigma_RV	GCCGAGGCGATGTCGATCTC
gvp_FW	CGACGTGCTGGAGCTTGTCT
gvp_RV	CAGGATCTCGATGCCGACCA
dpsA_FW	GCACCTGACCCTCAAGCAC
dpsA_RV	GTCGTCCCAC TTCCGCTC
ecrA_FW	GGCACCCGCAACGACACC
ecrA_RV	CCAGCAGCGTCCAGCCGA

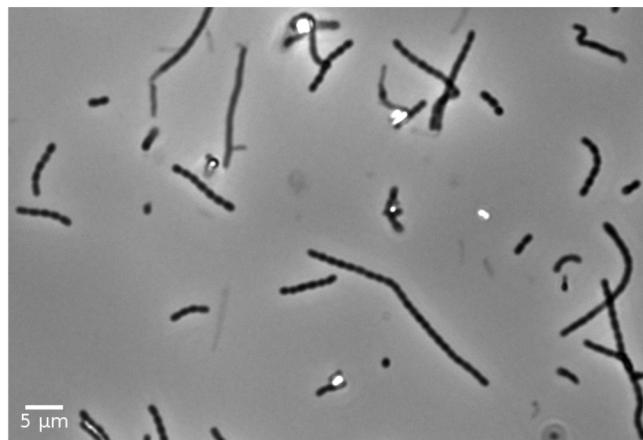
fab_FW	TGGAGCGGACGATGGACAC
fab_RV	GCCAGGTCGGAGGTGAGG
lsr_FW	CGTGTCTGGTCACTCTTTG
lsr_RV	GGCATTGGCTTCGTTAGGT
nsdA_FW	GCCCACCGCCCAGAAGTA
nsdA_RV	GCTGGAGAGGATGTAGGAGCC
sigB_FW	CGACGACTTCCACGCCCT
sigB_RV	ATGTCCTCTGGGTGAGTCCT
rfsA_FW	TTCCGGCTCACCCGTACG
rfsA_RV	TCGACCAGCAGCAGACCC
hrdBRT_f	TGCTCTCCTGGACCTCATC
hrdBRT_r	GTAGCCCTTGGTAGTCGAA
hupA_pSET_FW	GCTTGGGCTGCAGGTCGACTCGAACTACATCATCCCCAG
hupA_pSET_RV	CGATATCGCGCGCGCCGCGCTACTGCCCTGGCGGCTTC



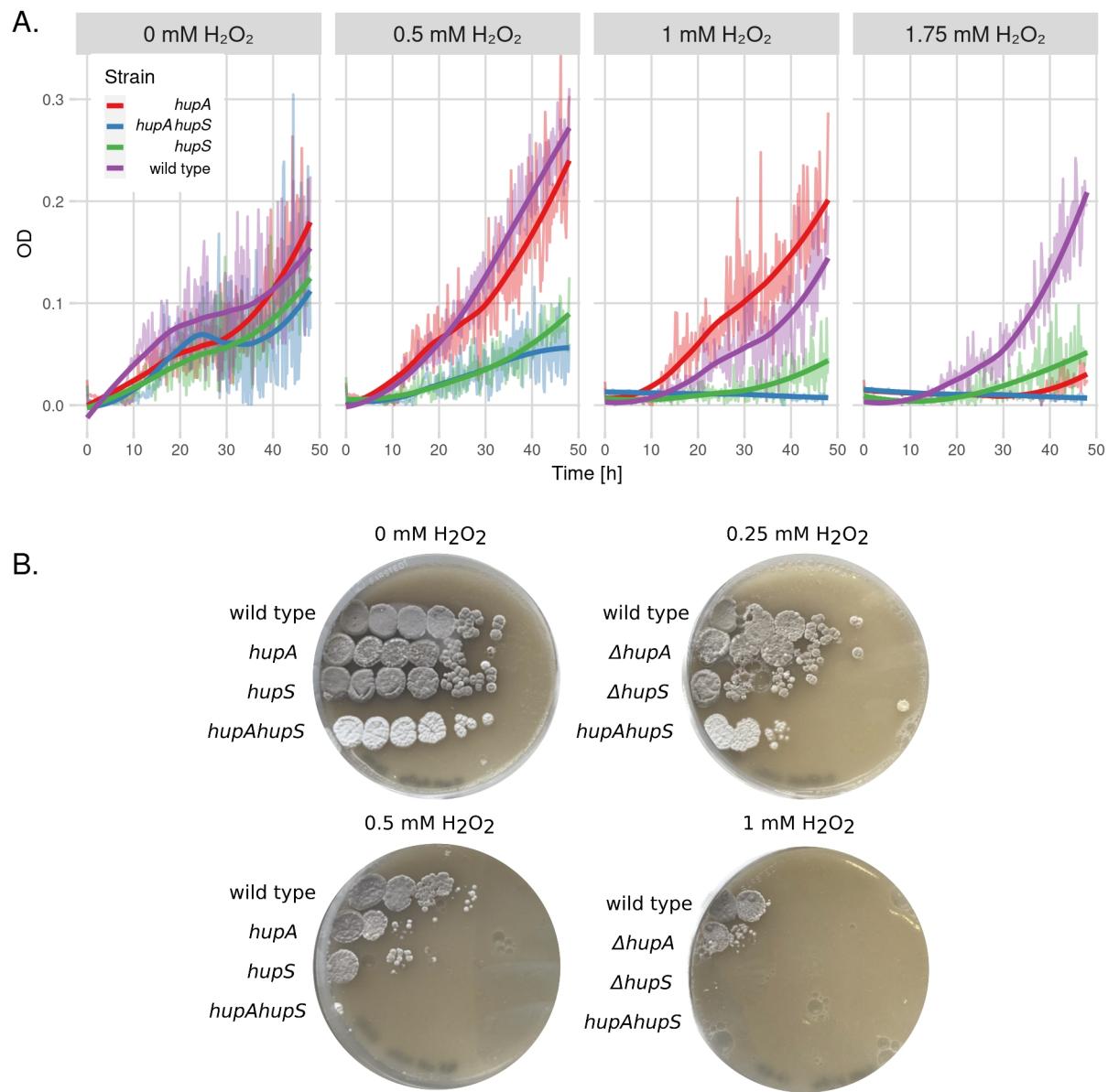
**B.** Wild type



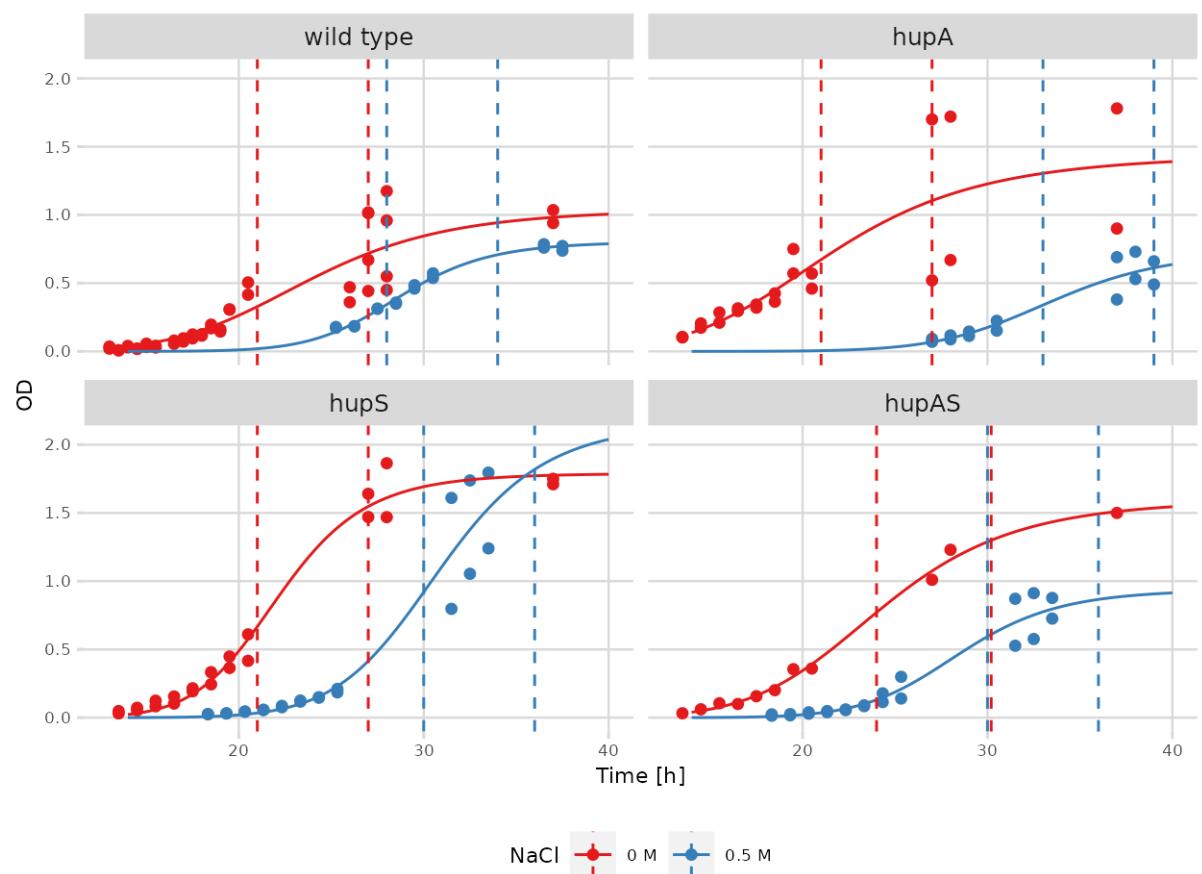
$\Delta hupA \Delta hupS$



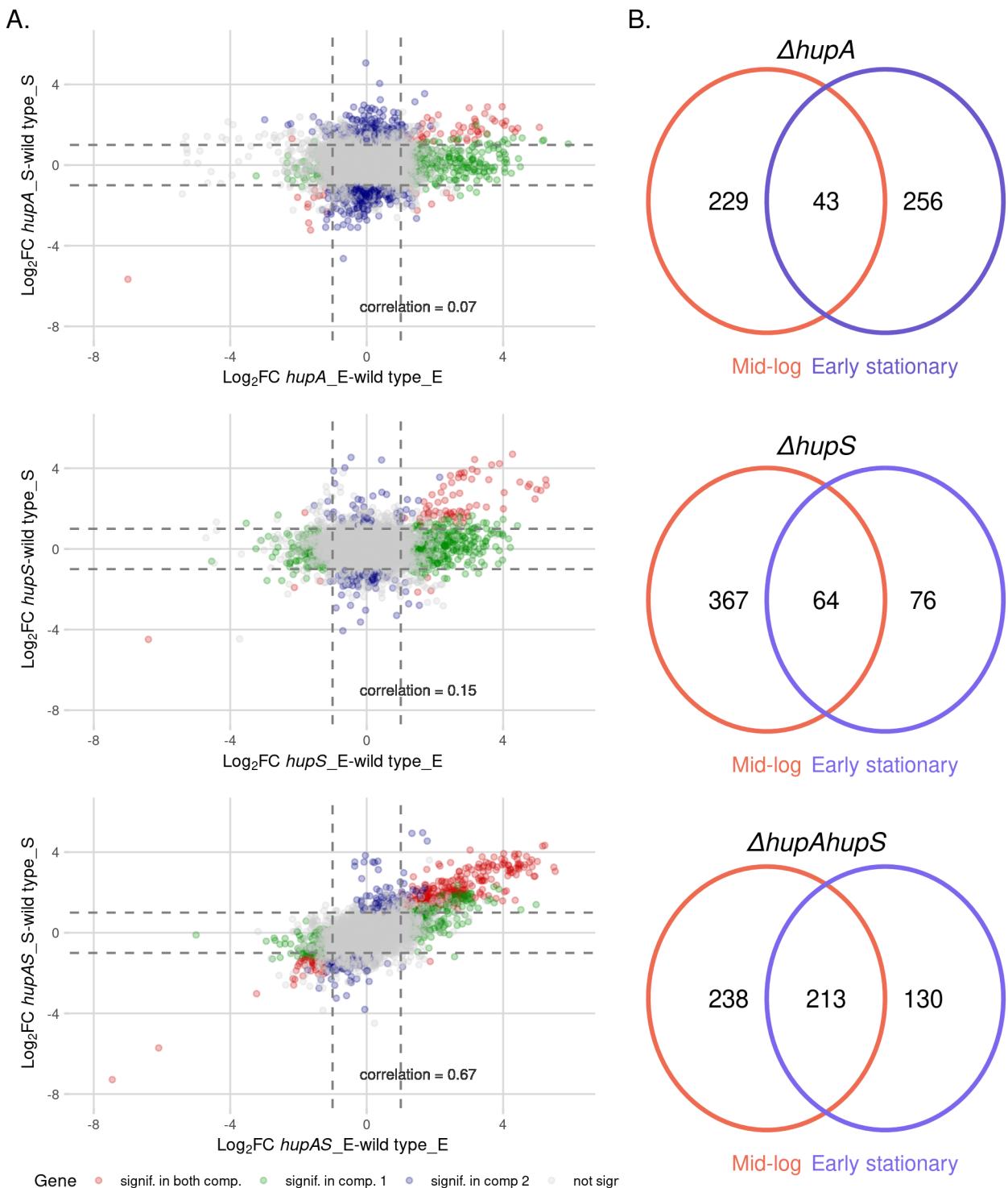
**Figure S1**



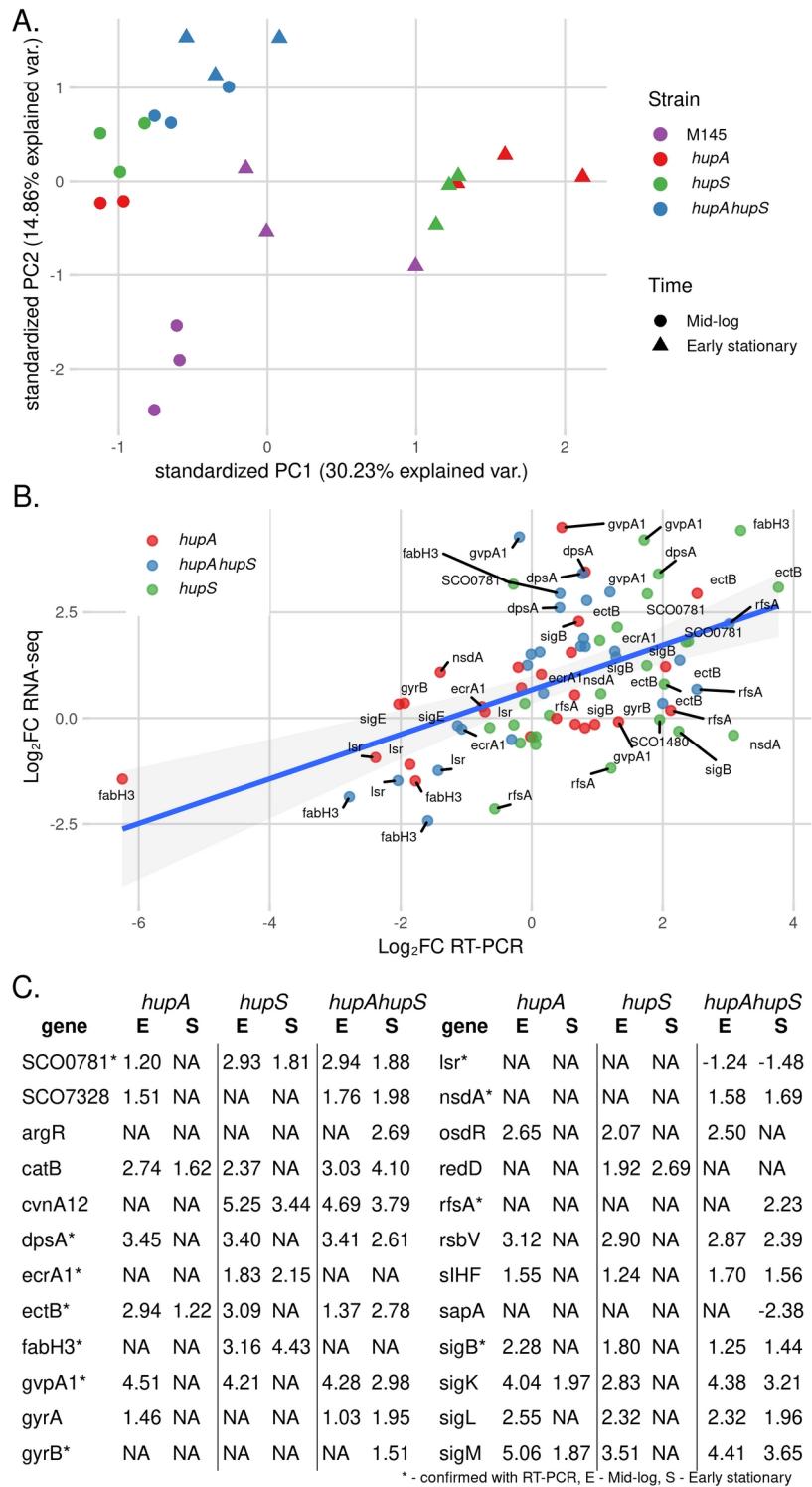
**Figure S2**



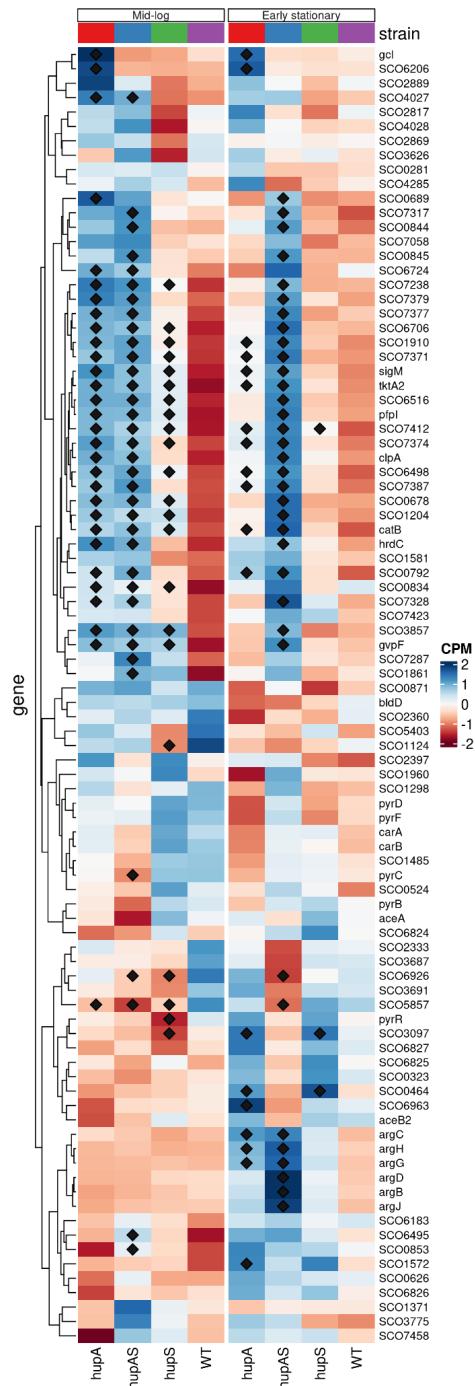
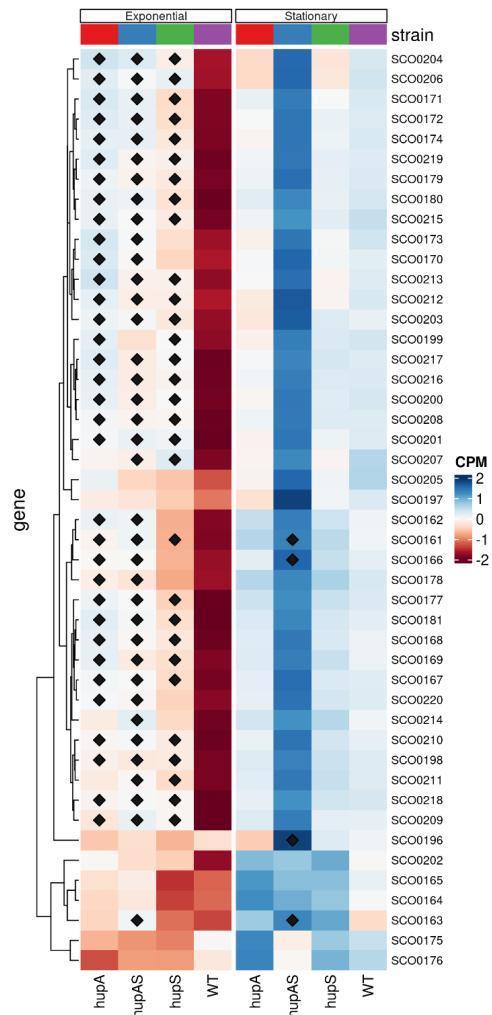
**Figure S3**

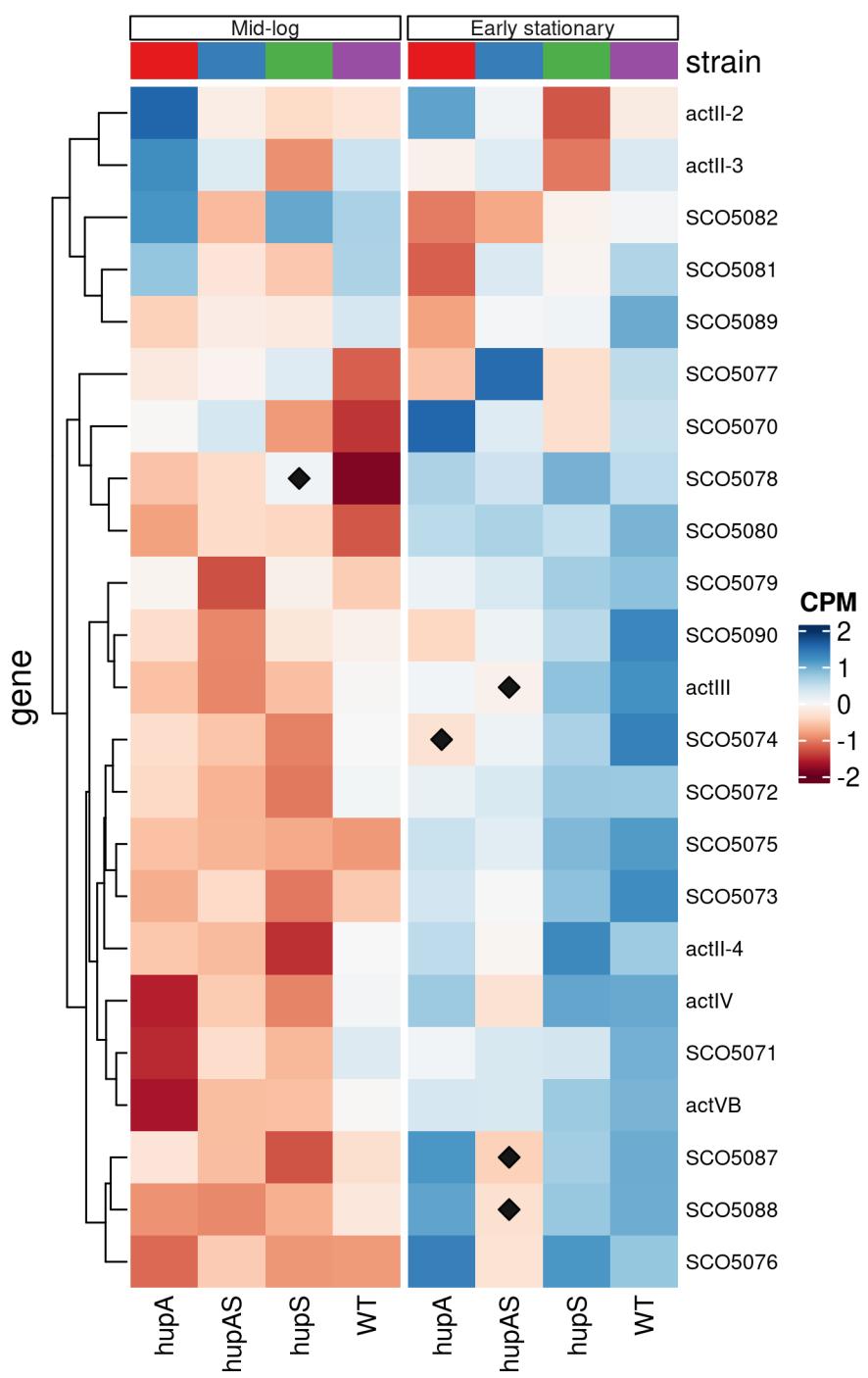


**Figure S4**

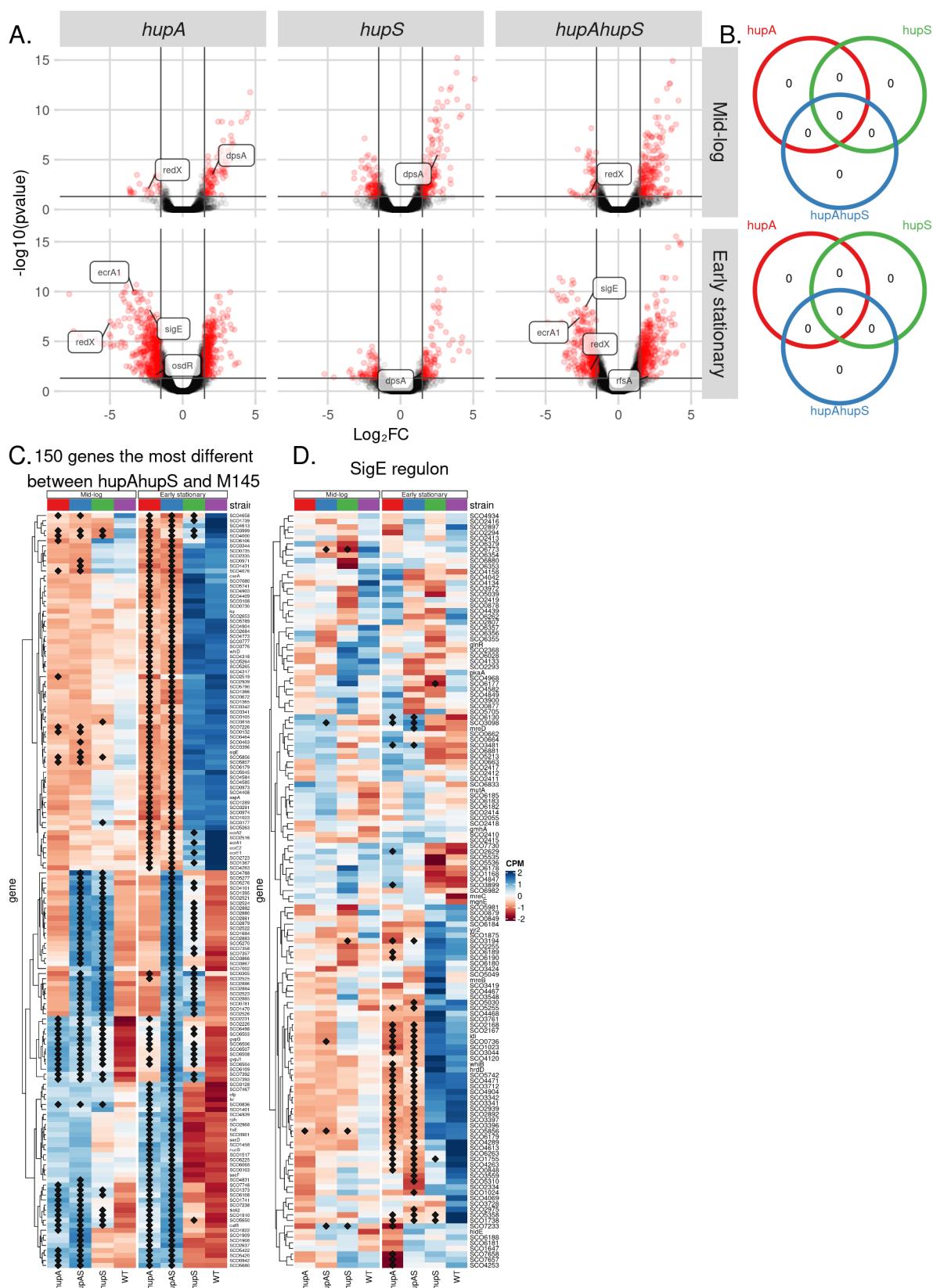


**Figure S5**

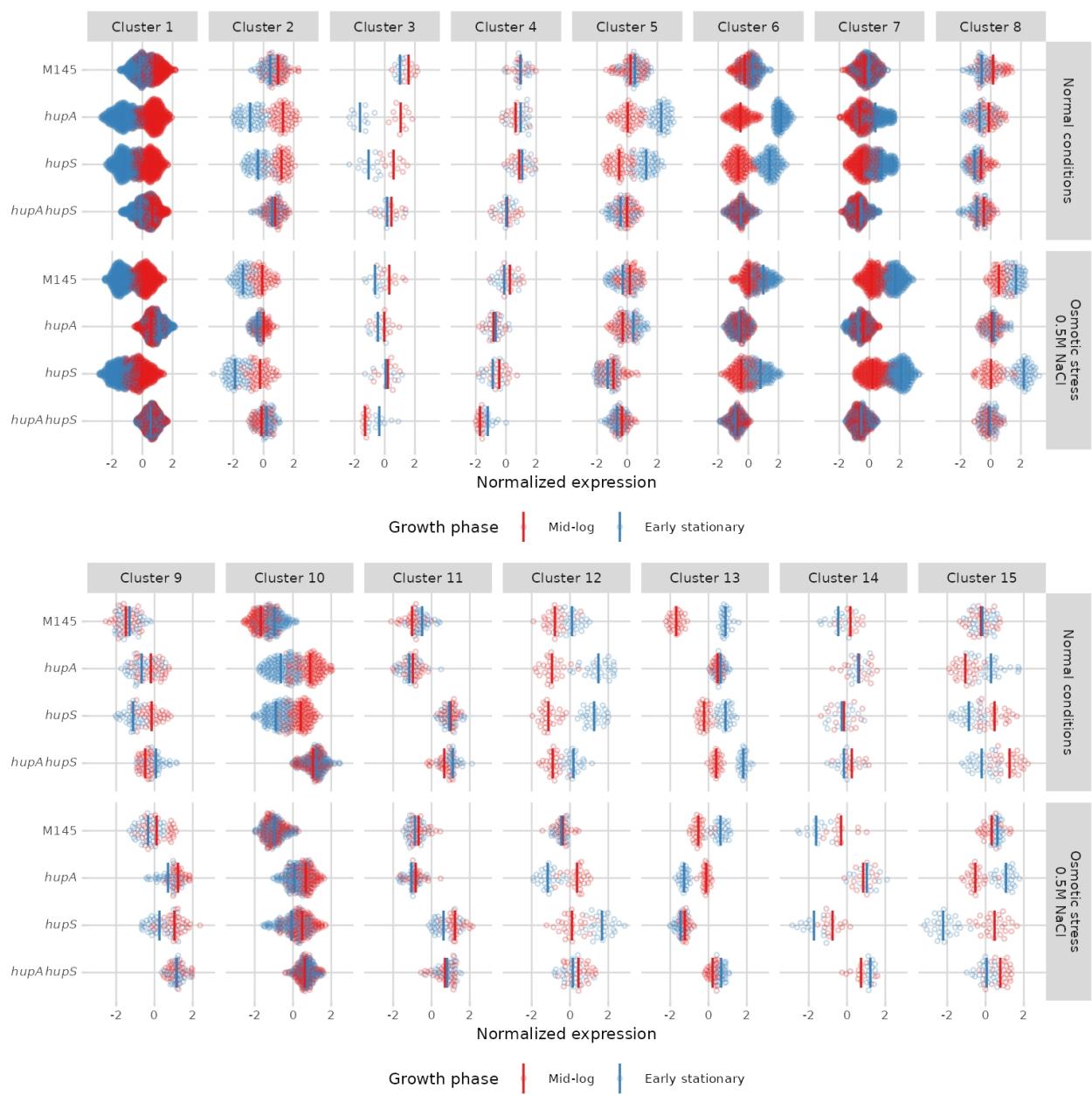
**A.****ArgR regulon****B.****OsdR regulon****Figure S6**



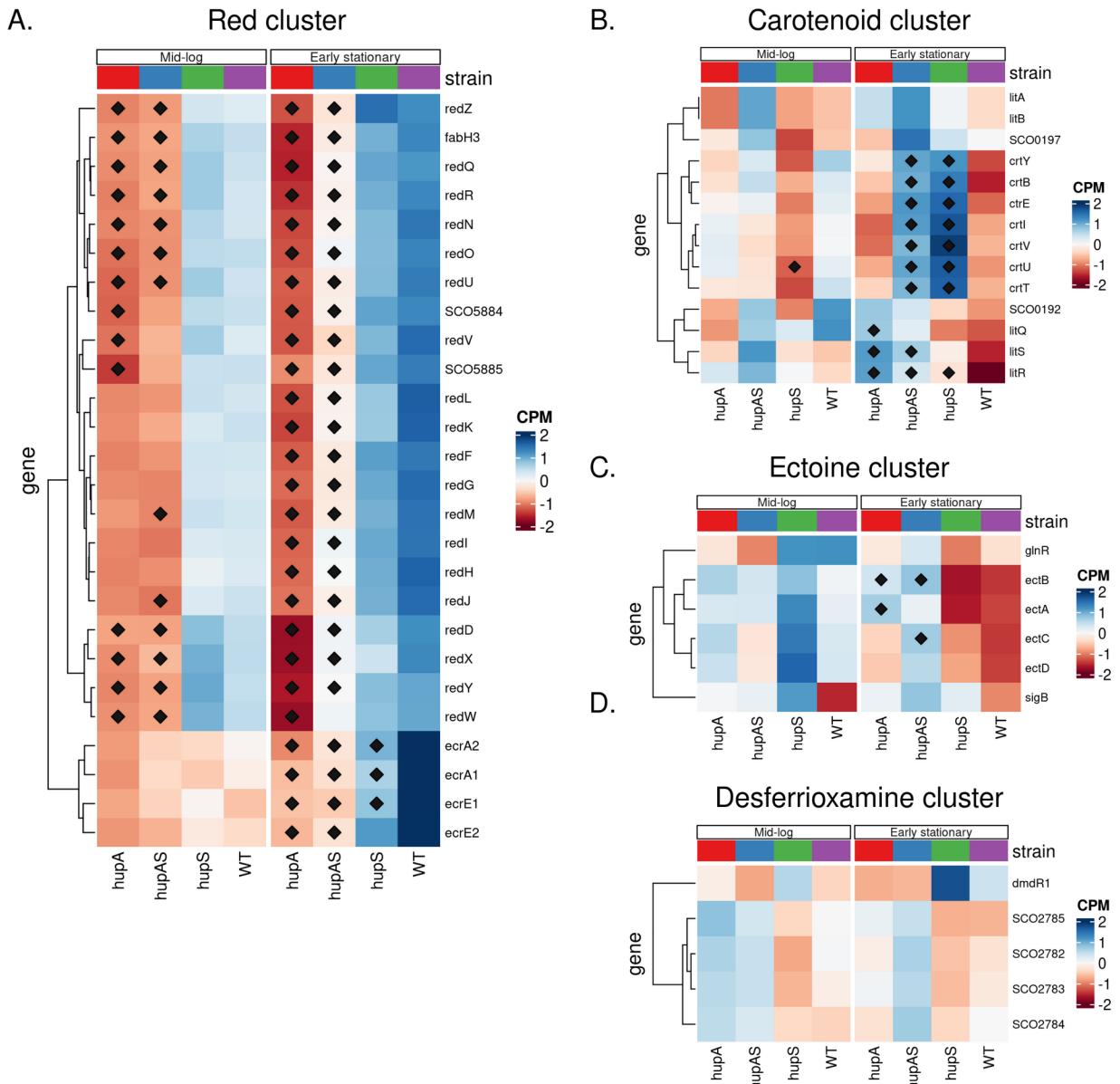
**Figure S7**



**Figure S8**



**Figure S9**



**Figure S10**