

1 **Table S1.** Primers used in this study

Primer ^a	Oligo sequence (5' to 3') ^b	Purpose ^c
Mutagenesis		
Pr1097	<u>gagatctaga</u> accgtttgatttttaattgataatg	Janus cassette For
Pr1098	gagact <u>cgagc</u> ctttccttatgcttttgac	Janus cassette Rev
Pr1836	<u>gagagtcgac</u> accgtttgatttttaattgataatg	Janus cassette For
Pr1837	gagac <u>catggc</u> ctttccttatgcttttgac	Janus cassette Rev
Pr2841	actacatcacaatggctgcc	<i>pdxT</i> upstream For
Pr2842	aaat <u>ctagacc</u> gattttcatctattttcc	<i>pdxT</i> upstream Rev
Pr2843	aaact <u>cgagttg</u> agattgaatttctcaac	<i>pdxT</i> downstream For
Pr2844	accagttttgctctatgctc	<i>pdxT</i> downstream Rev
Pr2845	gaagcctttgaacgtcttg	<i>pdxS</i> upstream For
Pr2846	aaat <u>ctagac</u> caccacctgagcatctg	<i>pdxS</i> upstream Rev
Pr2847	aaact <u>cgagt</u> tattaatgaaaatgaaatcc	<i>pdxS</i> downstream For
Pr2848	tggcctatggttcgaccac	<i>pdxS</i> downstream Rev
Pr2919	gcgatgatcttgcaagagggaat	<i>pdxR</i> upstream For
Pr2920	cgcg <u>ctcgac</u> acggatagaaggcaggcgactac	<i>pdxR</i> upstream Rev
Pr2921	cgcg <u>ccatggc</u> acagtaactgggcttggga	<i>pdxR</i> downstream For
Pr2922	aacactgacaagtaactggatg	<i>pdxR</i> downstream Rev
Complementation		
Pr2967	tttt <u>gtcgaca</u> actacatcacaatggctgc	<i>pdxS</i> promoter and ORF For
Pr2838	cggg <u>gatcc</u> gttctgcaaaggccccttgc	<i>pdxS</i> promoter and ORF Rev
Pr2968	tttt <u>gtcgaca</u> taaaaggccagccatcaagg	<i>pdxR</i> promoter and ORF For
Pr2969	cgcg <u>gatcc</u> gcagagttggatgctatcc	<i>pdxR</i> promoter and ORF Rev
Overexpression		
Pr2862	cgcg <u>gctagc</u> atgactgaaaatcggtatga	<i>pdxS</i> ORF For
Pr2863	cgcg <u>gatcc</u> cttgcaaggccaatattcc	<i>pdxS</i> ORF Rev
Pr2864	cgcg <u>catatg</u> aaaatcggaatattggc	<i>pdxT</i> ORF For
Pr2865	cgcg <u>gatcca</u> atacatcgctattgttta	<i>pdxT</i> ORF Rev
Pr2917	cgcg <u>ccatgg</u> agaaacaaagcaagtacaaagag	<i>pdxR</i> ORF For
Pr2918	cgcg <u>aagctt</u> tccaattctgcttttaaatagtttaaaca	<i>pdxR</i> ORF Rev
Heterogeneous expression in <i>E. coli</i> mutant		
JY161	cgcaatattaaaagttaacccttcgacc	<i>E. coli crp</i> promoter For
JY170	cgggctagcgggtacctctagacatgcgcttatcctctg	<i>E. coli crp</i> promoter Rev
Pr2835	cggg <u>aattc</u> atgaaaatcggaatattggc	<i>pdxT</i> ORF For
Pr2854	cgggctagcaagttgagaaatcaatctc	<i>pdxT</i> ORF Rev
Pr2855	cgg <u>tctaga</u> atgactgaaaatcggtatg	<i>pdxS</i> ORF For
Pr2838	cggg <u>gatcc</u> gttctgcaaaggccccttgc	<i>pdxS</i> ORF Rev
Northern blot		
Pr2926	cttagccactactggaactaac	<i>pdxS</i> probe
Pr2782	actttcactctcacactcg	16S rRNA probe

2 ^a*pdxST* double mutant was generated with *pdxS* upstream primers and *pdxT* downstream primers.

3 ^bThe restriction site is underlined if presents in an oligo.

4 ^cTwo pairs of primers flanking the Janus cassette were used for combination of different
5 restriction sites. For, forward primer; Rev, reverse primer.

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6 **FIGURE LEGENDS**

7 **Figure S1.** Western blot analyses to verify the deletion of *pdxT* and *pdxS*. WT (D39) and its
8 derivatives, including $\Delta pdxT$ (ST2675), $\Delta pdxS$ (ST2676) and $\Delta pdxST$ (ST2677), were initially
9 grown in complete CDM and then incubated for 4 h in depleted CDM. The protein samples were
10 analyzed with the indicated antibodies (right side). Blotting with antibody against SPD_1063
11 was served as a loading control.

12 **Figure S2.** Growth of bacterial strains on M9 plates in the presence or absence of 0.1 mM PLP.
13 ec022 is the *E. coli* WT strain. ec048 and ec053 are *E. coli* $\Delta pdxA$ and $\Delta pdxJ$, respectively.
14 ST2688 and ST2689 express pneumococcal *pdxS* in $\Delta pdxA$ and $\Delta pdxJ$, respectively. Strains
15 ec061 and ec065 contain the control vector in $\Delta pdxA$ and $\Delta pdxJ$, respectively. Images of plates
16 were taken after incubation for 18 h at 37 °C

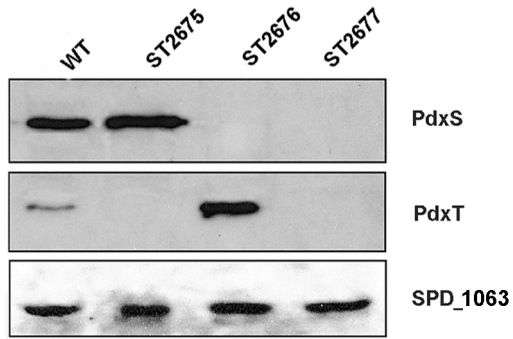
17 **Figure S3.** (A) Putative GntR family proteins in *S. pneumoniae*. All these proteins have a
18 wHTH motif as a signature of GntR family proteins. Three proteins have an UbiC transcription
19 regulator-associated (UTRA) domain. SPD_1225 has an AAT domain. (B) Verification of
20 deletion and complementation of *pdxS* and *pdxR*, respectively, using PCR. Internal primers of
21 *pdxR* and *pdxS* were used for PCR. Bacterial DNA samples isolated from the indicated strains,
22 including WT (D39), ST2726 ($\Delta pdxR$), ST2786 ($\Delta pdxR/pdxR+$), ST2787 ($\Delta pdxR$ /vector control),
23 ST2676 ($\Delta pdxS$), ST2784 ($\Delta pdxS/pdxS+$) and ST2785 ($\Delta pdxS$ /vector control), were used as
24 templates.

25 **Figure S4.** Growth curve of ST2676 (A) and ST2726 (B) in complete CDM. The bacterial
26 strains tested include WT (D39), ST2676 ($\Delta pdxS$), ST2784 ($\Delta pdxS/pdxS+$) and ST2785
27 ($\Delta pdxS$ /vector control), ST2726 ($\Delta pdxR$), ST2786 ($\Delta pdxR/pdxR+$), ST2787 ($\Delta pdxR$ /vector

28 control). The bacterial growth was monitored hourly at OD_{620} . Data shown are the means of
29 three repeat experiments. Error bars denote SEM.

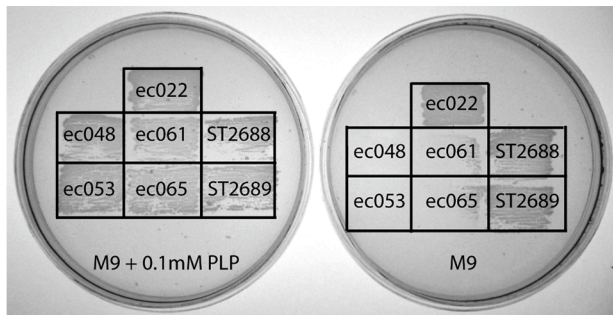
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30 FIG. S1



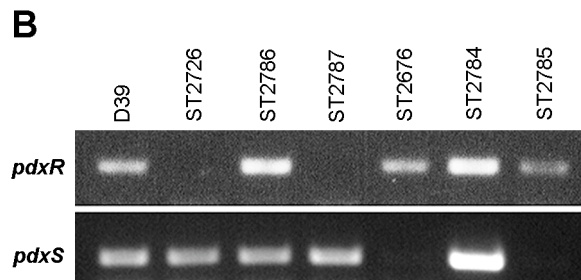
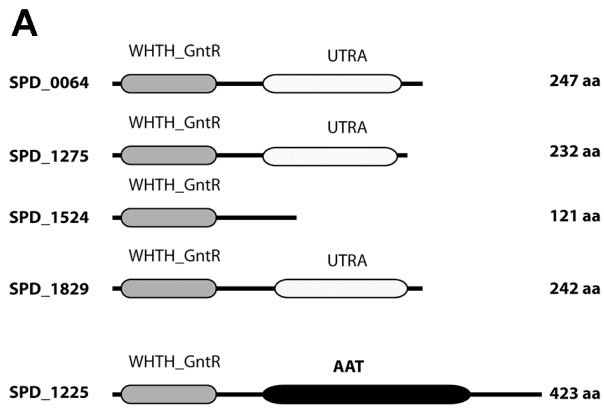
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32 FIG. S2



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34 FIG. S3



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