

SUPPLEMENTAL MATERIAL

Insight into the dual function of lipid phosphate phosphatase PgpB involved in two essential cell-envelope metabolic pathways in *Escherichia coli*

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Table S1. Plasmids

pMAK <i>bacA</i>	Cam ^R	4
p <i>TrcHis60</i>	Amp ^R	33
p <i>TrcHis30</i>	Amp ^R	33
p <i>TrcBac30</i>	p <i>TrcHis30</i> derivative for overproduction of N-His ₆ -BacA	3
p <i>TrcH60ybjG</i>	p <i>TrcHis60</i> derivative for overproduction of C-His ₆ -YbjG	This study
p <i>TrcH60lpxT</i>	p <i>TrcHis60</i> derivative for overproduction of C-His ₆ -LpxT	5
p <i>TrcH60pgpB</i>	p <i>TrcHis60</i> derivative for overproduction of C-His ₆ -PgpB	6
p <i>TrcH30ynbD</i>	p <i>TrcHis30</i> derivative for overproduction of N-His ₆ -YnbD	This study
p <i>TrcH60pgpA</i>	p <i>TrcHis60</i> derivative for overproduction of C-His ₆ -PgpA	This study
p <i>TrcH60pgpC</i>	p <i>TrcHis60</i> derivative for overproduction of C-His ₆ -PgpC	This study
pKD4	PCR amplification of Kan ^R cassette for gene deletion	31
pCP20	Resistance cassette removal by Flp recombinase expression	31
pKD46	Resistance cassette insertion by lambda red recombinase expression	31
pA727	p <i>TrcH60pgpB</i> variant with PgpB K97A mutation	This study
pA730	p <i>TrcH60pgpB</i> variant with PgpB ΔV100 mutation	This study
pA733	p <i>TrcH60pgpB</i> variant with PgpB R104A mutation	This study
pA737	p <i>TrcH60pgpB</i> variant with PgpB P105A mutation	This study
pA741	p <i>TrcH60pgpB</i> variant with PgpB A158S mutation	This study
pA745	p <i>TrcH60pgpB</i> variant with PgpB F159A mutation	This study
pA762	p <i>TrcH60pgpB</i> variant with PgpB P160A mutation	This study
pA766	p <i>TrcH60pgpB</i> variant with PgpB S161A mutation	This study
pA771	p <i>TrcH60pgpB</i> variant with PgpB G162A mutation	This study
pA749	p <i>TrcH60pgpB</i> variant with PgpB G162D mutation	This study
pA798	p <i>TrcH60pgpB</i> variant with PgpB H163A mutation	This study
pA802	p <i>TrcH60pgpB</i> variant with PgpB T164A mutation	This study
pB244	p <i>TrcH60pgpB</i> variant with PgpB H150A mutation	This study
pA807	p <i>TrcH60pgpB</i> variant with PgpB S200A mutation	This study
pA812	p <i>TrcH60pgpB</i> variant with PgpB R201A mutation	This study
pA817	p <i>TrcH60pgpB</i> variant with PgpB L202A mutation	This study
pA822	p <i>TrcH60pgpB</i> variant with PgpB G205A mutation	This study
pA828	p <i>TrcH60pgpB</i> variant with PgpB H207A mutation	This study
pA833	p <i>TrcH60pgpB</i> variant with PgpB W208A mutation	This study
pA838	p <i>TrcH60pgpB</i> variant with PgpB P209A mutation	This study
pA843	p <i>TrcH60pgpB</i> variant with PgpB D211A mutation	This study
pA847	p <i>TrcH60pgpB</i> variant with PgpB L212A mutation	This study
pA852	p <i>TrcH60pgpB</i> variant with PgpB A215G mutation	This study
pA857	p <i>TrcH60pgpB</i> variant with PgpB S219A mutation	This study
pB089	p <i>TrcH60pgpB</i> variant with PgpB R6A mutation	This study
pB093	p <i>TrcH60pgpB</i> variant with PgpB R7A mutation	This study

pB232	<i>pTrcH60pgpB</i> variant with PgpB H57A mutation	This study
pB115	<i>pTrcH60pgpB</i> variant with PgpB R69A mutation	This study
pB099	<i>pTrcH60pgpB</i> variant with PgpB R71A mutation	This study
pB103	<i>pTrcH60pgpB</i> variant with PgpB K73A mutation	This study
pB107	<i>pTrcH60pgpB</i> variant with PgpB R181A mutation	This study
pB111	<i>pTrcH60pgpB</i> variant with PgpB R182A mutation	This study
pB118	<i>pTrcH60pgpB</i> variant with PgpB R183 mutation	This study
pB121	<i>pTrcH60pgpB</i> variant with PgpB R181A-R182A mutations	This study
pB124	<i>pTrcH60pgpB</i> variant with PgpB R182A-R183A mutations	This study
pB127	<i>pTrcH60pgpB</i> variant with PgpB R232A mutation	This study
pB194	<i>pTrcH60pgpB</i> variant with PgpB Δ 236-254 (Δ C ^{ter}) deletion	This study
pB877	<i>pTrcH60pgpB</i> variant with PgpB E154A mutation	This study
pB881	<i>pTrcH60pgpB</i> variant with PgpB E154A–H163A mutations	This study

Table S2. Phosphatase activities in membranes of *E. coli* mutants.

Substrate	Relative activities (% of WT)	
	C ₅₅ -PP	PGP
BW25113	100 +/- 2	100 +/- 3
BW Δ <i>pgpB</i>	96 +/- 11	7 +/- 8
BW <i>pgpB</i> -single	7 +/- 3	73 +/- 20

The C₅₅-PP and PGP phosphatase activities of BW25113, BW Δ *pgpB* and BW*pgpB*-single strains were measured and normalized by the quantity of proteins in each membrane extracts. The relative activities as compared to the WT strain membrane extracts are indicated. Each value is the mean of three independent measurements.

Table S3. Quantitative RT-PCR analysis of *pgpB* transcripts.

Plasmid	IPTG	PgpB expression
		(ratio of chromosomal expression)
none	-	1
<i>pTrcH60pgpB</i>	-	33 +/- 6
	+	2003 +/- 362

BW25113 cells carrying either no plasmid or *pTrcH60pgpB* were grown in 2YT medium in the presence or not of 1 mM IPTG up to A_{600nm} = 0.5. The total RNA was extracted and quantitative RT-PCR analysis was performed using appropriate primers (Table S4) and reference genes for normalization.

Table S4. PrimersGene inactivation

Inact1- <i>ynbD</i>	ACGTTTTTCGATGGCGGTGCGTCGTGATAACTGAACGTCGGAACGT	<i>ynbD</i> inactivation
Inact2- <i>ynbD</i>	TAGATATTCAGTCCACATCTCAATCCACTTACCTGTTTTCCATA	

Gene cloning^a

<i>ybjGNcoI</i>	GCGCCCATGGTGGAAAATTTGAATCTCTCTCTATTC	Cloning of <i>ybjG</i> in pTrcHis60
<i>ybjGBglII</i>	CTTTTTAGCAGAGATCTGTACGCACCCAGCC	
<i>ynbDBamHI</i>	GCGCGGATCCCTACAAGGCGCTGGCTGGTTATTGTTGC	Cloning of <i>ynbD</i> in pTrcHis30
<i>ynbDHindIII</i>	GCGCAAGCTTTTACCTGTTTTCCATAATCTCAGC	
<i>pgpBNcoI</i>	GCGCCCATGGCCATGCGTTCGATTGCCAGACGTACC	Cloning of <i>pgpB</i> in pTrcHis60
<i>pgpBBglII</i>	GCGCAGATCTACTTTCTTGTCTCGTTGCGCTAT	
<i>pgpBBglII-ΔC^{ter}</i>	CGCGAGATCTTGGCCCAAAATTCGTTGCGC	Cloning of <i>pgpB</i> ΔC ^{ter} in pTrcHis60
<i>pgpANcoI</i>	GCGCCCATGGTGACCATTTTGCCACGCCATAAAG	Cloning <i>pgpA</i> in pTrcH60
<i>pgpABglII</i>	GCGCAGATCTCGACAGAATACCCAGCGGCCAGTG	
<i>pgpCNcoI</i>	GCGCCCATGGTGGCAACTCACGAGCGTCGTGTGGTG	Cloning of <i>pgpC</i> in pTrcHis60
<i>pgpCBglII</i>	GCGCAGATCTTTCCAGTTGCTGGAGTTCACCGCG	

pgpB site-directed mutagenesis^b

K97A	GTAAATCCTGGATCGCAGACAAAGTCCAGG
ΔV100	CCTGGATCAAAGACAAACAGGAACCACGACC
R104A	CAAAGTCCAGGAACCAGCACCTTTTGTATCTGG
P105A	GTCCAGGAACCACGAGCTTTTGTATCTGGCTG
E154A	CACACTGGCAGAAAGCGACGGGGTTTGCCTTTC
A158S	GAAAGAGACGGGGTTTAGCTTTCCTTCCGGTC
F159A	GAGACGGGGTTTGCCGCTCCTTCCGGTCACAC
P106A	CGGGGTTTGCCTTTGCTTCCGGTCACACG
S161A	GGGTTTGCCTTTCCTGCCGGTCACACGATG
G162A	GTTTGCCTTTCCTTCCGCTCACACGATGTTTG
G162D	GTTTGCCTTTCCTTCCGATCACACGATGTTTG
H163A	GCCTTTCCTTCCGGTGCCACGATGTTTGCTGC
T164A	CTTTCCTTCCGGTCACGCGATGTTTGCTGCC
S200A	CGGGAGTCATGGGAGCCCGCCTGCTGCTCGG
R201A	GGAGTCATGGGAAGCGCCCTGCTGCTCGGG
L202A	GTCATGGGAAGCCGCGCTGCTCGGGATGC
G205A	GCCGCCTGCTGCTCGCGATGCATTGGCCACGC
H207A	CTGCTGCTCGGGATGGCTTGGCCACGCGATCTG
W208A	CTGCTCGGGATGCATGCGCCACGCGATCTGG
P209A	CTCGGGATGCATTGGGCACGCGATCTGGTAG
D211A	GCATTGGCCACGCGCTCTGGTAGTAGCTACG
L212A	CATTGGCCACGCGATGCGGTAGTAGCTACGTTG
A215G	GCGATCTGGTAGTAGGTACGTTGATTTTCGTG

S219A	GTAGCTACGTTGATTGCGTGGGCGCTGGTGG
R6A	GCGTTCGATTGCCGCACGTACCGCAGTGG
R7A	GTTCGATTGCCAGAGCTACCGCAGTGGGAG
H57A	GGCGTCATTACAGCTTTGATTTTATTCGG
R69A	GTTTCTCTGGTGTCTGGCTTTTCGCATTAAGG
R71A	GGTGTCTGCGTTTTGCCATTAAGGCTGCC
K73A	CTGCGTTTTTCGCATTGCGGCTGCCTTTGTAT
R181A	GTTTGCTGTGGCCGGCTCGGCGAACGTTAAC
R182A	GCTGTGGCCGCGTGC GCGAACGTTAAC
R183A	GTGGCCGCGTCGGGCAACGTTAACCATTG
R181-182A	GGTTTGCTGTGGCCGGCTGCGCGAACGTTAACCAT
R182-183A	GCTGTGGCCGCGTGC GCGAACGTTAACCATTGC
R232A	CGTGGCTTGC GCAAGCAATTTGTGGGCC

Quantitative PCR

198-pgpB-F	GTGTCTGCGTTTTTCGCATTA
320-pgpB-R	ACAAAAGGTCGTGGTTCCTG
394-gyrA-F	TGAACGGTTCTTCCGGTATC
635-gyrA-R	ATGTGTTCCATCAGCCCTTC
1163-ffh-F	GCGCTAAGCCAGAAATCATC
1304-ffh-R	ATCCGCCCTTCTTCATTTT
284-rrsA-F	CAGCCCACTGGAACTGAGA
400-rrsA-R	CCGAAGGCCTTTTTCATACA

^aThe restriction site used for the insertion of the amplicon in the expression vector is indicated in the primer's name and it is underlined in the sequence. ^bOf the two primers used for site-directed mutagenesis, only one sequence is indicated, the other sequence being its reverse complement.

Table S5. Functional complementation of thermosensitive strains by PgpB variants

Plasmid	protein	BWTetra-TsbacA		BWPGPTs	
		- IPTG	+ IPTG	- IPTG	+ IPTG
pTrcHis60	none	-	-	-	-
pTrcH60pgpB	PgpB	+	+	+	+
N-terminus					
pB089	R6A	+	+	+	+
pB093	R7A	-	+	+	+
TM 2					
pB232	H57A	+	+	+	+
Cytoplasmic loop 1					
pB115	R69A	+	+	+	+
pB099	R71A	+	+	+	+
pB103	K73A	+	+	+	+
Cytoplasmic loop 2					
pB107	R181A	+	+	+	+
pB111	R182A	+	+	+	+
pB118	R183	+	+	+	+
pB121	R181A-R182A	+	+	+	+
pB124	R182A-R183A	+	+	+	+
C-terminus					
pB127	R232A	+	+	+	+
pB194	$\Delta 236-254(\Delta C_{\text{ter}})$	+	+	+	+

The plasmids carrying a copy of *pgpB* (wild-type or variants) were tested for their ability to restore the growth at 42°C of thermosensitive strains BWTetra-TsbacA and BWPGPTs. +, normal growth at 42°C; - no growth at 42°C.

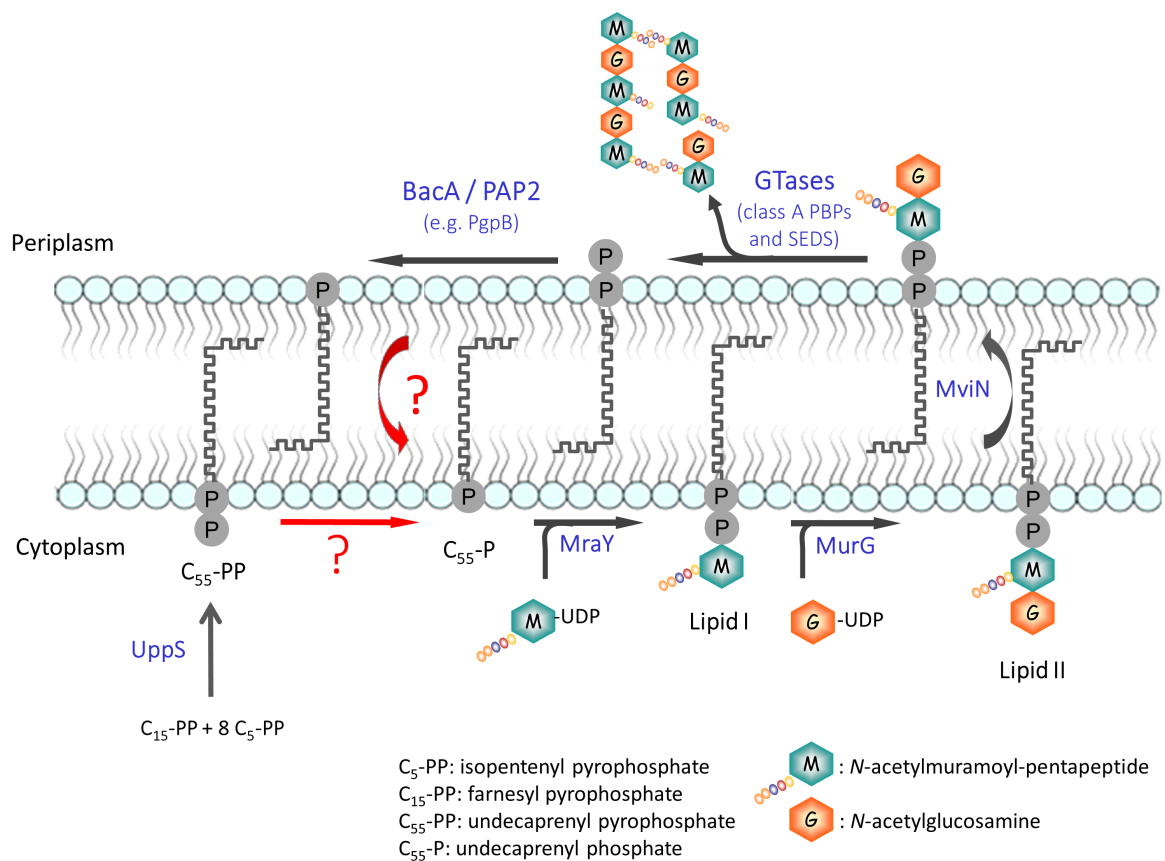


Figure S1. Schematic representation of C₅₅-P metabolism and peptidoglycan biosynthesis. The C₅₅-PP is *de novo* synthesized by UppS at the cytosolic face of the membrane; thereafter it is dephosphorylated, by a yet unknown enzyme, to yield the C₅₅-P lipid carrier. The MraY and MurG enzymes catalyze the successive transfers of phospho-*N*-acetylmuramoyl-pentapeptide and *N*-acetylglucosamine moieties from nucleotide precursors to C₅₅-P, yielding lipid II. The lipid II is flipped to the periplasmic face of the membrane, where glycosyltransferases (GTases) transfer the glycan-peptide moiety to the nascent polymer. GTases release C₅₅-PP as a by-product, which is then recycled through a dephosphorylation step by BacA or PAP2 enzymes and a flip back to the inner side, by a yet unknown mechanism, to enter a new cycle of peptidoglycan biosynthesis.

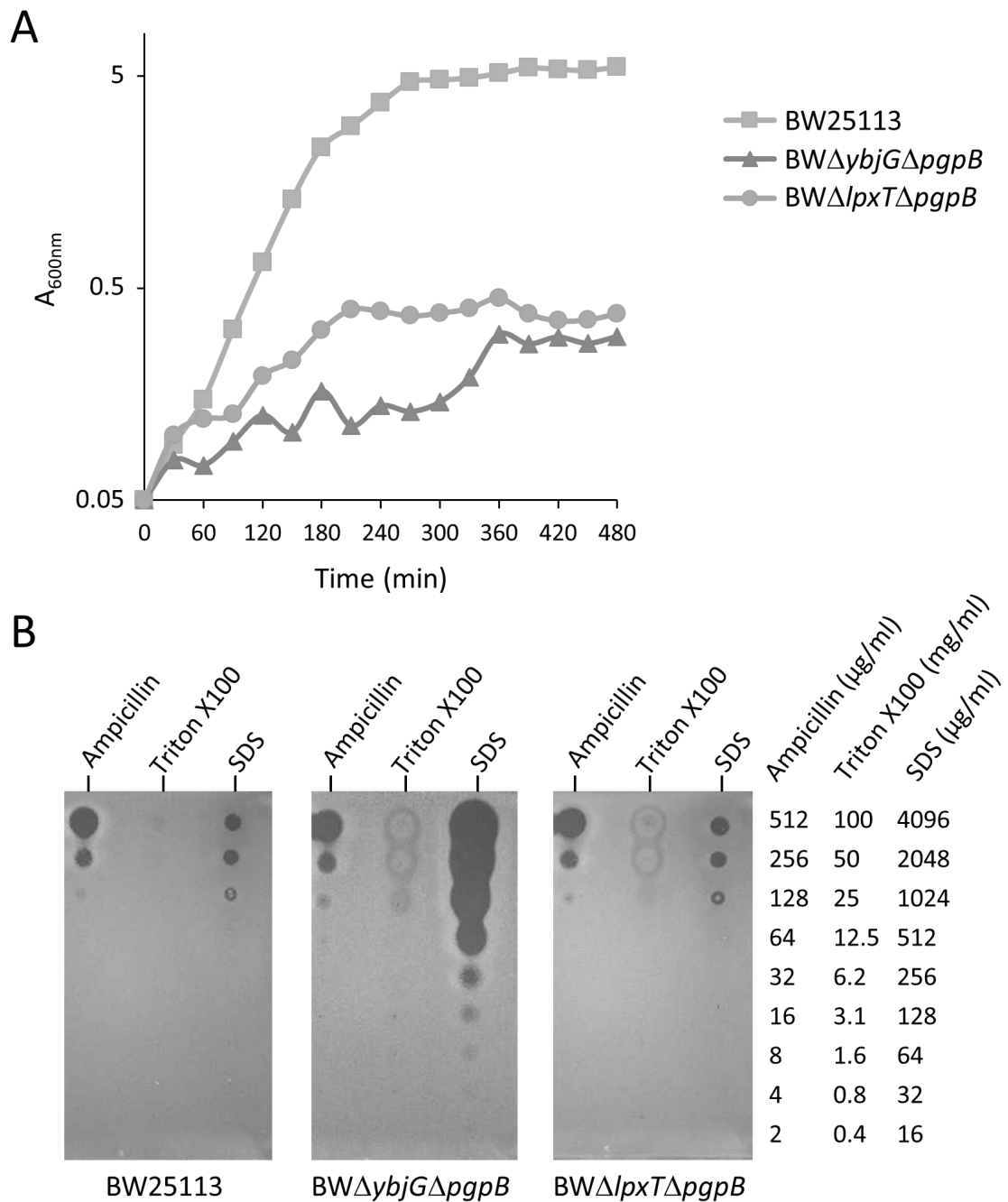


Figure S2. Multiple inactivation of PAP2s differentially sensitizes the cells to small hydrophobic compounds. (A) Cell growth was monitored at 37°C in 2YT medium containing 20 mg/ml of DOC. (B) 2YT-agar plates were overlaid with the indicated strains and 5 μl of serial dilutions (as indicated on the left) of ampicillin, Triton X100 or SDS were dropped on these plates.

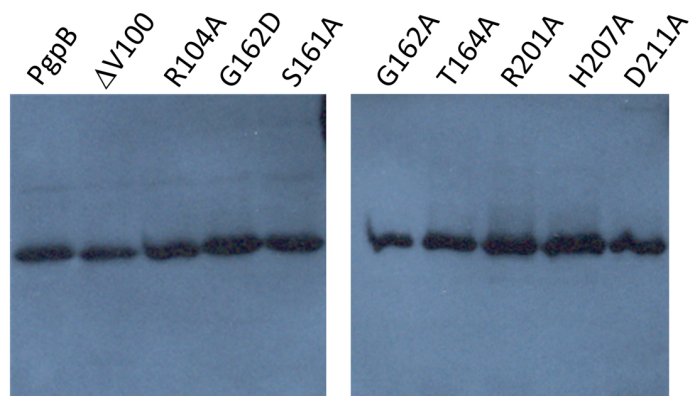


Figure S3. Expression of PgpB WT and variants presenting a default of complementation. Immunoblotting analysis of membrane fractions isolated from BW25113 cells producing PgpB and its variants from the *pTrcHis60* vector. Total DDM-solubilized membrane proteins (5 μ g) were separated by SDS-PAGE and analyzed by immunoblotting with anti-His antibodies.

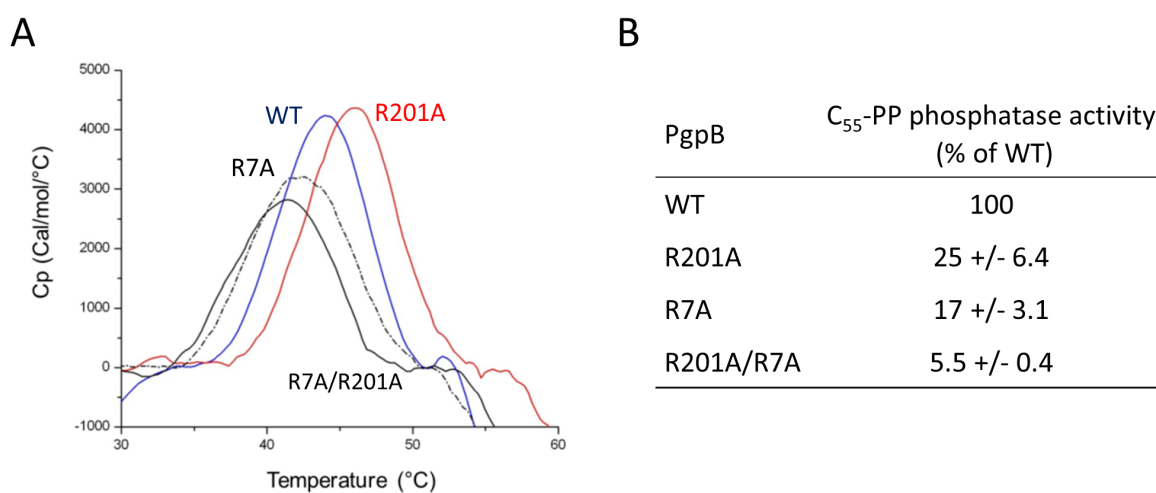


Figure S4. Thermal stability and activity of PgpB variants. (A) DSC measurements were performed with proteins at a concentration of 0.5 mg/ml in potassium phosphate buffer, pH 6.0. (B) The C_{55} -PP phosphatase activity of the variants is expressed as a percentage of the WT PgpB activity.