

Supporting information to

Non-competitive binding of PpiD and YidC to the SecYEG translocon expands the global view on the SecYEG interactome in *E. coli*

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File content:

Table S1 (separate data file): Complete list of putative SecYEG interacting proteins identified by label free proteomics.

Table S2: List of oligonucleotides used for PCR reactions

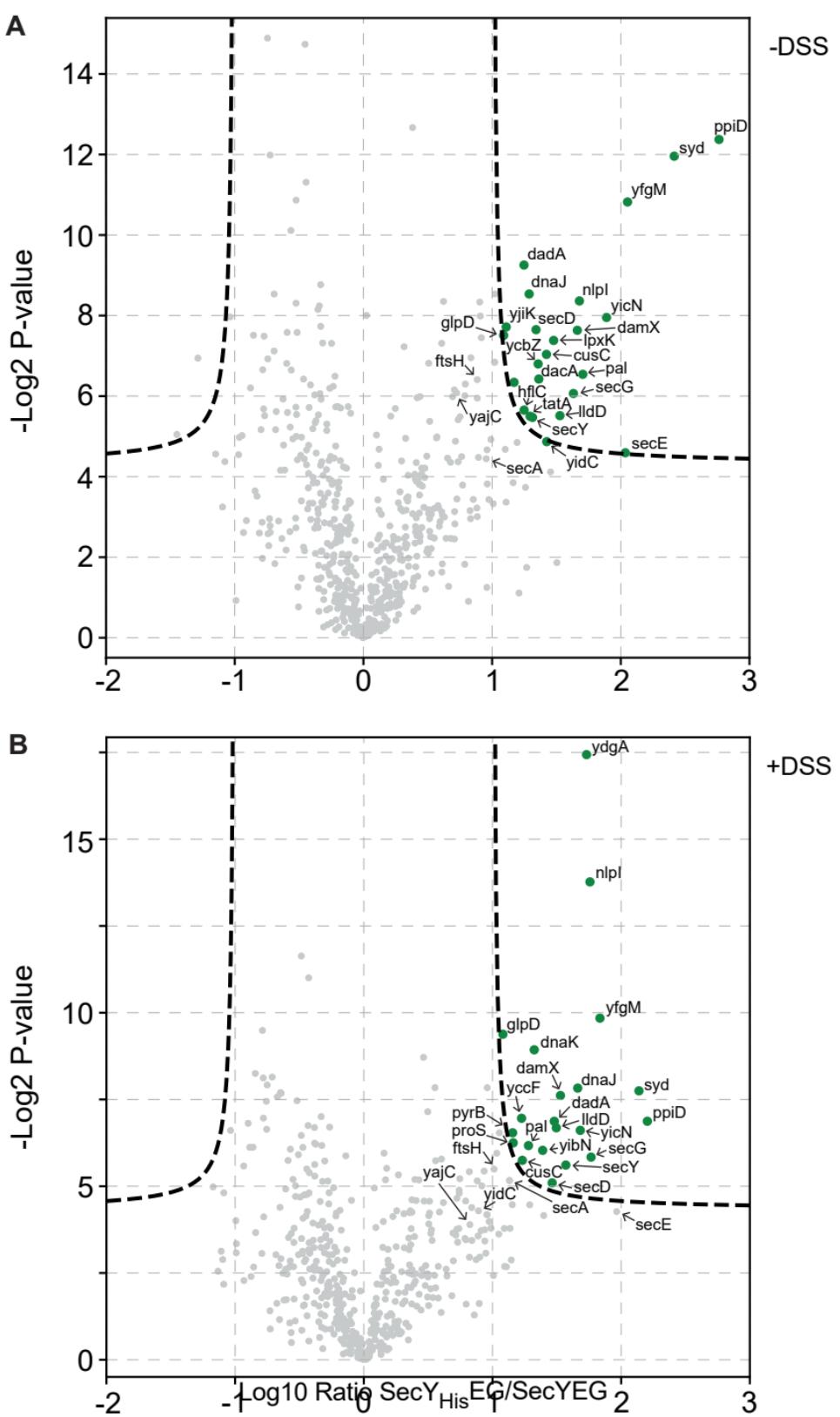
Figure S1: Significant interactors of the SecYEG in *E. coli*.

Figure S2: PpiD antibodies recognize both PpiD and YfgM

Table S1 (separate Data file) : Complete List of putative SecYEG interacting proteins identified by label-free proteomics with and without DSS-cross-linking. When indicated, YidC was co-expressed with SecY_{His}EG.

Table S2: List of oligonucleotides used for PCR reactions.

Primer name	5' – oligonucleotides – 3'	Purpose of use
SecY_plug_removal_for	GCTCTCAGCCGTGCTTCTATCTTGCTC	Removal of SecY plug domain
SecY_plug_removal_rev	GATGGTGCCTCGCTGTTGCTCAAGC	
YidC_cutpBadlong_for	CGCGATCTAGAGGGCTAGCAGGAGGAATTAC	Cloning of pTrc99a-SecY _{His} EG-YidC
YidC_cutpBadlong_rev	CGGTCGTCGACTCAGGATTTTCTTCGCG	
YidC_HiR10_for	GGCAGCGGAGATT CGC	Cloning of pTrc99a-SecY _{His} EG-YidC
YidC_HiR10_rev	CATCGTTAGTGTCTCCATGGTG	
SecY_I61_for	ACCATCTAGGAGATGTTAAC	Construction of SeY plug (61-70) - pBpA mutants
SecY_I61_rev	GCCTCGCTGTTGCTC	
SecY_E62_for	CGAGGCACCATCATTAGATGTTAACATGTT	
SecY_E62_rev	GAACATGTTAACATCTAAATGATGGTGCCTC	
SecY_M63_for	GAGGCACCATCATTGAGTAGTTAACATGTTCTCTGG	
SecY_M63_rev	CCAGAGAACATGTTAAACTACTCAATGATGGTGCCTC	
SecY_F64_for	CACCATCATTGAGATGTAGAACATGTTCTCTGGTG	
SecY_F64_rev	CACCAAGAACATGTTCTACATCTCAATGATGGTG	
SecY_N65_for	CATCATTGAGATGTTTAGATGTTCTCTGGTGGTG	
SecY_N65_rev	CACCACCAGAGAACATCTAAAACATCTCAATGATG	
SecY_M66_for	CATTGAGATGTTAACTAGTTCTCTGGTGGTGCTC	
SecY_M66_rev	GAGCACCACCAGAGAACATGTTAACATCTCAATG	
SecY_F67_for	AACATGTTAGTCTGGTGGTGCT	
SecY_F67_rev	AAACATCTCAATGATGGTGCC	
SecY_S68_for	GATGTTAACATGTTCTAGGGTGGTGCTCTCAGCC	
SecY_S68_rev	GGCTGAGAGCACCAACCCTAGAACATGTTAACATC	
SecY_G69_for	GTTAACATGTTCTCTAGGGTGGCTCTCAGCCGTG	
SecY_G69_rev	CACGGCTGAGAGCACCAACCCTAACAGAACATGTTAAC	
SecY_G70_for	CATGTTCTCTGGTTAGGCTCTCAGCCGTG	
SecY_G70_rev	CACGGCTGAGAGCCTAACAGAACATG	



Supporting Figure 1: Significant interactors of SecYEG in *E. coli*. (A) SecYEG or SecYHisEG were expressed in *E. coli*, and subsequently, SecYEG/SecYHisEG were purified via metal affinity chromatography and subjected to digestion using trypsin followed by LC-MS analyses. Volcano plot representing negative logarithmic p-values from the t-test across replicate experiments ($n=4$) plotted against the mean log₁₀ intensity ratio of the respective protein in the SecYHisEG (His) / SecYEG (Ctl) pull-downs. A hyperbolic curve (black dashed line) separates specific interactors of the SecYEG complex marked by green dots from unspecifically copurified proteins (grey dots). The curves are defined by a bend of 0.25, p-value threshold of 0.05 and log₁₀ ratio limit of 10. Parameters were chosen such that a false positive rate <0.01 was obtained with the left branch (ratios <1) in (A) and (B). (B) As in (A) but protein-protein-interactions were stabilized by on-column cross-linking using the chemical cross-linker DSS. A complete list of all proteins is given in Supporting Table S1.

MC4100 (wt)	$\Delta ppiD$	$\Delta yfgM$	$\Delta ppiD/yfgM$
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kDa

PpiD →

-70

-55

-25

-15

YfgM →



Supporting Figure 2: PpiD antibodies recognize both PpiD and YfgM. Cells of the indicated *E. coli* strains were TCA precipitated and after SDS-PAGE/Western-blotting decorated with α -PpiD antibodies. Indicated are PpiD and YfgM and the corresponding gel areas are identical to those displayed in Fig. 5F