Table S1. Bacterial strains and plasmids used in this study.

Name	Description	Reference
P. aeruginosa		
PA01	Wild-type P. aeruginosa	[1]
ΔyfiR	PA01 with <i>yfiR</i> deleted	[2]
ΔyfiNR	PA01 with <i>yfiNR</i> deleted	[2]
ΔyfiBNR	PA01 with <i>yfiBNR</i> deleted	[2]
ΔyfiBNR Tn7::yfiNR	ΔyfiBNR with yfiNR genes inserted at the Att::Tn7 locus	[2]
PA01 yfiR-M2	ΔyfiR with yfiR-flag inserted at the Att::Tn7 locus	[2]
ΔyfiBN yfiR-M2	ΔyfiBNR with yfiR-flag inserted at the Att::Tn7 locus	[2]
ΔyfiN yfiR-M2	ΔyfiNR with yfiR-flag inserted at the Att::Tn7 locus	[2]
PA01 Tn7::yfiRN-flag	PA01 with yfiRN-flag inserted at the Att::Tn7 locus	This study
ΔyfiNR Tn7::yfiRN-flag	ΔyfiNR with yfiRN-flag inserted at the Att::Tn7 locus	This study
ΔdsbA	PA01 with <i>dsbA</i> deleted	This study
ΔdsbA ΔyfiBNR	ΔyfiBNR with dsbA deleted	This study
ΔdsbA ΔyfiB	ΔdsbA ΔyfiBNR with yfiNR genes inserted at the Att::Tn7 locus	This study
wspR::Tn5	Tn5 insertion in wspR	This study
ΔdsbA wspR::Tn5	ΔdsbA with Tn5 insertion in wspR	This study
Clin110	SCV strain isolated from CF sputum	[2]
Clin163	Smooth strain isolated from CF sputum. Descendent of Clin110	This study
Δ <i>yfiBNR</i> Tn <i>7</i> :: <i>yfiBNR</i>	ΔyfiBNR with yfiBNR genes inserted at the Att::Tn7 locus. yfiN present	This study
Clinical yfiN alleles	in four alleles containing combinations of G173D and $\Delta$ 255-7 mutations	
SCV20265	SCV strain isolated from CF sputum	[3]
WT20265	Smooth precursor of SCV20265, isolated from CF sputum	[3]
E. coli		
DH5α	endA1, hsdR17( $r_K$ - $m_K$ +), supE44, recA1, gyrA (Nal $^r$ ), relA1, $\Delta$ (laclZYA-	[4]
	argF)U169, deoR, Φ80dlacΔ(lacZ)M15	
DY330	W3110 ΔlacU169 gal490 λcl857 Δ(cro-bioA)	[5]
XL1-red	F- endA1 gyrA96(nalR) thi-1 relA1 lac glnV44 hsdR17(rK- mK+) mutS	Stratagene
	mutT mutD5 Tn10	
Plasmids		
pME6032	Tet <sup>R</sup> , P <sub>K</sub> , 9.8 kb pVS1 derived shuttle vector	[6]
pME6032-Gm	Gm <sup>R</sup> , pME6032 based vector, Tet cassette replaced with <i>aacCl</i>	This study
	gentamycin resistance cassette	
pME- <i>araC</i>	Tet <sup>R</sup> , pME6032 with the lac-promoter and <i>lacI</i> replaced with the <i>araC</i> -	[2]
	pBAD promoter fragment from pBAD18s	
pME- <i>araC-yfiR</i>	Tet <sup>R</sup> , pME- <i>araC</i> containing YfiR under control of ara-promoter as EcoRI-	[2]
	BgIII fragment	
pME6032- <i>yfiB</i>	pME6032 with <i>yfiB</i> as <i>EcoRI-BgII</i> fragment	[2]
pBV-PA5295	pBV-MCS4 with PA5295 under the control of vanillate promoter	[2]

pBV-PA5295 <sub>E328A</sub>	pBV-MCS4 with PA5292 <sub>E328A</sub> under the control of vanillate promoter	[2]
Activated yfiB mutants	20 pME6032-yfiB variants with activating mutations in yfiB (see Table 3)	This study
pGm- <i>yfiprom-N</i>	pME6032-Gm with <i>yfiN</i> as <i>Bam</i> HI-blunt fragment, <i>yfiN</i> expressed from	This study
	the native promoter	
pGm-yfiprom-N -	pME6032-Gm with yfiN alleles as BamHI-Ncol fragments, containing	This study
constructs	single point mutations	
YfiR-insensitive YfiN	20 pGm-yfiprom-N variants with YfiR-insensitive mutations in yfiN (see	This should
mutants	Table 1 and Fig.7C)	This study
pMR- <i>yfiR-flag</i>	pMR20-yfiR-M2 with lac promoter replaced by FRT scar sequence	This study
Compensatory YfiR	14 pMR-yfiR-flag variants with compensatory mutations in yfiR (see	This study
mutants	Table 2 and Fig. 7C)	This study
pME- <i>yfiB-PG</i> -	pME6032- <i>yfiB</i> with <i>yfiB</i> D102A,G105A (± F48S, L43P) as <i>Eco</i> RI- <i>Asc</i> I	This street.
constructs	fragments	This study
pME- <i>yfiB-LA</i> -	pME6032 containing yfiR residues 1-42 fused to yfiB (± D102A,G105A)	This study
constructs	residues 26-169 as <i>EcoRI-BgII</i> fragment	This study
pME- <i>yfiB</i> short/long	nMFC022 with ufile short/long linker allales as FeeDI Ball fragments	-1:
constructs	pME6032 with <i>yfiB</i> short/long linker alleles as <i>EcoRI-BgII</i> fragments	This study
nNAE will constructs	pME6032 with yfiB alleles as EcoRI-Bg/II fragments, containing single	This study
pME- <i>yfiB</i> constructs	point mutations	
pME-dsbA	pME6032 with dsbA as EcoRI-KpnI fragment	This study
pME3087	Tet <sup>R</sup> , suicide vector for allelic replacement; <i>ColE1</i> -replicon, <i>IncP-1</i> , <i>Mob</i>	[7]
pME3087-Δ <i>dsbA</i>	pME3087 with dsbA deletion cassette as HindIII-BamHI fragment	This study
pMR20	Tet <sup>R</sup> , low copy number and broad host-range vector	[8]
pMR20- <i>yfiBNR</i>	pMR20 with <i>yfiBNR</i> as <i>Hind</i> III- <i>Bam</i> HI fragment	[2]
pMR20- <i>yfiR-M2</i>	pMR20 with <i>yfiR-flag</i> as <i>Hind</i> III- <i>Bam</i> HI fragment	[2]
pMR20- <i>yfiR-M2</i> -	pMR20 with yfiR-flag alleles as HindIII-BamHI fragments, containing	This study
constructs	single point mutations	
pMR20-yfiRN-flag	pMR20 with yfiRN-flag as HindIII-BamHI fragment	This study
pUC18T-mini-Tn7T-Gm	Amp <sup>R</sup> , Gm <sup>R</sup> , Tn7 insertion vector	[9]
pTn <i>7-yfiR-M2</i>	pUC18T-mini-Tn7T with yfiR-flag as HindIII-BamHI fragment	[2]
pTn <i>7-yfiRN-flag</i>	pUC18T-mini-Tn7T with <i>yfiRN-flag</i> as <i>Hind</i> III- <i>Bam</i> HI fragment	This study
pTn <i>7-yfiBNR</i>	Vector for complementation of entire yfiBNR operon at Tn7 locus	[2]
.T. 7 . S.D.V. C4.72D. D.	Vector for complementation of yfiBNR operon with yfiN G173D	This study
pTn <i>7-yfiBN-G173D-R</i>	mutation	
pTn <i>7-yfiBN-Δ255-7-R</i>	Vector for complementation of <i>yfiBNR</i> operon with <i>yfiN</i> Δ255-7	This study
	mutation	
pTn <i>7-yfiBN20265-R</i>	Vector for complementation of <i>yfiBNR</i> operon with SCV20265 <i>yfiN</i>	This study
	allele	
pTNS2	Amp <sup>R</sup> , helper plasmid for Tn7 integration events	[9]
pFLP2	Amp <sup>R</sup> , FRT cassette excision vector	[10]

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