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Strain	Genotype	Reference
BT5482	WT *B. thetaiotaomicron* (gentamycin^R)	Lab stock.
BT5482 Δ*tdk*	As BT5482 with Δ*tdk*	(Koropatkin et al., 2008)
SM135 AKA HPX$^-$	As BT5482 Δ*tdk* Δ*katE1* Δ*ahpC1* Δ*rbr1-1* Δ*rbr2-1*	(Mishra & Imlay, 2013)
WM6026	*E. coli laclq*, *rrnB3, DElacZ4787, hsdR514, DE(araBAD)567, DE(rhaBAD)568, rph-1 att-lambda::pAE12-del (oriR6K/cat::frt5), del 4229(dapA)::frt, del(endA)::frt, uidA(delMlul)::pir(wt), attHK::pJK1006::del1/2(del oriR6K-cat::frt5, del trfA::frt)*	(Blodgett, 2007)
BW19851	*E. coli rp4-2(Km::Tn7, Tc::Mu-1) ΔuidA::pir+ recA1 hsdR17 creC510 endA1 thiE1*	(Metcalf, 1994)
Top10	*F– mcrA Δ(mrr-hsdRMS-mcrBC) φ80lacZΔM15 ΔlacX74 recA1 araD139 Δ(araA-leu)7697 galU galK rpsL endA1 nupG*	Invitrogen #V43001
MK462	As TOP10 with pBAD/His A/*pfor* of *B. thetaiotaomicron*	This study
MK466	As BW19851 with pExchange-*pfl2955* (BT_2955)	This study
MK470	As BW19851 with pExchange-*fdx* (BT_2414))	This study
MK474	As BW19851 with pExchange-*pfor* (BT_1747)	This study
MK482	As BW19851 with pExchange-*rnf* (BT_0616 to BT_0622 in *rnf* operon)	This study
MK486	As BW19851 with pExchange-*hyd3472* (BT_3472)	This study
MK490	As BW19851 with pExchange-*hyd1834* (BT_1834)	This study
MK494	As BT5482 Δ*tdk* with Δ*pfl2955* (BT_2955)	MK466 X BT5482 Δ*tdk*
MK500	As BT5482 Δ*tdk* with Δ*hyd3472* (BT_3472)	MK486 X BT5482 Δ*tdk*
MK508	As BT5482 Δ*tdk* with Δ*pfor* (BT_1747)	MK474 X BT5482 Δ*tdk*
MK520	As BT5482 Δ*tdk* with Δ*rnf* (BT_0616 to BT_0622 in *rnf* operon)	MK482 X BT5482 Δ*tdk*
MK526	As BW19851 with pExchange-*pfl4738* (BT_4738)	This study
MK532	As BT5482 Δ*tdk* with Δ*fdx* (BT_2414)	MK470 X BT5482 Δ*tdk*
MK540	As BT5482 Δ*tdk* with Δ*hyd1834* (BT_1834)	MK622 X BT5482 Δ*tdk*
MK550	As BT5482 Δ*tdk* with Δ*pfl4738* (BT_4738)	MK526 X BT5482 Δ*tdk*
MK556	As SM135 with Δ*pfl4738* (BT_4738)	MK526 X SM135
MK562	As SM135 with Δ*pfl2955* (BT_2955)	MK466 X SM135

MK570	As SM135 with Δ*hyd1834* (BT_1834)	MK490 X SM135
MK592	As SM135 with Δ*pfor* (BT_1747)	MK474 X SM135
MK580	As SM135 with Δ*hyd3472* (BT_3472)	MK486 X SM135
MK584	As SM135 with Δ*rnf* (BT_0616 to BT_0622 in *rnf* operon)	MK482 X SM135
MK616	As WM6026 with pExchange-*fdx* (BT_2414)	This study
MK624	As WM6026 with pExchange-*pfor* (BT_1747)	This study
MK626	As WM6026 with pExchange-*pfl4738* (BT_4738)	This study
MK628	As WM6026 with pExchange-*pfl2955* (BT_2955)	This study
MK630	As MK508 with Δ*pfl2955* (BT_2955)	MK628 X MK508
MK635	As MK550 with Δ*pfor* (BT_1747)	MK624 X MK550
MK644	As MK494 with Δ*pfl4738* (BT_4738)	MK626 X MK494
MK654	As SM135 with Δ*fdx* (BT_2424)	MK616 X SM135
MK622	As WM6026 with pExchange-*hyd1834* (BT_1834)	This study
LZ01 AKA SOD-	As BT5482 Δ*tdk* with Δ*sod1* (BT_0655)	(Lu et al., 2018)
LZ200 AKA pSOD	As BT5482 Δ*tdk* with pNLY-*B.thetaiotaomicron sod1* (BT_0655)	lab stock
LZ62	As BT5482 Δ*tdk* with Δ*fum* (BT_2256)	(Lu et al., 2018)

Supplementary Table 1. The strains used in this study.

Plasmid	Genotype	Reference
pExchange-*tdk*	Derivative of pKNOCK-*bla-ermGb* carrying cloned *tdk*	(Koropatkin et al., 2008)
pNLY-$sodB_{BT}$	pNLY plasmid constitutively expressing B. thetaiotaomicron *sod* from its own promoter and RBS (520 bp upstream of the gene)	(Lu et al., 2018)
pBAD/HisA-$pfor_{BT}$	pBAD/HisA plasmid (Invitrogen, catalog number V430-01) overexpressing *pfor* gene from *B. thetaiotaomicron*	This study

Supplementary Table 2. The plasmids used in this study.

Primer		Sequence
***pfor*-deletion**	F1	5'TTGT**TCTAGA**ACCTGCAGACAGGAATGAAATTCCG3'
	R1	5'TAGTCGAAGAAGGAGTGATGGG3'
	F2	5'ATCACTCCTTCTTCGACTAGAAACAATATCCTGCTGAAGC G3'
	R2	5'TTGT**GGATCC**GTATTATGAAAGCTACCGATCAGGCCAT3'
***pfl4738*-deletion**	F1	5'TTGT**TCTAGA**TCACGCTTCTTTTCCAGGGCCTTTC3'
	R1	5'TCTTCTGCCAACGCTTCGAGGC3'
	F2	5'TCGAAGCGTTGGCAGAAGAGCTATACGATGCCATGGAGC3'
	R2	5'TTGT**GGATCC**ATCCTTGGGAACGGATTAAAGAATGCCC3'
***pfl2955*-deletion**	F1	5'TTGT**TCTAGA**ACCGCAACTCCAGTATATTGAAGAC3'
	R1	5'CCACCCGTCCATAGTCATCTTC3'
	F2	5'GATGACTATGGACGGGTGGAAGATCCTGCCGGATATCCGGAA3'
	R2	5'TTGT**GGATCC**ACTGAGCGAAACTCCCAGATAATCAATTC3'
***fdx*-deletion**	F1	5'TTGT**TCTAGA**TTTCGATTCGGAAAGCAAT3'
	R1	5'TTCGTTCGGCAAAAATAAAGC3'
	F2	5'TATTTTTGCCGAACGAAGTTTGCCCGTCTGAAGCTATT3'
	R2	5'TTGT**GTCGAC**CGAACATATCAATTATATCGGTGCACAT3'
***rnf*-deletion**	F1	5'TTGT**TCTAGA**CAATCTGAAATATATTGCGCAG3'
	R1	5'CGATTCTCACCGACAGATGAG3'
	F2	5'TCTGTCGGTGAGAATCGACCCTGGTGAAGATTCCGAA3'
	R2	5'TTGT**GTCGAC**ACCGGAAGCTATTGTATCACGGATAA3'
***hyd1834*-deletion**	F1	5'TTGT**TCTAGA**CCTCTTTATCACAAACTCCGGCA3'
	R1	5'CTCAATAGGGAGACGGTCAATTTTT3'
	F2	5'ACCGTCTCCCTATTGAGACGACATTGTTTCAGGACGTC3'
	R2	5'TTGT**GTCGAC**TCAGACGGAAGATGGATAATAAAAGGA3'
***hyd3472*-deletion**	F1	5'TTGT**TCTAGA**GAAAACAAGTATTACGAGTTCCCGGA3'
	R1	5'AAAAGGCAGAAAGCCAGGGA3'
	F2	5'CTGGCTTTCTGCCTTTTTGCTCTGCCTATTGGGTCAC3'
	R2	5'TTGT**GTCGAC**GCATCAAAAGTTCCGCTTCATAGAAAGTACG3'
***pfor*-overexpression**	F	5'ATA**CTCGAG**ACTAAACAGAAGAAATTCATCAC3'
	R	5'ATA**GGTACC**TTATTCAGCTGTATCAGCTCC3'

Supplementary Table 3. The primers used in this study. The bold regions are restriction enzyme cut sites.