

pKSMSufl amber mutant	Primer 1 (5'→3')	Primer 2 (5'→3')
F22Bpa	C CGT GGG CAA CCG CTG <u>TAG</u> ATG ACT GTA CAA CGT GC	GC ACG TTG TAC AGT CAT <u>CTA</u> CAG CGG TTG CCC ACG G
F167Bpa	ATC CAG GAT AAA CGG CTG GAT AAC <u>TAG</u> GGT ACG CCA GAA T	A TTC TGG CGT ACC <u>CTA</u> GTT ATC CAG CCG TTT ATC CTG GAT
L301Bpa	CTG CGC CCA ACC GGC <u>TAG</u> CTG CCG CTG GTC ACA	TGT GAC CAG CGG CAG <u>CTA</u> GCC GGT TGG GCG CAG
M311Bpa	C ACA GAC AGT CTT CCG <u>TAG</u> CGC TTG CTG CCA ACT	AGT TGG CAG CAA GCG <u>CTA</u> CGG AAG ACT GTC TGT G
L341Bpa	CCG GGT ATT AAT GGA CAG <u>TAG</u> TGG GAC GTC AAC CGT AT	AT ACG GTT GAC GTC CCA <u>CTA</u> CTG TCC ATT AAT ACC CGG
W414Bpa	TT TAT TTC GGT CAG CCT TCC <u>TAG</u> GCG CAC TTC C	G GAA GTG CGC <u>CTA</u> GGA AGG CTG ACC GAA ATA AA
W113pBpa	G AAC GCA GCT ACT CTG <u>TAG</u> TAT CAC GCC AAT ACT C	G AGT ATT GGC GTG ATA <u>CTA</u> CAG AGT AGC TGC GTT C
W134Bpa	C GGC CTT GCC GGA ATG <u>TAG</u> CTG GTG GA	TC CAC CAG <u>CTA</u> CAT TCC GGC AAG GCC G
L206Bpa	C TGG GTG CGC TTG CGA <u>TAG</u> CTG AAC GCG TCG	CGA CGC GTT CAG <u>CTA</u> TCG CAA GCG CAC CCA G
F234Bpa	G ATT TCT GGC GAT CAG GGA <u>TAG</u> CTG CCT GCT CC	GG AGC AGG CAG <u>CTA</u> TCC CTG ATC GCC AGA AAT C

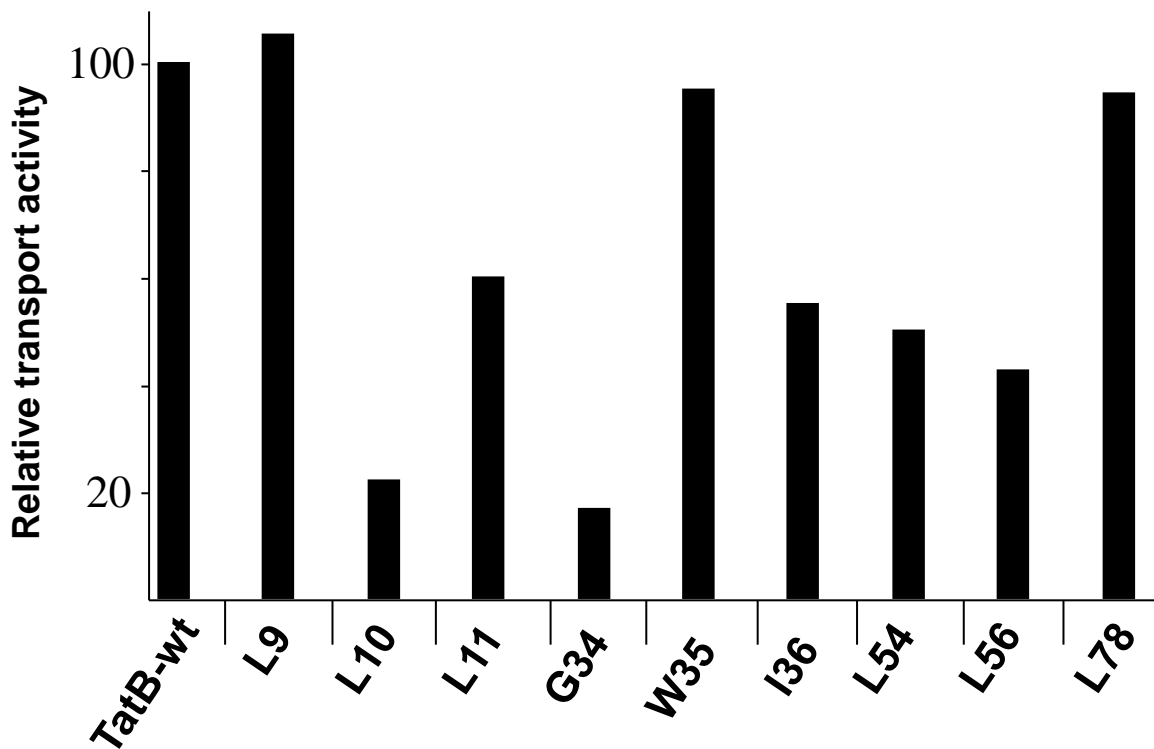
pET28aTorA-PhoA amber mutant	Primer 1 (5'→3')	Primer 2 (5'→3')
I124Bpa	GCG CTG GGC GTC GAT <u>TAG</u> CAC GAA AAA GAT CAC	GTG ATC TTT TTC GTG <u>CTA</u> ATC GAC GCC CAG CGC
W220Bpa	GCA ACC GCC GGT GAA <u>TAG</u> CAG GGA AAA ACG CTG	CAG CGT TTT TCC CTG <u>CTA</u> TTC ACC GGC GGT TGC
I279Bpa	ACG TAC CAC GGC AAT <u>TAG</u> GAC AAG CCC GCA GTT ACC	GGT AAC TGC GGG CTT GTC <u>CTA</u> ATT GCC GTG GTA CGT
L417Bpa	CAT ACC GGT AGT CAG <u>TAG</u> CGT ATT GCG GCG TAT	ATA CGC CGC AAT ACG <u>CTA</u> CTG ACT ACC GGT ATG
I99Bpa	CCT GCA AAA AAT ATT ATT TTG CTG <u>TAG</u> GGC GAT GGG ATG GGG GAT TCG G	C CGA ATC CCC CAT CCC ATC GCC <u>CTA</u> CAG CAA AAT AAT ATT TTT TGC AGG
V196	GGT CTG GCG ACC GGT AAC <u>TAG</u> TCT ACC GCA GAG TTG CAG	CTG CAA CTC TGC GGT AGA <u>CTA</u> GTT ACC GGT CGC CAG ACC

p8737-TatB amber mutant	Primer 1 (5'→3')	Primer 2 (5'→3')
L9Bpa	ATC GGT TTT AGC GAA <u>TAG</u> CTA TTG GTG TTC ATC ATC GGC CTC	GAG GCC GAT GAT GAA CAC CAA TAG <u>CTA</u> TTC GCT AAA ACC GAT
L10Bpa	GAT ATC GGT TTT AGC GAA CTG <u>TAG</u> TTG GTG TTC ATC ATC GGC CTC	GAG GCC GAT GAT GAA CAC CAA <u>CTA</u> CAG TTC GCT AAA ACC GAT ATC
L11Bpa	C GGT TTT AGC GAA CTG CTA <u>TAG</u> GTG TTC ATC ATC GG	CC GAT GAT GAA CAC <u>CTA</u> TAG CAG TTC GCT AAA ACC G
G34Bpa	CG GTA AAA ACG GTA GCG <u>TAG</u> TGG ATT CGC GCG TTG CG	CG CAA CGC GCG AAT CCA <u>CTA</u> CGC TAC CGT TTT TAC CG
W35Bpa	ACG GTA GCG GGC <u>TAG</u> ATT CGC GCG TTG	CAA CGC GCG AAT <u>CTA</u> GCC CGC TAC CGT
I36Bpa	ACG GTA GCG GGC TGG <u>TAG</u> CGC GCG TTG CGT TCA	TGA ACG CAA CGC GCG <u>CTA</u> CCA GCC CGC TAC CGT
L54Bpa	AAC GAA CTG ACC CAG GAG <u>TAG</u> AAA CTC CAG GAG TTT CAG G	CTG AAA CTC CTG GAG TTT <u>CTA</u> CTC CTG GGT CAG TTC GTT

L56Bpa	GAA CTG ACC CAG GAG TTA AAA <u>TAG</u> CAG GAG TTT CAG GAC AGT CTG	CAG ACT GTC CTG AAA CTC CTG <u>CTA</u> TTT TAA CTC CTG GGT CAG TTC
L78Bpa	CT AAC CTG ACG CCC GAA <u>TAG</u> AAA GCG TCG ATG GAT G	C ATC CAT CGA CGC TTT <u>CTA</u> TTC GGG CGT CAG GTT AG

pKSMSufI ΔSP	Primer 1 (5'→3')	Primer 2 (5'→3')
SufI A27L	CCC CTG AAG GCC AGC <u>CTA</u> GCC GGG CAA CAG	CTG TTG CCC GGC <u>TAG</u> GCT GGC CTT CAG GGG
SufI S26L&A27L	CCC CTG AAG GCC <u>CTC</u> CTA GCC GGG CAA CAG	CTG TTG CCC GGC TAG <u>GAG</u> GGC CTT CAG GGG
SufI S26L&A27L&Q30L	CCC CTG AAG GCC <u>CTC</u> CTA GCC GGG CTA CAG	CTG TAG CCC GGC TAG <u>GAG</u> GGC CTT CAG GGG

Transport activity of *tatB* amber mutant vesicles



pSufI was synthesized in the presence of one of the indicated membrane vesicles. INVs were reisolated and then incubated in the presence of an ATP-regenerating system. The amount of PK-protected pSufI and SufI was determined, the translocation efficiency quantified and normalized with respect to that of wt-INVs (for further details, see Materials and Methods).