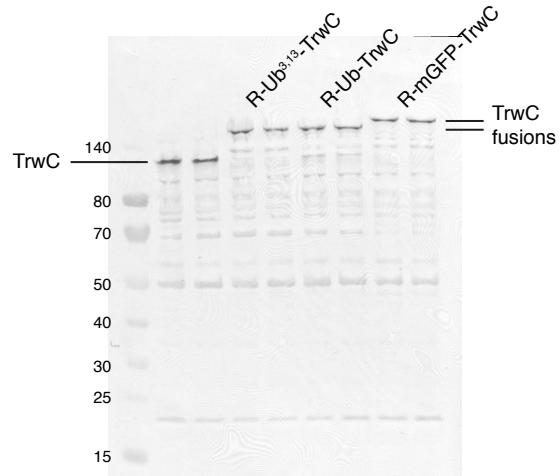


## Supplemental material

**Figure S1.** Western blot of donor cells carrying R388 plasmid encoding wild-type or modified TrwC as indicated detected using self-made polyclonal  $\alpha$ -TrwC antibody.



Shown are duplicates for each construct. The vast majority of the expressed TrwC consists of the full-length protein. There are no additional or stronger degradation bands in the R-Ub<sup>3,13</sup>-TrwC construct in comparison to R-Ub-TrwC.

**Supplementary table 1. Plasmids used in this study**

Plasmids	Description	Source
pRSFDuet-1	Vector carrying two T7 promoters, RSF replicon, <i>lacI</i> gene and kanamycin resistance gene (KnR)	Novagen
pRSF-oriT	pRSFDuet-1 vector containing R388 <i>oriT</i> sequence and lacking part of <i>lacI</i> gene and the first promoter region	This study
pUC18	High copy number cloning vector	Agilent Technologies
pUC-oriT	pUC18 vector containing R388 <i>oriT</i> sequence	This study
pBAD-B1-B10Strep	pBADM11 vector containing the genes <i>trwN/virB1</i> , <i>trwL/virB2</i> , <i>trwM/virB3</i> , <i>trwK/virB4</i> , <i>trwJ/virB5</i> , <i>trwI/virB6</i> , <i>trwH/virB7</i> , <i>trwG/virB8</i> , <i>trwF/virB9</i> , and <i>trwE/virB10</i> , with a Strep tag fused to TrwE C-terminus	(1)
pASK-B1-B10Strep-B11	pASK-IBA3C vector containing the R388 genes <i>virB1-virB11</i> , with a Strep tag fused to VirB10 C-terminus	(2)
pBAD-B1-B10Strep-B11	Identical to pBAD-B1-B10Strep, but containing <i>virB11</i> downstream of <i>virB10Strep</i>	This study
pBAD/MCS	Expression vector carrying <i>araBAD</i> promoter, <i>araC</i> gene and ampicillin resistance gene (AmpR)	EMBL
pBAD-ABCHis	pBAD/MCS vector containing the R388 genes <i>trwA</i> , <i>trwB</i> , and <i>trwC</i> with a (Gly)3 linker and a hexahistidine tag fused to the TrwC C-terminus	This study
pBAD-ABC-T4SS	Identical to pBAD-B1-B10Strep-B11, but containing the genes <i>trwA</i> , <i>trwB</i> , and <i>trwC</i> and a consensus <i>E. coli</i> rbs sequence upstream of <i>virB1</i> ; fused to the TrwC C-terminus is a hexahistidine tag separated by a (Gly)3 linker	This study
pBABE-Cre-ERT2	Vector encoding Cre recombinase	(3)
pBAD-AB_Cre-C	Identical to pBAD-ABCHis but containing Cre recombinase fused to the TrwC N-terminus with a (Gly)3Ser(Gly)2 linker	This study
pBAD-AB_R-Cre-H	Identical to pBAD-ABCHis but containing Cre recombinase inserted between the TrwC relaxase (R; aa 1-312) and helixase (H; aa 1-313) domains with (Gly)3SerGly linkers at both sides	This study
pBAD-ABC-Cre	Identical to pBAD-ABCHis but containing (Gly)3-mGFP-(Gly)3-His6 fused to the TrwC C-terminus	This study
pBAD-AB_Cre-C_T4SS	Identical to pBAD-AB_Cre-C but containing a consensus <i>E. coli</i> rbs sequence followed by <i>virB1-virB11</i> genes downstream of <i>trwC</i> (TrwE has a Strep tag fused to the C-terminus)	This study
pBAD-AB_R-Cre-H_T4SS	Identical to pBAD-AB_R-Cre-H but containing a consensus <i>E. coli</i> rbs sequence followed by <i>virB1-virB11</i> genes downstream of <i>trwC</i> (TrwE has a Strep tag fused to the C-terminus)	This study
pBAD-ABC-Cre_T4SS	Identical to pBAD-ABC-Cre but containing a consensus <i>E. coli</i> rbs sequence followed by <i>virB1-virB11</i> genes	This study

	downstream of <i>trwC</i> (TrwE has a Strep tag fused to the C-terminus)	
pBAD-AB_Cre(Y324F)-C	Identical to pBAD-AB_Cre-C but containing Y324F mutation in Cre	This study
pETZt	Expression vector carrying <i>lacZ</i> gene, kanamycin resistance gene (KnR), and T7 promoter encoding a hexahistidine tag followed by Z tag and TEV protease cleavage site	EMBL
pETZt-TrwC	pETZt vector encoding TrwC with an N-terminal hexahistidine tag followed by Z tag and TEV cleavage site	This study
pETZt-mGFP-TrwC	pETZt vector encoding TrwC with an N-terminal hexahistidine tag followed by Z tag, TEV cleavage site, and mGFP	This study
pFastBacHT-mGFP	A modified pFastBacHTa vector encoding monomeric GFP (the K221L mutation of eGFP; Zacharias et al. 2002)	(4)
pBAD-ABC-mGFP	Identical to pBAD-ABC-Cre but containing mGFP instead of Cre recombinase	This study
pBAD-AB_R-mGFP-H	Identical to pBAD-AB_R-Cre-H but containing mGFP instead of Cre recombinase	This study
pBAD-AB_mGFP-C	Identical to pBAD-AB_Cre-C but containing mGFP instead of Cre recombinase	This study
pET17-Ub	pET17b vector encoding ubiquitin (Ub)	(5)
pET17-UbI3G,I13G	pET17b vector encoding ubiquitin mutant; Ile at amino acid positions 3 and 13 mutated to Gly	This study
pBAD-AB_Ub-C	Identical to pBAD-AB_Cre-C but containing Ub instead of Cre recombinase	This study
pBAD-AB_Ub3,13-C	Identical to pBAD-AB_Ub-C apart from ubiquitin Ile to Gly mutations at amino acid positions 3 and 13	This study
pBAD-AB_Cre-Ub-C	Identical to pBAD-ABCHis but containing Cre-(Gly)3SerGly-Ub-(Gly)3Ser(Gly)2 fused to the TrwC N-terminus	This study
pBAD-AB_Cre-Ub3,13-C	Identical to pBAD-AB_Cre-Ub-C apart from ubiquitin Ile to Gly mutations at amino acid positions 3 and 13	This study
pBAD-AB_Cre-mGFP-C	Identical to pBAD-AB_Cre-Ub-C but containing mGFP instead of Ub	This study
pBAD-AB_Cre-Cre(Y324F)-C	Identical to pBAD-AB_Cre-Ub-C but containing Cre(Y324F) instead of Ub	This study
pBAD-AB_Cre-Ub-C_T4SS	Identical to pBAD-AB_Cre-Ub-C but containing a consensus E. coli rbs sequence followed by <i>virB1-virB11</i> genes downstream of <i>trwC</i> (TrwE has a Strep tag fused to the C-terminus)	This study
pBAD-AB_Cre-Ub3,13-C_T4SS	Identical to pBAD-AB_Cre-Ub3,13-C but containing a consensus E. coli rbs sequence followed by <i>virB1-virB11</i> genes downstream of <i>trwC</i> (TrwE has a Strep tag fused to the C-terminus)	This study
pBAD-AB_Cre-mGFP-C_T4SS	Identical to pBAD-AB_Cre-mGFP-C but containing a consensus E. coli rbs sequence followed by <i>virB1-virB11</i> genes downstream of <i>trwC</i> (TrwE has a Strep tag fused to the C-terminus)	This study

pBAD-AB_Cre-Cre(Y324F)-C_T4SS	Identical to pBAD-AB_Cre-Cre(Y324F)-C but containing a consensus E. coli rbs sequence followed by <i>virB1-virB11</i> genes downstream of <i>trwC</i> (TrwE has a Strep tag fused to the C-terminus)	This study
pBAD-AB_R-Ub-H	Identical to pBAD-ABCHis but containing (Gly)3Ser(Gly)3-Ub-(Gly)3SerGly inserted between the TrwC relaxase (R; aa 1-312) and helixase (H; aa 313-966) domains	This study
pBAD-AB_R-Ub3,13-H	Identical to pBAD-AB_R-Ub-H apart from ubiquitin Ile to Gly mutations at amino acid positions 3 and 13	This study
pBAD-AB_R-Ub-C	pBAD/MCS vector containing the genes <i>trwA</i> , <i>trwB</i> , and modified <i>trwC</i> ; TrwC contains a duplicated relaxase domain (R; aa 1-305) followed by (Gly)3-His6-GlySer(Gly)3-Ub-(Gly)3SerGly at its N-terminus and the second relaxase domain lacks the first methionine	This study
pBAD-AB_R-Ub3,13-C	Identical to pBAD-AB_R-Ub-C apart from ubiquitin Ile to Gly mutations at amino acid positions 3 and 13	This study
pBAD-AB_R-mGFP-C	Identical to pBAD-AB_R-Ub-C but containing mGFP instead of ubiquitin	This study
pBAD-ABC-Ub-H	pBAD/MCS vector containing the genes <i>trwA</i> , <i>trwB</i> , and modified <i>trwC</i> ; TrwC contains (Gly)3Ser(Gly)3-Ub-(Gly)3SerGly followed by a duplicated helicase domain (H; aa 313-966) and a hexahistidine tag at its C-terminus	This study
pBAD-ABC-Ub3,13-H	Identical to pBAD-ABC-Ub-H apart from ubiquitin Ile to Gly mutations at amino acid positions 3 and 13	This study
pBAD-ABC-mGFP-H	Identical to pBAD-ABC-Ub-H but containing mGFP instead of ubiquitin	This study
pEL04	Vector containing the <i>cm-sacB</i> cassette to be used for positive/negative selection in recombineering. Positive selection using chloramphenicol, and negative selection against the <i>sacB</i> gene using sucrose.	NCI at Frederik
R388_R-Ub-TrwC	R388 plasmid encoding modified TrwC; TrwC contains a duplicated relaxase domain (R; aa 1-305) followed by (Gly)3-His6-GlySer(Gly)3-Ub-(Gly)3SerGly at its N-terminus and the second relaxase domain lacks the first methionine	This study
R388_R-Ub3,13-TrwC	Identical to R388_R-Ub-TrwC apart from ubiquitin Ile to Gly mutations at amino acid positions 3 and 13	This study
R388_R-mGFP-TrwC	Identical to R388_R-Ub-TrwC but containing mGFP instead of ubiquitin	This study
R388_TrwC-Ub-H	R388 plasmid encoding modified TrwC; TrwC contains (Gly)3Ser(Gly)3-Ub-(Gly)3SerGly followed by a duplicated helicase domain (H; aa 313-966) and a hexahistidine tag at its C-terminus	This study
R388_TrwC-Ub3,13-H	Identical to R388_TrwC-Ub-H apart from ubiquitin Ile to Gly mutations at amino acid positions 3 and 13	This study

R388_TrwC-mGFP-H	Identical to R388_TrwC-Ub-H but containing mGFP instead of ubiquitin	This study
R388_R-Ub-H	R388 plasmid encoding modified TrwC; TrwC contains (Gly)3Ser(Gly)3-Ub-(Gly)3SerGly inserted between the TrwC relaxase (R; aa 1-312) and helixase (H; aa 313-966) domains	This study
R388_R-Ub3,13-H	Identical to R388_R-Ub-H apart from ubiquitin Ile to Gly mutations at amino acid positions 3 and 13	This study
R388_R-mGFP-H	R388 plasmid encoding modified TrwC; TrwC contains mGFP inserted between the TrwC relaxase (R; aa 1-312) and helixase (H; aa 1-313) domains with (Gly)3SerGly linkers at both sides	This study

**Supplementary table 2. Primers used in this study.** Restriction enzyme sites are shown in bold.

Name of the primers	Sequence of the primers	Name of the construct cloned using the primers	Description
BssHII-oriT fwd BssHII-oriT rev	<b>TAGCGCGCCACTCATTCTGCAT</b> CATTGTAGCACCATC <b>TAGCGCGCCACCGCCTCGTCCTC</b> CAAAAGTG	pRSF-oriT	primers used to amplify <i>oriT</i> from R388 plasmid
HindIII-oriT fwd Sacl-oriT rev	<b>TATAAGCTTCTCATTCTGCATCA</b> TTGTAGCACCATC <b>TATGAGCTCCGCCTCGTCCTCCA</b> AAAGTG	pUC-oriT	primers used to amplify <i>oriT</i> from pRSF-oriT plasmid
Strep-B11 fwd B11 rev	GCCACCCGCAGTCGAAAAATAAG G CTCTCTTATTTTCGCCCTGCAGGT CGACCTCG	pBAD-B1-B10Strep-B11	primers used to amplify <i>Strep-virB11</i> from pASK-B1-B10Strep-B11 plasmid
Ncol-TrwA fwd TrwC-His rev	AGGAATTAACCATGGCACTAGGCG ACCCCATC GATGATGATGATGATGACCGCCAC CCCTCCGGCCTCCATGC	pBAD-ABCHis	primers used to amplify <i>trwABC</i> while adding a hexahistidine tag to TrwC C-terminus
pBAD/MC S fwd pBAD/MC S rev	ATCATCATCATCATCATTAATGAGC TCCGTCGACAAGCTTGC CCATGGTTAACCTCCCTGTTAGC	pBAD-ABCHis	primers used to amplify pBAD/MCS vector for inserting <i>trwABC</i> and adding a hexahistidine tag to TrwC C-terminus
TrwA fwd His-rbs rev	GAGGAATTAACCATGGCACTAGG TCTGCCAGTGCCATGGTTATT <b>CC</b> <b>TCCT</b> TGTCGACGGAGCTCATTAAT GATG	pBAD-ABC-T4SS	primers used to amplify <i>trwABCHis</i> from pBAD-ABCHis plasmids (including all plasmids containing modified TrwC) and adding rbs (red) downstream of <i>trwCHis</i>
C-mGFP fwd C-mGFP rev	AGGGGTGGCGGTGTGAGCAAGGG CGAGGAGC TGATGATGATGCCGCCCTCCCTTG TACAGCTC	pBAD-ABC-mGFP	primers used to amplify <i>mGFP</i> from pFastBacHT-mGFP plasmid
mGFP-His fwd mGFP-TrwC rev	CGGGCATCATCATCATCATCATTAA TGAGC CACACCGCCACCCCTTCC	pBAD-ABC-mGFP	primers used to linearize pBAD-ABCHis plasmid
mGFP int fwd mGFP int rev	<b>GGCGGTGGGAGCGGGTGAGCAA</b> GGGCGAG GCCGCTGCCACCTCCCTGTACAG CTCGTCC	pBAD-AB_R-mGFP-H	primers used to amplify <i>mGFP</i> from pBAD-ABC-mGFP plasmid

mGFP-H fwd mGFP-R rev	GGAGGTGGCAGCGGCATGCCCTC CGACGAGG ACCGCTCCCACCGCCGAATGAATG GGCCTTTTCTG	pBAD- AB_R- mGFP-H	primers used to linearize pBAD-ABCHis plasmid
N-mGFP fwd N-mGFP rev	GACTATCTAATGGTGAGCAAGGGC GAGGAGC CATACCGCCGCTACCGCCTCCCTT GTACAGCTC	pBAD- AB_mGFP- C	primers used to amplify <i>mGFP</i> from pBAD-ABC-mGFP plasmid
mGFP- TrwC fwd mGFP- TrwB rev	GGTAGCGGGCGGTATGCTCAGTCA CATGGTATTGACC CACCAATTAGATAGTCCCCTAACAA AAGG	pBAD- AB_mGFP- C	primers used to linearize pBAD-ABCHis plasmid
Ub I3G Ub I13G	CAGGGTCTTCACGAAG <u>CC</u> CTGCAT ATGTATATCTCC CTCCACTTCGAGAG <u>TG</u> CCGGTCTT ACCAGTCAG	pET17- Ubl3G,I13 G	primers for multi site-directed mutagenesis that incorporate ubiquitin I3G and I13G mutations (underlined)
Ub int fwd Ub int rev	GGCGGTGGGAGCGGGTGGGGGCAT GCAGATCTCGTGAAGACCCCTGAC GCCGCTGCCACCTCCCCCACCTCT GAGACGGAGG	pBAD- AB_R-Ub- H	primers used to amplify Ub sequence from pET17-Ub plasmid
Ub-H fwd Ub-R rev	GGAGGTGGCAGCGGCATGCCCTC CGACGAGG ACCGCTCCCACCGCCGAATGAATG GGCCTTTTCTG	pBAD- AB_R-Ub- H	primers used to linearize pBAD-ABCHis plasmid
Ub3,13 int fwd Ub int rev	GGCGGTGGGAGCGGGTGGGGGCAT GCAGGGCTTCGTGAAGACCCCTGA C GCCGCTGCCACCTCCCCCACCTCT GAGACGGAGG	pBAD- AB_R- Ub3,13-H	primers used to amplify Ubl3G,I13G sequence from pET17-Ubl3G,I13G plasmid
Ub-H fwd Ub-R rev	GGAGGTGGCAGCGGCATGCCCTC CGACGAGG ACCGCTCCCACCGCCGAATGAATG GGCCTTTTCTG	pBAD- AB_R- Ub3,13-H	primers used to linearize pBAD-ABCHis plasmid
N-Ub fwd N-Ub rev	ATCTAATGCAGATCTCGTGAAGA CCCTGAC GCCGCTACCGCCTCCCCCACCTCT GAGACGGAGG	pBAD- AB_Ub-C	primers used to amplify Ub sequence from pET17-Ub plasmid
Ub-TrwC fwd Ub-TrwB rev	GGAGGCGGTAGCGGGCGGTATGCT CAGTCACATGGTATTGACC AGATCTGCATTAGATAGTCCCCTC AACAAAGG	pBAD- AB_Ub-C	primers used to linearize pBAD-ABCHis plasmid
N-Ub3,13 fwd N-Ub rev	ATCTAATGCAGGGCTTCGTGAAGA CC GCCGCTACCGCCTCCCCCACCTCT GAGACGGAGG	pBAD- AB_Ub3,13 -C	primers used to amplify Ubl3G,I13G sequence from pET17-Ubl3G,I13G plasmid
Ub-TrwC fwd Ub3,13- TrwB rev	GGAGGCGGTAGCGGGCGGTATGCT CAGTCACATGGTATTGACC AGCCCTGCATTAGATAGTCCCCTC AACAAAGG	pBAD- AB_Ub3,13 -C	primers used to linearize pBAD-ABCHis plasmid
C-Cre fwd NC-Cre rev	GGTCCAATTACTGACCGTACAC C	pBAD- ABC-Cre	primers used to amplify Cre sequence

	CGCCTCCATGCCATCTTCCAGCA G		from pBABE-Cre-ERT2 plasmid
Cre-His fwd Cre-TrwC rev	ATGGCGATGGAGGCAGGCATCAT CATC CAGTAAATTGGAACCGCCACCCCT TCCG	pBAD-ABC-Cre	primers used to amplify the portion of the pBAD-ABC-mGFP plasmid lacking mGFP
Cre int fwd Cre int rev	GGTTCCAATTACTGACCGTACAC C GCCACCTCCATGCCATCTTCCAG CAG	pBAD-AB_R-Cre-H	primers used to amplify Cre sequence from pBABE-Cre-ERT2 plasmid
Cre-H fwd Cre-R rev	GGCGATGGAGGTGGCAGCGGCAT G CAGTAAATTGGAACCGCTCCCACC GCCGAATG	pBAD-AB_R-Cre-H	primers used to amplify the portion of the pBAD-AB_R-mGFP-H plasmid lacking mGFP
N-Cre fwd NC-Cre rev	CTAATGTCCAATTACTGACCGTAC ACC CGCCTCCATGCCATCTTCCAGCA G	pBAD-AB_Cre-C	primers used to amplify Cre sequence from pBABE-Cre-ERT2 plasmid
Cre-TrwC fwd Cre-TrwB rev	ATGGCGATGGAGGCAGGTAGCGGC GGTATG TAAATTGGACATTAGATAGTAGTCCCCT CAACAAAGG	pBAD-AB_Cre-C	primers used to amplify the portion of the pBAD-AB_mGFP-C plasmid lacking mGFP
Cre Y324F sense Cre Y324F antisense	CCAATGTAAATATTGTCATGA <u>ACTT</u> TATCCGTAA <u>CTGG</u> CCAGGTTACGGATA <u>AAAGTT</u> CATGA CAATATTACATTGG	pBAD-AB_Cre(Y324F)-C	primers for site-directed mutagenesis that incorporate Cre Y324F mutation (underlined)
TrwC fwd TrwC rev	CATGCTCAGTCACATGGTATTGAC C ATCTCATTACCTTCCGGCCTCCAT GC	pETZt-TrwC	primers used to amplify <i>trwC</i> from pBAD-ABCHis plasmid
pETZt fwd TrwC-pETZt rev	GGAAGGTAATGAGATCCGGCTGCT AACAAAGC ATGTGACTGAGCATGGCGCCCTGA AAATAAAGATTTC	pETZt-TrwC	primers used to linearize pETZt vector
mGFP fwd TrwC rev	GCCATGGTGAGCAAGGGCGAG ATCTCATTACCTTCCGGCCTCCAT GC	pETZt-mGFP-TrwC	primers used to amplify <i>mGFP-trwC</i> from pBAD-AB_mGFP-C plasmid
pETZt fwd mGFP-pETZt rev	GGAAGGTAATGAGATCCGGCTGCT AACAAAGC CTTGCTCACCATGGCGCCCTGAAA ATAAAAGATTTC	pETZt-mGFP-TrwC	primers used to linearize pETZt vector
Cre-Ub fwd TrwC int rev	GGGAGCGGTGGGGCATGCAGAT CTTCGTGAAGACCTG GCATAGGCATGATCGACGTGC	pBAD-AB_Cre-Ub-C	primers used to amplify Ub-TrwC sequence from pBAD-AB_Ub-C plasmid

TrwC int fwd linker-Cre rev	CGATCATGCCTATGCGACGAC GCCCCCACCGCTCCACCATCGC CATCTTCAGCAG	pBAD- AB_Cre- Ub-C pBAD- AB_Cre- Ub3,13-C pBAD- AB_Cre- mGFP-C pBAD- AB_Cre- Cre(Y324F)- -C	primers used to linearize pBAD-AB_Cre-C plasmid
Cre- Ub3,13 fwd TrwC int rev	GGGAGCGGTGGGGCATGCAGG GCTTCGTGAAGACC GCATAGGCATGATCGACGTGC	pBAD- AB_Cre- Ub3,13-C	primers used to amplify Ub3,13-TrwC sequence from pBAD-AB_Ub3,13-C plasmid
Cre- mGFP fwd TrwC int rev	GGGAGCGGTGGGGCGTGAGCAA GGGCGAGGAGC GCATAGGCATGATCGACGTGC	pBAD- AB_Cre- mGFP-C	primers used to amplify mGFP-TrwC sequence from pBAD-AB_mGFP-C plasmid
Cre-Cre fwd TrwC int	GGGAGCGGTGGGGCTCCAATT ACTGACCGTACACCAAAATTGC GCATAGGCATGATCGACGTGC	pBAD- AB_Cre- Cre(Y324F)- -C	primers used to amplify Cre(Y324F)-TrwC sequence from pBAD-AB_Cre(Y324F)-C plasmid
pBAD fwd His-R rev	CGGAAGGTAATGAGCTCCGTCGAC AAGCTTG GTGATGATGGTGATGACCGCCACC TGAACCGCCCTTCTCCCC	pBAD- AB_R-Ub- C pBAD- AB_R- Ub3,13-C	primers used to linearize pBAD-ABCHis plasmid lacking helicase domain and the hexahistidine tag
His-Ub fwd linker-Ub rev	CATCACCATCATCACCATGGGAGC GGTGGGGGCATG GCCGCTGCCACCTCCCCACCTCT GAGACGGAGGACC	pBAD- AB_R-Ub- C pBAD- AB_R- Ub3,13-C	primers used to amplify Ub/Ub3,13 sequence from pBAD-AB_R-Ub-H plasmid
linker- TrwC fwd TrwC rev 2	GGAGGTGGCAGCGGCCTCAGTCA CATGGTATTGACCCGACAG GCTCATTACCTTCCGGCCTCCATG CC	pBAD- AB_R-Ub- C pBAD- AB_R- Ub3,13-C	primers used to amplify trwC from pBAD-ABCHis plasmid
linker- TrwC 2 mGFP- His rev	GGAGGTGGCAGCGGCCTCAGTCA CATGGTATTGACC CTCGCCCTTGCTCACGCCACCCACC GCTCCCAGT	pBAD- AB_R- mGFP-C	primers used to linearize pBAD-AB_R-Ub-C plasmid lacking ubiquitin sequence
mGFP fwd 2	GTGAGCAAGGGCGAGGAGC GCCGCTGCCACCTCCCTTG	pBAD- AB_R- mGFP-C	primers used to amplify mGFP sequence from pBAD-

mGFP rev 2			AB_R-mGFP-H plasmid
TrwC-linker fwd TrwB rev	CGGAAGGGCGGTGGGAGCGGT GG CCTCAACAAAGGCCGGTTGC	pBAD-ABC-Ub-H pBAD-ABC-Ub3,13-H	primers used to amplify Ub/Ub3,13-H-pBAD-AB from pBAD-AB_R-Ub/Ub3,13-H plasmid
TrwB fwd TrwC rev 3	CGGCCTTGTGAGGGGACTATC CCACCGCCCCCTTCCGGCCTCCATG	pBAD-ABC-Ub-H pBAD-ABC-Ub3,13-H	primers used to amplify <i>trwC</i> from pBAD-ABCHis plasmid
mGFP fwd 2 TrwB rev	GTGAGCAAGGGCGAGGAGC CCTCAACAAAGGCCGGTTGC	pBAD-ABC-mGFP-H	primers used to linearize pBAD-AB_R-mGFP-H plasmid lacking the relaxase domain sequence
TrwB fwd mGFP-linker rev	CGGCCTTGTGAGGGGACTATC CTCGCCCTTGCTCACGCCACC GCTCCCAC	pBAD-ABC-mGFP-H	primers used to amplify <i>trwC</i> from pBAD-ABC-Ub-H plasmid

## **Supplemental References**

- 1- Low, H. H., Gubellini, F., Rivera-Calzada, A., Braun, N., Connery, S., Dujeancourt, A., Lu, F., Redzej, A., Fronzes, R., Orlova, E. V., and Waksman, G. (2014) Structure of a type IV secretion system. *Nature* **508**, 550-553
- 2- Redzej, A., Ukleja, M., Connery, S., Trokter, M., Felisberto-Rodrigues, C., Cryar, A., Thalassinos, K., Hayward, R. D., Orlova, E. V., and Waksman, G. (2017) Structure of a VirD4 coupling protein bound to a VirB type IV
- 3- Lang, S., Gruber, K., Mihajlovic, S., Arnold, R., Gruber, C. J., Steinlechner, S., Jehl, M. A., Rattei, T., Frohlich, K. U., and Zechner, E. L. (2010) Molecular recognition determinants for type IV secretion of diverse families of conjugative relaxases. *Molecular microbiology* **78**, 1539-1555
- 4- Trokter, M., Mücke, N., Surrey, T. (2012) Reconstitution of the human cytoplasmic dynein complex. *Proc. Natl. Acad. Sci.* **109**, 20895-900
- 5- Hospenthal, M. K., Freund, S. M., Komander, D. (2013) Assembly, analysis and architecture of atypical ubiquitin chains. *Nat. Struct. Mol. Biol.* **20**, 555-65