

1 **Engineering Gram-negative Microbial Cell Factories Using Transposon Vectors**

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Running Head: Transposon vectors for metabolic engineering

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1 Abstract

2

3 The construction of microbial cell factories *à la carte* largely depends on specialized molecular biology
4 and synthetic biology tools, needed to re-program bacteria for modifying their existing functions or for
5 bestowing them with new-to-Nature tasks. In this protocol, we document the use of a series of broad-
6 host-range mini-Tn5 vectors for the delivery of gene(s) into the chromosome of Gram-negative bacteria
7 and the generation of saturated, random mutagenesis libraries for studies of gene function. The
8 application of these tailored mini-transposon vectors, which could be also used for chromosomal
9 engineering of a wide variety of Gram-negative microorganisms, is demonstrated in the platform
10 environmental bacterium *Pseudomonas putida* KT2440.

11

12 **Key words:** Mini-transposon, Tn5 transposon, *Pseudomonas putida*, *Escherichia coli*, Synthetic
13 biology, Metabolic engineering, Microbial cell factory, Genome editing

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1 1. Introduction

2

3 *Mini-transposon vectors* allow for the stable insertion of foreign DNA into the chromosome of many
4 types of Gram-negative bacterial targets (1,2). Tn5-derived elements (3) present clear advantages over
5 the use of their plasmid-based counterparts for the random interruption of gene(s), or for the
6 introduction and expression of heterologous genes into several bacterial species. These features
7 include, but are not limited to, (i) maintenance of the corresponding transgenes without antibiotic
8 selective pressure, (ii) long-term stability of the constructs and re-usability of the functional DNA parts,
9 and, furthermore, (iii) mini-Tn5-based vectors admit cloning and chromosomal delivery of considerably
10 long DNA fragments (which would be cumbersome to manipulate in other DNA delivery tools). As the
11 transposase gene *tnpA* is lost following each transposition event (4,5), one added value of mini-Tn5
12 vectors is the possibility to use them recursively in the same microbial host, provided that they bear
13 different selection markers. Since the TnpA transposase tends to act in *cis* (6), it promotes the insertion
14 of DNA sequences borne by the plasmid irrespective of any previous DNA insertions in a given
15 chromosome. These features allow for the delivery and integration of various DNA cargoes into the
16 same target genome. However, the original layout of such mini-transposon vectors was not exempt of
17 downsides. One of them is the unavoidable inheritance of long, non-functional DNA fragments
18 stemming from the intricate cutting-and-pasting DNA methods available at the time when the original
19 vectors were constructed. These procedures were also afflicted by the presence of an excessive and
20 inconvenient number of non-useful restriction sites scattered along the plasmids, and the suboptimal
21 transposition machinery encoded therein.

22

23 Martínez-García *et al.* (7) thoroughly revisited the original mini-Tn5 transposon vector concept. The
24 most attractive features of the mini-Tn5-aided mutagenesis procedure have been enhanced while
25 each of its drawbacks [identified along >20 years of use in many independent laboratories worldwide
26 (8)] has been eliminated. The functional modules that constitute the vector (including the mosaic
27 elements, MEs) have been edited to minimize the length of the corresponding DNA fragments,
28 improving their functionality and making them entirely modular and exchangeable. The final product
29 was the entirely synthetic plasmid construct termed pBAM1 (*born-again mini-transposon*). This design
30 was soon followed by a series of synthetic, modular broad-host-range mini-Tn5 plasmids derived from
31 pBAM1. These vectors, termed pBAMDs vectors (9), enable the possibility of easy cloning and

1 subsequent chromosomal insertion of functional DNA cargoes with three different and interchangeable
2 antibiotic resistance markers. Another set of pBAM1-derivative plasmids, termed pBELs and pBEXs
3 vectors (**10**), were designed to exploit the possibility of delivering DNA cargoes under the control of
4 regulated gene expression modules (i.e., LacI^O/P_{trc} in pBELs vectors or XylS/P_m in pBEXs vectors).
5 Furthermore, the antibiotic-resistance determinants in the mini-transposon modules of the pBELs and
6 pBEXs vectors can be removed by means of the FLP recombinase from *Saccharomyces cerevisiae*
7 (**11**). In all the cases presented above, the functional parts of the mini-transposon vectors can be easily
8 swapped by digestion with the appropriate restriction enzymes, allowing for the easy shuffling of each
9 DNA element as needed. Finally, the multiple cloning site of all the mini-transposon vectors share the
10 same set of restriction sites, which eases the subcloning of DNA cargoes by making them compatible
11 with plasmids from the Standard European Vector Architecture (SEVA) initiative (**12,13**).
12

13 The expansion of the available mini-transposon tools is a step forward in our efforts to purposely
14 engineer microbial cell factories, mainly based on environmental bacteria. *Pseudomonas putida*
15 KT2440 is a robust host for strong oxidative bioreactions (**14-16**), it exhibits the GRAS (i.e., generally
16 recognized as safe) status (**17**), and it has the inherent ability to grow on a wide range of (often,
17 difficult-to-degrade) substrates (**18-21**). Re-wiring its extant genetic features to extend its metabolic
18 potential –or even introducing new-to-Nature functions– is a task continuously undergoing in our
19 laboratory. In the present protocol, we detail all the experimental steps needed to either (i) construct
20 random mutant libraries by mini-Tn5 insertions to explore gene-function relationships, or (ii) deliver a
21 DNA cargo into a target chromosome, with the option of FLP-catalyzed removal of the antibiotic
22 resistance determinant.
23

23

24 **2. Materials**

25

26 **2.1. Bacterial strains and plasmids**

27

28 The bacterial strains and plasmids and vectors used in this protocol are described in Table 1 and 2,
29 respectively.
30

1 2.2. Culture media and reagents preparation

2

3 Unless otherwise stated, all the culture medium components and chemicals described below were
4 purchased from Sigma-Aldrich Co. (St. Louis, MO, USA). Whenever appropriate, please follow all
5 waste disposal regulations when disposing waste materials.

6

- 7 1. LB medium is used as the nutrient-rich culture medium in routine cultivations of both *P. putida* and
8 *Escherichia coli*. The components of LB medium (10 g of tryptone, 5 g of yeast extract, and 5 g of
9 NaCl) were dissolved and brought up to 1 L with deionized H₂O, and sterilized by autoclaving (20
10 min at 121°C and 1.05 kg/cm²). This culture medium can be indefinitely stored at room
11 temperature protected from light.
- 12 2. Nutritional selection is employed as a general strategy to counter-select for *P. putida*. M9 minimal
13 medium, supplemented with sodium citrate at 0.2% (w/v) as the sole carbon source (see **Note 1**)
14 and MgSO₄ at 2 mM, is used for this purpose since *E. coli* cannot grow on citrate. A 10× stock of
15 M9 salts is prepared by dissolving 42.5 g of Na₂HPO₄·2H₂O, 15 g of KH₂PO₄, 2.5 g of NaCl, and 5
16 g of NH₄Cl in deionized H₂O up to a final volume of 500 mL. A 20% (w/v) sodium citrate solution is
17 prepared by dissolving 20 g of anhydrous sodium citrate in deionized H₂O up to 100 mL, and a 1 M
18 MgSO₄ solution is prepared by dissolving 12 g of anhydrous MgSO₄ in deionized H₂O up to 100
19 mL. All these solutions are separately sterilized by autoclaving as indicated above, and can be
20 indefinitely stored at room temperature. Components are mixed and diluted as appropriate with
21 deionized H₂O to prepare M9 minimal medium immediately prior to use.
- 22 3. To elaborate solid media [containing 1.5% (w/v) agar], add 15 g of bacteriological agar (e.g.,
23 BactoAgar™; Becton-Dickinson Diagnostics Co., Sparks, MD, USA) to 1 L of LB medium and
24 autoclave it as indicated above. In the case of M9 minimal medium plates, prepare a 1.6% (w/v)
25 agar suspension in deionized H₂O, autoclave it separately from the other medium stock solutions,
26 and then mix an adequate amount of this suspension with the rest of the M9 minimal medium
27 components to reach a final agar concentration of 1.4% (w/v). Antibiotics and other additives are
28 added when the molten agarized medium reaches ca. 50°C. Distribute the agarized culture media
29 in plastic Petri dishes (25 mL of molten culture medium per 90-mm Petri dish), and let the medium
30 solidify at room temperature. Culture medium plates are prepared freshly immediately prior to use,
31 but they can be stored at 4°C (ideally for no longer than 1 week, especially if antibiotics were

1 added to the agarized culture medium).

- 2 4. All the antibiotics needed for bacterial selection in this protocol are prepared as concentrated stock
3 solutions in deionized H₂O at the concentrations indicated, sterilized by filtration (0.45 μm), and
4 stored at -20 °C for several months (see **Note 2**). The concentration of the stock solutions is as
5 follows: ampicillin (Ap), 150 mg/mL; kanamycin (Km), 50 mg/mL; streptomycin (Sm), 50 mg/mL;
6 gentamicin (Gm), 10 mg/mL; and carbenicillin (Cb), 500 mg/mL. The chloramphenicol (Cm) stock
7 solution is prepared at 30 mg/mL in 100% (v/v) ethanol. Unless indicated otherwise, all the
8 antibiotic stock solutions are considered to be 1000× concentrated.
- 9 5. To prepare 1× phosphate-buffered saline (PBS; 8 mM Na₂HPO₄, 1.5 mM KH₂PO₄, 3 mM KCl, and
10 137 mM NaCl, pH = 7.0), dissolve 1.44 g of Na₂HPO₄, 0.24 g of KH₂PO₄, 0.2 g of KCl, and 8 g of
11 NaCl in 800 mL of deionized H₂O, adjust the pH at 7.0 by dropwise addition of 6 N HCl, and bring
12 the volume to 1 L with deionized H₂O. Sterilize this buffer by autoclaving as indicated above, and
13 store it at room temperature.
- 14 6. In order to obtain electrocompetent cells of *P. putida*, prepare a 300 mM sucrose solution by
15 dissolving 10.27 g of sucrose in deionized H₂O up to a final volume of 100 mL, sterilize by filtration
16 (0.45 μm), and keep at room temperature.
- 17 7. To wash and prepare bacterial cells for mating, dilute the 1 M MgSO₄ solution described above
18 with sterile deionized H₂O to obtain a 10 mM MgSO₄ solution.
- 19 8. To maintain bacteria as frozen stocks, use 20% (v/v) glycerol in LB medium. Prepare this solution
20 by adding 118 mL of 85% (v/v) glycerol to 382 mL of LB medium, sterilize by filtration (0.45 μm),
21 and keep this solution at room temperature protected from light.

22

23 **2.3. DNA and general molecular biology techniques**

24

- 25 1. To purify plasmids from bacteria, we normally use the QIAprep Spin Miniprep™ kit (Qiagen Inc.,
26 Valencia, CA, USA) by following the manufacturer's instructions.
- 27 2. Colony PCR is routinely used as the source of template DNA for amplifications. Fresh bacterial
28 colonies (i.e., incubated for <24 h) are taken straight from the agar plate with a sterile toothpick
29 and dispersed into the PCR reaction tube containing deionized H₂O (see **Note 3**).
- 30 3. GoTaq™ Flexi DNA polymerase (Promega Corp., Madison, WI, USA) is routinely used for PCR
31 amplifications; however, any other DNA polymerase can be used for this purpose by following the

1 specific manufacturer's indications.

2 4. Prepare a 10 mM stock solution of deoxynucleotide triphosphates (dNTPs) containing equimolar
3 amounts of dATP, dCTP, dGTP, and dTTP in milliQ water (resistivity ≥ 18 M Ω /cm at 25°C); and
4 store the solution at -20 °C for up to 8-12 months.

5 5. The NucleoSpin™ Gel and PCR clean-up kit (Macherey-Nagel GmbH & Co. KG, Düren, Germany)
6 or the ExoSAP-IT™ PCR product cleanup kit (USB Molecular Biology, Affymetrix Ltd., Santa Clara,
7 CA, USA) are used for purification of PCR products.

8 6. Agarose gel electrophoresis is routinely used to identify, quantify, and purify DNA fragments. Basic
9 protocols are given elsewhere (22), and have to be adjusted for each specific purpose depending
10 on the application.

11

12 ***2.4. Primers used to localize and sequence the mini-transposon insertion point***

13

14 All the oligonucleotides needed for arbitrary PCR amplifications are indicated below, along with the
15 mini-transposon vectors used for the chromosomal insertions. Primers are purchased from Sigma-
16 Aldrich Co., as desalted, lyophilized DNA, and resuspended in the appropriate volume of milliQ H₂O to
17 obtain 5 μ M primer solutions. These solutions are aliquoted and stored at -20°C for several months.

18

19 ***2.4.1. Primers for arbitrary PCR amplifications***

20

21 Common to all vectors (i) ARB6: 5'-GGC ACG CGT CGA CTA GTA CMN NNN NNN NNA CGC C-3'.

22 This primer is used in arbitrary PCR, round 1 (23). In this sequence, *N*
23 represents any nucleotide.

24 (ii) ARB2: 5'-GGC ACG CGT CGA CTA GTA C-3'. This primer is used in
25 arbitrary PCR, round 2 (23).

26

27 ***2.4.2. Primers specific to the ME-O end of each mini-transposon***

28

29 pBAMD1-2 (i) ME-O-Km-Ext-F: 5'-CGT CTG TTT CAG AAA TAT GGC AT-3'. This primer
30 is used in arbitrary PCR, round 1 (9).

31 (ii) ME-O-Km-Int-F: 5'-ATC TGA TGC TGG ATG AAT TTT TC-3'. This primer

- 1 is used in arbitrary PCR, round 2, and for sequencing to map the integration
 2 point (9).
 3
- 4 pBAMD1-4 (i) ME-O-Sm-Ext-F: 5'-CTT GGC CTC GCG CGC AGA TCA G-3'. This primer
 5 is used in arbitrary PCR, round 1 (9).
 6 (ii) ME-O-Sm-Int-F: 5'-CAC CAA GGT AGT CGG CAA AT-3'. This primer is
 7 used in arbitrary PCR, round 2, and for sequencing to map the integration
 8 point (9).
 9
- 10 pBAMD1-6 (i) ME-O-Gm-Ext-F: 5'-GCA CTT TGA TAT CGA CCC AAG T-3'. This primer
 11 is used in arbitrary PCR, round 1 (9).
 12 (ii) ME-O-Gm-Int-F: 5'-TCC CGG CCG CGG AGT TGT TCG G-3'. This primer
 13 is used in arbitrary PCR, round 2, and for sequencing to map the integration
 14 point (9).
 15
- 16 pBELG and pBELK (i) pBEL-ME-O-Ext-F: 5'-CTG CGA CAT CGT ATA ACG TTA CTG GTT TC-3'.
 17 This primer is used in arbitrary PCR, round 1 (10).
 18 (ii) pBEL-ME-O-Int-F: 5'-GGG CGC TAT CAT GCC ATA CCG-3'. This primer
 19 is used in arbitrary PCR, round 2 (10).
 20
- 21 pBEXG and pBEXK (i) pBEX-ME-O-Ext-F: 5'-CTT CTT ACA TTT GGG ACG CTT CGC TG-3'.
 22 This primer is used in arbitrary PCR, round 1 (10).
 23 (ii) pBEX-ME-O-Int-F: 5'-CCT TCC GAC ACC CTG CGT CAA TG-3'. This
 24 primer is used in arbitrary PCR, round 2 (10).
 25
- 26 ***2.5.3. Primers specific to the ME-I end of each mini-transposon***
 27
- 28 pBAMD1-2 (i) pBAM-ME-I-Ext-R: 5'-CTC GTT TCA CGC TGA ATA TGG CTC-3'. This
 29 primer is used in arbitrary PCR, round 1 (7).
 30 (ii) pBAM-ME-I-Int-R: 5'-CAG TTT TAT TGT TCA TGA TGA TAT A-3'. This
 31 primer is used in arbitrary PCR, round 2, and for sequencing to map the

1 integration point (7).

2

3 pBAMD1-4 (i) ME-I-Sm-Ext-R: 5'-ATG ACG CCA ACT ACC TCT GAT A-3'. This primer
4 is used in arbitrary PCR, round 1 (9).

5 (ii) ME-I-Sm-Int-R: 5'-TCA CCG CTT CCC TCA TGA TGT T-3'. This primer
6 is used in arbitrary PCR, round 2, and for sequencing to map the integration
7 point (9).

8

9 pBAMD1-6 (i) ME-I-Gm-Ext-R: 5'-GTT CTG GAC CAG TTG CGT GAG-3'. This primer
10 is used in arbitrary PCR, round 1 (9).

11 (ii) ME-I-Gm-Int-R: 5'-GAA CCG AAC AGG CTT ATG TCA-3'. This primer
12 used in arbitrary PCR, round 2, and for sequencing to map the integration
13 point (9).

14

15 *2.4.4. Primers specific to the mini-transposon vector backbone*

16

17 Oligonucleotides annealing to specific sequences in SEVA vectors (12,13) are routinely used to detect
18 the presence of the Tn5-bearing mini-transposon vector backbone in transconjugant cells. If the
19 plasmid is present in such cells, a PCR amplification using primers PS4 [5'-CCA GCC TCG CAG AGC
20 AGG-3'] and PS5 [5'-CCC TGC TTC GGG GTC ATT-3'] generates a DNA amplicon of 225 bp.

21

22 *2.4.5. Primers to confirm elimination of antibiotic resistances by ectopic expression of the FLP* 23 *recombinase*

24

25 When employing the specialized pBELs or pBEXs mini-Tn5 vectors (10), the presence or the absence
26 of antibiotic resistance genes can be easily assessed by colony PCR with primers cFRT-Ab-R (5'-GAG
27 AAT AGG AAC TTC GGA ATA GG-3') in combination with either cKm-F (5'-CGG AAT GCT ATG CAG
28 ACG-3', when using pBELK or pBEXK) or cGm-F (5'-CCC GTA TGC CCA ACT TTG-3', when using
29 pBELG or pBEXG). The corresponding expected amplicon lengths are 736 bp (for pBELG and pBEXG)
30 and 535 bp (for pBELK and pBEXK).

31

1 2.5. Other laboratory material and equipment and standard procedures

2

3 1. Bacteria are routinely grown aerobically in 10-mL plastic test tubes (e.g., 16×100 mm, round
4 bottom tubes; E&K Scientific Products Inc., Santa Clara, CA, USA) containing 3 mL of the
5 corresponding liquid culture medium. *P. putida* KT2440 is incubated at 30°C and *E. coli* strains are
6 incubated at 37°C with rotary agitation at 170 rpm.

7 2. For electroporation, 2-mm gap width cuvettes (e.g., Gene Pulser™/Micropulser™ electroporation
8 cuvettes; Bio-Rad Laboratories Inc., Hercules, CA, USA) and a bacterial electroporation system
9 (e.g., MicroPulser™; Bio-Rad Laboratories Inc.) are used.

10 3. For filter-assisted bacterial matings, we recommend the use of mixed cellulose esters filter disks of
11 0.45-µm pore-size and 25-mm diameter (EMD Millipore Corp., Billerica, MA, USA), Millipore
12 SX0002500 Swinnex™ Syringae filter holders for 25-mm diameter filters, 20-mL Luer-lock™ tip
13 syringes (Becton-Dickinson Diagnostics Co.), and blunt-end filter forceps (e.g., XX6200006P
14 forceps; EMD Millipore Corp.) to manipulate the filter discs.

15 4. Thermocycler (e.g., T100™ Thermal Cycler; Bio-Rad Laboratories Inc.).

16 5. Sterile, round plastic Petri dishes (e.g., Nunc™ Lab-Tek™ Petri dishes; Thermo Fisher Scientific
17 Inc., Waltham, MA, USA), either of regular size (i.e., diameter = 90 mm) or bigger plates (i.e.,
18 diameter = 140 mm), for screening of colonies or during the generation of random mutant libraries.

19 6. To spread bacteria onto agar plates, use 5-10 sterile 3-mm diameter glass beads (VWR
20 International, Radnor, PA, USA). Once glass beads are used, immediately dispose them off into a
21 flask with 70% (v/v) ethanol. To recover the glass beads for subsequent use, rinse them twice with
22 deionized H₂O, let them dry overnight at room temperature, and sterilize the beads by autoclaving.

23

24 3. Methods

25

26 The mini-transposon vectors described in the present protocol have two principal uses: (i) the
27 generation of libraries of random mutants to correlate a particular and observable phenotype to a
28 specific gene (**1,7,8,24**); or (ii) the introduction of heterologous gene(s) randomly into the chromosome
29 of a target Gram-negative bacterium (**2,9,10**). In the first application, a typical random mutagenesis
30 protocol involves three steps: (i) *delivery* of the non-replicative plasmid bearing the mini-transposon
31 into a recipient strain, (ii) *selection* of transconjugants carrying the transposon, and (iii) *storing* the

1 library for future uses or directly *identifying* the insertion point of the mini-transposon in the genome of
2 selected recipient cells. In the second case, when the objective is to introduce heterologous DNA into a
3 bacterial genome, a previous step is included in which the gene (or genes) of interest has to be cloned
4 into the multiple cloning site of the mini-Tn5 delivery plasmid. Steps (i), (ii), and (iii) are then followed
5 as described in the specific procedure below. Finally, when using any of the pBELs or pBEXs mini-Tn5
6 plasmids (**10**) to deliver a DNA cargo under a controlled expression system, a final extra step is added
7 *ad libitum* to remove the antibiotic marker of the mini-Tn5 cassette.

8

9 The first step of the protocol (i.e., introduction of the mini-transposon plasmid into the recipient strain),
10 could be done either by mating or electroporation. Specific protocols to perform either delivery
11 technique are described below, and a specific application example, in which plasmid pBAMD1-2 (**9**)
12 was employed to generate a library of random insertion mutants of *P. putida* KT2440, is discussed.
13 The main steps of the protocol are summarized in Fig. 1.

14

15 **3.1. Mini-Tn5 delivery into *P. putida* by conjugation**

16

17 Whenever possible, we recommend to use this delivery method since it is more efficient than
18 electroporation of plasmids. Conjugation requires cell contact to transfer DNA from a donor cell to a
19 recipient strain. To establish such intimate contact, donor bacteria produce the conjugative pilus (i.e., a
20 type IV secretion system) that ultimately retracts, bringing both cells together. A number of proteins of
21 the donor bacterial cell form a bridge between both donor and recipient cells forming a mating pair (i.e.,
22 Mpf proteins, for *mating pair formation*). Then, the *relaxosome* (i.e., a complex formed by a relaxase
23 and auxiliary proteins) recognizes the origin of transfer (*oriT*) sequence and move one strand of the
24 target DNA to the recipient cell [for a review on the biology behind this process, please see Zechner *et*
25 *al.* (**25**) and Ilangovan *et al.* (**26**) and references therein].

26

27 To perform a conjugation experiment, one just needs to bring together the donor cell (i.e., bearing the
28 mini-transposon plasmid), the recipient cell (i.e., the target bacterium), and a helper bacterial strain to
29 assist and catalyze the mating process. The mating helper is an *E. coli* strain that provides the
30 conjugation machinery. Typically, this molecular machinery is derived from the IncP α plasmid RP4
31 (also known as RK2 or RP1), and involves the mobilization (*mob*) and transfer (*tra*) functions (**27,28**),

1 supplied *in trans*. There are two basic types of helper strains, which express the *mob/tra* functions
2 either (i) in a plasmid (e.g., *E. coli* HB101 carrying plasmid pRK600; Table 1 and 2), or (ii) integrated in
3 the genome (e.g., *E. coli* S17-1 λ *pir*, *E. coli* SM10 λ *pir*, or *E. coli* MFD λ *pir*; Table 1). When using the
4 former type of helper *E. coli* strain, the user needs to include three bacterial strains in the mating
5 process (i.e., setting up a triparental mating). A triparental mating offers the possibility of changing the
6 donor *E. coli* strain (which should contain the *pir* gene as a λ *pir* lysogen, e.g., *E. coli* CC118 λ *pir* or
7 DH5 α λ *pir*) in order to favor counter selection of transconjugants as needed. In the case of performing
8 a biparental mating, the protocol is exactly the same as per the triparental mating procedure, but using
9 only two bacterial strains in the mixture (i.e., the recipient strain and the mobilizing donor cell, that in
10 addition to the mini-Tn5 plasmid also has the *mob/tra* functions integrated in the genome).

11

12 As indicated above, in the present protocol we describe a triparental mating using *P. putida* KT2440 as
13 the recipient strain and pBAMD1-2 as the mini-Tn5 delivery plasmid (9). In order to properly generate a
14 random mutagenesis library when working with other recipient bacterial species, it is important to
15 perform several previous tests to determine the optimal experimental conditions for successful DNA
16 transfer, since the expected number of transconjugant colonies depends on several factors such as the
17 nature of the recipient species, the initial amount of recipient cells, the mixing ratio of recipient to donor
18 cells, and the mating incubation time. With the help of these prior experiments, the user should be able
19 to set the appropriate experimental conditions and to estimate the number of plates needed to obtain a
20 saturated library.

21

- 22 1. To prepare the mating mixture, grow the following strains overnight as indicated:
- 23 (i) **Donor:** *E. coli* CC118 λ *pir* (carrying plasmid pBAMD1-2) grown in LB medium added with Ap at
24 150 μ g/ml. Incubate for 18 h at 37°C with rotary agitation. These cells bear the mobilizable and
25 non-replicative plasmid with the Tn5 mini-transposon (see **Note 4**).
- 26 (ii) **Mating helper:** *E. coli* HB101 (carrying plasmid pRK600) grown in LB medium added with Cm
27 at 30 μ g/ml. Incubate for 18 h at 37°C with rotary agitation. This bacterium provides the plasmid
28 with the mobilization (*mob*) and transfer (*tra*) functions, encoded in plasmid pRK600.
- 29 (iii) **Recipient:** *P. putida* KT2440 grown in LB medium at 30°C with rotary agitation (see **Note 5**).
- 30 2. Measure the optical density at 600 nm (OD₆₀₀) of the bacterial cultures and adjust the bacterial
31 suspensions to an OD₆₀₀ of 1 with PBS in a final volume of 1 mL in a 1.5-mL Eppendorf tube.

- 1 3. Centrifuge the cultures at $7200\times g$ for 2 min at room temperature, discard the supernatant, and re-
2 suspend the sediment in 1 mL of 10 mM MgSO_4 to wash the cells.
- 3 4. Mix the three bacterial suspensions in a 1:1:1 ratio (i.e., 150 μL of each suspension) in a test tube
4 containing 4.55 mL of 10 mM MgSO_4 . The final OD_{600} should be ≈ 0.03 (see **Note 6**).
- 5 5. Pass the 5-mL cell suspension through a filter disk (0.45- μm pore-size, 25-mm diameter) using a
6 20-mL sterile syringe (see **Note 7**). Discard the flow-through and laid the filter, in sterile conditions,
7 onto an LB medium agar plate (cells facing up). Incubate the plate containing the filter (lid facing
8 up) at 30 °C during the desired mating time (4 h, 6 h, or even 24 h) (see **Note 8**).
- 9 6. Gently take the filter from the agar plate with tweezers [blunt-end filter forceps, previously sterilized
10 by quickly dipping them in 70% (v/v) ethanol and flaming] and place it in a 10-mL test tube
11 containing 5 mL of 10 mM MgSO_4 .
- 12 7. Re-suspend the cells in the mating mixture from the filter by vigorous vortexing (at least for 1 min)
13 and plate appropriate dilutions (see **Note 9**) onto M9 minimal medium plus citrate at 0.2% (w/v)
14 and Km at 50 $\mu\text{g}/\text{mL}$ (i.e., selective culture medium for transconjugant *P. putida* cells harboring the
15 mini-transposon) (see **Notes 10 and 11**).

16

17 3.2. *Mini-Tn5 delivery into P. putida by electroporation*

18

19 If no other choice is available (or just for cases when a DNA cargo is to be integrated into a target
20 genome, where higher frequencies are not that important as they are for the construction of mutant
21 libraries) then electrotransformation is the preferred alternative, mainly due to the fastness and
22 simplicity of the protocol thereof. This technique is based in the transient permeabilization of the cell
23 membrane, that allows for the entry of DNA after applying a high electric field (**24,29,30**).

24

- 25 1. Inoculate a 100-mL Erlenmeyer flask containing 20 mL of LB medium with *P. putida* KT2440 from
26 a fresh LB medium agar plate (or directly from a frozen stock, by scrapping the surface of the stock
27 with a sterile toothpick). Let the cells grow overnight (e.g., 18-24 h) aerobically (170 rpm) at 30°C.
- 28 2. Transfer the saturated culture to a 50-mL Falcon tube and centrifuge it at $3220\times g$ for 10 min at
29 room temperature.
- 30 3. Discard the supernatant, add 10 mL of 300 mM sucrose and softly resuspend the cell sediment;
31 then, centrifuge the suspension at $3220\times g$ for 10 min at room temperature.

- 1 4. Remove the supernatant and add 1 mL of 300 mM sucrose, resuspend the cells, and transfer the
2 suspension to a 2-mL sterile Eppendorf tube. Centrifuge at $7200\times g$ for 3 min at room temperature.
- 3 5. Remove the supernatant, add 800 μL of 300 mM sucrose, resuspend the cells, and centrifuge the
4 suspension at $7200\times g$ for 3 min at room temperature. Repeat this washing step once more.
- 5 6. Remove the supernatant and add 500 μL of 300 mM sucrose to resuspend the sediment and to
6 obtain a concentrated cell suspension (after the final resuspension step, the concentration of
7 electrocompetent bacteria should be $\approx 5\times 10^{10}$ cells/mL).
- 8 8. Transfer 100 μL of the electrocompetent cell suspension to a 1.5-mL sterile Eppendorf tube and
9 add ≈ 500 ng of plasmid pBAMD1-2 (in < 10 μL). Pipet the plasmid DNA-cell suspension mix to a
10 2-mm gap width electroporation cuvette. Care has to be taken to avoid the formation of bubbles at
11 this step, which would reduce the overall efficiency of the electroporation process.
- 12 9. Place the cuvette in the MicroPulser™ apparatus, set the electroporation program to EC2, and
13 proceed to electroporate. With these working conditions and using an optimum electric pulse (a
14 single pulse of 2.5 kV with a field strength of 12.5 kV/cm), a time constant (τ) between 4 and 5 ms
15 should be obtained.
- 16 10. Immediately after the electric shock, add 900 μL of LB medium to the cuvette and then transfer the
17 cells to a sterile 1.5-mL Eppendorf tube. Incubate the cells aerobically for 3 h at 30 °C.
- 18 11. Spread dilutions of the cell suspension obtained in the step above onto LB medium agar plates
19 containing Km at 50 $\mu\text{g/mL}$. Since no *E. coli* cells are used in this procedure, there is no need for
20 nutritional selection as performed in mating experiments.

21

22 **3.3. Isolation and mapping the mini-transposon genomic insertion landing sites**

23

- 24 1. If specifically looking for particular phenotypes, select interesting colonies based on a trait (e.g.,
25 morphology or color) different from that observed in the wild-type cells, and streak them with a
26 sterile toothpick onto both (i) M9 minimal medium plates containing 0.2% (w/v) citrate and 50
27 $\mu\text{g/mL}$ Km, and (ii) M9 minimal medium plates containing 0.2% (w/v) citrate and 500 $\mu\text{g/mL}$ Ap.
28 This process is aimed to differentiate between genuine transposition events (i.e., Km^R colonies)
29 from spurious mini-Tn5 plasmid co-integration incidents (i.e., Km^R and Ap^R colonies). Incubate the
30 plates overnight at 30°C.
- 31 2. Select Km^R and Ap^S clones. Also, use colony PCR amplifications with oligonucleotides PS4 and

- 1 PS5 to confirm the absence of the delivery plasmid backbone.
- 2 3. Re-streak selected colonies several times onto M9 minimal medium plates containing 0.2% (w/v)
- 3 citrate and 50 $\mu\text{g/mL}$ Km to make sure of working with pure isolated clones.
- 4 5. Make a frozen stock in 20% (v/v) glycerol in LB medium of the selected mutants and store the
- 5 resulting stocks at -80°C . Bacterial frozen stocks can be prepared by growing the cells of interest
- 6 onto LB medium plates (with the appropriate antibiotics as necessary) overnight, and adding 2 mL
- 7 of 20% (v/v) glycerol in LB medium thereafter. Cells are gently scrapped from the surface by using
- 8 a sterile glass rod (i.e., a Drigalski spatula). One mL of the resulting suspension is then transferred
- 9 into a cryotube (e.g., a 1.8-mL Nunc™ CryoTubes™ cryogenic vial, round bottom). Cells can be
- 10 stored at -80°C under these conditions for several years without significant loose of viability,
- 11 provided that the bacterial stock is not repeatedly frozen and thawed.
- 12 6. Take mutant clones from the frozen stock and streak the cells onto LB medium agar plates
- 13 containing 50 $\mu\text{g/mL}$ Km. Grow the cells overnight at 30°C .
- 14 7. In order to genetically analyze the transconjugants, firstly choose one of the mini-transposon ends
- 15 (i.e., ME-I or ME-O) to determine its insertion place in the genome and then perform arbitrarily-
- 16 primed colony PCR (**31**). The DNA sequence of the primers needed to perform arbitrarily primed
- 17 PCR amplifications when using the different mini-Tn5 plasmids is described in *Section 2.4*. (see
- 18 **Note 12**).
- 19 8. Prepare a PCR reaction mix on ice as per the following recipe. Note that most of the components
- 20 indicated in the recipe are provided along with the commercial *Taq* DNA polymerase. Thoroughly
- 21 vortex each concentrated solution before pipetting into the PCR reaction mix.
- 22 - 5 μL of 5 \times Green or Colorless Go*Taq*™ reaction buffer
- 23 - 1.5 μL of 25 mM MgCl_2
- 24 - 0.5 μL dNTPs (10 mM)
- 25 - 0.5 μL of dimethyl sulfoxide (when performing amplifications from high G+C DNA templates)
- 26 - 1 μL of 5 μM arbitrary primer
- 27 - 1 μL of 5 μM mini-transposon primer (i.e., ME-I or ME-O)
- 28 - 0.2 μL of 5 U/ μL Go*Taq*™ Flexi DNA polymerase
- 29 9. Aliquot 15.3 μL of sterile deionized H_2O into each PCR tube.
- 30 10. Transfer fresh colonies from agar plates directly into the PCR reaction tube with a sterile toothpick.
- 31 11. Distribute 9.7 μL of the PCR reaction mix into each PCR tube.

- 1 12. The primers needed for round 1 of the arbitrarily primed PCR amplification are ARB6 together with
2 the external ME-I or ME-O primers (i.e., either ME-I-Ext or ME-O-Ext).
- 3 13. The settings for round 1 of the arbitrarily primed PCR amplification are as follows:
4 - 5 min at 95°C
5 - 30 s at 95°C, 30 s at 30°C, and 1.5 min at 72°C (6×)
6 - 30 s at 95°C, 30 s at 45°C, and 1.5 min at 72°C (30×)
- 7 14. Directly take 1 µL of the PCR after running round 1 (i.e., no need to check for positive
8 amplifications in an agarose gel) and use it as the template for round 2 of arbitrary PCR. In this
9 round, use primer ARB2 together with the internal ME-I or ME-O primers (i.e., either ME-I-Int or
10 ME-O-Int) (see **Note 13**). Prepare the PCR reaction mix for round 2 as indicated in step 8 above.
- 11 15. The settings for round 2 of the arbitrarily primed PCR amplification are as follows:
12 - 1 min at 95°C
13 - 30 s at 95°C, 30 s at 52°C, and 1.5 min at 72°C (30×)
14 - 4 min at 72°C
- 15 16. Clean up the PCR product from the second round of the arbitrary PCR amplification using either
16 the NucleoSpin™ Gel and PCR clean-up kit or the ExoSAP-IT™ PCR product cleanup kit.
- 17 17. Send the DNA product to sequence (**32,33**) with the ME internal primer used in round 2 of the
18 arbitrary PCR.
- 19 18. Analyze the sequencing results. Start by identifying the DNA sequence of the mini-transposon end
20 (i.e., either ME-I or ME-O) (see **Note 14**), and then trim that part and select the rest of the DNA
21 sequence. Use the BlastN program (**34**), available on-line at www.pseudomonas.com/blast/set
22 (**35**), to map the precise genomic coordinates of the mini-transposon insertion (see **Note 15**).
- 23 19. Once an interesting mutant is spotted, in which the phenotype-gene has been identified, it is
24 always recommended to complement that mutant back with the identified gene(s) to rule out the
25 occurrence of polar effects, since mini-Tn5 insertions are known to alter the expression of
26 neighbouring genes (**36,37**).

27

28 **3.4. Eliminating the antibiotic resistance marker of specialized mini-Tn5 vectors**

29

30 When using any of the pBELs or pBEXs mini-Tn5 vectors (Table 2) to introduce heterologous DNA
31 under the control of an expression system (i.e., $\text{LacI}^{\text{O}}/P_{trc}$ or XylS/P_m) (**10**), the genes conferring

1 resistance to Km (*aphA*) or Gm (*aacC1*) in these transposons can be removed as they are flanked by
2 FLP recombinase target (*FRT*) sequences (**38**). This layout offers the possibility to the user of
3 eliminating that marker by means of ectopic expression of the FLP recombinase from *Saccharomyces*
4 *cerevisiae* using plasmid pFLP2 (Table 2). The expression of the FLP recombinase in plasmid pFLP2
5 is driven by the strong, rightward λ promoter (located within the *FLP-cI857* intergenic region) and is
6 regulated by the temperature-sensitive, *cI857*-encoded λ repressor (**39**).

7

- 8 1. Select a transconjugant *P. putida* clone in which the insertion place of the mini-transposon has
9 been successfully localized.
- 10 2. Introduce plasmid pFLP2 (see **Note 16**) into this selected clone by either mating or electroporation
11 as described above.
- 12 3. Plate the cells on M9 minimal medium plates added with sodium citrate at 0.2% (w/v) and Cb at
13 500 $\mu\text{g}/\text{mL}$. Incubate the plates overnight at 30°C. If no discernible colonies are observed after this
14 incubation period, try lowering the Cb concentration to 350 $\mu\text{g}/\text{mL}$.
- 15 4. Select two or three independent colonies and re-streak them on M9 minimal medium plates added
16 with sodium citrate at 0.2% (w/v) and Cb at 500 $\mu\text{g}/\text{mL}$. Incubate the plates overnight at 30°C.
- 17 5. Pick single colonies and check for Km or Gm sensitivity and Cb resistance in LB medium plates
18 containing these antibiotics. Double check for the removal of the antibiotic gene by colony PCR
19 using the primers described in *Section 2.4.5*. (i.e., *cFRT-Ab-R* and either *cKm-F* or *cGm-F*). Such
20 PCR should give no amplification. If possible, use primers annealing within the gene(s) delivered in
21 the mini-transposon cassette to conduct a colony PCR of the antibiotic-sensitive clone to make
22 sure that the gene(s) of interest have been stably inserted into the target chromosome.
- 23 6. Cure plasmid pFLP2 from the selected clone by performing several (at least three) cycles of
24 growth in LB medium without any antibiotic.
- 25 7. Plate cells onto M9 minimal medium plates added with 0.2% (w/v) sodium citrate.
- 26 8. Pick single colonies and double re-streak onto M9 minimal medium plates added with 0.2% (w/v)
27 sodium citrate plus Cb at 500 $\mu\text{g}/\text{mL}$.
- 28 9. Select Cb-sensitive clones and store them as frozen stocks at -80°C .

1 3.5. Preparing and storing a mutant library of mini-Tn5 insertions in *P. putida*

2

3 After obtaining a random mutagenesis library, it is always useful to save it for later analyses. The steps
4 below indicate the procedure to store the library after introduction of plasmid pBAMD1-2 in strain
5 KT2440 as explained in the preceding sections.

6

- 7 1. Spread dilutions of the triparental mating mixture onto selective agar plates in order to obtain an
8 approximate number of ≈ 3000 transconjugant colonies per plate in a regularly sized (i.e., 90 mm)
9 Petri dish. Estimate the number of plates needed to obtain a non-saturated mutant library as
10 indicated by Liberati *et al.* (37).
- 11 2. Add 2.5 mL of LB medium containing 20% (v/v) glycerol to each overnight-incubated plate and,
12 with the aid of an inoculation loop or a Drigalski spatula gently scrap the cells from the agar
13 surface. Tilt the plate and collect 1-mL of the bacterial suspension with a micropipet (see **Note 17**).
- 14 3. Mix the liquid content collected from all the plates, aliquot the resulting suspension in several
15 cryotube vials, and store the library as a series of frozen stocks at -80°C .

16

17 4. Notes and troubleshooting

18

- 19 1. The appropriate culture medium composition has to be defined to select against *E. coli*
20 donor/mating helper cells when using other bacterial species as the target strain. As a general rule,
21 try to make use of specific carbon sources in which only the recipient strain grows or take
22 advantage of the auxotrophies of the *E. coli* donor/mating helper cells [e.g., most of the laboratory
23 *E. coli* strains need thiamine-HCl to grow (40)].
- 24 2. Avoid repeated freezing and thawing of antibiotic solutions as they may lose effectiveness. We
25 routinely distribute the stock solutions in 0.5-mL working aliquots that are used just a couple of
26 times before discarding them.
- 27 3. If no amplification is obtained through colony PCR, genomic DNA can be isolated with a
28 commercial kit (e.g., UltraClean™ Microbial DNA isolation kit; MoBio Laboratories Inc., Carlsbad,
29 CA, USA) and used as the template for amplifications.
- 30 4. It is very important to grow the donor bacterial strain in the presence of the antibiotic for which the
31 plasmid backbone carries a resistance gene (e.g., Ap) to avoid inadvertent selection of transposed

- 1 donor cells. Note that there is a vector derived from pBAM1 which carries a promoterless *gfp* gene
2 (plasmid pBAM1-GFP), which allows for the visual inspection of successful *gene::gfp* fusions after
3 the transposition event.
- 4 5. In some cases, incubating the recipient strain at high temperatures (40°C-42°C) for a few hours
5 before mating is known to increase the efficiency of the process by inactivating its endogenous
6 DNA restriction machinery.
- 7 6. Different ratios of the bacterial strains to be included in the triparental mating could also be tested if
8 needed (e.g., by increasing the amount of donor cells). To do this, simply adjust the volume of
9 each bacterial suspension appropriately with 10 mM MgSO₄ to bring the final volume to 5 mL, and
10 proceed as indicated. In the case of integrating DNA cargoes into the bacterial genome, where
11 there is no need of the high numbers of transconjugant colonies usually required for random
12 mutant libraries, one can use just 100 μL of each overnight cultures (i.e., without adjusting the
13 OD₆₀₀ of the individual cultures).
- 14 7. If a filter system for bacterial matings is not available, one can simply mix the three bacterial strains
15 in a 1.5-mL Eppendorf tube (e.g., 150 μL of each bacterial suspension adjusted at OD₆₀₀ = 1),
16 centrifuge the cells at 7200×g, discard the supernatant, and resuspend the sediment in 25 μL of 10
17 mM MgSO₄ (i.e., a small buffer volume to maximize cell contact). The 25-μL mating mix can be
18 laid onto a 0.45-μm filter disc onto an LB medium plate, or be directly spotted onto the surface of
19 an LB medium plate. In the later case, cells can easily recovered using an inoculation loop and
20 resuspended in 10 mM MgSO₄ before plating on a selective culture medium.
- 21 8. When creating non-saturated random mutant libraries it is better to use shorter incubation times to
22 maintain cell divisions of transconjugants to a minimum.
- 23 9. Depending on the purpose of the experiment, different Petri dishes sizes could be used to recover
24 more transconjugant cells per plate. Adjust the plating volume accordingly.
- 25 10. It is also important to plate (i) the donor strain, (ii) the mating helper, and (iii) the recipient strain
26 onto the selective culture medium used to recover transconjugants. These three bacterial strains
27 should not grow in the selective culture medium (i.e., they are used as negative controls).
- 28 11. Take into account that different mini-Tn5-bearing plasmids need other antibiotics (e.g., Sm at 80
29 μg/mL or Gm at 10 μg/mL in the case of plasmids pBAMD1-4 and pBAMD1-6, respectively) to
30 select for positive transconjugants.
- 31 12. Note that the specific ME primers have to be chosen depending on the mini-Tn5 plasmid used for

- 1 insertions and on the selected ME-end.
- 2 13. Clean-up the PCR products obtained after the first round of arbitrarily primed PCR with a
3 commercial kit to eliminate unbound primers in the case of experiencing problems (e.g., no
4 amplification in the second round of arbitrarily primed PCR).
- 5 14. If the insertion of transposon cannot be unequivocally mapped using the protocol and primers
6 suggested here, select other set of arbitrary primers, such as primer ARB1 (**23**) or even others as
7 described by Das *et al.* (**31**). Alternatively, a new custom arbitrary primer could be designed by
8 changing the five nucleotides at the 3'-end of the oligonucleotide sequence to match the G+C
9 content of the recipient bacterial strain, thereby increasing the frequency of appearance of that
10 motif in the target genome.
- 11 15. For other target bacterial species, use the BlastN tool against the genome of the desired recipient
12 strain. If the complete genome sequence of your favorite microorganism is not available, perform a
13 more general BlastN search in order to identify homologous genes or sequences in related
14 species. Specific primers could then be designed on the basis of these results to sequence the
15 exact locus in which the mini-Tn5 insertion has occurred.
- 16 16. Other plasmids can be used for the ectopic expression of the FLP recombinase, such as plasmid
17 pBBFLP (**41**). The procedure to be followed in this case is essentially the same as the one
18 described in the main protocol, but using tetracycline (at 15 $\mu\text{g}/\text{mL}$) instead of Cb to select for the
19 presence of the pBBFLP helper plasmid. If the insertion of pBELs or pBEXs vectors is carried out
20 in *E. coli*, plasmid pCP20 (**38**) is recommended for the FLP-dependent removal of antibiotic-
21 resistance determinants after transposition.
- 22 17. It is a good procedure to perform several independent matings in order to yield a representative
23 random mutant library.
- 24

25 Acknowledgments

26

27 The work described in this protocol was supported by the CAMBIOS Project of the Spanish Ministry of
28 Economy and Competitiveness (RTC-2014-1777-3), the ST-FLOW (FP7-KBBE-2011-5-289326),
29 EVOPROG (FP7-ICT-610730), ARISYS (ERC-2012-ADG-322797), and EmPowerPutida (EU-H2020-
30 BIOTEC-2014-2015-6335536) Contracts of the European Union, and the PROMPT Project of the

1 Autonomous Community of Madrid (CAM-S2010/BMD-2414). The authors declare that there is no
2 conflict of interest. All the bacterial strains and plasmids described are available upon request.

3

4 References

5

- 6 1. de Lorenzo V, Herrero M, Jakubzik U, Timmis KN (1990) Mini-Tn5 transposon derivatives for
7 insertion mutagenesis, promoter probing, and chromosomal insertion of cloned DNA in Gram-
8 negative eubacteria. *J Bacteriol* 172: 6568-6572.
- 9 2. de Lorenzo V, Timmis KN (1994) Analysis and construction of stable phenotypes in Gram-negative
10 bacteria with Tn5- and Tn10-derived minitransposons. *Methods Enzymol* 235: 386-405.
- 11 3. Reznikoff WS (2008) Transposon Tn5. *Annu Rev Genet* 42: 269-286.
- 12 4. Berg DE (1989) Transposon Tn5. In: Berg DE, Howe MM, editors. *Mobile DNA*, Washington, D.C.:
13 American Society for Microbiology Press. pp. 185-210.
- 14 5. Reznikoff WS (2006) Tn5 transposition: a molecular tool for studying protein structure-function.
15 *Biochem Soc Trans* 34: 320-323.
- 16 6. Phadnis SH, Sasakawa C, Berg DE (1986) Localization of action of the IS50-encoded transposase
17 protein. *Genetics* 112: 421-427.
- 18 7. Martínez-García E, Calles B, Arévalo-Rodríguez M, de Lorenzo V (2011) pBAM1: an all-synthetic
19 genetic tool for analysis and construction of complex bacterial phenotypes. *BMC Microbiol* 11: 38.
- 20 8. de Lorenzo V, Herrero M, Sánchez JM, Timmis KN (1998) Mini-transposons in microbial ecology
21 and environmental biotechnology. *FEMS Microbiol Ecol* 27: 211-224.
- 22 9. Martínez-García E, Aparicio T, de Lorenzo V, Nikel PI (2014) New transposon tools tailored for
23 metabolic engineering of Gram-negative microbial cell factories. *Front Bioeng Biotechnol* 2: 46.
- 24 10. Nikel PI, de Lorenzo V (2013) Implantation of unmarked regulatory and metabolic modules in
25 Gram-negative bacteria with specialised mini-transposon delivery vectors. *J Biotechnol* 163: 143-
26 154.
- 27 11. Schweizer HP (2003) Applications of the *Saccharomyces cerevisiae* Flp-FRT system in bacterial
28 genetics. *J Mol Microbiol Biotechnol* 5: 67-77.
- 29 12. Martínez-García E, Aparicio T, Goñi-Moreno A, Fraile S, de Lorenzo V (2014) SEVA 2.0: an update
30 of the Standard European Vector Architecture for de-/re-construction of bacterial functionalities.
31 *Nucleic Acids Res* 43: D1183-D1189.

- 1 13. Silva-Rocha R, Martínez-García E, Calles B, Chavarría M, Arce-Rodríguez A, et al. (2012) The
2 Standard European Vector Architecture (SEVA): a coherent platform for the analysis and
3 deployment of complex prokaryotic phenotypes. *Nucleic Acids Res* 41: D666-D675.
- 4 14. Nikel PI, Martínez-García E, de Lorenzo V (2014) Biotechnological domestication of
5 pseudomonads using synthetic biology. *Nat Rev Microbiol* 12: 368-379.
- 6 15. Nikel PI, de Lorenzo V (2014) Robustness of *Pseudomonas putida* KT2440 as a host for ethanol
7 biosynthesis. *New Biotechnol* 31: 562-571.
- 8 16. Benedetti I, de Lorenzo V, Nikel PI (2016) Genetic programming of catalytic *Pseudomonas putida*
9 biofilms for boosting biodegradation of haloalkanes. *Metab Eng* 33: 109-118.
- 10 17. Timmis KN (2002) *Pseudomonas putida*: a cosmopolitan opportunist *par excellence*. *Environ*
11 *Microbiol* 4: 779-781.
- 12 18. Nikel PI, Chavarría M, Fuhrer T, Sauer U, de Lorenzo V (2015) *Pseudomonas putida* KT2440
13 strain metabolizes glucose through a cycle formed by enzymes of the Entner-Doudoroff, Embden-
14 Meyerhof-Parnas, and pentose phosphate pathways. *J Biol Chem* 290: 25920-25932.
- 15 19. Nikel PI, Kim J, de Lorenzo V (2014) Metabolic and regulatory rearrangements underlying glycerol
16 metabolism in *Pseudomonas putida* KT2440. *Environ Microbiol* 16: 239-254.
- 17 20. Nikel PI, Romero-Campero FJ, Zeidman JA, Goñi-Moreno A, de Lorenzo V (2015) The glycerol-
18 dependent metabolic persistence of *Pseudomonas putida* KT2440 reflects the regulatory logic of
19 the GlpR repressor. *mBio* 6: e00340-00315.
- 20 21. Nikel PI, Silva-Rocha R, Benedetti I, de Lorenzo V (2014) The private life of environmental
21 bacteria: pollutant biodegradation at the single cell level. *Environ Microbiol* 16: 628-642.
- 22 22. Makovets S (2013) Basic DNA electrophoresis in molecular cloning: a comprehensive guide for
23 beginners. *Methods Mol Biol* 1054: 11-43.
- 24 23. Pratt LA, Kolter R (1998) Genetic analysis of *Escherichia coli* biofilm formation: roles of flagella,
25 motility, chemotaxis and type I pili. *Mol Microbiol* 30: 285-293.
- 26 24. Martínez-García E, de Lorenzo V (2012) Transposon-based and plasmid-based genetic tools for
27 editing genomes of Gram-negative bacteria. *Methods Mol Biol* 813: 267-283.
- 28 25. Zechner EL, Lang S, Schildbach JF (2012) Assembly and mechanisms of bacterial type IV
29 secretion machines. *Philos Trans R Soc Lond B Biol Sci* 367: 1073-1087.
- 30 26. Ilangovan A, Connery S, Waksman G (2015) Structural biology of the Gram-negative bacterial
31 conjugation systems. *Trends Microbiol* 23: 301-310.

- 1 27. Álvarez-Martínez CE, Christie PJ (2009) Biological diversity of prokaryotic type IV secretion
2 systems. *Microbiol Mol Biol Rev* 73: 775-808.
- 3 28. Babic A, Guérout AM, Mazel D (2008) Construction of an improved RP4 (*RK2*)-based conjugative
4 system. *Res Microbiol* 159: 545-549.
- 5 29. Iwasaki K, Uchiyama H, Yagi O, Kurabayashi T, Ishizuka K, et al. (1994) Transformation of
6 *Pseudomonas putida* by electroporation. *Biosci Biotechnol Biochem* 58: 851-854.
- 7 30. Choi KH, Kumar A, Schweizer HP (2006) A 10-min method for preparation of highly
8 electrocompetent *Pseudomonas aeruginosa* cells: application for DNA fragment transfer between
9 chromosomes and plasmid transformation. *J Microbiol Methods* 64: 391-397.
- 10 31. Das S, Noe JC, Paik S, Kitten T (2005) An improved arbitrary primed PCR method for rapid
11 characterization of transposon insertion sites. *J Microbiol Methods* 63: 89-94.
- 12 32. Zimmermann J, Voss H, Schwager C, Stegemann J, Ansorge W (1988) Automated Sanger
13 dideoxy sequencing reaction protocol. *FEBS Lett* 233: 432-436.
- 14 33. Shendure JA, Porreca GJ, Church GM, Gardner AF, Hendrickson CL, et al. (2011) Overview of
15 DNA sequencing strategies. *Curr Prot Mol Biol* 96: 7.1.1-7.1.23.
- 16 34. Altschul SF, Gish W, Miller W, Myers EW, Lipman DJ (1990) Basic local alignment search tool. *J*
17 *Mol Biol* 215: 403-410.
- 18 35. Winsor GL, Lam DK, Fleming L, Lo R, Whiteside MD, et al. (2011) *Pseudomonas* Genome
19 Database: improved comparative analysis and population genomics capability for *Pseudomonas*
20 genomes. *Nucleic Acids Res* 39: D596-D600.
- 21 36. Berg DE, Weiss A, Crossland L (1980) Polarity of Tn5 insertion mutations in *Escherichia coli*. *J*
22 *Bacteriol* 142: 439-446.
- 23 37. Liberati NT, Urbach JM, Miyata S, Lee DG, Drenkard E, et al. (2006) An ordered, nonredundant
24 library of *Pseudomonas aeruginosa* strain PA14 transposon insertion mutants. *Proc Natl Acad Sci*
25 *USA* 103: 2833-2838.
- 26 38. Cherepanov PP, Wackernagel W (1995) Gene disruption in *Escherichia coli*: Tc^R and Km^R
27 cassettes with the option of Flp-catalyzed excision of the antibiotic-resistance determinant. *Gene*
28 158: 9-14.
- 29 39. Hoang TT, Karkhoff-Schweizer RR, Kutchma AJ, Schweizer HP (1998) A broad-host-range Flp-
30 FRT recombination system for site-specific excision of chromosomally-located DNA sequences:
31 application for isolation of unmarked *Pseudomonas aeruginosa* mutants. *Gene* 212: 77-86.

- 1 40. Bachmann BJ (1996) Derivations and genotypes of some mutant derivatives of *Escherichia coli* K-
2 12. In: Neidhardt FC, Curtiss III R, Ingraham JL, Lin ECC, Low Jr KB, et al., editors. *EcoSal-
3 Escherichia coli and Salmonella: cellular and molecular biology*, Washington, D.C.: American
4 Society for Microbiology Press. pp. 2460-2488.
- 5 41. de las Heras A, Carreño CA, de Lorenzo V (2008) Stable implantation of orthogonal sensor circuits
6 in Gram-negative bacteria for environmental release. *Environ Microbiol* 10: 3305-3316.
- 7 42. Herrero M, de Lorenzo V, Timmis KN (1990) Transposon vectors containing non-antibiotic
8 resistance selection markers for cloning and stable chromosomal insertion of foreign genes in
9 Gram-negative bacteria. *J Bacteriol* 172: 6557-6567.
- 10 43. Miller VL, Mekalanos JJ (1988) A novel suicide vector and its use in construction of insertion
11 mutations: osmoregulation of outer membrane proteins and virulence determinants in *Vibrio
12 cholerae* requires *toxR*. *J Bacteriol* 170: 2575-2583.
- 13 44. de Lorenzo V, Cases I, Herrero M, Timmis KN (1993) Early and late responses of TOL promoters
14 to pathway inducers: identification of postexponential promoters in *Pseudomonas putida* with *lacZ-
15 tet* bicistronic reporters. *J Bacteriol* 175: 6902-6907.
- 16 45. Ferrières L, Hémary G, Nham T, Guérout AM, Mazel D, et al. (2010) Silent mischief: bacteriophage
17 Mu insertions contaminate products of *Escherichia coli* random mutagenesis performed using
18 suicidal transposon delivery plasmids mobilized by broad-host-range RP4 conjugative machinery.
19 *J Bacteriol* 192: 6418-6427.
- 20 46. Boyer HW, Roulland-Dussoix D (1969) A complementation analysis of the restriction and
21 modification of DNA in *Escherichia coli*. *J Mol Biol* 41: 459-472.
- 22 47. Worsey MJ, Williams PA (1975) Metabolism of toluene and xylenes by *Pseudomonas putida
23 (arvilla)* mt-2: evidence for a new function of the TOL plasmid. *J Bacteriol* 124: 7-13.
- 24 48. Bagdasarian M, Lurz R, Rückert B, Franklin FC, Bagdasarian MM, et al. (1981) Specific-purpose
25 plasmid cloning vectors. II. Broad host range, high copy number, RSF1010-derived vectors, and a
26 host-vector system for gene cloning in *Pseudomonas*. *Gene* 16: 237-247.
- 27 49. Ditta G, Stanfield S, Corbin D, Helinski DR (1980) Broad host range DNA cloning system for Gram-
28 negative bacteria: construction of a gene bank of *Rhizobium meliloti*. *Proc Natl Acad Sci USA* 77:
29 7347-7351.

- 1 50. Kessler B, de Lorenzo V, Timmis KN (1992) A general system to integrate *lacZ* fusions into the
- 2 chromosomes of Gram-negative eubacteria: regulation of the *P_m* promoter of the TOL plasmid
- 3 studied with all controlling elements in monocopy. *Mol Gen Genet* 233: 293-301.
- 4

1 **Table 1.** Bacterial strains used in this protocol.

2

Bacterial strain	Description and genotype	Relevant characteristics ^a	Reference
<i>Escherichia coli</i>			
CC118 λ <i>pir</i>	Cloning host for plasmids containing an R6K origin of replication; $\Delta(ara-leu)$ <i>araD</i> Δ <i>lacX174 galE galK phoA20 thi-1 rpsE rpoB argE</i> -(Am) <i>recA1</i> , λ <i>pir</i> lysogen	Sp ^R , Rif ^R , Thi ⁻ , Leu ⁻	(42)
SM10 λ <i>pir</i>	Cloning and mobilizing host for plasmids containing an R6K origin of replication; F- <i>thi-1 thr leu tonA lacY glnV recA::RP4-2-Tc::Mu</i> , λ <i>pir</i> lysogen	Km ^R , Thi ⁻ , Thr ⁻ , Leu ⁻	(43)
S17-1 λ <i>pir</i>	Cloning and mobilizing host for plasmids containing an R6K origin of replication; F- <i>recA1 endA1 thiE1 pro-82 creC510 hsdR17 RP4-2-Tc::Mu-Km::Tn7</i> , λ <i>pir</i> lysogen	Sm ^R /Sp ^R , Tp ^R , Thi ⁻ , Pro ⁻	(44)
MFD λ <i>pir</i>	Cloning and mobilizing Mu-free host for plasmids containing an R6K origin of replication; F- λ - <i>ilvG rfb-50 rph-1 RP4-2-Tc::[ΔMu1::aac(3)IV ΔaphA Δnic35 ΔMu2::zeo] ΔdapA::(erm-<i>pir</i>) ΔrecA</i>	Apra ^R , Zeo ^R , Erm ^R , DAP ⁻	(45)
HB101	Mating helper strain; F- λ - <i>hsdS20(r_B⁻ m_B⁻) recA13 leuB6(Am) araC14 Δ(gpt-<i>proA</i>)62 lacY1 galK2(Oc) xyl-5 mtl-1 thiE1 rpsL20 glnX44(AS)</i>	Sm ^R , Thi ⁻ , Leu ⁻ , Pro ⁻	(46)
<i>Pseudomonas putida</i>			
KT2440	Wild-type strain; derivative of strain mt-2 (47) cured of the TOL plasmid pWW0	Prototroph	(48)

3

4 ^a Antibiotic and auxotrophy markers: Apra, apramycin; Erm, erythromycin; Km, kanamycin; Rif,

1 rifampicin; Sp, spectinomycin; Sm, streptomycin; Tp, trimethopim; Zeo: zeocin; DAP,
2 diaminopimelic acid; Leu, leucine; Thi, thiamine (vitamin B1); Thr, threonine; and Pro, proline.
3 Please note that not all these features are used in the experiments described in the present
4 protocol.

1 **Table 2.** Plasmids used in this protocol.

2

Plasmid	Description and relevant characteristics ^a	Reference
pRK600	Helper plasmid used for conjugation; <i>oriV</i> (ColE1), RK2(<i>mob</i> ⁺ <i>tra</i> ⁺); derivative of plasmid pRK2013 (49); Cm ^R	(50)
pBAMD1-2	Mini-Tn5 delivery plasmid; <i>oriV</i> (R6K), <i>oriT</i> ; Ap ^R , Km ^R	(9)
pBAMD1-4	Mini-Tn5 delivery plasmid; <i>oriV</i> (R6K), <i>oriT</i> ; Ap ^R , Sm ^R /Sp ^R	(9)
pBAMD1-6	Mini-Tn5 delivery plasmid; <i>oriV</i> (R6K), <i>oriT</i> ; Ap ^R , Gm ^R	(9)
pBAM1	Mini-Tn5 delivery plasmid; <i>oriV</i> (R6K), <i>oriT</i> ; Ap ^R , Km ^R	(7)
pBAM1-GFP	Mini-Tn5 delivery plasmid to create random <i>gene::gfp</i> fusions by insertion; <i>oriV</i> (R6K), <i>oriT</i> ; Ap ^R , Km ^R	(7)
pBELK	Mini-Tn5 delivery plasmid for inserting a DNA cargo under the control of the LacI ^Q / <i>P</i> _{trc} expression system; <i>oriV</i> (R6K), <i>oriT</i> ; Ap ^R , Km ^R	(10)
pBELG	Mini-Tn5 delivery plasmid for inserting a DNA cargo under the control of the LacI ^Q / <i>P</i> _{trc} expression system; <i>oriV</i> (R6K), <i>oriT</i> ; Ap ^R , Gm ^R	(10)
pBEXK	Mini-Tn5 delivery plasmid for inserting a DNA cargo under the control of the XylS/ <i>P</i> _m expression system; <i>oriV</i> (R6K), <i>oriT</i> ; Ap ^R , Km ^R	(10)
pBEXG	Mini-Tn5 delivery plasmid for inserting a DNA cargo under the control of the XylS/ <i>P</i> _m expression system; <i>oriV</i> (R6K), <i>oriT</i> ; Ap ^R , Gm ^R	(10)
pFLP2	Helper plasmid used to eliminate antibiotic markers flanked by <i>FRT</i> sequences; <i>oriV</i> (pRO1600), RK2(<i>mob</i> ⁺ <i>tra</i> ⁺), <i>oriT</i> , λ <i>P</i> _R :: <i>FLP</i> , λ <i>cl857</i> , <i>sacB</i> ; Cb ^R	(39)

3

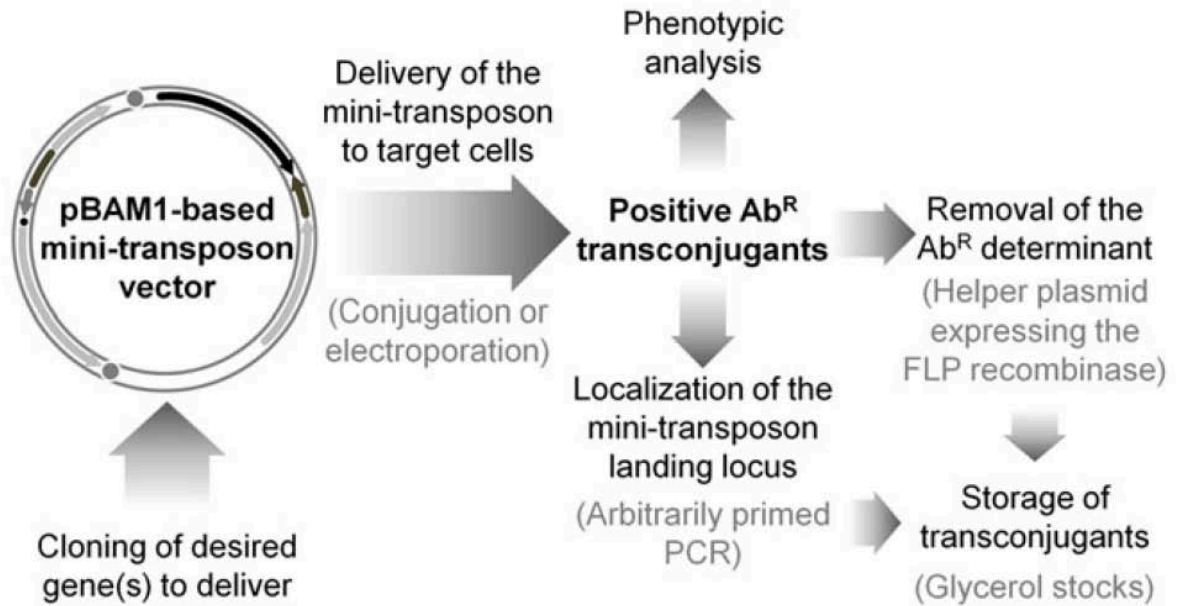
4 ^a Antibiotic markers: Ap, ampicillin; Km, kanamycin; Cm, chloramphenicol; Sm, streptomycin; Sp,
5 spectinomycin; Gm, gentamicin; Cb, carbenicillin.

6

Figure

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FIG. 1. Outline of the procedure described in this protocol.



5
6

7 Mini-transposon vectors can be used for delivering gene(s) into a target chromosome in virtually any
8 Gram-negative bacterium, as well as to obtain random mutant libraries. Ab^R, antibiotic resistance.