

# UltraCAST: A Flexible All-In-One Suicide Vector for Modifying Bacterial Genomes Using a CRISPR-Associated Transposon

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## **Abstract**

CRISPR-associated transposons (CASTs) are RNA-guided mobile genetic elements that are widespread in bacterial genomes. Here, we describe the UltraCAST, a suicide vector with the *Vibrio cholerae* Type I-F CAST system and Golden Gate assembly sites with fluorescent protein gene dropouts for guide RNA and a mini-transposon cargo cloning. We show an example of UltraCAST genome editing by disrupting a gene in the chromosome of *Serratia symbiotica* CWBI-2.3<sup>T</sup>, a culturable relative of aphid endosymbionts. The UltraCAST can be used to flexibly insert DNA into specific genomic sites and facilitates testing this genome editing platform in non-model bacterial species that lack genetic tools.

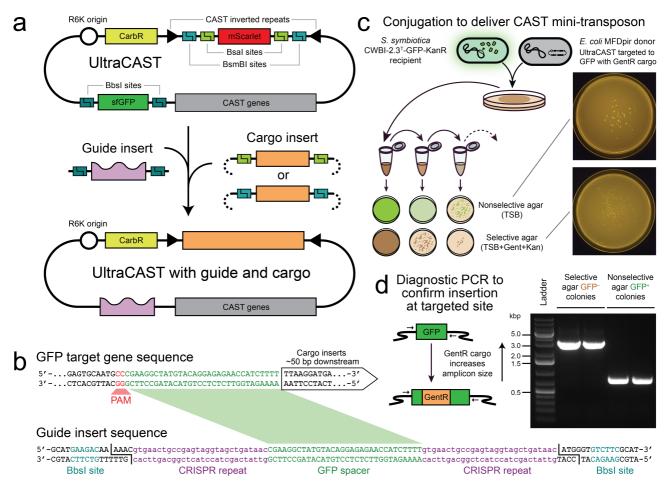


Figure 1. UltraCAST editing of the Serratia symbiotica genome:

(a) UltraCAST plasmid map and steps for cloning guide and cargo inserts. In the example, we replaced the sfGFP dropout with a guide sequence targeting GFP using BbsI Golden Gate assembly and replaced the mScarlet dropout with a gentamicin resistance cassette (GentR) using BsaI Golden Gate assembly. (b) Guide insert sequence design. Oligos are annealed to create the double-stranded piece of DNA encoding the guide RNA flanked with BbsI restriction sites. The sequence shown was designed for targeting the GFP gene integrated into the chromosome of *S. symbiotica* CWBI-2.3<sup>T</sup>-GFP-KanR. (c) Delivery of the UltraCAST suicide plasmid from a donor *E. coli* strain to the recipient bacterium through conjugation followed by selection for insertion of the cargo cassette. In the example, loss of GFP expression from the *S. symbiotica* recipient in most cells on the selective plate indicates a high frequency of the on-target insertion event. Plates were visualized using a blue light transilluminator. (d) Diagnostic PCR to verify insertion at the targeted site. In the example, PCR amplicons with primers flanking the targeted GFP gene in the *S. symbiotica* chromosome show the

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expected increase in size after insertion in two different colony picks from the selective plate relative to the size in colonies with no insertion from a nonselective plate.

## **Description**

Metagenomic studies have discovered diverse bacterial communities in the environment and plant and animal microbiomes, but most of these bacteria have never been experimentally characterized. Even when they can be cultured, developing reliable genetic tools for non-model bacteria can be a daunting and time-consuming process of trial and error (Elston et al. 2023; Gilbert et al. 2023). Transposons are mobile genetic elements that are often among the first tools available for mutagenesis and genome editing of new microbial species. Most transposons either insert nonspecifically at many sites throughout a genome (e.g., Tn5) or insert into one conserved site (e.g., Tn7) (de Lorenzo et al. 1990; Craig 1991; Reznikoff 1993). To insert a DNA cargo or edit a specific site in a bacterial genome, homology-based recombination methods are typically used (Sharan et al. 2009; Lariviere et al. 2024), but these may require expressing phage recombinases or working out DNA delivery methods.

CRISPR-associated transposons (CASTs) are RNA-guided mobile elements related to Tn7 that are widespread in bacteria (Peters et al. 2017; Rybarski et al. 2021; Klompe et al. 2022; Park et al. 2022; Faure et al. 2023). As with other transposons, they can be engineered into mini-transposons in which a cargo, located within the terminal inverted repeats of the transposon, can be mobilized by expressing transposon genes in *trans* (Strecker et al. 2019; Klompe et al. 2019; Vo et al. 2021; Klompe et al. 2022; Gelsinger et al. 2024). Refactored VcCASTs based on the type I-F system from *Vibrio cholerae* HE-45 have been shown to function in *Escherichia coli* and *Klebsiella, Pseudomonas*, and *Ralstonia* species (Klompe et al. 2019; Vo et al. 2021; Rubin et al. 2022; Gelsinger et al. 2024). VcCAST mini-transposons have >95% specificity for inserting their cargo at a site ~50 bp downstream of a match to their guide RNA sequence with a protospacer-adjacent motif (PAM) (Klompe et al. 2019; Vo et al. 2021). They may insert their cargo in either a forward or reverse orientation, with a higher preference for one orientation with certain guide RNAs and shorter cargos (Klompe et al. 2019). VcCAST mini-transposons have been used to insert >10-kb DNA cargos (Klompe et al. 2019), and they can likely mobilize even larger cargos, as the natural sizes of some type I-F CASTs surpass 80 kb (Klompe et al. 2022).

Here we present the UltraCAST, a VcCAST plasmid that improves on a previous design (Hu et al. 2024). The UltraCAST is a suicide vector with an R6K origin of replication that includes two Golden Gate compatible fluorescent protein dropout sequences for cloning in DNA inserts encoding a guide and cargo (**Fig. 1a**). A double-stranded DNA fragment encoding the guide RNA is cloned in using BbsI in place of the sfGFP dropout (**Fig. 1b**). The mScarlet dropout site can accept a cargo sequence containing either BsmBI or BsaI cut sites that generate overhangs compatible with either Stage 1 or Stage 2 assembly according to existing synthetic biology standards (Lee et al. 2015; Leonard et al. 2018). Guide and cargo cloning can be done in either order, facilitating use cases where a researcher wants to target different sites and/or species with the same cargo or when different cargos are to be inserted at the same site in one genome.

The assembled UltraCAST is delivered to the strain of interest through conjugation from a pir<sup>+</sup> *E. coli* donor strain. Since the R6K plasmid cannot replicate in recipient cells that are *pir*<sup>-</sup> or otherwise incompatible with this origin of replication, CAST-mediated insertions of the mini-transposon cargo are selected in a single step. Unlike VcCAST configurations that rely on one or more replicative plasmids (Klompe et al. 2019; Vo et al. 2021; Gelsinger et al. 2024), this means that editing can take place in non-model bacterial strains for which no compatible plasmid origins are known and that there is no need to cure cells of the plasmid after editing is complete.

We initially verified that the refactored UltraCAST system functioned by using it to insert a chloramphenicol resistance cassette into the *lacZ* gene of *E. coli* REL606, the ancestor of a >35-year long-term evolution experiment (Lenski et al. 1991, Barrick et al. 2023). We have also used the UltraCAST to insert a gentamicin resistance cassette into various genes in *Serratia symbiotica* CWBI-2.3<sup>T</sup>, a culturable gut symbiont of aphids that is related to strains that have evolved into vertically inherited intracellular endosymbionts (Sabri et al. 2011; Renoz et al. 2021). To date, we have used the UltraCAST to disrupt *aroG*, *esaR*, *esaI*, *leuB*, *sctN*, *cyaA*, *ridA*, and *pheA* in this species to study how these gene knockouts affect bacterial growth and aphid colonization.

This protocol describes how to use the UltraCAST to insert a DNA cargo at a specific position in the genome of a recipient bacterium. We demonstrate the steps and show expected results for inserting a ~2-kb gentamicin resistance cassette into the chromosome of *S. symbiotica* CWBI-2.3<sup>T</sup>-GFP-KanR (Perreau et al. 2021). The on-target efficiency of insertion of the CAST mini-transposon into GFP in *S. symbiotica* was ~2.4% (**Fig. 1c**). A diagnostic PCR verified that the gentamicin resistance cassette was successfully integrated into the sfGFP gene (**Fig. 1d**). This protocol can be modified for other targets and cargos. This blueprint and the flexible UltraCAST plasmid should facilitate using this system to genetically engineer other non-model bacteria.

## Methods



- 1. Design the DNA sequence encoding the CAST guide RNA by scanning the region being targeted for a 5'-CC-3' protospacer adjacent motif (PAM). Use the 32 bases following the PAM for the guide sequence. The CAST will insert  $\sim$ 50 bp downstream of the 3' end of where the guide matches the genome. For the GFP example, we used 5'-CGAAGGCTATGTACAGGAGAGAACCATCTTTT-3'.
- 2. Append sequences that will be transcribed into the CRISPR array repeats (lowercase) and flanks needed for BbsI Golden Gate assembly (uppercase). Add 5'-GCATGAAGACAAAAACgtgaactgccgagtaggtaggtgataac-3' before your guide sequence and 5'-gtgaactgccgagtaggtaggtgataacATGGGTGTCTTCGCAT-3' after it. Order the full 120-base sequence and its reverse complement for DNA synthesis.
- 3. Resuspend oligos in nuclease-free water to  $100~\mu M$ . Dilute and combine them in duplex buffer. We used a  $10~\mu M$  final concentration of each oligo. Anneal the oligos to create a double-stranded DNA fragment by running this program in a thermocycler:  $95^{\circ}C$  (2 min), ramp to  $12^{\circ}C$  by  $1^{\circ}C$  per 30 s,  $12^{\circ}C$  (hold).
- 4. Clone the guide into the UltraCAST vector using BbsI Golden Gate assembly. We used 2.5 nM vector and 25 nM guide insert in a 20  $\mu$ L reaction with 20 U NEB BbsI-HF enzyme and 800 U T4 DNA ligase in 1× rCutSmart buffer supplemented with 0.1 mM of ATP and ran this thermocycler program: 30 cycles alternating between 37°C (1 min) and 16°C (1 min), then final 60°C (5 min) and 12°C (hold) steps.
- 5. Transform the Golden Gate reaction into E. coli pir $^+$  cells and plate on LB agar with 100  $\mu$ g/mL carbenicillin (LB-Carb agar) for selection. Incubate overnight at 37 $^{\circ}$ C.
- 6. Pick one or more GFP negative colonies by inspecting plates on a blue light transilluminator. Grow liquid cultures in LB-Carb overnight at 37°C with shaking, miniprep plasmids, and verify that one has the intended guide by sequencing.

**Troubleshooting**: If there are no GFP negative colonies after transformation, try increasing the time at both 37°C and 16°C for each Golden Gate assembly cycle to 5 min and the number of cycles to 45.

#### Clone a cargo into the UltraCAST

- 1. The UltraCAST has a cargo site that accepts either a Type 2-4 YTK/BTK construct via BsaI Golden Gate assembly or a ConLS-ConRE construct via BsmBI Golden Gate assembly (Lee et al. 2015; Leonard et al. 2018). Miniprep part plasmids or create PCR products with compatible restriction sites and overhangs for cloning into this site.
- 2. Clone cargo into the UltraCAST using BsaI or BsmBI Golden Gate assembly. Use the same thermocycler conditions as for BbsI assembly to add the guide (see above). For the example, we performed BsaI assembly with 2.5 nM of UltraCAST vector and 25 nM of a Type 2-4 gentamicin resistance (GentR) cassette part plasmid in a 20  $\mu$ L NEBridge (BsaI-HF v2) kit reaction.
- 3. Transform the Golden Gate reaction into E. coli pir $^+$  cells and plate on LB-Carb agar supplemented with additional antibiotic for a resistance cassette in the cargo, if applicable. We used 20  $\mu$ g/mL gentamicin. Incubate overnight at 37°C.
- 4. Pick one or more mScarlet negative colonies by inspecting plates on a blue light transilluminator. Grow liquid cultures in LB-Carb overnight at 37°C with shaking, miniprep plasmids, and verify that one has the intended cargo by sequencing.

**Alternatives**: We created an equivalent Type 2-4 kanamycin resistance (KanR) cassette that can be used for organisms and cases where selection with this antibiotic is preferred. The guide and cargo inserts can be cloned in either order to facilitate making a family of plasmids with the same cargo and different guides or vice versa.

#### Prepare the *E. coli* conjugation donor strain

- 1. Transform the final UltraCAST plasmid with a guide and cargo into a pir<sup>+</sup> *E. coli* conjugation donor strain. Plate on LB-Carb plates supplemented with any additional nutrients needed for auxotrophic strains to grow. Incubate overnight at 37°C. We used *E. coli* MFDpir electrocompetent cells and cultured them with 0.3 mM 2,6-diaminopimelic acid (DAP).
- 2. Pick colonies and grow in at least 5 mL of LB with the same supplements overnight with shaking at 37°C.

**Pause Point:** These cultures can be frozen at –80°C with 15% w/w glycerol and later revived with overnight growth for conjugation.

**Alternative**: *E. coli* ST18 can be used as the pir<sup>+</sup> conjugation donor strain. In this case, supplement media with 50  $\mu$ g/mL 5-aminolevulinic acid (ALA) instead of DAP.

## Conjugate into the recipient strain

1. Grow a culture of the recipient strain. Time its growth so that a saturated culture or agar plate that can be scraped is available at the same time as a freshly grown overnight culture of the  $E.\ coli$  donor strain. We used  $Serratia\ symbiotica$  CWBI-2.3<sup>T</sup>-GFP-KanR revived from a glycerol stock and grown in 5 mL TSB with shaking at 25°C for ~48 h.



- 2. Spin down donor and recipient cultures at  $6800 \times g$  for 3 min. Wash each tube of cells three successive times with  $1 \times PBS$  by removing supernatant and spinning down at  $6800 \times g$  for 1 min. Resuspend the washed pellets in  $1 \times PBS$ , diluting as necessary to reach an OD600 value of  $\sim$ 1 for each strain.
- 3. Mix together a total volume of 1 mL of washed donor and recipient cells. The ratio between the two strains should be adjusted so that it includes more of the slower-growing strain (usually the recipient). Spin down at  $10,000 \times g$  for 1 min. Resuspend the combined pellet in  $100 \mu L$  of  $1 \times PBS$ . Plate the entire solution in a single drop on an agar plate on which both strains can grow. It should be supplemented with any nutrient the *E. coli* donor needs to grow and should not contain the antibiotic that will be used later to select for successful insertion of the cargo into the genome of the recipient. Once dry, incubate the plate under conditions where both strains can grow for long enough that there is some growth of the recipient. For the example, we used a 1:5 ratio of *E. coli* to *S. symbiotica*, plated on TSB-DAP, and incubated at 25°C overnight.
- 4. Scrape cells and resuspend in 1 mL of  $1 \times PBS$ . Make a 10-fold dilution series in  $1 \times PBS$  out to a  $10^7$  dilution. Plate on selective and non-selective agar that lacks the nutrient needed by the auxotrophic donor. The selective agar should include the antibiotic for the resistance gene encoded in the CAST cargo. Incubate under growth conditions for the recipient strain. We used TSB-Gent-Kan agar and incubated at  $25^{\circ}$ C for 96 h. The gentamicin was to select for cargo insertion. The kanamycin was added because our *S. symbiotica* recipient strain already had this resistance marker in its genome.
- 5. Use diagnostic PCR reactions and/or genome resequencing to confirm insertion of the transposon cassette at the targeted site. In the example, successful knockouts were verified based on an increase in the size of a PCR amplicon spanning the insertion site and loss of GFP expression due to the mini-transposon cargo inserting into and inactivating this gene.

Reagents

Supplier/Catalog # or Recipe
Sigma Aldrich (D8537)
Avantar TransforMax EC100D pir+ (75927-934)
Ferrières et al. 2010
German Collection of Microorganisms and Cell Cultures (DSM 22074)
GoldBio (G-400-5)
GoldBio (K-120-5)
GoldBio (C-103-5)
Sigma Aldrich (D1377-5G)
Sigma Aldrich (A7793-1G)
Addgene (236111)
Addgene (236187)
Addgene (236186)
Integrated DNA Technologies (Ultramer)
100 mM Potassium Acetate, 30 mM HEPES, pH 7.5

Fisher Bioreagents (10153193)
Tisher Dioreagents (10133133)
For 1L: 10 g Tryptone, 5 g Yeast Extract, 10 g Sodium Chloride
For 1L: 10 g Tryptone, 5 g Yeast Extract, 10 g Sodium Chloride, 15 g Agar
For 1L: 30 g of BD Bacto Tryptic Soy Broth (DF0370-17-3)
For 1L: 30 g of BD Bacto Tryptic Soy Broth (DF0370-17-3), 15 g Agar
Perreau et al. 2021
ZymoPure II Plasmid Midiprep Kit (D4200)
New England Biolabs (R3539S)
New England Biolabs (E1601L)
New England Biolabs (E1602L)
New England Biolabs (M0202L)
New England Biolabs (B6004S)
New England Biolabs (P0756S)

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## References

Barrick JE, Blount ZD, Lake DM, Dwenger JH, Chavarria-Palma JE, Izutsu M, Wiser MJ. 2023. Daily transfers, archiving populations, and measuring fitness in the long-term evolution experiment with *Escherichia coli*. Journal of Visualized Experiments 198: e65342. DOI: 10.3791/65342

Craig NL. 1991. Tn7: a target site-specific transposon. Molecular Microbiology 5: 2569-2573. DOI:  $\frac{10.1111}{j.1365-2958.1991.tb01964.x}$ 

de Lorenzo V, Herrero M, Jakubzik U, Timmis KN. 1990. Mini-Tn5 transposon derivatives for insertion mutagenesis, promoter probing, and chromosomal insertion of cloned DNA in gram-negative eubacteria. Journal of Bacteriology 172: 6568-6572. DOI: 10.1128/jb.172.11.6568-6572.1990

Elston KM, Phillips LE, Leonard SP, Young E, Holley JaC, Ahsanullah T, et al., Barrick JE. 2023. The Pathfinder plasmid toolkit for genetically engineering newly isolated bacteria enables the study of *Drosophila*-colonizing *Orbaceae*. ISME Communications 3: 49. DOI: 10.1038/s43705-023-00255-3

Faure G, Saito M, Benler S, Peng I, Wolf YI, Strecker J, et al., Zhang F. 2023. Modularity and diversity of target selectors in Tn7 transposons. Molecular Cell 83: 2122-2136.e10. DOI: <u>10.1016/j.molcel.2023.05.013</u>

Ferrières L, Hémery Gl, Nham T, Guérout AM, Mazel D, Beloin C, Ghigo JM. 2010. Silent mischief: Bacteriophage Mu insertions contaminate products of *Escherichia coli* random mutagenesis performed using suicidal transposon delivery plasmids mobilized by broad-host-range RP4 conjugative machinery. Journal of Bacteriology 192: 6418-6427. DOI: 10.1128/JB.00621-10



Gelsinger DR, Vo PLH, Klompe SE, Ronda C, Wang HH, Sternberg SH. 2024. Bacterial genome engineering using CRISPR-associated transposases. Nature Protocols 19: 752-790. DOI: <u>10.1038/s41596-023-00927-3</u>

Gilbert C, Brumwell SL, Crits-Christoph A, Kang SC, Martin-Moldes Z, Alsharif W, et al., Ostrov N. 2023. A scalable framework for high-throughput identification of functional origins of replication in non-model bacteria. bioRxiv DOI: 10.1101/2023.05.19.541510

Hu K, Chou CW, Wilke CO, Finkelstein IJ. 2024. Distinct horizontal transfer mechanisms for type I and type V CRISPR-associated transposons. Nature Communications 15: 6653. DOI: <u>10.1038/s41467-024-50816-w</u>

Klompe SE, Vo PLH, Halpin-Healy TS, Sternberg SH. 2019. Transposon-encoded CRISPR–Cas systems direct RNA-guided DNA integration. Nature 571: 219-225. DOI: <u>10.1038/s41586-019-1323-z</u>

Klompe SE, Jaber N, Beh LY, Mohabir JT, Bernheim A, Sternberg SH. 2022. Evolutionary and mechanistic diversity of Type I-F CRISPR-associated transposons. Molecular Cell 82: 616-628.e5. DOI: <u>10.1016/j.molcel.2021.12.021</u>

Lariviere PJ, Ashraf AHMZ, Navarro-Escalante L, Leonard SP, Miller LG, Moran NA, Barrick JE. 2024. One-step genome engineering in bee gut bacterial symbionts. mBio 15: e0139224. DOI: <u>10.1128/mbio.01392-24</u>

Lee ME, DeLoache WC, Cervantes B, Dueber JE. 2015. A highly characterized yeast toolkit for modular, multipart assembly. ACS Synthetic Biology 4: 975-986. DOI: 10.1021/sb500366v

Lenski RE, Rose MR, Simpson SC, Tadler SC. 1991. Long-term experimental evolution in *Escherichia coli*. I. Adaptation and divergence during 2,000 generations. The American Naturalist 138: 1315-1341. DOI: <u>10.1086/285289</u>

Leonard SP, Perutka J, Powell JE, Geng P, Richhart DD, Byrom M, et al., Barrick JE. 2018. Genetic engineering of bee gut microbiome bacteria with a toolkit for modular assembly of broad-host-range plasmids. ACS Synthetic Biology 7: 1279-1290. DOI: <a href="https://doi.org/10.1021/acssynbio.7b00399">10.1021/acssynbio.7b00399</a>

Park JU, Tsai AWL, Chen TH, Peters JE, Kellogg EH. 2022. Mechanistic details of CRISPR-associated transposon recruitment and integration revealed by cryo-EM. Proceedings of the National Academy of Sciences of the United States of America 119: e2202590119. DOI: 10.1073/pnas.2202590119

Peters JE, Makarova KS, Shmakov S, Koonin EV. 2017. Recruitment of CRISPR-Cas systems by Tn7-like transposons. Proceedings of the National Academy of Sciences of the United States of America 114: E7358-E7366. DOI: 10.1073/pnas.1709035114

Perreau J, Patel DJ, Anderson H, Maeda GP, Elston KM, Barrick JE, Moran NA. 2021. Vertical transmission at the pathogen-symbiont interface: *Serratia symbiotica* and aphids. mBio 12: e00359-21. DOI: <u>10.1128/mBio.00359-21</u>

Renoz Fo, Foray V, Ambroise Jrm, Baa-Puyoulet P, Bearzatto B, Mendez GL, et al., Hance T. 2021. At the gate of mutualism: Identification of genomic traits predisposing to insect-bacterial symbiosis in pathogenic strains of the aphid symbiont *Serratia symbiotica*. Frontiers in Cellular and Infection Microbiology 11: 660007. DOI: 10.3389/fcimb.2021.660007

Reznikoff WS. 1993. The Tn5 transposon. Annual Review of Microbiology 47: 945-964. DOI: 10.1146/annurev.mi.47.100193.004501

Rubin BE, Diamond S, Cress BF, Crits-Christoph A, Lou YC, Borges AL, et al., Doudna JA. 2022. Species- and site-specific genome editing in complex bacterial communities. Nature Microbiology 7: 34-47. DOI: <u>10.1038/s41564-021-01014-7</u>

Rybarski JR, Hu K, Hill AM, Wilke CO, Finkelstein IJ. 2021. Metagenomic discovery of CRISPR-associated transposons. Proceedings of the National Academy of Sciences of the United States of America 118: e2112279118. DOI: 10.1073/pnas.2112279118

Sabri A, Leroy P, Haubruge E, Hance T, Frère I, Destain J, Thonart P. 2011. Isolation, pure culture and characterization of *Serratia symbiotica* sp. nov., the R-type of secondary endosymbiont of the black bean aphid *Aphis fabae*. International Journal of Systematic and Evolutionary Microbiology 61: 2081-2088. DOI: 10.1099/ijs.0.024133-0

Sharan SK, Thomason LC, Kuznetsov SG, Court DL. 2009. Recombineering: a homologous recombination-based method of genetic engineering. Nature Protocols 4: 206-223. DOI: <u>10.1038/nprot.2008.227</u>

Strecker J, Ladha A, Gardner Z, Schmid-Burgk JL, Makarova KS, Koonin EV, Zhang F. 2019. RNA-guided DNA insertion with CRISPR-associated transposases. Science 365: 48-53. DOI: <a href="https://doi.org/10.1126/science.aax9181">10.1126/science.aax9181</a>

Vo PLH, Ronda C, Klompe SE, Chen EE, Acree C, Wang HH, Sternberg SH. 2021. CRISPR RNA-guided integrases for high-efficiency, multiplexed bacterial genome engineering. Nature Biotechnology 39: 480-489. DOI: <u>10.1038/s41587-020-00745-v</u>



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