Supplementary File 2: bacterial strains used in this work.

**GENERAL STRAINS**

|  |
| --- |
|  |
| **Strain** | **Genotype** | **Use** | **Reference** |
| DH5α | *supE44 ΔlacU169* (Φ80*lacZ’ ΔM15) ΔargF hsdR17 recA1 endA1 gyrA96 thi-1 relA1* | General | Lab strain |
| UB5120 | F-pro met *recA56* gyrA [NalR] | General | Martinez and de la Cruz. 1990 |
| β2163 | MG1655:: *Δ*dapA::(erm-pir)RP4-2-T c::Mu [KmR] | Donor strain in conjugation | Demarre *et al*. 2005 |
| B36 | MG1655 Δ*dapA* recA269::Tn10 attB::*attI1WT*-*attI1STOP*-*dapA* [SpR] | Enrichment cycles and recombination tests | Escudero *et al*. 2016 |

**CONJUGATION ASSAYS**

|  |
| --- |
| **Donor strains** |
| **Strain** | **Genotype** | **Reference** |
| 4137 | β2163/pSW23T::*attCaadA7* | Bouvier *et al.* 2005 |
| 2714 | β2163/pSW23T::*attI1* | Demarre *et al*. 2005 |
| H203 | β2163/p6944 (pSW23T:: *attCaadA7-*VCR2/1) | Strain from this work.Plasmid p6944 from Bouvier *et al*. 2009 |
| **Receptor strains** |
| **Strain** | **Genotype** | **Parental allele** | **Additional mutations** | **Resident plasmids** |
| 9669 | DH5α |  | Wild type *intI1* | P929 (pSU38Δ::*attI1*) |
| H294 | “ | *intI1* | T118S | “ |
| H291 | “ | “ | D161G | “ |
| H297 | “ | “ | G320D | “ |
| H306 | “ | “ | T118S, D161G | “ |
| H303 | “ | “ | T118S, G320D | “ |
| H300 | “ | “ | D161G, G320D | “ |
| H309 | “ | “ | T118S, D161G, G320D | “ |
| B903 | “ | “ | T118S, D161G, H162Q, K219R, G320D | “ |
| B904 | “ | “ | T118S, D161G, H162Q, K219R, D299E, G320D, | “ |
| B905 | “ | “ | T118S, D161G, H162Q, K219R, D299E, G319E, G320D, A329T | “ |
| A283 | “ |  | Wild type *alt1l.e.* | “ |
| B506 | “ | *alt1l.e.* | D161, G319E | “ |
| B507 | “ | “ | R12G, 3xSynon (Glu62+Leu113+Pro181) | “ |
| B654 | “ | “ | K219R | “ |
| B655 | “ | “ | H162Q | “ |
| B656 | “ | “ | T96I, G320R, R337C | “ |
| B657 | “ | “ | V318G | “ |
| B658 | “ | “ | H48Q, A85V, D97G, M145V, G269E, L316 | “ |
| B965 | “ | “ | T118S, D161G, G320D, H162Q, K219R, D299E, G319E, A329T | “ |
| A286 | “ |  | Wild type *alt2l.e.* | “ |
| B504 | “ | *alt2l.e.* | D299E, A329T, R324G, Synon (G417A) | “ |
| B505 | “ | “ | D299E, A329T | “ |
| B659 | “ | “ | D299E, A329T, V323A | “ |
| **Recipient strains for *attC* x *attC* excision assays with p6944** |
| 3938 | “ |  | Wild type *intI1* | None |
| H315 | “ | *intI1* | T118S | None |
| H312 | “ | “ | D161G | None |
| H318 | “ | “ | G320D | None |
| 7617 | “ | “ | T118S, D161G | None |
| H324 | “ | “ | T118S, G320D | None |
| H321 | “ | “ | D161G, G320D | None |
| 7618 | “ | “ | T118S, D161G, G320D | None |
| B877 | “ | “ | T118S, D161G, H162Q, K219R, G320D | None |
| B901 | “ | “ | T118S, D161G, H162Q, K219R, D299E, G320D, | None |
| B902 | “ | “ | T118S, D161G, H162Q, K219R, D299E, G319E, G320D, A329T | None |

**CHROMOSOMAL ASSAYS**

|  |  |  |  |  |
| --- | --- | --- | --- | --- |
| **Strain** | **Genotype** | **Parental allele** | **Additional mutations** | **Observations** |
| C006 | B36 | *intI1* | Wild type allele |  |
| C007 | “ | *alt1l.e.* | Wild type allele |  |
| C008 | “ | *intI1* | *intI18mut*: contains the same 8 mutations as B905 | Not referred to by strain number but rather as *XXX8mut* |
| C009 | “ | *alt1l.e.* | *alt18mut*: contains the same 8 mutations as B905 |
| C010 | “ | *alt2l.e.* | *alt28mut*: contains the same 8 mutations as B905 |
| C325 | “ | *intI18mut* | *intI18mut* + H133Q, V315M,T329M |  |
| C326 | “ | “ | *intI18mut* + Y220N, A321G, 1Syn (G357A), A+49T (3’UTR) |  |
| C492 | “ | “ | *intI18mut* + H133Q |  |
| C493 | “ | “ | *intI18mut* + V315M |  |
| C494 | “ | “ | *intI18mut* + T329M |  |
| C495 | “ | “ | *intI18mut* + Syn (G357A) |  |
| C496 | “ | “ | *intI18mut* + Y220N |  |
| C497 | “ | “ | *intI18mut* + A321G |  |
| C498 | “ | “ | *intI18mut* + A+49 (3’UTR) |  |
| C499 | “ | “ | *intI18mut* + Y220N, Syn (G357A) |  |
| C500 | “ | “ | *intI18mut* + Y220N, A+49 (3’UTR) |  |
| C118 | “ | *alt18mut* | *alt18mut* + V315A |  |
| C119 | “ | “ | *alt18mut* + L216W, V315A |  |
| C120 | “ | “ | *alt18mut* + P221S, V315A |  |
| C121 | “ | “ | *alt18mut* + D241E, D320N |  |
| C122 | “ | “ | *alt18mut* + E103K, E130K, 2 Syn |  |
| C123 | “ | “ | *alt18mut* + S173R |  |
| C124 | “ | “ | *alt18mut* + D320N |  |
| C125 | “ | “ | *alt18mut* + Q102R + S173R |  |
| D472 | “ | “ | *alt18mut* + E103K, 1 Syn |  |
| D473 | “ | “ | *alt18mut* + E130K, 1 Syn |  |
| C126 | “ | *alt28mut* | *alt28mut* + V315A, A-4T (5’UTR) |  |
| C127 | “ | “ | *alt28mut* + E103A, C-2T (5’UTR) |  |
| C128 | “ | “ | *alt28mut* + D320N |  |

**EPISTASIS PURIFICATION**

|  |  |  |  |  |
| --- | --- | --- | --- | --- |
| **Sequence** | **Strain** | **Genotype** | **Reaction** | **Additional mutations on the basis of the *intI1* allele** |
| *intI18mut* | C008 | B36 | *attI* x *attI* | T118S, D161G, H162Q, K219R, D299E, G319E, G320N, A329T |
| E450 | B413 | *attC* x *attC* |
| E458 | B415 | *attI* x *attC* |
| E526 | B37 | *attI* x *attI**inverted orientation (double clivage)* |
| Seq 3 | E393 | B36 | *attI* x *attI* | E103K, T118S, H162Q, S173R, K219R, Y220N, G319D, G320S, |
| E451 | B413 | *attC* x *attC* |
| E459 | B415 | *attI* x *attC* |
| E527 | B37 | *attI* x *attI**inverted orientation (double cleavage)* |
| Seq 5 | E394 | B36 | *attI* x *attI* | E103K, T118S, H162Q, S173R, K219R, Y220N, G319D, G320N, |
| E452 | B413 | *attC* x *attC* |
| E460 | B415 | *attI* x *attC* |
| E528 | B37 | *attI* x *attI**inverted orientation (double cleavage)* |
| Seq 7 | E395 | B36 | *attI* x *attI* | E103K, T118S, H162Q, S173R, K219R, G319E, G320N, A329T |
| E453 | B413 | *attC* x *attC* |
| E461 | B415 | *attI* x *attC* |
| E529 | B37 | *attI* x *attI**inverted orientation (double cleavage)* |
| Seq 19 | E396 | B36 | *attI* x *attI* | E103K, T118S, H162Q, S173R, K219R, Y220N, G319D, G320N, A329T |
| E530 | B37 | *attI* x *attI**inverted orientation (double cleavage)* |
| Seq 23 | E397 | “ | *attI* x *attI* | E103K, T118S, D161G, H162Q, S173R, K219R, Y220N, G319D, G320S |
| E531 | B37 | *attI* x *attI**inverted orientation (double cleavage)* |
| Seq 31 | E398 | “ | *attI* x *attI* | E103K, T118S, H162Q, S173R, M179I, K219R, Y220N, V315A, G319D, G320N, A329T |
| E532 | B37 | *attI* x *attI**inverted orientation (double cleavage)* |