

Table S1. *R. sphaeroides* and *E. coli* strains used in this study

| Strains | Relevant features | References |
|---|--|-------------------------------|
| <i>R. sphaeroides</i> | | |
| 2.4.1 | <i>Rhodobacter sphaeroides</i> wild type | (van Niel, 1944) |
| 2.4.1Δ <i>iscR</i> | Sp ^r , <i>iscR</i> deletion strain | This study |
| 2.4.1Δ <i>iscR</i> _pBBR <i>iscR</i> | 2.4.1Δ <i>iscR</i> harbouring pBBR <i>iscR</i> | This study |
| 2.4.1-pBE::P _{iscR} ::eCFP | Km ^r , 2.4.1 containing pBE::P _{iscR} ::eCFP, used for <i>in vivo</i> reporter system | This study |
| 2.4.1Δ <i>iscR</i> -pBE::P _{iscR} ::eCFP | 2.4.1Δ <i>iscR</i> containing pBBR <i>iscR</i> , Km ^r , used for <i>in vivo</i> reporter system | This study |
| <i>E. coli</i> | | |
| JM109 | Host strain for cloning procedures | (Yanisch-Perron et al., 1985) |
| S17-1 | Strain for diparental conjugation, tra ⁺ | (Simon et al., 1986) |
| M15 | Strain for recombinant protein expression | Qiagen |
| M15-pQE30:: <i>iscR</i> | M15 containing pQE30:: <i>iscR</i> , Ap ^r , used for IscR overexpression | This study |
| M15-pQE30:: <i>iscR</i> _H93A | M15 containing pQE30:: <i>iscR</i> _H93A, Ap ^r , used for IscR overexpression | This study |
| M15-pQE30:: <i>iscR</i> _H121A | M15 containing pQE30:: <i>iscR</i> _H121A, Ap ^r , used for IscR overexpression | This study |
| M15-pQE30:: <i>iscR</i> _H127A | M15 containing pQE30:: <i>iscR</i> _H127A, Ap ^r , used for IscR overexpression | This study |
| M15-pQE30:: <i>iscR</i> _C142A | M15 containing pQE30:: <i>iscR</i> _C142A, Ap ^r , used for IscR overexpression | This study |
| M15-pQE30:: <i>iscR</i> _P143A | M15 containing pQE30:: <i>iscR</i> _P143A, Ap ^r , used for IscR overexpression | This study |
| M15-pQE30:: <i>iscR</i> _H93A/H121A/H127A | M15 containing pQE30:: <i>iscR</i> _H93A/H121A/H127A, Ap ^r , used for IscR overexpression | This study |

Sp^r, spectinomycin-resistant; Ap^r, ampicillin-resistant; Km^r, kanamycin resistant; when required, antibiotics were used in the following concentrations: spectinomycin (10 µg·ml⁻¹) and kanamycin (25 µg·ml⁻¹) for *R. sphaeroides*; ampicillin (200 µg·ml⁻¹) for *E. coli*

Table S2. Plasmids used in this study

| Plasmid names | Relevant features | Source |
|---------------------------------------|---|------------------------|
| pBBR1-MCS-2 | Km ^r , broad-host-range cloning vector | (Kovach et al., 1995) |
| pBBR <i>iscR</i> | Km ^r , pBBR1-MCS-2 containing <i>iscR</i> gene | This study |
| pET28(a) | Km ^r , high expression vector | Novagen |
| pBE | Km ^r , Derivate of pBBR1-MCS-2 | This study |
| pBE4352 | Km ^r , pBE containing RSP_4352 promoter | This study |
| pBE4352::eCFP:eCFP | Km ^r , pBE4352 containing two eCFP | This study |
| pBE::P _{iscR} ::eCFP | Km ^r , pBBR containing <i>iscR</i> fragment for <i>cfp</i> fusion | This study |
| pHP45Ω | Sp ^r , source of Ω-Sp ^r cassette | (Prentki et al., 1991) |
| pPHU281 | Tc ^r , lacZ' mob(RP4) | (Hubner et al., 1991) |
| pPHUΔ2.4.1 <i>iscR</i> | Tc ^r , pPHU281 containing <i>iscR</i> gene with flanking sites | This study |
| pPHUΔ2.4.1 <i>iscR</i> ::ΩSp | Tc ^r , Sp ^r , pPHU281Δ2.4.1 <i>iscR</i> containing spectinomycin cassette | This study |
| pJET1.2 | Ap ^r , 2.97 kb, PCR cloning vector | Fermentas |
| pQE30 | Ap ^r , 3.4 kb, 6xHis-tag overexpression vector | Qiagen |
| pQE30:: <i>iscR</i> | Ap ^r , pQE30 containing <i>iscR</i> fragment for overexpression | This study |
| pQE30:: <i>iscR</i> _H93A | Ap ^r , pQE30:: <i>iscR</i> containing a mutation of amino acid 93 | This study |
| pQE30:: <i>iscR</i> _H121A | Ap ^r , pQE30:: <i>iscR</i> containing a mutation of amino acid 121 | This study |
| pQE30:: <i>iscR</i> _H127A | Ap ^r , pQE30:: <i>iscR</i> containing a mutation of amino acid 127 | This study |
| pQE30:: <i>iscR</i> _H93A/H121A/H127A | Ap ^r , pQE30:: <i>iscR</i> containing mutations of amino acids 93, 121 and 127 | This study |
| pQE30:: <i>iscR</i> _C142A | Ap ^r , pQE30:: <i>iscR</i> containing a mutation of amino acid 142 | This study |
| pQE30:: <i>iscR</i> _P143A | Ap ^r , pQE30:: <i>iscR</i> containing a mutation of amino acid 143 | This study |

Sp^r, spectinomycin-resistant; Ap^r, ampicillin-resistant; Tc^r, tetracycline-resistant; Km^r, kanamycin resistant

Table S3. Oligonucleotides used in this study

| Name | Sequence | Purpose |
|---------------|------------------------------|--|
| RT_RSP_0040_A | TCGAACGACACCAACACC | Forward primer for RSP_0040 (<i>fliS</i>) real-time RT-PCR |
| RT_RSP_0040_B | CACCTTCATGCCGTTGAA | Reverse primer for RSP_0040 (<i>fliS</i>) real-time RT-PCR |
| RT_RSP_0257_A | ACACCTACGGCAACTTCC | Forward primer for RSP_0257 (<i>pufL</i>) real-time RT-pCR |
| RT_RSP_0257_B | ATCGAGTAGCCGACCAGA | Reverse primer for RSP_0257 (<i>pufL</i>) real-time RT-PCR |
| RT_RSP_0288_A | CGATCAGGCAGGTGGGTGGT | Forward primer for RSP_0288 (<i>bchL</i>) real-time RT-PCR |
| RT_RSP_0288_B | CGTCGAGGTCCGGCATGT | Reverse primer for RSP_0288 (<i>bchL</i>) real-time RT-PCR |
| RT_RSP_0443_A | GGTGGAAAGAGACGCTCAA | Forward primer for RSP_0443 (<i>iscR</i>) real-time RT-PCR |
| RT_RSP_0443_B | ATAGACATGCCCGAGAC | Reverse primer for RSP_0443 (<i>iscR</i>) real-time RT-PCR |
| RT_RSP_0920_A | GCGAGCTTCGAGGAACTG | Forward primer for RSP_0920 (<i>exbB</i>) real-time RT-PCR |
| RT_RSP_0920_B | GTCTGGCTCTCGCAGATG | Reverse primer for RSP_0920 (<i>exbB</i>) real-time RT-PCR |
| RT_RSP_1545_A | CTTCGAGCGCAGAT | Forward primer for RSP_1545 real-time RT-PCR |
| RT_RSP_1545_B | GCCGAGGAACATCG | Reverse primer for RSP_1545 real-time RT-PCR |
| RT_RSP_1547_A | GTCTGCCACTGCATGGCAT | Forward primer for RSP_1547 (<i>bfd</i>) real-time RT-PCR |
| RT_RSP_1547_B | GTTATTCCCTCGGGTTG | Reverse primer for RSP_1547 (<i>bfd</i>) real-time RT-PCR |
| RT_RSP_1669_A | ATCGCGGAAGAGACCCAGAG | Forward primer for RSP_1669 (<i>rpoZ</i>) real-time RT-PCR |
| RT_RSP_1669_B | GAGCAGCGCCATCTGATCCT | Reverse primer for RSP_1669 (<i>rpoZ</i>) real-time RT-PCR |
| RT_RSP_2913_A | CATCAGCCTCGGCAACAC | Forward primer for RSP_2913 (<i>afuA</i>) real-time RT-PCR |
| RT_RSP_2913_B | TGGTTCGTCTCGCGTAG | Reverse primer for RSP_2913 (<i>afuA</i>) real-time RT-PCR |
| RT_RSP_3416_A | CGAGATGATCCGGCT | Forward primer for RSP_3416 real-time RT-PCR |
| RT_RSP_3416_B | GTCGGTCGCGGAATA | Reverse primer for RSP_3416 real-time RT-PCR |
| RT_RSP_3417_A | AGACACGCAAGAGCG | Forward primer for RSP_3417 real-time RT-PCR |
| RT_RSP_3417_B | GCCACCGATTCCACA | Reverse primer for RSP_3417 real-time RT-PCR |
| RT_RSP_4275_A | ACACCTCCTTCAGCT | Forward primer for RSP_4275 (<i>fecI</i>) real-time RT-PCR |
| RT_RSP_4275_B | AGACGCAGGAAGATG | Reverse primer for RSP_4275 (<i>fecI</i>) real-time RT-PCR |
| RT_RSP_6006_A | ATGACCGACCCGATGGAG | Forward primer for RSP_6006 (<i>hemP</i>) real-time RT-PCR |
| RT_RSP_6006_B | AGTAGATCTGCCGTCAG | Reverse primer for RSP_6006 (<i>hemP</i>) real-time RT-PCR |
| RT_RSP_6020_A | CGCGAGGTCAAGGTGAT | Forward primer for RSP_6020 (<i>feoA2</i>) real-time RT-PCR |
| RT_RSP_6020_B | TGCAGCGGAGTGACGAAG | Reverse primer for RSP_6020 (<i>feoA2</i>) real-time RT-PCR |
| suf1RT_A | CGAACTGCACGGCTTCACCT | Forward primer for RT-PCR |
| suf1RT_B | AGGCGCCATTGTCGAGATAG | Reverse primer for RT-PCR |
| suf2RT_A | CGACCTCGAGGCCGCGATCAAAGG | Forward primer for RT-PCR |
| suf2RT_B | GCGAAGTCCATCTCGATCTGGTTTC | Reverse primer for RT-PCR |
| suf3RT_A | CTCGGAGAGGAAGAGGCCGGCTC | Forward primer for RT-PCR |
| suf3RT_B | GTGCCGACCGAGACCGAGTCGAAG | Reverse primer for RT-PCR |
| suf4RT_A | AGACGCAGGTGATGCAGATG | Forward primer for RT-PCR |
| suf4RT_B | GAACCGCAGTCTCTCCAGT | Reverse primer for RT-PCR |
| suf5RT_A | TCTCGCGCATGTCTATGTC | Forward primer for RT-PCR |
| suf5RT_B | CGTCGCATTCCAGTCGAGAT | Reverse primer for RT-PCR |
| 0443_upA | GAGCCGCCAATTCCGGGTGC | Forward primer for RSP_0443 cloning |
| 0443_upB | GAGCGCGGGATCCACCACGCG | Reverse primer for RSP_0443 cloning |
| 0443_downA | GCGTGCAGGATCCTGACCAACCG | Forward primer for RSP_0443 cloning |
| 0443_downB | GAGACAGCCTCAAGCTTCACGTCG | Reverse primer for RSP_0443 cloning |
| IscR_complA | TAATCTAGAACCATCCACCTGGCG | Forward primer for RSP_0443 complementation |
| IscR_complB | CGCTGGATCCTCACTCTGGC | Reverse primer for RSP_0443 complementation |
| IscR_His_fwd | GGCGGGAACCGGGGATCCAAACTCTC | Forward primer for IscR overexpression |
| IscR_His_rev | CGACAGCCAAGCTTCAGTCCTCGTC | Forward primer for IscR overexpression |
| IscR_repA | TCTAGAAAATCACTCGGGCATCGC | Forward primer RSP_0443 for <i>in vivo</i> reporter system |
| IscR_repB | GGATCCTTCCGGTCCCCCAAATC | Reverse primer RSP_0443 for <i>in vivo</i> reporter system |
| IscR_H93A_A | GCGATGGCTACCGGTGCTGGTGCAAGTG | Forward primer inserting a mutation to H93 |
| IscR_H93A_B | GAGACGGTCAATCGATGGCTACCGGTG | Reverse primer inserting a mutation to H93 |
| IscR_H121A_A | CTCGCGGCTGTCTATGTCCTCTGCAC | Forward primer inserting a mutation to H121 |
| IscR_H121A_B | GTGGGAGGGCCTCTCGGGCTGTCTAT | Reverse primer inserting a mutation to H121 |
| IscR_H127A_A | CTTCCTGGCCCAGACCCGTCTGCGAC | Forward primer inserting a mutation to H127 |
| IscR_H127A_B | GCATGTCTATGTCCTCTGGCCCAGACC | Reverse primer inserting a mutation to H127 |
| IscR_C142A_A | GCGTCCAGCCCCGGCGGTGCCGGCGCTG | Forward primer inserting a mutation to C142 |
| IscR_C142A_B | CAAAAACGAGATCGTCCAGCCCCGGCG | Reverse primer inserting a mutation to C142 |
| IscR_P143A_A | GCGTCCAGCCCCGGCGGTGCCGGCGCTG | Forward primer inserting a mutation to P143 |
| IscR_P143A_B | AAACGAGATCGTCCATGCGCGGGGTG | Reverse primer inserting a mutation to P143 |
| iscR_up_fwd | CGCGCGTAATCTTGACAAAAACG | Forward primer for generation of the <i>iscR</i> promoter region |
| iscR_up_rev | CGACACGTCGACAAGCGAGACAAG | Reverse primer for generation of the <i>iscR</i> promoter region |
| hemP_up_fwd | CGCATAAGTCGACCGAAAGAATCAG | Forward primer for generation of the <i>hemP</i> promoter region |
| hemP_up_rev | CCGTCGACAAGGATCCGGGCC | Reverse primer for generation of the <i>hemP</i> promoter region |

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