

## *Supplementary Material for*

Elucidation of sequence-function relationships for an improved biobutanol *in vivo* biosensor in *E. coli*

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### Supplementary Figures

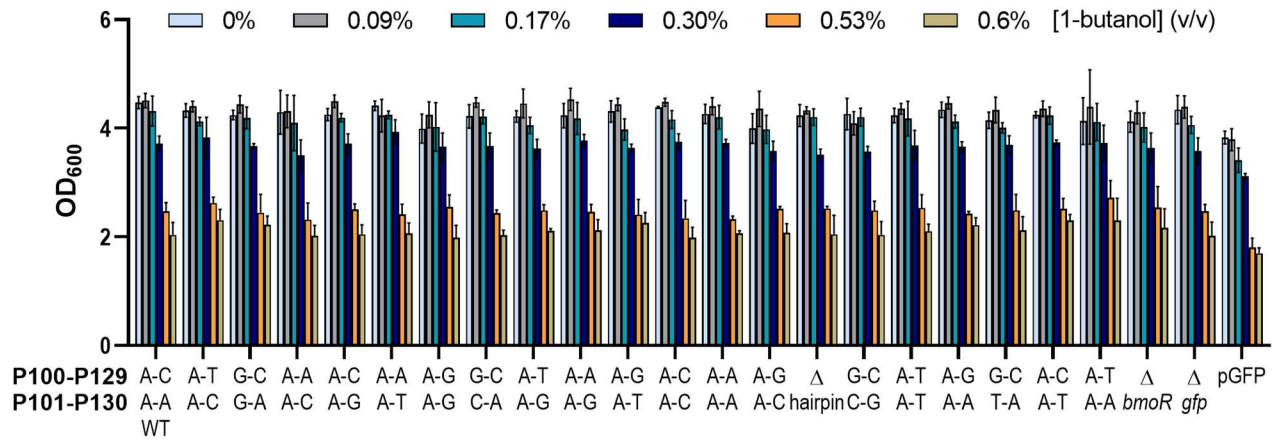
A

P<sub>BMO</sub> Promoter Sequence

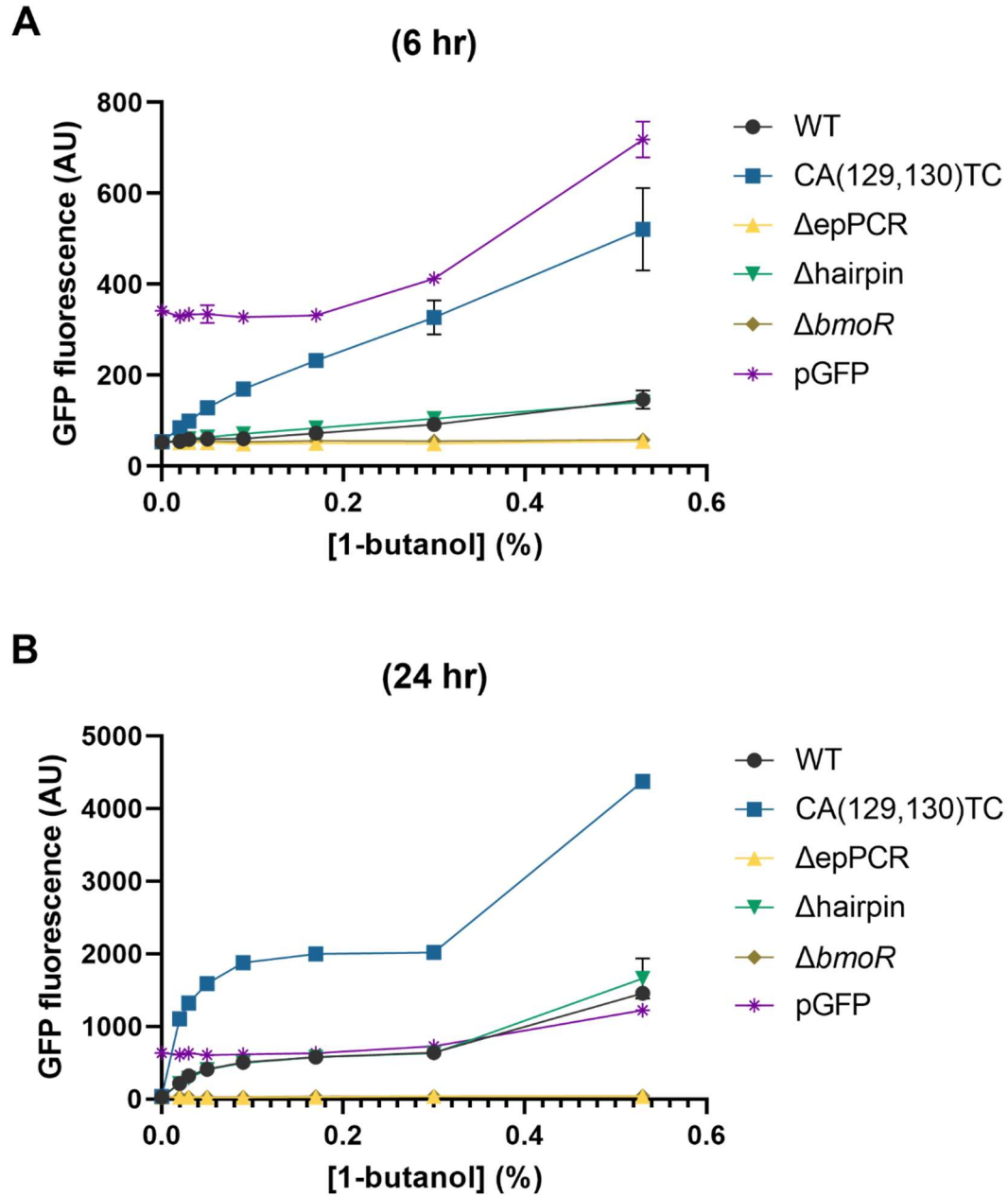
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**B**

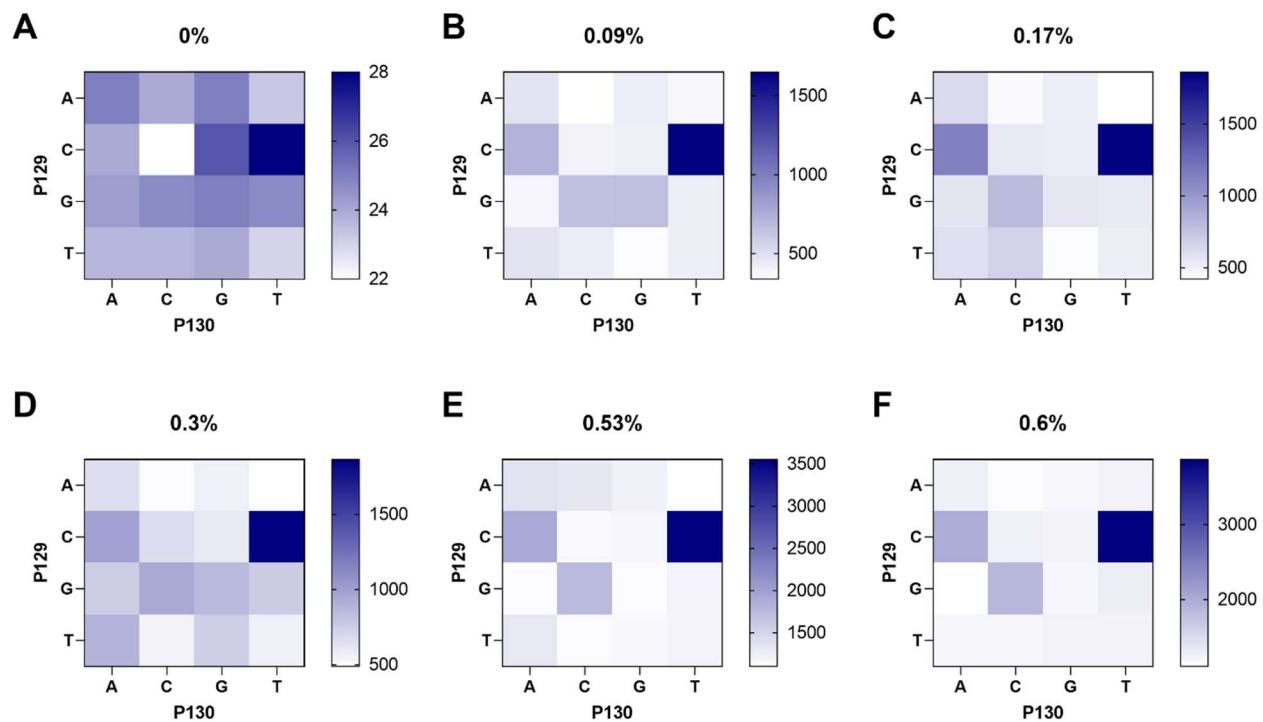
**Supplementary Figure 1. (a)**  $P_{BMO}$  promoter sequence with annotations (error-prone PCR region, putative operator,  $\sigma^{54}$  binding site, +1 TSS) **(b)** Colony PCR confirmation of deletion of the hairpin in  $P_{BMO}$  promoter on 1.6% agarose gel lane 1 – 1 kb+ DNA ladder (NEB); lane 2 –  $P_{BMO}$  WT promoter (413 bp); lane 3 –  $P_{BMO} \Delta$ hairpin (376 bp).



**Supplementary Figure 2.** OD<sub>600</sub> of P<sub>BMO</sub> hairpin mutants. Cell density decrease as the 1-butanol concentration increases.



**Supplementary Figure 3.** Response curves of  $P_{BMO}$  hairpin mutants at **(a)** 6 hours and **(b)** 24 hours post-butanol induction.



**Supplementary Figure 4.** Heat maps of the GFP fluorescence of site-specific mutations at P129 and P130 in  $P_{BMO}$  hairpin mutants. Site-specific dependence of  $P_{BMO}$  hairpin mutation sites P129 and P130 on the right half of the fifteen  $P_{BMO}$  hairpin mutants and the wild-type hairpin when induced with (a) 0%, (b) 0.09%, (c) 0.17%, (d) 0.3%, (e) 0.53%, and (f) 0.6% 1-butanol (v/v). Two-way ANOVA.

## Supplementary Tables

**Supplementary Table 1.** Mutational Bias Analysis of the unsorted P<sub>BMO</sub> library (R=A/G, Y=T/C).

| Mutation Types |          | Mutation Frequency |
|----------------|----------|--------------------|
| Transition     | R→R, Y→Y | 56.4%              |
|                | A→G, T→C | 18.0%              |
|                | G→A, C→T | 38.4%              |
| Transversion   | R→Y, Y→R | 43.6%              |
|                | A→T, T→A | 16.4%              |
|                | A→C, T→G | 2.3%               |
|                | G→C, C→G | 6.5%               |
|                | G→T, C→A | 18.4%              |

**Supplementary Table 2.** Bacterial strains and plasmids used in this study.

| Plasmid                    | Description                                                                                                                               | Strain                 | Purpose                     | Source Reference             |
|----------------------------|-------------------------------------------------------------------------------------------------------------------------------------------|------------------------|-----------------------------|------------------------------|
| P <sub>SELECT#1</sub>      | P <sub>BmoR</sub> - <i>bmoR</i> , P <sub>BMO</sub> - <i>tetA-gfp</i>                                                                      | <i>E. coli</i> (DH1α)  |                             | (Dietrich, Shis et al. 2013) |
| P <sub>SELECT#2</sub>      | P <sub>BmoR</sub> - <i>bmoR</i> , P <sub>BMO</sub> - <i>tetA</i>                                                                          | <i>E. coli</i> (DH1α)  |                             | (Dietrich, Shis et al. 2013) |
| WTΔ <i>bmoR</i>            | ΔP <sub>BmoR</sub> - <i>bmoR</i> , P <sub>BMO</sub> - <i>gfp</i>                                                                          | <i>E. coli</i> (NEB5α) | negative control            | This study                   |
| WTΔ <i>gfp</i>             | P <sub>BmoR</sub> - <i>bmoR</i> , P <sub>BMO</sub> -Δ <i>gfp</i>                                                                          | <i>E. coli</i> (NEB5α) | negative control            | This study                   |
| Δ <i>hairpin</i>           | P <sub>BmoR</sub> - <i>bmoR</i> , P <sub>BMO</sub> Δ <i>hairpin-gfp</i>                                                                   | <i>E. coli</i> (NEB5α) | hairpin characterization    | This study                   |
| ΔepPCR                     | P <sub>BmoR</sub> - <i>bmoR</i> , P <sub>BMO</sub> Δ247-nt- <i>gfp</i>                                                                    | <i>E. coli</i> (NEB5α) | negative control            | This study                   |
| pGFP                       | pUC19 vector; LacI-P <sub>lac</sub> - <i>gfp</i>                                                                                          | <i>E. coli</i> (NEB5α) | positive control            | This study                   |
| CA(129,130)TCΔ <i>bmoR</i> | ΔP <sub>BmoR</sub> - <i>bmoR</i> , P <sub>BMO</sub> - <i>gfp</i> with A <sub>100</sub> A <sub>101</sub> T <sub>129</sub> C <sub>130</sub> | <i>E. coli</i> (NEB5α) | negative control            | This study                   |
| WT                         | P <sub>BmoR</sub> - <i>bmoR</i> , P <sub>BMO</sub> - <i>gfp</i>                                                                           | <i>E. coli</i> (NEB5α) | wild-type control           | This study                   |
| CA(129,130)AA              | P <sub>BmoR</sub> - <i>bmoR</i> , P <sub>BMO</sub> - <i>gfp</i> with A <sub>100</sub> A <sub>101</sub> A <sub>129</sub> A <sub>130</sub>  | <i>E. coli</i> (NEB5α) | P129, P130 characterization | This study                   |

|               |                                                                                                                                             |                           |                                |            |
|---------------|---------------------------------------------------------------------------------------------------------------------------------------------|---------------------------|--------------------------------|------------|
| CA(129,130)AC | P <sub>BmoR</sub> - <i>bmoR</i> , P <sub>BMO</sub> - <i>gfp</i> with A <sub>100</sub><br>A <sub>101</sub> A <sub>129</sub> C <sub>130</sub> | <i>E. coli</i><br>(NEB5α) | P129, P130<br>characterization | This study |
| CA(129,130)AG | P <sub>BmoR</sub> - <i>bmoR</i> , P <sub>BMO</sub> - <i>gfp</i> with A <sub>100</sub><br>A <sub>101</sub> A <sub>129</sub> G <sub>130</sub> | <i>E. coli</i><br>(NEB5α) | P129, P130<br>characterization | This study |
| CA(129,130)AT | P <sub>BmoR</sub> - <i>bmoR</i> , P <sub>BMO</sub> - <i>gfp</i> with A <sub>100</sub><br>A <sub>101</sub> A <sub>129</sub> T <sub>130</sub> | <i>E. coli</i><br>(NEB5α) | P129, P130<br>characterization | This study |
| CA(129,130)CC | P <sub>BmoR</sub> - <i>bmoR</i> , P <sub>BMO</sub> - <i>gfp</i> with A <sub>100</sub><br>A <sub>101</sub> C <sub>129</sub> C <sub>130</sub> | <i>E. coli</i><br>(NEB5α) | P129, P130<br>characterization | This study |
| CA(129,130)CG | P <sub>BmoR</sub> - <i>bmoR</i> , P <sub>BMO</sub> - <i>gfp</i> with A <sub>100</sub><br>A <sub>101</sub> C <sub>129</sub> G <sub>130</sub> | <i>E. coli</i><br>(NEB5α) | P129, P130<br>characterization | This study |
| CA(129,130)CT | P <sub>BmoR</sub> - <i>bmoR</i> , P <sub>BMO</sub> - <i>gfp</i> with A <sub>100</sub><br>A <sub>101</sub> C <sub>129</sub> T <sub>130</sub> | <i>E. coli</i><br>(NEB5α) | P129, P130<br>characterization | This study |
| CA(129,130)GA | P <sub>BmoR</sub> - <i>bmoR</i> , P <sub>BMO</sub> - <i>gfp</i> with A <sub>100</sub><br>A <sub>101</sub> G <sub>129</sub> A <sub>130</sub> | <i>E. coli</i><br>(NEB5α) | P129, P130<br>characterization | This study |
| CA(129,130)GC | P <sub>BmoR</sub> - <i>bmoR</i> , P <sub>BMO</sub> - <i>gfp</i> with A <sub>100</sub><br>A <sub>101</sub> G <sub>129</sub> C <sub>130</sub> | <i>E. coli</i><br>(NEB5α) | P129, P130<br>characterization | This study |
| CA(129,130)GG | P <sub>BmoR</sub> - <i>bmoR</i> , P <sub>BMO</sub> - <i>gfp</i> with A <sub>100</sub><br>A <sub>101</sub> G <sub>129</sub> G <sub>130</sub> | <i>E. coli</i><br>(NEB5α) | P129, P130<br>characterization | This study |
| CA(129,130)GT | P <sub>BmoR</sub> - <i>bmoR</i> , P <sub>BMO</sub> - <i>gfp</i> with A <sub>100</sub><br>A <sub>101</sub> G <sub>129</sub> T <sub>130</sub> | <i>E. coli</i><br>(NEB5α) | P129, P130<br>characterization | This study |
| CA(129,130)TA | P <sub>BmoR</sub> - <i>bmoR</i> , P <sub>BMO</sub> - <i>gfp</i> with A <sub>100</sub><br>A <sub>101</sub> T <sub>129</sub> A <sub>130</sub> | <i>E. coli</i><br>(NEB5α) | P129, P130<br>characterization | This study |
| CA(129,130)TC | P <sub>BmoR</sub> - <i>bmoR</i> , P <sub>BMO</sub> - <i>gfp</i> with A <sub>100</sub><br>A <sub>101</sub> T <sub>129</sub> C <sub>130</sub> | <i>E. coli</i><br>(NEB5α) | P129, P130<br>characterization | This study |
| CA(129,130)TG | P <sub>BmoR</sub> - <i>bmoR</i> , P <sub>BMO</sub> - <i>gfp</i> with A <sub>100</sub><br>A <sub>101</sub> T <sub>129</sub> G <sub>130</sub> | <i>E. coli</i><br>(NEB5α) | P129, P130<br>characterization | This study |
| CA(129,130)TT | P <sub>BmoR</sub> - <i>bmoR</i> , P <sub>BMO</sub> - <i>gfp</i> with A <sub>100</sub><br>A <sub>101</sub> T <sub>129</sub> T <sub>130</sub> | <i>E. coli</i><br>(NEB5α) | P129, P130<br>characterization | This study |
| AA(100,101)TG | P <sub>BmoR</sub> - <i>bmoR</i> , P <sub>BMO</sub> - <i>gfp</i> with T <sub>100</sub><br>G <sub>101</sub> C <sub>129</sub> A <sub>130</sub> | <i>E. coli</i><br>(NEB5α) | P129, P130<br>characterization | This study |
| AA(100,101)CG | P <sub>BmoR</sub> - <i>bmoR</i> , P <sub>BMO</sub> - <i>gfp</i> with C <sub>100</sub><br>G <sub>101</sub> C <sub>129</sub> A <sub>130</sub> | <i>E. coli</i><br>(NEB5α) | P129, P130<br>characterization | This study |
| AA(100,101)GG | P <sub>BmoR</sub> - <i>bmoR</i> , P <sub>BMO</sub> - <i>gfp</i> with G <sub>100</sub><br>G <sub>101</sub> C <sub>129</sub> A <sub>130</sub> | <i>E. coli</i><br>(NEB5α) | P129, P130<br>characterization | This study |

|                               |                                                                                                                               |                                    |                                |            |
|-------------------------------|-------------------------------------------------------------------------------------------------------------------------------|------------------------------------|--------------------------------|------------|
| AACA(100,101,129,130)<br>CGCG | $P_{BmoR}$ - <i>bmoR</i> , $P_{BMO}$ - <i>gfp</i> with C <sub>100</sub><br>G <sub>101</sub> C <sub>129</sub> G <sub>130</sub> | <i>E. coli</i><br>(NEB5 $\alpha$ ) | P129, P130<br>characterization | This study |
| AACA(100,101,129,130)<br>GATC | $P_{BmoR}$ - <i>bmoR</i> , $P_{BMO}$ - <i>gfp</i> with G <sub>100</sub><br>A <sub>101</sub> T <sub>129</sub> C <sub>130</sub> | <i>E. coli</i><br>(NEB5 $\alpha$ ) | P129, P130<br>characterization | This study |
| WT <sub>FLIP</sub>            | $P_{BmoR}$ - <i>bmoR</i> , $P_{BMO}$ - <i>gfp</i> with T <sub>100</sub><br>G <sub>101</sub> T <sub>129</sub> T <sub>130</sub> | <i>E. coli</i><br>(NEB5 $\alpha$ ) | P129, P130<br>characterization | This study |
| CA(129,130)TC <sub>FLIP</sub> | $P_{BmoR}$ - <i>bmoR</i> , $P_{BMO}$ - <i>gfp</i> with G <sub>100</sub><br>A <sub>101</sub> T <sub>129</sub> T <sub>130</sub> | <i>E. coli</i><br>(NEB5 $\alpha$ ) | P129, P130<br>characterization | This study |
| G(21)A                        | $P_{BmoR}$ - <i>bmoR</i> , $P_{BMO}$ - <i>gfp</i> with A <sub>21</sub>                                                        | <i>E. coli</i><br>(NEB5 $\alpha$ ) | Validation                     | This study |
| C(32)G                        | $P_{BmoR}$ - <i>bmoR</i> , $P_{BMO}$ - <i>gfp</i> with G <sub>21</sub>                                                        | <i>E. coli</i><br>(NEB5 $\alpha$ ) | Validation                     | This study |
| G(48)C                        | $P_{BmoR}$ - <i>bmoR</i> , $P_{BMO}$ - <i>gfp</i> with C <sub>48</sub>                                                        | <i>E. coli</i><br>(NEB5 $\alpha$ ) | Validation                     | This study |
| T(52)G                        | $P_{BmoR}$ - <i>bmoR</i> , $P_{BMO}$ - <i>gfp</i> with G <sub>52</sub>                                                        | <i>E. coli</i><br>(NEB5 $\alpha$ ) | Validation                     | This study |
| G(77)C                        | $P_{BmoR}$ - <i>bmoR</i> , $P_{BMO}$ - <i>gfp</i> with C <sub>77</sub>                                                        | <i>E. coli</i><br>(NEB5 $\alpha$ ) | Validation                     | This study |
| C(129)T                       | $P_{BmoR}$ - <i>bmoR</i> , $P_{BMO}$ - <i>gfp</i> with T <sub>129</sub>                                                       | <i>E. coli</i><br>(NEB5 $\alpha$ ) | Validation                     | This study |
| A(130)C                       | $P_{BmoR}$ - <i>bmoR</i> , $P_{BMO}$ - <i>gfp</i> with C <sub>130</sub>                                                       | <i>E. coli</i><br>(NEB5 $\alpha$ ) | Validation                     | This study |
| T(131)G                       | $P_{BmoR}$ - <i>bmoR</i> , $P_{BMO}$ - <i>gfp</i> with G <sub>131</sub>                                                       | <i>E. coli</i><br>(NEB5 $\alpha$ ) | Validation                     | This study |
| C(133)T                       | $P_{BmoR}$ - <i>bmoR</i> , $P_{BMO}$ - <i>gfp</i> with T <sub>133</sub>                                                       | <i>E. coli</i><br>(NEB5 $\alpha$ ) | Validation                     | This study |
| T(135)G                       | $P_{BmoR}$ - <i>bmoR</i> , $P_{BMO}$ - <i>gfp</i> with G <sub>135</sub>                                                       | <i>E. coli</i><br>(NEB5 $\alpha$ ) | Validation                     | This study |
| C(162)G                       | $P_{BmoR}$ - <i>bmoR</i> , $P_{BMO}$ - <i>gfp</i> with G <sub>162</sub>                                                       | <i>E. coli</i><br>(NEB5 $\alpha$ ) | Validation                     | This study |
| C(185)G                       | $P_{BmoR}$ - <i>bmoR</i> , $P_{BMO}$ - <i>gfp</i> with G <sub>185</sub>                                                       | <i>E. coli</i><br>(NEB5 $\alpha$ ) | Validation                     | This study |
| G(186)C                       | $P_{BmoR}$ - <i>bmoR</i> , $P_{BMO}$ - <i>gfp</i> with C <sub>186</sub>                                                       | <i>E. coli</i><br>(NEB5 $\alpha$ ) | Validation                     | This study |
| G(188)T                       | $P_{BmoR}$ - <i>bmoR</i> , $P_{BMO}$ - <i>gfp</i> with G <sub>188</sub>                                                       | <i>E. coli</i><br>(NEB5 $\alpha$ ) | Validation                     | This study |
| C(190)T                       | $P_{BmoR}$ - <i>bmoR</i> , $P_{BMO}$ - <i>gfp</i> with T <sub>190</sub>                                                       | <i>E. coli</i><br>(NEB5 $\alpha$ ) | IHF                            | This study |
| C(193)G                       | $P_{BmoR}$ - <i>bmoR</i> , $P_{BMO}$ - <i>gfp</i> with G <sub>193</sub>                                                       | <i>E. coli</i><br>(NEB5 $\alpha$ ) | IHF                            | This study |
| G(196)A                       | $P_{BmoR}$ - <i>bmoR</i> , $P_{BMO}$ - <i>gfp</i> with A <sub>196</sub>                                                       | <i>E. coli</i><br>(NEB5 $\alpha$ ) | Validation                     | This study |
| G(200)A                       | $P_{BmoR}$ - <i>bmoR</i> , $P_{BMO}$ - <i>gfp</i> with A <sub>200</sub>                                                       | <i>E. coli</i><br>(NEB5 $\alpha$ ) | Validation                     | This study |



|                                        |                                                                                                                                                                            |                        |            |            |
|----------------------------------------|----------------------------------------------------------------------------------------------------------------------------------------------------------------------------|------------------------|------------|------------|
| A(202)T                                | P <sub>BmoR</sub> - <i>bmoR</i> , P <sub>BMO</sub> - <i>gfp</i> with T <sub>202</sub>                                                                                      | <i>E. coli</i> (NEB5α) | IHF        | This study |
| G(205)A                                | P <sub>BmoR</sub> - <i>bmoR</i> , P <sub>BMO</sub> - <i>gfp</i> with A <sub>205</sub>                                                                                      | <i>E. coli</i> (NEB5α) | Validation | This study |
| GG(186,188)CT                          | P <sub>BmoR</sub> - <i>bmoR</i> , P <sub>BMO</sub> - <i>gfp</i> with C <sub>186</sub> T <sub>188</sub>                                                                     | <i>E. coli</i> (NEB5α) | Validation | This study |
| TCT(131,133,135)GTG                    | P <sub>BmoR</sub> - <i>bmoR</i> , P <sub>BMO</sub> - <i>gfp</i> with G <sub>131</sub> T <sub>133</sub> G <sub>135</sub>                                                    | <i>E. coli</i> (NEB5α) | Validation | This study |
| CA(129,130)TC//G(21)A                  | P <sub>BmoR</sub> - <i>bmoR</i> , P <sub>BMO</sub> - <i>gfp</i> with A <sub>100</sub> A <sub>101</sub> T <sub>129</sub> C <sub>130</sub> A <sub>21</sub>                   | <i>E. coli</i> (NEB5α) | Validation | This study |
| CA(129,130)TC//C(32)G                  | P <sub>BmoR</sub> - <i>bmoR</i> , P <sub>BMO</sub> - <i>gfp</i> with A <sub>100</sub> A <sub>101</sub> T <sub>129</sub> C <sub>130</sub> G <sub>32</sub>                   | <i>E. coli</i> (NEB5α) | Validation | This study |
| CA(129,130)TC//G(48)C                  | P <sub>BmoR</sub> - <i>bmoR</i> , P <sub>BMO</sub> - <i>gfp</i> with A <sub>100</sub> A <sub>101</sub> T <sub>129</sub> C <sub>130</sub> C <sub>48</sub>                   | <i>E. coli</i> (NEB5α) | Validation | This study |
| CA(129,130)TC//T(52)G                  | P <sub>BmoR</sub> - <i>bmoR</i> , P <sub>BMO</sub> - <i>gfp</i> with A <sub>100</sub> A <sub>101</sub> T <sub>129</sub> C <sub>130</sub> G <sub>52</sub>                   | <i>E. coli</i> (NEB5α) | Validation | This study |
| CA(129,130)TC//G(77)C                  | P <sub>BmoR</sub> - <i>bmoR</i> , P <sub>BMO</sub> - <i>gfp</i> with A <sub>100</sub> A <sub>101</sub> T <sub>129</sub> C <sub>130</sub> C <sub>77</sub>                   | <i>E. coli</i> (NEB5α) | Validation | This study |
| CA(129,130)TC//T(131)G                 | P <sub>BmoR</sub> - <i>bmoR</i> , P <sub>BMO</sub> - <i>gfp</i> with A <sub>100</sub> A <sub>101</sub> T <sub>129</sub> C <sub>130</sub> G <sub>131</sub>                  | <i>E. coli</i> (NEB5α) | Validation | This study |
| CA(129,130)TC//C(133)T                 | P <sub>BmoR</sub> - <i>bmoR</i> , P <sub>BMO</sub> - <i>gfp</i> with A <sub>100</sub> A <sub>101</sub> T <sub>129</sub> C <sub>130</sub> T <sub>133</sub>                  | <i>E. coli</i> (NEB5α) | Validation | This study |
| CA(129,130)TC//T(135)G                 | P <sub>BmoR</sub> - <i>bmoR</i> , P <sub>BMO</sub> - <i>gfp</i> with A <sub>100</sub> A <sub>101</sub> T <sub>129</sub> C <sub>130</sub> G <sub>135</sub>                  | <i>E. coli</i> (NEB5α) | Validation | This study |
| CA(129,130)TC//C(162)G                 | P <sub>BmoR</sub> - <i>bmoR</i> , P <sub>BMO</sub> - <i>gfp</i> with A <sub>100</sub> A <sub>101</sub> T <sub>129</sub> C <sub>130</sub> G <sub>162</sub>                  | <i>E. coli</i> (NEB5α) | Validation | This study |
| CA(129,130)TC//C(185)G                 | P <sub>BmoR</sub> - <i>bmoR</i> , P <sub>BMO</sub> - <i>gfp</i> with A <sub>100</sub> A <sub>101</sub> T <sub>129</sub> C <sub>130</sub> G <sub>185</sub>                  | <i>E. coli</i> (NEB5α) | Validation | This study |
| CA(129,130)TC//G(186)C                 | P <sub>BmoR</sub> - <i>bmoR</i> , P <sub>BMO</sub> - <i>gfp</i> with A <sub>100</sub> A <sub>101</sub> T <sub>129</sub> C <sub>130</sub> C <sub>186</sub>                  | <i>E. coli</i> (NEB5α) | Validation | This study |
| CA(129,130)TC//G(188)T                 | P <sub>BmoR</sub> - <i>bmoR</i> , P <sub>BMO</sub> - <i>gfp</i> with A <sub>100</sub> A <sub>101</sub> T <sub>129</sub> C <sub>130</sub> T <sub>188</sub>                  | <i>E. coli</i> (NEB5α) | Validation | This study |
| CA(129,130)TC//G(196)A                 | P <sub>BmoR</sub> - <i>bmoR</i> , P <sub>BMO</sub> - <i>gfp</i> with A <sub>100</sub> A <sub>101</sub> T <sub>129</sub> C <sub>130</sub> A <sub>196</sub>                  | <i>E. coli</i> (NEB5α) | Validation | This study |
| CA(129,130)TC//G(200)A                 | P <sub>BmoR</sub> - <i>bmoR</i> , P <sub>BMO</sub> - <i>gfp</i> with A <sub>100</sub> A <sub>101</sub> T <sub>129</sub> C <sub>130</sub> A <sub>200</sub>                  | <i>E. coli</i> (NEB5α) | Validation | This study |
| CA(129,130)TC//G(205)A                 | P <sub>BmoR</sub> - <i>bmoR</i> , P <sub>BMO</sub> - <i>gfp</i> with A <sub>100</sub> A <sub>101</sub> T <sub>129</sub> C <sub>130</sub> A <sub>205</sub>                  | <i>E. coli</i> (NEB5α) | Validation | This study |
| CA(129,130)TC//G,G(186,188)C,T         | P <sub>BmoR</sub> - <i>bmoR</i> , P <sub>BMO</sub> - <i>gfp</i> with A <sub>100</sub> A <sub>101</sub> T <sub>129</sub> C <sub>130</sub> C <sub>186</sub> T <sub>188</sub> | <i>E. coli</i> (NEB5α) | Validation | This study |
| CA(129,130)TC//T,C,T(131,133,135)G,T,G | P <sub>BmoR</sub> - <i>bmoR</i> , P <sub>BMO</sub> - <i>gfp</i> with A <sub>100</sub> A <sub>101</sub> T <sub>129</sub> G <sub>131</sub> T <sub>133</sub> G <sub>135</sub> | <i>E. coli</i> (NEB5α) | Validation | This study |
| Δhairpin//G(21)A                       | P <sub>BmoR</sub> - <i>bmoR</i> , P <sub>BMO</sub> Δhairpin- <i>gfp</i> with A <sub>21</sub>                                                                               | <i>E. coli</i> (NEB5α) | Validation | This study |
| Δhairpin//C(32)G                       | P <sub>BmoR</sub> - <i>bmoR</i> , P <sub>BMO</sub> Δhairpin- <i>gfp</i> with G <sub>32</sub>                                                                               | <i>E. coli</i> (NEB5α) | Validation | This study |
| Δhairpin//G(48)C                       | P <sub>BmoR</sub> - <i>bmoR</i> , P <sub>BMO</sub> Δhairpin- <i>gfp</i> with C <sub>48</sub>                                                                               | <i>E. coli</i> (NEB5α) | Validation | This study |
| Δhairpin//T(52)G                       | P <sub>BmoR</sub> - <i>bmoR</i> , P <sub>BMO</sub> Δhairpin- <i>gfp</i> with G <sub>52</sub>                                                                               | <i>E. coli</i> (NEB5α) | Validation | This study |

|                                   |                                                                                                               |                                 |            |            |
|-----------------------------------|---------------------------------------------------------------------------------------------------------------|---------------------------------|------------|------------|
| $\Delta$ hairpin//G(77)C          | $P_{BmoR}$ - <i>bmoR</i> , $P_{BMO}$ $\Delta$ hairpin- <i>gfp</i> with C <sub>77</sub>                        | <i>E. coli</i> (NEB5 $\alpha$ ) | Validation | This study |
| $\Delta$ hairpin//C(162)G         | $P_{BmoR}$ - <i>bmoR</i> , $P_{BMO}$ $\Delta$ hairpin- <i>gfp</i> with G <sub>162</sub>                       | <i>E. coli</i> (NEB5 $\alpha$ ) | Validation | This study |
| $\Delta$ hairpin//C(185)G         | $P_{BmoR}$ - <i>bmoR</i> , $P_{BMO}$ $\Delta$ hairpin- <i>gfp</i> with G <sub>185</sub>                       | <i>E. coli</i> (NEB5 $\alpha$ ) | Validation | This study |
| $\Delta$ hairpin//G(186)C         | $P_{BmoR}$ - <i>bmoR</i> , $P_{BMO}$ $\Delta$ hairpin- <i>gfp</i> with C <sub>186</sub>                       | <i>E. coli</i> (NEB5 $\alpha$ ) | Validation | This study |
| $\Delta$ hairpin//G(188)T         | $P_{BmoR}$ - <i>bmoR</i> , $P_{BMO}$ $\Delta$ hairpin- <i>gfp</i> with T <sub>188</sub>                       | <i>E. coli</i> (NEB5 $\alpha$ ) | Validation | This study |
| $\Delta$ hairpin//G(196)A         | $P_{BmoR}$ - <i>bmoR</i> , $P_{BMO}$ $\Delta$ hairpin- <i>gfp</i> with A <sub>196</sub>                       | <i>E. coli</i> (NEB5 $\alpha$ ) | Validation | This study |
| $\Delta$ hairpin//G(200)A         | $P_{BmoR}$ - <i>bmoR</i> , $P_{BMO}$ $\Delta$ hairpin- <i>gfp</i> with A <sub>200</sub>                       | <i>E. coli</i> (NEB5 $\alpha$ ) | Validation | This study |
| $\Delta$ hairpin//G(205)A         | $P_{BmoR}$ - <i>bmoR</i> , $P_{BMO}$ $\Delta$ hairpin- <i>gfp</i> with A <sub>205</sub>                       | <i>E. coli</i> (NEB5 $\alpha$ ) | Validation | This study |
| $\Delta$ hairpin//G,G(186,188)C,T | $P_{BmoR}$ - <i>bmoR</i> , $P_{BMO}$ $\Delta$ hairpin- <i>gfp</i> with C <sub>186</sub> C <sub>188</sub>      | <i>E. coli</i> (NEB5 $\alpha$ ) | Validation | This study |
| IHF-5'-half                       | $P_{BmoR}$ - <i>bmoR</i> , $P_{BMO}$ - <i>gfp</i> with GCGTTCGCTCCCGCGGC <sub>174-190</sub>                   | <i>E. coli</i> (NEB5 $\alpha$ ) | IHF        | This study |
| IHF-3'-half                       | $P_{BmoR}$ - <i>bmoR</i> , $P_{BMO}$ - <i>gfp</i> with GCCGCGGGTGTACCGTT <sub>191-207</sub>                   | <i>E. coli</i> (NEB5 $\alpha$ ) | IHF        | This study |
| IHF-5' & 3'                       | $P_{BmoR}$ - <i>bmoR</i> , $P_{BMO}$ - <i>gfp</i> with GCGTTCGCTCCCGCGGCGCC GCGGGTGTACCGTT <sub>174-207</sub> | <i>E. coli</i> (NEB5 $\alpha$ ) | IHF        | This study |
| IHF-pspA                          | $P_{BmoR}$ - <i>bmoR</i> , $P_{BMO}$ - <i>gfp</i> with TCAATCAGATCTTTATAAAT CAAAAAGATAAAAA <sub>174-207</sub> | <i>E. coli</i> (NEB5 $\alpha$ ) | IHF        | This study |
| IHF-glnHp2                        | $P_{BmoR}$ - <i>bmoR</i> , $P_{BMO}$ - <i>gfp</i> with TTTGCCGCATCTCGAAAAAT CAATGAATTACTCG <sub>174-207</sub> | <i>E. coli</i> (NEB5 $\alpha$ ) | IHF        | This study |
| IHF- $\lambda$ H'                 | $P_{BmoR}$ - <i>bmoR</i> , $P_{BMO}$ - <i>gfp</i> with GCCAAAAAAGCATTGCTTAT CAATTGTTGCACC <sub>174-207</sub>  | <i>E. coli</i> (NEB5 $\alpha$ ) | IHF        | This study |

**Supplementary Table 3.** Primers used to construct plasmids in this study. Mutations and overhangs hifi assembly primer regions are in lower case letters.

| Plasmid                     | Name  | Forward Primer (5'-->3')                   | Name  | Reverse Primer (5'-->3')               |
|-----------------------------|-------|--------------------------------------------|-------|----------------------------------------|
| WT $\Delta bmoR$            | NK155 | CCACAGATAGTAGGTGCTG                        | NK212 | AGGTGGCACTTTTCGGG                      |
| WT $\Delta gfp$             | NK107 | CTACAAATAAGGATCCTAACT<br>CGAGTCTAGACC      | NK129 | TTGTGTGTTCTGCTGTCCGTAG                 |
| $\Delta$ hairpin            | NK218 | TTGCCACACCCAACCGGA                         | NK219 | AATCTTCCGCGCTGTCCG                     |
| $\Delta$ epPCR              | NK210 | TCTTTAACGTGTAACACACG                       | NK160 | GACCTTGAGGTGACCTTG                     |
| CA(129,130)TC $\Delta bmoR$ | NK155 | CCACAGATAGTAGGTGCTG                        | NK212 | AGGTGGCACTTTTCGGG                      |
| WT $\Delta bmoR$            | NK155 | CCACAGATAGTAGGTGCTG                        | NK212 | AGGTGGCACTTTTCGGG                      |
| pGFP                        | NK109 | ctttactcatAGCTGTTTCCT<br>GTGTGAAATTGTTATCC | NK110 | ctacaaataaAAGGGCCTCGTG<br>ATACGCCT     |
|                             | NK111 | cgaggcccttTTATTTGTAGA<br>GCTCATCCATG       | NK112 | ggaaacagctATGAGTAAAGGA<br>GAAGAACTTTTC |
| CA(129,130)AA               | NK1   | CCTCGGGCTGaaTCCTTGCCAC                     | NK16  | CACGCACGCTCGGGCTGT                     |
| CA(129,130)AC               | NK2   | CCTCGGGCTGacTCCTTGCCACACC                  | NK16  | CACGCACGCTCGGGCTGT                     |
| CA(129,130)AG               | NK3   | CCTCGGGCTGagTCCTTGCCACACCAAC               | NK16  | CACGCACGCTCGGGCTGT                     |
| CA(129,130)AT               | NK4   | CCTCGGGCTGatTCCTTGCCACACC                  | NK16  | CACGCACGCTCGGGCTGT                     |
| CA(129,130)CC               | NK5   | CCTCGGGCTGccTCCTTGCCAC                     | NK16  | CACGCACGCTCGGGCTGT                     |
| CA(129,130)CG               | NK6   | CCTCGGGCTGcgTCCTTGCCAC                     | NK16  | CACGCACGCTCGGGCTGT                     |
| CA(129,130)CT               | NK7   | CCTCGGGCTGctTCCTTGCCAC                     | NK16  | CACGCACGCTCGGGCTGT                     |
| CA(129,130)GA               | NK8   | CCTCGGGCTGgaTCCTTGCCAC                     | NK16  | CACGCACGCTCGGGCTGT                     |
| CA(129,130)GC               | NK9   | CCTCGGGCTGgcTCCTTGCCACACC                  | NK16  | CACGCACGCTCGGGCTGT                     |
| CA(129,130)GG               | NK10  | CCTCGGGCTGggTCCTTGCCACACCAAC               | NK16  | CACGCACGCTCGGGCTGT                     |
| CA(129,130)GT               | NK11  | CCTCGGGCTGgtTCCTTGCCACACC                  | NK16  | CACGCACGCTCGGGCTGT                     |
| CA(129,130)TA               | NK12  | CCTCGGGCTGtaTCCTTGCCAC                     | NK16  | CACGCACGCTCGGGCTGT                     |
| CA(129,130)TC               | NK13  | CCTCGGGCTGtcTCCTTGCCACACCAAC               | NK16  | CACGCACGCTCGGGCTGT                     |
| CA(129,130)TG               | NK14  | CCTCGGGCTGtgTCCTTGCCACACCAAC               | NK16  | CACGCACGCTCGGGCTGT                     |
| CA(129,130)TT               | NK15  | CCTCGGGCTGttTCCTTGCCACACCAAC               | NK16  | CACGCACGCTCGGGCTGT                     |
| AA(100,101)TG               | NK119 | GAAGATTGGATgCAGCCCGAG<br>CGTGCGTGCC        | NK121 | CGCGCTGTCCGCCGAGGT                     |
| AA(100,101)CG               | NK120 | GAAGATTGGAcgCAGCCCGAG<br>CGTGCGTGCC        | NK121 | CGCGCTGTCCGCCGAGGT                     |
| AA(100,101)GG               | NK123 | GAAGATTGGANNcAGCCCGAG<br>CGTGCGTGCC        | NK121 | CGCGCTGTCCGCCGAGGT                     |
| AACA(100,101,129,130)CGCG   | NK120 | GAAGATTGGAcgCAGCCCGAG<br>CGTGCGTGCC        | NK121 | CGCGCTGTCCGCCGAGGT                     |

|                               |       |                                           |       |                                           |
|-------------------------------|-------|-------------------------------------------|-------|-------------------------------------------|
| AACA(100,101,129,130)GATC     | NK331 | gtgcctcgggctgtcTCCTTG<br>CCACACCCAACC     | NK332 | gcacgctcgggctgtcTCCAAT<br>CTTCCGCGCTGTC   |
| WT <sub>FLIP</sub>            | NK292 | cacgctcgggctgtttccTTG<br>CCACACCCAACCGGA  | NK293 | cgtgcctcgggctgcatccAAT<br>CTTCCGCGCTGTCCG |
| CA(129,130)TC <sub>FLIP</sub> | NK292 | cacgctcgggctgtttccTTG<br>CCACACCCAACCGGA  | NK294 | cgtgcctcgggctgtctccAAT<br>CTTCCGCGCTGTCCG |
| G(21)A                        | NK226 | aGCTGCTCATGCTCCTGTGCGC                    | NK241 | GCAGCACCTACTATCTGTGGG                     |
| C(32)G                        | NK227 | GGCTGCTCATGgTCCTGTGCGC                    | NK241 | GCAGCACCTACTATCTGTGGG                     |
| G(48)C                        | NK228 | GGTAGCcCGCTGTTACGCGAC<br>C                | NK242 | GCGACAGGAGCATGAGCAG                       |
| T(52)G                        | NK229 | GGTAGCGCGCgGTTACGCGAC                     | NK242 | GCGACAGGAGCATGAGCAG                       |
| G(77)C                        | NK230 | CCCGGACCTCcCGGACAGCG                      | NK243 | GGCGGTTCGCGTAACAGCG                       |
| C(129)T                       | NK12  | CCTCGGGCTGtaTCCTTGCCA<br>C                | NK16  | CACGCACGCTCGGGCTGT                        |
| A(130)C                       | NK05  | CCTCGGGCTGccTCCTTGCCA<br>C                | NK16  | CACGCACGCTCGGGCTGT                        |
| T(131)G                       | NK231 | GGGCTGCAgCCTTGCCACACC                     | NK244 | GAGGCACGCACGCTCGGG                        |
| C(133)T                       | NK232 | GGGCTGCATctTTGCCACACC<br>C                | NK244 | GAGGCACGCACGCTCGGG                        |
| T(135)G                       | NK233 | GGGCTGCATCCTgGCCACACC<br>C                | NK244 | GAGGCACGCACGCTCGGG                        |
| C(162)G                       | NK234 | TTCGTCGGACgGCTCGACATT<br>CGC              | NK245 | TCCGGTTGGGTGTGGCAAG                       |
| C(185)G                       | NK235 | CCGCcGGAGCGAACGCGAATG<br>TC               | NK246 | CGCCGCGGGTGTACCGTTG                       |
| G(186)C                       | NK236 | CCGgGGGAGCGAACGCGAATG<br>TC               | NK246 | CGCCGCGGGTGTACCGTTG                       |
| G(188)T                       | NK237 | CaGCGGGAGCGAACGCGAATG<br>TC               | NK246 | CGCCGCGGGTGTACCGTTG                       |
| C(190)T                       | NK325 | gccgcgggtgtaccgttGCGT<br>TACAGATGTACCCTTC | NK326 | accgcgggagcgaacgcGAATG<br>TCGAGCGGTCCGA   |
| C(193)G                       | NK327 | gccgcgggtgtaccgttGCGT<br>TACAGATGTACCCTTC | NK328 | gccgcgggagcgaacgcGAATG<br>TCGAGCGGTCCGA   |
| G(196)A                       | NK238 | CGCCGCaGGTGTACCGTTGCG                     | NK247 | CCGCGGGAGCGAACGCG                         |
| G(200)A                       | NK239 | CGCCGCGGGTaTACCGTTGCG                     | NK247 | CCGCGGGAGCGAACGCG                         |
| A(202)T                       | NK329 | gccgcgggtgttccgttGCGT<br>TACAGATGTACCCTTC | NK330 | gccgcgggagcgaacgcGAATG<br>TCGAGCGGTCCGA   |
| G(205)A                       | NK240 | CGCCGCGGGTGTACCaTTGCG<br>TTACAG           | NK247 | CCGCGGGAGCGAACGCG                         |
| GG(186,188)CT                 | NK253 | CaGgGGGAGCGAACGCGAATG<br>TC               | NK246 | CGCCGCGGGTGTACCGTTG                       |
| TCT(131,133,135)GT<br>G       | NK254 | TCGGGCTGCAgCtTgGCCACA<br>CCCAACC          | NK252 | GGCACGCACGCTCGGGCT                        |
| CA(129,130)TC//G(21)<br>A     | NK13  | CCTCGGGCTGtcTCCTTGCCA<br>CACCCAAC         | NK16  | CACGCACGCTCGGGCTGT                        |
| CA(129,130)TC//C(32)<br>G     | NK13  | CCTCGGGCTGtcTCCTTGCCA<br>CACCCAAC         | NK16  | CACGCACGCTCGGGCTGT                        |
| CA(129,130)TC//G(48)<br>C     | NK13  | CCTCGGGCTGtcTCCTTGCCA<br>CACCCAAC         | NK16  | CACGCACGCTCGGGCTGT                        |
| CA(129,130)TC//T(52)<br>G     | NK13  | CCTCGGGCTGtcTCCTTGCCA<br>CACCCAAC         | NK16  | CACGCACGCTCGGGCTGT                        |

|                                        |       |                                           |       |                                          |
|----------------------------------------|-------|-------------------------------------------|-------|------------------------------------------|
| CA(129,130)TC//G(77)C                  | NK13  | CCTCGGGCTGtcTCCTTGCCA<br>CACCCAAC         | NK16  | CACGCACGCTCGGGCTGT                       |
| CA(129,130)TC//T(131)G                 | NK248 | GGGCTGTcCCTTGCCACACC                      | NK244 | GAGGCACGCACGCTCGGG                       |
| CA(129,130)TC//C(133)T                 | NK249 | GGGCTGTCTcTTGCCACACC<br>C                 | NK244 | GAGGCACGCACGCTCGGG                       |
| CA(129,130)TC//T(135)G                 | NK250 | GGGCTGTCTCCTgGCCACACC<br>C                | NK244 | GAGGCACGCACGCTCGGG                       |
| CA(129,130)TC//C(162)G                 | NK13  | CCTCGGGCTGtcTCCTTGCCA<br>CACCCAAC         | NK16  | CACGCACGCTCGGGCTGT                       |
| CA(129,130)TC//C(185)G                 | NK13  | CCTCGGGCTGtcTCCTTGCCA<br>CACCCAAC         | NK16  | CACGCACGCTCGGGCTGT                       |
| CA(129,130)TC//G(186)C                 | NK13  | CCTCGGGCTGtcTCCTTGCCA<br>CACCCAAC         | NK16  | CACGCACGCTCGGGCTGT                       |
| CA(129,130)TC//G(188)T                 | NK13  | CCTCGGGCTGtcTCCTTGCCA<br>CACCCAAC         | NK16  | CACGCACGCTCGGGCTGT                       |
| CA(129,130)TC//G(196)A                 | NK13  | CCTCGGGCTGtcTCCTTGCCA<br>CACCCAAC         | NK16  | CACGCACGCTCGGGCTGT                       |
| CA(129,130)TC//G(200)A                 | NK13  | CCTCGGGCTGtcTCCTTGCCA<br>CACCCAAC         | NK16  | CACGCACGCTCGGGCTGT                       |
| CA(129,130)TC//G(205)A                 | NK13  | CCTCGGGCTGtcTCCTTGCCA<br>CACCCAAC         | NK16  | CACGCACGCTCGGGCTGT                       |
| CA(129,130)TC//G,G(186,188)C,T         | NK13  | CCTCGGGCTGtcTCCTTGCCA<br>CACCCAAC         | NK16  | CACGCACGCTCGGGCTGT                       |
| CA(129,130)TC//T,C,T(131,133,135)G,T,G | NK251 | TCGGGCTGTcGcTgGCCACA<br>CCCAACC           | NK252 | GGCAGGCACGCTCGGGCT                       |
| $\Delta$ hairpin//G(21)A               | NK226 | aGCTGCTCATGCTCCTGTcGC                     | NK241 | GCAGCACCTACTATCTGTGGG                    |
| $\Delta$ hairpin//C(32)G               | NK227 | GGCTGCTCATGgTCCTGTcGC                     | NK241 | GCAGCACCTACTATCTGTGGG                    |
| $\Delta$ hairpin//G(48)C               | NK228 | GGTAGCcCGCTGTTACGCGAC<br>C                | NK242 | GCGACAGGAGCATGAGCAG                      |
| $\Delta$ hairpin//T(52)G               | NK229 | GGTAGCGCGcGTTACGCGAC                      | NK242 | GCGACAGGAGCATGAGCAG                      |
| $\Delta$ hairpin//G(77)C               | NK230 | CCCGGACCTCcGCGGACAGCG                     | NK243 | GGCGGTTCGCGTAACAGCG                      |
| $\Delta$ hairpin//C(162)G              | NK234 | TTCGTcCGGACgGCTCGACATT<br>CGC             | NK245 | TCCGGTTGGGTGTGGCAAG                      |
| $\Delta$ hairpin//C(185)G              | NK235 | CCGCcGGAGCGAACGCGAATG<br>TC               | NK246 | CGCCGCGGGTGTACCGTTG                      |
| $\Delta$ hairpin//G(186)C              | NK236 | CCGgGGGAGCGAACGCGAATG<br>TC               | NK246 | CGCCGCGGGTGTACCGTTG                      |
| $\Delta$ hairpin//G(188)T              | NK237 | CaGCGGGAGCGAACGCGAATG<br>TC               | NK246 | CGCCGCGGGTGTACCGTTG                      |
| $\Delta$ hairpin//G(196)A              | NK238 | CGCCGCaGGTGTACCGTTGCG                     | NK247 | CCGCGGGAGCGAACGCG                        |
| $\Delta$ hairpin//G(200)A              | NK239 | CGCCGCGGGTaTACCGTTGCG                     | NK247 | CCGCGGGAGCGAACGCG                        |
| $\Delta$ hairpin//G(205)A              | NK240 | CGCCGCGGGTGTACCaTTGCG<br>TTACAG           | NK247 | CCGCGGGAGCGAACGCG                        |
| $\Delta$ hairpin//G,G(186,188)C,T      | NK253 | CaGgGGGAGCGAACGCGAATG<br>TC               | NK246 | CGCCGCGGGTGTACCGTTG                      |
| IHF-5'-half                            | NK319 | aatcaaggtgttccattGCGT<br>TACAGATGTACCCTTC | NK320 | gccgcgggagcgaacgcGAATG<br>TCGAGCGGTCCGA  |
| IHF-3'-half                            | NK321 | gccgcgggtgtaccgttGCGT<br>TACAGATGTACCCTTC | NK322 | ataaggatatacgaacccGAATG<br>TCGAGCGGTCCGA |
| IHF-5' & 3'                            | NK323 | aatcaaggtgttccattGCGT<br>TACAGATGTACCCTTC | NK324 | ataaggatatacgaacccGAATG<br>TCGAGCGGTCCGA |

|            |       |                                           |       |                                         |
|------------|-------|-------------------------------------------|-------|-----------------------------------------|
| IHF-pspA   | NK313 | aatcaaaaagataaaaaGCGT<br>TACAGATGTACCCTTC | NK314 | tataaagatctgattgaGAATG<br>TCGAGCGGTCCGA |
| IHF-glnHp2 | NK315 | aatcaatgaattactcgGCGT<br>TACAGATGTACCCTTC | NK316 | tttcgagatgcggcaaaGAATG<br>TCGAGCGGTCCGA |
| IHF-λH'    | NK317 | tatcaatttgttgaccGCGT<br>TACAGATGTACCCTTC  | NK318 | agcaatgcttttttggcGAATG<br>TCGAGCGGTCCGA |

**Supplementary Table 4.** Library generation, Sanger & NGS, and Additional Primers. Mutations and overhangs HiFi assembly primer regions are in lower case letters.

| Name                               | Name  | Forward Primer (5'-->3')                                     | Name  | Reverse Primer (5'-->3')                                      |
|------------------------------------|-------|--------------------------------------------------------------|-------|---------------------------------------------------------------|
| Sanger sequencing primer 1         |       |                                                              | NK22  | GCATTGAACACCATAAGAGAAAGTA<br>GTGACA                           |
| Sanger sequencing primer 2         |       |                                                              | NK129 | TTGTGTGTTCTGCTGTCGGTAG                                        |
| correcting BmoR Alanine->Valine 31 | NK222 | GACATCAGCCaCTTTCCCCGG                                        | NK223 | GTTTTGCGCTCGTGGACC                                            |
| epPCR library generation           | NK161 | ctcaagggtcacctcaagggtcCC<br>ACAGATAGTAGGTGCTG                | NK211 | ggcgtgtgttacacgttaaagaAGG<br>GTACATCTGTAACGC                  |
| epPCR library Vector               | NK210 | TCTTTAACGTGTAACACACG                                         | NK160 | GACCTTGAGGTGACCTTG                                            |
| Amplicon Illumina partial adapters | NK224 | ACACTCTTTCCCTACACGACGC<br>TCTTCCGATCTccacagatagt<br>aggtgctg | NK225 | GACTGGAGTTCAGACGTGTGCTCTT<br>CCGATCTggcgtgtgttacacgtta<br>aag |

## References

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