

1 Supplementary materials for

2 **Metabolic engineering of a probiotic *Saccharomyces boulardii***

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7 (Running title: Metabolic engineering of *Saccharomyces boulardii*)

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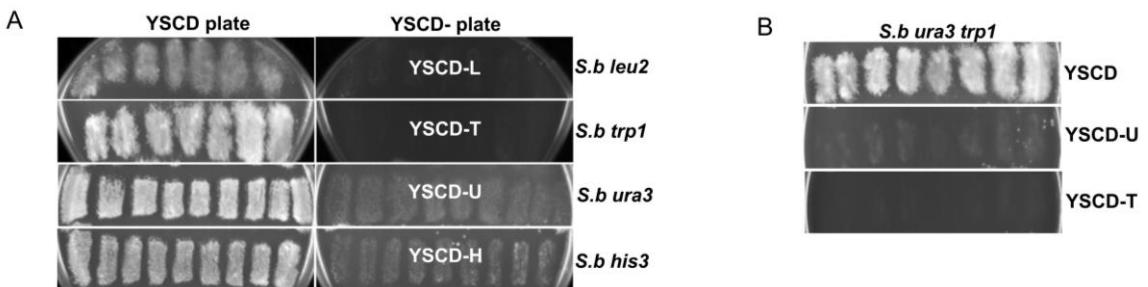
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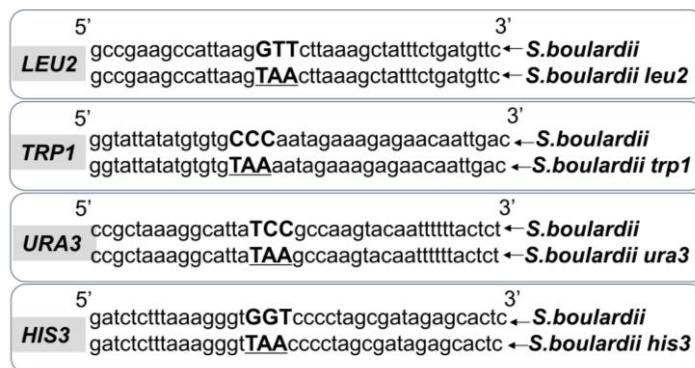
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24 **Fig. S1** The screen of auxotrophic markers inactivation. (A) The screen of single
25 deletion of each auxotrophic markers; (B) Identification of *ura3 trp1* double
26 inactivation using tandem guide RNA. *S. b* is short for *Saccharomyces boulardii*.

27 YSCD: yeast synthetic complete medium with 20 g/L glucose as a carbon source;
28 YSCD-Leu: YSCD medium minus leucine; YSCD-Trp: YSCD medium minus
29 tryptophan; YSCD-Ura: YSCD medium minus uracil; YSCD-His: YSCD medium
30 minus histidine.

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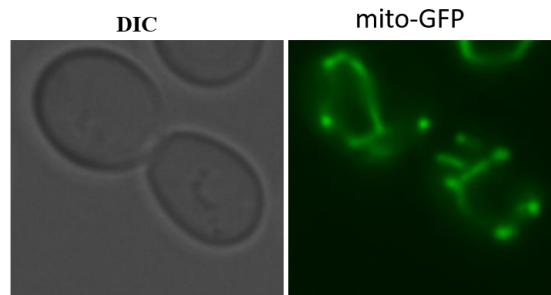


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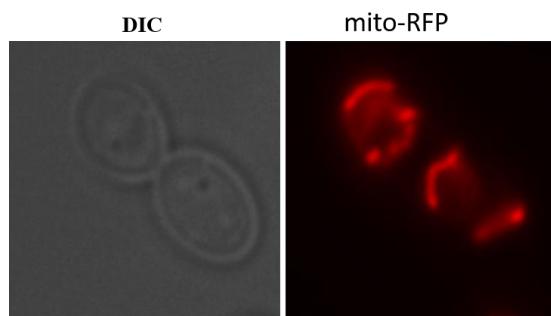
34 **Fig. S2** The sequencing results of four marker genes *LEU2*, *TRP1*, *URA3* and *HIS3*
35 indicated that the targeted nucleotides were replaced by the stop codon (TAA) as
36 expected.

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38



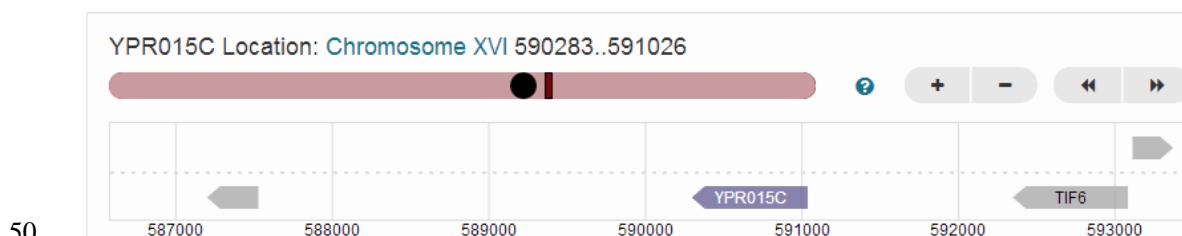
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40 **Fig. S3.** Targeted localization of the green fluorescent protein (GFP) and red
41 fluorescent protein (RFP) in mitochondria of *S. boulardii* expressing GFP and mRuby
42 with a mitochondrial targeting sequence by introducing pVT100U-mtGFP and
43 pVT100U-mt-mRuby plasmid, respectively. The left panel is differential interference
44 contrast (DIC) images; right panel is GFP and RFP fluorescence.
45

46 **Intergenic site CS8 for CRISPR-Cas9 based integration.**

47 The 2.7 kb non-coding region between YPR015C and a CDS of unknown function
48 from chromosome XVI was chosen for further target gene insertion. The map and
49 genome sequence from Yeastgenome.org are listed as follows:



51 <http://www.yeastgenome.org/locus/S000006219/overview>

96 aacgttaacattttacaattttagctcgactactatgtttaagtaatccaaagggaactatttgttatcccagaattttaca
97 ttctgtttcttagttcataaacaatgaatacctattgaatggatagaattctgacttgatttacgagttatttgcacattgc
98 caaagacatctcgtttttccctctacaacccaatgaggaggctagccaggcgtgcggccaaaaaaatagccaataaag
99 cgcaacttcgt**ATGTCTTGCATTTTCCGCTGACTTGGAGTAGAGTACA**
100 **GCTGCGCGGAGTCGCGTATTACAAATCTGTACTTGCATTCTGTGCAT**
101 **ACGTGAGGAAAAAGCAGCACCTGTTGTGAAAAAAAGATAAATTCTTTT**
102 **TTTCATTTCTATCCTCAAAGGAAAATCTTCTGCGAAATTCTAGC**
103 **CGCTCTTCAGTTCCGAAGTAAAGTCGCTGAGGGCGAAAAACAACTTT**
104 **ATTGCCCACACGCTGTTAGGCTCCAGGGAGCAAAAGCGCAATTG**
105 **GGAATAAGGTTCTCAATTGAATGGTTGCTCGCCTACAGCATTAAc**
106 cagaaggtaattgtatcttgtatgtccgctaacagatctgtccattaattgttattttctcaggtaggtggctgagccacttgt
107 agtagctccctaccattactttggcctgtagttatggcttgatttcttcatgacaatccggttagcaaccctgcgaag
108 ggcttttggaaaaaaaggcgtggcaagatctgtat.....

109

110 Human lysozyme with chicken signal peptide.

111 The gBlock sequence of human lysozyme with a chicken-lysozyme signal sequence

112 based on previous report (1, 2) is listed as follows:

113

114 NNNN: enzyme site for the confirmation of insertion of gBlock into plasmid

115 nnnn: Chicken-lysozyme signal sequence

116 nnnn: Starting codon

117 nnnn: Stop codon

118

119 GGTACCA**atgaggtcttgc**taatcttggtgcttgc~~ttcctggctgtctgggg~~aaggtttcgaacgttgtaa
120 tggccagaacttgaagagattggatggacggttaccgttatctttggctactgatgtttggccaagtggaaat
121 ctggttacaacactagagactactaactacaacgcccgtgaccgttactgactacggtatcttccaaattaactctagatact
122 ggtgtacgcacggtaagactccaggcgccgttaacgcctgtcagttgttctgttgcagacaacatcgctgac
123 gccgttgcctgtgctaagagagtcgttagagaccacaaggatcagagcttgggtgc~~ttggcgt~~acagatgtcaaaac
124 agagacgtcagacaatacgttcaaggttgtgtgtctaa**GTCGAC**

125

126 References:

- 127 **1. Muraki M, Jigami Y, Tanaka H, Harada N, Kishimoto F, Agui H, Ogino S, Nakasato S.** 1986.
128 Expression of Synthetic Human Lysozyme Gene in Escherichia coli. Agricultural and
129 Biological Chemistry **50**:713-723.

130 **2. Jigami Y, Muraki M, Harada N, Tanaka H.** 1986. Expression of synthetic human-lysozyme gene in
131 *Saccharomyces cerevisiae*: use of a synthetic chicken-lysozyme signal sequence for secretion
132 and processing. Gene **43**:273-279.