

## Supplementary Materials

### **Coupling metabolic addiction with negative autoregulation to improve strain stability and pathway yield**

Yongkun Lv<sup>1,2,3</sup>, Yang Gu<sup>1,3</sup>, Jingliang Xu<sup>2</sup>, Jingwen Zhou<sup>3</sup> and Peng Xu<sup>1\*</sup>

<sup>1</sup> Department of Chemical, Biochemical and Environmental Engineering, University of Maryland Baltimore County, Baltimore, MD 21250, USA.

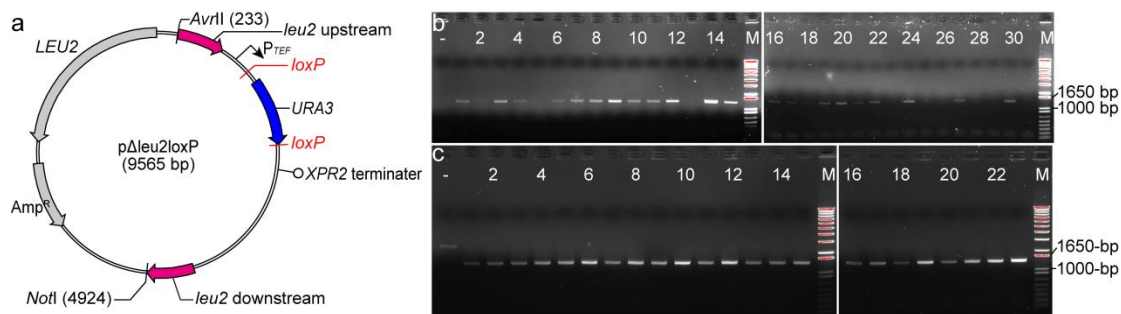
<sup>2</sup> School of Chemical Engineering, Zhengzhou University, Zhengzhou, Henan 450001, China.

<sup>3</sup> National Engineering Laboratory for Cereal Fermentation Technology, Jiangnan University, Wuxi, Jiangsu 214122, China.

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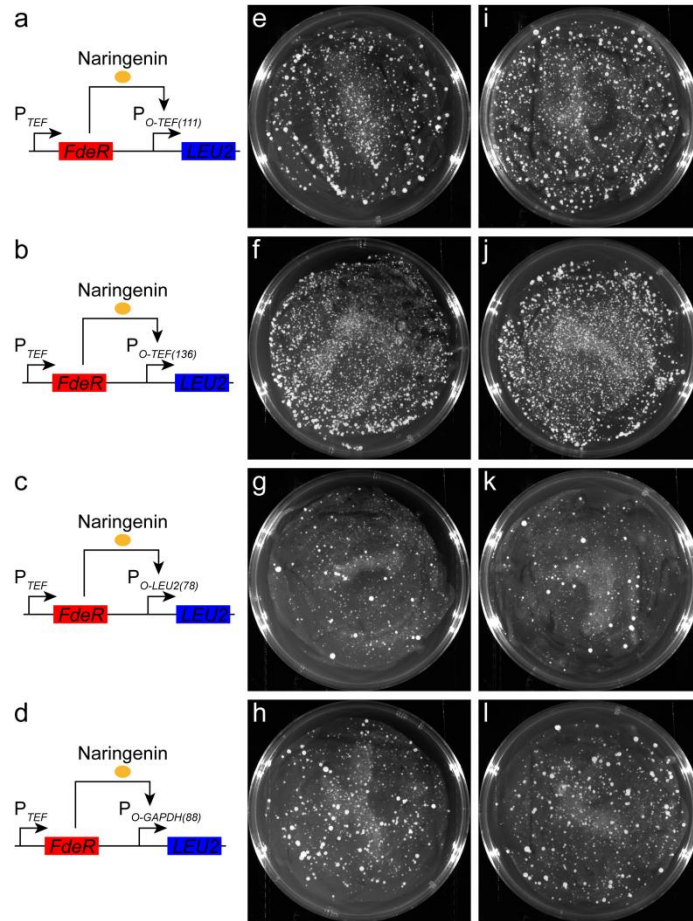
\*Corresponding author Tel: +1(410)-455-2474; fax: +1(410)-455-1049. E-mail address: zhoujw1982@jiangnan.edu.cn (JWZ), and pengxu@umbc.edu (PX).

## Supplementary figures

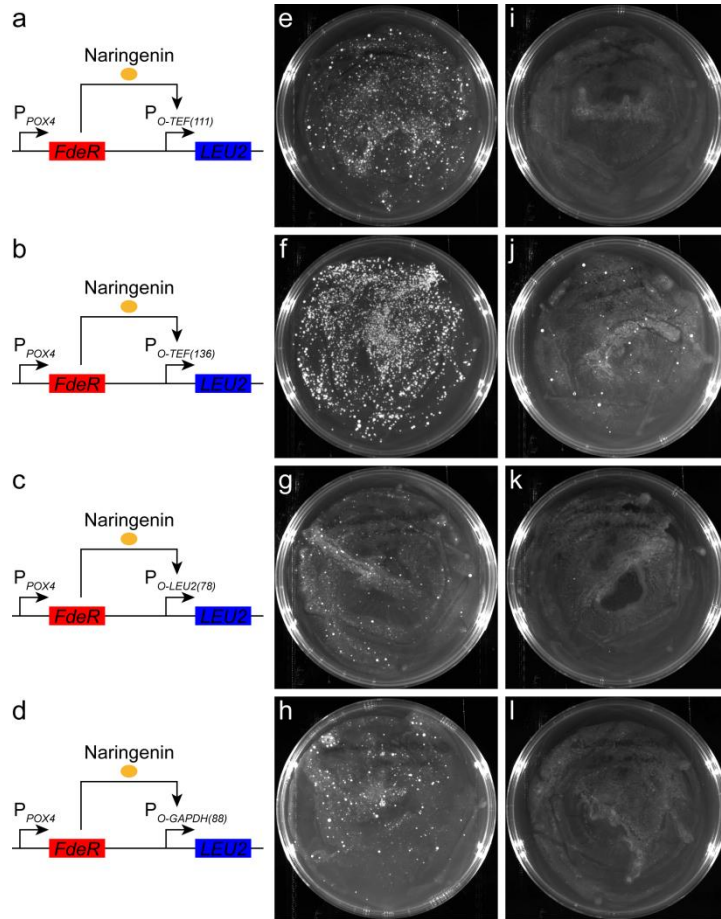


**Supplementary Fig. 1** Site-directed integration at *leu2* loci and *URA3* marker curation.

**a** Structure of plasmid pΔleu2loxP. *leu2* upstream and *leu2* downstream refer to the up and down homologous sequence of *leu2* loci. The *loxP* sites were used for *URA3* marker curation. **b** Colony PCR analysis of site-directed integration at *leu2* site. The colony PCR was carried out using primer pair *leu2\_Inte F/leu2\_Inte R*. The positive colony will yield a 1368-bp band, while the negative colony will not yield any specific band. **c** Colony PCR analysis of *URA3* marker curation. The colony PCR was carried out using primer pair *26srDNA2s F/XPR2\_Seq*. The positive colony will yield a 1533-bp band, while the negative colony will yield a 2432-bp band.

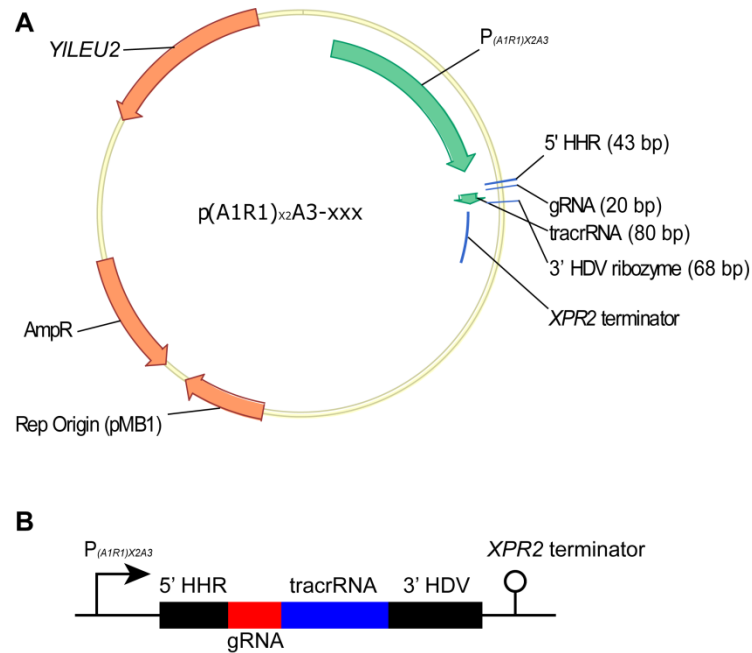


**Supplementary Fig. 2** Analysis of naringenin addiction circuits on CSM-Leu and CSM-Leu+Naringenin plates. **a-d** Components of naringenin addiction circuits containing promoters  $P_{O-TEF(111)}$ ,  $P_{O-TEF(136)}$ ,  $P_{O-LEU2(78)}$ , and  $P_{O-GAPDH(88)}$ . **e-h** The control set. Growth of *Y. lipolytica* transformants containing promoters  $P_{O-TEF(111)}$ ,  $P_{O-TEF(136)}$ ,  $P_{O-LEU2(78)}$ , and  $P_{O-GAPDH(88)}$  on CSM-Leu plates. Pictures were taken 3 days after transformation. **i-l** Growth of *Y. lipolytica* transformants containing promoters  $P_{O-TEF(111)}$ ,  $P_{O-TEF(136)}$ ,  $P_{O-LEU2(78)}$ , and  $P_{O-GAPDH(88)}$  on CSM-Leu+Naringenin plates. Pictures were taken 3 days after transformation.



**Supplementary Fig. 3** Analysis of complete and incomplete naringenin addiction circuits in naringenin producing strain NarPro/ASC\_Rep. **a-d** Components of naringenin addiction circuits containing promoters  $P_{O-TEF(111)}$ ,  $P_{O-TEF(136)}$ ,  $P_{O-LEU2(78)}$ , and  $P_{O-GAPDH(88)}$ . FdeR was expressed using weaker promoter  $P_{POX4}$ . **e-h** Growth of NarPro/ASC\_Rep containing complete addiction circuits on leucine drop-out CSM-leu plates. *LEU2* expression was controlled by  $P_{O-TEF(111)}$ ,  $P_{O-TEF(136)}$ ,  $P_{O-LEU2(78)}$ , and  $P_{O-GAPDH(88)}$ , respectively. Pictures were taken 3 days after transformation. **i-l** Negative control set. Growth of NarPro/ASC\_Rep containing incomplete addiction circuits without FdeR on leucine drop-out CSM-leu plates. *LEU2* expression was controlled by  $P_{O-TEF(111)}$ ,  $P_{O-TEF(136)}$ ,  $P_{O-LEU2(78)}$ , and  $P_{O-GAPDH(88)}$ , respectively. Pictures were taken 3 days

after transformation.



**Supplementary Fig. 4. Genetic structure of plasmid  $p(A1R1)_{x2}A3-xxx$ .** “xxx” here refers to gRNA. For instance,  $p(A1R1)_{x2}A3-FAS1-2$  refers to  $p(A1R1)_{x2}A3$  containing gRNA FAS1-2 generating components. **A.** Overall structure of plasmid  $p(A1R1)_{x2}A3-xxx$ . **B.** Structure of the gRNA generating components. The 5' HHR and 3' HDV ribozyme sites were used for generating mature gRNAs.

## Supplementary tables

**Supplementary Table 1 gRNA sequences and targeting locations**

gRNA	Sequence (5'-3')	Targeting locations
FAS1-1	ACTCACCAAATCAAATAATG	UTR
FAS1-2	TTTTGCTACAGGAAACAGCG	Intron
FAS1-3	GGTGCAGTTGATGTACAGAG	CDS
FAS2-1	AAACCTGAGCCACAAATCAG	Intron
FAS2-2	GGGCATCACCACCAGCAGCG	CDS
FAS2-3	GAAGGGGACAACAATCAGAG	CDS
FabD-1	GCAGCATTCTTCCCCGACA	CDS
FabD-2	GAGGCGTTGGATACCACCTG	CDS
FabD-3	CACATGATATACACAAAGCA	Intron

## Supplementary Table 2 Plasmids used in this paper

Plasmid	Annotation
pYLXP'	A frozen stock of our Lab <sup>1</sup> .
pYLXP'-Nluc	A frozen stock of our Lab <sup>2</sup> .
pYLXP'-dCas9	A frozen stock of our Lab.
pYLXP'-Cre	A frozen stock of our Lab <sup>3</sup> .
pYLXP'-URA3-loxP	A frozen stock of our Lab <sup>3</sup> .
pΔleu2loxP	Integration at <i>leu2</i> site.
pΔxpr2loxP	Integration at <i>xpr2</i> site.
pΔleu2loxP-POX2(1591)-Nluc	Integrating POX2(1591)-Nluc at <i>leu2</i> site.
pΔleu2loxP-A1R1A3-Nluc	Integrating A1R1A3-Nluc at <i>leu2</i> site.
pΔleu2loxP-(A1R1) <sub>x2</sub> A3-Nluc	Integrating (A1R1) <sub>x2</sub> A3-Nluc at <i>leu2</i> site.
pΔleu2loxP-FAS1-1-dCas9	Integrating gRNA FAS1-1 and dCas9 circuits at <i>leu2</i> site.
pΔleu2loxP-FAS1-2-dCas9	Integrating gRNA FAS1-2 and dCas9 circuits at <i>leu2</i> site.
pΔleu2loxP-FAS1-3-dCas9	Integrating gRNA FAS1-3 and dCas9 circuits at <i>leu2</i> site.
pΔleu2loxP-FAS2-1-dCas9	Integrating gRNA FAS2-1 and dCas9 circuits at <i>leu2</i> site.
pΔleu2loxP-FAS2-2-dCas9	Integrating gRNA FAS2-2 and dCas9 circuits at <i>leu2</i> site.
pΔleu2loxP-FAS2-3-dCas9	Integrating gRNA FAS2-3 and dCas9 circuits at <i>leu2</i> site.
pΔleu2loxP-FabD-1-dCas9	Integrating gRNA FabD-1 and dCas9 circuits at <i>leu2</i> site.
pΔleu2loxP-FabD-2-dCas9	Integrating gRNA FabD-2 and dCas9 circuits at <i>leu2</i> site.
pΔleu2loxP-FabD-3-dCas9	Integrating gRNA FabD-3 and dCas9 circuits at <i>leu2</i> site.
pΔleu2loxP-FAS1-2-FAS2-1-dCas9	Integrating gRNAs FAS1-2 and FAS2-1 and dCas9 circuits at <i>leu2</i> site.
pΔleu2loxP-FAS1-2-FabD-1-dCas9	Integrating gRNAs FAS1-2 and FabD-1 and dCas9 circuits at <i>leu2</i> site.
pΔleu2loxP-FAS2-1-FabD-1-dCas9	Integrating gRNAs FAS2-1 and FabD-1 and dCas9 circuits at <i>leu2</i> site.
pΔleu2loxP-FAS1-2-FAS2-1-FabD-1-dCas9	Integrating gRNAs FAS1-2, FAS2-1, and FabD-1 and dCas9 circuits at <i>leu2</i> site.
pOTEF(111)	Replacing <i>TEF</i> promoter in pYLXP'2 with <i>OTEF(111)</i> hybrid promoter.
pOTEF(136)	Replacing <i>TEF</i> promoter in pYLXP'2 with <i>OTEF(136)</i> hybrid promoter.
pOLEU2(78)	Replacing <i>TEF</i> promoter in pYLXP'2 with <i>OLEU2(78)</i> hybrid promoter.
pOGAPDH(88)	Replacing <i>TEF</i> promoter in pYLXP'2 with <i>OGAPDH(88)</i> hybrid promoter.
pOTEF(111)-Nluc	Placing <i>Nluc</i> under the control of hybrid promoter <i>OTEF(111)</i> .
pOTEF(136)-Nluc	Placing <i>Nluc</i> under the control of hybrid promoter <i>OTEF(136)</i> .

pOLEU2(78)-Nluc	Placing <i>Nluc</i> under the control of hybrid promoter <i>OLEU2(78)</i> .
pOGAPDH(88)-Nluc	Placing <i>Nluc</i> under the control of hybrid promoter <i>OGAPDH(88)</i> .
pOTEF(111)-LEU2	Placing <i>LEU2</i> under the control of hybrid promoter <i>OTEF(111)</i> .
pOTEF(136)-LEU2	Placing <i>LEU2</i> under the control of hybrid promoter <i>OTEF(136)</i> .
pOLEU2(78)-LEU2	Placing <i>LEU2</i> under the control of hybrid promoter <i>OLEU2(78)</i> .
pOGAPDH(88)-LEU2	Placing <i>LEU2</i> under the control of hybrid promoter <i>OGAPDH(88)</i> .
pYLP'-FdeR	Placing <i>FdeR</i> under the control of promoter <i>TEF</i> .
pYaliJ1-FdeR	Placing <i>FdeR</i> under the control of promoter <i>POX4</i> .
pYaliL1-FdeR	Placing <i>FdeR</i> under the control of promoter <i>IDP2</i> .
pOTEF(111)-FdeR	Placing <i>FdeR</i> under the control of hybrid promoter <i>OTEF(111)</i> .
pOTEF(136)-FdeR	Placing <i>FdeR</i> under the control of hybrid promoter <i>OTEF(136)</i> .
pOLEU2(78)-FdeR	Placing <i>FdeR</i> under the control of hybrid promoter <i>OLEU2(78)</i> .
pOGAPDH(88)-FdeR	Placing <i>FdeR</i> under the control of hybrid promoter <i>OGAPDH(88)</i> .
pOTEF(111)-Nluc-FdeR	pOTEF(111)-Nluc containing <i>FdeR</i> under the control of <i>TEF</i> promoter.
pOTEF(136)-Nluc-FdeR	pOTEF(136)-Nluc containing <i>FdeR</i> under the control of <i>TEF</i> promoter.
pOLEU2(78)-Nluc-FdeR	pOLEU2(78)-Nluc containing <i>FdeR</i> under the control of <i>TEF</i> promoter.
pOGAPDH(88)-Nluc-FdeR	pOGAPDH(88)-Nluc containing <i>FdeR</i> under the control of <i>TEF</i> promoter.
pOTEF(111)-LEU2-FdeR	pOTEF(111)-LEU2 containing <i>FdeR</i> under the control of <i>TEF</i> promoter.
pOTEF(136)-LEU2-FdeR	pOTEF(136)-LEU2 containing <i>FdeR</i> under the control of <i>TEF</i> promoter.
pOLEU2(78)-LEU2-FdeR	pOLEU2(78)-LEU2 containing <i>FdeR</i> under the control of <i>TEF</i> promoter.
pOGAPDH(88)-LEU2-FdeR	pOGAPDH(88)-LEU2 containing <i>FdeR</i> under the control of <i>TEF</i> promoter.
pOTEF(111)-LEU2-POX4-FdeR	pOTEF(111)-LEU2 containing <i>FdeR</i> under the control of <i>POX4</i> promoter.
pOTEF(136)-LEU2-POX4-FdeR	pOTEF(136)-LEU2 containing <i>FdeR</i> under the control of <i>POX4</i> promoter.
pOLEU2(78)-LEU2-POX4-FdeR	pOLEU2(78)-LEU2 containing <i>FdeR</i> under the control of <i>POX4</i> promoter.
pOGAPDH(88)-LEU2-POX4-FdeR	pOGAPDH(88)-LEU2 containing <i>FdeR</i> under the control of <i>POX4</i> promoter.
pOTEF(111)-LEU2-IDP2-FdeR	pOTEF(111)-LEU2 containing <i>FdeR</i> under the control of <i>IDP2</i> promoter.
pOTEF(136)-LEU2-IDP2-FdeR	pOTEF(136)-LEU2 containing <i>FdeR</i> under the control of <i>IDP2</i>



	promoter.
pOLEU2(78)-LEU2-IDP2-FdeR	pOLEU2(78)-LEU2 containing <i>FdeR</i> under the control of <i>IDP2</i> promoter.
pOGAPDH(88)-LEU2-IDP2-FdeR	pOGAPDH(88)-LEU2 containing <i>FdeR</i> under the control of <i>IDP2</i> promoter.
pOTEF(136)-LEU2-OTEF(111)-FdeR	pOTEF(136)-LEU2 containing <i>FdeR</i> under the control of <i>OTEF(111)</i> promoter.
pOTEF(136)-LEU2-OTEF(136)-FdeR	pOTEF(136)-LEU2 containing <i>FdeR</i> under the control of <i>OTEF(136)</i> promoter.
pOTEF(136)-LEU2-OLEU2(78)-FdeR	pOTEF(136)-LEU2 containing <i>FdeR</i> under the control of <i>OLEU2(78)</i> promoter.
pOTEF(136)-LEU2-OGAPDH(88)-FdeR	pOTEF(136)-LEU2 containing <i>FdeR</i> under the control of <i>OGAPDH(88)</i> promoter.
pΔxpr2loxP-OTEF(111)-Nluc-FdeR	Integrating OTEF(111)-Nluc-FdeR at <i>xpr2</i> site. Used for testing <i>OTEF(111)</i> promoter.
pΔxpr2loxP-OTEF(136)-Nluc-FdeR	Integrating OTEF(136)-Nluc-FdeR at <i>xpr2</i> site. Used for testing <i>OTEF(136)</i> promoter.
pΔxpr2loxP-OLEU2(78)-Nluc-FdeR	Integrating OLEU2(78)-Nluc-FdeR at <i>xpr2</i> site. Used for testing <i>OLEU2(78)</i> promoter.
pΔxpr2loxP-OGAPDH(88)-Nluc-FdeR	Integrating OGAPDH(88)-Nluc-FdeR at <i>xpr2</i> site. Used for testing <i>OGAPDH(88)</i> promoter.
pΔxpr2loxP-OTEF(111)-LEU2	Integrating OTEF(111)-LEU2 at <i>xpr2</i> site. Used as control.
pΔxpr2loxP-OTEF(136)-LEU2	Integrating OTEF(136)-LEU2 at <i>xpr2</i> site. Used as control.
pΔxpr2loxP-OLEU2(78)-LEU2	Integrating OLEU2(78)-LEU2 at <i>xpr2</i> site. Used as control.
pΔxpr2loxP-OGAPDH(88)-LEU2	Integrating OGAPDH(88)-LEU2 at <i>xpr2</i> site. Used as control.
pΔxpr2loxP-OTEF(111)-LEU2-FdeR	Integrating OTEF(111)-LEU2-FdeR at <i>xpr2</i> site.
pΔxpr2loxP-OTEF(136)-LEU2-FdeR	Integrating OTEF(136)-LEU2-FdeR at <i>xpr2</i> site.
pΔxpr2loxP-OLEU2(78)-LEU2-FdeR	Integrating OLEU2(78)-LEU2-FdeR at <i>xpr2</i> site.
pΔxpr2loxP-OGAPDH(88)-LEU2-FdeR	Integrating OGAPDH(88)-LEU2-FdeR at <i>xpr2</i> site.
pΔxpr2loxP-OTEF(111)-LEU2-POX4-FdeR	Integrating OTEF(111)-LEU2-POX4-FdeR at <i>xpr2</i> site.
pΔxpr2loxP-OTEF(136)-LEU2-POX4-FdeR	Integrating OTEF(136)-LEU2-POX4-FdeR at <i>xpr2</i> site.

pΔxpr2loxP-OLEU2(78)-LEU2-POX4-FdeR	Integrating OLEU2(78)-LEU2-POX4-FdeR at <i>xpr2</i> site.
pΔxpr2loxP-OGAPDH(88)-LEU2-POX4-FdeR	Integrating OGAPDH(88)-LEU2-POX4-FdeR at <i>xpr2</i> site.
pΔxpr2loxP-OTEF(111)-LEU2-IDP2-FdeR	Integrating OTEF(111)-LEU2-IDP2-FdeR at <i>xpr2</i> site.
pΔxpr2loxP-OTEF(136)-LEU2-IDP2-FdeR	Integrating OTEF(136)-LEU2-IDP2-FdeR at <i>xpr2</i> site.
pΔxpr2loxP-OLEU2(78)-LEU2-IDP2-FdeR	Integrating OLEU2(78)-LEU2-IDP2-FdeR at <i>xpr2</i> site.
pΔxpr2loxP-OGAPDH(88)-LEU2-IDP2-FdeR	Integrating OGAPDH(88)-LEU2-IDP2-FdeR at <i>xpr2</i> site.
pΔxpr2loxP-OTEF(136)-LEU2-OTEF(111)-FdeR	Integrating OTEF(136)-LEU2-OTEF(111)-FdeR at <i>xpr2</i> site.
pΔxpr2loxP-OTEF(136)-LEU2-OTEF(136)-FdeR	Integrating OTEF(136)-LEU2-OTEF(136)-FdeR at <i>xpr2</i> site.
pΔxpr2loxP-OTEF(136)-LEU2-OLEU2(78)-FdeR	Integrating OTEF(136)-LEU2-OLEU2(78)-FdeR at <i>xpr2</i> site.
pΔxpr2loxP-OTEF(136)-LEU2-OGAPDH(88)-FdeR	Integrating OTEF(136)-LEU2-OGAPDH(88)-FdeR at <i>xpr2</i> site.

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### Supplementary Table 3 Primers used in this paper

Primer	Sequence (5'-3')
leu2-up F	ATCCCTAAATTTGATGAAAGCCTAGGGCATAAAATGTGGAGAAGAAATC
leu2-up R	CCAACCCGGTCTCTGTCGTCTGTGGATGTGTGTGGTTGTATG
leu2-down F	AGCTTTACCGCAGCAGATCCGTCGTTTCTACGACGCATTGATGG
leu2-down R	CACTATTGGCCTATGCGGCCGCTGGCACTGAGCTCGTCTAACGG
xpr2-up F	ATCCCTAAATTTGATGAAAGCCTAGGGACGAGGACTCGTCCAACGG
xpr2-up R	CCAACCCGGTCTCTGTCGTACCCGAGAAGATCAGCGTTCTGTG
xpr2-down F	AGCTTTACCGCAGCAGATCCGACGCCAACACCAAGCTGGT
xpr2-down R	CACTATTGGCCTATGCGGCCGCGGACTCAGTAATAAGAGCCTCG
Nluc F	ACCAGCACTTTTTGCAGTACTAACCGCAGGTCTTCACACTCGAAGATTTCCG
Nluc R	CATAGCACGCGTGTAGATACTTACGCCAGAATGCGTTCGCAC
LEU2 F	ACCAGCACTTTTTGCAGTACTAACCGCAGGAACCCGAAACTAAGAAGACCAAGA
LEU2 R	CATAGCACGCGTGTAGATACTTATACACTAGCGGACCCTGCCGGT
pPOX2(1591) F	CTAAATTTGATGAAAGCCTAGGGACGACGATATTCCGGTCCCGAAACCC
pPOX2(1591) R	CTGCCTCTGAAACTCACCATTCTAGAGGCGTCGTTGCTTGTGTG
A3 F	CGAATGGTACGATTCCGCCACAATTGGACATGTTTGTTCGATC
A1R1 R	GATCGGAAAAACAAACATGTCCAATTGTGGCGGAATCGTACCATTCCG
A1R1_R1	GGTTTCGGGACCGGAATATCTGGCGGAATCGTACCATTCCG
A1R1_F2	GATATTCCGGTCCCGAAACCC
A1R1_R2	TGGCGGAATCGTACCATTCCG
gRNA F	TCCCTAAATTTGATGAAAGCCTAGGGACGACGATATTCCGGTCC
gRNA R	CCTTTTATCAGACATAGTCGACTCCTCCGTTATTGTCTCGCTAGC
gRNA_FAS1-1 F	GACGAAACGAGTAAGCTCGTCACTACCAAATCAAATAATGGTTTTAGAGCTAGAAATAG
gRNA_FAS1-1 R	ACGAGCTTACTCGTTTCGTCTCACGGACTCATCAGACTCACCTGCGGTTAGTACTGCAA
gRNA_FAS1-2 F	GACGAAACGAGTAAGCTCGTCTTTTGCTACAGGAAACAGCGGTTTTAGAGCTAGAAATAG
gRNA_FAS1-2 R	ACGAGCTTACTCGTTTCGTCTCACGGACTCATCAGTTTTGCCTGCGGTTAGTACTGCAA
gRNA_FAS1-3 F	GACGAAACGAGTAAGCTCGTCCGGTGCAGTTGATGTACAGAGTTTTAGAGCTAGAAATAG
gRNA_FAS1-3 R	ACGAGCTTACTCGTTTCGTCTCACGGACTCATCAGGGTGCAGTTCGCGGTTAGTACTGCAA
gRNA_FAS2-1 F	GACGAAACGAGTAAGCTCGTCAAACCTGAGCCACAAATCAGGTTTTAGAGCTAGAAATAG
gRNA_FAS2-1 R	ACGAGCTTACTCGTTTCGTCTCACGGACTCATCAGAAACCTCTGCGGTTAGTACTGCAA
gRNA_FAS2-2 F	GACGAAACGAGTAAGCTCGTCCGGCATCACCACCAGCAGCGGTTTTAGAGCTAGAAATAG
gRNA_FAS2-2 R	ACGAGCTTACTCGTTTCGTCTCACGGACTCATCAGGGGCATCTGCGGTTAGTACTGCAA

gRNA_FAS2-3 F	GACGAAACGAGTAAGCTCGTCGAAGGGGACAACAATCAGAGGTTTTAGAGCTAGAAATAG
gRNA_FAS2-3 R	ACGAGCTTACTCGTTTCGTCTCACGGACTCATCAGGAAGGGCTGCGGTTAGTACTGCAA
gRNA_FabD-1 F	GACGAAACGAGTAAGCTCGTCGCAGCATTCTCCCCGGACAGTTTTAGAGCTAGAAATAG
gRNA_FabD-1 R	ACGAGCTTACTCGTTTCGTCTCACGGACTCATCAGGCAGCACTGCGGTTAGTACTGCAA
gRNA_FabD-2 F	GACGAAACGAGTAAGCTCGTCGAGGCGTTGGATAACACCTGGTTTTAGAGCTAGAAATAG
gRNA_FabD-2 R	ACGAGCTTACTCGTTTCGTCTCACGGACTCATCAGGAGGCGCTGCGGTTAGTACTGCAA
gRNA_FabD-3 F	GACGAAACGAGTAAGCTCGTCCACATGATATACACAAAGCAGTTTTAGAGCTAGAAATAG
gRNA_FabD-3 R	ACGAGCTTACTCGTTTCGTCTCACGGACTCATCAGCACATGCTGCGGTTAGTACTGCAA
TEF_F2	GGGTATAAAAGACCACCGTCCCC
26srDNA2s F	ATCCCTAAATTTGATGAAAGCCTAGGCAGACACTGCGTCGCTCCGTCC
XPR2_Seq	GGTGTTGGACTCAGTAATAAGAGCC
leu2_Inte F	GTGTGCACTCCAACTTTTACAC
xpr2_Inte F	AGCCGTGTTTCGTGACGCAATC
leu2_Inte R	GATCATGCACACATAAGGTCC
FAS1 <sub>rt</sub> F <sup>1</sup>	GTCTCTGTATGGTCTGTGTCTTG
FAS1 <sub>rt</sub> R	GAGTGAAAGGAGAGGTGATG
FAS2 <sub>rt</sub> F	CACTCTCCCTTTTCTCCACATC
FAS2 <sub>rt</sub> R	AGCAACCTCAAGACCATCG
FabD <sub>rt</sub> F	GCATCTCAAAGCCTCCAAAC
FabD <sub>rt</sub> R	TGAAATCGGGACGGATCTTG
ACT1 <sub>rt</sub> F	GGTATCGTTCTTGACTCTGGTG
ACT1 <sub>rt</sub> R	AGGTAGTCGGTAAGATCTCGG

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1. The subscripts “<sub>rt</sub>” refers to primers used for qRT-PCR.

## References

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2. Wong, L.; Engel, J.; Jin, E.; Holdridge, B.; Xu, P., YaliBricks, a versatile genetic toolkit for streamlined and rapid pathway engineering in *Yarrowia lipolytica*. *Metab. Eng. Commun.* **2017**, *5*, 68-77.
3. Lv, Y.; Edwards, H.; Zhou, J.; Xu, P., Combining 26s rDNA and the Cre-loxP system for iterative gene integration and efficient marker curation in *Yarrowia lipolytica*. *ACS Synth. Biol.* **2019**.