

Supplementary Information

Metabolic Engineering and Genome-Wide Adaptive Evolution for Efficient Reduction of Glycerol in Industrial *Saccharomyces cerevisiae*

Na Xu ¹, Hui Chen ¹, Yan Zhang ¹, Yuxian Yang ¹, Yasi Wang ¹, Bei Liao ¹, Nan Peng ² and Xiaosong Gu ^{1,*}

¹ The Hubei Provincial Key Laboratory of Yeast Function, Yichang 443003, China

² National Key Laboratory of Agricultural Microbiology, Hubei Hongshan Laboratory, College of Life Science and Technology, Huazhong Agricultural University, Wuhan 430070, China

* Corresponding author. E-mail: xiaosonggu@yeah.net (X.G.); Tel.: +86-13007123941 (X.G.)

Supplementary Figures

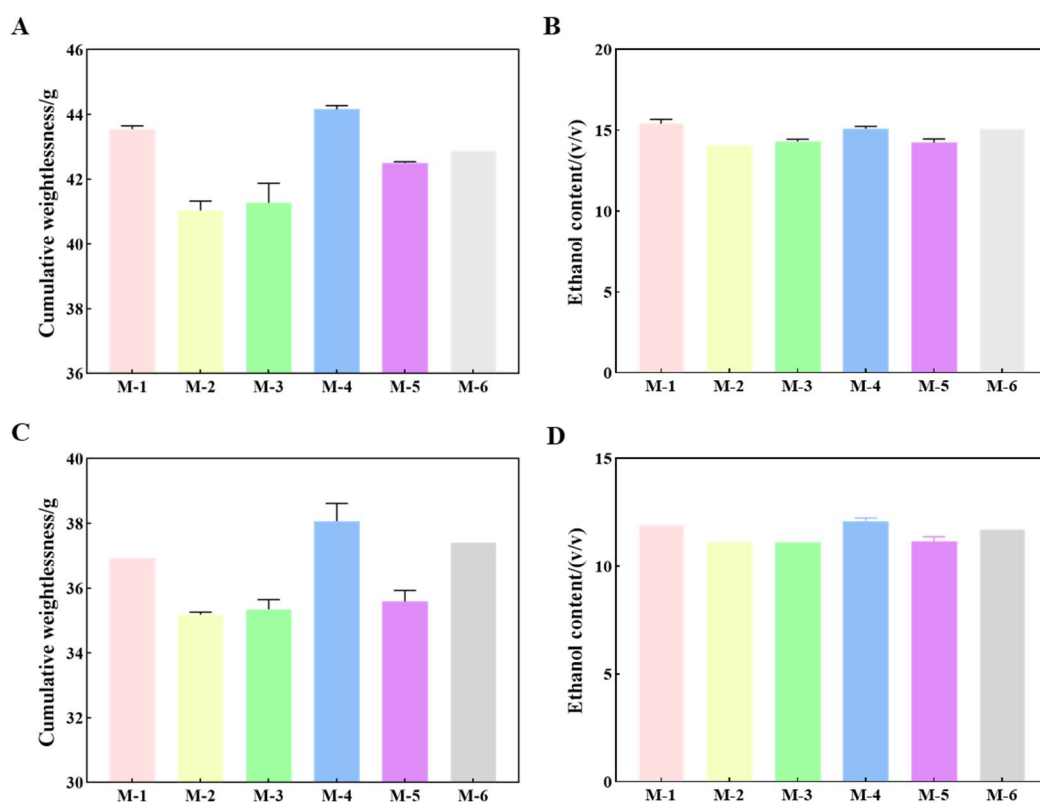


Figure S1. Cumulative weightlessness and ethanol yield in engineered strains. (A) Difference in cumulative weightlessness at 36°C. (B) Difference in ethanol yield at 36°C. (C) Difference in cumulative weightlessness at 39°C. (D) Difference in ethanol yield at 39°C.

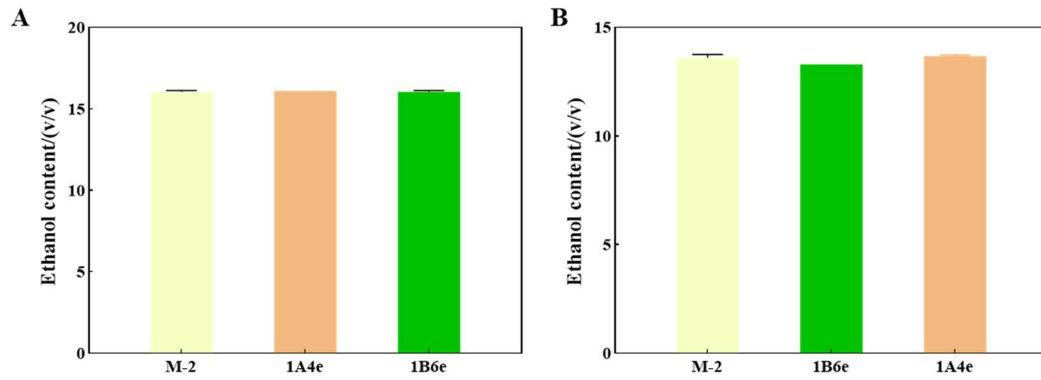


Figure S2. The ethanol yield of the evolved strains and the parental strain. (A) Ethanol yield of the evolved strains and the parental strain fermented at 35°C. (B) Ethanol yield of the evolved strains and the parental strain fermented at 38°C.

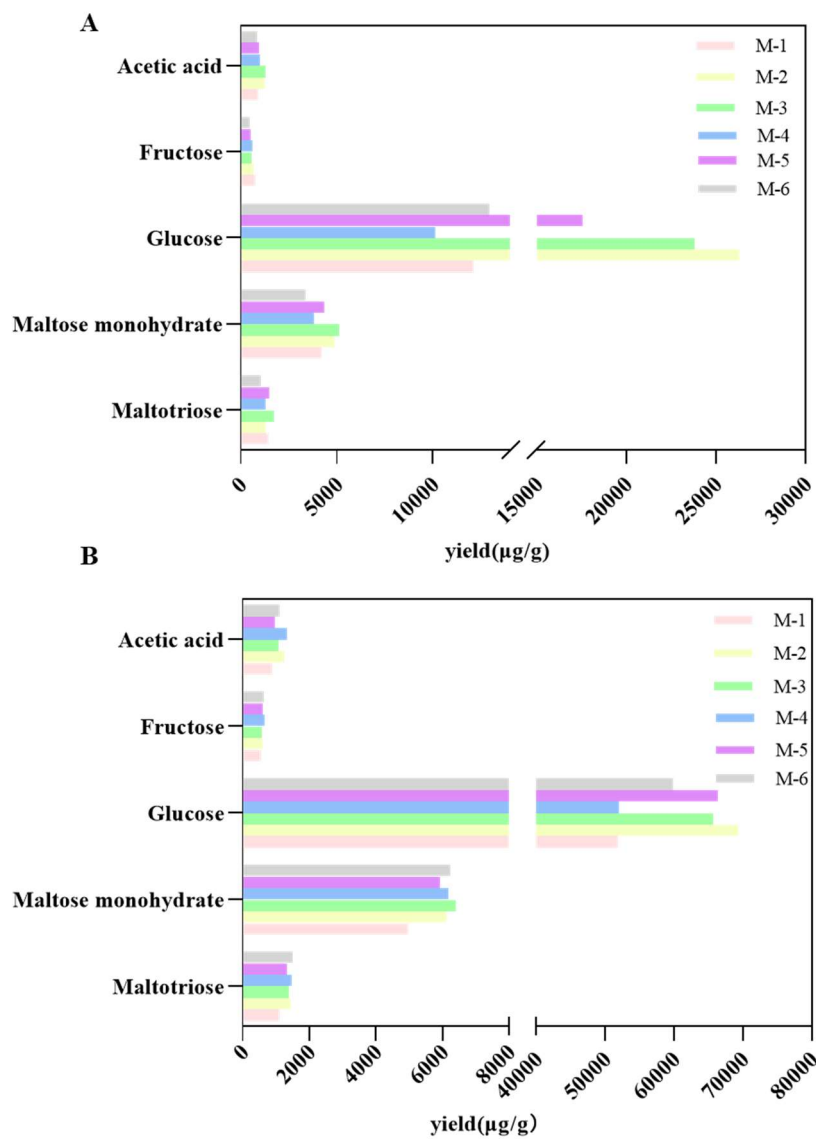


Figure S3. Residual sugar and acetic acid in fermentation of engineered strains. (A) Residual sugar and acetic acid in fermentation of engineered strains at 36°C. (B) Residual sugar and acetic acid in fermentation of engineered strains at 39°C.

Supplementary Tables

Supplementary Table S1. A list of plasmids constructed in this study.

Name	Description	Source
PSCM-H-gRNA-site5	<i>S. cerevisiae</i> M, HygB, <i>SNR52p</i> -site5-gRNA	
PSCM-H-gRNA-site10	<i>S. cerevisiae</i> M, HygB, <i>SNR52p</i> -site10-gRNA	
PSCM-G-gRNA- site19	<i>S. cerevisiae</i> M, G418, <i>SNR52p</i> -site19-gRNA ^a	
PSCM-G-gRNA- site21	<i>S. cerevisiae</i> M, G418, <i>SNR52p</i> -site21-gRNA	
PSCM-H-gRNA- <i>FPSp</i>	<i>S. cerevisiae</i> M, HygB, <i>SNR52p</i> - <i>FPSp</i> -gRNA	
PSCM-H-gRNA- <i>ΔADH2</i>	<i>S. cerevisiae</i> M, HygB, <i>SNR52p</i> - <i>ΔADH2</i> -gRNA	
PSCM-H-gRNA- <i>Δade2</i>	<i>S. cerevisiae</i> M, HygB, <i>SNR52p</i> - <i>Δade2</i> -gRNA	
pHCas9	2μ; <i>AmpR</i> ; <i>TEF1p</i> - <i>Cas9</i> - <i>CYC1t</i>	This study
PUC- <i>TEF1p</i> - <i>Rim15</i> - <i>ADH1t</i>	2μ; <i>AmpR</i> ; <i>TEF1p</i> - <i>Rim15</i> - <i>ADH1t</i>	This study
PUC- <i>PRK</i> - <i>Cbbm</i>	2μ; <i>AmpR</i> ; <i>PGKp</i> - <i>PRK</i> - <i>ADH1t</i> - <i>CIT2p</i> - <i>Cbbm</i> - <i>CYC1t</i>	This study
PUC- <i>EroES</i> - <i>EroEL</i>	2μ; <i>AmpR</i> ; <i>GAP1p</i> - <i>GroES</i> - <i>CYC1t</i> - <i>TEF1p</i> - <i>GroEL</i> - <i>ADH1t</i>	This study
pSpdCas9- <i>poll5M</i> -gRNA- <i>mre11</i>	2μ; <i>KanR</i> ; <i>TEFp</i> - <i>dCas9</i> - <i>poll5M</i> , tRNA ^{phe} -HDV- <i>SNR52p</i> - <i>mre11</i> -gRNA	This study
pSpdCas9- <i>poll5M</i> -gRNA- <i>msh2</i>	2μ; <i>KanR</i> ; <i>TEFp</i> - <i>dCas9</i> - <i>poll5M</i> , tRNA ^{phe} -HDV- <i>SNR52p</i> - <i>msh2</i> -gRNA	This study

Note: ^a HygB, Hygromycin B; G418, Geneticin; Amp, Ampicillin; Kan, Kanamycin.

Supplementary Table S2. A list of primers used in this study.

Oligo	Sequences (5'-3')
Site5-gRNA-F	gatcgctaataaagaggtaacggt
Site5-gRNA-R	aaacaccgttacctctttattagc
Site10-gRNA-F	gatccatgagcagccactgtatcg
Site10-gRNA-R	aaaccgatacagtggtgctcatg
Site19-gRNA-F	gatcgttgaaatataagtaaccct
Site19-gRNA-R	aaacaggggtacttatattcaac
Site21-gRNA-F	gatacaggactaaataacaccagg
Site21-gRNA-R	aaaccctgggtttatttagtcctg
<i>ΔADH2</i> -gRNA-F	gatctggcataccgaccaaaccgg
<i>ΔADH2</i> -gRNA-R	aaaccggtttggtcgctatcca
<i>FPSp</i> -gRNA-F	gatccataaccgttaggaaggtacg
<i>FPSp</i> -gRNA-R	aaaccgtaccttctacgggtatg
<i>Δade2</i> -gRNA-F	gatcgtataatgtccagagttgtg
<i>Δade2</i> -gRNA-R	aaaccacaactctggacattatac
HR-UP-site5	ACATTAGATTGGAATTAGAGCTTAAGTGGTACAACTAGGAtaagcgtccttctgtggttaga
HR-DW-site5	TATTCGGCTTAAAGTGCTTAACATCGGTGCCACACAATAGcaaaattaaagccttcgagcgtc
HR-UP-site10	CCTCTATGTGACGCTGTGTATTCTTTGTTGTAGTTATGCTgagcgacctcatgctatac
HR-DW-site10	TGTACGCTATACATTTACGTGCTGAGCTCCTAGGAAAGCTgcaaattaaagccttcgag
HR-UP-site19	TGAAAATTGTGCCGATATTCAAGACTAAGAGATGTACAaccacacaccatagcttcaa
HR-DW-site19	AAATCGACATGTTAATGATCTTACGACAGAGTAGTTTATGgagcgacctcatgctatac
HR-UP-site21	ATAGCACGAGAAAAAACTCGTTCAACTAAGTCTAGACACAgagcgacctcatgctatac
HR-DW-site21	AGAGCTGTTTGAGAGCTCGTAGGCTTTGTTGTTAAGAGAGcaaaattaaagccttcgag

Supplementary Table S3. A list of heterologous genes coding sequences used in this study.

Gene	Coding sequences (5'-3')
<i>PRK</i>	ATGGCTGTTTGTACTGTTTATACTATTCTACTACAACCTCATTGGGTTCTTCTTTCAATCAA AACAAATAAGCAAGTTTCTTCAATTATAAAAAGATCTTCATCTTCTAATAACACATTGTTTAC AACTAGACCATCTTATGTTATTACTTGTCTCAACAGCAAACCTATTGTTATTGGTTTGGCTG CTGATTCTGGTTGTGGTAAATCTACTTTTATGAGAAGATTGACTTCTGTTTTTGGTGGTGCT GCTGAACCACCAAAAGGTGGTAATCCTGATTCTAATACTTTGATTTCTGATACAACAACCTG TTATCTGTTTGGACGATTTTCATTCTTTGGACAGAAAATGGTAGAAAAGTTGAAAAAGTTAC TGCTTTGGATCCAAAAGCTAATGATTTTGATTTGATGTATGAACAAGTTAAGGCTTTGAAA GAAGGTAAAGCTGTTGATAAACCAATTATAATCATGTTTCTGGTTTGTGGATCCACCTGA ATTGATTCAACCACCAAAAATTTGGTTATTGAAGGTTTGCATCCAATGTACGATGCAAGA GTTAGAGAATTGTTGGACTTTTCAATCTACTTGGACATTTCTAATGAGGTCAAATTTGCTTG GAAAATTCAAAGAGATATGAAAGAAAGAGGTCACCTTTTGAATCTATTAAGGCTTCTATT GAATCTAGAAAACCTGATTTTGATGCTTATATTGATCCACAAAAACAACATGCTGATGTTGT TATTGAAGTTTGGCAACTGAATTGATTCCTGATGACGATGAAGGTAAAGTCTTAAGAGTC AGAATGATCCAAAAAGAAGGTGTCAAATCTTCAATCCTGTTTATTTGTTTGATGAAGGTT CTACTATTTCTTGGATTCCATGTGGTAGAAAATTGACTTGTCTTATCCTGGTATTAATTTT CTTATGGTCTGATACTTTTATGGTAATGAAGTTACTGTTGTTGAAATGGATGGAATGTTT GATAGGTTGGATGAATTGACTACGTTGAATCACATTTATCAAACCTGTCAACTAAATTTTA TGGTGAAGTCACACAACAATGTTGAAGCATCAAACCTTTCTGGTTCTAACAATGGTAC TGGTTTCTTTCAAACCTATTATTGGTTTAAAGATCAGAGATTGTTTGAACAATTGGTTGCTT CTAGATCTACTGCTACTGCTACTGCTGCTAAAGCTTAA
<i>Cbbm</i>	ATGGATCAATCTGCTAGATATGCTGATTTGTCTTTGAAAAGAAGAAGATTGATTAAAGGTG GTAGACATATTTTGGTTGCTTATAAAATGAAACCAAAATCTGGTTATGGTTATTTGGAAGCT GCAGCTCATTTTGTCTGCTGAATCTTCTACTGGTACTAATGTTGAAGTTTCTACTACTGATGA TTTTACTAAAGGTGTTGATGCTTTGGTTTATTATATTGATGAGGCTTCTGAAGATATGAGAAT TGCTTATCCATTGGAATTGTTTGATAGAAATGTTACTGATGGTAGATTTATGTTGGTTTCTTT TTTGACTTTGGCTATTGGTAATAATCAAGGTATGGGTGATATTGAACATGCTAAAATGATTG ATTTTATGTTCTGAAAGATGATTCAAATGTTTGATGGTCTGCTACTGATATTCTAATT TGTGGAGAATTTGGGTAGACCTGTTGTTAATGGTGGTTATATTGCTGGTACTATTATTA CCAAAATTGGGTTTGGACCTGAACCATTTGCTAAAGCTGCTTATCAATTTTGGTTGGGTG GTGATTTTATTAATAATGATGAACCACAAGGTAATCAAGTTTTTGTCCATTGAAAAAAGT TTTGCCATTGGTTTATGATGCTATGAAAAGAGCTCAAGATGATACTGGTCAAGTAAATTTG TTTCTATGAATATTACTGCTGATGATCATTATGAAATGTGTGCTAGAGCTGATTATGCTTTGG AAGTTTTTGGTCTGATGCTGATAAATTAGCTTTTTTGGTTGATGGTTATGTTGGTGGTCT GGTATGGTTACTACTGCTAGAAAGACAATATCCTGGTCAATATTGCATTATCATAGAGCTGG TCATGGTGCTGTTACTTCTCCATCTGCTAAAAGAGGTTATACTGCTTTTGTTTTGGCTAAAA TGCTAGATTGCAAGGTGCTTCTGGTATTATGTTGGTACTATGGGTTATGGTAAAATGGAA GGTGAAGGTGATGATAAAATTATTGCTTATATGATTGAAAGAGATGAATGTCAAGGTCCTG TTTATTTTCAAAAATGGTATGGTATGAAACCACTACTCCAATTATTTCTGGTGGTATGAAT GCTTTGAGATTGCCTGGTTTTTTTGAATAATTGGGTGATGGTAATGTTATTAATACTGCTGG TGGAGGTTCTTATGGTCATATTGATTCTCCTGCTGCTGGTGCTATTTCTTTGAGACAATCTTA TGAATGTTGGAACAAGGTGCTGATCCAATTGAATTTGCTAAAGAACATAAAGAATTTGCT AGAGCTTTTGAATCTTTTCCAAAAGACGCTGATAAGTTGTTTCTGTTGGAGAGAAAAAT TGGGTGTTCAATCTTAA
<i>GroE S</i>	ATGAATATTAGACCATTCATGATAGAGTTATTGTTAAAAGAAAAGAAGTTGAACTAAAT CTGCTGGTGGTATTGTTTTGACTGGTTCTGCTGCAGCTAAATCTACTAGAGGTGAAGTTTT GGCTGTTGGTAATGGTAGAATTTTGGAAAATGGTGAAGTTAAACCATTTGGATGTTAAAGTC GGTGATATTGTTATTTTCAATGATGGTTATGGTGTCAAATCTGAGAAAATTGATAATGAAGA AGTTTTGATCATGTCTGAATCTGATATTTGGCTATTGTTGAAGCTTAA
<i>GroE L</i>	ATGGCTGCTAAAGATGTTAAATTTGGTAATGATGCTAGAGTTAAATGTTGAGAGGTGTTA ATGTTTTGGCTGATGCTGTAAAGTTACTTTGGGTCCAAAAGGTAGAAATGTTGTTTTGGA TAAATCTTTTGGTGCTCCAACCTATTACTAAAGATGGTGTCTTCTGTTGCTAGAGAAATTGAAT TGGAAGATAAATTTGAAAATATGGGTGCTCAAATGGTTAAAGAAGTTGCTTCTAAAGCTAA TGATGCTGTGGTGATGGTACTACAAGTCTACTGTTTTGGCTCAAGCTATTACTGAAG GTTTGAAAGCTGTTGCTGCTGGTATGAATCCAATGGATTGAAAAGAGGTATTGATAAAGC TGTTACTGCAGCTGTTGAAGAATTGAAGGCTTTGTCTGTTCCATGTTCTGATTCTAAAGCT ATTGCTCAAGTTGGTACTATTTCTGCTAATTCTGATGAAACTGTTGGTAAATTGATTGCTGA AGCTATGGATAAAGTTGGTAAAGAAGGTGTTATTACTGTTGAAGATGGTACTGGTTTGCAA GATGAATTGGATGTTGTTGAAGGTATGCAATTTGATAGAGGTTATTTGTCTCCATATTTTATT AATAAACCTGAACTGGTGCTGTTGAGTTGGAATCTCCATTATCTTGTGGCTGATAAAA AAATCTCAATATTAGAGAAATGTTGCCTGTTTTGGAAGCTGTTGCTAAAGCTGTTAAACC ATTGTTGATTATTGCTGAAGATGTTGAAGGTGAGGCTTTGGCTACTTTGGTTGTTAATACTA TGAGAGGTATTGTTAAAGTTGCTGCTGTTAAAGCTCCTGGTTTTGGTGATAGAAGAAAAGC TATGTTGCAAGATATTGCTACTTTGACTGGTGGTACTGTTATTTCTGAGGAGATTGGTATGG

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