

Adaptive Evolution Relieves Nitrogen Catabolite Repression and Decreases Urea Accumulation in Cultures of the Chinese Rice Wine Yeast Strain *Saccharomyces cerevisiae* XZ-11

Weiping Zhang^{†,‡}, Yan Cheng[†], Yudong Li[§], Guocheng Du[†], Guangfa Xie[¶], Huijun Zou[¶], Jingwen Zhou^{*,†,‡}, and Jian Chen^{*,†}

[†] Key Laboratory of Industrial Biotechnology, Ministry of Education and School of Biotechnology, Jiangnan University, 1800 Lihu Road, Wuxi, Jiangsu 214122, China;

[‡] National Engineering Laboratory for Cereal Fermentation Technology, Jiangnan University, 1800 Lihu Road, Wuxi, Jiangsu 214122, China

[§] Department of Bioengineering, School of Food Sciences and Biotechnology, Zhejiang Gongshang University, Hangzhou 310018, China

[¶] College of Shaoxing Rice Wine, Zhejiang Shuren University, Shaoxing 312028, China

^{*} Zhejiang Guyuelongshan Shaoxing Wine Company, 13 Yangjiang Road, Shaoxing, Zhejiang, China.

* Corresponding authors.

Jian Chen, Jingwen Zhou

Mailing address: School of Biotechnology, Jiangnan University, 1800 Lihu Road, Wuxi, Jiangsu 214122, China.

Phone: +86-510-85918312, Fax: +86-510-85918309

E-mail: zhoujw1982@jiangnan.edu.cn, jchen@jiangnan.edu.cn.

SUPPORTING INFORMATION

Table S1. Primers used in the study

Primers	Sequences (5'-3')
GLN3-F	CTAATAACTTAATGCGTCACAA
GLN3-R	GGTAATATCGGAACAAACAGA
GAT1-F	CTGATATGAATATGACTATGAAC
GAT1-R	TCCAGAACATGATGATCTATTG
ARG1-F	ACTCATTGTCACACCAA
ARG1-R	GATCTGATGTACTCACACTCT
ARG3-F	GGTGAAGAATTGCGAACAG
ARG3-R	CAGGCCACAGAGACAAGTTC
ARG4-F	GCCTCAGAAGAAGAAATGC
ARG4-R	TCATCAAGAACATCCTGTCAGAT
CAR1-F	GAAACAAACGGTGAAGGT
CAR1-R	TGTAGCAGGAATGTATAATGG
DAL80-F	GGAGTTGGTGACAGACCTATAC
DAL80-R	TAGTTCCACTTCCAGTTGCTT
GZF3-F	GCACATAACTGGACAACTT
GZF3-R	GACCGCTTGATTCTATATTACA
TOR1-F	CTTGTGGTAGGTTGTC
TOR1-R	TGTTAGCAGTTCCAGTAGTAT
TOR2-F	TTCTTGCTGTTGATACTCTG
TOR2-R	ATAGTTGCCAGTCTTGA
URE2-F	CTGGTGATGGAATTAGACA
URE2-R	GAAACGACTTGTGACATT
DUR1,2-F	AAGAAGTAATGGTGGTGAAG
DUR1,2-R	AAGTAATAAACTGGCTCATCA
DUR3-F	AACGATAAGGAACAAAGAAGAA
DUR3-R	CCAAGGTTAGGCTCAATCAAT
ACT1-F	TGGATTCTGGTATGTTCT
ACT1-R	GTCATATAGGAGGTTATGG

Table S2 Genetic variants in the genome of adaptive evolution improved strains 4B, 7H, and 10G

(See Excel file Table S2.xlsx)

Table S3 Fast evolved genes in three evolved strains

(See Excel file Table S3.xlsx)

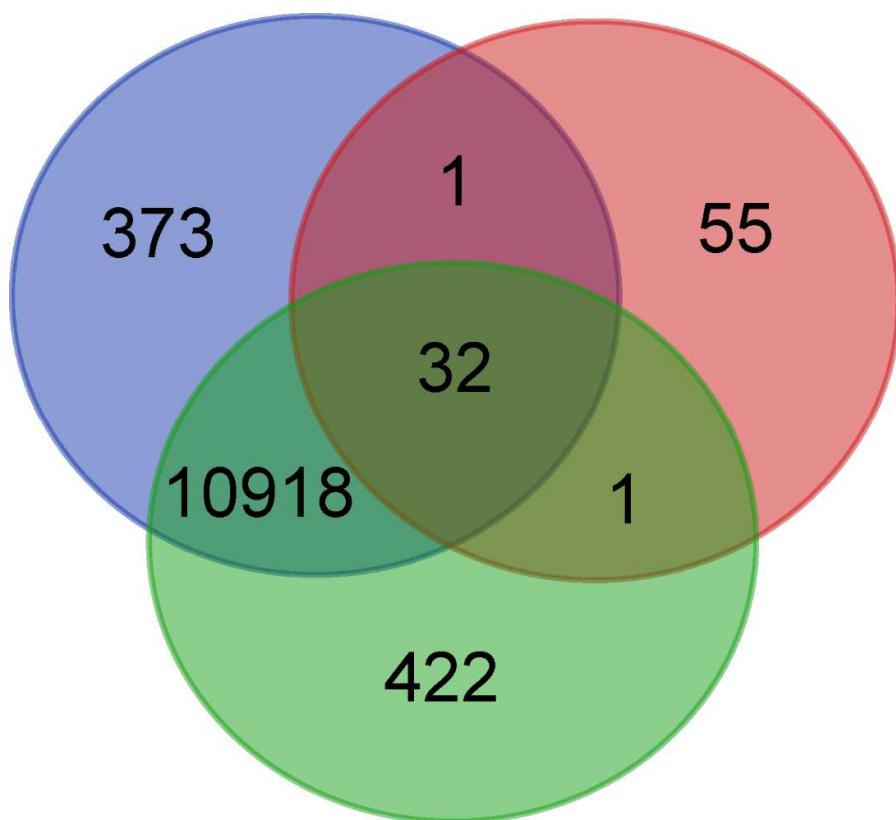


Figure S1 Comparative analysis of the variants among three evolved strains.

The red circle stands for strain 4H; the blue circle stands for strain 7H, and the green circle stands for strain 10G.