

Supplementary materials

Cofactor engineering for efficient production of α -farnesene by rational modification of NADPH and ATP regeneration pathway in *Pichia pastoris*

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Supplementary_data_1:

Plasmid construction process

Ligation of fragments and vectors using the ClonExpress II One Step Cloning Kit (Vazyme Biotech, Nanjing, China). The fragments of homologous recombination include homologous fragments of 15-20 bp. The constructed plasmid was amplified using *E. coli* JM109. sgRNA design using <http://chopchop.cbu.uib.no/> website.

Construction of strains

The gene integration and insertion of *Pichia pastoris* engineering strains were carried out by electroporation, and Cre/LoxP system was used for the recovery of screening resistance, in which the gene expressing Cre enzyme used an episomal plasmid, and antibiotic-free culture was used to eliminate the plasmid. The homologous fragment used for knockout and insertion in the CRISPR/CAS9 system is about 1000 bp.

Supplementary_data_2:

Table S1 Primers used for expression vectors in this study

Primers named	Primers (5'-3')
PGAP-Z-F	TATGCTATACGAACGGTAATCATGTAATTAGTTATGTCACG
PGAP-Z-R	GGATCCGCACAAACGAAGGTCTCACTTAATCTTCTGTACTC
Lox71-F	TACCGTTCGTATAATGTATGCTATACGAAGTTATGGATCCCCACACACCA TAG
Lox66-R	TACCGTTCGTATAGCATACATTATACGAAGTTATATTAAGGGTTCTCGAGA GCTC
HIS4-F	CTTCGTTTGTGCGGATCCTTAAATAAGTCCCAGTTTCTC
HIS4-R	CATACATTATACGAACGGTAATGACATTTCCCTTGCTACC
PIS1-F	GTCATGCATGAGATCAGATCGTAACGAGGCTAAAAGTTTTTGC
PIS1-R	GAATTCCTCGTTTTCGAATGCAGGTGGACTATCTAGAGACAAG
PGPM1-F	TCATGCATGAGATCAGATC AAGGTTTATCTGACTGTTGCGCAC
PGPM1-R	GAATTCCTCGTTTTCGATGTTTGTGTTGTGTAATTGAAAGTT
PMET3-F	TCATGCATGAGATCAGATCAATTCAGGCAACAGGACC
PMET3-R	GAATTCCTCGTTTTCGATTCTTTCTGAGTTGGTTTCC
PPGK1-F	TCATGCATGAGATCAGATCTACCCAGCCGATCACGCCTGC
PPGK1-R	GAATTCCTCGTTTTCGAGGGCCCAAGTTGGTACCCAGCC
ZWF1-F	TCGAAACGAGGAATTCATGACCGATACGAAAGCCG
ZWF1-R	GGCCGCCGCGGCTCGAGTTACATCTTGTGCAGCACATC
sol3-F	TCGAAACGAGGAATTCATGGTACAAATCTATTCCT
sol3-R	CGGCCGCCGCGGCTCGAGTCAGTATTTCGAAGTAGAAAC
Gnd2-F	TCGAAACGAGGAATTCATGGTTGAAGCAACAGGAG

Gnd2-R	GGCCGCCGCGGCTCGAGTTAAGCATCGTAGGTACTGG
Rpel-F	TCGAAACGAGGAATTCATGGTTAAACCTATTATTGC
Rpel-R	GGCCGCCGCGGCTCGAGTCATAAGTTGGGTTTTGGTTTG
cPOS5-F	TCGAAACGAGGAATTCATGTTTGTGAGGGTTAAATTG
cPOS5-R	GGCCGCCGCGGCTCGAG TTAATCATTATCAGTCTGTCTC
APRT-F	TCGAAACGAGGAATTCATGTCAACCGCTTCTCTG
APRT-R	GGCCGCCGCGGCTCGAGTCATTCTGCTGCCCCGA
PGI-sg	TGTTCACTTCGTATCCAACG
GPD1-sg	GTTGACGGATATTATCAACG