Supporting information

Supplementary Table 1. Strains and plasmids

Strain	Relevant genotype	Source
E. coli		
DH5a	F- λ - endA1 glnV44 thi-1 recA1 relA1 gyrA96 deoR nupG Φ 80dlacZ Δ M15 Δ (lacZYA-argF) U169, hsdR17(rK-mK+)	TaKaRa (Dalian China)
S. cerevisiae		
BY4741	MATa his $3\Delta 1$ leu $2\Delta 0$ met $15\Delta 0$ ura $3\Delta 0$	ATCC
BY4742	MATα his3 Δ 1 leu2 Δ 0 lys2 Δ 0 ura3 Δ 0	ATCC
JX1	BY4741/pYX212-GGPPS-P2A-CARRP	This study
JX2	BY4741/pYX212-CARB-P2A-CARRP	This study
JX3	BY4741/pYX212-CARRP-P2A-CARB	This study
JX4	BY4741/pYX212-GGPPS-P2A-CARB-P2A-CARRP	This study
JX5	BY4741/pYX212-CARB-P2A-GGPPS-P2A-CARRP	This study
JX6	BY4741/pYX212-CARRP-P2A-GGPPS-P2A-CARB	This study
JX7	BY4741/pYX212-GGPPS-P2A-CARRP-P2A-CARB	This study
JX8	BY4741/pYX212-CARB-P2A-CARRP-P2A-GGPPS	This study
JX9	BY4741/pYX212-CARRP-P2A-CARB-P2A-GGPPS	This study
Plasmid	Description	Source
pUC18	Amp ^R	TaKaRa
pYX212	Expression vector, URA3 selection marker for S. cerevisiae, Amp ^R	Zhou <i>et al.</i> 2012
pJX1	pYX212-GGPPS-P2A-CARRP	This study
pJX2	pYX212-CARB-P2A-CARRP	This study
pJX3	pYX212-CARRP-P2A-CARB	This study
pJX4	pYX212-GGPPS-P2A-CARB-P2A-CARRP	This study
pJX5	pYX212-CARB-P2A-GGPPS-P2A-CARRP	This study
pJX6	pYX212-CARRP-P2A-CARB-P2A-GGPPS	This study
pJX7	pYX212-GGPPS-P2A-CARRP-P2A-CARB	This study
pJX8	pYX212-CARB-P2A-CARRP-P2A-GGPPS	This study
pJX9	pYX212-CARRP-P2A-CARB-P2A-GGPPS	This study
pJX11	pUC18-GGPPS-P2A-CARB-P2A-CARRP	This study
pJX12	pUC18-CARB-P2A-GGPPS-P2A-CARRP	This study
pJX13	pUC18-GGPPS-P2A-CARB-P2A-CARRP	This study

Supplementary Table 2. Primers used in this study

Primer name	Sequence $(5'-3')$			
ADH2-EcoRI-GGPPS-F	ctgacagctgccacgtctccgacagtcacagaattcatgctcacctcgtccaagtcgatc			
CCDDC D24 DamIII D	catggatccagggccagggttctcttcgacgtcgccagcctgcttgaggagcgagaagttggtag			
GGPPS-P2A-BamHI-K	cgcccgagccgtgatggtgatggtggtgttcgtt			
BamHI-CarB-F	cctggatccatgtcggatcagaagaagcacatc			
Corp D2A Van I D	catggtaccagggccagggttctcttcgacgtcgccagcctgcttgaggagcgagaagttggtag			
Сагв-Р2А-крш-к	cgcccgagccgtgatggtggtggtggtggatgcg			
KpnI-CarRP-F	cctggtaccatgtcgattttgacctacctcgag			
CarDD D2A UindIII D	cata agettagggccagggttctcttcgacgtcgccagcctgcttgaggagcgagaagttggtag			
Carkf-f2A-filliulli-K	cgcccgagccgtgatggtggtggtggtcgac			
ADH2-EcoRI-CarB-F	ctgacagctgccacgtctccgacagtcacagaattcatgtcggatcagaagaagcacatc			
DIA RamHI D	catggatccagggccagggttctcttcgacgtcgccagcctgcttgaggagcgagaagttggtag			
I ZA-Dalli II-K	cgcccgagccgtgatggtgatggtggg			
BamHI-GGPPS-F	cctggatccatgctcacctcgtccaagtcgatc			
P2A-KpnI-R	catggtaccagggccagggttctcttcgacgtc			
ADH2-EcoRI-CarRP-F	ctgacagctgccacgtctccgacagtcacagaattcatgtcgattttgacctacct			
KpnI-CarB-F	cctggtaccatgtcggatcagaagaagcacatc			
P2A-HindIII-Hyg-R	ctcgaccgacgtcgccgtgagctccggcataagcttagggccagg			
CarD D2A UindIII D	cata agett agg gc cag gg tt ct ct tc g acg tc g cc ag cc t g ct g ag g ag			
Cal D-F 2A-minum-K	cgcccgagccgtgatggtgatggtggtggatgcg			
GGPPS-dB-CarRP-F	ggacgtcctctccgtcaacgaa			
GGPPS-dB-CarRP-R	acaagtggaactcgaggtaggtcaaaatcgacatggatccagggccagggttctcttc			
CarB-dS-CarRP-F	cagaactcgaacgacatccgcatc			
CarRP-dS-CarB-F	gatcaaatcgttcttcgtcgac			
CarRP-dS-CarB-R	caatcaccacgatgtgcttcttctgatccgacatggatccagggccagggttctcttc			
CarRP1373-F	agatgggcttggtcctcc			
RP-CarB-pYX-R	taagcttgtgggccctaggatccatggctagtgatggtgatggtggtggatgcggatgtcgtt			
RP-GGPPS-pYX-R	taagettgtgggccctaggatccatggctagtgatggtgatggtgttcgttgacggagag			
CarB1279-F	cgagcacgaggaggtcaa			

Time (min)	Buffer A%	Buffer B%
0	90	10
10	83	17
20	75	25
25	75	25
27	50	50
35	30	70
40	30	70
42	25	75
45	90	10
55	90	10

Supplementary Table 3. Gradient method for carotenoid analysis

buffer A [methanol/H₂O (92:8, v/v) and 10 mM ammonium acetate], buffer B (methyl *tert*-butyl ether).

Referce

1. Zhou Y, Gao W, Rong Q, *et al.* 2012. Modular pathway engineering of diterpenoid synthases and the mevalonic acid pathway for miltiradiene production. *J Am Chem Soc* **134**: 3234-3241.