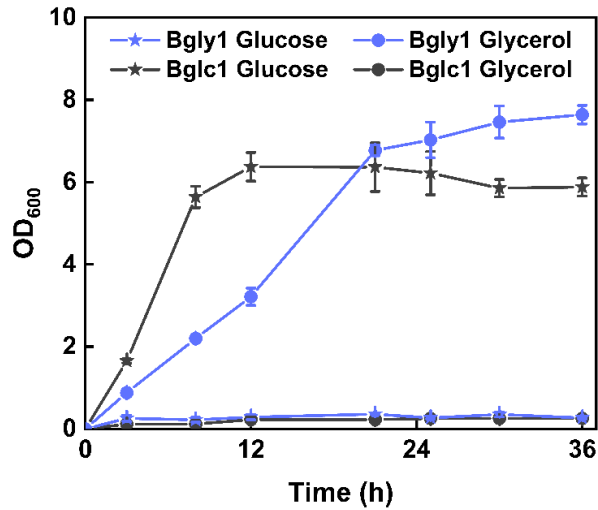
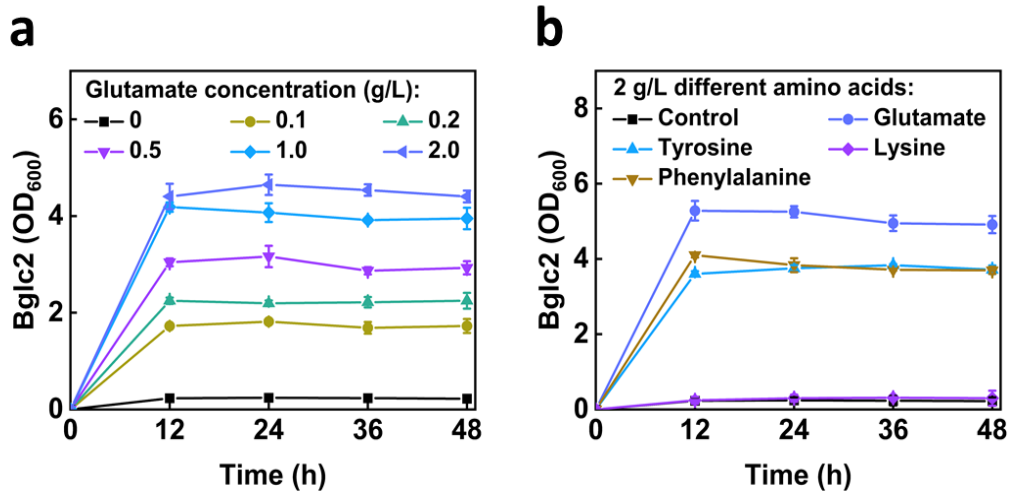


**Design of stable and self-regulated microbial consortia for chemical
synthesis**

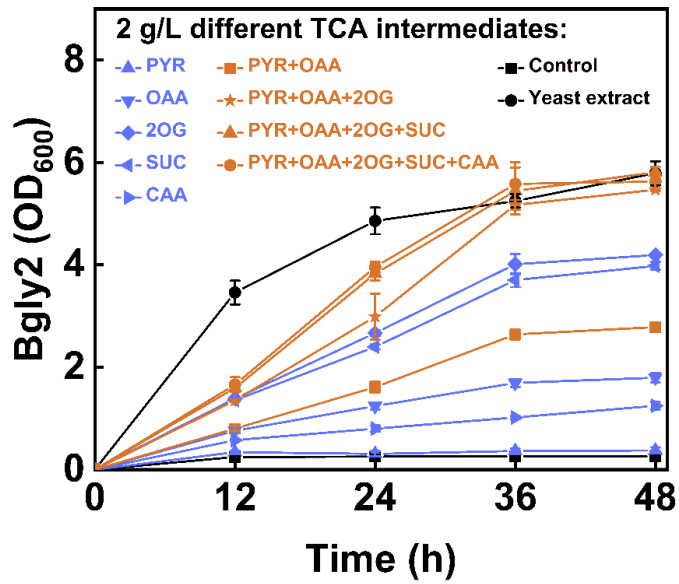
Li et. al



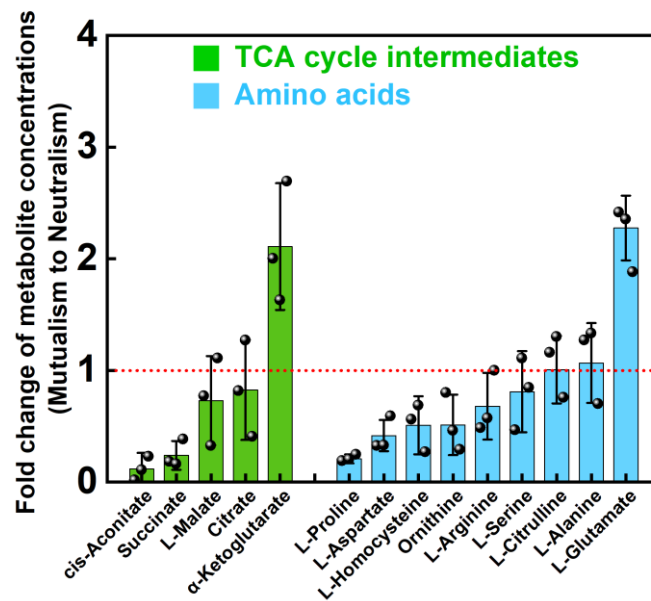
Supplementary Fig. 1. Growth of glycerol-utilizing strain Bgly1 and glucose-utilizing strain Bglc1 in glucose and glycerol media, respectively. Data shown are mean \pm s.d. (n = 3 independent experiments). Source data are provided as a Source Data file.



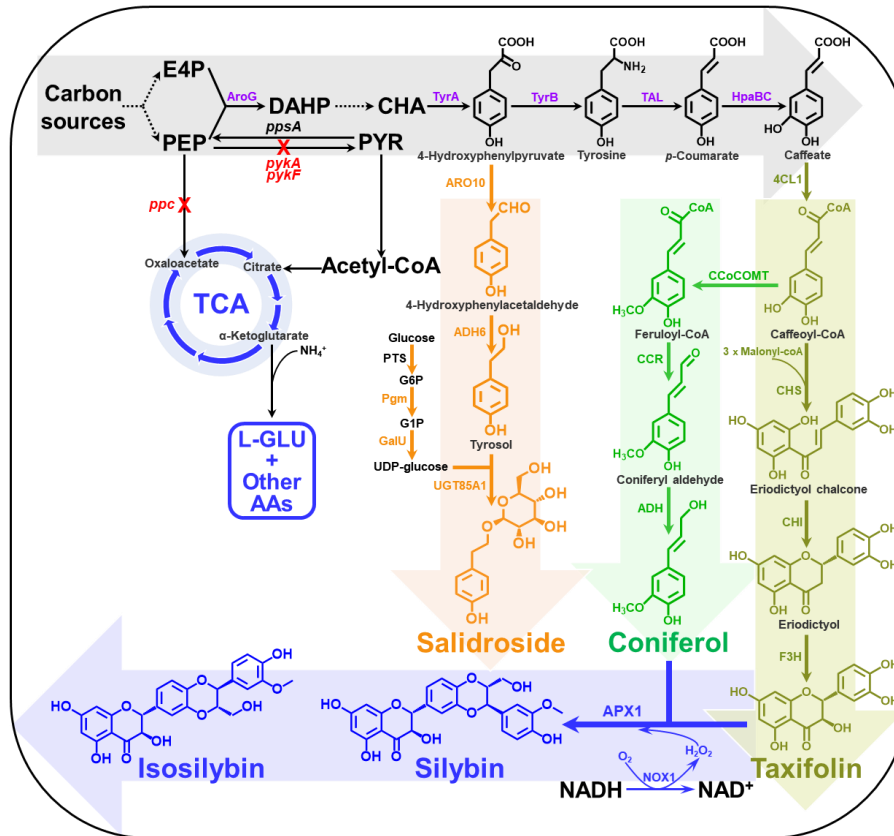
Supplementary Fig. 2. Growth of Bglc2 in culture media supplemented with different amino acids. a Growth with glutamate at different concentrations, **b** Growth with 2 g/L of each amino acid. Data shown are mean \pm s.d. (n = 3 independent experiments). Source data are provided as a Source Data file.



Supplementary Fig. 3. Growth curves of Bgly2 in culture media supplemented with different combinations of carboxylic acids at 2 g/L each. Data shown are mean \pm s.d. (n = 3 independent experiments). Source data are provided as a Source Data file.

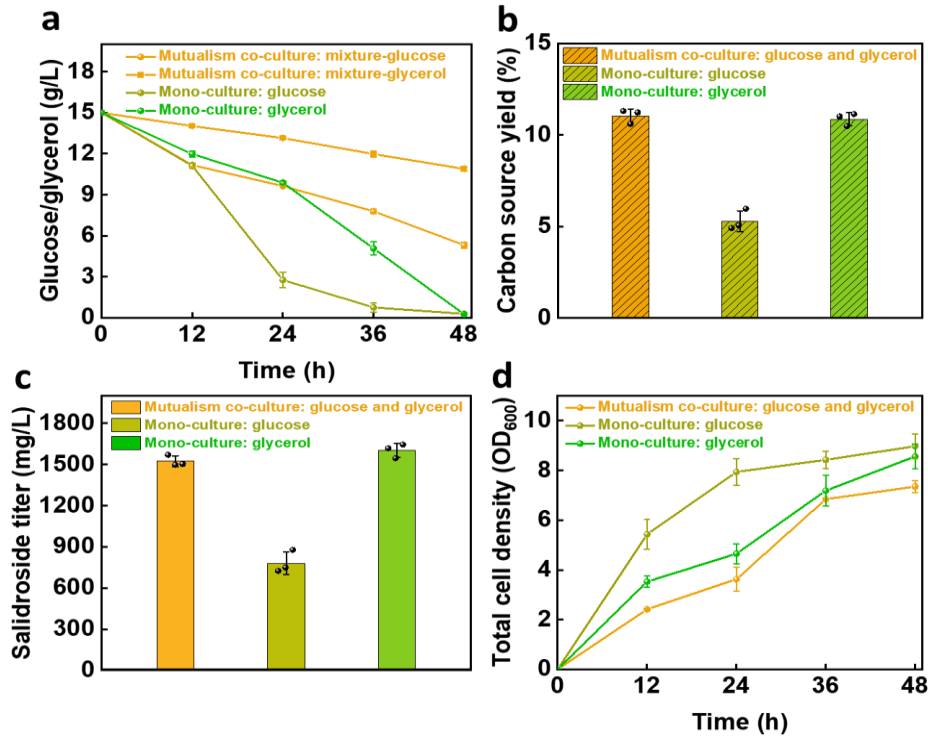


Supplementary Fig. 4. Metabolomic analysis of the supernatants of the neutral and the mutualistic cocultures. Samples were taken at 36h. Data shown are mean \pm SD (n=3 independent experiments). Source data are provided as a Source Data file.

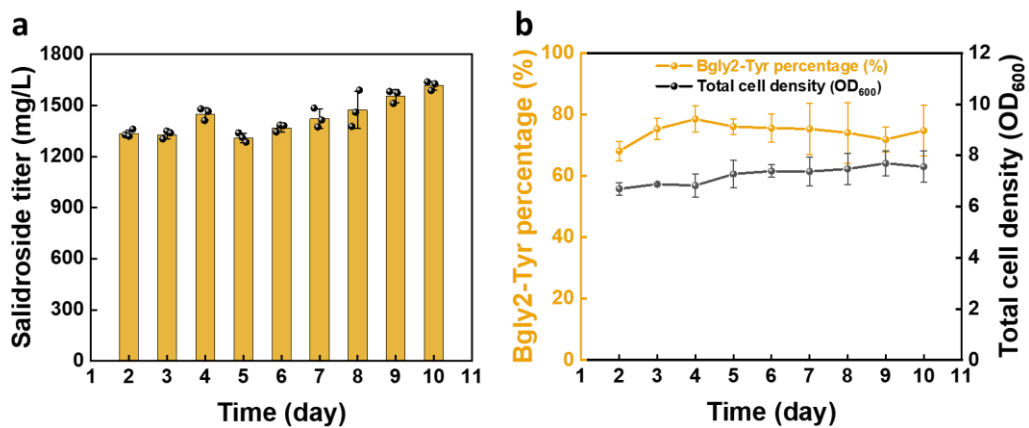


Supplementary Fig. 5. Biosynthetic pathways of salidroside, coniferol and silybin/isosilybin.

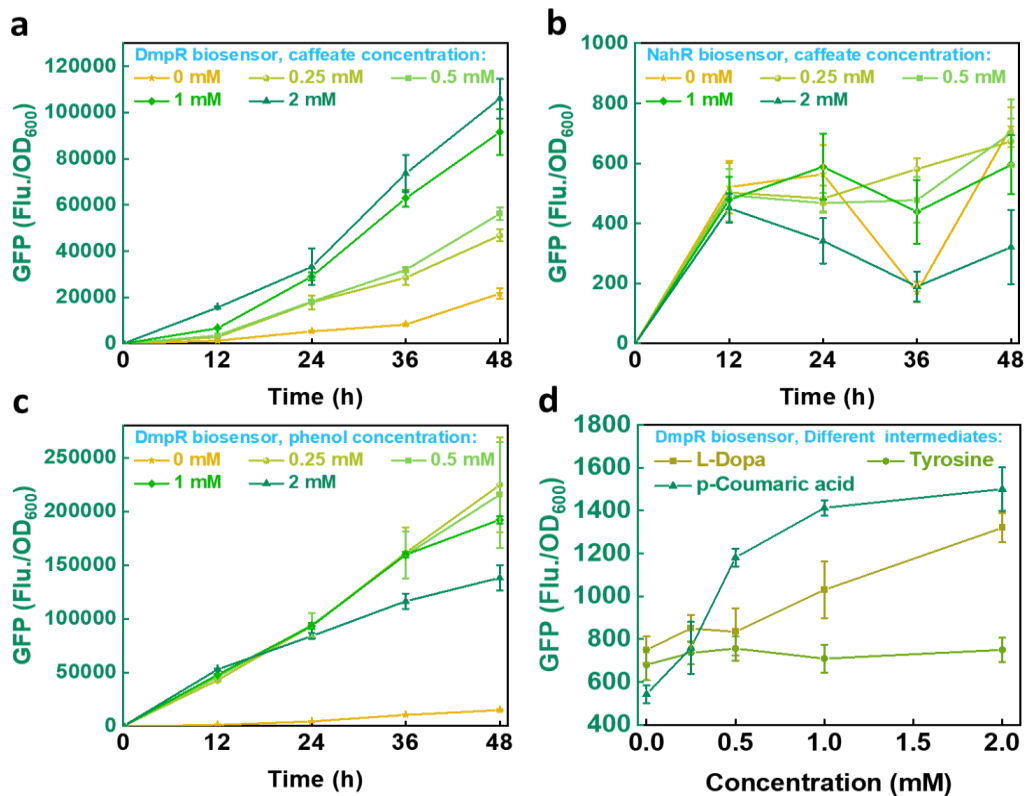
Genes: *ppc* encodes phosphoenolpyruvate carboxylase; *pykA/F* encodes pyruvate kinase; *ppsA* encodes phosphoenolpyruvate synthetase; Enzymes: AroG, 3-deoxy-7-phosphoheptulonate synthase; TyrA, prephenate dehydrogenase; TyrB, aromatic amino acid transaminase; TAL, tyrosine ammonia lyase; HpaBC, 4-hydroxyphenylacetic acid 3-hydroxylase; ARO10, ketoacid decarboxylase; ADH6, alcohol dehydrogenase; UGT85A1, glycosyltransferase; Pgm, phosphoglucomutase; GalU, UDP-glucose pyrophosphorylase; 4CL1, 4-coumarate-CoA ligase; CCoAOMT, caffeoyl-CoA O-methyltransferase; CCR, cinnamoyl-CoA reductase; ADH, alcohol dehydrogenase; CHS, chalcone synthase; CHI, chalcone isomerase; F3H, flavanone 3-hydroxylase; APX1, ascorbate peroxidase 1; NOX1, NADH oxidase. Metabolites: PEP, phosphoenolpyruvate; E4P, erythrose 4-phosphate; DAHP, 3-deoxy-7-phosphoheptulonate; CHA, chorismate; PYR, pyruvate; L-GLU, L-glutamate; G6P, glucose-6-phosphate; G1P, glucose-1-phosphate; UDP-glucose, uridine diphosphate glucose.



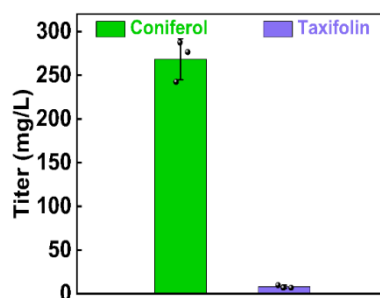
Supplementary Fig. 6. Comparison of salidroside production in the monoculture and the coculture. **a** Curves of carbon sources consumption, **b** Salidroside yields, **c** Salidroside titers, **d** Curves of cell growth. Strain BW-Sal was used in the monoculture while strains Bgly2-Tyr/Bglc2-Sal were used in the coculture. The IIR of the coculture is 1:1. Data shown are mean \pm SD (n=3 independent experiments). Source data are provided as a Source Data file.



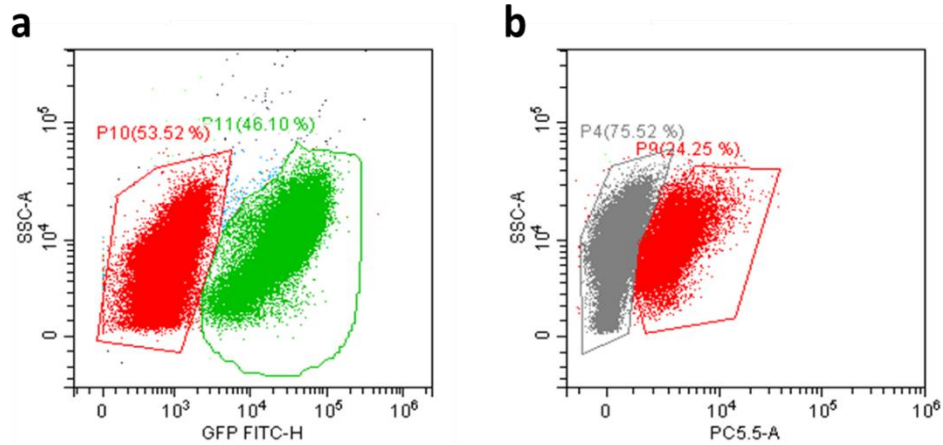
Supplementary Fig. 7. Continuous passage cultivation of the mutualistic coculture. **a** Salidroside titers, and **b** Population composition of each subculture. Data shown are mean \pm SD (n=3 independent experiments). Source data are provided as a Source Data file.



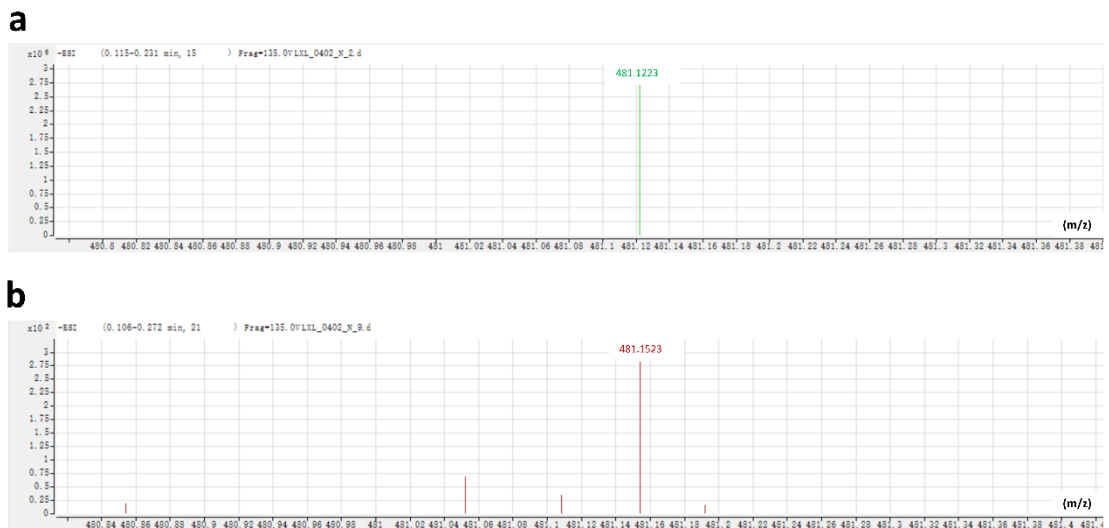
Supplementary Fig. 8. Responsiveness of DmpR and NahR biosensors to different phenolic compounds. a DmpR to caffeate, b NahR to caffeate, c DmpR to phenol and d DmpR to L-dopa, tyrosine and *p*-coumaric acid. Data shown are mean \pm s.d. (n = 3 independent experiments). Source data are provided as a Source Data file.



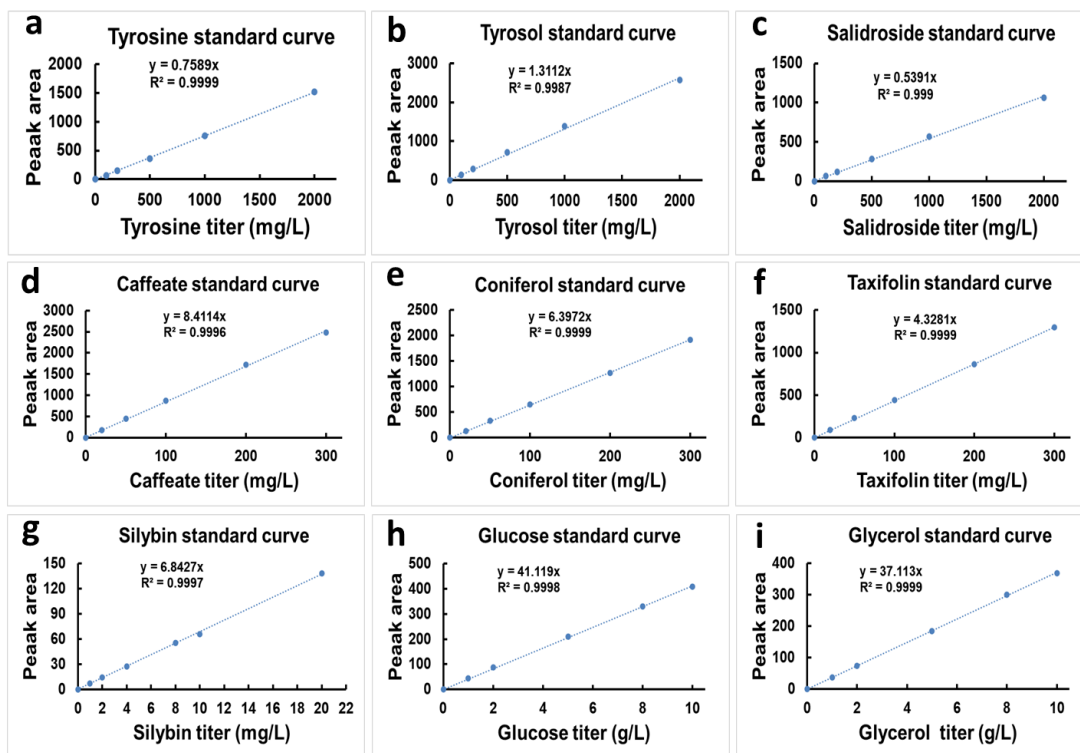
Supplementary Fig. 9. Production of coniferol and taxifolin from caffeate. Strains BW21513 (pZE-CA4C) and BW21513 (pZE-CCF4) were used. Caffeate (500 mg/L) was fed to the cell cultures, and samples were taken at 48 h. The Data shown are mean \pm s.d. (n = 3 independent experiments). Source data are provided as a Source Data file.



Supplementary Fig. 10. Two examples of the flow cytometry results. a Analysis of the population composition with green and red fluorescence. **b** Analysis of the population composition with/without red fluorescence. SSC-A, Side Scatter-Area; FITC-H, Fluoresce in Isothiocyanate-Height; PC5.5-A, Phycoerythrin Cyanin 5-Area.



Supplementary Fig. 11. ESI-MS results of silybin/isosilybin. a The standard, **b** The sample. Source data are provided as a Source Data file.



Supplementary Fig. 12. Standard curves of the related compounds in HPLC. a Tyrosine, **b** Tyrosol, **c** Salidroside, **d** Caffeate, **e** Coniferol, **f** Taxifolin, **g** Silybin/isosilybin, **h** Glucose and **i** Glycerol. Source data are provided as a Source Data file.

Supplementary Table 1. Strains used in this study.

Strains	Description	Source
BW25113	<i>rrnBT14 ΔlacZWJ16 hsdR514 ΔaraBADAH33 ΔrhaBADLD78</i>	Coli genome stock center
BW25113ΔpykAF	BW25113ΔpykAΔpykF	1
Bgly1	BW25113ΔpykAΔpykFΔptsGΔglkΔmanXYZ	This study
Bglc1	BW25113Δglpk	This study
Bgly2	BW25113ΔpykAΔpykFΔptsGΔglkΔmanXYZΔppc	This study
Bglc2	BW25113ΔglpkΔgdhAΔgltBD	This study
Bgly1-Tyr	Bgly1, pZE12-luc and pCS-TPTA and pSA-mcherry	This study
Bglc1-Sal	BGlc1, pZE-ugt85A1 and pCS-pg and pSA-AA	This study
Bgly2-Tyr	Bgly2, pZE12-luc and pCS-TPTA and pSA-mcherry	This study
Bglc2-Sal	Bglc2, pZE-ugt85A1 and pCS-pg and pSA-AA	This study
BW-Sal	BW25113ΔpykAF, pZE-ugt85A1 and pCS-TPTA-pg and pSA-AA	This study
Bgly2-Caf	Bgly2, pZE-TAL and pCS-TH and pSA-mcherry	This study
Bglc2-Con	Bglc2, pZE-CA4C and pCS27 and pSA-eGFP	This study
Bglc2-ConDmpR	Bglc2, pZE-CA4C and pCS27 and pSA-P _{dmp} -GS-P _{J23101} -dmpR	This study
Bglc2-Sil	Bglc2, pZE-CA and pCS-CF4C-P _{J23101} -CA and pSA-P _{dmp} -GS-P _{J23101} -DN	This study
Bglc2-Con(b)	Bglc2, pZE12-luc and pCS-4C-P _{J23101} -CA and pSA-eGFP	This study
Bglc2-Sil(b)	Bglc2, pZE-CA and pCS-CF4 and pSA-P _{dmp} -GS-P _{J23101} -DN	This study

Supplementary Table 2. Plasmids used in this study.

Plasmids	Description	Source
pZE12-luc	P _L lacO1, <i>colE</i> ori, <i>luc</i> , <i>Amp</i> ^R	1
pCS27	P _L lacO1, <i>P15A</i> ori, <i>Kan</i> ^R	1
pSA74	P _L lacO1, <i>pSC101</i> ori, <i>CI</i> ^R	1
pCS-TPTA	pCS27, <i>tyrA</i> , <i>ppsA</i> , <i>tktA</i> and <i>aroG</i> from <i>E. coli</i>	1
pSA-mcherry	pSA74, <i>mcherry</i>	This study
pZE-ugt85A1	pZE12-luc, <i>ugt85A1</i> from <i>Arabidopsis thaliana</i>	This study
pCS-pg	pCS27, <i>pgm</i> and <i>galU</i> from <i>E. coli</i>	This study
pSA-AA	pSA74, <i>aro10</i> and <i>adh6</i> from <i>Saccharomyces cerevisiae</i>	This study
pZE-TAL	pZE12-luc, <i>tal</i> from <i>Rhodobacter glutinis</i>	2
pCS-TH	pCS27, <i>tyrA</i> , <i>ppsA</i> , <i>tktA</i> , <i>aroG</i> and <i>hpaBC</i> from <i>E. coli</i>	1
pZE-CCF4	pZE12-luc, <i>chs</i> from <i>Petunia x hybrida</i> , <i>chi</i> from <i>Medicago sativa</i> , <i>f3h</i> from <i>Camellia sinensis</i> and <i>4cl1</i> from <i>A. thaliana</i>	This study
pZE-CA4C	pZE12-luc, <i>ccr</i> from <i>Leucaena leucocephala</i> , <i>adh</i> from <i>S.cerevisiae</i> , <i>4cl1</i> , <i>ccoamt</i> from <i>A. thaliana</i>	2
pSA-eGFP	pSA74, <i>egfp</i>	This study
pSA- P _{J23101} -dmpR	pSA74, P _{J23101} promoter, <i>dmpR</i> from <i>Pseudomonas CF600</i>	This study
pSA- P _{J23101} -nahR	pSA74, P _{J23101} promoter, <i>nahR</i> from <i>Pseudomonas putida</i>	This study
pZE-P _{dmp} -eGFP	pZE12-luc, <i>egfp</i> with P _{dmp} promoter	This study
pZE-P _{nah} -eGFP	pZE12-luc, <i>egfp</i> with P _{nah} promoter	This study
pSA-P _{dmp} -GS-P _{J23101} -dmpR	pSA74, <i>gdhAssrA</i> from <i>E. coli</i> with P _{dmp} promoter and <i>dmpR</i> with P _{J23101} promoter	This study
pSA-P _{dmp} -GS-P _{J23101} -DN	pSA74, <i>gdhAssrA</i> with P _{dmp} promoter, <i>dmpR</i> and <i>nox1</i> (from <i>Lactococcus lactis</i>) with P _{J23101} promoter	This study
pZE-APX1	pZE12-luc, <i>apx1</i> from <i>Silybum marianum</i>	This study
pCS-P _{lac} -CF4C-P _{J23101} -CA	pCS27, <i>chi</i> , <i>f3h</i> , <i>4cl1</i> and <i>ccoamt</i> with P _{lac} promoter, <i>ccr</i> and <i>adh</i> with P _{J23101} promoter	This study
pZE-CHS	pZE12-luc, <i>chs</i>	This study
pCS-CF4	pCS27, <i>chi</i> , <i>f3h</i> and <i>4cl1</i>	This study
pZE-CA	pZE12-luc, <i>chs</i> and <i>apx1</i>	This study
pCS-4C-P _{J23101} -CA	pCS27, <i>4cl1</i> and <i>ccoamt</i> with P _{lac} promoter, <i>ccr</i> and <i>adh</i> with P _{J23101} promoter	This study
pSA-P _{J23101} -NOX1	pSA74, <i>nox1</i> with P _{J23101} promoter	This study

Supplementary references

1. Li, X. et al. Establishing an artificial pathway for efficient biosynthesis of hydroxytyrosol. *ACS Synth Biol* **7**, 647-654 (2018).
2. Chen, Z., Sun, X., Li, Y., Yan, Y. & Yuan, Q. Metabolic engineering of *Escherichia coli* for microbial synthesis of monolignols. *Metab Eng* **39**, 102-109 (2017).