

## Electronic Supplementary Information (ESI)

### **Bypassing the bottlenecks in the shikimate and methylerythritol phosphate pathways for enhancing the production of natural products from methane in *Methylotuvimicrobium alcaliphilum* 20Z**

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Supplementary Figures

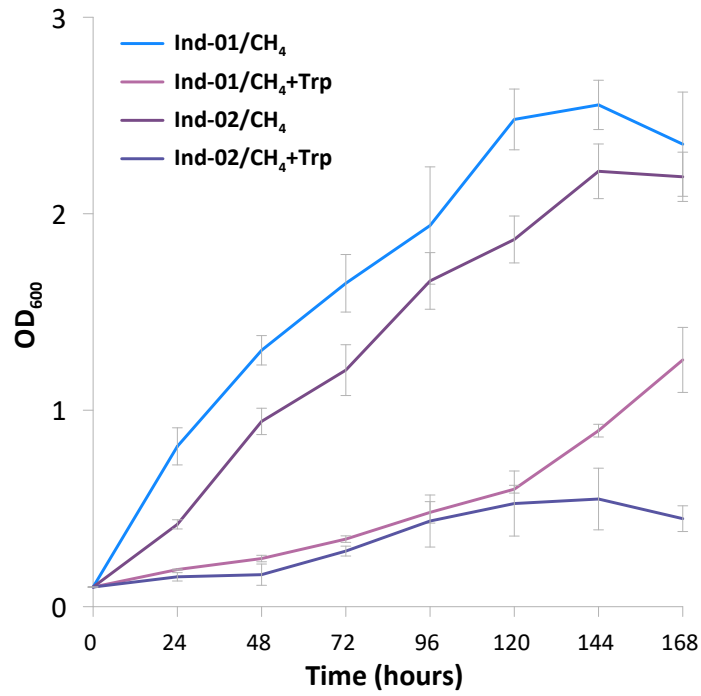


Figure S1. Growth profiles of Ind-01 and Ind-02 strains in medium supplied with methane or methane and 0.5 g/L of tryptophan.

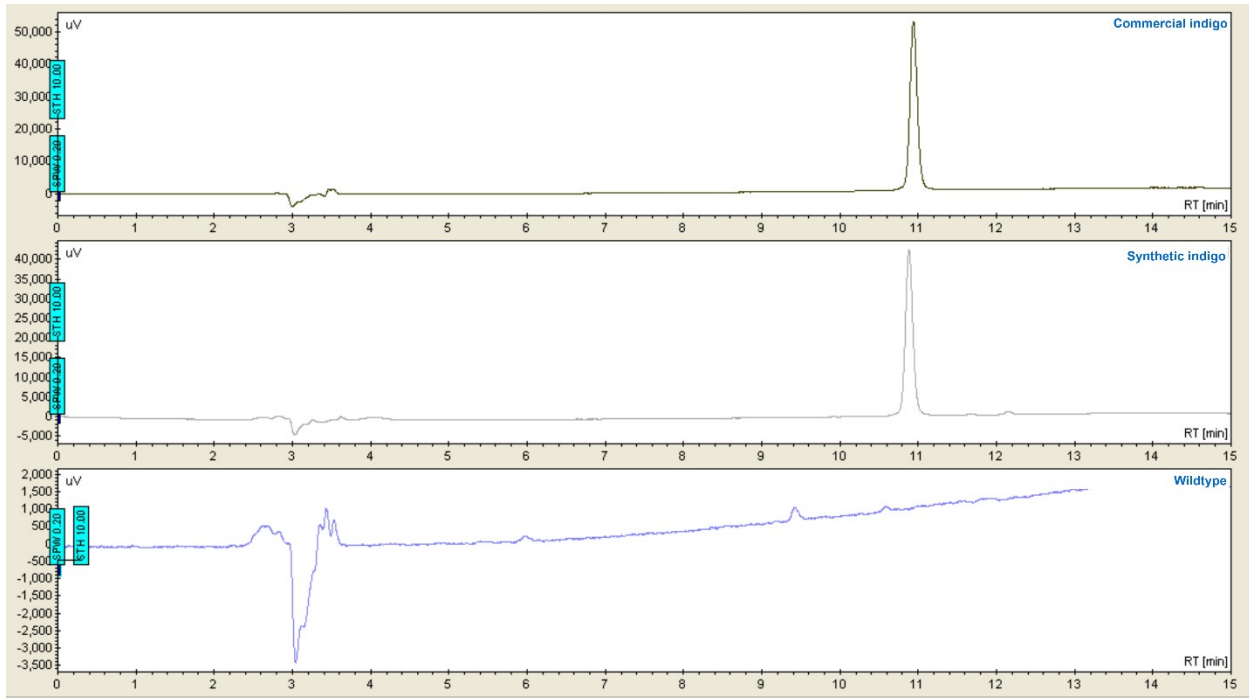


Figure S2. LC chromatograms of commercial indigo and synthetic indigo produced from Ind-03 strain.

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      10      20      30      40      50      60      70      80
C. glutamicum  MEMVMKMKRVAIIGAGPSSGIAQLRAFESAEKQGHIEPELVCFEKQDTWGGQWNYSWRTGTDSYGEPVHSSMYRNLWSNGP
M. aminisulfidivorans  ----MATRIAILGAGPSSGMAQLRAFQSAQEKGAIEPELVCFEKQADWGGQWNYTWRTGLDENGEPVHSSMYRYLWSNGP
Clustal Consensus  .*:**:*:*****:*****:*:.* *****  *****:**** *. *****  *****

      110     120     130     140     150     160     170     180
C. glutamicum  SYPPREVLWDYIAGRRAKKSNEKYIKFAHVVRWVSFDEATKLFVTVTENLRTGETSSDITYDNVIVGAGHFSFPNVPHFDG
M. aminisulfidivorans  SYPPREVLWDYIKGRVEKAGVRKYIRENTAVRHVEFNEDSQTFVTVQDHTDITIYSEEFDYVVCCTGHFSTFYVPEFEG
Clustal Consensus  ***** **.:*:*:*:*  .** *:*:* :. *****: *  * : : * * : :***** * *.:*

      210     220     230     240     250     260     270     280
C. glutamicum  ADKDLLIGASYSIEDIGTQAYKMGARSVTFYSRNPNGYEWPEEMTELPLVERFDGSEVHFVNGEKRKVDIVVFCGTGYL
M. aminisulfidivorans  KDKTVLLVGSYSIEDIGSQCYKYGAKKLISCYRTAPMGYKWPENWDERPNLVRVDTENAYFADGSSEKVDAILLCTGYI
Clustal Consensus  ** :*:*.*****:*:* **.:. :.*: *****:***: * * : * * .:.*:.*.*** :.:*****:

      310     320     330     340     350     360     370     380
C. glutamicum  DTLYRGVVSSEANNQLFWLGAQDQWLTFNMFDAQAWYVRDVILGRVALPSKEAQRNHMQWLSRFEGLKSENDQIDFQCDY
M. aminisulfidivorans  LNLKGVVWEDNPKFFYIGMQDQWYSFNMFDAQAWYARDVIMGRLLPSPKEEMKADSMARREKELTLVTAEMEYTYQGDY
Clustal Consensus  .**:* ** * :.:*:* ***** :*****:*****:*:***** :. * .: * : : : * **

      410     420     430     440     450     460     470
C. glutamicum  ILKGWVKSKEEDILNRYDYTYTSVMTGTTSEVHHTFWMIELDDSLERYLSEFPQDEARQVYRKVKVRDKA
M. aminisulfidivorans  TFLKWKHHKKNIMTFRDHSYRSIMTGTMAPKHHTFWIDALDDSLAYLSDKSE-----IPVAKEA
Clustal Consensus  : * : *:*:.*:.*:* *:* **.: ***** : :*****: ***** ***: .* * :.*

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Figure S3. Sequence alignment of Fmo from *C. glutamicum* and *M. aminisulfidivorans*.

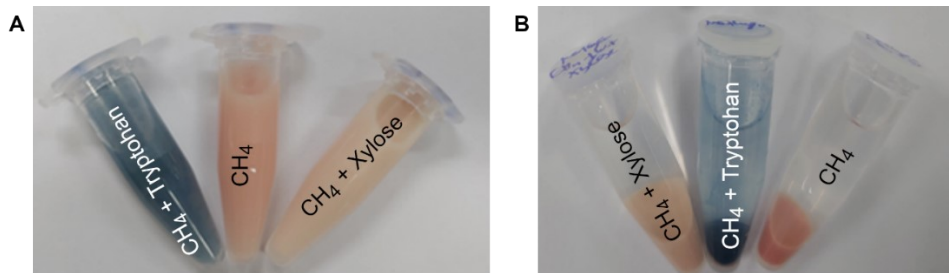


Figure S4. Culture broth of Ind-04 strain in methane, methane plus xylose and methane plus tryptophan before (A) and after (B) centrifugation.

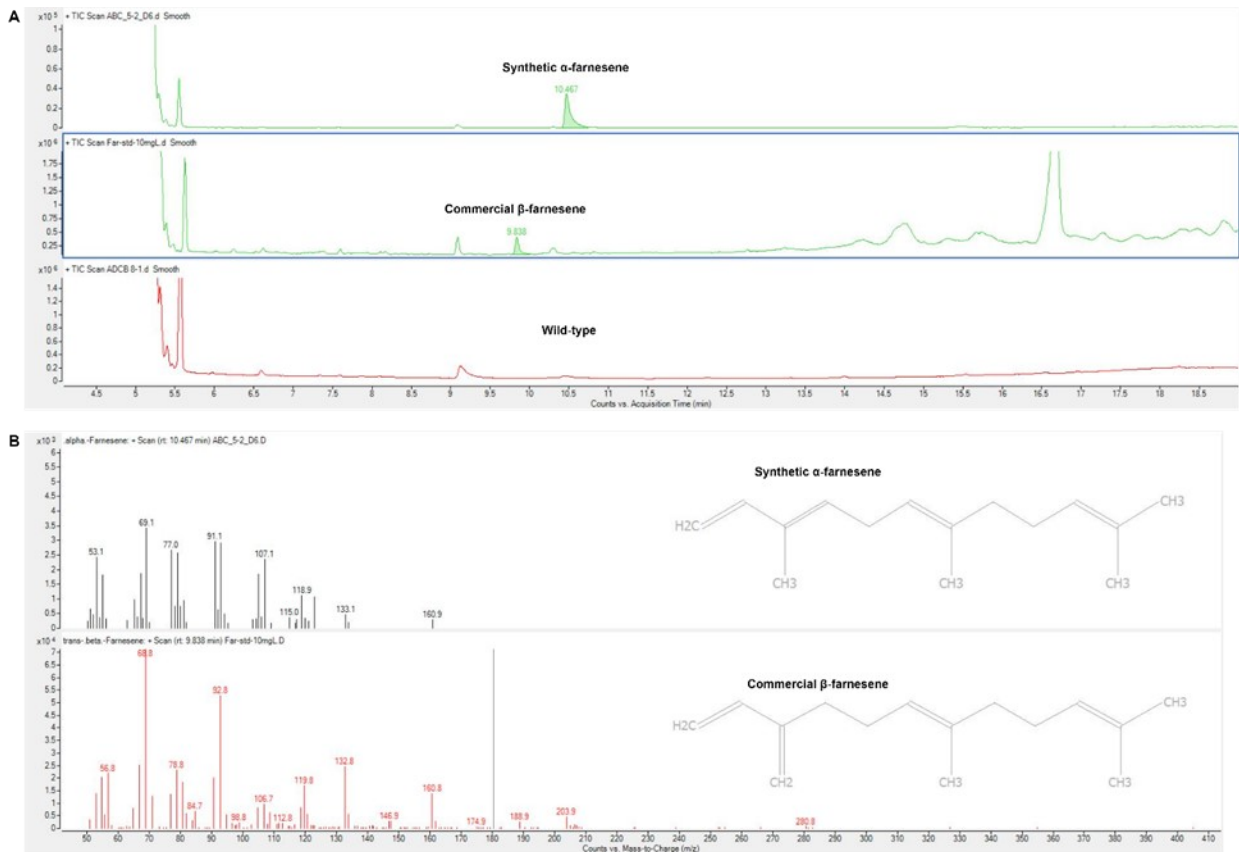


Figure S5. GC chromatograms (A) and mass spectrum chromatograms (B) of synthetic  $\alpha$ -farnesene and commercial  $\beta$ -farnesene.

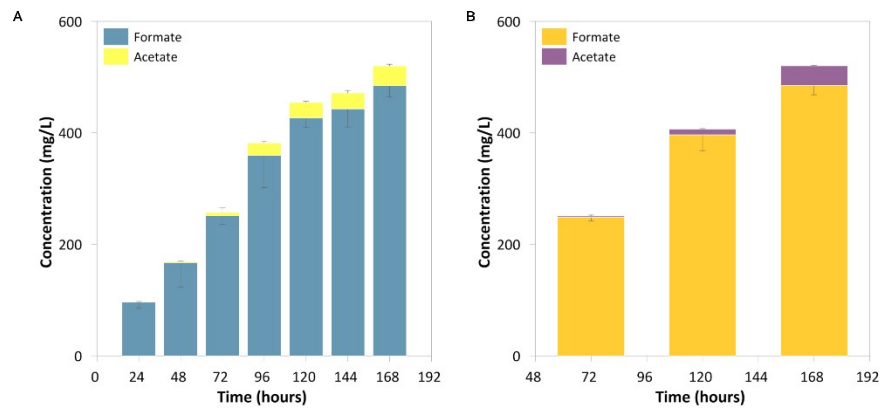


Figure S6. Excretion of organic acids including formate and acetate during cultivation of engineered strains Ind-05 (A) and FAR-02 (B) on methane.

## Supplementary Tables

Table S1. Strains and plasmids used in this study

Strain	Characteristic	Reference
<i>Escherichia coli</i> DH5 $\alpha$		Novagen
<i>Corynebacterium glutamicum</i>		KCTC
<i>Methylophaga aminisulfidivorans</i>		KCTC
<i>Methylotuvimicrobium alcaliphilum</i> 20Z	Used as host strain	DMSZ
Ind-01	<i>M. alcaliphilum</i> 20Z harboring pFMO-Cg plasmid	This study
Ind-02	<i>M. alcaliphilum</i> 20Z harboring pFMO-Ma plasmid	This study
Ind-03	<i>M. alcaliphilum</i> 20Z harboring pFMO-Ma-A plasmid	This study
Ind-04	<i>M. alcaliphilum</i> 20Z harboring pFMO-Ma-T plasmid	This study
Ind-05	<i>M. alcaliphilum</i> 20Z harboring pFMO-Ma-AT plasmid	This study
Ind-06	<i>M. alcaliphilum</i> 20Z harboring pFMO-Ma-AT plasmid with chromosomally integrated xylose-utilizing pathway genes	This study
FAR-01	<i>M. alcaliphilum</i> 20Z harboring pAFS plasmid	This study
FAR-02	<i>M. alcaliphilum</i> 20Z harboring pAFS-nDXP plasmid	This study

FAR-03	<i>M. alcaliphilum</i> 20Z harboring pAFS-nDXP plasmid with chromosomally integrated xylose-utilizing pathway gene	This study
<b>Plasmid</b>		
pAWP89	oriV oriT trfA ahp dTomato pTac	Puri et al. 2015 <sup>1</sup>
pCM351-XYL	pCM351-(FR)glgA containing construct for integrating $P_{tac}$ - <i>xyfAB-rpe</i> into chromosome	Nguyen et al. 2021 <sup>2</sup>
pBs-02	pAWP89-based backbone carrying <i>AgBS-ispA-dxs-ispG-ribB-DSAG-dxr</i>	Nguyen et al. 2021 <sup>3</sup>
pET-28a(+)	Expression vector with N-terminal His-tag	Novagen
pET-tna	pET-28a carrying <i>tnaA</i>	This study
pET-tna-Mfmo	pET-28a carrying <i>tnaA</i> and <i>fmo</i> from <i>M. aminisulfidivorans</i>	This study
pET-tna-Cfmo	pET-28a carrying <i>tnaA</i> and <i>fmo</i> from <i>C. glutamicum</i>	This study
pET-Trp	pET-28a carrying <i>aroG</i> <sup>D146N</sup> and <i>trpE</i> <sup>S40F</sup>	Pham et al. 2022 <sup>4</sup>
pET-indigo-trpE	pET-28a carrying <i>tna-Mfmo-trpE</i> <sup>S40F</sup>	This study
pET-indigo-aroG-trpE	pET-28a carrying <i>tna-Mfmo-aroG</i> <sup>D146N</sup> - <i>trpE</i> <sup>S40F</sup>	This study
pFMO-Cg	pAWP89-based backbone carrying $P_{tac}$ promoter and <i>tna-Cfmo</i> amplified from pET-28a-tna-Cfmo plasmid	This study
pFMO-Ma	pAWP89-based backbone carrying $P_{tac}$ promoter and <i>tna-Mfmo</i> amplified from pET-28a-tna-Mfmo plasmid	This study

pFMO-Ma-A	pAWP89-based backbone carrying $P_{tac}$ promoter and <i>tna-Mfmo-aroG</i> <sup>D146N</sup> amplified from pET-28a-indigo-aroG-trpE plasmid	This study
pFMO-Ma-T	pAWP89-based backbone carrying $P_{tac}$ promoter and <i>tna-Mfmo-trpE</i> <sup>S40F</sup> amplified from pET-28a-indigo-trpE plasmid	This study
pFMO-Ma-AT	pAWP89-based backbone carrying $P_{tac}$ promoter and <i>tna-Mfmo-aroG</i> <sup>D146N</sup> - <i>trpE</i> <sup>S40F</sup> amplified from pET-28a-indigo-aroG-trpE plasmid	This study
pAFS	pAWP89-based backbone carrying $P_{tac}$ promoter <i>afs</i> and <i>ispA</i>	This study
pAFS-nDXP	pAWP89-based backbone carrying $P_{tac}$ promoter <i>afs</i> and <i>ispA</i> and <i>ribB-DSAG-dxr</i> amplified from pBs-02 plasmid	This study

Table S2. Primers used in this study

Primers	Primer sequences (5'-3')	Description
pAWP89_fw	TAGTTGTCGGGAAGATGCG	For amplifying pAWP89 backbone, in which promoter $P_{tac}$ and SD were kept.
pAWP89_rv	AGCTGTTTCTGTGTGAATA	
afs_fw	aggtattcacacaggaacagctATGGAATTTCTGGGTTTCAT	For amplifying <i>afs</i> gene from synthetic <i>afs</i> fragment to ligate to <i>ispA</i> and pAWP89 backbone, resulting in pAFS plasmid.
afs_rv	agaggtactcagagaCTAGTTAACCAACGGTTG	
ispA_fw	caaccgttggttaactagGATCTGAGTACCTCTAGAAAATAAGGAGCAATCCAATG AGTAACGCACTGAAAG	For amplifying <i>ispA</i> gene from genomic DNA of <i>M. alcaliphilum</i> 20Z to ligate to <i>afs</i> and pAWP89 backbone, resulting in pAFS plasmid.
ispA_rv	gcatcttcccacaactaTTAATGATCTCGCTGGAT	
pAWP89_fw	TAGTTGTCGGGAAGATGCG	For amplifying pAFS backbone, in which promoter $P_{tac}$ , SD, <i>ispA</i> , and <i>afs</i> genes were kept.
pAWP89_Far_rv	TTAATGATCTCGTGGATAATA	
ribBC_fw	atccagcgagatcattaaTTCACACAGGAAACAGCTATGAATCAGACGCTACTT	For amplifying <i>ribB-DSAG-dxr</i> fragment from pBs-02 plasmid, which was constructed by our group in previous study (Nguyen et al. 2021) <sup>3</sup> , to ligate to pAFS backbone, resulting in pAFS-nDXP plasmid.
ribBC_rv	gcatcttcccacaactaTTAACGCTTAAGTTCTTCGAC	
tnaA_fw	gacagcaaatgggtcgcgATGAAAACTTTAAACATCTCC	For amplifying <i>tnaA</i> gene from genomic DNA of <i>E. coli</i> to ligate to pET-28a at BamHI restriction site, resulting in pET-28a-tna plasmid.

tnaA_rv	cgacggagctcgaattcgTAAACTTCTTTAAGTTTTGCG	
Mfmo_fw	acttaaagaagttaacgTATTCACACAGGAAACAGCTATGGCAACTCGTATTGCG	For amplifying <i>fmo</i> gene from genomic DNA of <i>M. aminisulfidivorans</i> to ligate to pET-28a-tna at SacI restriction site, resulting in pET-28a-tna-Mfmo plasmid.
Mfmo_rv	gcttgtcgacggagctcgTTAAGCTTCTTTAGCCACAG	
Cfmo_fw	acttaaagaagttaacgTATTCACACAGGAAACAGCTATGAAGAATAAGCGCGTT	For amplifying <i>fmo</i> gene from genomic DNA of <i>C. glutamicum</i> to ligate to pET-28a-tna at SacI restriction site, resulting in pET-28a-tna-Cfmo plasmid.
Cfmo_rv	gcttgtcgacggagctcgTTAGGCTTTATCGCGGAC	
trpE_fw	gacagcaaatgggtcgcgTATTCACACAGGAAACAGCTATGCAAACACAAAAACC GACTCT	For amplifying <i>trpE<sup>fbr</sup></i> from pET-Trp to ligate to pET-tna-Mfmo at NotI restriction site, resulting in pET-28a-Ma-T plasmid.
trpE_rv	gcttgtcgacggagctcgTCAGAAAGTCTCTGTGC	
aroG_fw	gacagcaaatgggtcgcgTATTCACACAGGAAACAGCTATGAATTATCAGAACGA CGA	For amplifying <i>aroG<sup>fbr</sup>-trpE<sup>fbr</sup></i> from pET-Trp to ligate to pET-tna-Mfmo at NotI restriction site, resulting in pET-28a-Ma-AT plasmid.
trpE_rv	gcttgtcgacggagctcgTCAGAAAGTCTCTGTGC	
tnaA_89_fw	ttcacacaggaaacagctATGGAAAACCTTTAAACATCTCC	For amplifying <i>tnaA-Mfmo</i> cluster gene from pET-28a-tna-Mfmo plasmid to ligate to pAWP89 backbone, resulting in pFMO-Ma plasmid.
Mfmo_89_rv	gcatcttcccgacaactaTTAAGCTTCTTTAGCCACAG	
tnaA_89_fw	ttcacacaggaaacagctATGGAAAACCTTTAAACATCTCC	For amplifying <i>tnaA-Cfmo</i> cluster gene from pET-28a-tna-Cfmo plasmid to ligate to pAWP89 backbone, resulting in pFMO-Cg plasmid.
Cfmo_89_rv	gcatcttcccgacaactaTTAGGCTTTATCGCGGAC	
tnaA_89_fw	ttcacacaggaaacagctATGGAAAACCTTTAAACATCTCC	For amplifying <i>tnaA-Mfmo-aroG</i> cluster gene from pET-28a-indigo-aroG-trpE plasmid to ligate to pAWP89 backbone, resulting in pFMO-Ma-AT plasmid.
aroG_89_rv	gcatcttcccgacaactaTTACCCGCGACGCGCTTTTA	
tnaA_89_fw	ttcacacaggaaacagctATGGAAAACCTTTAAACATCTCC	For amplifying <i>tnaA-Mfmo-trpE</i> cluster gene from pET-28a-indigo-trpE plasmid to ligate to pAWP89 backbone, resulting in pFMO-Ma-AT plasmid.
trpE_89_rv	gcatcttcccgacaactaTCAGAAAGTCTCTGTGC	
tnaA_89_fw	ttcacacaggaaacagctATGGAAAACCTTTAAACATCTCC	For amplifying <i>tnaA-Mfmo-aroG-trpE</i> cluster gene from pET-28a-indigo-aroG-trpE plasmid to ligate to pAWP89 backbone, resulting in pFMO-Ma-AT plasmid.
trpE_89_rv	gcatcttcccgacaactaTCAGAAAGTCTCTGTGC	

Homologous sequences used for Gibson Assembly are indicated by lower-case. Ribosome binding sites are underlined.

Table S3. Sequence of homologous genes used in this study



Gene	Sequence
<i>tnaA (E. coli K12)</i>	<p>ATGGAAAACCTTTAAACATCTCCCTGAACCGTTCCGCATTCTGTTATTGAGCCAGTAAAACGTACCCTCGCGCTTATCGTGAAGAGGCAATTATTAATCCGGTAT  GAACCCGTTCTGCTGGATAGCGAAGATGTTTTATCGATTTACTGACCGACAGCGGCACCGGGGCGGTGACGCAGAGCATGCAGGCTGCGATGATGCGCGGCG  ACGAAGCCTACAGCGGCAGTCTGACTACTATGCGTTAGCCGAGTCACTGAAAAATATCTTCGTTATCAATACACCATTCCGACTACCAGGGCCGTGGCGCAG  AGCAAATCTATATCCGGTACTGATTA AAAAACGCGAGCAGGAAAAAGGCTGGATCGCAGCAAAATGGTGGCGTTCTAACTATTTCTTGATACCACGCAGG  GCCATAGCCAGATCAACGGCTGTACCGTGCCTAACGTCTATATCAAAGAAGCCTTCGATACGGGCGTGCCTACGACTTTAAAGGCAACTTTGACCTTGAGGGAT  TAGAACGCGGTATTGAAGAAGTTGGTCCGAATAACGTGCCGTATATCGTTGCAACCATACCAGTAACTCTGCAGGTGGTCAGCCGGTTTCACTGGCAAACCTAA  AAGCGATGTACAGCATCGCGAAGAAATACGATATTCCGGTGGTAATGGACTCCGCGCGCTTTGCTGAAAACGCCTATTTCAATTAAGCAGCGTGAAGCAGAATACA  AAGACTGGACCATCGAGCAGATCACCCGCGAAACCTACAAATATGCCGATATGCTGGCGATGTCCGCCAAGAAAGATGCGATGGTGCCGATGGGCGGCGCTGCTG  TGATGAAAGACGACAGCTTCTTTGATGTGTACACCGAGTGCAGAACCCTTTGCGTGGTGCAGGAAGGCTTCCCGACATATGGCGGCTAGAAGGCGGCGCGAT  GGAGCGTCTGGCGGTAGGTCTGTATGACGGCATGAATCTCGACTGGCTGGCTTATCGTATCGCGCAGGTACAGTATCTGGTGCATGGTCTGGAAGAGATTGGCG  TTGTCTGCCAGCAGGCGGCGGTACGCGGCATTCTGTTGATGCCGTAACCTGTTGCCGCATATCCCGCAGACCAGTCCCGGCAACAGGCCTGGCCTGCGAG  CTGTATAAAGTCGCCGTATCCGTGCGGTAGAAATTGGCTCTTCTGTTAGGCCGATCCGAAAACCGGTAAACAACCTGCCATGCCGGCTGAACTGCTGCGTT  TAACCATTCCGCGCGCAACATATACTCAAACACATATGGACTTCATTATTGAAGCCTTTAAACATGTGAAAGAGAACGCGGCGAATATTAAGGATTAACCTTTAC  GTACGAACCGAAAGTATTGCGTCACTTACCAGCAAACCTAAAGAAGTTTAA</p>
<i>Cfmo (C. glutamicum)</i>	<p>ATGGAGATGGTTATGAAGAATAAGCGCGTTGCGATTATTGGTGCAGGTCCGAGTGGTATCGCTCAGTTGAGGGCGTTTGTAGTCTGCTGAAAAGCAGGGTCAATGA  GATCCCTGAGCTGGTGTGTTTTGAAAAGCAGGATACTGGGGTGGGCGAGTGAATTAATCTTGGCGCACGGGAACAGACTCTTATGGTGAGCCTGTGCACTCAA  GTATGTACCGAAACCTGTGGTCAAACCGTCCGAAGGAAGTCTCGAATTTGCTGAGTACAGCTTCGATGAGCACTTCGAAAGCCAATTTCTTACCCTCCACG  TGAAGTGTGGGATTACATTGCAGGTCGTGCAAAGAAGTCAACGTTGAGAAGTATCAAGTTCCGCGCATGTTGTTGCTGGGTGAGTTTTGATGAGGCCAC  CAAGCTGTTACCGTGACGGTGGAGAACCTCCGACCGGTGAGACCAGCAGTGTACTTATGACAACGTGATTGTTGGCGCTGGACACTTCAGTTTCCCGAACGT  CCCTCACTTTGATGGTGTGGAGACTTCCCAAGTCCAGATCATGCATGCTCACGAGTCCGTTGGTGCAGAGGCTGTTGCTGACAAGGATATTTGCTGATTGGTGCA  AGTTATTCTGCGGAAGATATCGGTACCCAGGCGTACAAGATGGGTGCTCGTTCCGTTGACTTCTTACCCTCAAACCAATGGGGTATGAGTGGCCTGAAGAG  ATGACTGAGCTTCTTTGGTTGAGCGTTTCGATGGCTCCGAGGTTCACTTTGTCAATGGTAAAAGCGCAAGGTTGACATCGTGGTGTCTGTACTGGTACTTAC  ACCATTACCATTTATGCCGTCTGAGCTGACTTTAAGCTCACCAACAACCTGTACCCGGATACGCTTTATCGTGGCGTGGTGTCCGAGGCTAATAACCAGCTGTT  TGTTGGGCGCTCAGGATCAGTGGCTGACGTTCAACATGTTTGTGCTCAGGCTGGTATGTTCCGCGATGTCATTTGGGTGCGTGGCTCTTCCCTTCAAGGAGG  CGCAGCGCAATCATATGGATCAGTGGCTGTCACGTTTTGAGGGTTTGAAGTCTGAGAATGATCAGATTGATTCCAGTGCGATTACGTTGAGGACCTCATTGACCA  GACCGATTACCCTTCTGTTGATCTGAAGGAAGTTGCAATATCTTGAAGGGCTGGGTGAAGTCAAGGAGGAGGATATCCTCAACTACCGGATTACACCTACAC  GTCCGTGATGACTGGCACTACCTCTGTTGAGCACCACACTCCGTGGATGATTGAGTTGGATGATTCTTGGAGCGTTACCTCAGCGAGCCTCAGGAAGATGAAGC  TCGTGAGTTTACCCTGGCAAGAAAGTCCGCGATAAAGCCTAA</p>
<i>Mfmo aminosulfidivorans</i>	(M. 221 724 891 868) <p>ATGGCAACTCGTATTGCGATACTGGTGCAGGCCAAGTGGTATGGCACAACCTCAGAGCATTCCAATCCGCCAGGAAAAAGGTGCTGAGATCCCTGAACTCGTT  TGTTTTGAAAAACAAGCTGATTGGGGCGGCCAGTGAATACACATGGCGCACTGGTTAGATGAAAATGGCGAACCTGTTATAGCAGTATGTATCGTATCTG  TGGTCAAACGGCCGAAAAGAATGCTTGAATTTGCTGATTACACGTTTACGAAACACTTTGGTAAGCCCATCGCTCTTATCCACCCCGTGAAGTCTTATGGGACT  ATATTAAGGCCGTGTTGAAAAAGCCGGCGTCAGAAAATATATCCGTTTTAATACCGCTGTTGTCATGTTGAATTCACGAAGACAGCCAAACTTTTACCGTTAC  CGTGCAGGACCATACTACTGACACAATTACTCTGAAGAGTTTACTATGTTGTCTGTTGACCGGTCACCTCTCAACACCTTACGTGCCTGAATTTGAAGGCTTTG  AAAATTTGGTGGCCGATTCTGCATGCCATGACTCCGTGACGCATTAGAATTTAAAGACAAAACCTGATTACTGGTGGCAGCAGTACTCAGCTGAAGATAT  CGGCTACAATGTTATAAATACGGCGCGAAAAACTGATCAGCTGCTACCGTACCGCACCGATGGGTTATAAATGGCCTGAAAACCTGGGATGAAAGACCCAACT  GGTTCGTGTTGATACTGAAAACGCTTATTTGCCGATGTTTATCAGAAAAAGTCGATGCGATTATCTGTGTACCGGTTATATCCATCACTTCCCTTCTCAATGA</p>

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*aroG*<sup>fbr(D146N)</sup>

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GGGAGCCGCTGGCTACGGTAAGAGCATCACCGATGCCTGCATCGGCTGGGAAGATACCGATGCTCTGTTACGTCAACTGGCGAATGCAGTAAAGCGCGTCCG  
GGTAA

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*trpE*<sup>fbr(S40F)</sup>

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*afs* (codon-optimized)

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*ispA* (*M. alcaliphilum*  
20Z)

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*ribB*<sup>G1135 3</sup>

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*dxr* (*M. alcaliphilum*  
20Z)

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Table S4. The production of indigo and farnesene in previous studies. (n.d: not determined)

Host strain	Product	Substrate	Culture conditions	Titer	Production yield	References
<i>E. coli</i>	Indigo	Tryptophan (1.0 g/L)	Optimized medium (3.55 g/L NaCl, 5.12 g/L yeast extract); Batch culture	307.4 mg/L	n.d	Dai et al. <sup>5</sup>
<i>E. coli</i>	Indigo	Tryptophan	Tryptophan medium (2 g/L tryptophan, 5 g/L yeast extract, 10 g/L NaCl); Continuous fermentation, 3000 L	911 mg/L	0.234 mg/g tryptophan	Han et al. <sup>6</sup>
<i>E. coli</i>	Indigo	Glycerol (5 g/L) Tryptophan (0.5 g/L)	M9 medium; Consortium cultivation; Batch fermentation.	104.3 mg/L	n.d	Chen et al. <sup>7</sup>
<i>M. alcaliphilum</i> 20Z	Indigo	Methane (176 mg)	NMS medium; Fed-batch culture, 50 mL	3.5 µg/L	0.994 ug/g methane	This study
<i>M. alcaliphilum</i> 20Z	Indigo	Methane (135 mg) Xylose (641 mg)	NMS medium; Semi fed-batch culture, 50 mL	6.3 µg/L	0.421 ug/g substrates	This study
<i>S. cerevisiae</i>	Farnesene	Cane syrup	Culture medium (15 g/L NH <sub>4</sub> H <sub>2</sub> PO <sub>4</sub> , trace element, vitamin); Fed-batch fermentation, 250 mL	130 g/L	143 mg/g glucose	Meadows et al. <sup>8</sup>
<i>Yarrowia lipolytica</i>	Farnesene	Glucose	Modified YPD medium (50 g/L glucose, 20 g/L tryptone, 10 g/L yeast extract); Fed-batch fermentation, 800 mL	25.55 g/L	n.d	Liu et al. <sup>9</sup>
<i>S. elongatus</i> PCC 7942	Farnesene	CO <sub>2</sub>	BG-11 medium (10 mM MOPS, 5% CO <sub>2</sub> , 100 µE/(m <sup>2</sup> s) continuous fluorescent light); Fed-batch fermentation, 100 mL	12.99 mg/L	n.d	N.P. et al. <sup>10</sup>
<i>M. alcaliphilum</i> 20Z	Farnesene	Methane (140 mg)	NMS medium; Fed-batch culture, 50 mL	48.98 mg/L	17.49 mg/g methane	This study
<i>M. alcaliphilum</i> 20Z	Farnesene	Methane (130 mg) Xylose (740 mg)	NMS medium; Semi fed-batch culture, 50 mL	91.55 mg/L	5.26 mg/g substrates	This study

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