

Supporting information

Table S1 Plasmids used in the study

Plasmid	Description	Reference
pZElac	<i>E. coli</i> vector, amp ^R	1
pZElac-his	The pZElac inserted with the T5 promoter/lacO::6Xhis, amp ^R	2
pZElac_GDH	pZElac inserted with <i>E.coli gdhA</i> wild type, amp ^R	This work
pZElac_GDH F18	pZElac inserted with <i>E.coli gdhA</i> F18 mutant, amp ^R	This work
pZElac_GDH F18TA	pZElac inserted with <i>E.coli gdhA</i> F18 with T195A mutation, amp ^R	This work
pZElac_GDH F18TV	pZElac inserted with <i>E.coli gdhA</i> F18 with T195V mutation, amp ^R	This work
pZElac_GDH F18TL	pZElac inserted with <i>E.coli gdhA</i> F18 with T195L mutation, amp ^R	This work
pZElac_GDH F18TF	pZElac inserted with <i>E.coli gdhA</i> F18 with T195F mutation, amp ^R	This work
pZElac_aspC	pZElac inserted with <i>E.coli aspC</i> , amp ^R	This work
pZElac_tyrB	pZElac inserted with <i>E.coli tyrB</i> , amp ^R	This work
pZElac_pheDH	pZElac inserted with <i>Thermoactinomyces intermedius pdh</i> , amp ^R	This work
pZElac_hisGDH	for his tag- <i>E.coli</i> GDH expression, amp ^R	This work
pZElac_hisF18	for his tag- <i>E.coli</i> GDH F18 mutant expression, amp ^R	This work
pZElac_hisF18TA	for his tag- <i>E.coli</i> GDH F18 T195A expression, amp ^R	This work
KAVS library	pZElac vector harboring <i>E. coli gdhA</i> with K92, A166, V377 and S380 residues randomized by NNK codon	This work

Reference:

1. Zhang, K., Li, H., Cho, K. M., Liao, J. C. 2010. Expanding metabolism for total biosynthesis of the nonnatural amino acid L-homoalanine. *Proc Natl Acad Sci U S A* 107: 6234-6239
2. Li, H., Liao, J. C. 2013. Engineering a cyanobacterium as the catalyst for the photosynthetic conversion of CO₂ to 1,2-propanediol. *Microb Cell Fact* 12: 4

Table S2 Primers used in the study

Primer	sequence (5'-3')
GDH Acc65I fwd	agaaaggtaccatggatcagacatattctctggagtc
GDH XbaI rev	gatgcctctagattaaatcacaccctgcgccagcatcg
aspC Acc65I fwd	gagaaaggtaccatgtttgagaacattaccgccgctc
aspC XbaI rev	gatgcctctagattacagcactgccacaatcgcttcg
tyrB Acc65I fwd	gagaaaggtaccatgtttcaaaaagtgacgcctacg
tyrB XbaI rev	gatgcctctagattacatcaccgcagcaaacgcctttgc
PheDH Acc65I fwd	gagaaaggtaccatgataagaatgggaagcatgaaaatc
PheDH XbaI rev	atgcctctagattacctccttgcgctgttgcgggg
T195A fwd	caacaataccgcctgcgtcttcgctgggtaagggcctttcatttggcg
T195A rev	cgccaaatgaaaggcccttaccgcgaagacgcaggcggtattgttg
T195L fwd	caacaataccgcctgcgtcttcctgggtaagggcctttcatttggcg
T195L rev	cgccaaatgaaaggcccttaccaggaagacgcaggcggtattgttg
T195V fwd	caacaataccgcctgcgtcttcgtgggtaagggcctttcatttggcg
T195V rev	cgccaaatgaaaggcccttaccacgaagacgcaggcggtattgttg
T195F fwd	caacaataccgcctgcgtcttctcggtaagggcctttcatttggcg
T195F rev	cgccaaatgaaaggcccttaccgaagaagacgcaggcggtattgttg
K92 fwd	gcagttcagctctgccatcgcccgctacnnkggcggtatgcgcttccatccgtcagttaa
K92 rev	ttaactgacggatggaagcgcataccgcmnnngtacgggccgatggcagagctgaactgc
A166 fwd	gccacctgggcgcggataccgacgttcggnkkggtgatacgggggttggtggtcgtgaag
A166 rev	cttcacgaccaccaaccccgatcaccmnnncggaacgtcggatccgcgccaggtggc
VS fwd	gtaaagcggctaatactggtggcnnkgctacannkggcctggaaatggcaciaaacgctg
VS rev	cagcgttttgtgccatttcaggccmnnntgtacmnnngccaccagcattagccgctttac

Cofactor specificity of GDH variants

E. coli GDH is naturally NADPH-dependent. To determine the cofactor specificity of the engineered GDH variants toward the final target 2-OPBA, enzyme assays were performed using purified proteins (Figure S1). The results suggested that both GDH F18 and F18 T195A are highly NADPH-specific for this reaction. The NADH-dependent activity was very minimal.

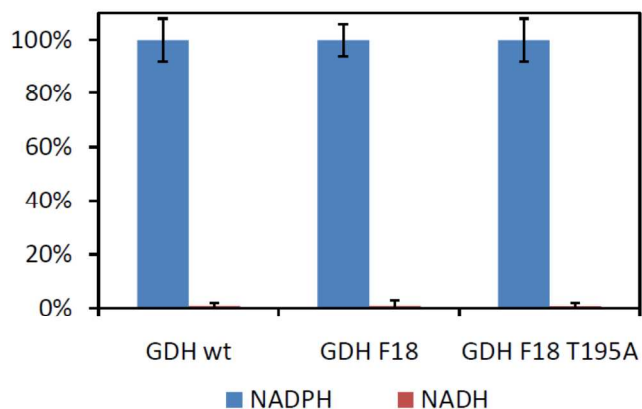


Figure S1 Cofactor specificity of GDH variants. 2-OPBA was the keto acid substrate used in the enzyme assays. The error bars represent the standard deviation of three repeats. (n=3)

Performance of the second generation GDH variant, F18 T195A, in keto acid feeding experiments

To test the enzyme's performance in converting the keto acid 2-OPBA to L-homophenylalanine inside the cells, GDH F18 T195A was overexpressed in *E. coli* and the keto acid feeding experiments were performed as described in the "methods" session. The results showed that F18 T195A produced higher level of L-homophenylalanine than the first generation variant F18. The increase was relatively small (~10%) but significant.

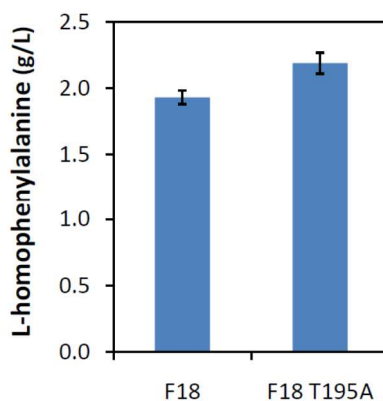


Figure S2 Performance of F18 T195A GDH variant in the keto acid feeding experiments. The error bars represent the standard deviation of three repeats. (n=3)