

Table S1 Strains and plasmids used in this study.

	Description	Source
Plasmids		
pEZ15Asp	Shuttle vector contains <i>Z. mobilis</i> origin and <i>E. coli</i> origin p15A; Biobrick-compatible, <i>Spe</i> ^R	Yang et al., 2016b
pEZ-Pt	pEZ15A containing a <i>phaCAB</i> operon from <i>C. necator</i> H16 driven by inducible promoter <i>Ptet</i> , <i>Spe</i> ^R	This work
pEZ-Pg	pEZ15A containing <i>phaCAB</i> operon driven by <i>Pgap</i> , <i>Spe</i> ^R	This work
pEZ-Pg ^{N1}	pEZ15A containing <i>ZMO1329 (ppnK)</i> gene and <i>phaCAB</i> operon driven by promoter <i>Pgap</i> , <i>Spe</i> ^R	This work
pEZ-Pg ^{N2}	pEZ15A containing <i>ZMO0367 (zwf)</i> gene and <i>phaCAB</i> operon driven by promoter <i>Pgap</i> , <i>Spe</i> ^R	This work
pE39p-Pe ^{EUP}	pEZ15A containing <i>ada</i> from <i>Dickeya zaeae</i> and <i>adh2</i> gene from <i>Saccharomyces cerevisiae</i> driven by stronger promoter <i>Peno</i> , <i>Kana</i> ^R	This work
pL2R-Pt	The <i>Ptet</i> -driven <i>phaCAB</i> operon was inserted into ZM4 chromosome <i>ZMO0038</i> using pL2R editing plasmid, <i>Spe</i> ^R	This work
pL2R-Pg	The <i>Pgap</i> -driven <i>phaCAB</i> operon was inserted into ZM4 chromosome <i>ZMO0038</i> using pL2R editing plasmid, <i>Spe</i> ^R	This work
pL2R-Flo	pL2R with the deletion of a nucleotide thymine in <i>ZMO1082</i> result the integration of <i>ZMO1082</i> with <i>ZMO1083</i> , <i>Spe</i> ^R	This work
Strains		
DH5 α	<i>E. coli</i> for plasmid construction	Lab stock
Trans110	<i>E. coli</i> for plasmid demethylation	Lab stock
ZM4	<i>Zymomonas mobilis</i> subsp. <i>mobilis</i> ZM4 strain	Lab stock
ZM4Pt	<i>Z. mobilis</i> ZM4 containing plasmid pEZ-Pt, <i>Spe</i> ^R	This work
ZM4Pg	<i>Z. mobilis</i> ZM4 containing plasmid pEZ-Pg, <i>Spe</i> ^R	This work
ZM4Pg ^{N1}	<i>Z. mobilis</i> ZM4 containing plasmid pEZ-Pg ^{N1} , <i>Spe</i> ^R	This work
ZM4Pg ^{N2}	<i>Z. mobilis</i> ZM4 containing plasmid pEZ-Pg ^{N2} , <i>Spe</i> ^R	This work
ZMPt	<i>Ptet</i> -driven <i>phaCAB</i> operon inserted into ZM4 chromosome	This work
ZMPg	<i>Pgap</i> -driven <i>phaCAB</i> operon inserted into ZM4 chromosome	This work
ZMPt ^g	ZMPt containing plasmid pEZ-Pg, <i>Spe</i> ^R	This work
ZMPg ^g	ZMPg containing plasmid pEZ-Pg, <i>Spe</i> ^R	This work
ZMPt ^{N1}	ZMPt containing plasmid pEZ-Pg ^{N1} , <i>Spe</i> ^R	This work
ZMPt ^{N2}	ZMPt containing plasmid pEZ-Pg ^{N2} , <i>Spe</i> ^R	This work
ZMPt ^{EUP}	ZMPt containing plasmid pE39p-Pe ^{EUP} , <i>Kana</i> ^R	This work
ZMPt ^{N1-EUP}	ZMPt containing plasmids pEZ-Pg ^{N1} and pE39p-Pe ^{EUP} , <i>Spe</i> ^{R, <i>Kana</i>^R}	This work
ZMPt ^{N2-EUP}	ZMPt containing plasmids ZM4Pg ^{N2} and pE39p-Pe ^{EUP} , <i>Spe</i> ^{R, <i>Kana</i>^R}	This work
ZMPt-Flo	Deletion of a nucleotide thymine in <i>ZMO1082</i> resulting in the integration of <i>ZMO1082</i> with <i>ZMO1083</i>	This work
ZMPt-Flo ^{N1}	ZMPt-Flo containing plasmid pEZ-Pg ^{N1} , <i>Spe</i> ^R	This work
ZMPt-Flo ^{N2}	ZMPt-Flo containing plasmid pEZ-Pg ^{N2} , <i>Spe</i> ^R	This work
ZMPt-Flo ^{N1-EUP}	ZMPt-Flo containing pEZ-Pg ^{N1} and pE39p-Pe ^{EUP} , <i>Spe</i> ^{R, <i>Kana</i>^R}	This work
ZMPt-Flo ^{N2-EUP}	ZMPt-Flo containing pEZ-Pg ^{N2} and pE39p-Pe ^{EUP} , <i>Spe</i> ^{R, <i>Kana</i>^R}	This work

Table S2 Primers used in this study. Font underlined indicates homology arms.

Name	Sequence (5'—3')	Usage
Primers for construction of PHB		
Ptet-CAB-F	<u>GATCTCCCGGATCC</u> ATGGCCACCGGCAAAG	Clone and expression
Pgap-CAB-F	<u>CTTATAAGTTAGGAGAATAAAC</u> ATGGCCACCGGCAAAG	Clone and expression
phaC-R	<u>CTTACTTTCTCTAGATTATGGT</u> TAGGCTTAGCTTAACATAACGACCAG	Clone and expression
phaA-F	<u>CCATAATCTAGAGAAAG</u> TAAGCACATGACCGATTTGTCATTGTCTCTG	Clone and expression
phaA-R	<u>GTGCTTACTTTCTCTAGATTATGGT</u> TATTGCGTTAACCGGCCAAAG	Clone and expression
phaB-F	<u>CTAGAGAAAGTAAGC</u> CACATGACCCAACGTATTGCCTATC	Clone and expression
phaB-R	<u>GGCCGCTACTAGTT</u> AACCCATATGCAAGCCACCATT	Clone and expression
ppnK-R	<u>TCCTTCTCCTCTTTAAC</u> ATAAGCGAAATTGTTCGCG	Clone and expression
ppnK-phaC-F	<u>GTTAAAAAGAGGAGAAAGGATCTCC</u> ATGGCCACCGGCAAAG	Clone and expression
zwf-F	ATGACAAATACCGTTCGACGATG	Clone and expression
zwf-R	<u>GAGATCCTTCTCCTCTTT</u> CAGTCATAACAGTTACTCCATCAC	Clone and expression
zwf-phaC-F	<u>AAAGAGGAGAAAGGATCTCC</u> CATGGCCACCGGCAAAGG	Clone and expression
Peno-F	TGTCTATACTCCAGTTACTCAATACGTAACAATAATCAGTTATCCTAAC	Clone and expression
Peno-R	<u>ATCGAAACCTTCTTAAAATCTTT</u> AGACG	Clone and expression
Peno-ada-F	<u>AAAGATT</u> TAAGAAAGGTTCGATATGGAACATAGCGTTATTGAACC	Clone and expression
ada-R	<u>CTTCTAGACCCTGTGATT</u> TAAGCAATACGAAACATATCAACCAAAAC	Clone and expression
adh2-F	<u>ATCACAGGGTCTAGAAGGAGGTC</u> GAATGAGCATTCCGGAAACCC	Clone and expression
adh2-R	<u>CGAGATCTATGGGACGT</u> TATTAGAGGTATCAACACATAACGACC	Clone and expression
Primers for gene editing		
0038-gr1-F	<u>GAAAGCGTCCAGCAA</u> AATACGCCCTTATTGATGAA	0038gRNA-F
0038-gr1-R	GAAC <u>TTCAATAGAAGGCGT</u> ATTTGCTGGACGC	0038gRNA-R
0038-up-F	<u>CACCA</u> GCTACCGTCTGCTTTGCCAACAAAGCG	Generate pL2R- <i>phaCAB</i> -0038
0038-up-R	TCACGCCCGACGCCAGACGGGATTAGAAATTGTGCG	Generate pL2R- <i>phaCAB</i> -0038
0038-up-Ptet-F	<u>GGCGTCGGCGTGAT</u> TAAGACCCACTTCACATTAAAGTTGTTCTAAC	Generate pL2R- <i>phaCAB</i> -0038
phab-down-R	<u>CGTCTATCTGAAT</u> TTAACGATTAACCCATATGCAAGCCACC	Generate pL2R- <i>phaCAB</i> -0038
0038-down-F	<u>TCGTTAA</u> ATATTCAAGATAGACGGAGATAATAACGGGAGAGAGGTG	Generate pL2R- <i>phaCAB</i> -0038
0038-down-R	<u>GCTCGAGATCTGAT</u> ATCAACTAACAGATCAACC	Generate pL2R- <i>phaCAB</i> -0038
0038-up-Pgap-F	<u>GGCGTCGGCGTGAG</u> TTCGATCAACAAACCGAATCCTATC	Generate pL2R- <i>phaCAB</i> -0038
Flo-gr-F	<u>GAAAGCTTTATGGTGGTTGCTG</u> TTCCGCTACCGCT	FlogRNA-F
Flo-gr-R	GAAC <u>AGCGGTAGCGGAACAGCAACCACCATAAGAGC</u>	FlogRNA-R
Flo-up-F	<u>ACCA</u> GCTACCGTCTTAACCTCATATCGCGTACAAGAAGAAG	Generate pL2R-Flo-401
Flo-up-R	<u>AGCAACCACCATAAGAGC</u> GTGCAACGGTAATCAAGCAAAGCAATG	Generate pL2R-Flo-401
Flo-down-F	GCTTTATGGTGGTTGCTGTTCCG	Generate pL2R-Flo-401
Flo-down-R	<u>GCTCGAGATCTGAT</u> ATCACTGTACAGTCAAAATGCAGACTAAC	Generate pL2R-Flo-401
Primers for gene detection		
15A-fwd	GGCAAAGCCACCTATTAG	Plasmid detection
15A-rev	CACTTCACTGACACCCCTCAT	Plasmid detection

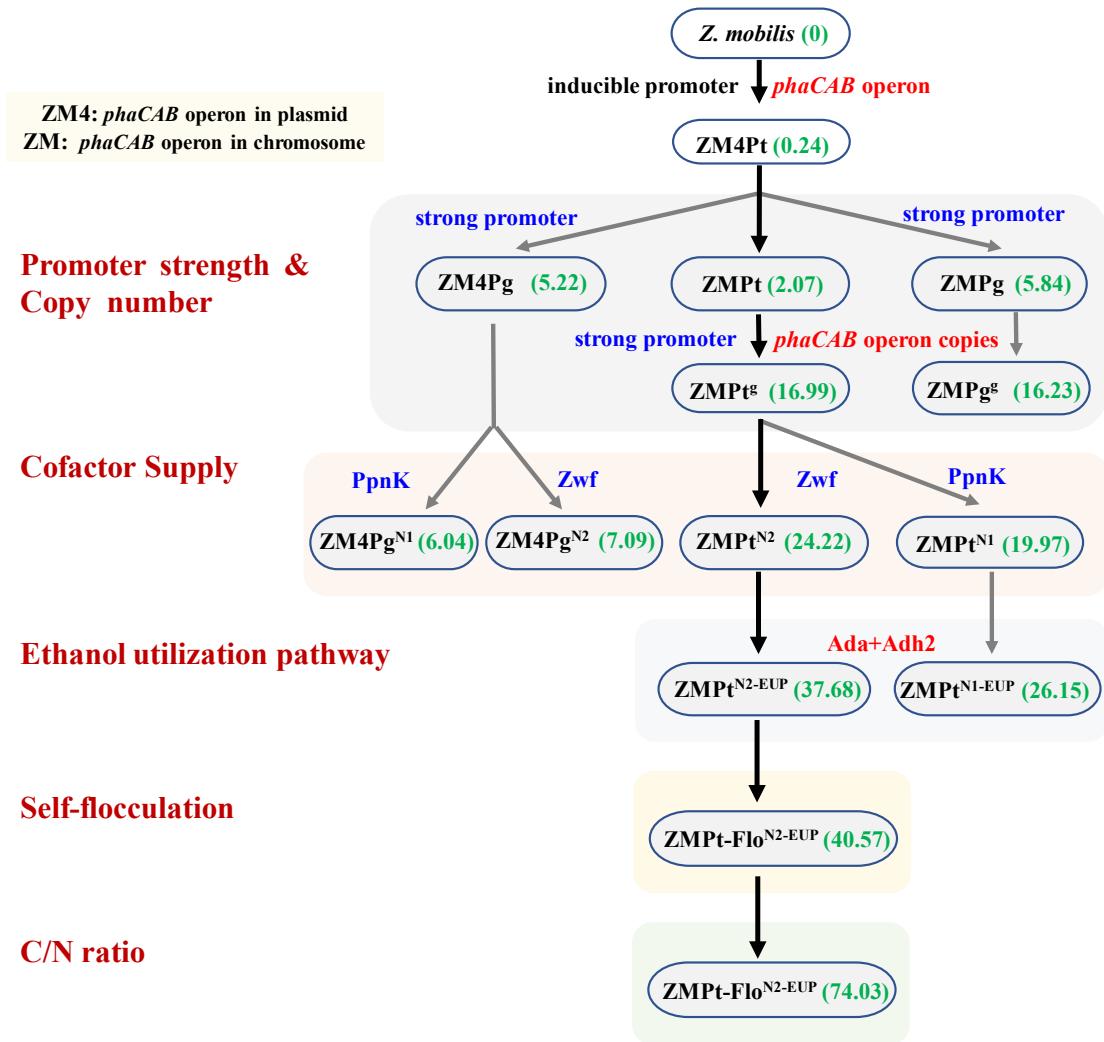


Fig S1. Overview of the strain constructed in *Z. mobilis* for PHB production. Native genes overexpressed or promoter replacement are shown in blue font, while heterologous genes are shown in red font. The PHB content of dry cell weight (%DCW) in recombinant *Z. mobilis* are shown in green font.