

Supporting Information

Enhancing Production of Medium-chain-length Polyhydroxyalkanoates from *Pseudomonas* sp. SG4502 by *tac* Enhancer Insertion

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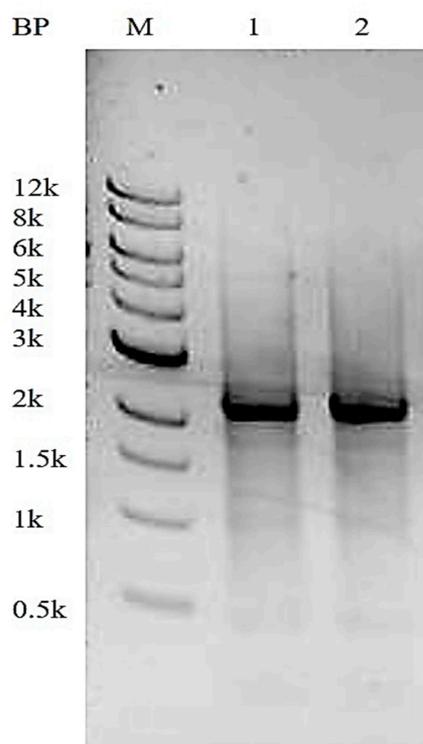


Figure S1. Electrophoretic results of target fragment *C1ZC2* M: DNA Marker GsDL10001, 1,2: PCR product of *C1ZC2* gene.

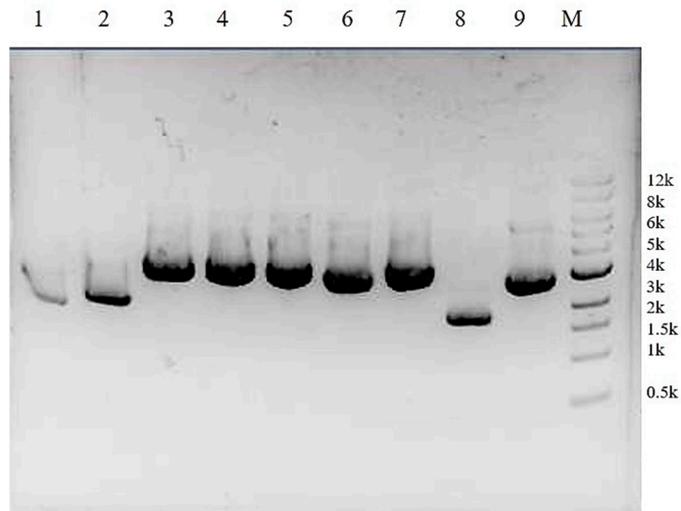


Figure S2. Electrophoretic results of recombinant plasmid pUC19-C1ZC2. 1: pUC19 plasmid; 2-9: recombinant plasmid pUC19-C1ZC2; M: DNA Marker GsDL10001.

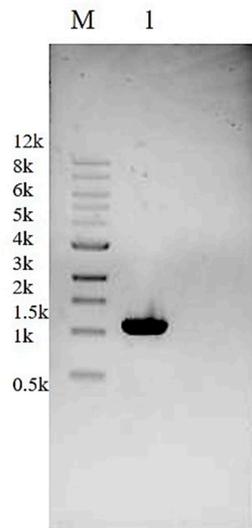


Figure S3. PCR electrophoresis results of *smr* gene. M: DNA Marker GsDL10001; 1: PCR product of *smr* gene.

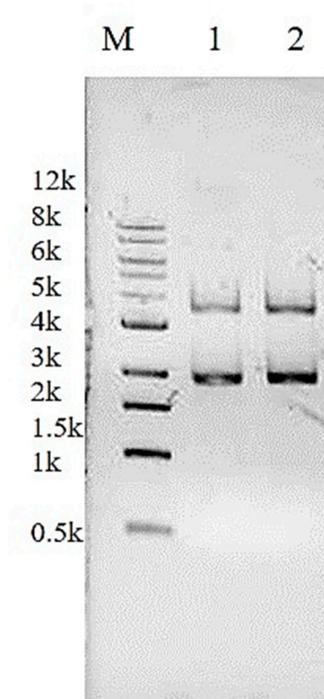


Figure S4. Double enzyme digestion electrophoresis results of recombinant plasmid pUC19-*C1ZC2-smr*. M: DNA Marker GsDL10001; 1,2: Double enzyme digestion products of recombinant plasmid pUC19-*C1ZC2-smr*.

Range 1: 107 to 898 [Graphics](#) [Next Match](#) [Previous Match](#)

Score	Expect	Identities	Gaps	Strand
1463 bits(792)	0.0	792/792(100%)	0/792(0%)	Plus/Minus
Query 1	TTATTTGCCGACTACCTTGGTGATCTCGCCCTTTCACGTAGTGGCAAAATCTTCCAAC	60		
Sbjct 898	TTATTTGCCGACTACCTTGGTGATCTCGCCCTTTCACGTAGTGGCAAAATCTTCCAAC	839		
Query 61	ATCTGCGCGAGGCCAAGCGATCTTCTTTGTCCAAAGTAAGCCGTGTAGCTTCAAG	120		
Sbjct 838	ATCTGCGCGAGGCCAAGCGATCTTCTTTGTCCAAAGTAAGCCGTGTAGCTTCAAG	779		
Query 121	TATGACGGGCTGATACTGGCCCGCAGGCGCTCCATTGCCAGTCGGCAGCGACATCCTT	180		
Sbjct 778	TATGACGGGCTGATACTGGCCCGCAGGCGCTCCATTGCCAGTCGGCAGCGACATCCTT	719		
Query 181	CGGCGCGATTTGCCGGTTACTGCGCTGTACCAATGCCGGACAACTAAGCACTACATT	240		
Sbjct 718	CGGCGCGATTTGCCGGTTACTGCGCTGTACCAATGCCGGACAACTAAGCACTACATT	659		
Query 241	TCGCTCATCGCAGCCAGTCGGGCGCGAGTTCCTTAGCGTTAAGGTTTCATTAGCGC	300		
Sbjct 658	TCGCTCATCGCAGCCAGTCGGGCGCGAGTTCCTTAGCGTTAAGGTTTCATTAGCGC	599		
Query 301	CTCAATAGATCCTGTTCAAGAACCGGATCAAGAGTTCCTCCGCCGTGGACCTACCAA	360		
Sbjct 598	CTCAATAGATCCTGTTCAAGAACCGGATCAAGAGTTCCTCCGCCGTGGACCTACCAA	539		
Query 361	GGCAACGCTATGTTCTTCTGCTTTTGTGACGAGATAGCCAGATCAATGTCGATCGTGGC	420		
Sbjct 538	GGCAACGCTATGTTCTTCTGCTTTTGTGACGAGATAGCCAGATCAATGTCGATCGTGGC	479		
Query 421	TGGCTCGAAGTACCTGCAAGAAATGTCATTGGCGTCCATTCTCAAATTCAGTTCGCG	480		
Sbjct 478	TGGCTCGAAGTACCTGCAAGAAATGTCATTGGCGTCCATTCTCAAATTCAGTTCGCG	419		
Query 481	CTTAGCTGGATAACCCACGGAAATGATGTCGTGTCACAAATGGTACTTCTACAGC	540		
Sbjct 418	CTTAGCTGGATAACCCACGGAAATGATGTCGTGTCACAAATGGTACTTCTACAGC	359		
Query 541	GCGGAGAATCTGCTCTCTCCAGGGGAAGCCGAAGTTTCCAAAAGGTCGTTGATCAAAGC	600		
Sbjct 358	GCGGAGAATCTGCTCTCTCCAGGGGAAGCCGAAGTTTCCAAAAGGTCGTTGATCAAAGC	299		
Query 601	TGCGCGGTTGTTTCATCAAGCCTTACGTCACCGTAACCAAGCAATCAATACACTGTG	660		
Sbjct 298	TGCGCGGTTGTTTCATCAAGCCTTACGTCACCGTAACCAAGCAATCAATACACTGTG	239		
Query 661	TGGCTTCAGGCGCCATCCACTGGCGAGCGTACAAAATGTACGGCCAGCAAGCTCGGTT	720		
Sbjct 238	TGGCTTCAGGCGCCATCCACTGGCGAGCGTACAAAATGTACGGCCAGCAAGCTCGGTT	179		
Query 721	GAGATGGCGCTGATGACGCCCAACTCTGTAGTTGAGTCGATACTTCGGCGATCAC	780		
Sbjct 178	GAGATGGCGCTGATGACGCCCAACTCTGTAGTTGAGTCGATACTTCGGCGATCAC	119		
Query 781	CGCTTCCCTCAT 792			
Sbjct 118	CGCTTCCCTCAT 107			

Figure S5. The sequencing results of recombinant plasmid pUC19-*C1ZC2-smr*. (the accession number of the nucleotide sequences was GenBank: AB448740.1).

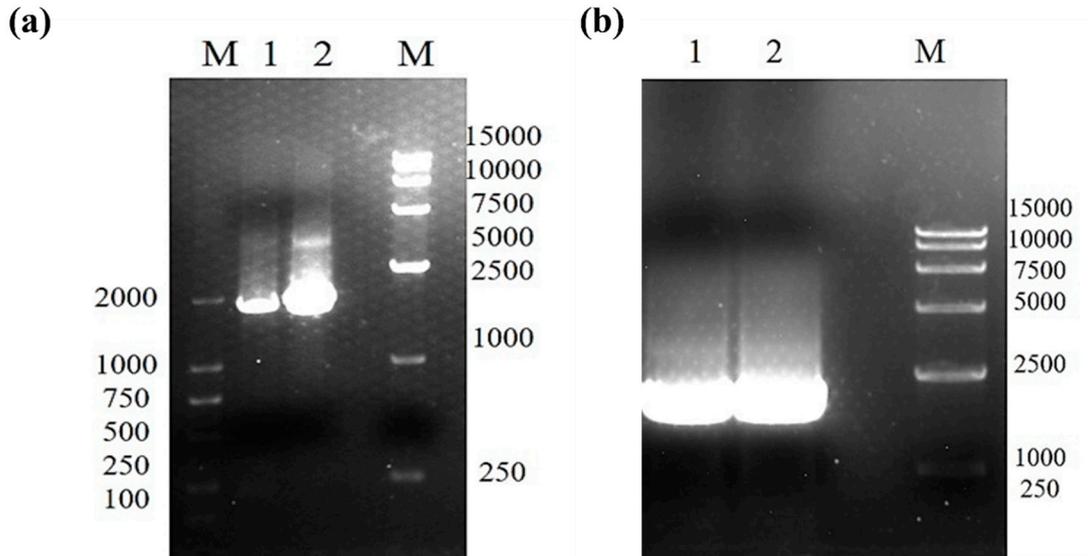


Figure S6. Electrophoretic target genes map. (a) *phaC1*, the left M: DNA Marker DL2000, 1-2: the PCR product of *phaC1*, the right M: DNA Marker DL15000; (b) *phaC2*, 1-2: the PCR product of *phaC2*, M: DNA Marker DL15000.

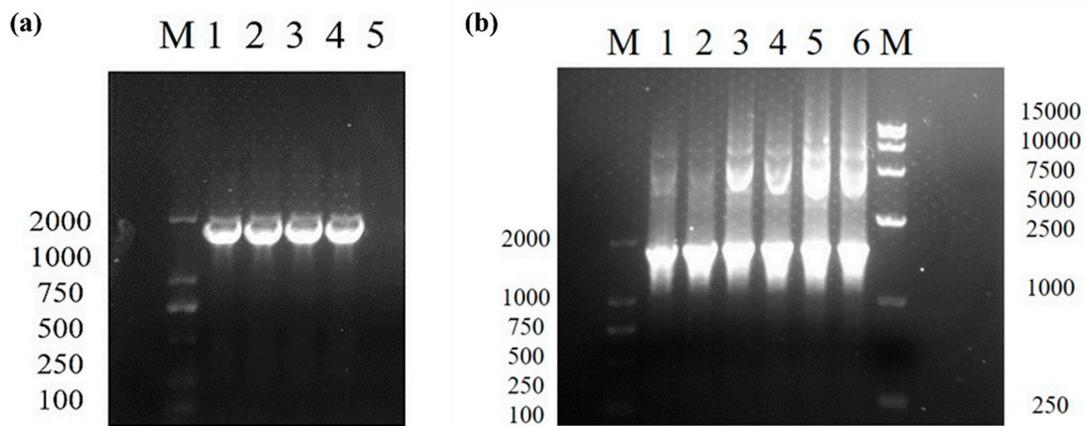


Figure S7. Electrophoresis results of PCR identification. (a) Recombinant plasmid *pk18-phaC1*, M: DNA Marker DL2000, 1: PCR product of *phaC1* gene, 2-4: PCR product of *pk18-phaC1*, 5: PCR product of *pk18* empty vector; (b) Recombinant plasmid *pk18-phaC2*, M: DNA Marker DL2000, 1: PCR product of *phaC2* gene, 2-6: PCR product of *pk18-phaC2*, M: DNA Marker DL15000.

(a)

Score	Expect	Method	Identities	Positives	Gaps
1150 bits(2976)	0.0	Compositional matrix adjust	559/559(100%)	559/559(100%)	0/559(0%)

Query 1 MNKIAEDLQRQASEHTLSLFPVVLGRGKDLLSSTRQVLQALRQFLNSTRVHAFPGVQLK 60
 Sbjct 1 MNKIAEDLQRQASEHTLSLFPVVLGRGKDLLSSTRQVLQALRQFLNSTRVHAFPGVQLK 60

Query 61 NVLLQADLRFEDGRRFADFAWSHNPLTKRYMQLYLAWRQELDWDIENHLPQDISRG 120
 Sbjct 61 NVLLQADLRFEDGRRFADFAWSHNPLTKRYMQLYLAWRQELDWDIENHLPQDISRG 120

Query 121 HFVILNLTALAFSLSLANPAAKRRFFETGGKSLDGLGHLAKLDVVRGGFLSQVNEAF 180
 Sbjct 121 HFVILNLTALAFSLSLANPAAKRRFFETGGKSLDGLGHLAKLDVVRGGFLSQVNEAF 180

Query 181 EVGKRLALTEGAVVFRNDLLELQVRFNTEQVHARPLLVPQIKRFVFDLSPDSLVR 240
 Sbjct 181 EVGKRLALTEGAVVFRNDLLELQVRFNTEQVHARPLLVPQIKRFVFDLSPDSLVR 240

Query 241 FALRSGLQTFILSVRNPTKQREWGLSTYIEALKEAVEAVLAIQSTDLNMLGACSGGIT 300
 Sbjct 241 FALRSGLQTFILSVRNPTKQREWGLSTYIEALKEAVEAVLAIQSTDLNMLGACSGGIT 300

Query 301 TAALLGHTAARGEQPIHALTLLVSVLDTETQPSLFDVEQTEAAKRSEYQAGVLEGRN 360
 Sbjct 301 TAALLGHTAARGEQPIHALTLLVSVLDTETQPSLFDVEQTEAAKRSEYQAGVLEGRN 360

Query 361 LAKLFAWRFNDLIRWYVWVNYLLGRQFPADLWVWVNDITRLPATLHGLIELFKTNPL 420
 Sbjct 361 LAKLFAWRFNDLIRWYVWVNYLLGRQFPADLWVWVNDITRLPATLHGLIELFKTNPL 420

Query 421 FRPGALEVCQTFIDLKQVSDLCVAVGVDHITPWEACTRSARLFGGSEFVLSRSGHIQ 480
 Sbjct 421 FRPGALEVCQTFIDLKQVSDLCVAVGVDHITPWEACTRSARLFGGSEFVLSRSGHIQ 480

Query 481 AILNFPQWPKARPHIYNGELPTEPKAWQENATKQIDSWLWQWQLIERSGPIKAPGKL 540
 Sbjct 481 AILNFPQWPKARPHIYNGELPTEPKAWQENATKQIDSWLWQWQLIERSGPIKAPGKL 540

Query 541 GNRQVFSGEAAPGVYHER 559
 Sbjct 541 GNRQVFSGEAAPGVYHER 559

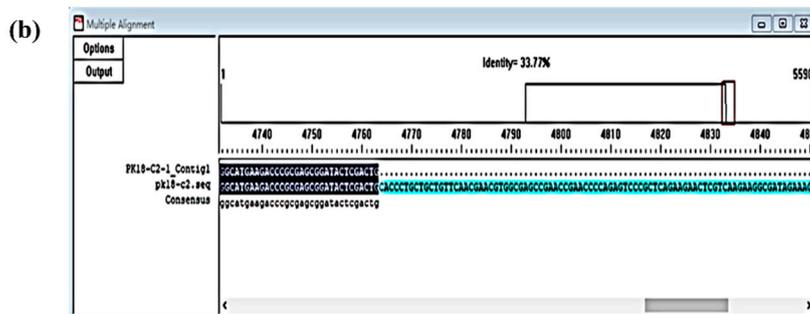


Figure S8. Comparison of sequencing results of recombinant plasmid (a) pK18-*phaC1*, (b) pK18-*phaC2*.

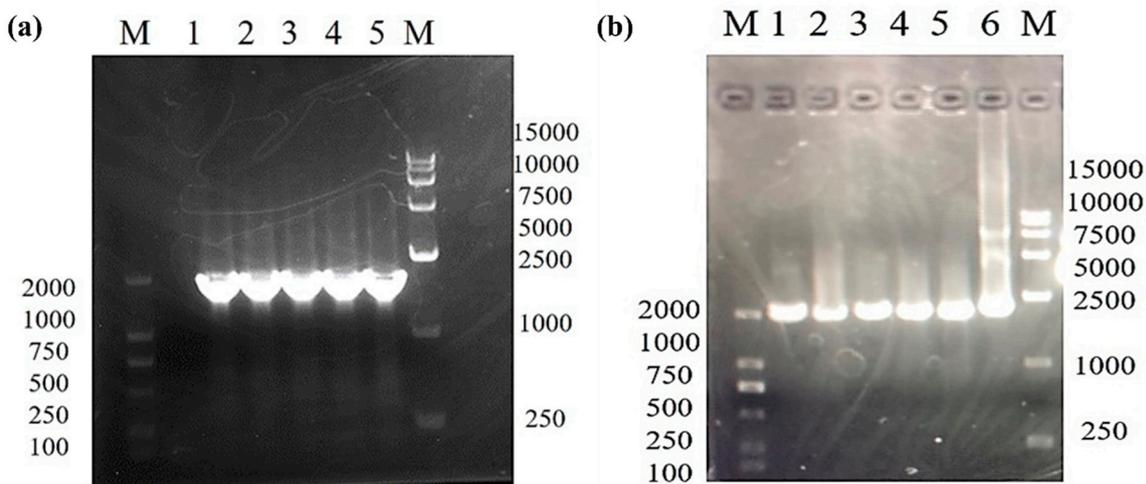


Figure S9. Electrophoresis results of PCR identification. (a) *Pseudomonas* sp. SG4502+ *tac-phaC1*, M: DNA Marker DL2000, 1-5: PCR product of *Pseudomonas* sp. SG4502+*tac-phaC1*, M: DNA Marker DL15000; (b) *Pseudomonas* sp. SG4502+ *tac-phaC2*, M: DNA Marker DL2000, 1-6: PCR product of *Pseudomonas* sp. SG4502+*tac-phaC2*, M: DNA Marker DL15000.

Table S1. Strains and plasmids used in this study.

Strains and plasmids	Related Information	Source
Strains		
<i>E. coli</i> DH5 α	Used for the construction of overexpressed plasmids; <i>F</i> -, ϕ 80 <i>dlacZAM15</i> Δ (<i>lacZYA-arg</i>) <i>U169</i> , <i>deoR</i> , <i>recA1</i> , <i>endA1</i> , <i>hsdR17</i> (<i>rK</i> +, <i>mk</i> +), <i>phoA</i> , <i>supE44</i> , λ -, <i>thi-1</i> , <i>gyrA96</i> , <i>relA1</i>	This laboratory
<i>E. coli</i> JM109	Construction for knocking out plasmids; <i>recA1</i> , <i>endA1</i> , <i>gyrA96</i> , <i>thi-1</i> , <i>hidR17</i> , <i>supE44</i> , <i>relA1</i> , Δ (<i>lac</i> - <i>proAB</i>)/ <i>F'</i> [<i>traD36</i> , <i>proAB</i> +, <i>lac 1q</i> , <i>lacZAM15</i>]	This work
Plasmids		
pK18	Starting plasmid containing strong promoter <i>tac</i> , Used as a gene booster; Km ^R	This work
pUC19	Starting plasmid, used as gene knockout; Ap ^R	This laboratory
pCDFDuet-1	The starting plasmid, which is used to provide the <i>Smr</i> gene required for knockout; Sm ^R	This laboratory
pK18- <i>phaC1</i>	Recombinant plasmids for overexpression of the <i>phaC1</i> gene; Km ^R	This work
pK18- <i>phaC2</i>	Recombinant plasmids for overexpression of the <i>phaC1</i> gene; Km ^R	This work
pUC19- <i>C1ZC2</i>	Recombinant plasmids, knocked out intermediate plasmids containing the <i>phaC1-phaZ-phaC2</i> gene; Ap ^R ; Sm ^R	This work
pUC19- <i>C1ZC2-smr</i>	Recombinant plasmid, knockout intermediate plasmid contain- ing the <i>phaC1-phaZR-smr-phaZF-phaC2</i> gene; Ap ^R ; Sm ^R	This work

Table S2. Primers used in this study.

Primers	Description	Source
phaC1-F	CGGATCCCCGGGTACCGGGACAACGGAGCGTCGTAG	This work
phaC1-R	ACGAATTCGAGCTCGGCGGAACACGAAGGGGCTGGG	This work
phaC2-F	CTTGGTCGGTCATTTGGCAATCTGCAGCAGGCAGTC	This work
phaC2-R	GGACTCTGGGGTTTCGGTTCGGCTCGCCACGTTTCGTT	This work
tac-phaC1-F	GATAAGCCCGGATCCCCGGGTACCGGGACA	This work
tac-phaC1-R	ACGAATTCGAGCTCGGCGGAACACGAAGGGGCTGGG	This work
tac-phaC2-F	CGGTCATTTGGCAATCTGCAGCAGG	This work
tac-phaC2-R	TCGCTCTCGTCGATCCACT	This work
q-phaC1-F	TGTTCCGCAACGACCTGCTA	This work
q-phaC1-R	GAATCGCACCAGGCTCTTGTC	This work
q-phaC2-F	CGTCCGTTTCGCCGATCCCA	This work
q-phaC2-R	TCGCTCTCGTCGATCCACT	This work
q-phaZ-F	CCTTCGTGTTCCGCACCAT	This work
q-phaZ-R	ACGTCGAAGGCATCACCTC	This work
pC1ZC2-F	GACGGCCAGTGAATTGGGAACAACGACACCACGCGCCTGCC	This work
pC1ZC2-R	TGATTACGCCAAGCTAGCAGGTCGTCGATCAGGTGGCGCAG	This work
Smr-F	GACCCTGCGCACCGCGACATAAGCGGCTATTTAACGACCC	This work
Smr-R	TGGAGGTAGCGCCGCGCCGAGTCTCACGCCCGGAGCGTA	This work