

TABLE S1 Strains, plasmids, and oligonucleotide primers used in this study

| Strains | Description | Sources |
|----------------------------|---|--|
| <i>C. beijerinckii</i> | | |
| NCIMB 8052 | Wild type | Lab stock |
| <i>pta</i> ::int(17) | Group II intron inserted at 17/18a bp of <i>pta</i> ^a | This study |
| <i>buk</i> ::int(532) | Group II intron inserted at 532/533s bp of <i>buk</i> ^b | This study |
| <i>E. coli</i> | | |
| Turbo | High efficiency chemically competent cell | NEB ^c |
| Top10 | Electrocomp TM competent cell | Invitrogen ^d |
| Plasmids | Description & Relevant characteristics | References |
| pTJ1 | Ap ^r , Em ^r , <i>C. beijerinckii</i> - <i>E. coli</i> shuttle vector ^e | (1) |
| pJIR750ai | L1.LtrB-ΔORF intron and ltrA, targeting <i>C.perfringens</i> alpha toxin gene | (2) |
| pCR-XL-TOPO [®] | one-step cloning vector for long PCR products | Invitrogen ^d |
| pYW1 | pTJ1 integrated with intron+ltrA, and <i>ptb</i> promoter | This study |
| pYW1- <i>pta</i> | pYW1 with intron targeting <i>pta</i> | This study |
| pYW1- <i>buk</i> | pYW1 with intron targeting <i>buk</i> | This study |
| Primers | Sequence (5'-3') | Description |
| Intron-F | CCCGGGATAATTATCCTTACCAGCCCATAGG | XmaI |
| Intron-R | CTTGTAGATATGACGACAGGAAGAGT | |
| <i>ptb</i> -p-F | GTATATCAAGCTTGAAGATATATTATATTACGTTC GTGTTGTGA | HindIII |
| <i>ptb</i> -p-R | GTATATCACTAGTAATCAATGCTATGAATATTTCT TTATACCTT | SpeI |
| <i>pta</i> -17/18a-IBS | AAAACCCGGGATAATTATCCTTAGCAGTCCATA TTGTGCGCCAGATAGGGTG | XmaI |
| <i>pta</i> -17/18a-EBS1d | CAGATTGTACAAATGTGGTGATAACAGATAAGT CCATATTTTTAACTTACCTTTCTTTGT | BsrGI |
| <i>pta</i> -17/18a-EBS2 | TGAACGCAAGTTTCTAATTTTCGATTACTGCTCG ATAGAGGAAAGTGTCT | |
| EBS universal | CGAAATTAGAACTTGCGTTCAGTAAAC | Universal primer for knockout re-targeting |
| <i>pta</i> -17/18a-F | TTGTTATTTTACAGGAGTATTGTCTGA | Confirmation of knockout for <i>pta</i> |
| <i>pta</i> -17/18a-R | TCTTTCCTCGTTTCCCTCTG | Confirmation of knockout for <i>pta</i> |
| <i>buk</i> -532/533s-IBS | AAAACCCGGGATAATTATCCTTAATGAACACCT AAGTGCGCCAGATAGGGTG | XmaI |
| <i>buk</i> -532/533s-EBS1d | CAGATTGTACAAATGTGGTGATAACAGATAAGT CACCTAAATTAACCTTTCTTTGT | BsrGI |
| <i>buk</i> -532/533s-EBS2 | TGAACGCAAGTTTCTAATTTTCGGTTTTTCATCCG ATAGAGGAAAGTGTCT | |
| <i>buk</i> -532/533s-F | GCTTCTAACCTTGGCGGAAT | Confirmation of knockout for <i>buk</i> |
| <i>buk</i> -532/533s-R | AACGCCTCCACCCATATGTA | Confirmation of |

| | | |
|-----------------------|-------------------------|--|
| pYW1-cure-F | GCTTTCGTTTCGTTCCCATATA | knockout for <i>buk</i> Detection of plasmid curing |
| pYW1-cure-R | TGAATTGATTGGGCCCTCTA | Detection of plasmid curing |
| Intron-SB-F | CCTATGGGAACGAAACGAAA | Amplification of Southern blotting probe |
| Intron-SB-R | CGAGTACTCCGTACCCTTGC | Amplification of Southern blotting probe |
| <i>buk</i> -qPCR-U-F | TGTTGGTAGAGGTGGAATGCT | q-RT-PCR, upstream of intron insertion |
| <i>buk</i> -qPCR-U-R | GCATGAGGTCCTTGAACCTCC | q-RT-PCR, upstream of intron insertion |
| <i>buk</i> -qPCR-D-F | GGAAAAGGTGGATTTGTTGGT | q-RT-PCR, downstream of intron insertion |
| <i>buk</i> -qPCR-D-R | ACTGGGGCTATCCATTCAACT | q-RT-PCR, downstream of intron insertion |
| <i>bukII</i> -qPCR-F | TAGTTGGAAGAGGTGGCCTTT | q-RT-PCR, for <i>bukII</i> |
| <i>bukII</i> -qPCR-R | CTGGCACTCCTGAAAGTCTTG | q-RT-PCR, for <i>bukII</i> |
| <i>bukIII</i> -qPCR-F | CACAAACGATGCTAGGGTTGTA | q-RT-PCR, for <i>bukIII</i> |
| <i>bukIII</i> -qPCR-R | AGCATTTTCATCTTCACCTGGAT | q-RT-PCR, for <i>bukIII</i> |
| <i>16s</i> -qPCR-F | CCGCTAACGCATTAAGTATTCC | q-RT-PCR, for 16s RNA as endogenous control |
| <i>16s</i> -qPCR-R | CTTAACCCAACATCTCACGACA | q-RT-PCR, for 16s RNA as endogenous control |

^a 17/18a, between 17 and 18 bp in the antisense direction

^b 532/533s, between 532 and 533 bp in the sense direction

^c New England Biolabs Inc., Ipswich, MA

^d Invitrogen Inc., Grand Island, NY

^e Ap, ampicillin; Em, erythromycin

REFERENCES

1. Jesse TW. 2003. Genetic characterization and manipulation of solvent-producing clostridia. PhD dissertation. University of Illinois at Urbana-Champaign, Urbana, IL.
2. Chen Y, McClane BA, Fisher DJ, Rood JI, Gupta P. 2005. Construction of an alpha toxin gene knockout mutant of *Clostridium perfringens* type A by use of a mobile group II intron. *Appl. Environ. Microbiol.* 71:7542-7547.

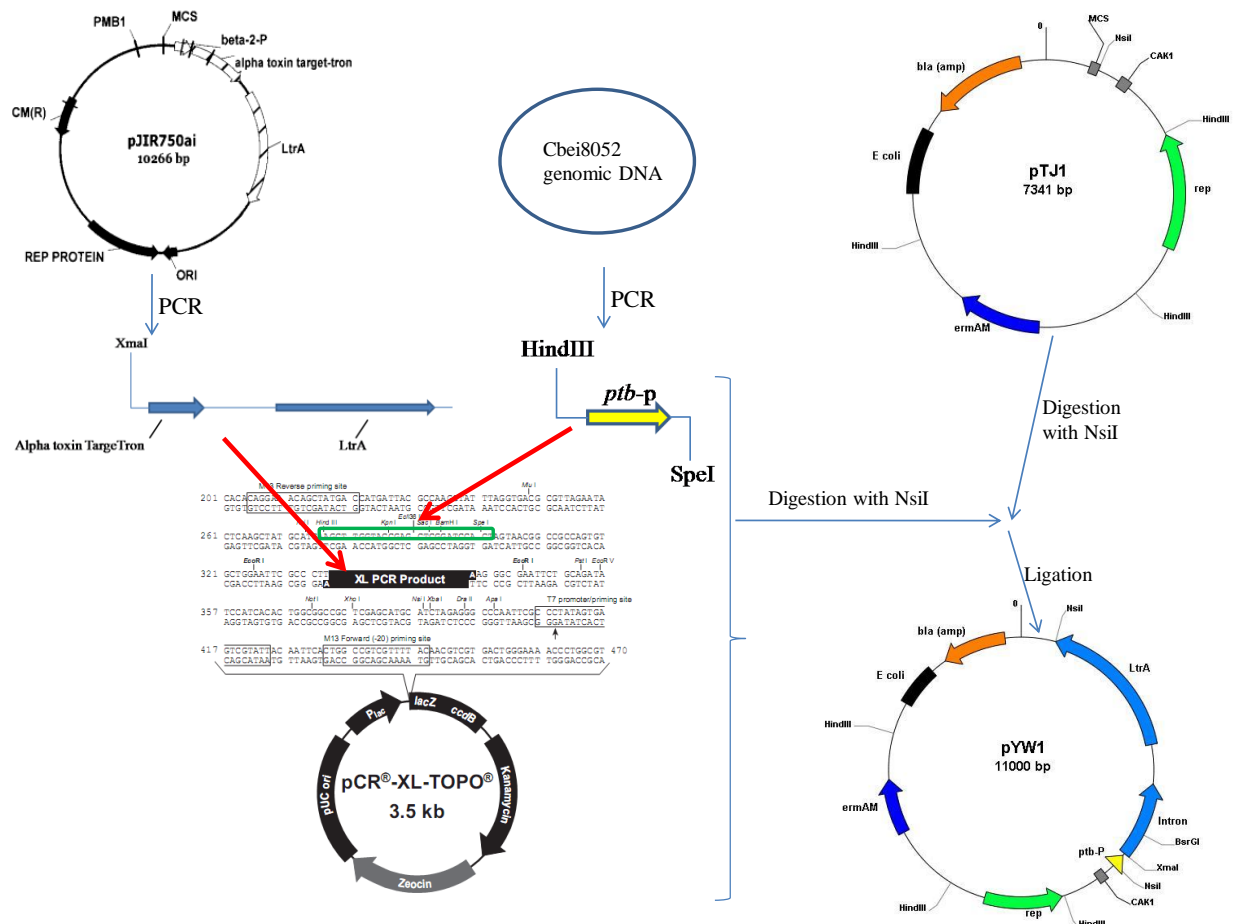


FIG S1 The schematic diagram for construction of the Targetron plasmid pYW1 for gene knockout in *C. beijerinckii*. The DNA fragment containing the LtrB intron and LtrA gene in pJIR750ai (1) was first amplified by PCR, with an *Xma*I restriction enzyme site introduced for later intron re-targeting purposes. The PCR product was cloned into the cloning site of pCR-XL-TOPO[®] vector (Invitrogen, Grand Island, NY). The *ptb* promoter from *C. beijerinckii* 8052 genomic DNA was amplified by PCR with introduction of the *Hind*III and *Spe*I sites into the fragment. The PCR product was integrated into the *Hind*III and *Spe*I sites of the pCR-XL-TOPO[®] vector containing the intron fragment. The resulting cloning vector was digested with *Nsi*I, and ligated into pTJ1 digested with the same enzyme. The plasmid with the insertion at the desired direction was verified by sequencing and named pYW1 for later intron re-targeting and gene disruption.

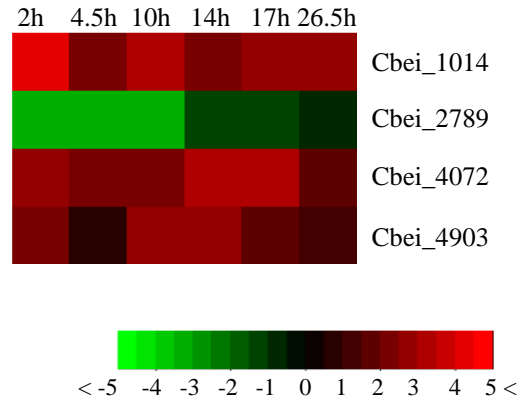


FIG S2 The dynamic transcriptional profiles of lactate dehydrogenase genes over the batch fermentation process based on the RNA-Seq data published earlier (2).

REFERENCES

1. Chen Y, McClane BA, Fisher DJ, Rood JI, Gupta P. 2005. Construction of an alpha toxin gene knockout mutant of *Clostridium perfringens* type A by use of a mobile group II intron. *Appl. Environ. Microbiol.* 71:7542-7547.
2. Wang Y, Li X, Mao Y, Blaschek H. 2012. Genome-wide dynamic transcriptional profiling in *Clostridium beijerinckii* NCIMB 8052 using single-nucleotide resolution RNA-Seq. *BMC Genomics* 13:102.