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Tandem gene duplication selected by activation of horizontally transferred gene in bacteria

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Supplementary figures

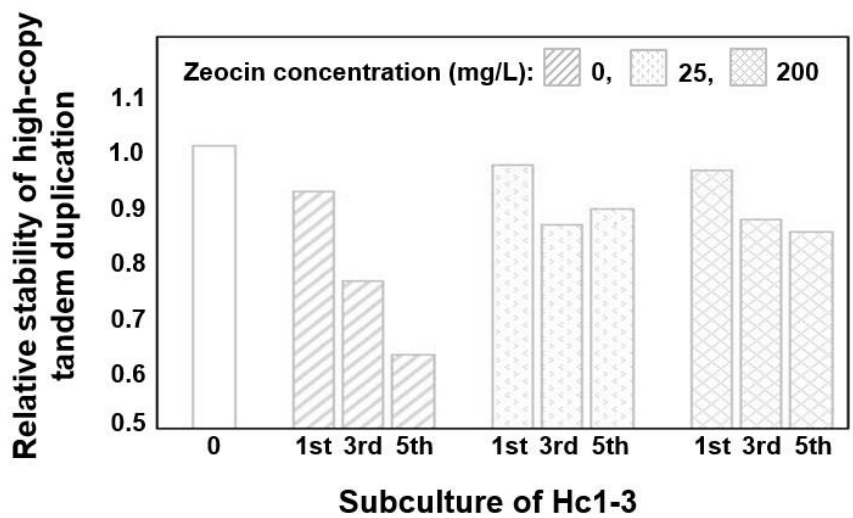


Figure S1 Stability of tandem-duplication copy number of Hc1 during the continuous subculture. Based on Figure 2D, “intensity of high-copy-tandem-duplication band (top) / intensity of plasmid backbone band (bottom)” was used to determine relative stability of high-copy-tandem-duplication [relative to starting culture (0)]. The intensity of visualized band was quantified by software ImageJ.

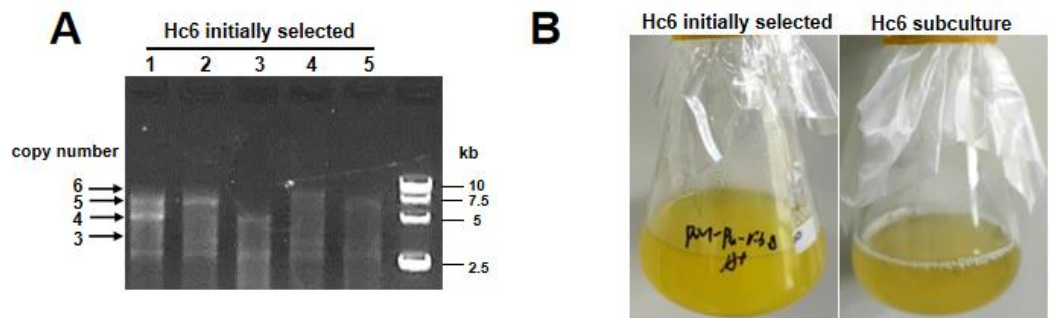


Figure S2 Hc6 initially selected had higher copy number of *ribA* and synthesized more vitamin B2. (A) Restriction analysis of Hc6 initially selected from 200 mg/L zeocin. Plasmid of Hc6 initially selected from 200 mg/L zeocin was restricted by *KpnI* and *AscI*, indicating the copy number of Hc6 initially selected could reach up to 6. (B) Comparison between Hc6 initially selected and its subculture. Under the same circumstance of cultivation, Hc6 initially selected is more yellow than its subculture, indicating vitamin B2 biosynthesis of higher-copy Hc6 strain is more than that of 3-copies Hc6 strain.

Supplementary tables

Table S1 List of oligonucleotides used in this study.

Primers for the vector construction

Target sequence	Forward primer (5'-3')	Reverse primer (5'-3')
<i>Prn</i> promoter	GCTCCCCCGCCGTCGTTCAA	ATGTATATCTCCTTCTTAAA
<i>lac</i> operon left flanking sequence	AGAAAAACCACCCTGGCGCC	GGTGCCTAATGAGTGAGCTA
<i>lac</i> operon right flanking sequence	GTTTCTTTATGGCAGGGTGA	ATTCGGGATTTTCGGCGCTCC
sp110promoter	GAAAATATCAAAAATGTGTT	CTCCAATTATATCATACATT

Primers for the identification of Ev strains

Target sequence	Forward primer (5'-3')	Reverse primer (5'-3')
Ev1	AGTCCTAGGGACTATGCTAGC	ACTCTAGTAGAGAGCGTTCACC
Ev2	ATACCGGCTCCCCCGCCGTCG	ACTCTAGTAGAGAGCGTTCACC
Ev3	GAAAATATCAAAAATGTGTTG	AAAAAAAACGCCCGGCTTTCAC

Primers for the generation of hybridization probes

Gene	Forward primer (5'-3')	Reverse primer (5'-3')
<i>Ble</i>	AGGACGACTTCGCCGGTGTG	CGGCTGCTCGCCGATCTCGGT

Primers for real-time qPCR and qRT-PCR.

Gene	Forward primer (5'-3')	Reverse primer (5'-3')
<i>Ble</i>	AGGACGACTTCGCCGGTGTG	CGGCTGCTCGCCGATCTCGGT
<i>Spec</i>	ATGTTTGGATCAGGAGTTGA	CCACGGTACCATTCTTGCT
<i>bioA</i>	GTGATGCCGAAATGGTTGCC	GCGGTCAGACGCTGCAACTG
<i>GFP</i>	AGCTGACCCTGAAGTTCATCTG	TGTAGTTGTACTCCAGCTTGTG
<i>GAPDH</i>	GAAAGCGAAAGGCGCAGAAA	TTGCCATCCAGAGTGTCGTC

