

The gRNA-miRNA-gRNA ternary cassette combining CRISPR/Cas9 with
RNAi approach strongly inhibits hepatitis B virus replication

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Supplementary materials

Table S1. Oligo-nucleotides sequences for constructing gRNA-miR-HBV-gRNA cassette

Name	Sequences
Pri-miR-31(22+21)/miR-HBV	Top: 5'-TCGAGctgtaactcggaactggagaggGGTGAAGCGAAGTGCAC ACGGgttgaactgggaacgACGTGTGCACATCGATTCACGGCtttctgtctgaca gcagcttG-3' Bottom: 5'-GATCCaagctgctgtcagacaggaaaGCCGTGAATCGATGTG CACACGTcgttcccagttcaacCCGTGTGCACTTCGCTTCACCcctctccagttcc gagttacagC-3'
Pri-miR-31(30+31)/miR-HBV	Top: 5'-TCGAGtattgctcctgtaactcggaactggagaggGGTGAAGCGAAGT GCACACGGgttgaactgggaacgACGTGTGCACATCGATTCACGGCtttctgt tctgacagcagcttggtacctccG-3' Bottom: 5'-GATCCggaggtagccaagctgctgtcagacaggaaaGCCGTGAA TCGATGTGCACACGTcgttcccagttcaacCCGTGTGCACTTCGCTTCACC cctctccagttccgagttacaggagcaataC-3'
gRNA3	Top:5'-CACCGCAAGCCTCCAAGCTGTGCCT-3' Bottom:5'-AAACAGGCACAGCTTGGAGGCTTGC-3'
gRNA4	Top: 5'-CACCGCGAGGGAGTTCTTCTTCTAG-3'

Bottom: 5'-AAACCTAGAAGAAGAAGACTCCCTCGC-3'

Table S2. The synthesized gene sequences for constructing gRNA-miR-HBV-gRNA ternary cassette and gRNA-gRNA binary cassette

Name	Sequences
Pri-miR-31(38+40)/miR-HBV	5'-agaCTCGAGagggatggtattgctcctgtaactcggaactggagaggGGTG AAGCGAAGTGCACACGGgttgaactgggaacgACGTGTGCACATCG ATTCACGGCtttctgtctgacagcagcttggtacctccgctcctgttcGGATCCgag- 3'
Pri-miR-31(38+40)/miR-HBVm	5'-agaCTCGAGagggatggtattgctcctgtaactcggaactggagaggGGTG AAGCGAATGGCATAGCGgttgaactgggaacgAGCTATGCCACATC GATTCACGGCtttctgtctgacagcagcttggtacctccgctcctgttcGGATCCgag -3'
Pri-miR-31(51+51)/miR-HBV-gRNA1	5'-agaCTCGAGcataacaacgaagaggatggtattgctcctgtaactcggaact ggagaggGGTGAAGCGAAGTGCACACGGgttgaactgggaacgACGTG TGCACATCGATTACGGCtttctgtctgacagcagcttggtacctccgctcctgttc ctccttgtcttGGATCCgagCCTGCTGGTGGCTCCAGTTCGTTTTAGA GCTAGAAATAGCAAGTTAAAATAAGGCTAGTCCGTTATCAA CTTGAAAAAGTGGCACCGAGTCGGTGCtttttaccAAGCTTgca-3'
Pri-miR-31(51+51)/miR-HBV-gRNA2	5'-agaCTCGAGcataacaacgaagaggatggtattgctcctgtaactcggaact ggagaggGGTGAAGCGAAGTGCACACGGgttgaactgggaacgACGTG TGCACATCGATTACGGCtttctgtctgacagcagcttggtacctccgctcctgttc ctccttgtcttGGATCCgagGGTTGCGTCAGCAAACACTGTTTTAGA GCTAGAAATAGCAAGTTAAAATAAGGCTAGTCCGTTATCAA CTTGAAAAAGTGGCACCGAGTCGGTGCtttttaccAAGCTTgca-3'
U6-gRNA2	5'-agaCTCGAGtttttgaggccatttcccatgattccttcatattgcatatacagata caaggctgttagagagataattggaattaattgactgtaaacacaaagatattagtacaaaatacgtg acgtagaaagtaataatttctgggtagttgacgttttaaattatgttttaaattggactatcatatgctt accgtaactgaaagtatttcgatttctggctttatatatcttGTGGAAAGGACGAAAC ACCGTTGCGTCAGCAAACACTGTTTTAGAGCTAGAAATAG

CAAGTTAAAATAAGGCTAGTCCGTTATCAACTTGAAAAAGT
GGCACCGAGTCGGTGCtttttaccAAGCTTgca-3'

Table S3. Target sequences of the HBV-specific gRNAs in HBV genome

gRNA No.	Sequence (N ₁₉₋₂₀ NGG, 5'-3')	Genotype
1	CCTGCTGGTGGCTCCAGTTC	A/B/C/D
2	AGTGTTTGCTGACGCAACC	A/B/C/D
3	CAAGCCTCCAAGCTGTGCCT	A/B/C/D
4	CTAGAAGAAGAACTCCCTCG	A/B/C/D

Table S4. The full DNA sequences of all gRNAa-miR-HBV-gRNAb cassettes

Name	Sequences
3-H51-2	5'-GCAAGCCTCCAAGCTGTGCCTGTTTTAGAGCTAGAAATAGCAAG TTAAAATAAGGCTAGTCCGTTATCAACTTGAAAAAGTGGCACCG AGTCGGTGCCTCGAGcataacaacgaagaggatggtattgctcctgtaactcggaactgg agaggGGTGAAGCGAAGTGCACACGGgttgaactggaacgACGTGTGCA CATCGATTCACGGCtttctgtctgacagcagcttggctacctcctgctgttctcctgtcttG GATCCgagGGTTGCGTCAGCAAACACTGTTTTAGAGCTAGAAATA GCAAGTTAAAATAAGGCTAGTCCGTTATCAACTTGAAAAAGTGG CACCGAGTCGGTGCtttttaccAAGCTT-3'
3-H38-2	5'-GCAAGCCTCCAAGCTGTGCCTGTTTTAGAGCTAGAAATAGCAAG TTAAAATAAGGCTAGTCCGTTATCAACTTGAAAAAGTGGCACCG AGTCGGTGCCTCGAGagggatggtattgctcctgtaactcggaactggagaggGGTGA AGCGAAGTGCACACGGgttgaactggaacgACGTGTGCACATCGATTC ACGGCtttctgtctgacagcagcttggctacctcctgcttGGATCCgagGGTTGCG TCAGCAAACACTGTTTTAGAGCTAGAAATAGCAAGTTAAAATAA GGTAGTCCGTTATCAACTTGAAAAAGTGGCACCGAGTCGGTGCt tttttaccAAGCTT-3'

3-M38-2 5'-GCAAGCCTCCAAGCTGTGCCTGTTTTAGAGCTAGAAATAGCAAG
TTAAAATAAGGCTAGTCCGTTATCAACTTGAAAAAGTGGCACCG
AGTCGGTGCCTCGAGGagggatggtattgctcctgtaactcggactggagaggGGTG
AAGCGAATGGCATAGCGgttgaactgggaacgAGCTATGCCAATCGATT
CACGGCtttctgtctgacagcagcttggtacctccgctcctgttcGGATCCgagGGTTGC
GTCAGCAAACACTGTTTTAGAGCTAGAAATAGCAAGTTAAAATA
AGGCTAGTCCGTTATCAACTTGAAAAAGTGGCACCGAGTCGGTG
CtttttaccAAGCTT-3'

3-H30-2 5'-GCAAGCCTCCAAGCTGTGCCTGTTTTAGAGCTAGAAATAGCAAG
TTAAAATAAGGCTAGTCCGTTATCAACTTGAAAAAGTGGCACCG
AGTCGGTGCCTCGAGtattgctcctgtaactcggactggagaggGGTGAAGCGA
AGTGCACACGGgttgaactgggaacgACGTGTGCACATCGATTCACGGCtt
tctgtctgacagcagcttggtacctccGGATCCgagGGTTGCGTCAGCAAACAC
TGTTTTTAGAGCTAGAAATAGCAAGTTAAAATAAGGCTAGTCCGT
TATCAACTTGAAAAAGTGGCACCGAGTCGGTGCtttttaccAAGCTT-
3'

3-H22-2 5'-GCAAGCCTCCAAGCTGTGCCTGTTTTAGAGCTAGAAATAGCAAG
TTAAAATAAGGCTAGTCCGTTATCAACTTGAAAAAGTGGCACCG
AGTCGGTGCCTCGAGctgtaactcggactggagaggGGTGAAGCGAAGTG
CACACGGgttgaactgggaacgACGTGTGCACATCGATTCACGGCtttctgtc
tgacagcagcttGGATCCgagGGTTGCGTCAGCAAACACTGTTTTAGAG
CTAGAAATAGCAAGTTAAAATAAGGCTAGTCCGTTATCAACTTG
AAAAAGTGGCACCGAGTCGGTGCtttttaccAAGCTT-3'

4-H38-1 5'-GCGAGGGAGTTCTTCTTAGTTTTAGAGCTAGAAATAGCAAG
TTAAAATAAGGCTAGTCCGTTATCAACTTGAAAAAGTGGCACCG
AGTCGGTGCCTCGAGagggatggtattgctcctgtaactcggactggagaggGGTGA
AGCGAAGTGCACACGGgttgaactgggaacgACGTGTGCACATCGATTC
ACGGCtttctgtctgacagcagcttggtacctccgctcctgttcGGATCCgagCCTGCTG
GTGGCTCCAGTTCGTTTTAGAGCTAGAAATAGCAAGTTAAAATA
AGGCTAGTCCGTTATCAACTTGAAAAAGTGGCACCGAGTCGGTG
CtttttaccAAGCTT-3'

The red letters were the sequences of gRNA, the sequences of HBV specific short guide RNA were highlighted in green for gRNAa and in yellow for gRNAb; The blue letters were the sequences of pri-miR31 and anti-HBV pri-miR-31 mimic, the sequences of miRNA were highlighted in dark blue; The sequences of XhoI and HindIII restriction endonucleases cutting sites were shown in bold fonts.

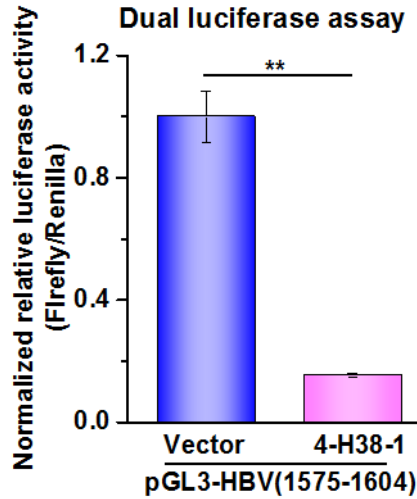


Figure S1. A dual-luciferase assay was conducted to analyze the level of miR-HBV produced by 4-H38-1 cassette. pGL3-HBV (1575-1604), PRL-TK and the expression plasmid containing 4-H38-1 cassette were co-transfected into HuH7 cells, and the dual-luciferase assay was performed. Data was shown as mean \pm SD of 5 independent experiments.

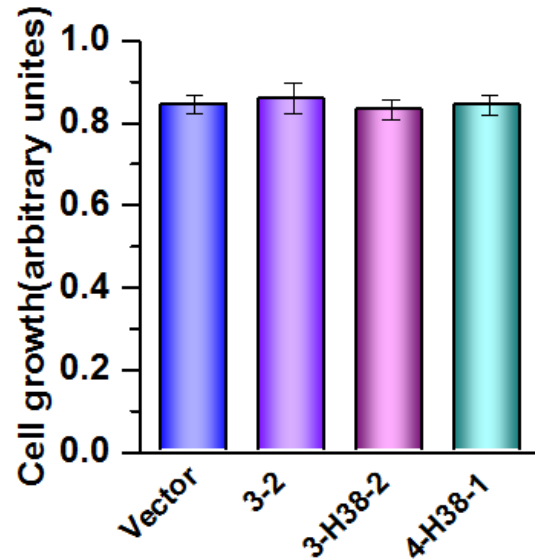


Figure S2. The cytotoxicity of 3-2 binary and gRNA-miR-HBV-gRNA ternary cassettes was examined using an MTT assay. Data was shown as mean \pm s.e.m. of 5 independent experiments.

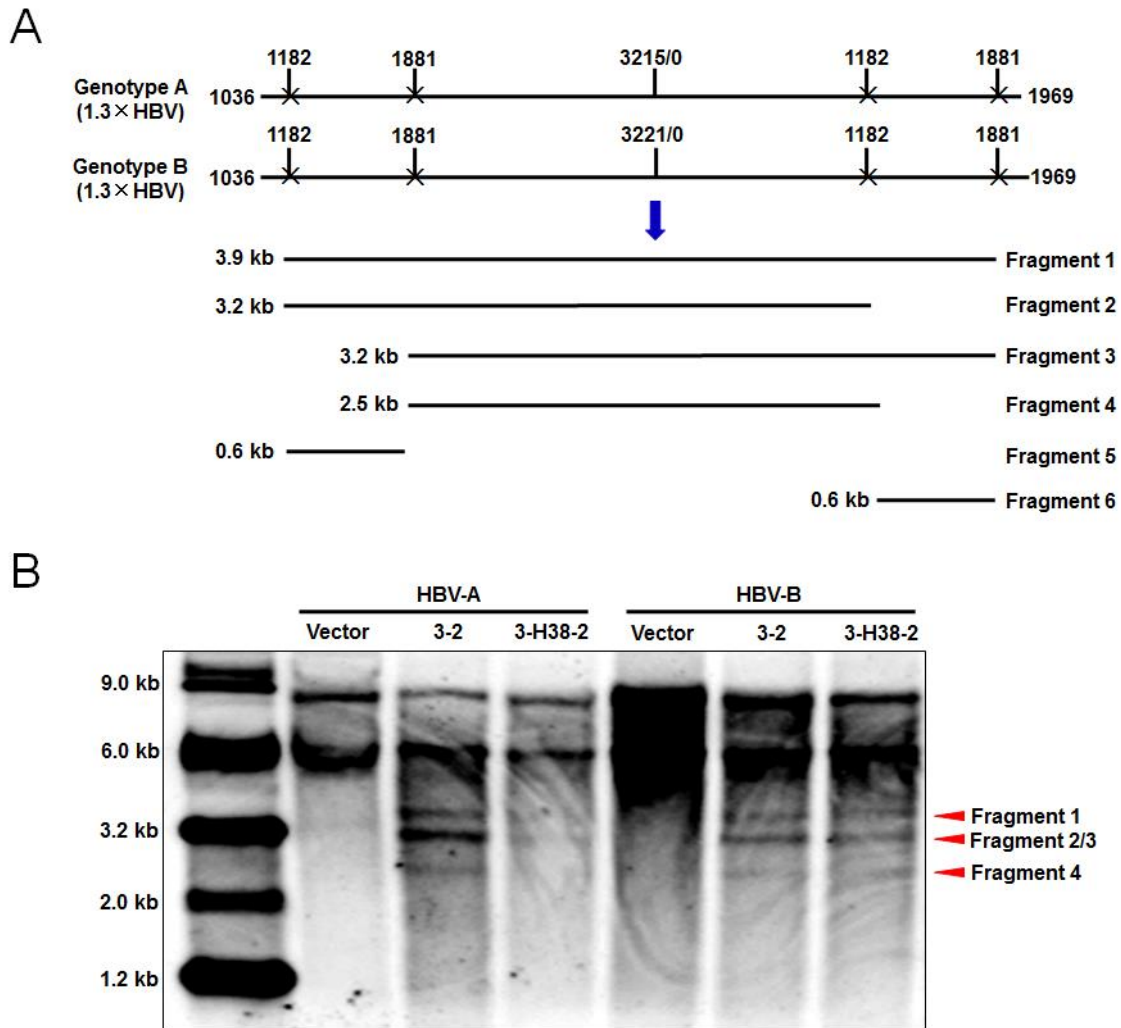


Figure S3. The HBV-specific gRNAs-mediated destructions of HBV genome were detected by Southern Blot. (A) Schematic illustration of Cas9/gRNA-mediated destruction of HBV genome (1.3×HBV expression plasmid). (B) The Cas9/gRNA-mediated destructions of HBV genome (1.3×HBV expression plasmid) were detected by Southern Blot. Vector group is the PX458 plasmid (the backbone of 3-2 and 3-H38-2 expression plasmid) was cotransfected with pGEM-HBV1.3A or pGEM-HBV1.3B into HuH7 cells.