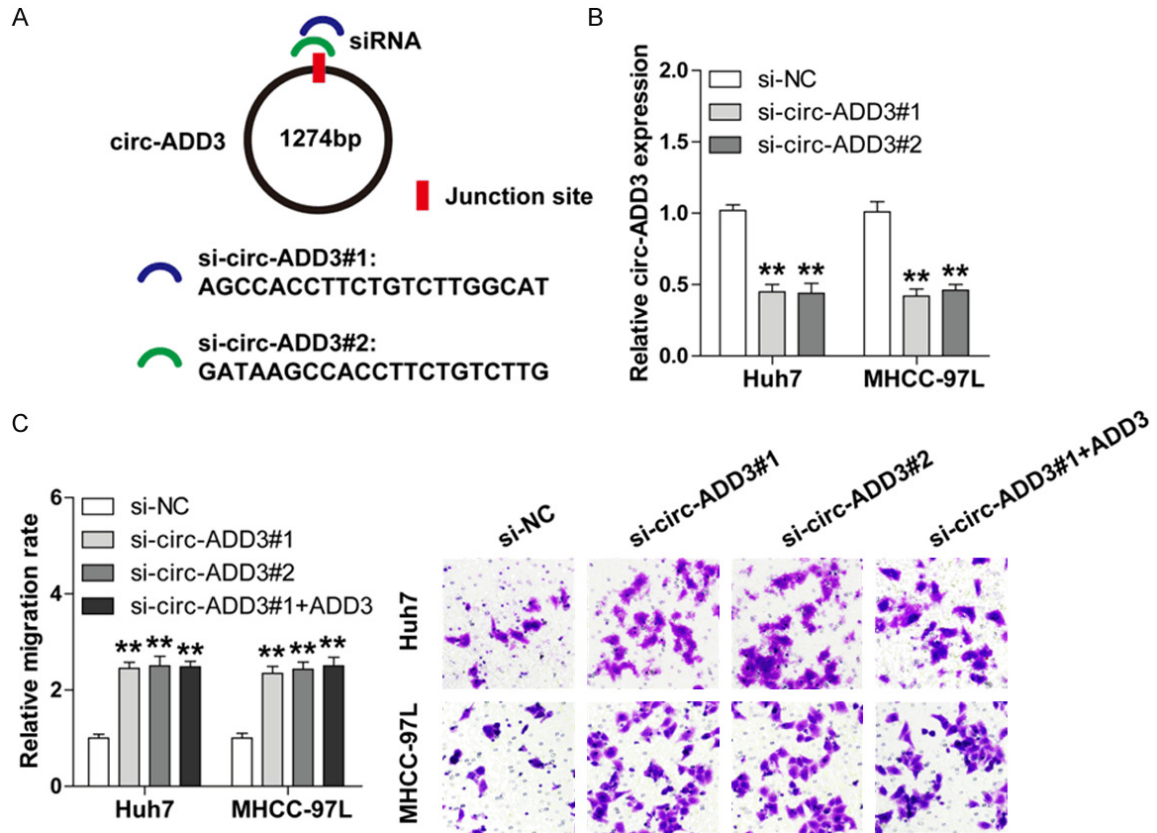


Circular RNA circ-ADD3 in hepatocellular carcinoma



Supplementary Figure 1. Knockdown of circ-ADD3 promotes HCC cell migration. A. The sequences of two siRNAs targeting the junction site of circ-ADD3. B. qRT-PCR analysis of circ-ADD3 expression in Huh7 and MHCC-97L cells transfected with the indicated siRNAs. C. Transwell migration assay in Huh7 and MHCC-97L cells transfected with the indicated siRNAs and ADD3 expression vector. ** $P < 0.01$.

Circular RNA circ-ADD3 in hepatocellular carcinoma

Protein:

MGQTGKKSEKGPVCWRKRVKSEYMRLRQLKRFRRRADEVKSMFSSNRQKILERTEILNQEW
KORRIQPVHILTSVSSLRGTRECSVTSDLDFPTQVIPLKTLNAVASVPIMYSWSPLQQNF
MVEDETVLHNIPYMGDEVLDQDGTFFIELIKNYDGKVHGDRECGFINDEIFVELVNLGQ
YNDDDDDDDDGDDPEEREKQKQLEDHRDDKESRPPRKFP SDKIFEAISSMFPDKGTAEEL
KEKYKELTEQQLPGALPPECTPNIDGPNKSVQREQLSHSFHTLFCRRCFKYDCFLHRKC
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GRLPNNSSRPSTPTINVLKSDTSDREAGTETGGENNDKEEEEKKDETSSSSEANSRCQ
TPIKMKPIEPPENVEWSGAEASMFRLVIGTYD NFC AIARLIGTKTCRQVYEFVKESS
IIAPAPAEDVDTPPRKKRKHRLWAAHCRKIQLKKGSSNHVYNYQPCDHPRQPCDSSCP
CVIAQNFCEKFCQCSSECQNRFPGCRCKAQCNKQCPCYLAVRECDPDLCLTCGAADHW
SKNVSCKNCSIQRGSKKHLLLAPSDVAGWGIFIKDPVQKNEFISEYCGEISQDEADRRG
KYVDKYMCSFLFNLNDFVVDATRKGNKIRFANHSVNPNCYAKVMMVNGDHRIGIFAKRA
IQTGEELFFDYRYSQADALKYVGIEREMEIP

EZH2
protein

RNA:

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CUCUCAUUCGAGAGAAGCCUAGGCACAAGAGUGAUGUGGAAAUCCAGCAACUGUGACUG
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AUCGAUAUGACUUGAAAACAGCAGGACCACAUCUCAGUUGCUUGCUGGAAUUGUUGUGG
AUAAGCCACCUUCU

circ-ADD3

Interaction probabilities

Prediction using RF classifier 0.8

Prediction using SVM classifier 0.96

Supplementary Figure 2. The interaction probability between circ-ADD3 and EZH2 was predicted by RPISeq online software based on RF and SVM classifiers (probabilities > 0.5 were considered "positive").

Circular RNA circ-ADD3 in hepatocellular carcinoma

Protein:

MEDYTKIEKIGEGTYGVVYKGRHKTTGQVWAMKKIRLESEEEGVPSTAIRESLLKELRH
 PNIVSLQDVLMDQSRLYLIFEFLSMDLKKYLDSSIPGQYMDSSLVKSYLEYLQILQGIVFCH
 SRRVLRDLKPNQLLIDDKGTIKLADFLARAFGIPIRVYTHEVTLWYRSPEVLLGSAR
 YSTPVDIWSIGTIFAEATKKPLFHGDSEIDLFRIFRALGTPNNEVWPEVESLQDYKNT
 FPKWKPGSLASHVKNLDENGLDLLSKMLIYDPAKRISGKMLNHPYFNDLDNQIKKM

CDK1p
 rotein

RNA:

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 GAUGGGCACACCUUGGCAAAUACCUAUAUCUCAGUAAGAAUAAGUAAGGAGCAAGACCACA
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 UCAAUAUAAUAGGAGAAGUGGUUUGACCAGGGAAGUACCAAUUUUGAAAAUUGACCAUACAG
 GAUUCAGUCCCAUGCUGCAAUUAUUAACACGUCCUGAUGUUUAGUGUGUGAUACACA
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 GUGGAGGAGGUGUGAAUUGGGUUCUCAAAAAUUGGAAGGUUGGCGAAAUUGAGUUUG
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 CUCUCAUUCGAGAGAAGCCUAGGCACAAGAGUGAUGUGGAAUCCAGCAACUGGACUG
 CUUUUUCUUUGAAGACGAUACAGUGCCACUCUCUCCUCUCAAAUACAUGGCACAGAGGC
 AACAGCGUGAAAAACAAGAUGGCUGAACUCACCAAAUACUUACAUGAAAGUGAAUGUGC
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 AUAAGCCACCUUCU

circ-ADD3

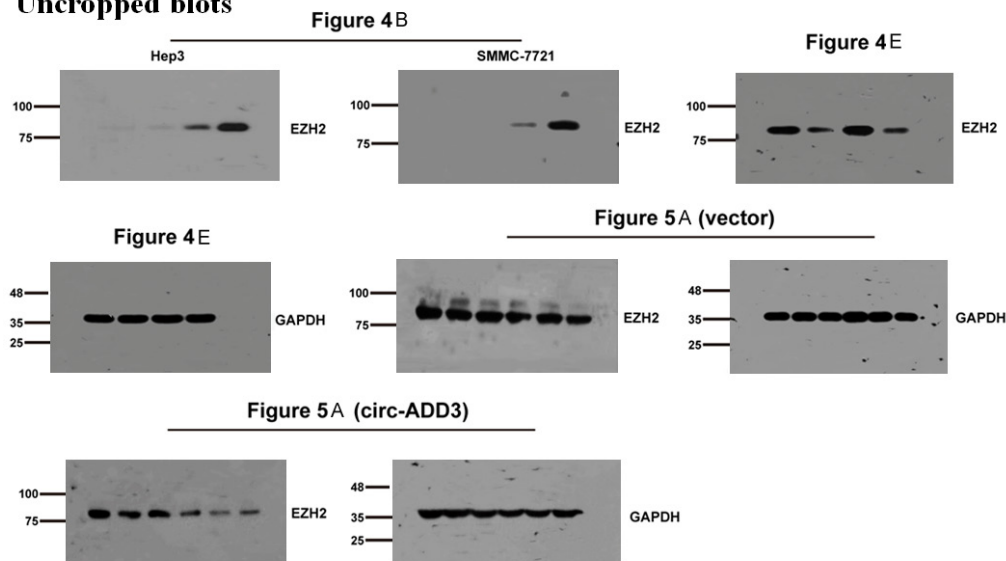
Interaction probabilities

Prediction using RF classifier 0.85

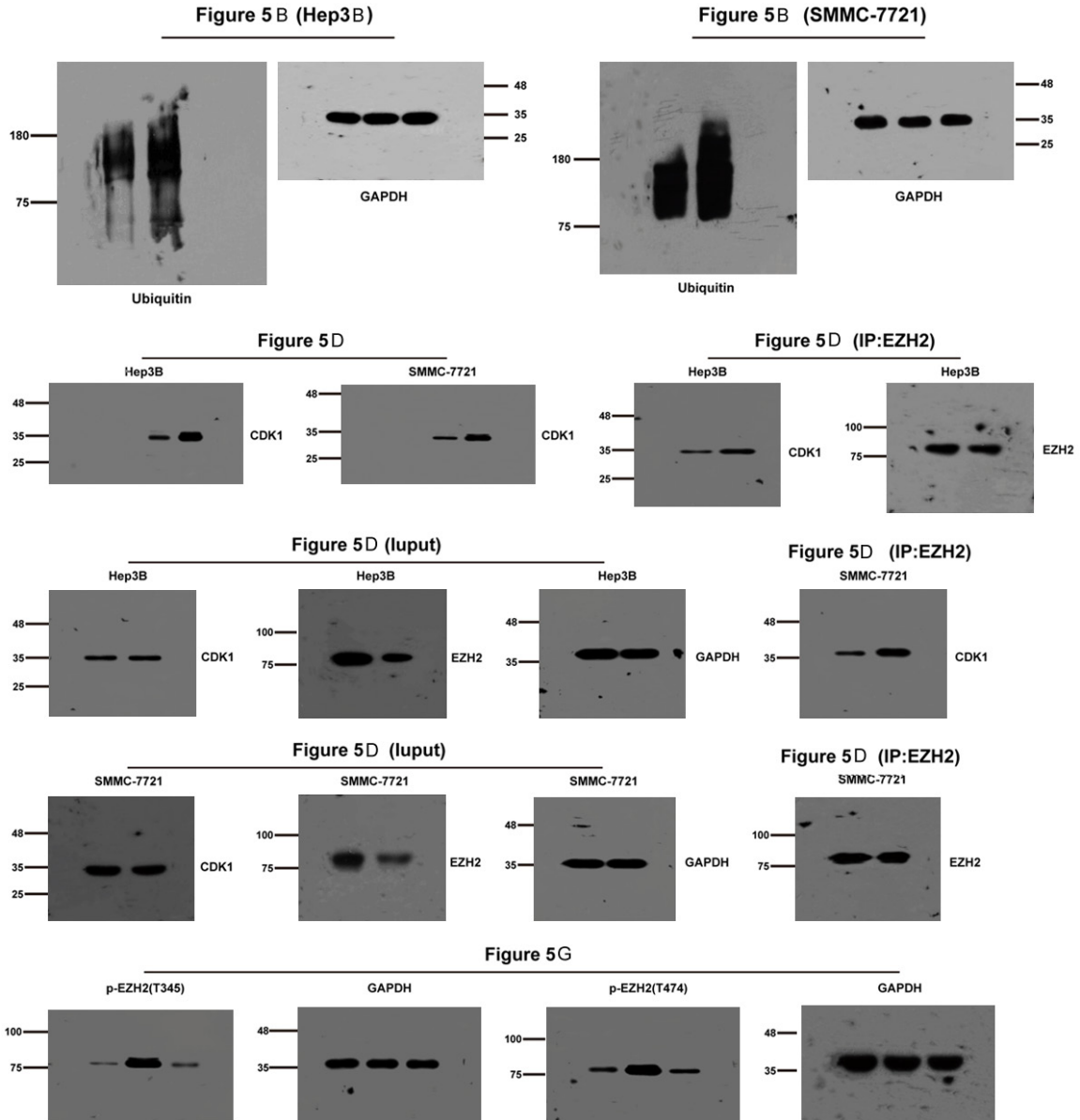
Prediction using SVM classifier 0.96

Supplementary Figure 3. The interaction probability between circ-ADD3 and CDK1 was predicted by RPISeq online software based on RF and SVM classifiers (probabilities > 0.5 were considered "positive").

Uncropped blots



Circular RNA circ-ADD3 in hepatocellular carcinoma



Supplementary Figure 4. The amino acid sequence alignment between hsa-circ-ADD3 and homologous mus-circ-ADD3. The data showed 87% similarity between them.