

## Supplementary Material

### FTO-mediated autophagy promotes progression of clear cell renal cell carcinoma via regulating SIK2 mRNA stability

**Table S1.** Sequences of primers.

**Table S2.** Primary antibodies.

**Figure S1.** Immunofluorescence staining with mRFP-GFP-LC3 in ccRCC cell lines.

**Figure S2.** IGFBP2 and ALKBH5 expression in OSRC-2 and 786-O cells treated with rapamycin.

**Figure S3.** The time-dependent receiver operating characteristic (ROC) analysis for the FTO risk score, the clinical variables risk score, and the combined FTO and clinical variables risk scores in the ccRCC cohort.

**Figure S4.** m<sup>6</sup>A peak distribution in the 5'UTR, 3'UTR, and CDS across the entire set of mRNA transcripts.

**Figure S5.** MeRIP-seq data revealed that m<sup>6</sup>A abundance in the SIK2 mRNA was markedly increased during autophagy flux.

**Figure S6.** Gene Ontology pathway and Kyoto Encyclopedia of Genes and Genomes pathway analyses of differentially expressed genes in autophagy-induced OSRC-2 cells compared with normal OSRC-2 cells.

**Figure S7.** Representative images of metastatic lung tumors and hematoxylin and eosin staining, and quantification of lung tumors.

**Figure S8.** Representative IHC staining images of N-cadherin and E-cadherin and quantification of IHC score.

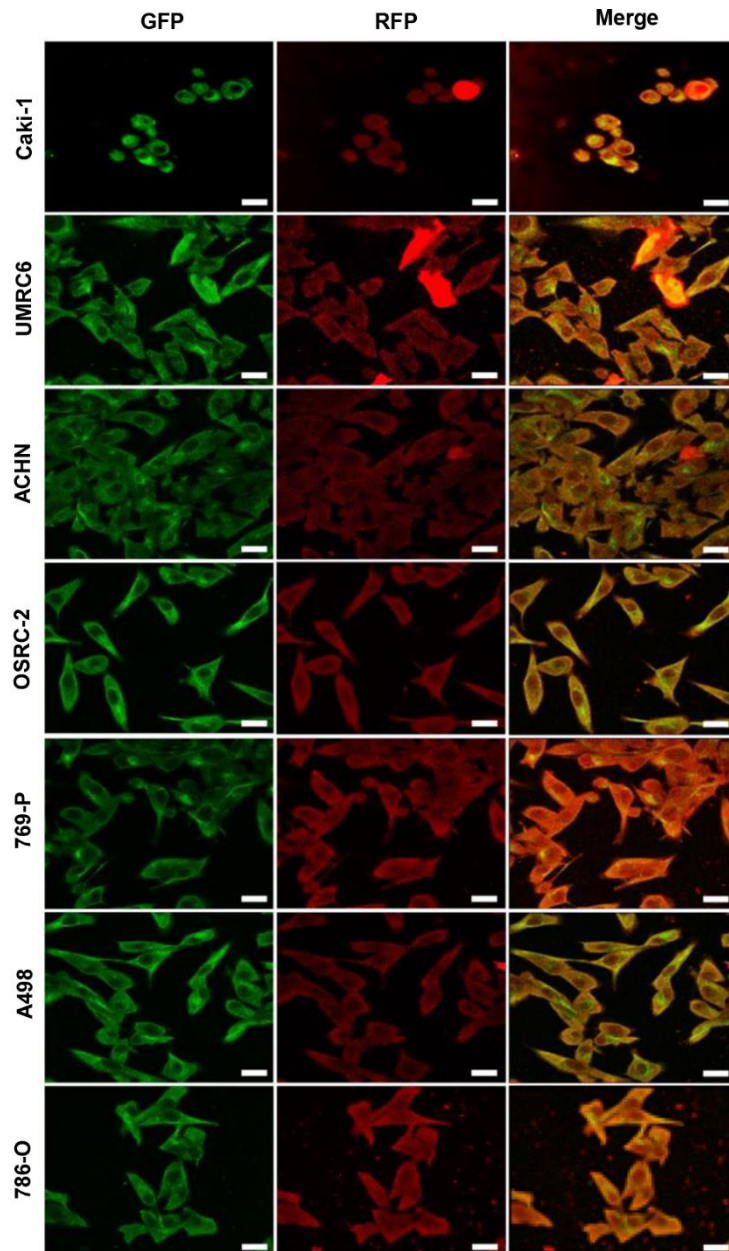
**Table S1.** Sequences of primers

Gene Symbol	Sequences
FTO	Forward 5'- ATTCTATCAGCAGTGGCAGC -3' Reverse 5'- GGATGCGAGATACCGGAGTG -3'
ULK1	Forward 5'- CCTCGGTCAGGAAATCCCA -3' Reverse 5'- GAGCTGTGCAGGAAGCCA -3'
ATG3	Forward 5'- CCCAGGATGCAGAATGTGA -3' Reverse 5'- TCCAGCTGCCACAACTCTT -3'
ATG4B	Forward 5'- TCCATCGCTGTGGGGTTTTT -3' Reverse 5'- TCCAGTCGCTCTACATCAGA -3'
ATG4D	Forward 5'- ACACATCCTCAGGAAAGCCG -3' Reverse 5'- TGACCACAGACTTCCACTCG-3'
ATG5	Forward 5'- GCAACTCTGGATGGGATTGC -3' Reverse 5'- AGGTCTTTCAGTCGTTGTCTGAT -3'
ATG7	Forward 5'- AGCATCCAGAAGGGGGCTA -3' Reverse 5'- TGCAGCAATGTAAGACCAGTCA -3'
ATG10	Forward 5'- ACATTCCAACGTTATTGTGCAG -3' Reverse 5'- GCTCGAAAGCCTCCTCCAT -3'
ATG12	Forward 5'- TGCTGGAGGGGAAGGACTTA -3' Reverse 5'- CCATCACTGCCAAAACACTCA -3'
ATG13	Forward 5'- AGCCCGGAACCACTCTT -3' Reverse 5'- TCACTTGGACAGTCTTGAGGG -3'
ATG14	Forward 5'- TACGTGGCTGTGGAGCG -3' Reverse 5'- CCTTCCATAGCTTTTAACACTTCTT -3'
ATG16L1	Forward 5'- TTGGCCCAGAACTACAGGC -3' Reverse 5'- CCAGTTGAGCTAACTCCCA -3'
SIK2	Forward 5'- CTGAAGCCAGGCGAAAATTCT -3' Reverse 5'- TTGCCAGCAGTTCACCACTTT -3'
PIK3C2A	Forward 5'- GGTTCCGATCCTCTGCGTT -3' Reverse 5'- GGTTCCGATCCTCTGCGTT -3'
RNF34	Forward 5'- GGAGAGCTTATGGATGGAGACC -3' Reverse 5'- GGTTCCGATCCTCTGCGTT -3'
IGF2BP1	Forward 5'- TTGGGACGCCATCAGTACCTA -3' Reverse 5'- TTGGCTAACTCTCTACGACTCT -3'
IGF2BP2	Forward 5'- TGCACATCCCCAACTGTGAC -3' Reverse 5'- TGTAGAAGAGATGACACTCGGG -3'
IGF2BP3	Forward 5'- AGACACACTGAATCACCTGAAGT -3' Reverse 5'- AGGGCGACACTGCTTTTTTCTT -3'
YTHDF1	Forward 5'- GGGGACAAGTGGGTCTCAAG -3' Reverse 5'- AGGGTGTGCTGTGAAAGC -3'
YTHDF2	Forward 5'- GTTGGTAGCGGGTCCACTACT -3' Reverse 5'- GGTCTTCAGTTTAGGTTGCTGT -3'
YTHDF3	Forward 5'- GGTGTATTTAGTCAACCTGGGG -3'

	Reverse 5'- AAGAGAACTAGGTGGATAGCCAT -3'
METTL3	Forward 5'- TTGTCTCCAACCTTCCGTAGT -3' Reverse 5'- CCAGATCAGAGAGGTGGTGTAG-3'
METTL16	Forward 5'- CTCTGACGTGTA CTCTCCTAAGG -3' Reverse 5'- TACCAGCCATTCAAGGTTGCT -3'
WTAP	Forward 5'- CTTCCAAGAAGGTTTCGATTGA -3' Reverse 5'- TCAGACTCTCTTAGGCCAGTTAC -3'
VIRMA	Forward 5'- CCTTCCTGGGCTAGTGCAAA -3' Reverse 5'- TTTTGTCTTTTGGTACAGCCAT -3'
ZC3H13	Forward 5'- TCTGATAGCACATCCCGAAGA -3' Reverse 5'- CAGCCAGTTACGGCACTGT -3'
RBM15	Forward 5'- ACGACCCGCAACAATGAAG -3' Reverse 5'- GGAAGTCGAGTCCTCACCAC -3'
RBM15B	Forward 5'- ACCTGGACCACAGCGTATCT -3' Reverse 5'- GGGTTGCGACCAATCACTC -3'
RBMX	Forward 5'- CTTCAGGACCAGTTCGCAGTA -3' Reverse 5'- TCACGACCACTTGAGTAGAGAT -3'
YTHDC1	Forward 5'- CTTCTGATGAGCAAGGGAACAA -3' Reverse 5'- GGCCTCACTTCGAGTGT CATAA -3'
YTHDC2	Forward 5'- CAAAACATGCTGTTAGGAGCCT -3' Reverse 5'- CCACTTGTCTTGCTCATT TCCC -3'
HNRNPC	Forward 5'- TCCTCCTCCTATTGCTCGGG -3' Reverse 5'- AGCCACTTTTGCCCTTCG -3'
FMR1	Forward 5'- ACTTACGGCAAATGTGTGCCA -3' Reverse 5'- GCAGACTCCGAAAGTGCATGT -3'
LRPPRC	Forward 5'- CGGAGGACTACTGAGCCCA -3' Reverse 5'- AGCGGCAGGTATCATTAAAACT -3'
HNRNPA2B1	Forward 5'- TGGAGGTAGCCCCGGTTATG -3' Reverse 5'- GGACCGTAGTTAGAAGGTTGCT -3'
ALKBH5	Forward 5'- ATGCACCCCGGTTGGAAAC -3' Reverse 5'- GACTTGCGCCAGTAGTTCTCA -3'

**Table S2.** Primary antibodies

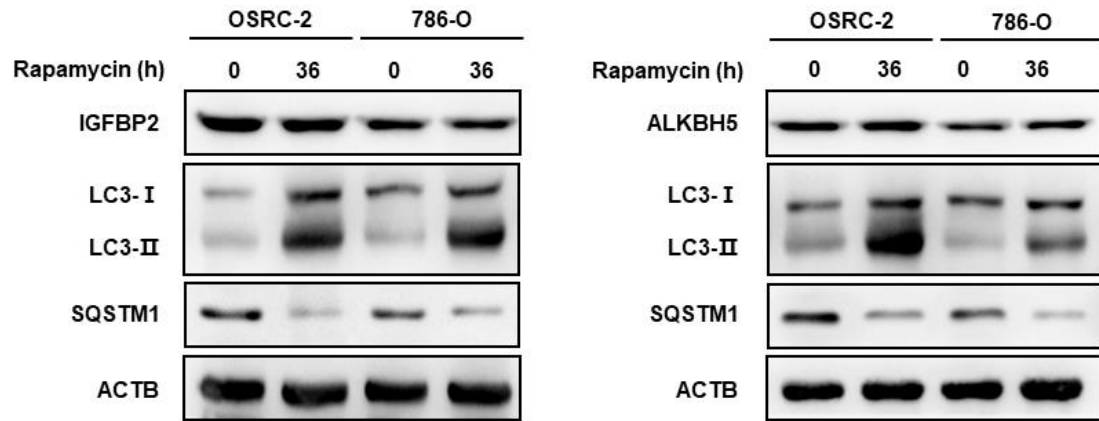
Primary antibodies	Manufacturer, Catalogue number
Rabbit anti-m <sup>6</sup> A	Synaptic Systems, 202003
Rabbit anti-LC3- I/II	Abcam, ab192890
Mouse anti-GFP	Abcam, ab1218
Rabbit anti-RFP	Abcam, ab185921
Rabbit anti-SQSTM1	Abcam, ab109012
Rabbit anti-ACTB	Abcam, ab8227
Rabbit anti-FTO	Abcam, ab126605
Rabbit anti-GAPDH	Abcam, ab9485
Rabbit anti-Ki67	Abcam, ab15580
Rabbit anti-SIK2	Abcam, ab245211
Rabbit anti-IGF2BP2	Abcam, ab128175
Rabbit anti-ATG5	Cell Signaling Technology, 9980S
Rabbit anti-ATG7	Cell Signaling Technology, 8558S
Rabbit anti-IGFBP2	Proteintech, 11065-3-AP
Rabbit anti-ALKBH5	Proteintech, 16837-1-AP
Rabbit anti-PIK3C2A	Proteintech, 22028-1-AP
Rabbit anti-N-cadherin	Proteintech, 66219-1-Ig
Rabbit anti-E-cadherin	Proteintech, 20874-1-AP
Rabbit anti-RNF34	ABclonal, A8517



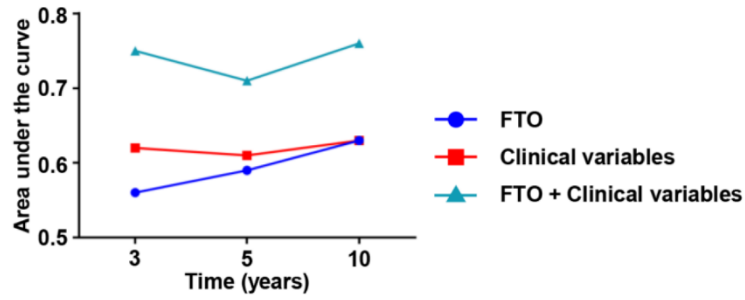
**Figure S1.** Immunofluorescence staining with mRFP-GFP-LC3 in ccRCC cell lines.

Red puncta signify autolysosomes and yellow puncta signify autophagosomes. Scale

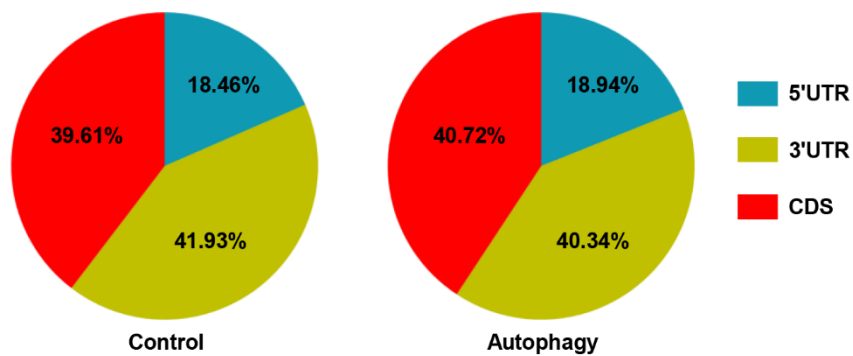
bar, 10  $\mu\text{m}$ .



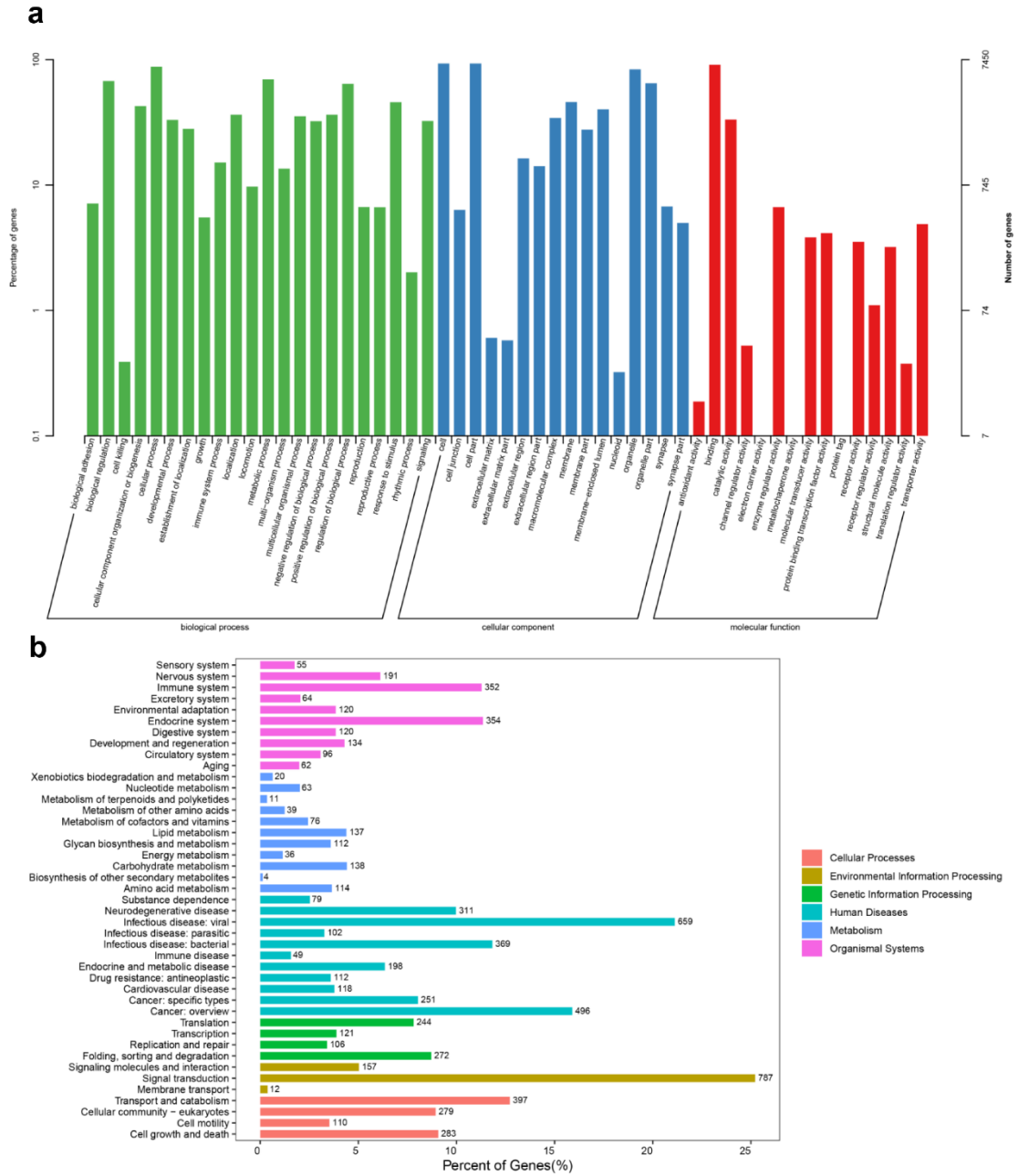
**Figure S2.** IGFBP2 and ALKBH5 expression in OSRC-2 and 786-O cells treated with rapamycin.



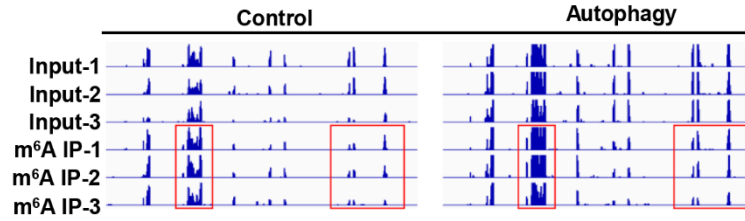
**Figure S3.** The time-dependent receiver operating characteristic (ROC) analysis for the FTO risk score, the clinical variables risk score, and the combined FTO and clinical variables risk scores in the ccRCC cohort.



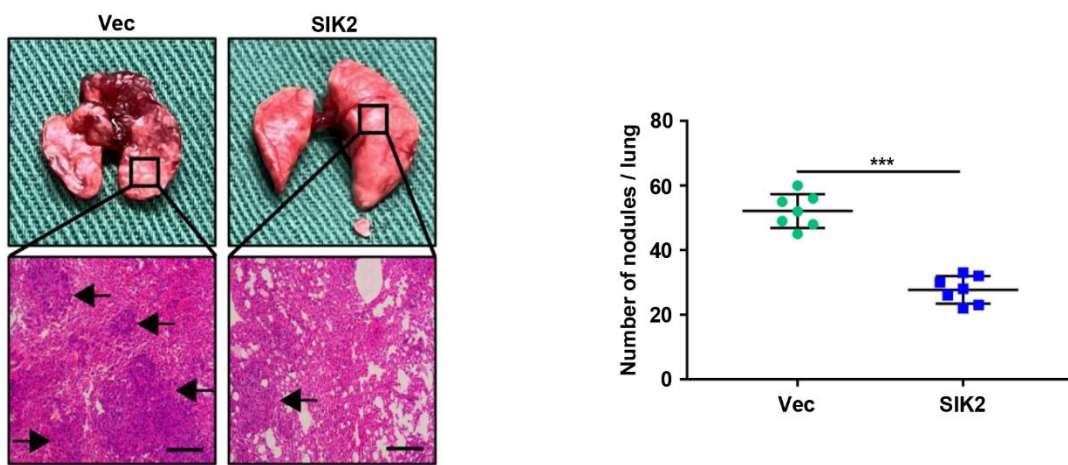
**Figure S4.** m<sup>6</sup>A peak distribution in the 5'UTR, 3'UTR, and CDS across the entire set of mRNA transcripts.



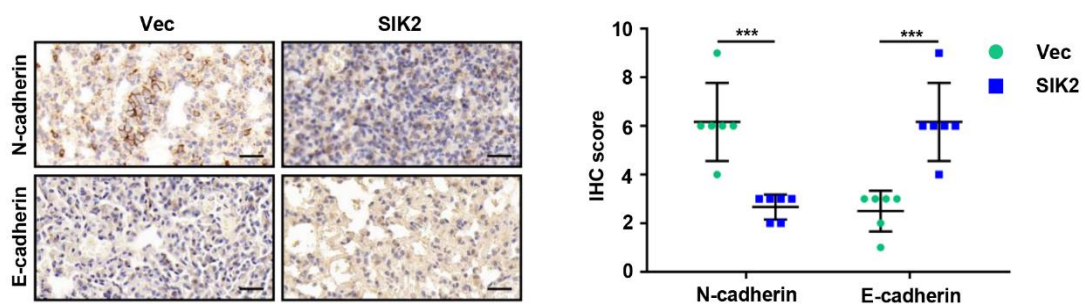




**Figure S6.** MeRIP-seq data revealed that m<sup>6</sup>A abundance in the SIK2 mRNA was markedly increased during autophagy flux.



**Figure S7.** Representative images of metastatic lung tumors and hematoxylin and eosin (H&E) staining (left panel), and quantification of lung tumors (right panel).



**Figure S8.** Representative IHC staining images of N-cadherin and E-cadherin (scale bar, 100µm, left panel) and quantification of IHC score (right panel). Error bars, SEM;

\*\*\*,  $P < 0.001$ .