

Supplementary materials

The m6A reader YTHDC2 promotes the pathophysiology of temporal lobe epilepsy by modulating SLC7A11-dependent glutamate dysregulation in astrocytes

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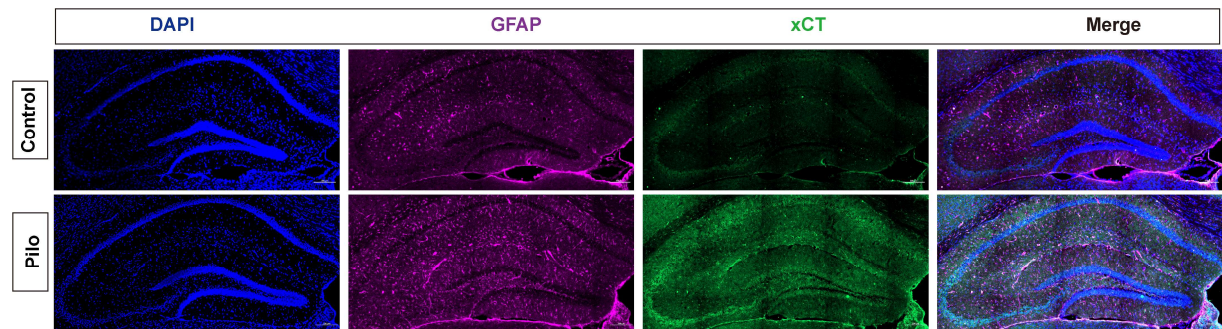


Figure S1. Representative immunofluorescence image of GFAP and xCT co-localization in the mouse hippocampus (scale bar = 200 μm)

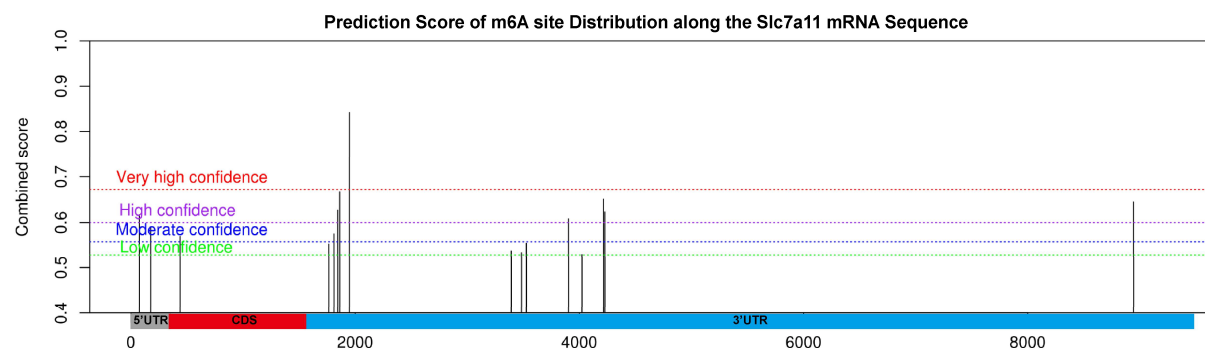


Figure S2. Predicted m6A sites in SLC7A11 mRNA identified by the SRAMP algorithm.

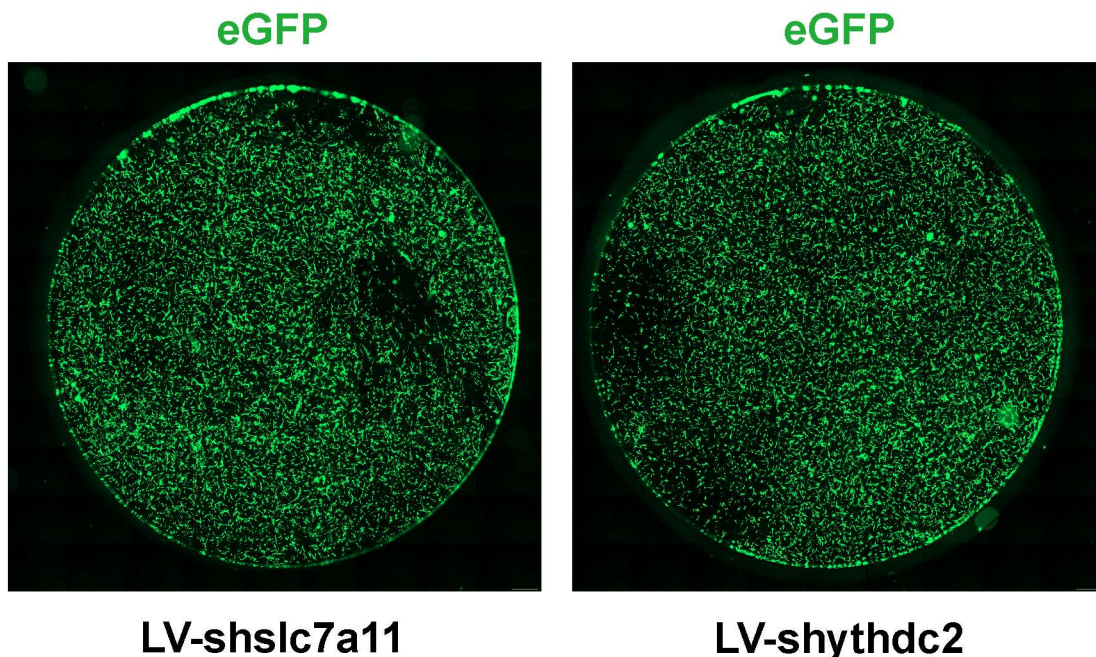


Figure S3. Representative immunofluorescence image of primary astrocytes transfected with

lentivirus.

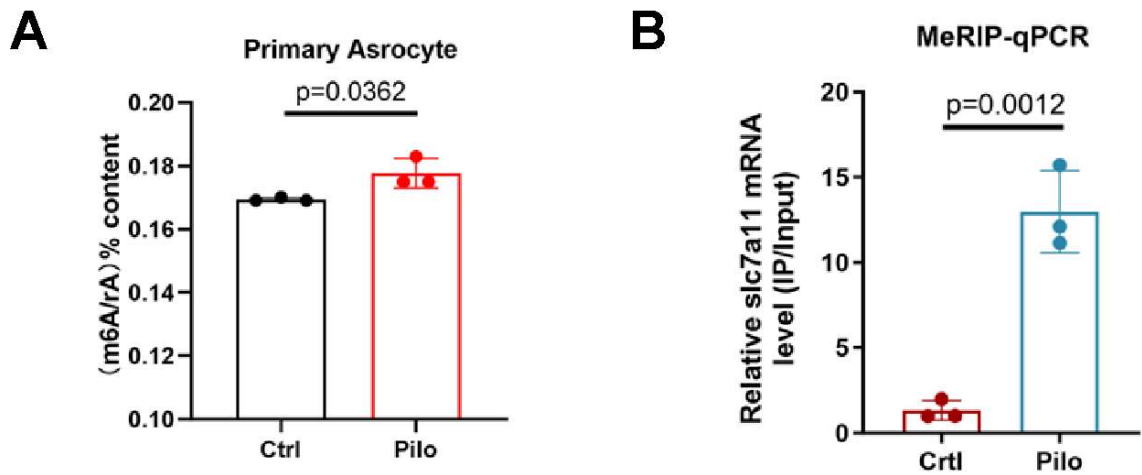


Figure S4. (A) Methylation modification of m6A in primary astrocytes after stimulation with pilocarpine and in the control group ($p = 0.0362$, $n = 3$). (B) MeRIP-qPCR analysis of hippocampal SLC7A11 m6A methylation modification in the pilocarpine and control groups ($p = 0.0012$, $n = 3$); two-sided unpaired Student's *t*-tests.

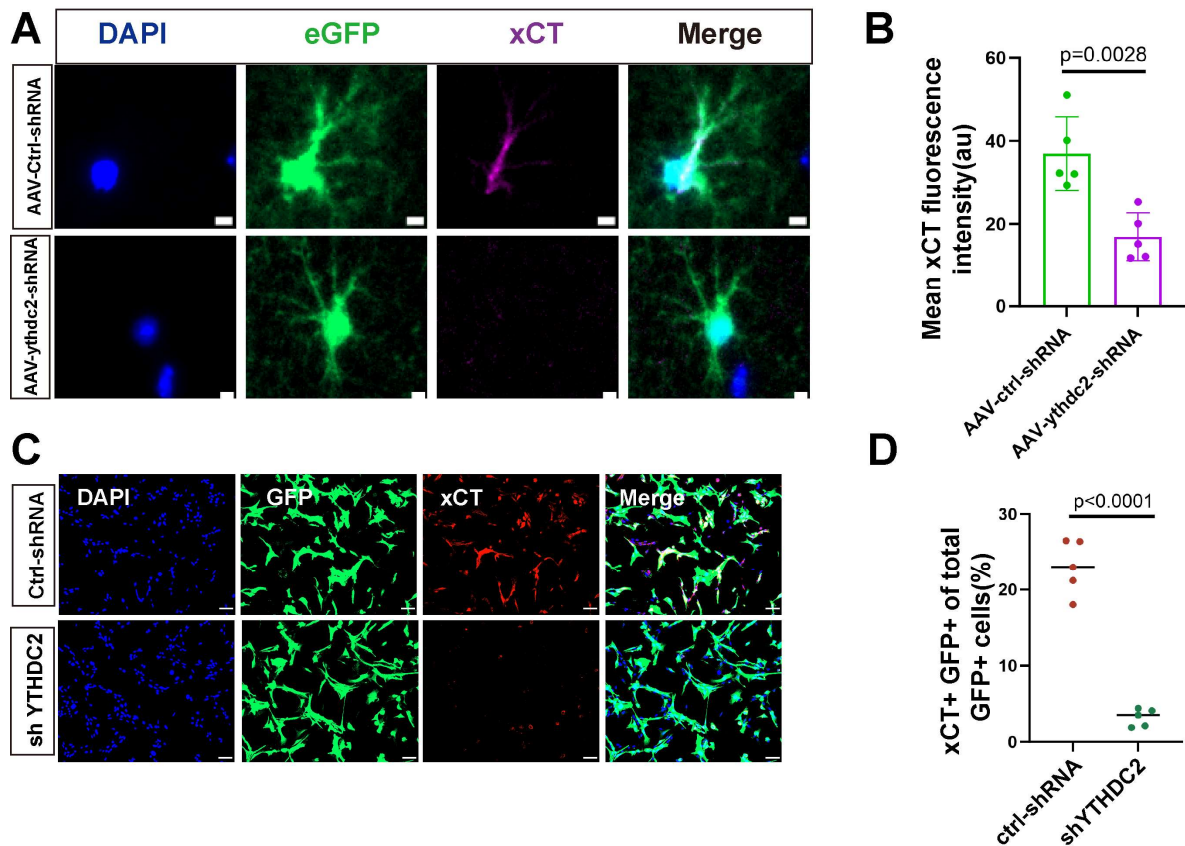


Figure S5. (A, B) Immunofluorescence staining of xCT protein fluorescence intensity in astrocytes of the YTHDC2-knockdown group and control groups ($p = 0.0028$, $n = 5$). scale bar = 2 μ m. (C, D) Cell-immunofluorescence of xCT expression after pilocarpine stimulation in the YTHDC2-knockdown group and control groups ($p < 0.0001$, $n = 5$). scale bar = 50 μ m. two-sided unpaired Student's *t*-tests (B, D).

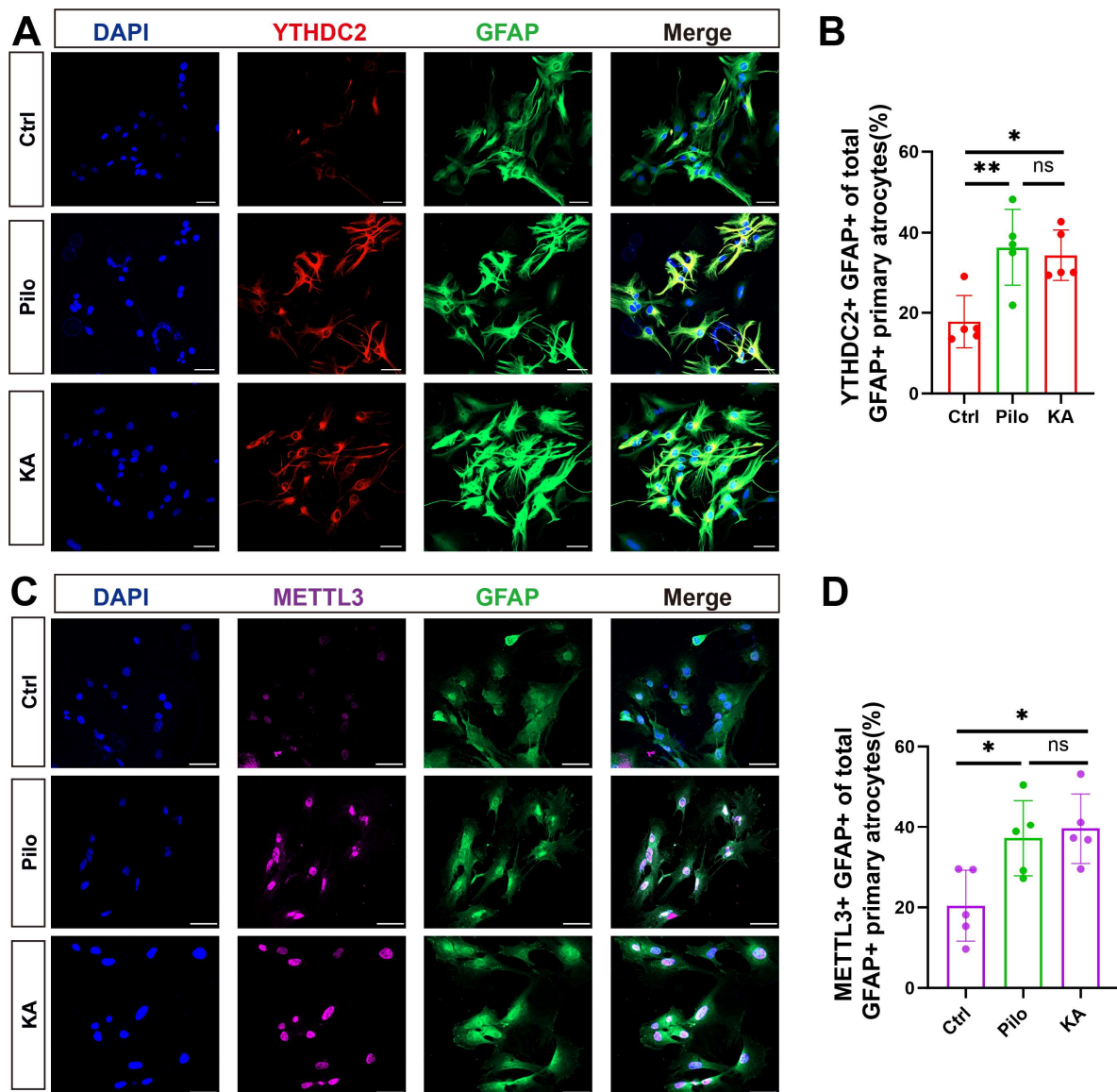


Figure S6 (A, B) Immunofluorescence staining of YTHDC2 in astrocytes treated with pilocarpine ($p = 0.0058$, $n = 5$) or KA ($p=0.0115$, $n = 5$). Scale bar = 50 μm . **(C, D)** Immunofluorescence staining of METTL3 in astrocytes treated with pilocarpine ($p = 0.0298$, $n = 5$) or KA ($p = 0.0141$, $n = 5$). scale bar = 50 μm . one-way ANOVA with Bonferroni's post hoc test (B, D).

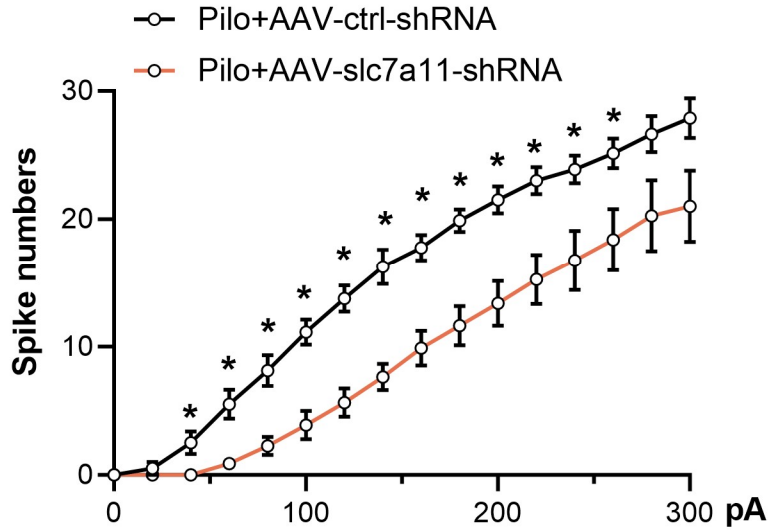


Figure S7 A range of current injections versus the number of spikes between AAV-ctrl-shRNA and AAV-slc7a11-shRNA group; two-side unpaired Student's *t*-test. * $p < 0.05$

Table S1 Human participant information

Patients	Sex	Age	Side	Histology	Experiment
TLE1	Male	23	Left	Hippocampal sclerosis	MRS
TLE2	Male	35	Left	Hippocampal sclerosis	MRS
TLE3	Female	29	Right	Hippocampal sclerosis	MRS
TLE4	Male	35	Left	Hippocampal sclerosis	MRS, IF
TLE5	Female	32	Right	Hippocampal sclerosis	IF
TLE6	Female	38	Left	Hippocampal sclerosis	IF
TLE7	Male	30	Right	Hippocampal sclerosis	IF
TLE8	Female	44	Left	Hippocampal sclerosis	IF
Normal1	Female	21	Right	-	IF
Normal2	Male	28	Left	-	IF
Normal3	Female	24	Right	-	IF
Normal4	Male	26	Right	-	IF
Normal5	Male	16	Left	-	IF

Table S2. Primer sequences used for qPCR

Gene	Forward Primer (5'-3')	Reverse Primer (5'-3')
<i>Slc7a11</i>	ATCTCCCCCAAGGGCATACT	GCATAGGACAGGGCTCCAAA
<i>Actb</i>	TATAAAACCCGCGCGCA	TCATCCATGGCGAACTGGTG
<i>Ythdc2</i>	ACGGTGACCAGAGAGAAATGG	GGTCATCATTGCATGAGCTGT
<i>Ythdc1</i>	CGGGAGGAGAAAGATGGGGA	TGTCGCTTGGTGTCAGTAGAC
<i>Mettl3</i>	CTTGCCATCTCTACGCCAGA	TCATGGCAGACAGCTTGGAG
<i>Mettl4</i>	AAGCGAGGGTTTGTTCCTCGA	AAAGGTGAGTGTCCGTCAGG
<i>Mettl14</i>	CTTGGGAGAGTATGCTTGCG	CCTTTGATCCCCATCAGGCA
<i>Alkbh5</i>	GTGACTGTGCTCAGTGGGTA	TTCCAATCGCGGTGCATCTA
<i>Fto</i>	GGGACATCGAGACACCAGGA	AGGTGCCTGTTGAGCACTCT
<i>Ythdf1</i>	CCGTGATACACAGGAGGTGC	TCGATTCTGTCTTTCCTTACGCA
<i>Ythdf2</i>	CAGGCAAGGCCGAATAATGC	TCTCCGTTGCTCAGTTGTCC

<i>Ythdf3</i>	ACTAGCGTGGATCAGAGACCT	GGAAATCCAATGGACGGAGC
<i>Nrf2</i>	GCCCTCAGCATGATGGACTT	AACTTGTACCGCCTCGTCTG
<i>Atf4</i>	CCTATAAAGGCTTGCGGCCA	GATTCGTGAAGAGCGCCAT
<i>Bap1</i>	TTGCAAACCTATGAGGCCTGTCT	TTGGCCAGCATTCCTTCCTGA
<i> Tp53</i>	ATCCTGGCTGTAGGTAGCGA	CCATGGCAGTCATCCAGTCT

Knockdown sequences of adeno-associated virus (AAV)

1. GOSV0334324 MIR155(*Slc7a11*) *3 CV258: GFAP-EGFP-MIR155

CTGGAGGCTTGCTGAAGGCTGTATGCTGATAACCTGGAGACAGCGAACAGTTTTG
GCCACTGACTGACTGTTTCGCTCTCCAGGTTATCAGGACACAAGGCCTGTTACTAGC
ACTCACATGGAACAAATGGCCCGGGCCCTTTAAACTGGAGGCTTGCTGAAGGCTG
TATGCTGAAATCAGTCCTGCGACTGCCAGTTTTGGCCACTGACTGACTGGCAGTCA
GGACTGATTTAGGACACAAGGCCTGTTACTAGCACTCACATGGAACAAATGGCC
CTTAATTA ACTGGAGGCTTGCTGAAGGCTGTATGCTGAATAGTGCCGGGATGAAGA
GAGTTTTGGCCACTGACTGACTCTCTTCACCGGCACTATTAGGACACAAGGCCTG
TTACTAGCACTCACATGGAACAAATGGCCC

2. GOSV0334325 MIR155(*Ythdc2*) *3 CV258: GFAP-EGFP-MIR155

CTGGAGGCTTGCTGAAGGCTGTATGCTGTGCAATGGCATTCTCAAGAGTGTTTTGG
CCACTGACTGACACTCTTGAATGCCATTGCA CAGGACACAAGGCCTGTTACTAGCA
CTCACATGGAACAAATGGCCCGGGCCCTTTAAACTGGAGGCTTGCTGAAGGCTGT
ATGCTGTACAAAGCATAGCTGCCCGTTGTTTTGGCCACTGACTGACAACGGGCATA
TGCTTTGTA CAGGACACAAGGCCTGTTACTAGCACTCACATGGAACAAATGGCCCT
TAATTA ACTGGAGGCTTGCTGAAGGCTGTATGCTGTAAACCTGCAGACTGTTCTTC
GTTTTGGCCACTGACTGACGAAGA AACTGCAGGTTTACAGGACACAAGGCCTGT
TACTAGCACTCACATGGAACAAATGGCCC