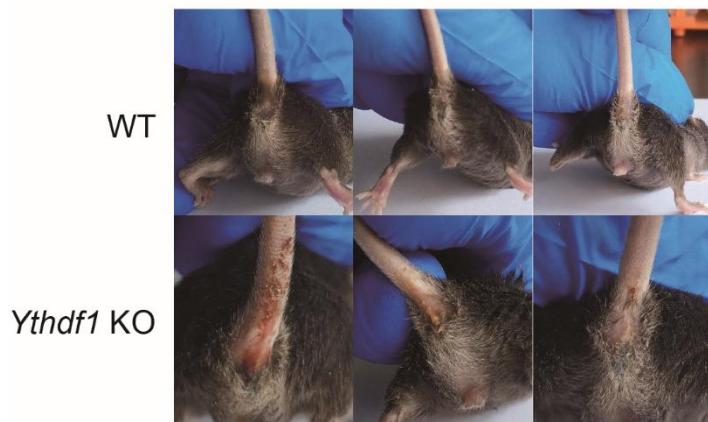


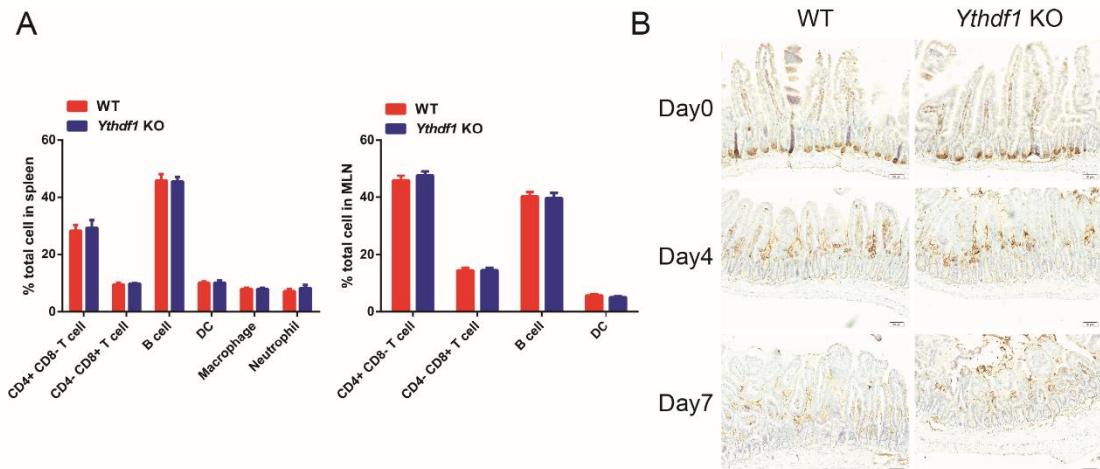
## Supplementary information

# ***Ythdf1* deficiency exacerbates colitis but prevents colitis-associated cancer in mice**

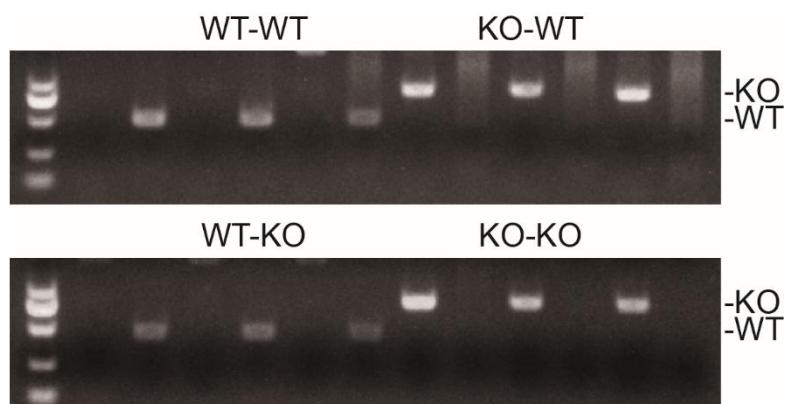
Zhen Zhang, Jie Zhang, Chenzhong Xu, Yuping Ning, Baohua Liu



**Figure S1.** Images showing representative diarrhoea and bloody stools in *Ythdf1* KO mice at day 7 following DSS challenge. Three mice per genotype were analysed.

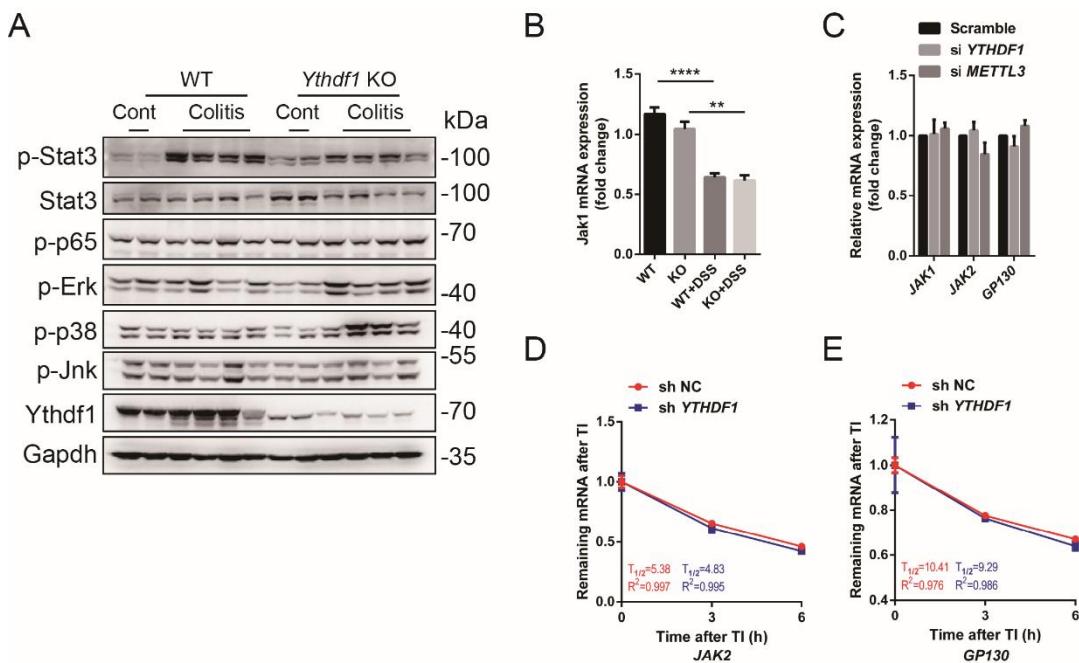


**Figure S2.** (A) Flow cytometry analysis of immune cell frequency in spleen (left) and MLN (right) after the induction of colitis in WT ( $n = 5$ ) or KO ( $n = 4$ ) mice; (B) Lysozyme expression in ileum sections at different time points after exposure to DSS, as determined with IHC (scale bar, 50  $\mu$ m).

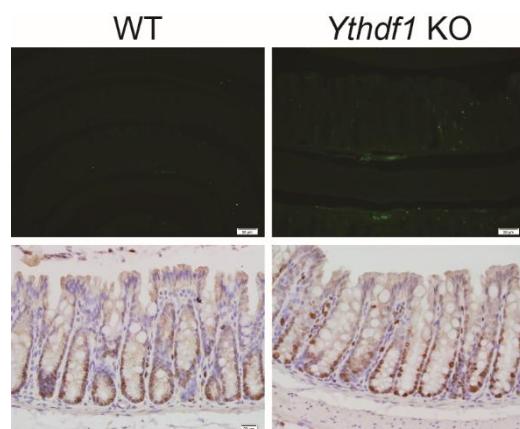


**Figure S3.** The results of genotyping peripheral blood cells isolated from the indicated 4 groups of chimeric mice with bone marrow reconstitution. WT-WT: WT mice reconstituted with WT bone marrow cells; KO-WT: WT mice reconstituted with *Ythdf1* KO bone marrow cells; WT-KO: *Ythdf1* KO mice

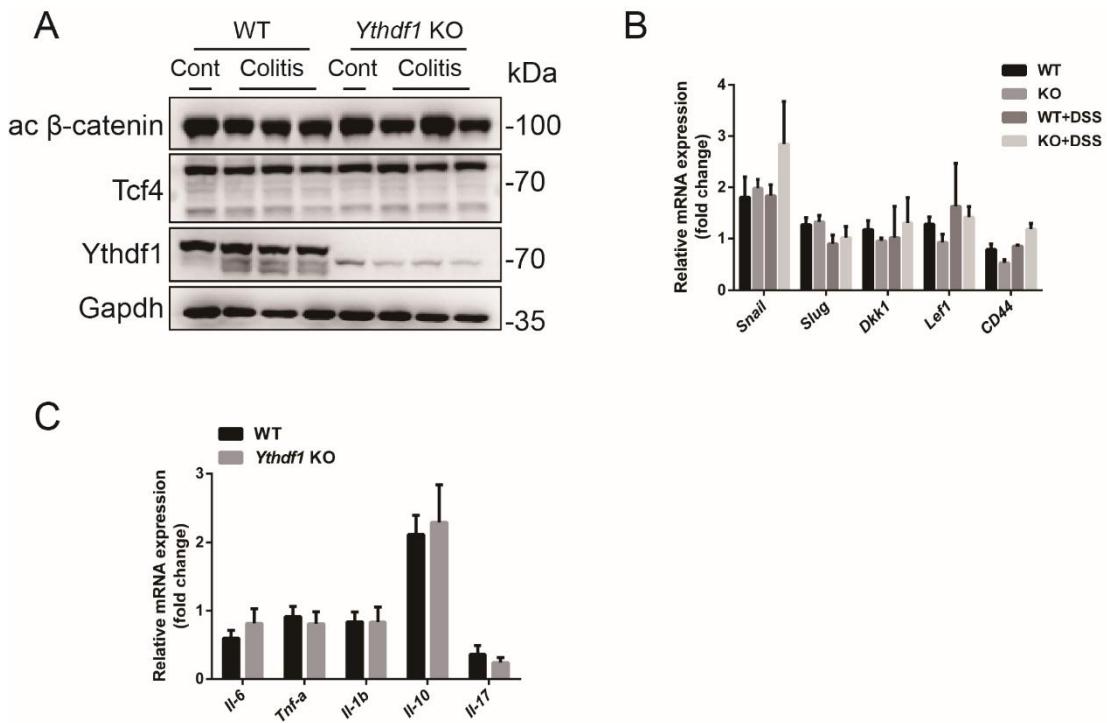
reconstituted with WT bone marrow cells; KO-KO: *Ythdf1* KO mice reconstituted with *Ythdf1* KO bone marrow cells. Three mice per group were analysed.



**Figure S4.** (A) Western blot analysis of Stat3, NF-κB, and MAPK pathways in colon epithelial cells collected from control and DSS-induced colitis mouse models. One lane represents one sample from an individual mouse; (B) qRT-PCR analysis of *Jak1* mRNA level in colon epithelial cells before and after the induction of colitis ( $n = 5$  for WT and WT+DSS,  $n = 3$  for KO and KO+DSS); (C) qRT-PCR analysis of *JAK1*, *JAK2* and *GP130* expression in HCT116 cells treated with scramble or si*YTHDF1* siRNA; (D-E) RNA lifetime for *JAK2* and *GP130* in shNC and sh*YTHDF1*-transfected HCT116 cells. TI, transcription inhibition;  $t_{1/2}$  means RNA lifetime. Data are presented as the means  $\pm$  s.e.m.



**Figure S5.** Representative TUNEL-stained apoptotic cells (upper; scale bar, 50  $\mu$ m) and IHC analysis of Ki-67 (lower; scale bar, 20  $\mu$ m) in WT and *Ythdf1* KO colon sections prior to colitis.



**Figure S6.** (A) Western blot analysis of activated  $\beta$ -catenin and Tcf4 in control and DSS-induced colitis samples. One lane represents one sample from an individual mouse. The protein lysates are the same as those used in Figure 4A, and the *Ythdf1* blot is identical to that shown in Figure 4A; (B) qRT-PCR analysis of the expression of Wnt/ $\beta$ -catenin target genes before and after the induction of colitis in colon epithelial cells ( $n = 3$  for each group); (C) qRT-PCR analysis of genes encoding inflammatory cytokines in WT and *Ythdf1* KO colon tumour tissues ( $n = 6$  for each group). Data are presented as the means  $\pm$  s.e.m.

**Table S1.** Primer sequences used for qRT-PCR and siRNA oligos.

Gene name (mouse)	Sense (5'-3')	Anti-sense (5'-3')
Il-6	CTGCAAGAGACTTCCATCCAGTT	GGGAAGGCCGTGGTTGTC
Tnf- $\alpha$	TCAAGGACTCAAATGGGCTTTC	TGCAGAACTCAGGAATGGACAT
Cxcl1	ACTGCACCCAAACCGAAGTC	TGGGGACACCTTTAGCATCTT
Ccl2	TAAAAACCTGGATCGAACCAAA	GCATTAGCTTCAGATTACGGGT
Il-10	CTTACTGACTGGCATGAGGATCA	GCAGCTCTAGGAGCATGTGG
Il-17	GCTCCAGAAGGCCCTCAGA	CTTCCCTCCGCATTGACA
Bclxl	GCTGCATTGTTCCCGTAGAG	GTTGGATGCCACCTATCTG
Survivin	CAGGGGAGTGCTTCTATGC	TACCGAGAACGAGCCTGATT
Ccnd1	TGACTGCCGAGAAGTTGTGC	CTCATCCGCCTCTGGCATT
c-Myc	CCCTATTTCATCTGCGACGAG	GAGAAGGACGTAGCGACCG
Jak1	AGTGCAGTATCTCTCCTCTG	GATTGGTTGGAGCGTACCC
Jak2	GGAATGGCCTGCCTTACAATG	TGGCTCTATCTGCTTCACAGAAT
Bcl2	ATGCCTTGTGGAACTATATGGC	GGTATGCACCCAGAGTGATGC
Snail	CACACGCTGCCCTGTGTCT	GGTCAGAAAAGCACGGTT
Slug	CAGCGAACTGGACACACACA	ATAGGGCTGTATGCTCCGAG
CD44	AGCAAGAAGGGCGAGTATAG	CTGTTCAGTAGGAAGGTAGGTG
Lef1	GCCACCGATGAGATGATCCC	TTGATGTCGGCTAAGTCGCC
Dkk1	CAGTGCCACCTGAACTCAGT	CCGCCCTCATAGAGAACTCC
Gapdh	TGGCCTCCGTGTTCCCTAC	GAGTTGCTGTTGAAGTCGCA

Gene name (human)	Sense (5'-3')	Anti-sense (5'-3')
JAK1	CTTGCCCTGTATGACGAGAAC	ACCTCATCCGGTAGTGGAGC
JAK2	TCTGGGGAGTATGTTGCAGAA	AGACATGGTGGGTGGATACC
GP130	CGGACAGCTTGAACAGAAATGT	ACCATCCCACACACCTCA
GAPDH	ACAACTTGGTATCGTGGAAAGG	GCCATCACGCCACAGTTTC
Gene name (human)	siRNA sequences (5'-3')	
YTHDF1	GCUCCAUAAGUACUCCAU	
METTL3	AGGAGCCAGCCAAGAAATCAA	