### **Supplementary information**

# Distribution and characterization of extrachromosomal circular DNA in colorectal cancer

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#### Supplementary Fig. 1-3

**Supplementary Fig. 1 Length distribution of eccDNA in colorectal cancer and normal tissues. a** Histogram of the reads mapping rate of eccDNA in each tumor group and normal group. **b-f** Cumulative frequency plots of eccDNA in tumor group and normal group.

**Supplementary Fig. 2 Chromosomal distribution and origin of genes highlyexpressed both in Circle-seq and RNA-seq. a** Heatmap of 219 genes highly-expressed both in Circle-seq and RNA-seq in each group. **b** Genomic distribution of 219 genes highly-expressed both in Circle-seq and RNA-seq.

**Supplementary Fig. 3 Verification of eccDNA in colorectal cancer cell lines. a** Gel electrophoresis result of the total DNA. HG: genome DNA of HCT116. HLE: total DNA with linear DNA elimination of HCT116. LG: genome DNA of LoVo. LLE: total DNA with linear DNA elimination of LoVo. **b** Schematic representation of the sequencing results of the PCR products of gene TP53I3 in samples after RCA with matching the sequence in chr2. **c** DNA electrophoresis of inverse PCR for junction sites of synthetic eccDNAs to prove transfection effectivity in HCT116. G: genome DNA. LE: total DNA with linear DNA elimination. RCA: total DNA with linear DNA elimination after RCA.





#### **Supplementary Fig. 2**



#### b

Supplementary Fig. 3



# Supplementary table 1-5

## Supplementary Table 1

## Sequences of inverse primers<sup>1</sup> and PCR reaction<sup>2</sup>

Gene	Primer	Sequences	PCR conditions
ERBB2	FRD:	5'-GCCGTGGCTCACACCTGTAATC-3'	
	REV:	5'-AGCAAGTTCCCAGACCTCCAGG-3'	1.98°C for 2 min
МҮСВРАР	FRD:	5'-TTCGGCAGAGTGGCGGCAAT-3'	2. 35 cycles of:
	REV:	5'-AGAGCAAACAACCCAGCAGAGC-3'	00°C for 10 and
TP53I3	FRD:	5'-AACCTCTTGGCGGGCGGATT-3'	98 C 101 10 sec
	REV:	5'-TGATCCTCTTCGGCGTTGTCCT-3'	60°C for 15 sec
EGFR	FRD:	5'-ACCTCCATCAGTGGCGATCTCC-3'	
	REV:	5'-GCCGCAATGTGGACAATACAGG-3'	72°C for 10 sec
SLC7A1	FRD:	5'-GGCAGGCTTGCTTTAGCATTGA-3'	2 70% 6 5 .
	REV:	5'-TGTGAGAGGCATGTTCGTGTCT-3'	$3.72^{\circ}$ for 5 min
SLC29A1	FRD:	5'-CCTGCCCTGGGTGGTATGAACT-3'	4. 4°C for save
	REV:	5'-GCTAAAGGCTTGGTTCGGAGGC-3'	

<sup>1</sup> Oligonucleotide primers were designed by Primer Premier 6.0

<sup>2</sup> PCR reaction was performed on ProFlex<sup>TM</sup> PCR System (Thermo Fisher)

#### eccβ-actin: 358bp

#### eccSLC7A1-exon: 335bp

CCCCACCACCCTGCAATTTCCCCGTCCTCGGAACTGTCTCACGAGACCACCAGCGC ACAGACACGAACATGCCTCTCACACACCCCAGAACGCAGCTGTGCTGGAGTTGGGC ACAAAGGCCCCTCACTCCCTGTCTTCATGGTGTGGGGTTCTTGGCTGCCAACAGAACT GTCAGCCTTTATTAAGAAGGAAGGGTCTGTGTAAAAATGGGATGTGTGAATCAGAA GAGGGGCCCTGGCACAGAGCCAGGCAGGCTTGCTTTAGCATTGAGTCCTGCGTCGC GCAATGCCCAGGACCTTGCCTGTTAAGTCTGGGTGGGGCCCCCTATGGGGGCATGT

#### eccSLC7A1-CpG: 382bp

#### eccSLC7A1-5'UTR: 387bp

#### eccSLC7A1-3'UTR: 385bp

Name	Primer	Sequences	PCR conditions
eccβ-actin	FRD:	5' CCAGAGGCGTACAGGGATAGCA 3'	
	REV:	5' GCGGGAAATCGTGCGTGACAT 3'	1.98°€ for 2 min
eccSLC7A1-exon	FRD:	5' CCCCACCACCCTGCAATTTCCC 3'	2. 50 cycles of:
	REV:	5' TGTGAGAGGCATGTTCGTGTCT 3'	98°C for 10 sec
eccSLC7A1-CpG	FRD:	5' CACTGGTCCAAGTTGCCGTCAG 3'	60°C for 15 sec
	REV:	5' CAGAAACCACCTGCGTCCACAC 3'	72°C for 10 sec
eccSLC7A1- 5'UTR	FRD:	5' TTGCTGCAAGATCGAGGCTGTC 3'	2 72°C for 5 min
	REV:	5' AGAGGAGTGGAGGCTGCGGTTA 3'	5. 72 C 101 5 mm
eccSLC7A1- 3'UTR	FRD:	5' CAAGTGTAGTTGTCGTCCACCA 3'	4.4°C for save
	REV:	5' CACAACTCTTAATAATGGCTCC 3'	

Sequences of primers<sup>1</sup> and PCR reaction<sup>2</sup>

<sup>1</sup> Oligonucleotide primers were designed by Primer Premier 6.0

<sup>2</sup> PCR reaction was performed on ProFlex<sup>TM</sup> PCR System (Thermo Fisher)

Name	Primer	Sequences	PCR conditions
eccβ-actin	FRD:	5' GCTCATTGCCAATGGTGATGAC 3'	
	REV:	5' GCTATCCAGGCTGTGCTATCCC 3'	1.08℃ for 2 min
eccSLC7A1-exon	FRD:	5' GGCAGGCTTGCTTTAGCATTGA 3'	2. 50 cycles of:
	REV:	5' ACATGCCCCATAGGGGGCCCCAC 3'	98°C for 10 sec
eccSLC7A1-CpG	FRD:	5' CCCGGAGAACCGGCTGCAGAGC 3'	60°C for 15 sec
	REV:	5' GGTTTGCTGTGGGACCGCCCTG 3'	72°C for 10 sec
eccSLC7A1- 5'UTR	FRD:	5' GCACTGCTGATGAAACCTGGCG 3'	2 72°C for 5 min
	REV:	5' GTTGCTGTTCAGAGCTGGAATA 3'	5. 72 C 101 5 mm
eccSLC7A1- 3'UTR	FRD:	5' TTCTTCTACTGAACATGGAGCC 3'	4.4°C for save
	REV:	5' GGAAGCCATAGAATTTATTCGA 3'	

Sequences of inverse primers<sup>1</sup> and PCR reaction<sup>2</sup>

<sup>1</sup> Oligonucleotide primers were designed by Primer Premier 6.0

<sup>2</sup> PCR reaction was performed on ProFlex<sup>TM</sup> PCR System (Thermo Fisher)

Gene	Primer	Sequences	RT-qPCR conditions
SLC7A1	FRD:	5'GCCTGTGCTATGGCGAGTTT3'	1. 95℃ for 3 min
	REV:	5'ACGCTTGAAGTACCGATGATGTA3'	2. 50 cycles of: 95℃ for 5 sec
xβ-actin	FRD:	5'TGCTGTCCCTGTATGCCTCTGG3'	60°C for 30 sec
	REV:	5'GGAACCGCTCGTTGCCAATAGT3'	3. 95°C for 5 sec
GAPDH	FRD:	5'CGGATTTGGTCGTATTGGG3'	60℃ for 1 min
	REV:	5'CTGGAAGATGGTGATGGGATT3'	4. 50°C for 30 sec

## Sequences of primers<sup>1</sup> and RT-qPCR reaction<sup>2</sup>

<sup>1</sup> Oligonucleotide primers were designed by Primer Premier 6.0

<sup>2</sup> RT-qPCR reaction was performed on ROCHE LightCycler480 System (Rotor gene

6000 Software, Sydney, Australia).