

Table SI. Function of circRNAs in gastric cancer.

First author, year	circRNA	Dysregulation	Function	Mechanism	(Refs.)
Zhang <i>et al</i> , 2019	circNRIP1	Up	miRNA sponge	circNRIP1 sponges miR-149-5p to affect the expression of AKT1	(1)
Zhu <i>et al</i> , 2019	circMLLT10	Up	miRNA sponge	circMLLT10 acts as a miR-509-3-5p sponge to attenuate its suppressive effect on target GINS4	(2)
Xie <i>et al</i> , 2020	circSHKBP1	Up	miRNA sponge	circSHKBP1 regulates the miR-582-3p/HUR/VEGF pathway, and suppresses HSP90 degradation	(3)
Rong <i>et al</i> , 2019	circPSMC3	Down	miRNA sponge	Acting as a competitive endogenous RNA through sponging miR-296-5p	(4)
Huang <i>et al</i> , 2019	circAKT3	Up	miRNA sponge	circAKT3 could promote PIK3R1 expression by sponging miR-198	(5)
Wang <i>et al</i> , 2019	circOSBPL10	Up	miRNA sponge	Via a circOSBPL10/miR-136-5p/WNT2 axis	(6)
Wang <i>et al</i> , 2019	circLMTK2	Up	miRNA sponge	circLMTK2 functions through the miR-150-5p/c-Myc axis	(7)
Zhu <i>et al</i> , 2019	circNHSL1	Up	miRNA sponge	circNHSL1 acts through the miR-1306-3p/SIX1/Vimentin axis	(8)
Liang <i>et al</i> , 2019	hsa_circ_006100	Up	miRNA sponge	hsa_circ_006100 functions via miR-195/GPRC5A signaling	(9)
Lin <i>et al</i> , 2020	circRNA_100876	Up	miRNA sponge	circRNA_100876 exerts its effects through miR-665/YAP1 signaling	(10)
Peng <i>et al</i> , 2020	circCUL2	Down	miRNA sponge	circCUL2 functions through miR-142-3p/ROCK2-mediated autophagy activation	(11)
Zhang <i>et al</i> , 2019	circDLST	Up	miRNA sponge	Sponging miR-502-5p to activate the NRAS/MEK1/ERK1/2 signaling	(12)
Liu <i>et al</i> , 2018	circYAP1	Down	miRNA sponge	circYAP1 functions by targeting the miR-367-5p/p27Kip1 axis	(13)
Wang <i>et al</i> , 2020	circR-RNF111	Up	miRNA sponge	Binding to miR-27b-3p	(14)
Cai <i>et al</i> , 2020	circ_0000267	Up	miRNA sponge	Modulating the miR-503-5p/HMGA2 axis	(15)
Liu <i>et al</i> , 2018	circ-SERPINE2	Up	miRNA sponge	Through the regulation of miR-375/YWHAZ	(16)
Deng <i>et al</i> , 2020	circRHOBTB3	Down	miRNA sponge	Acting as a sponge for miR-654-3p	(17)
Cai <i>et al</i> , 2020	circRNA_0005529	Up	miRNA sponge	Regulating miR-527/Spl axis	(15)
Li <i>et al</i> , 2020	circ_0008035	Up	miRNA sponge	Upregulating EIF4A1 through sponging miR-599	(18)
Luo <i>et al</i> , 2020	circCCDC9	Down	miRNA sponge	circCCDC9 functions through the miR-6792-3p/CAV1 axis	(19)
Zhang <i>et al</i> , 2021	circHN1	Up	miRNA sponge	Binding to miR-1248 and miR-375	(20)
Wu <i>et al</i> , 2020	circRNA_0005075	Down	miRNA sponge	Regulating miR-431/p53/epithelial-mesenchymal transition axis in gastric cancer	(21)
Du <i>et al</i> , 2019	circ_PRMT5	Up	miRNA sponge	Targeting the miR-145/miR-1304/MYC axis	(22)
Peng <i>et al</i> , 2020	circ-ITCH	Down	miRNA sponge	Through the Wnt/β-catenin signaling pathway by sequestering miR-17	(23)
Wang <i>et al</i> , 2020	circ_0000190	Down	miRNA sponge	Inhibiting the miR-1252/PAK3 pathway	(24)
Xia <i>et al</i> , 2020	circSMC3	Up	miRNA sponge	Through miR-4720-3p/TJP1 axis	(25)
Zhang <i>et al</i> , 2020	circDUSP16	Up	miRNA sponge	Sponging the miR-145-5p/ IVNS1ABP axis	(26)

Ma <i>et al</i> , 2020	hsa_circ_0004872	Down	miRNA sponge	Via the miR-224/Smad4/ADAR1 successive regulatory circuit	(27)
Xia <i>et al</i> , 2020	circPDZD8	Up	miRNA sponge	Regulating CHD9 via sponging miR-197-5p	(28)
Liu <i>et al</i> , 2020	circ_OXCT1	Down	miRNA sponge	Through the circ-OXCT1/miR-136/SMAD4 axis	(29)
Wang <i>et al</i> , 2020	circ_0027599	Down	miRNA sponge	PHLDA1 is regulated by circ_0027599/miR-101	(30)
Zhang <i>et al</i> , 2019	circCACTIN	Up	miRNA sponge	Sponging miR-331-3p and regulating TGFBR1 expression	(31)
Liang <i>et al</i> , 2020	circNRIP1	Up	miRNA sponge	By miR-182/ROCK1 axis	(32)
Zhou <i>et al</i> , 2020	circ_002117	Down	miRNA sponge	Upregulating HERPUD1 through miR-370 inhibition	(33)
Dai <i>et al</i> , 2019	circGRAMD1B	Down	miRNA sponge	Sponging miR-130a-3p and regulating PTEN and p21 expression	(34)
Lin <i>et al</i> , 2020	circRIMS	Up	miRNA sponge	As a sponge for hsa-miR-148a-5p and hsa-miR-218-5p	(35)
Cai <i>et al</i> , 2019	circHECTD1	Up	miRNA sponge	Targeting miR-1256 and activating β -catenin/c-Myc signaling	(36)
Wang <i>et al</i> , 2020	circ_0001023	Up	miRNA sponge	Regulating the miR-409-3p/PHF10 axis	(37)
Zhang <i>et al</i> , 2020	hsa_circ_0023642	Up	miRNA sponge	Sponging microRNA-223	(38)
Lai <i>et al</i> , 2019	circRNA0047905	Up	miRNA sponge	Acting as a sponge for miR-4516 and miR-1227-5	(39)
Wei <i>et al</i> , 2020	circRNA_104433	Up	miRNA sponge	Targeting miR-497-5p, which directly regulates CDC25A	(40)
Ma <i>et al</i> , 2020	circPIP5K1A	Up	miRNA sponge	Via miR-376c-3p/ZNF146 axis	(41)
Mo <i>et al</i> , 2020	hsa_circ_0000467	Up	miRNA sponge	Binding to miR-326-3p	(42)
Niu <i>et al</i> , 2020	hsa_circ_0001829	Up	miRNA sponge	Through miR-155-5p/SMAD2 axis	(43)
Wei <i>et al</i> , 2020	circHIPK3	Up	miRNA sponge	Sponging miR-107 and regulating BDNF expression	(44)
Wang <i>et al</i> , 2020	hsa_circ_0003159	Down	miRNA sponge	Regulating the miR-223-3p/NDRG1 axis	(45)
Chen <i>et al</i> , 2020	circ_0032821	Up	miRNA sponge	Regulating the miR-1236-3p/HMGB1 axis	(46)
Shen <i>et al</i> , 2020	hsa_circ_0005556	Up	miRNA sponge	Sponging miR-4270 to increase MMP19 expression	(47)
Cao <i>et al</i> , 2019	hsa_circ_0000291	Up	miRNA sponge	Targeting the miR-183/ITGB1 axis	(48)
Sun <i>et al</i> , 2020	circMAN2B2	Up	miRNA sponge	Regulating miR-145, as well as PI3K/AKT and JNK pathways	(49)
Li <i>et al</i> , 2020	circHIPK3	Up	miRNA sponge	Through regulation of the miR-876-5p/PIK3R1 axis	(50)
Liu <i>et al</i> , 2020	circ-NRIP1	Up	miRNA sponge	Modulating MYH9 via miR-186-5p	(51)
Hui <i>et al</i> , 2020	circNHSL1	Up	miRNA sponge	Through the miR-149-5p/YWHAZ axis	(52)
Liu <i>et al</i> , 2020	circ-MAT2B	Up	miRNA sponge	Regulating the miR-515-5p/HIF-1 α axis	(53)
Sun <i>et al</i> , 2018	circ-SFMBT2	Up	miRNA sponge	Sponging miR-182-5p to enhance CReB1 expression	(54)
Zhang <i>et al</i> , 2019	hsa_circ_0067997	Up	miRNA sponge	Regulating the miR-515-5p/XIAP axis	(55)
Zhang <i>et al</i> , 2020	circATXN7	Up	miRNA sponge	Sponging miR-4319 and regulating ENTPD4	(56)
Xue <i>et al</i> , 2019	hsa_circ_0081143	Up	miRNA sponge	Targeting the miR-646/CDK6 pathway	(57)
Wang <i>et al</i> , 2019	circNF1	Up	miRNA sponge	Absorbing miR-16	(58)
Jin <i>et al</i> , 2020	circHIPK3	Up	miRNA sponge	Via the miR-653-5p/miR-338-3p-NRP1 axis	(59)
Chen <i>et al</i> , 2019	hsa_circ_0092306	Up	miRNA sponge	Regulating the pathway of miR-197-3p/PRKCB in MKN-45 cells	(60)

Qu <i>et al</i> , 2020	circFLNA	Up	miRNA sponge	Regulating the miR-646/PFKFB2 axis	(61)
Ma <i>et al</i> , 2021	circPTPN22	Up	miRNA sponge	Through the competitive binding of miRNA to inhibit the epithelial-mesenchymal transition pathway	(62)
Zhu <i>et al</i> , 2020	circKIAA0907	Down	miRNA sponge	Via the miR-452-5p/KAT6B axis	(63)
Lin <i>et al</i> , 2020	circCYFIP2	Up	miRNA sponge	Via the miR-1205/E2F1 axis	(64)
Li <i>et al</i> , 2021	hsa_circ_0023409	Up	miRNA sponge	Activating the IRS4/PI3K/AKT pathway by acting as a sponge for miR-542-3p	(65)
Mi <i>et al</i> , 2020	circ 0000144	Up	miRNA sponge	Modulating GPRC5A expression by acting as an miR-623 sponge	(66)
Jin <i>et al</i> , 2020	circ_C16orf62	Down	miRNA sponge	By the miR-421/TUBB2A axis	(67)
Guo <i>et al</i> , 2021	circREPS2	Down	miRNA sponge	Via miR-558/RUNX3/β-catenin signaling	(68)
Pu <i>et al</i> , 2020	circCUL3	Up	miRNA sponge	Accelerating the Warburg effect progression through regulating the STAT3/HK2 axis	(69)
Yue <i>et al</i> , 2021	circ_0004104	Up	miRNA sponge	Regulating the miR-539-3p/RNF2 axis	(70)
Yu <i>et al</i> , 2021	circNEK9	Up	miRNA sponge	Targeting the miR-409-3p/MAP7 axis	(71)
Yang <i>et al</i> , 2021	circ_0044516	Up	miRNA sponge	Modulating the miR-149-5p/HuR axis	(72)
Wang <i>et al</i> , 2021	circ_PGPEP1	Up	miRNA sponge	Sponging miR-1297 and regulating E2F3	(73)
Xu <i>et al</i> , 2020	hsa-circ-0007766	Up	miRNA sponge	Regulating GDF15 via the hsa-circ-0007766/miR-1233-3p/GDF15 axis	(74)
Cao <i>et al</i> , 2021	circLMO7	Up	miRNA sponge	Acting as an miR-30a-3p sponge and affecting the WNT2/β-catenin pathway	(75)
Wang <i>et al</i> , 2021	circ_SMAD4	Up	miRNA sponge and interaction with protein	Activating CTNNB1-dependent WNT/β-catenin signaling by interacting with miR-1276 and TCF4	(76)
Wang <i>et al</i> , 2021	circ_ITCH	Down	miRNA sponge	Regulating the miR-199a-5p/Klotho axis	(77)
Xu <i>et al</i> , 2021	circ_0081146	Up	miRNA sponge	Sponging miR-144 and upregulating HMGB1	(78)
Hua <i>et al</i> , 2021	circ_0006282	Up	miRNA sponge	Regulating miR-144-5p/tyrosine 3-monooxygenase/tryptophan 5-monooxygenase activation protein β axis	(79)
Wang <i>et al</i> , 2021	hsa_circ_0009172	Down	miRNA sponge	Regulation of miR-485-3p-mediated NTRK3	(80)
Deng <i>et al</i> , 2021	circVAPA	Up	miRNA sponge	Regulating the miR-125b-5p/STAT3 axis	(81)
Li <i>et al</i> , 2021	circ_002059	Down	miRNA sponge	Via the miR-182/MTSS1 axis	(82)
Xia <i>et al</i> , 2021	circFAM73A	Up	miRNA sponge and interaction with protein	Through the miR-490-3p/HMGA2 positive feedback loop and HNRNPK-mediated β-catenin stabilization	(83)
Bu <i>et al</i> , 2021	circAFF2	Up	miRNA sponge	Via miR-6894-5p/ANTXR 1 signaling	(84)
Chen <i>et al</i> , 2021	circACC1	Up	miRNA sponge	Via the circACC1/miR-29c-3p/FOXP1 network	(85)

Li <i>et al</i> , 2021	circ-sirt1	Down	miRNA sponge	Sponging miR-132-3p/miR-212-3p and upregulating sirt1 expression	(86)
Han <i>et al</i> , 2021	circ_0027599	Down	miRNA sponge	Via modulation of the miR-21-5p/RUNX1 axis	(87)
Ye <i>et al</i> , 2021	hsa_circ_0001874	Down	miRNA sponge	Sponging miR-593-5p, miR-103a-3p and miR-107	(88)
Lu <i>et al</i> , 2021	circVPS33B	Up	miRNA sponge	Regulating the miR-873-5p/HNRNPK axis	(89)
Fan <i>et al</i> , 2021	circ_CORO1C	Up	miRNA sponge	Modulating miR-138-5p/KLF12 axis	(90)
Lu <i>et al</i> , 2021	circHECTD1	Up	miRNA sponge	Via the miR-137/PBX3 axis	(91)
Zhao <i>et al</i> , 2021	circATP2B1	Up	miRNA sponge	Decreasing the suppression of PKM2 by miR-326-3p/miR-330-5p	(92)
Wang <i>et al</i> , 2021	circBFAR	Up	miRNA sponge	Sponging miR-513a-3p/hexokinase 2 axis	(93)
Li <i>et al</i> , 2021	circ_0044366	Up	miRNA sponge	Targeting the miR-29a/VEGF axis	(94)
Gao <i>et al</i> , 2021	hsa_circ_0000117	Up	miRNA sponge	Sponging miR-337-3p and increasing STAT3 expression	(95)
Wang <i>et al</i> , 2021	hsa_circRNA_100269	Down	miRNA sponge	Inactivating the PI3K/Akt axis	(96)
Xu <i>et al</i> , 2021	circTMC5	Up	miRNA sponge	Targeting miR-361-3p/RABL6	(97)
Hare <i>et al</i> , 2021	hsa-circ_0000064	Up	miRNA sponge	Modulating the miR-621/SYF2 axis	(98)
Qiang <i>et al</i> , 2021	circCSNK1G1	Up	miRNA sponge	Via the miR-758/ZNF217 axis	(99)
Wu <i>et al</i> , 2021	circALPL	Up	miRNA sponge	Sponging miR-127, thus upregulating MTDH	(100)
Jiang <i>et al</i> , 2021	CDR1as	Down	miRNA sponge	Sponging miR-876-5p to upregulate GNG7 expression	(101)
Wang <i>et al</i> , 2021	circRNA_100290	Up	miRNA sponge	Via the miR-29b-3p/ITGA11 axis	(102)
Liang <i>et al</i> , 2021	hsa_circ_0110389	Up	miRNA sponge	Upregulating SORT1 via sponging miR-127-5p and miR-136-5p	(103)
Yang <i>et al</i> , 2021	circHIPK3	Up	miRNA sponge	Through the miR-637/AKT1 pathway	(104)
Qiu <i>et al</i> , 2022	circTHBS1	Up	miRNA sponge	Sponging miR-204-5p to promote the expression of INHBA	(105)
Yang <i>et al</i> , 2019	circ-HuR	Down	Interaction with protein	Inhibiting CNBP transactivation	(106)
Zang <i>et al</i> , 2022	circEIF4G3	Down	Interaction with protein	Through the regulation of δ-catenin protein stability and the miR-4449/SIK1 axis	(107)
Ma <i>et al</i> , 2022	circARID1A	Up	Interaction with protein	Forming a circARID1A-IGF2BP3-SLC7A5 RNA-protein ternary complex	(108)
Ding <i>et al</i> , 2019	circ-DONSON	Up	Regulating transcription	Recruiting the NURF complex to initiate SOX4 expression	(109)
Jie <i>et al</i> , 2020	circMRPS35	Down	Chromatin remodeling	Recruiting KAT7 to govern histone modification	(110)
Jiang <i>et al</i> , 2021	circMAPK1	Down	Encoding protein	Suppressing activation of MAPK signaling	(111)

miR/miRNA, microRNA; circRNA, circular RNA.

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Table SII. circRNAs as biomarkers in gastric cancer.

First author, year	circRNA	Sample	Dysregulation	Function	Sensitivity, %	Specificity, %	AUC	Cutoff value	Clinical association	(Refs.)
Tang <i>et al</i> , 2018	circ-KIAA1244	Tissues	Down	Independent prognostic indicator	77.42	0.68	0.7481	1.443	TNM stage, lymphatic metastasis and overall survival time	(1)
Zhao <i>et al</i> , 2018	hsa_circ_0000181	Plasma Tissues	Down	Diagnosis biomarker	0.21 0.85	0.99 0.54	0.5820 0.7560	7.270 9.400	Differentiation and CEA Tumor diameter, lymphatic metastasis, distal metastasis and CA19-9	(2)
Rong <i>et al</i> , 2018	circ_0066444	Tissues	Up	Diagnosis biomarker	0.71	0.69	0.7328	-	Lymphatic metastasis	(3)
Cai <i>et al</i> , 2019	circSMARCA5	Tissues	Down	Diagnosis and prognosis biomarker	-	-	0.8060	-	Tumor differentiation, LNM, vascular invasion, and AJCC stage	(4)
Lu <i>et al</i> , 2019	hsa_circ_0000467	Tissues	Up	Diagnosis and prognostic biomarker	0.71	0.65	0.7900	-	Lymphatic invasion and TNM stage	(5)
Lu <i>et al</i> , 2019	hsa_circ_0006848	Plasma	Down	Diagnosis biomarker	-	-	0.7330	-	Histological grade and tumor size	(6)
Lu <i>et al</i> , 2019	hsa_circ_0067582	Tissues	Down	Diagnosis biomarker	0.55	0.75	0.6710	10.610	Tissue CEA level and stages	(7)
	hsa_circ_0005758	Tissues	Down	Diagnosis biomarker	0.75	0.68	0.7210	10.200	Tissue CEA level and perineural invasion	
Yang <i>et al</i> , 2019	hsa_circ_0005556	Tissues	Down	Diagnosis and prognosis biomarker	0.64	0.82	0.7730	-	Differentiation, TNM stage, and lymphatic metastasis	(8)
Wei <i>et al</i> , 2020	hsa_circRNA_1029 58	Tissues	Up	Diagnosis biomarker	0.61	86.00	0.7400		TNM stage	(9)
Shao <i>et al</i> , 2020	hsa_circ_0065149	Tissues	Down	Diagnosis and prognosis biomarker	0.79	0.62	0.7690	10.190	Tumor diameter, perineural invasion and overall survival time	(10)
Xu <i>et al</i> , 2020	circ_0004771	Plasma	Up	Diagnosis and prognosis	0.68	0.79	0.8310	-	Differentiation grade, Lymph node metastasis, TNM stage and	(11)

Yu <i>et al</i> , 2020	hsa_circ_0067582	Tissues	Down	biomarker Diagnosis and prognosis biomarker	0.67	0.61	0.6937	-	T stage Tumor diameter and carbohydrate antigen 19-9	(12)
Zhang <i>et al</i> , 2020	hsa_circ_0001811	Tissues	Down	Diagnosis biomarker	0.50	0.81	0.6580	-	CEA, tissue differentiation, and lymph node metastasis	(13)
		Plasma			0.91	0.52	0.7470	-	CA19-9, lymph node metastasis, distant metastasis, and age	
Zhang <i>et al</i> , 2020	circ_0026344	Tissues	Down	Diagnostic biomarker	-	-	-	-	Tumor size, lymph node metastasis, TNM stage, invasive depth and overall survival time	(14)
Tang <i>et al</i> , 2021	circ_0049447	Tissues	Down	Diagnosis biomarker	0.82	0.77	0.8380	-	Proliferation, migration, invasion and EMT	(15)
Yao and Xie, 2021	hsa_circ_0006470	Tissues	Down	Diagnosis biomarker	0.73	0.75	0.7830	10.740	TNM stage and invasion	(16)
Ye <i>et al</i> , 2021	hsa_circ_0001874	Tissues	Down	Diagnostic and prognosis biomarker	0.62	0.68	0.6730	-	Cell differentiation, tumor stage, invasion, lymphatic metastasis and CEA level	(17)

circRNA, circular RNA; AUC, area under the curve; TNM, Tumor-Node-Metastasis; CEA, carcinoembryonic antigen; AJCC, American Joint Committee on Cancer; EMT, epithelial-mesenchymal transition.

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