Diagnostic and prognostic value of circulating tumor DNA in gastric cancer: a meta-analysis

Supplementary Materials

Supplementary Table S1: Information of included genes and their main functions

	Full name	Gene function	Gene location	
APC	APC	Signal transduction	5q22.2	
BCL6B	B-cell CLL/lymphoma 6B	Signal transduction	17p13.1	
E-cadherin	E-cadherin	Adhesion/invasiveness	16q22.1	
MINT2	Amyloid beta precursor protein binding family A member 2	Signal transduction	15q11-q12	
OSR2	Odd-skipped related transciption factor 2	Apoptosis/Proliferation	8q22.2	
P16	Cyclin dependent kinase inhibitor 2A	Signal transduction	9p21	
PCDH10	Protocadherin 10	Adhesion/Invasiveness	4q28.3	
PPFIA3	PTPRF interacting protein alpha 3	Apoptosis/Proliferation	19q13.33	
RASSF1A	Ras association domain family member 1	Apoptosis/Proliferation	3p21.3	
RUNX3	Runt related transcription factor 3	Apoptosis/Proliferation	1p36	
SFRP1	Secreted frizzled related protein 1	Signal transduction	8p11.21	
SOX17	SRY-box 17	Adhesion/Invasiveness	8q11.23	
SPG20	Spastic paraplegia 20	Signal transduction	13q13.3	
TFPI2	Tissue factor pathway inhibitor-2	Signal transduction	7q22	
TIMP3	TIMP metallopeptidase inhibitor 3	Adhesion/Invasiveness	22q12.3	
VAV3	Vav guanine nucleotide exchange factor 3	Signal transduction	1p13.3	
XAF1	XIAP associated factor 1	Apoptosis/Proliferation	17p13.1	

	Risk of Bias				Applicability Concerns			
Study	Patients Selection	Index Test	Reference Standard	Flow and Time	Patients Selection	Index Test	Reference Standard	
WK Leung ⁸	L	L	U	L	Н	L	L	
Mohammad R.A.9	L	L	L	U	L	L	L	
Wang YC ¹⁰	L	L	U	L	L	L	L	
Chouhei.S ¹¹	L	L	L	L	L H		L	
Kenji H. ¹²	Н	Н	L	L	Н	L	L	
Ioanna B.13	Н	L	L	L	Н	L	L	
Yang QF ¹⁴	L	L	L	L	L	L	L	
Zhi QL ¹⁵	L	L	L	L	L	L	L	
Han J ¹⁶	L	L	L	L	L	L	L	
Wu YC ¹⁷	L	L	L	Н	L	L	L	
Yu JL ¹⁸	L	L	L	L	Н	L	L	
Zhang H ¹⁹	L	L	L	L	L	L	L	
Chang L ²⁰	L	L	L	L	L	L	L	
Ioanna B. (2015) 21	Н	L	L	Н	Н	L	L	
Charinya P ²²	L	L	L	L	L	L	L	
Li WH ²³	L	L	L	L	L	L	L	

Supplementary Table S2: Results of diagnosis quality assessment of included studies according to the QUADAS-2 tool criteria L: low risk of bias; H: high risk of bias; U: unclear risk of bias

Study	Selection(0-4)			Comparability		Outcome (0–3)		Total		
	REC	Snec	AE	OINP	SCB	SCA	AO	FU	AFC	
WK Leung ⁸	1	0	1	1	1	0	1	1	1	7
Mohammad R.A.9	1	1	1	1	1	1	0	0	0	6
Wang YC ¹⁰	1	1	1	0	1	0	1	0	0	5
Chouhei.S11	1	1	1	0	1	0	1	0	0	5
Kenji H. ¹²	1	1	1	0	1	1	0	0	0	5
Ioanna B. ¹³	1	1	1	1	1	0	1	1	0	7
Yang QF ¹⁴	1	1	1	0	1	0	1	1	0	6
Zhi QL ¹⁵	1	1	1	1	1	1	1	1	0	8
Han J ¹⁶	1	1	1	0	1	0	1	1	0	6
Wu YC ¹⁷	1	1	1	0	1	1	1	1	0	7
Yu JL ¹⁸	1	1	1	1	1	1	1	1	0	8
Zhang H ¹⁹	1	1	1	0	1	0	1	1	0	7
Chang L ²⁰	1	1	1	0	1	0	1	0	0	5
Ioanna B. (2015) ²¹	1	1	1	1	1	0	1	1	1	8
Charinya P ²²	1	1	1	1	1	1	1	1	0	8
Li WH ²³	1	1	1	0	1	0	1	0	0	5

Supplementary Table S3: The assessment of quality of each study according to the newcastleottawa scale

REC: representativeness of the exposed cohort; SNEC: selection of the nonexposed cohort; AE: ascertainment of exposure; OINP: outcome of interest not presented in the start of study; SCB: study controls for basic characteristics; SCA: study controls for additional factor; AO: assessment of outcome; FU: follow-up; AFC: adequacy of follow up.