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Diagnostic value of circN4BP2L2 in type I and type II epithelial ovarian cancer

Li Ning¹, Jinghe Lang², Bo Long^{2*} and Lingying Wu^{1*}

Abstract

Background CircN4BP2L2 was previously identified to be significantly decreased in epithelial ovarian cancer (EOC) and was associated with disease progression. The aim of this study was to evaluate the diagnostic value of plasma circN4BP2L2 using the unifying model of type I and type II EOC.

Methods A total of 540 plasma samples were obtained from 180 EOC patients, 180 benign ovarian cyst patients, and 180 healthy volunteers. CircN4BP2L2 was assessed using reverse transcription-quantitative polymerase chain reaction (RT-qPCR). Cancer antigen 125 (CA125) and human epididymis protein 4 (HE4) were assessed using enzymelinked immunosorbent assay (ELISA). Receiver operating curve (ROC), the area under the curve (AUC), sensitivity and specificity were estimated.

Results Low level of circN4BP2L2 was associated with advanced tumor stage (p < 0.01) in type I EOC. Decreased circN4BP2L2 was associated with lymph node metastasis (LNM) (p = 0.04) in type II EOC. The expression level of circN4BP2L2 in type I was similar to that in type II. CircN4BP2L2 could significantly separate type I or type II from benign or normal cohort (p < 0.01). Early-stage type I or type II EOC vs. benign or normal cohort could also be distinguished by circN4BP2L2 (p < 0.01).

Conclusion CircN4BP2L2 might serve as a promising diagnostic biomarker for both type I and type II EOC. The diagnostic safety for circN4BP2L2 in early-stage type I or type II EOC is also acceptable. Further large-scale well-designed studies are warranted to investigate whether circN4BP2L2 is specific for all histologic subgroups.

Keywords Epithelial ovarian cancer, Circular RNA, Human, Biological markers, Diagnosis

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Background

Ovarian cancer (OC) is the most lethal malignancy in female reproductive system [1]. The number of new cases and deaths of OC worldwide was estimated at 295,000 and 187,000 annually [2]. Despite recent progress in surgical skills and targeted drugs, most advanced patients still develop tumor recurrence in three years after OC diagnosis, and relapsed diseases were usually incurable [3]. The 5-year survival rate of OC patients is merely 40% [3]. Epithelial ovarian cancer (EOC) is the most common subtype, accounting for approximately 95% of all OC cases [4]. Due to lacking of specific early symptoms and reliable screening tests, over 70% of EOC patients



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Ning et al. BMC Cancer (2022) 22:1210 Page 2 of 12

are diagnosed at advanced stages [5]. Owing to the great heterogeneity in biological and molecular status, EOC is actually consisting of many different tumors [6]. Based on morphologic features, the four major histological subtypes of EOC are, serous, mucinous, endometrioid and clear cell [6]. EOC has been thought to arise from epithelial cells lining the surface of the ovary [7]. Recent findings suggest that EOC may also arise from the epithelium of fallopian tube and endometriosis [8].

Based on distinct morphological features and molecular genetics, EOC is further divided into type I and type II tumors [9]. Type I tumors include low-grade (G1) serous, low-grade (G1) endometrioid, all clear cell, mucinous and Brenner carcinomas [10]. These tumors are generally indolent, more often diagnosed at stage I (tumor confined to the ovary), harbor specific gene mutations such as KRAS, CTNNB1 and ARID1A, have relatively stable genome and rarely have TP53 mutation [9-11]. Type II tumors include high-grade (G2-G3) serous, high-grade (G2-G3) endometrioid, all undifferentiated carcinoma and malignant mixed mesodermal tumors (carcinosarcomas) [10]. These tumors are more aggressive, more often present at advanced stage, harbor highly unstable genome, have a high frequency of TP53 mutation had have molecular alterations that causing BRCA dysfunction via gene mutation or promoter methylation [9, 10, 12].

Cancer antigen 125 (CA125) is the first biomarker for OC approved by the US Food and Drug Administration (FDA). Nevertheless, CA125 is limited by its relatively low sensitivity and low specificity [13]. It has been reported that CA125 was elevated in only 50–60% of early-stage tumors [14]. Human epididymal protein 4 (HE4) is the second FDA-approved OC biomarker [15]. Although HE4 has a relatively high specificity since it does not increase in benign gynecological diseases, it is also limited by its relatively low sensitivity (approximately 70%) [16].

Circular RNAs (circRNAs) are a novel subtype of noncoding RNAs [17]. Their high stability, abundance, conservation and tissue-specificity make them promising biomarkers for disease detection [18]. A body of clinical trials have been performed using blood samples to evaluate the diagnostic value of circRNAs in various diseases, including cancer. Ouyang et al. [19] reported that plasma circ_002453 was significantly increased in patients with lupus nephritis and might serve as a potential biomarker for disease diagnosis. Huang et al. [20] reported that plasma circ_0001953 and circ_0009024 were remarkably elevated in patients with active tuberculosis and might represent novel diagnostic biomarkers. In Zhu et al's study [21], circ_0027089 exhibited high abundance in plasma and could distinguish hepatitis B virus-related hepatocellular carcinoma from cirrhosis or healthy controls. In Liu et al.'s study [22], a two-circular RNA signature was identified as a promising noninvasive diagnostic biomarker for lung adenocarcinoma.

Recent studies showed that circRNAs participated in EOC tumorigenesis and progression by regulating various processes, including cell proliferation, apoptosis, invasion and migration. Ding et al. [23] reported that circ_0072995 could promote cell proliferation and migration via modulating miR-147a/CKD6 axis in EOC. Gan et al. [24] reported that circMUC16 could promote autophagy of EOC via interaction with miR-199a and ATG13. Additionally, circRNAs are reported to serve as potential biomarkers for EOC diagnosis and prognosis. In Liu et al.'s study [25], circHIPK3 was upregulated and predicted a poor prognosis in EOC. In Hu et al.'s study [26], plasma circBNC2 represented a promising diagnostic biomarker for EOC.

In our previous experiment, we found that circN-4BP2L2 was significantly downregulated in EOC by performing circRNA-sequencing analysis [27]. Besides, low level of circN4BP2L2 was predictive of disease progression [27]. Functional experiments revealed that downregulation of circN4BP2L2 could promote cell migration and invasion in EOC cell line [28]. Furthermore, plasma circN4BP2L2 could significantly distinguish EOC from benign ovarian cyst and healthy controls [28]. In this study, we aimed to further evaluate the diagnostic value of circN4BP2L2 by the unifying model of type I and type II EOC.

Methods

Study population

A total of 559 women were initially enrolled in this study. Nineteen malignancies were excluded because of non-EOC (n=10, 6 granulosa cell tumors, 2 endodermal sinus tumors, and 2 dysgerminomas) and metastasis (n=9). The eligible study population (n=540) consisted of age and menopause-matched women with EOC (n=180), benign ovarian cyst (n=180), and healthy volunteers (n=180). The study protocol was approved by the local ethics committee at the National Cancer Center/National Clinical Research Center for Cancer/Cancer Hospital and Peking Union Medical College Hospital of Chinese Academy of Medical Sciences (CAMS), and was conducted in accordance with the Declaration of Helsinki. Written informed consents for sample collection and article publication were obtained from all enrolled women.

After surgery, the tumors were independently examined by two experienced pathologists for the diagnosis of EOC, tumor histology, tumor grade and tumor stage (I–IV), according to the International Federation of Gynecology and Obstetrics (FIGO) standards. The exclusion criteria were: ovarian borderline tumors; patients with preoperative chemotherapy, radiotherapy or target

Ning et al. BMC Cancer (2022) 22:1210 Page 3 of 12

therapy; and patients with other coexisting malignancies. The EOCs were further divided into type I and type II tumors.

Sample collection

Patients were prospectively and consecutively enrolled when admitted for surgery for suspicious malignancies or benign ovarian cysts at the Department of Gynecologic Oncology in Cancer Hospital and Peking Union Medical College Hospital of CAMS, between December 2015 and April 2021. Blood samples were obtained on the surgery day. After centrifugation (3000 rotations per minute for 5 min), plasma samples were separated and stored at $-80\,^{\circ}\mathrm{C}$ until use.

RNA preparation and quality assessment

Total RNA was extracted from 200 μ L of plasma samples using TRIzol reagent (Takara Bio, Nojihigashi, Kusatsu, Japan) according to the manufacturer's instructions. RNA concentration was measured using NanoDrop 1000 spectrophotometers, and RNA was set at an OD A260/280 ratio between 1.8 and 2.1 and an OD A260/230 ratio > 1.8.

RT-qPCR

RT-qPCR was conducted using PrimeScript™ RT reagent Kit with gDNA Eraser (Takara Bio, Nojihigashi, Kusatsu, Japan) and SYBR® Premix Ex Taq™ II (Tli RNaseHPlus) (Takara Bio, Nojihigashi, Kusatsu, Japan) according to the manufacturer's instructions. GAPDH was used as an internal reference gene. The RT-qPCR protocol included a denaturation step (95 °C for 30 s) and 40 cycles of denaturation (95 °C for 5 s) and annealing (60 °C for 40 s). The relative expression level was calculated using the $2^{-\Delta\Delta CT}$ method. The primer sequences were as follows: circN4BP2L2 (forward, 5'-CATG-GTGTGTCTCGAAAGAAG-3' and reverse, 5'-CTG-TACCCATC TTGATGGTGA-3') and GAPDH (forward, 5'-AACGTGTCAGTGGTGGACCTG-3' and reverse, 5'-GAGACCACCTGGTGCTCAGTG-3').

ELISA

ELISA analyses were conducted on coded plasma samples to measure the CA125 levels (Quantikine Human CA125 Immunoassay; R&D Systems, Minneapolis, USA) and HE4 concentrations (Quantikine Human HE4 Immunoassay; R&D Systems, Minneapolis, USA) according to the manufacturer's instructions.

Statistical analysis

SPSS 24.0 (SPSS Inc., Chicago, IL, USA) were used to conduct the statistical analyses. The statistical differences between two groups were compared by the Mann-Whitney test. The statistical differences among three or more groups were compared using the one-way ANOVA test.

Youden index (specificity+sensitivity-1) was used to calculate the cut-off value of circN4BP2L2 (Supplementary Table S1). The cut-off value for CA125 < 35 U/mL and for HE4 < 55.86 pmol/L were adopted. Cases with marker levels above (CA125 and HE4) or below (circN4BP2L2) threshold levels were considered to be positive. The probabilities for each marker were predicted by constructing the receiver operating characteristic (ROC) curves. The area under the curve (AUC) value, sensitivity (sen) and specificity (spe) were calculated for individual markers. A value of p<0.05 was considered to be statistically significant.

Results

Patient material

Of 540 women eligible for analysis, 180 had EOC, 180 had benign ovarian cyst, and 180 were healthy controls. The average age was 56 (range, 30-76) years old. The malignancies were further divided into type I (n=70, 39%) and type II (n=110, 61%) EOC based on histology and grade. Type I included low-grade serous (n=16; 9%), low-grade endometrioid (n=15; 8%), all clear cell (n=23; 13%) and all mucinous (n=14; 8%). Type II included high-grade serous (n=88; 50%), high-grade endometrioid (n=13; 7%), all undifferentiated (n=7; 4%) and all malignant mixed mesodermal tumors (n=2; 1%). Most patients were diagnosed at advanced stages (FIGO III+IV: 116/180, 65%). The benign cohort included endometriosis (n=104, 58%), serous cystadenoma (n=24, 13%), mucinous cystadenoma (n=29, 16%), simple cystadenoma (n=10, 6%) and mature teratoma (n=13, 7%) (Table 1).

CircN4BP2L2 level and clinicopathologic features of type I and type II EOC

The relationship between circN4BP2L2 level and clinicopathologic features of EOC was assessed in 180 patients (Table 2). The results showed that decreased circN4BP2L2 was associated with advanced FIGO stage (p<0.01), worse tumor grade (p=0.03) and LNM (p<0.01) in EOC patients. CircN4BP2L2 level had no correlation to age or histology in EOC patients. A total of 70 patients with type I tumors were enrolled to evaluate the relationship between circN4BP2L2 and clinicopathological parameters. The data showed that low expression level of circN4BP2L2 was associated with advanced FIGO stage (p<0.01), but not correlated to age, histology, tumor grade or LNM in type I EOC. A total of 110 type II EOC patients were included to investigate the association between circN4BP2L2 level and clinicopathological features. The results revealed that decreased circN4BP2L2 was associated with LNM (p=0.04), but not related to age, histology or tumor stage in type II EOC.

Ning et al. BMC Cancer (2022) 22:1210 Page 4 of 12

Table 1 The main clinicopathologic parameters of eligible women (n = 540)

	n (%)
Age, average, range ^a	56 (30–76)
Epithelial ovarian cancer ^b	
EOC type I	70/180 (39%)
EOC type II	110/180 (61%)
Histological subtype	
Serous carcinoma	104/180
Low-grade	(59%)
High-grade	16/180 (9%) 88/180 (50%)
Endometrioid carcinoma	28/180 (15%)
Low-grade	15/180 (8%)
High-grade	13/180 (7%)
Clear cell carcinoma	23/180 (13%)
Mucinous carcinoma	14/180 (8%)
Undifferentiated carcinoma	7/180 (4%)
Malignant mixed mesodermal tumor	2/180 (1%)
FIGO stage	
I	47/180 (26%)
II	17/180 (9%)
III	97/180 (54%)
IV	19/180 (11%)
Tumor grade	
High	54/180 (30%)
Moderate	35/180 (19%)
Poor	84/180 (47%)
Undifferentiated	7/180 (4%)
Benign ovarian cyst ^c	
Endometriosis	104/180
	(58%)
Serous cystadenoma	24/180 (13%)
Mucinous cystadenoma	29/180 (16%)
Simple cystadenoma	10/180 (6%)
Mature teratoma	13/180 (7%)

Notes: ^a Average age of 540 age-matched eligible women. ^b The clinic opathologic parameters of patients with epithelial ovarian cancer (n=180). ^c Histological subtype of patients with benign ovarian cyst (n=180)

Abbreviations: n, number; EOC, epithelial ovarian cancer; FIGO, International Federation of Gynecology and Obstetrics

Levels of circN4BP2L2, CA125 and HE4 in type I and type II EOC

The relative expression level of circN4BP2L2 was significantly lower in both type I and type II EOC, when compared to those in benign and normal cohort (p<0.01). Conversely, the relative expression level of CA125 was significantly higher in both type I and type II EOC, when in comparison with those in benign and normal cohort (p<0.01). However, when comparing with the benign and normal cohorts, the relative expression level of HE4 was higher in type II EOC (p<0.01), but not in type I cohort (p>0.05). Besides, CA125 and HE4 levels were significantly different between type I and type II cohorts (p<0.01), but not circN4BP2L2 (p=0.91). CA125 (p<0.01), but not circN4BP2L2 (p=0.21) or HE4

(p=0.99), was significant when benign cohort was compared to normal cohort (Fig. 1).

The diagnostic value of circN4BP2L2 in type I EOC

Both circN4BP2L2 and HE4 could separate type I EOC from benign cohort, whereas CA125 could only distinguish late-stage type I EOC from benign cohort. Besides, circN4BP2L2, CA125 and HE4 could all separate type I EOC from normal cohort (Table 3). The median value in Type I EOC for HE4 was 61 pmol/L, and decreased to 46 pmol/L and 44 pmol/L in benign and normal cohorts, respectively. The decrease was even more notable for CA125 where median value of CA125 ranged from 85 U/mL in type I EOC to 48 U/mL in benign and 29 U/mL in normal cohort. The median value in type I EOC for circN4BP2L2 was 21, and increased to 82 in both benign and normal cohort.

In discrimination between type I EOC and benign cohorts, the ROC AUC was high for circN4BP2L2 (0.84) (Fig. 2a), followed by HE4 (0.70) (Fig. 2c) and CA125 (0.55) (Fig. 2b). The sensitivity and specificity were higher for circN4BP2L2 (69%, 87%) than those for CA125 (59%, 32%) and HE4 (54%, 76%). When comparing early-stage tumors with benign cohort, the ROC AUC was high for circN4BP2L2 (0.79 circN4BP2L2; 0.62 HE4; 0.40 CA125) (Fig. 3a, i and e), the sensitivity was also high for circN-4BP2L2 (90% circN4BP2L2; 48% CA125; 38% HE4), and the specificity was high for HE4 (76% HE4; 56% circ-N4BP2L2; 32% CA125). When comparing late-stage tumors with benign cohort, the ROC AUC was high for circN4BP2L2 (0.92 circN4BP2L2; 0.80 HE4; 0.75 CA125) (Fig. 3c and k g), the sensitivity was also high for circN-4BP2L2 (87% circN4BP2L2; 77% HE4; 73% CA125), and the specificity was also high for circN4BP2L2 (94% circ-N4BP2L2; 76% HE4; 32% CA125).

When distinguishing type I EOC from normal cohort, the ROC AUC was high for circN4BP2L2 (0.87) (Fig. 2d), followed by CA125 (0.73) (Fig. 2e) and HE4 (0.70) (Fig. 2f). The sensitivity and specificity were higher for circN4BP2L2 (69%, 93%) than those for HE4 (54%, 82%) and CA125 (59%, 58%). In discrimination between earlystage tumors and normal cohort, the ROC AUC was high for circN4BP2L2 (0.81 circN4BP2L2; 0.64 CA125; 0.63 HE4) (Fig. 3b, f and j), the sensitivity was also high for circN4BP2L2 (55% circN4BP2L2; 48% CA125; 38% HE4), and the specificity was also high for circN4BP2L2 (93% circN4BP2L2; 82% HE4; 58% CA125). In discrimination between late-stage tumors and normal cohort, the ROC AUC was high for circN4BP2L2 (0.95 circN4BP2L2; 0.85 CA125; 0.80 HE4) (Fig. 3d h, Fig. 3 L), the sensitivity was also high for circN4BP2L2 (87% circN4BP2L2; 77% HE4; 73% CA125), and the specificity was also high for circN-4BP2L2 (98% circN4BP2L2; 82% HE4; 58% CA125).

Ning et al. BMC Cancer (2022) 22:1210 Page 5 of 12

Table 2 Correlation between plasma circN4BP2L2 expression level and clinicopathologic parameters of EOC (n = 180)

Parameter	EOC (n = 18	EOC (n = 180)					Type II EOC (n = 110)		
	n (%)	CircN- 4BP2L2, Mean±SD	р	n (%)	CircN- 4BP2L2, Mean±SD	р	n (%)	CircN- 4BP2L2, Mean±SD	р
Age			0.52			0.70			0.47
≤ 50y	54 (30%)	27.48 ± 21.15		17 (24%)	30.24 ± 24.74		37 (34%)	26.21 ± 19.52	
> 50y	126 (70%)	26.70 ± 22.51		53 (76%)	31.04 ± 24.92		73 (66%)	23.54 ± 20.19	
Histology			0.75			0.11			0.76
Serous	106 (59%)	27.01 ± 22.88		18 (26%)	40.11 ± 30.81		88 (80%)	28.53 ± 22.19	
Others	74 (41%)	26.81 ± 20.98		52 (74%)	27.64 ± 21.64		22 (20%)	23.30 ± 19.22	
FIGO stage			< 0.01*			< 0.01*			0.44
1+11	64 (36%)	35.19 ± 25.19		40 (57%)	39.19 ± 26.29		24 (22%)	39.19 ± 26.29	
III + IV	116 (64%)	22.38 ± 18.73		30 (43%)	19.73 ± 17.27		86 (78%)	19.73 ± 17.27	
Grade			0.03*			0.15			NA
G1	54 (30%)	32.69 ± 25.61		54 (77%)	32.69 ± 25.61		0 (0%)	NA	
G2+G3	126 (70%)	24.47 ± 19.96		16 (23%)	24.64 ± 20.88		110 (100%)	24.44±19.92	
LNM ^a			< 0.01*			0.70			0.04*
Yes	59 (40%)	19.71 ± 15.11		8 (14%)	19.27 ± 10.81		51 (57%)	19.78 ± 15.76	
No	89 (60%)	30.46 ± 23.00		50 (86%)	31.61 ± 24.55		39 (43%)	29.00 ± 21.07	

Notes: ^a The information for lymph node metastasis was only available in 148 EOC patients

Abbreviations: n, number; FIGO, International Federation of Gynecology and Obstetrics; G, grade; LNM, lymph node metastasis; SD, standard deviation; NA, not available

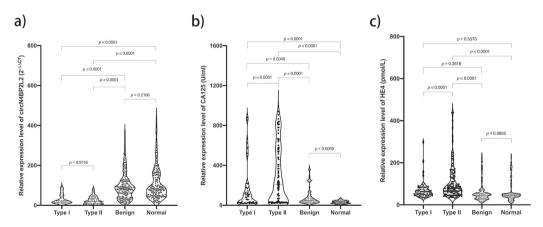


Fig. 1 Relative expression level of plasma circN4BP2L2 (a), CA125 (b) and HE4 (c) in type I EOC (n = 70), type II EOC (n = 110), benign ovarian cysts (n = 180) and normal controls (n = 180)

The diagnostic value of circN4BP2L2 in type II EOC

When separating type II EOC from benign or normal cohort, statistical significance was achieved between all groups, except for CA125 in distinguishing early-stage tumors from benign cohort (Table 4). The median value for circN4BP2L2 in type II EOC was 19, and increased to 82 in both benign and normal cohort. The median value for HE4 in type II EOC was 79 pmol/L, and decreased to 46 pmol/L and 44 pmol/L in benign and normal cohorts, respectively. The decrease was even more notable for CA125 where median value of CA125 ranged from 233 U/ mL in type II EOC cohort to 48 U/ mL in benign cohort and 29 U/ mL in normal cohort.

In discrimination between type II EOC and benign cohorts, the ROC AUC was high for circN4BP2L2 (0.88) (Fig. 2 h), followed by HE4 (0.79) (Fig. 2j) and CA125 (0.72) (Fig. 2i). The sensitivity and specificity were higher for circN4BP2L2 (93%, 71%) than those for HE4 (75%, 76%) and CA125 (72%, 32%). When comparing early-stage tumors with benign cohort, the ROC AUC was high for circN4BP2L2 (0.85 circN4BP2L2; 0.77 HE4; 0.46 CA125) (Fig. 4a, i and e), the sensitivity was also high for circN4BP2L2 (88% circN4BP2L2; 71% HE4; 46% CA125), and the specificity was high for HE4 (76% HE4; 67% circN4BP2L2; 32% CA125). When comparing late-stage tumors with benign cohort, the ROC AUC was high for circN4BP2L2 (0.88 circN4BP2L2; 0.79 CA125; 0.79 HE4)

Ning et al. BMC Cancer (2022) 22:1210 Page 6 of 12

Table 3 CircN4BP2L2, CA125 and HE4 expression levels according to histology, type and stage; ROC AUC, sensitivity, specificity, and significant difference in type I EOC vs. benign and normal cohorts

Type I EOC cohort		Benign c	ohort		Normal cohort						
	Median (range)	Median (range)	ROC AUC (95% CI)	Sen	Spe	P -value	Median (range)	ROC AUC (95% CI)	Sen	Spe	P - value
CircN4BP2L2	21 (7–95)	82 (1–367)	0.84 (0.79–0.89)	69%	87%	< 0.01*	82 (11–439)	0.87 (0.82–0.92)	69%	93%	< 0.01*
Early-stage ^a	25 (7-91)		0.79 (0.72-0.85)	90%	56%	< 0.01*		0.81 (0.75-0.88)	55%	93%	< 0.01*
Late-stage ^a	15 (11–95)		0.92 (0.86-0.97)	87%	94%	< 0.01*		0.95 (0.90-0.99)	87%	98%	< 0.01*
CA125	85 (16–5420)	48 (13–373)	0.55 (0.46–0.65)	59%	32%	0.20	29 (4–69)	0.73 (0.65–0.81)	59%	58%	< 0.01*
Early-stage ^a	33 (16–181)		0.40 (0.30-0.51)	48%	32%	0.06		0.64 (0.53-0.75)	48%	58%	< 0.01*
Late-stage ^a	506 (18–5420)		0.75 (0.62–0.88)	73%	32%	< 0.01*		0.85 (0.75–0.94)	73%	58%	< 0.01*
HE4	61 (28–298)	46 (12–237)	0.70 (0.63–0.76)	54%	76%	< 0.01*	44 (7–235)	0.70 (0.63–0.77)	54%	82%	< 0.01*
Early-stage ^a	50 (28-207)		0.62 (0.53-0.70)	38%	76%	0.02*		0.63 (0.54-0.71)	38%	82%	0.01*
Late-stage ^a	69 (43-298)		0.80 (0.73-0.87)	77%	76%	< 0.01*		0.80 (0.74-0.87)	77%	82%	< 0.01*

^a According to International Federation of Gynecology and Obstetrics staging

Abbreviations: EOC, epithelial ovarian cancer; ROC, receiver operating characteristic curve; AUC, area under curve; 95% CI, 95% confidence interval; Sen, sensitivity; Spe, specificity

(Fig. 4c g, Fig. 4k), the sensitivity was also high for circN-4BP2L2 (94% circN4BP2L2; 79% CA125; 77% HE4), and the specificity was high for HE4 (76% HE4; 69% circN-4BP2L2; 32% CA125).

When distinguishing type II EOC from normal cohort, the ROC AUC was high for circN4BP2L2 (0.91) (Fig. 2k), followed by CA125 (0.84) (Fig. 2 L) and HE4 (0.78) (Fig. 2 m). The sensitivity and specificity were higher for circN4BP2L2 (80%, 87%) than those for HE4 (75%, 82%) and CA125 (72%, 58%). In discrimination between earlystage tumors and normal cohort, the ROC AUC was high for circN4BP2L2 (0.89 circN4BP2L2; 0.75 HE4; 0.68 CA125) (Fig. 4b, j and f), the sensitivity was also high for circN4BP2L2 (83% circN4BP2L2; 71% HE4; 46% CA125), and the specificity was high for HE4 (82% HE4; 77% circ-N4BP2L2; 58% CA125). In discrimination between latestage tumors and normal cohort, the ROC AUC was high for circN4BP2L2 (0.92 circN4BP2L2; 0.89 CA125; 0.79 HE4) (Fig. 4d h, Fig. 4 L), the sensitivity was also high for circN4BP2L2 (83% circN4BP2L2; 79% CA125; 77% HE4), and the specificity was also high for circN4BP2L2 (87% circN4BP2L2; 82% HE4; 58% CA125).

Discussion

Traditionally, EOC is classified based on histologic features [6]. Recently, it has been proposed that EOC should be classified into slow-growing type I and more aggressive type II tumors according to both histologic features and molecular genetics [8]. Previous studies reported that CA125 and HE4 were highly representative diagnostic biomarkers for type II EOC; however, both markers showed limited diagnostic ability in all type I tumors and early-stage type II malignancies [10, 29]. In this study, we aimed to assess the diagnostic value of circN4BP2L2 in

type I and type II EOC. The blood samples of a cohort of 540 women that were prospectively and consecutively collected were used for analysis. Our data suggested that circN4BP2L2 had a good diagnostic ability in both type I and type II EOC. In comparison to the benign or normal cohort, circN4BP2L2 resulted in good diagnostic power with ROC AUC 0.84 and 0.87 in type I, and 0.88 and 0.91 in type II tumors, and impressively high AUC 0.95 for late-stage type I EOC in comparison to the normal cohort. Notably, circN4BP2L2 also had good diagnostic performance in early-stage tumors in both type I and type II. To our knowledge, circN4BP2L2 has not earlier been evaluated by dividing EOC into type I and type II tumors.

Consistent with our results, the diagnostic value of circRNAs has been previously proved in various malignancies. Plasma circFARSA is significantly upregulated in non-small cell lung cancer and might serve as a promising noninvasive diagnostic biomarker for this malignancy [30]. Circ_0001785 is significantly dysregulated in breast cancer and was closely associated with tumor stage and distant metastasis, plasma circ_0001785 had better diagnostic performance than CA15-3 and CEA in breast cancer patients [31]. Circ_0000190, which was previously discovered to be down-regulated in gastric cancer tissues by microarray screening, had better AUC, sensitivity and specificity than commonly used diagnostic biomarkers such as CA19-9 and CEA [32]. The expression level of circ_0003998 was significantly increased in patients with hepatocellular carcinoma, and plasma circ_0003998 had a good diagnostic accuracy in distinguishing cancer patients from hepatitis B patients (AUC=0.83) and healthy controls (AUC=0.89) [33]. Similar results have

Ning et al. BMC Cancer (2022) 22:1210 Page 7 of 12

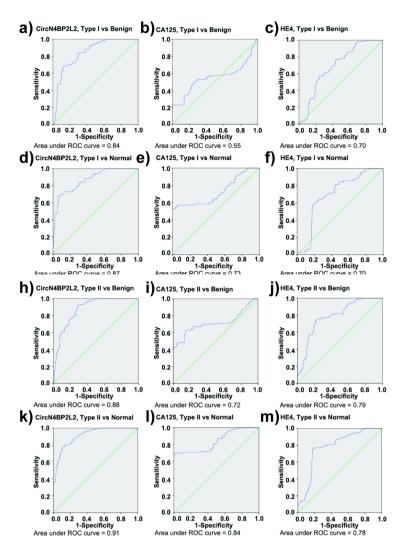


Fig. 2 ROC AUC for circN4BP2L2, CA125 and HE4 in type I and type II EOC. It contains ROC AUC for circN4BP2L2 comparing type I EOC with benign (a) and normal (d) cohorts; ROC AUC for CA125 comparing type I EOC with benign (b) and normal (e) cohorts; ROC AUC for HE4 comparing type I EOC with benign (c) and normal (f) cohorts; ROC AUC for circN4BP2L2 comparing type II EOC with benign (h) and normal (k) cohorts; ROC AUC for CA125 comparing type II EOC with benign (i) and normal (l) cohorts; ROC AUC for HE4 comparing type II EOC with benign (j) and normal (m) cohorts

also been achieved in colorectal cancer [34], esophageal squamous cell carcinoma [35] and prostate cancer [36].

The ultimate goal to improve EOC prognosis is to detect early-stage cases regardless of tumor subtype. In our dataset, early stage EOC comprised 35% of the malignant cohort, which is in accordance with previous studies [37, 38]. While CA125 and HE4 showed limited diagnostic value for recognizing early-stage tumors in both type I and type II EOC, circN4BP2L2 had a good diagnostic ability for early-stage cases. In comparison to the benign or normal cohort, circN4BP2L2 exhibited good diagnostic performance with ROC AUC 0.79 and 0.81 in early-stage type I, and 0.85 and 0.89 in early-stage type II EOC. Conventional techniques for detecting ovarian tumors include gynecological bimanual palpation, transvaginal ultrasound and computed tomography scanning [39]. It has been proposed that type I tumors, which are more

often localized in the pelvis and more generally of a larger size, are more easily to be detected at an earlier stage with conventional techniques than type II tumors, which are more often of invisible early lesions [40, 41]. Notably, our data revealed that circN4BP2L2 had even better diagnostic performance in distinguishing early-stage type II EOC than that in distinguishing early-stage type I EOC. These results suggested that circN4BP2L2 might serve as an adjunct to conventional techniques for detecting early stage EOC cases. Further large-scale clinical trials are warranted to evaluate the practicability of circN4BP2L2 for clinical application.

By assessing the association between the expression level of circN4BP2L2 and clinicopathological features of EOC, we found that low expression of circN4BP2L2 was correlated to advanced stage, poor differentiation and LNM. In stratified analysis, the data revealed that

Ning et al. BMC Cancer (2022) 22:1210 Page 8 of 12

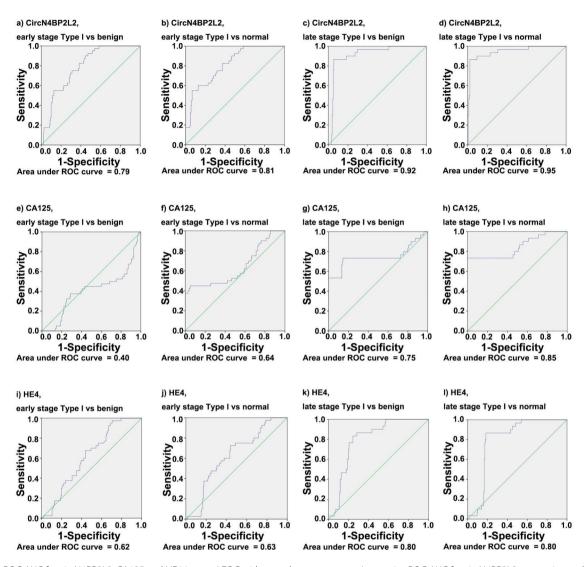


Fig. 3 ROC AUC for circN4BP2L2, CA125 and HE4 in type I EOC with regard to tumor stage. It contains ROC AUC for circN4BP2L2 comparing early-stage type I EOC with benign (a) and normal (b) cohorts; ROC AUC for circN4BP2L2 comparing late-stage type I EOC with benign (c) and normal (d) cohorts; ROC AUC for CA125 comparing early-stage type I EOC with benign (e) and normal (f) cohorts; ROC AUC for CA125 comparing late-stage type I EOC with benign (g) and normal (h) cohorts; ROC AUC for HE4 comparing early-stage type I EOC with benign (i) and normal (j) cohorts; and ROC AUC for HE4 comparing late-stage type I EOC with benign (k) and normal (l) cohorts

decreased circN4BP2L2 was respectively associated with advanced stage in Type I tumor and LNM in type II tumor. Additionally, our in vitro experiments revealed that decreased circN4BP2L2 might improve cell metastasis and invasion in EOC. Both clinical analyses and fundamental researches suggested that circN4BP2L2 participated in EOC development via regulating tumor cell metastasis. These results are in accordance with other published articles, where they concluded that circRNAs might promote EOC progression by regulating cell proliferation, apoptosis or migration [23–26]. With regard to the function mechanism of circRNAs in ovarian cancer, the most studied one was for circRNAs acting as microRNA sponges. It has been reported that circU-BAP2 [42], circPLEKHM3 [43], and circMYLK [44] could

respectively promote ovarian cancer development via regulating miR-382-5p, miR-9 and miR-652. Besides, circRNAs could participate in ovarian cancer development by targeting and regulating RNA-binding proteins. Liu et al. [45] reported that circ_0005276 was significantly upregulated in EOC and could aggravate the development of EOC by binding ADAM9. Chen et al. [46] discovered that circNOLC1 could promote EOC tumorigenesis and progression by binding ESRP1 and modulating the expression of CDK1 and RhoA. Furthermore, Yan et al. [47] reported that circITCH could inhibit the proliferation of ovarian cancer via downregulating lncRNA HULC. Despite the abovementioned mechanisms, the exact function mechanism of circN4BP2L2 is still

Ning et al. BMC Cancer (2022) 22:1210 Page 9 of 12

Table 4 CircN4BP2L2, CA125 and HE4 expression levels according to histology, type and stage; ROC AUC, sensitivity, specificity, and significant difference in type II EOC vs. benign and normal cohorts

Type II EOC cohort	Benign cohort						Normal cohort				
	Median (range)	Median (range)	ROC AUC (95% CI)	Sen	Spe	P -value	Median (range)	ROC AUC (95% CI)	Sen	Spe	P-value
CircN4BP2L2	19 (1–84)	82 (1–367)	0.88 (0.84–0.91)	93%	71%	< 0.01*	82 (11–439)	0.91 (0.88–0.94)	80%	87%	< 0.01*
Early-stage ^a	24 (1-74)		0.85 (0.79-0.92)	88%	67%	< 0.01*		0.89 (0.83-0.95)	83%	77%	< 0.01*
Late-stage ^a	19 (2-84)		0.88 (0.84-0.92)	94%	69%	< 0.01*		0.92 (0.89-0.95)	83%	87%	< 0.01*
CA125	233 (18–5801)	48 (13–373)	0.72 (0.65–0.79)	72%	32%	< 0.01*	29 (4–69)	0.84 (0.79–0.89)	72%	58%	< 0.01*
Early-stage ^a	30 (18–560)		0.46 (0.31-0.60)	46%	32%	0.49		0.68 (0.55-0.80)	46%	58%	< 0.01*
Late-stage ^a	398 (22–5801)		0.79 (0.72–0.86)	79%	32%	< 0.01*		0.89 (0.84–0.94)	79%	58%	< 0.01*
HE4	79 (32–438)	46 (12–237)	0.79 (0.73–0.84)	75%	76%	< 0.01*	44 (7–235)	0.78 (073–0.84)	75%	82%	< 0.01*
Early-stage ^a	88 (34-231)		0.77 (0.68-0.87)	71%	76%	< 0.01*		0.75 (0.65-0.85)	71%	82%	< 0.01*
Late-stage ^a	78 (32-438)		0.79 (0.74-0.85)	77%	76%	< 0.01*		0.79 (0.73-0.85)	77%	82%	< 0.01*

^a According to International Federation of Gynecology and Obstetrics staging

Abbreviations: EOC, epithelial ovarian cancer; ROC, receiver operating characteristic curve; AUC, area under curve; 95% CI, 95% confidence interval; Sen, sensitivity; Spe, specificity

unknown. Further laboratory experiments are needed to clarify how circN4BP2L2 functions in EOC progression.

Despite its rarity, type I EOC is a group of individually different tumors of clinical importance. Advanced mucinous and clear cell carcinomas are quite aggressive and have even higher mortality than type II EOC. Neither CA125 nor HE4 could effectively detect these type I tumors. Accordingly, our data revealed that the expression level of CA125 and HE4 in type II tumors were significantly higher than those in type I tumors, and both markers failed to separate type I tumors from benign or normal cohort. Notably, we found that the expression level of circN4BP2L2 in type I EOC was similar to that in type II EOC. When compared to benign or normal cohort, circN4BP2L2 was significantly decreased in both type I and type II tumors. The diagnostic performance of circN4BP2L2 was equally well in type I and type II. These results suggested that circN4BP2L2 might serve as an early biomarker for diagnosing all histology subtypes, including mucinous and clear cell tumors. However, due to limited number of type I cases, we were unable to conduct stratified analysis based on different histology. Further larger-scale clinical trials are needed to investigate the expression level of circN4BP2L2 in different histological subtype.

CA125 and HE4 are two FDA-approved EOC biomarkers. Instead of being used for disease detection, they are more usually applied for efficacy evaluating and recurrence monitoring. Except for relatively low sensitivity in early-stage cases, CA125 also had poor specificity in diagnosing EOC. It has been reported that, apart from EOC, elevated CA125 could also be tested in non-EOC gynecological malignancies such as endometrial cancer and cervical cancer, non-gynecological malignancies such as

lung cancer and breast cancer, benign gynecological diseases such as endometriosis, cardiopulmonary diseases such as myocardial infarction and chronic obstructive pulmonary disease, pregnant women, and even healthy women [48, 49]. Particularly, CA125 could elevate in approximately half of the patients with endometriosis [50]. Therefore, the value of CA125 in distinguishing EOC from benign ovarian cysts is often limited. In our series, more than half of the patients (104/180, 58%) in benign cohort had endometriosis, which might partly explain the poor diagnostic value of CA125 in separating EOC from benign cohort. While HE4 has relatively high specificity in diagnosing EOC, its sensitivity is also unsatisfactory [16]. Besides, controversies arise regarding whether the performance of HE4 could be affected by menopausal status. Some researchers reported that HE4 had better diagnostic value in post-menopausal ovarian cancer patients [51, 52], while others believed that HE4 performed equally well in both pre- and post-menopausal ovarian cancer patients [53, 54]. By and large, both markers showed limited value in diagnosing EOC.

This study has two main limitations. On one hand, the sample size was relatively small and we were unable to conduct stratified analysis based on tumor histology. On the other hand, the evaluation of circN4BP2L2 as a true diagnostic biomarker was limited in our cohort of patients with suspicious malignancies or benign ovarian cysts, formal study using a screening cohort is needed in the future.

Conclusion

Our results demonstrated that circN4BP2L2 might serve as a promising diagnostic biomarker for both type I and type II EOC. The diagnostic safety for circN4BP2L2 in Ning et al. BMC Cancer (2022) 22:1210 Page 10 of 12

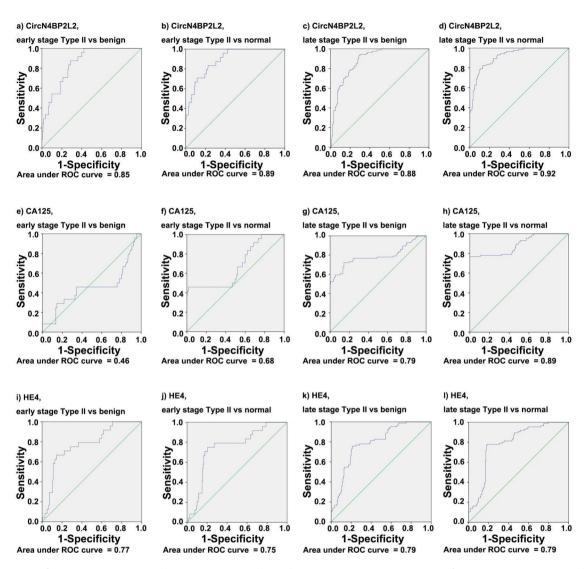


Fig. 4 ROC AUC for circN4BP2L2, CA125 and HE4 in type II EOC with regard to tumor stage. It contains ROC AUC for circN4BP2L2 comparing early-stage type II EOC with benign (a) and normal (b) cohorts; ROC AUC for circN4BP2L2 comparing late-stage type II EOC with benign (c) and normal (d) cohorts; ROC AUC for CA125 comparing early-stage type II EOC with benign (e) and normal (f) cohorts; ROC AUC for CA125 comparing late-stage type II EOC with benign (q) and normal (h) cohorts; ROC AUC for HE4 comparing early-stage type II EOC with benign (i) and normal (j) cohorts; and ROC AUC for HE4 comparing late-stage type II EOC with benign (k) and normal (l) cohorts

early-stage type I and type II EOC is also acceptable. Further large-scale well-designed studies are warranted to investigate whether circN4BP2L2 is specific for all histologic subgroups.

Abbreviations

Epithelial ovarian cancer EOC

RT-aPCR Reverse transcription-quantitative polymerase chain reaction

CA125 Cancer antigen 125 HE4 Human epididymis protein 4

ELISA Enzyme-linked immunosorbent assay

Receiver operating curve ROC AUC The area under the curve LNM Lymph node metastasis

OC ovarian cancer

grade G

FDA the US Food and Drug Administration

CircRNAs Circular RNAs CAMS Chinese Academy of Medical Sciences

FIGO The International Federation of Gynecology and Obstetrics

Sensitivity sen Specificity spe

CA15-3 Cancer antigen 15 – 3 CFA Carcinoembryonic antigen CA19-9 Cancer antigen 19-9

Supplementary Information

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Supplementary Material 1

Acknowledgements

Not applicable.

Ning et al. BMC Cancer (2022) 22:1210 Page 11 of 12

Authors' contributions

L Ning contributes to conception, design, analysis and interpretation of data, drafting and revising the manuscript. J Lang, Bo Long and L Wu contribute to revising the manuscript critically for important contents, and final approval of the manuscript.

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Not applicable.

Data availability

The datasets used and/or analyzed during the current study are available from the corresponding author on reasonable request.

Declarations

Ethics approval and consent to participate

The study protocol was approved by the local ethics committee at the National Cancer Center/National Clinical Research Center for Cancer/Cancer Hospital of CAMS and Peking Union Medical College Hospital of CAMS and was conducted in accordance with the Declaration of Helsinki. Written informed consent for taking the venous blood and tissues was obtained from all patients and healthy volunteers.

Consent for publication

Not applicable.

Competing interests

The authors declare that they have no competing interests.

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References

- Siegel RL, Miller KD, Fuchs HE, Jemal A. Cancer Statistics. 2021. CA: a cancer journal for clinicians. 2021;71(1):7–33.
- Bray F, Ferlay J, Soerjomataram I, Siegel RL, Torre LA, Jemal A. Global cancer statistics 2018: GLOBOCAN estimates of incidence and mortality worldwide for 36 cancers in 185 countries. CA: a cancer journal for clinicians. 2018;68(6):394–424.
- Wu L, Zhu J, Yin R, Wu X, Lou G, Wang J, et al. Olaparib maintenance therapy in patients with newly diagnosed advanced ovarian cancer and a BRCA1 and/or BRCA2 mutation: SOLO1 China cohort. Gynecol Oncol. 2021;160(1):175–81.
- Lheureux S, Gourley C, Vergote I, Oza AM. Epithelial ovarian cancer. Lancet (London England). 2019;393(10177):1240–53.
- Kim SI, Lee M, Kim HS, Chung HH, Kim JW, Park NH, et al. Effect of BRCA mutational status on survival outcome in advanced-stage high-grade serous ovarian cancer. J Ovarian Res. 2019;12.
- Prat J. New insights into ovarian cancer pathology. Annals of oncology: official journal of the European Society for Medical Oncology. 2012;23(Suppl 10):x111–117.
- Kindelberger DW, Lee Y, Miron A, Hirsch MS, Feltmate C, Medeiros F, et al. Intraepithelial carcinoma of the fimbria and pelvic serous carcinoma: Evidence for a causal relationship. Am J Surg Pathol. 2007;31(2):161–9.
- Shih le M, Kurman RJ. Ovarian tumorigenesis: a proposed model based on morphological and molecular genetic analysis. Am J Pathol. 2004;164(5):1511–8.
- Kurman RJ, Shih le M. Molecular pathogenesis and extraovarian origin of epithelial ovarian cancer-shifting the paradigm. Hum Pathol. 2011;42(7):918–31.
- Kristjansdottir B, Levan K, Partheen K, Sundfeldt K. Diagnostic performance of the biomarkers HE4 and CA125 in type I and type II epithelial ovarian cancer. Gynecol Oncol. 2013;131(1):52–8.
- 11. Watanabe T, Nanamiya H, Endo Y, Kojima M, Nomura S, Furukawa S, et al. Identification and clinical significance of somatic oncogenic mutations in epithelial ovarian cancer. J Ovarian Res. 2021;14(1):129.
- 12. Saleh A, Perets R. Mutated p53 in HGSC-From a Common Mutation to a Target for Therapy. Cancers (Basel). 2021;13(14).

- Zhang M, Cheng S, Jin Y, Zhao Y, Wang Y. Roles of CA125 in diagnosis, prediction, and oncogenesis of ovarian cancer. Biochim et Biophys acta Reviews cancer. 2021;1875(2):188503.
- Dochez V, Caillon H, Vaucel E, Dimet J, Winer N, Ducarme G. Biomarkers and algorithms for diagnosis of ovarian cancer: CA125, HE4, RMI and ROMA, a review. J Ovarian Res. 2019;12(1):28.
- Scaletta G, Plotti F, Luvero D, Capriglione S, Montera R, Miranda A, et al. The role of novel biomarker HE4 in the diagnosis, prognosis and followup of ovarian cancer: a systematic review. Expert Rev anticancer therapy. 2017:17(9):827–39.
- Granato T, Porpora MG, Longo F, Angeloni A, Manganaro L, Anastasi E. HE4 in the differential diagnosis of ovarian masses. Clin Chim acta; Int J Clin Chem. 2015;446:147–55.
- 17. Ebbesen KK, Hansen TB, Kjems J. Insights into circular RNA biology. RNA Biol. 2017;14(8):1035–45.
- Qu S, Zhong Y, Shang R, Zhang X, Song W, Kjems J, et al. The emerging landscape of circular RNA in life processes. RNA Biol. 2017;14(8):992–9.
- Ouyang Q, Huang Q, Jiang Z, Zhao J, Shi GP, Yang M. Using plasma circRNA_002453 as a novel biomarker in the diagnosis of lupus nephritis. Mol Immunol. 2018;101:531–8.
- Huang Z, Su R, Qing C, Peng Y, Luo Q, Li J. Plasma Circular RNAs hsa_ circ_0001953 and hsa_circ_0009024 as Diagnostic Biomarkers for Active Tuberculosis. Front Microbiol. 2018;9:2010.
- 21. Zhu K, Zhan H, Peng Y, Yang L, Gao Q, Jia H, et al. Plasma hsa_circ_0027089 is a diagnostic biomarker for hepatitis B virus-related hepatocellular carcinoma. Carcinogenesis. 2020;41(3):296–302.
- Liu XX, Yang YE, Liu X, Zhang MY, Li R, Yin YH, et al. A two-circular RNA signature as a noninvasive diagnostic biomarker for lung adenocarcinoma. J Transl Med. 2019;17(1):50.
- Ding J, Wang Q, Guo N, Wang H, Chen H, Ni G, et al. CircRNA circ_0072995 promotes the progression of epithelial ovarian cancer by modulating miR-147a/CDK6 axis. Aging (Albany NY). 2020;12(17):17209–23.
- 24. Gan X, Zhu H, Jiang X, Obiegbusi SC, Yong M, Long X, et al. CircMUC16 promotes autophagy of epithelial ovarian cancer via interaction with ATG13 and miR-199a. Mol cancer. 2020;19(1):45.
- Liu N, Zhang J, Zhang LY, Wang L. CircHIPK3 is upregulated and predicts a poor prognosis in epithelial ovarian cancer. Eur Rev Med Pharmacol Sci. 2018;22(12):3713–8.
- Hu Y, Zhu Y, Zhang W, Lang J, Ning L. Utility Of Plasma circBNC2 As A Diagnostic Biomarker In Epithelial Ovarian Cancer. OncoTargets and therapy. 2019:12:9715–23.
- Ning L, Long B, Zhang W, Yu M, Wang S, Cao D, et al. Circular RNA profiling reveals circEXOC6B and circN4BP2L2 as novel prognostic biomarkers in epithelial ovarian cancer. Int J Oncol. 2018;53(6):2637–46.
- 28. Ning L, Lang J, Wu L. Plasma circN4BP2L2 is a promising novel diagnostic biomarker for epithelial ovarian cancer. BMC Cancer. 2022;22(1):6.
- Lu D, Kuhn E, Bristow RE, Giuntoli RL 2nd, Kjær SK, Shih le M, et al. Comparison of candidate serologic markers for type I and type II ovarian cancer. Gynecol Oncol. 2011;122(3):560–6.
- Hang D, Zhou J, Qin N, Zhou W, Ma H, Jin G, et al. A novel plasma circular RNA circFARSA is a potential biomarker for non-small cell lung cancer. Cancer Med. 2018;7(6):2783–91.
- Yin WB, Yan MG, Fang X, Guo JJ, Xiong W, Zhang RP. Circulating circular RNA hsa_circ_0001785 acts as a diagnostic biomarker for breast cancer detection. Clin Chim acta; Int J Clin Chem. 2018;487:363–8.
- Chen S, Li T, Zhao Q, Xiao B, Guo J. Using circular RNA hsa_circ_0000190 as a new biomarker in the diagnosis of gastric cancer. Clin Chim acta; Int J Clin Chem. 2017;466:167–71.
- Qiao GL, Chen L, Jiang WH, Yang C, Yang CM, Song LN, et al. Hsa_ circ_0003998 may be used as a new biomarker for the diagnosis and prognosis of hepatocellular carcinoma. OncoTargets and therapy. 2019;12:5849–60.
- Ye DX, Wang SS, Huang Y, Chi P. A 3-circular RNA signature as a noninvasive biomarker for diagnosis of colorectal cancer. Cancer Cell Int. 2019;19:276.
- Zhang Y, Wang Q, Zhu D, Rong J, Shi W, Cao X. Up-regulation of circ-SMAD7 inhibits tumor proliferation and migration in esophageal squamous cell carcinoma. Biomed pharmacotherapy = Biomedecine pharmacotherapie. 2019:111:596–601.
- Kong Z, Wan X, Lu Y, Zhang Y, Huang Y, Xu Y, et al. Circular RNA circFOXO3 promotes prostate cancer progression through sponging miR-29a-3p. J Cell Mol Med. 2020;24(1):799–813.

Ning et al. BMC Cancer (2022) 22:1210 Page 12 of 12

- Jessmon P, Boulanger T, Zhou W, Patwardhan P. Epidemiology and treatment patterns of epithelial ovarian cancer. Expert Rev anticancer therapy. 2017;17(5):427–37.
- Fagotti A, Ferrandina MG, Vizzielli G, Pasciuto T, Fanfani F, Gallotta V, et al. Randomized trial of primary debulking surgery versus neoadjuvant chemotherapy for advanced epithelial ovarian cancer (SCORPION-NCT01461850). Int J Gynecol Cancer. 2020;30(11):1657–64.
- 39. Guo B, Lian W, Liu S, Cao Y, Liu J. Comparison of diagnostic values between CA125 combined with CA199 and ultrasound combined with CT in ovarian cancer. Oncol Lett. 2019;17(6):5523–8.
- Alcázar JL, Utrilla-Layna J, Mínguez J, Jurado M. Clinical and ultrasound features of type I and type II epithelial ovarian cancer. Int J Gynecol Cancer. 2013;23(4):680–4.
- 41. Yemelyanova AV, Cosin JA, Bidus MA, Boice CR, Seidman JD. Pathology of stage I versus stage III ovarian carcinoma with implications for pathogenesis and screening. Int J Gynecol Cancer. 2008;18(3):465–9.
- 42. Xu Q, Deng B, Li M, Chen Y, Zhuan L. circRNA-UBAP2 promotes the proliferation and inhibits apoptosis of ovarian cancer though miR-382-5p/PRPF8 axis. J Ovarian Res. 2020;13(1):81.
- Zhang L, Zhou Q, Qiu Q, Hou L, Wu M, Li J, et al. CircPLEKHM3 acts as a tumor suppressor through regulation of the miR-9/BRCA1/DNAJB6/KLF4/AKT1 axis in ovarian cancer. Mol cancer. 2019;18(1):144.
- Zhao Y, Hu Y, Shen Q, Chen Q, Zhu XJ, Jiang SS, et al. CircRNA_MYLK promotes malignant progression of ovarian cancer through regulating microRNA-652. Eur Rev Med Pharmacol Sci. 2020;24(10):5281–91.
- Liu ZH, Liu WJ, Yu XY, Qi XL, Sun CC. Circ_0005276 aggravates the development of epithelial ovarian cancer by targeting ADAM9. Eur Rev Med Pharmacol Sci. 2020;24(20):10375–82.
- Chen S, Wu W, Li QH, Xie BM, Shen F, Du YP, et al. Circ-NOLC1 promotes epithelial ovarian cancer tumorigenesis and progression by binding ESRP1 and modulating CDK1 and RhoA expression. Cell death discovery. 2021;7(1):22.

- 47. Yan H, Xiang H, Sun B, Feng F, Chen P. Circular RNA-ITCH Inhibits the Proliferation of Ovarian Carcinoma by Downregulating IncRNA HULC. Reproductive sciences (Thousand Oaks. Calif). 2020;27(1):375–9.
- Ghaemmaghami F, Karimi Zarchi M, Hamedi B. High levels of CA125 (over 1,000 IU/ml) in patients with gynecologic disease and no malignant conditions: three cases and literature review. Archives of gynecology and obstetrics. 2007;276(5):559–61.
- Wu HB, Shao K, Wang YC, Wang XC, Liu HL, Xie YT, et al. Research progress of CA125 and BDNF in serum of patients with acute myocardial infarction for predicting acute heart failure. Clin Hemorheol Microcirc. 2020;75(1):99–106.
- Hirsch M, Duffy J, Davis CJ, Nieves Plana M, Khan KS. Diagnostic accuracy of cancer antigen 125 for endometriosis: a systematic review and meta-analysis. Bjog. 2016;123(11):1761–8.
- 51. Kim B, Park Y, Kim B, Ahn HJ, Lee KA, Chung JE, et al. Diagnostic performance of CA 125, HE4, and risk of Ovarian Malignancy Algorithm for ovarian cancer. J Clin Lab Anal. 2019;33(1):e22624.
- 52. Zhang L, Chen Y, Wang K. Comparison of CA125, HE4, and ROMA index for ovarian cancer diagnosis. Curr Probl Cancer. 2019;43(2):135–44.
- Wei SU, Li H, Zhang B. The diagnostic value of serum HE4 and CA-125 and ROMA index in ovarian cancer. Biomedical Rep. 2016;5(1):41–4.
- Terlikowska KM, Dobrzycka B, Witkowska AM, Mackowiak-Matejczyk B, Sledziewski TK, Kinalski M, et al. Preoperative HE4, CA125 and ROMA in the differential diagnosis of benign and malignant adnexal masses. J Ovarian Res. 2016;9(1):43.

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