Prognostic value of serum CEA and CA19-9 levels in pancreatic ductal adenocarcinoma

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Abstract. The present study investigated the associations of serum carcinoembryonic antigen (CEA) and carbohydrate antigen 19-9 (CA19-9) levels with clinicopathological variables and survival outcomes in Libyan patients with pancreatic ductal adenocarcinoma (PDAC). The clinicopathological variables of 123 patients with PDAC registered at the National Cancer Institute in Misurata, Libya, between 2010 and 2018 were retrospectively analyzed. Blood samples from these patients were analyzed for serum CEA and CA19-9 levels before treatment by electrochemiluminescence immunoassay (double antibody sandwich ELISA) on a Roche cobas e 602 modules. The relationships between CA19-9 and CEA serum levels with clinicopathologic variables and survival outcomes were analyzed using the Kaplan-Meier method, log-rank test and Cox regression analyzes. Cut-off values for serum CEA and CA19-9 levels were 5 ng/ml and 400 U/ml, respectively. The median serum levels of all patients with PDAC for CEA and CA19-9 were 8 ng/ml (1.1-377 ng/ml) and 389 U/ml (1-10,050 U/ml), respectively. Tumors with higher serum CEA and CA19-9 levels were found in 63 and 48% of patients, respectively. Higher CEA and CA19-9 serum levels were significantly associated with more indicators of a malignant phenotype, including a surgically unresectable tumor, unevaluable lymph nodes, advanced stages and distant metastases. Regarding survival, patients with higher serum levels of the biomarkers CEA and CA19-9 had shorter overall survival rates (P<0.016 and (P<0.014, log-rank, respectively) and lower disease-free survival rates (P<0.002 and P<0.0001, log-rank, respectively). The present study demonstrated significant clinical and prognostic value of serum levels of biomarkers CEA and CA19-9 for Libyan patients with PDAC. Moreover, patients with PDAC with higher serum CEA and CA19-9 levels had more aggressive tumors, higher rates of disease recurrence and shorter overall survival rates and thus required more vigilant follow-up. Further multinational studies with larger PDAC cohorts are warranted to confirm these findings in terms of improved clinical decision making, more effective management and improved survival.

Introduction

Pancreatic ductal adenocarcinoma (PDAC) is one of the most aggressive malignancies and the fourth leading cause of cancer-related mortality (1). Despite advances in diagnostic and therapeutic strategies over the past two decades, the prognosis for patients with PDAC remains poor and the 5-year survival rate is ~6% (2,3). PDAC usually causes no detectable symptoms in early stages and the majority of patients are diagnosed in advanced stages with poor survival rates (4). The difficulty in identifying patients at high risk of metastasis and/or recurrence highlights the challenge of the unsatisfactory prognosis of patients with PDAC.

Computed tomography (CT) and magnetic resonance imaging (MRI) are currently the predominant imaging modalities for the preoperative diagnosis of patients with PDAC. These imaging techniques can classify pancreatic cancer into resectable, borderline resectable, locally advanced and metastatic (5). This radiological classification is based on local vascular invasion and the presence of parenchymatous and peritoneal metastases and allows patients to be classified for either upfront surgery or oncologic treatment (6,7). Surgical treatment of resectable pancreatic tumor remains the optimum treatment option for this cancer (8,9). Therefore, careful and appropriate preoperative staging is important in patients with pancreatic cancer.

As PDAC is a group of heterogeneous diseases with different biological and clinical characteristics, the identification of prognostic and predictive markers is clinically relevant for improved management of the disease. Some

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gene and protein candidates have been reported to be associated with the disease progression of PDAC (10,11). These biomarkers could also be useful to clarify molecular PDAC subtypes that appear anatomically similar to aiding clinical management (7,12).

In particular, serum tumor markers are non-invasive additional tools that are increasingly used in various cancers including PDAC, to advance understanding of cancer pathophysiology, improve molecular stratification and thus achieve improved outcomes. Carcinoembryonic antigen (CEA) and carbohydrate antigen 19-9 (CA19-9) are tumor markers associated with both diagnosis and prognosis of patients with PDAC (13-16). CEA, a glycoprotein with a molecular weight of 180-200 kDa, was first isolated from fetal colon and colorectal cancer tissue in 1965 (17). CEA is overexpressed in several cancers, including colon, breast, lung and thyroid cancers (18-20). Moreover, serum CEA level is elevated in 30-60% of patients with PDAC (21,22) and it is even suggested to be an independent predictor of poor survival rates in patients with PDAC (23).

CA 19-9 is a tumor-associated antigen with a half-life of 4-8 days, whose epitope has been shown to be the sialylated Lewis antigen (24). CA19-9 is the only biomarker currently recommended in the National Comprehensive Cancer Network guidelines for clinical use in PDAC (25). However, when analyzing serum CA19-9 levels, in a clinical setting, several constraints should be considered. This is because CA19-9 may not have sufficient sensitivity and specificity in patients with certain metastases or obstructive jaundice (26). Moreover, sialylated Lewis antigen-negative individuals, who comprise ~5-10% of the population, have low or no CA19-9 secretion (27). Although CA19-9 is not accurate enough to be used for screening asymptomatic individuals for PDAC, it is currently the most useful blood test for distinguishing pancreatic cancer from chronic or recurrent pancreatitis, with a sensitivity of 70-90% and a specificity of 68-91% (28,29). It is also one of the most important prognostic factors for both patients with resectable disease and those with unresectable disease (30,31). In patients with PDAC, high expression of CA19-9 was associated with positive nodal status, an advanced disease and low survival rates (32). Measurement of CA19-9 as a prognostic factor provides valuable information to support therapeutic decision making, as patients with high preoperative CA19-9 levels are expected to have early recurrence. An elevated tumor marker value also implies a high probability of residual disease after resection (33). Preoperative serum levels of CEA and CA19-9 can be helpful to identify a subgroup of patients with poor outcomes after surgery (34). Currently, there are no biomarkers that can be discovered in the time interval between carcinogenesis and invasion, when the disease is potentially curable, due to low sensitivity and specificity in early detectable small tumors (35). CEA and CA19-9 still are the most studied biological markers for establishing both diagnosis and prognosis in PDAC (16,36,37). CEA and CA19-9 as biological markers are independent predictive factors for the presence of advanced PDAC with a positive predictive value of 91.0% for the combination of CEA (>7.0 ng/ml) and CA 19-9 (>305 U/ml) in predicting the presence of advanced disease (37). The aim of this study was to investigate the serum levels of CEA and CA19-9 and their associations with clinicopathological variables and survival outcomes in Libyan patients with PDAC.

Patients and methods

Clinicopathological data. The study group consisted of 123 patients with PDAC diagnosed between 2010 and 2018 at the National Cancer Institute in Misurata, Libya. Biological markers included serum levels of CEA and CA19-9, were determined in all patients before any treatment. The present study was conducted under research ethics approval by ethical committee at the National Cancer Institute, Misurata (Ethical Approval Number: EAN 6/2021). Written informed consent was obtained from all patients for surgical treatment, pathologic examinations and investigations performed according to the institutional guidelines of the National Cancer Institute, Misurata, Libya. Complete demographic and clinicopathological data included age, gender, family history, tumor location and size, lymph node status, stage, histological grade, type of treatment and follow-up data. These clinicopathological data were obtained from the patients' records and are summarized in Table I. The mean age of the patients was 61.2 years (range, 19-90 years). Tumor staging of pancreatic adenocarcinoma was evaluated according to the TNM classification (38). Radiological staging by Computed Tomography (CT) and/or Magnetic Resonance Imaging (MRI) was performed in all patients to assess tumor resectability. The extent of the tumor (local and distant) at the time of diagnosis was confirmed by imaging [CT, MRI, or Positron Emission Tomography (PET)]. Lymph node status was also assessed radiologically and positivity for malignancy was confirmed by histopathology of surgically resected nodes.

The overall time to recurrence was estimated as the interval between the date of surgery and the date of recurrence (local and/or distant). Disease recurrence (local and distant) was confirmed by imaging (CT, MRI, or PET) performed when clinical symptoms suggestive of disease recurrence were present. Disease-free survival was defined as the time between initial treatment and last follow-up that patients survive without disease recurrence.

Treatment and follow-up. Approximately 21 patients were treated by pancreaticoduodenectomy (Whipple procedure), while palliative surgery was performed in three patients and no surgery was performed in 99 patients who had metastases at the time of diagnosis. However, fine needle aspiration biopsy with image guidance and endoscopic retrograde cholangiopancreatography were performed in these patients for histopathological diagnosis. In the National Cancer Institute in Misurata the following guidelines were established: Adjuvant combined chemotherapy (gemcitabine and oxaliplatin) was given to 18 patients while 2 patients received combined chemotherapy based on FOLFIRINOX (folinic acid, fluorouracil, irinotecan and oxaliplatin) and 81 patients received palliative chemotherapy with capecitabine and/or gemcitabine. In addition, 22 patients were not eligible for chemotherapy, so these patients did not receive chemotherapy. Patients were followed-up until death or the end of the observation period (until October 2018). The median duration

3

Table I. Clinicopathologica	l variables	of 12.	3 patients	with
pancreatic ductal adenocarc	inoma.			

Variables	Number of patients (n=123)	Percent (%)
Age (years)		
<50	24	19.5
≥50	99	80.5
Sex		
Male	73	59.3
Female	50	40.7
Family history		
Positive	6	4.9
Negative	117	95.1
Tumor site		
Head	96	78.0
Tail	15	12.2
Body	12	9.8
Surgical resectability		
Resectable	24	19.5
Unresectable	99	80.5
Tumor size	,,,	00.5
Tullior size	4	3.3
T1 T2	4 13	3.3 10.6
T2 T3	13	11.4
T4	1	0.8
Cannot be assessed	91	74.4
Lymph node status	51	7
Negative	12	9.8
Positive	12	9.3 11.4
Cannot be assessed	97	78.9
)1	70.7
M M0	27	22.0
M0 M1	27 96	78.0
	90	78.0
Histology grade	6	4.0
G1 G2	6 30	4.9 24.4
G2 G3	50 87	24.4 70.7
	07	70.7
Stage	10	0.9
Stage 1	12 12	9.8 9.8
Stage 2		
Stage 3 Stage 4	3 96	2.4 78.0
Stage 4	90	/ 0.0
Systemic treatment	20	16.2
Adjuvant chemotherapy	20	16.3
Palliative chemotherapy	81	65.9
No treatment	22	17.9
(supportive therapy)		

of follow-up was 6 months (range 1-41 months). At the end of the follow-up period, 91 patients (74%) had succumbed to pancreatic cancer.

CEA and CA19-9 measurement. Prior to each treatment, approximately 5 ml of peripheral fasting blood was drawn from the forearm veins. The blood was immediately taken to the central laboratory of the National Cancer Institute in Misurata and then routinely centrifuged for 10 min at speed of 1,792 x g at 20-22°C temperature. The serum samples were first stored at 4°C. Then they were placed in polypropylene vials and stored at -80°C. The concentrations of CEA and CA19-9 in serum were determined using an electrochemiluminescence immunoassay (double antibody sandwich ELISA, cat. nos. TM E-4131 and TM E-4531 respectively; Labor Diagnostika Nord GmbH & CoKG) on a Roche cobas e 602 modules (Roche Diagnostics). This technology uses a sandwich chemiluminescence immunoassay; Chemibeads contain a chemiluminescent dye and Sensibeads contain a photosensitizer dye. Biotinylated antibodies (1:100) and Chemibeads form sandwiches and immune complexes are formed by further addition of Sensibeads. A chemiluminescence reaction is initiated at 680 nm and finally the signal is detected at 612 nm (according to the manufacturer's instructions). The accuracy of internal and external quality controls was determined according to the guidelines of RiliBAeK (17). The detection limit and blank limit were as follows: CEA: 0.2 and 0.12 ng/ml, CA19-9: 2.0 and 1.0 U/ml, CA15-3: 1.0 and 0.3 U/ml, respectively. Roche's original ancillary reagents were used, including streptavidincoated magnetic beads, anti-CEA monoclonal antibody and biotinylated anti-CA19-9 and anti-CEA monoclonal antibody and Ru-labelled anti-CA19-9.

Statistical analysis. Continuous variables were calculated using SPSS 26.0 for Windows (IBM Corp.). Frequency tables were analyzed using the χ^2 or Fisher's exact tests to evaluate the power of association between categorical variables. Kaplan-Meier curves were constructed for survival rate analysis and differences between curves were analyzed using the log-rank test. Multivariate survival analysis for the outcome [overall survival and disease-free survival (DFS)] was performed using the proportional hazard Cox model in a backward stepwise manner with the log-likelihood ratio (L-R) significance test, using standard values for the entry and exclusion criteria. The cut-off point for CEA of 5 ng/ml and for CA19-9 of 400 U/ml was used to distinguish between high-expression and low-expression tumors as it provided the best results for prognosis prediction in this and other studies (39,40). The assumption of proportional hazards was controlled by log-minus-log (LML) survival plots. P<0.05 was considered to indicate a statistically significant difference.

Results

Patient demographic and clinicopathologic variables. The demographic and clinicopathologic variables are shown in Table I. The mean age of the patients was 61.2 years (range, 19-90 years) and the majority of patients (80.5%) were >50 years old. Regarding sex distribution, PDAC was commoner in men (59.3%). A total of 4.9% of patients had a family history of pancreatic cancer and in 96 patients the tumors were located in the head of the pancreas (87%). The majority of patients (80.5%) had distant metastases or locally advanced unresectable disease. The most common T stage

was Tx (74.4%), followed by T3, T2, T1 and T4 in decreasing frequency (11.4, 10.6 and 83.3 and 0.8%, respectively). A total of 14 patients (11.4%) had positive lymph nodes and negative lymph nodes were detected in 12 patients, while lymph node status could not be assessed in the majority of patients (78.9%). Most patients had high-grade tumors (70.7%). According to the AJCC staging system, 99 patients were at stage IV (78%), 3 patients were at stage III, 12 patients were at stage II and 12 patients were treated by radical surgery, while palliative surgery was performed in three patients and no surgical intervention was performed in 99 patients. Adjuvant chemotherapy was performed in 20 patients, 81 patients received palliative chemotherapy and 22 patients were not eligible for chemotherapy.

General description of CEA and CA19-9 expression profiles. CEA and CA19-9 expressions at cut point of (5 ng/ml and 400 U/ml, respectively) are shown in Table II. The median CEA expression was 8 ng/ml (mean 22.8; range 1.1-377 ng/ml). CEA expression was low in 46 samples (<5 ng/ml) and high in 77 samples (\geq 5 ng/ml). The median value of CA19-9 was 389 U/ml (mean 762.4; range 1-10050 U/ml). CA19-9 was low in 64 samples (<400 U/ml) and high in 59 samples (\geq 400 U/ml). CEA expression was more frequent in tumors with high CA19-9 than in patients with low CA19-9 (P<0.0001).

Correlation of CEA expression with clinicopathological variables. The significant correlations between CEA expression (<5 ng/ml vs. \geq 5 ng/ml) and clinicopathological variables are shown in Table III. High CEA expression was significantly associated with surgically unresectable tumor (P<0.020), unevaluable lymph node status (P<0.007), advanced stages (P<0.0001) and distant metastases (P<0.002). However, age at diagnosis, gender, family history, tumor site, tumor size and histological grade showed no significant relationship with CEA expression.

Correlation of CA19-9 expression with clinicopathological variables. The correlations between CA19-9 expression at the cut point of 400 U/ml and clinicopathological variables are shown in Table III. High CA19-9 expression was more common in patients with surgically unresectable tumor (P<0.02), unevaluable lymph node status (P<0.001), advanced stage (P<0.0001) and distant metastases (P<0.002). However, age at diagnosis, gender, family history, tumor site, tumor size and histological grade showed no significant association with CA19-9 expression.

Correlation of serum CEA and CA 19-9 expression patterns with patient survival outcomes. Univariate survival analyzes (survival rates) with CEA expression at a cut-off point of 5 ng/ml and CA19-9 at a cut-off point of 400 U/ml are shown in Table IV. The survival rate was 28.6% in patients with low CEA expression and 24.7% in patients with high expression profile (P<0.016). The low CA 19-9 expression group had an improved survival rate than the high expression group (31.3 and 20.3%, respectively) (P<0.014).

Kaplan-Meier survival curves for both CEA and CA19-9 levels showed that shorter survival was associated with high Table II. CEA and CA19-9 expression in Libyan pancreatic ductal adenocarcinoma. CEA level at cut point of 5 ng/ml and CA 19.9 level at cut point of 400 U/ml.

Biological variables	Number of patients (n=123)	Percent (%)
CEA level (ng/ml)		
<5	46	37.4
≥5	77	62.6
CA 19-9 level (U/ml)		
<400	64	52.0
≥400	59	48.0

CEA, carcinoembryonic antigen; CA19-9, carbohydrate antigen.

CEA and high CA19-9 levels (Fig. 1). On the other hand, patients with low CEA and CA19-9 levels were associated with a lower recurrence rate and therefore had longer disease-free survival (P<0.002 and P<0.0001, log rank, respectively, (Fig. 2). However, multivariate Cox regression analysis revealed that both serum CEA and CA19-9 levels were not independent markers of patient survival in relation to patient age, lymph node status, tumor grade and tumor stage.

In addition, the correlation of surgical and/or systemic treatment with clinicopathologic variables and patient outcomes was observed in the present study, as shown in Table V. Patients who underwent radical surgical treatment were associated with small tumor size (P<0.0001), negative lymph node status (P<0.0001), early stage (P<0.0001), low expression of CEA level (P=0.020) and low expression of CA19-9 level (P=0.020). Survival was 37.5% in patients with resectable tumor and 23.2% in patients with non-resectable tumor (P<0.0001). Analysis using Kaplan-Meier survival curves for surgical resectability showed that shorter survival was associated with non-resectable tumor patients (P<0.0001). The best survival was observed in patients receiving adjuvant chemotherapy compared with patients receiving palliative chemotherapy or supportive care only (35.0, 27.3 and 23.5%, respectively).

Discussion

PDAC is an aggressive heterogenous malignancy associated with a very poor prognosis and imposes an enormous burden on patients, their families and the healthcare system as a whole. Therefore, the identification and evaluation of diagnostic, prognostic and predictive markers is urgently needed for clinicians to achieve improved outcomes. Unfortunately, most patients are often diagnosed with metastatic disease or locally advanced, unresectable tumor (2-4,41). Although newer techniques have been used and target groups have been rigorously selected, screening programs are still ineffective for a large group and may be harmful, as noted in 2019 (42). Therefore, regular screening with endoscopic ultrasound and MRI/CT imaging is recommended for individuals who are at high risk for genetic diseases (43,44). Therefore, treatment plans are based on the extent of the tumor, performance

Clinicopathological		CE		CA19-9 expression (%)			
variable	Number	<5 ng/ml	≥5 ng/ml	P-value			P-value
Age /years				0.91			0.25
<50	24	37.5	62.5		62.5	37.5	
≥50	99	37.4	62.6		49.5	50.5	
Sex				0.38			0.91
Male	73	34.2	65.8		52.1	47.9	
Female	50	42.0	58.0		52.0	48.0	
Family history				0.07			0.08
Positive	6	16.7	83.3		33.3	66.7	
Negative	117	38.5	61.5		53.0	47.0	
Site				0.96			0.37
Head	96	37.5	62.5	0.90	54.2	45.8	0.57
Elsewhere	27	37.0	63.0		44.4	55.6	
Surgical resectability		0,10	0010	0.020		2210	0.020
Resectable	24	58.3	41.7	0.020	75.0	25.0	0.020
Unresectable	24 99	32.3	67.7		46.5	53.5	
	,,,	52.5	07.7	0.092	10.5	55.5	0.091
T stage T1	4	50.0	50.0	0.092	50.0	50.0	0.091
T1 T2	13	69.2	30.8		50.0 76.9	23.1	
T2 T3	13	42.9	57.1		70.9	28.6	
T4	1	0.00	100		0.00	100	
Tx	91	31.9	68.1		46.2	53.8	
	71	51.9	00.1	0.007	40.2	55.0	0.001
Lymph node status Positive	12	50.0	50.0	0.007	85.7	14.3	0.001
	12 14	83.3	30.0 16.7		66.7	14.5 33.3	
Negative Nx	97	29.9	70.1		45.4	55.5 54.6	
	91	29.9	70.1	0.26	45.4	54.0	0.(2
Histological grade	(22.2	((7	0.26	50.0	50.0	0.62
G1 G2	6 30	33.3	66.7		50.0	50.0	
		50.0	50.0		60.0	40.0 50.6	
G3	87	33.3	66.7		49.4	30.0	
Stage	<u>.</u>		26.2	<0.0001	00.0	165	<0.0001
Early stage	24	70.8	29.2		83.3	16.7	
Late stage	99	29.3	70.7		44.4	55.6	
Metastasis				0.002			0.002
MO	27	63.0	37.0		77.8	22.2	
M1	96	30.2	69.8		44.8	55.2	

Table III. The association between CEA and CA19-9 expressions with clinicopathological variables in pancreatic adenocarci-
noma cancer (n=123). Comparison between low CEA and CA19-9 expression group and high expression group.

Bold type indicates statistically significant values. CEA, carcinoembryonic antigen; CA19-9, carbohydrate antigen.

status of patients and various other clinical factors. However, not all patients benefit from conventional cancer therapy (45). Despite marked advances in treatment strategies, advanced PDAC cancer remains incurable and the goals of therapy range from relief of symptoms to prolonging survival. Of note, surgical treatment of a resectable tumor remains the best treatment option for this cancer (8,9). A number of studies suggest that multiple genes may correlate with PDAC patient outcomes (10,11). Moreover, tumor biology is heterogeneous in patients with PDAC with similar tumor anatomy (12,13). This highlights the importance of appropriate preoperative radiological PDAC staging and molecular and genetic stratification/classification to understand the biological behavior of this highly aggressive malignancy. Numerous studies have investigated the efficacy of various biological markers as diagnostic, predictive and prognostic markers in

			Survival analysis			
Variables	Threshold	No of patients	Median survival (months)	Mean survival (months)	Survival rate (percent)	P-value
All patients		123	6.00	8.55	26.0	
CEA level						0.016
	<5	46	8.00	11.02	28.6	
	≥5	77	5.00	7.27	24.7	
CA19-9 level						
	<400	64	7.00	9.89	31.3	0.014
	≥400	59	5.00	7.10	20.3	

Table IV. Univariate survival according to analysis of CEA expression (cut point of 5 ng/ml) and CA19-9 (cut point of 400 U/m) in Libyan pancreatic ductal adenocarcinoma (n=123).

Bold type indicates statistically significant values. CEA, carcinoembryonic antigen; CA19-9, carbohydrate antigen.

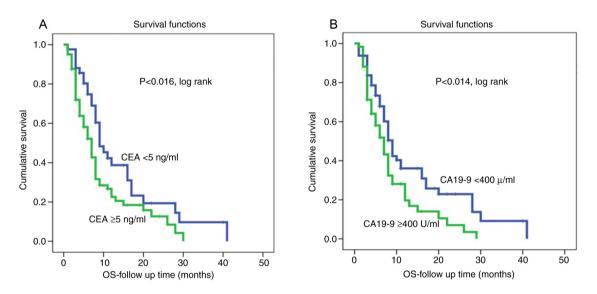


Figure 1. OS according to analysis of CEA expression and CA19-9 in pancreatic adenocarcinoma (Kaplan-Meier curves). (A) CEA level at cut point of 5 ng/ml. (B) CA 19-9 level at cut point of 400 U/ml. OS, overall survival; CEA, carcinoembryonic antigen; CA19-9, carbohydrate antigen.

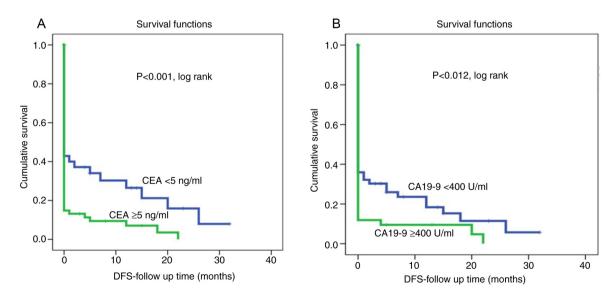


Figure 2. DFS according to analysis of CEA expression and CA19-9 in pancreatic adenocarcinoma (Kaplan-Meier curves). (A) CEA level at cut point of 5 ng/ml. (B) CA 19-9 level at cut point of 400 U/ml. DFS, disease free survival; CEA, carcinoembryonic antigen; CA19-9, carbohydrate antigen.

P-value 0.25				
0.25	Unresectable	Resectable	Number	Clinicopathological variable
				Age /years
	62.5	37.5	24	<50
	49.5	37.4	99	≥50
0.72				Sex
	79.5	20.5	73	Male
	82.0	18.0	50	Female
0.08				Family history
	100.0	0.00	6	Positive
	79.5	20.5	117	Negative
				Site
0.88	80.2	19.8	96	Head
	81,5	18.5	27	Elsewhere
<0.0001	,			T stage
	0.00	100.0	4	T1
	30.8	69.2	13	T2
	42.9	57.1	14	T3
	100.0	0.00	1	T4
	96.7	3.3	91	Тх
<0.0001				Ν
	78.6	21.4	12	Positive
	3.1	83.3	14	Negative
	96.9	3.1	97	Nx
0.62				Histological grade
0.02	16.7	83.3	6	G1
<0.0001				
\$0.0001	20.8	79.2	24	
0.020	2.112	5.11		-
0.020	60.6	30 /	46	
0.020	07.0	13.0	11	
0.020	71.0	28.1	61	
	63.3 90.8 20.8 94.9 69.6 87.0 71.9 89.8	33.5 36.7 9.2 79.2 5.1 30.4 13.0 28.1 10.2 mbryonic antigen; CA19-9,	30 87 24 99 46 77 64 59	G2 G3 Stage Early stage Late stage CEA level (ng/ml) $<5 \ge 5$ CA19-9 level (U/ml) $<400 \ge 400$

Table V. Surgical resectability accordance to demographic, clinicopathological and biological features in Libyan pancreatic ductal adenocarcinoma (n=123).

Bold type indicates statistically significant values. CEA, carcinoembryonic antigen; CA19-9, carbohydrate antigen.

PDAC (22,23,34,38,46). Among them, CEA and CA19-9 are the most commonly used tumor biomarkers.

The present study performed a detailed retrospective analysis of 123 patients with PDAC diagnosed and treated at the National Cancer Institute, Misurata, Libya. CEA and CA19-9 expression levels at cut-off points (5 ng/ml and 400 U/ml, respectively) were found to be the most promising discriminators of both clinicopathological variables and survival outcomes. In the cohort of present study, the median value of CEA in serum was 8 ng/ml and high CEA expression was detected in 63% of patients. The median value of CA19-9 in serum was 389 U/ml and high CA19-9 expression was detected in 48% of patients. In addition, the present study showed that the majority of Libyan patients had higher malignancy grade, such as locally advanced inoperable tumors, poorly undifferentiated tumors and distant metastases. Tumor size and lymph node status could not be determined in 75 and 79% of tumors, respectively. On the other hand, only 20% of patients had resectable tumors and 24 patients had early-stage PDAC at the time of diagnosis. These figures are consistent with other published data, which also reported that only 20% of patients diagnosed with PDAC were eligible for surgical resection and the majority of patients had advanced disease stage at the time of diagnosis (42,47-50).

Patients with high expression of CEA and/or CA19-9 are often associated with a worse prognosis. Therefore, increased expression of CEA and CA19-9 might indicate that the tumors are already at an advanced stage. Moreover, tumor progression was associated with higher levels of these tumor markers. Consistent with these findings, Lee *et al* (45) report that expression of CEA and CA19-9 is closely associated with tumor differentiation degree, lymph node metastasis and tumor progression. The aggressive behavior of PDAC, reflected in both unclear clinical symptoms at presentation and genetic heterogeneity of the tumor, may explain the advanced stages of the disease at diagnosis (41,45).

The present study also showed that the highly aggressive malignant phenotype of PDAC, manifested by surgically unresectable tumors, unevaluable lymph nodes, advanced stage and distant metastases, was significantly associated with high serum CEA and CA19-9 expression profiles. On the other hand, surgically resectable tumors, negative lymph nodes, early stage and low risk of metastasis were more common in the group with low serum CEA and CA19-9 expression. CEA expression was more frequent in tumors with high CA19-9 expression compared with those with low CA19-9 expression (P<0.0001). The most important finding of the present study was undoubtedly the significant correlation of CEA and CA19-9 expression with disease progression, especially overall survival and disease-free survival. The median follow-up time of the cohort study was six months and ~74% of patients had died of PDAC at the end of the follow-up period. Patients with low serum CEA and CA19-9 levels had a lower recurrence rate and lived longer than their counterparts with high serum levels. Analysis using Kaplan-Meier curves also showed that short survival was more common in the group with high CEA and CA19-9 levels, while the group with low CEA and CA19-9 levels had longer disease-free survival (P<0.002 and P<0.0001, respectively). These findings are consistent with the results of other previous studies (34,38,46), which indicated that Libyan patients with high expression of CEA and CA19-9 were associated with poor prognosis. Although associations between serum CEA and CA19-9 levels and treatment outcomes are found in numerous studies, some discrepancies are also reported. While some studies agreed with the findings of the present study and suggest that overexpression of CEA is associated with poor prognosis (6,38,46), others such as the study by Poruk et al (39) reported that CA19-9 is a useful biological marker in PDAC to predict disease extent, surgical resectability, disease progression and response to treatment. The present study showed that the combination of CEA and CA19-9 is more successful for prognosis prediction than a single biological marker (47-49). Therefore, Xu et al (50) showed that elevated CEA levels in combination with CA19-9 can significantly increase the prognostic efficacy of CA19-9. The present study confirmed this association and found that CEA expression was more frequent in tumors with high CA19-9 compared with patients with PDAC with low CA19-9. The present study showed that the prognostic value of CEA and CA19-9 was almost the same and both markers did not to act as independent prognosticators for the outcome of patients with PDAC. However, van Manen *et al* (37) reported that both biological markers (CEA and CA19-9) were independent predictors for an advanced PDAC cohort from the Netherlands and that the predictive power of CEA was higher than that of CA19-9. These discrepancies could be due to several factors, including cohort size, diagnostic techniques (imaging and/or biopsies), genomic background of the cohort (ethnicity) and/or stage at diagnosis. However, other reports have found that elevated CEA and advanced stage are independent factors for poor survival of patients with PDAC (51,52).

The present study confirmed the role of both CEA (>8 ng/ml) and CA19-9 (>389 U/ml) serum levels as key markers for multidisciplinary management of Libyan patients with PDAC to aid decision making. Despite their significant associations with survival outcomes, neither marker was informative enough to serve as an independent predictor for advanced patients with PDAC.

The median serum levels of CEA and CA19-9 for all PDAC tumors were 8 ng/ml and 389 U/ml, respectively. Tumors with higher serum CEA and CA19-9 levels were found in 62.6 and 48% of PDAC cases, respectively. Significantly, patients with PDAC with higher CEA and CA19-9 serum levels had aggressive tumor grade, higher recurrence rate and shorter survival time and should be treated carefully. The prognostic value of CEA and CA19-9 was almost equivalent but not sufficient to be an independent prognosticator of patient outcome. An extended multinational study with a larger cohort is needed to confirm the prognostic value of both serum levels and tissue expression of these two promising markers.

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Availability of data and materials

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

Authors' contributions

EE performed statistical analysis, designed the present study and drafted manuscript. MEd designed the study and collected demographic and clinicopathologic data and performed laboratory work and data analysis. OA, MEl and AJ collected and analyzed data and drafted the manuscript. MAS and MA performed data interpretation and analysis, drafting and proof reading and discussions. AB prepared the figures, reviewed the study, interpretated data and aided in drafting and proof reading of the manuscript. All authors critically reviewed and approved the final version of the manuscript. EE and MEl confirm the authenticity of all the raw data.

Ethics approval and consent to participate

The present study was conducted under research ethics approval by ethical committee at the National Cancer Institute, Misurata (Ethical Approval Number: EAN 6/2021). Written informed consent was obtained from all patients for surgical treatment, pathologic examinations and investigations performed according to the institutional guidelines of the National Cancer Institute, Misurata, Libya.

Patient consent for publication

Not applicable.

Competing interests

The authors declare that they have no competing interests.

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