

NIH Public Access

Author Manuscript

Hum Pathol. Author manuscript; available in PMC 2014 June 01

Published in final edited form as:

Hum Pathol. 2013 June ; 44(6): . doi:10.1016/j.humpath.2012.09.001.

Loss of PTEN Expression Is Associated with Recurrence and Poor Prognosis in Patients with Pancreatic Ductal Adenocarcinoma

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Abstract

Phosphatase and tensin homolog (PTEN) is a tumor suppressor in the AKT/mTOR pathway. Animal model studies have shown that loss of PTEN function is involved in the progression of pancreatic cancer. However, the prognostic significance of loss of PTEN expression in pancreatic cancer is unclear. PTEN expression was evaluated by immunohistochemistry on tissue microarrays consisting of multiple cores of 133 resected stage II pancreatic ductal adenocarcinomas. A PTEN expression score was calculated as the product of the percentage of positive tumor cells and the intensity of PTEN staining. We categorized PTEN expression for each tumor as retained (PTEN score>5) or loss (PTEN score 5). Thirty-four (25.6%) patients had tumors with loss of PTEN expression, and 99 (74.4%) had tumors with retained PTEN expression. Recurrence/metastasis was observed in 88.2% (30/34) of patients whose tumors showed loss of PTEN compared to 68.7% (68/99) of patients whose tumors showed retained PTEN (p=0.03). Patients whose tumors showed loss of PTEN had a shorter overall survival (median: 19.9±3.6 months) than patients whose tumors had retained PTEN (32.7 ± 5.0 months, p = 0.03). In a multivariate analysis, loss of PTEN expression was an independent prognostic factor for poor overall survival in patients with stage II PDA. No significant correlations between loss of PTEN expression and other clinicopathologic parameters were observed (p>0.05). Assessment of PTEN expression may be used as a prognostic marker for patients with resected pancreatic ductal adenocarcinoma.

DISCLOSURE/CONFLICTS OF INTEREST The authors declare no conflicts of interest.

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PTEN; Pancreatic cancer; Survival; Prognosis

INTRODUCTION

Pancreatic ductal adenocarcinoma is the fourth leading cause of cancer death in the United States (1, 2). Approximately 15–20% of patients present with localized disease, and the 5-year overall survival rate is less than 5% (3). Small tumor size, absence of lymph node metastases, and negative resection margins (R0) are independent prognostic factors predicting long term survival in patients with pancreatic ductal adenocarcinoma who have undergone pancreatectomy (4–6).

The development of pancreatic ductal adenocarcinoma represents the accumulation of successive mutations in proto-oncogenes and loss of tumor suppressor genes, typically beginning with activating mutations in *KRAS* followed by inactivation of tumor suppressor genes, including *p16/INK4A*, *p53*, and *SMAD4/DPC4* (7–11). *KRAS* mutations are found in greater than ninety-percent of pancreatic ductal adenocarcinoma and occur early in tumorigenesis (9–11). Once activated, KRAS functions through multiple signaling pathways, including the phosphoinositide 3-kinase/AKT (PI3K/AKT) pathway. Coinciding with *KRAS* activation, Asano and colleagues have demonstrated that activation of the PI3K/AKT pathway and its downstream effectors, NF- B and MYC, is essential for growth and survival of pancreatic ductal adenocarcinoma (12).

Phosphatase and tensin homolog (PTEN) is an important tumor suppressor encoded on chromosome 10q23.3 (13, 14). Loss of PTEN function has been implicated in the tumorigenesis of many human malignancies, including gliomas, endometrial cancers, thyroid cancers and pancreatic cancers. Loss of PTEN function can be due to multiple mechanisms, including biallelic or monoallelic deletions, mutations, gene silencing by promoter methylation or dysregulation of mRNA by microRNAs. In mouse model studies, pancreatic-specific knockout of Pten leads to ductal metaplasia from the expansion of centroacinar cells and the development of pancreatic ductal carcinoma (15). Single copy deletion of *Pten* in *Kras*^{G12D} mice (*Pdx1-Cre LSL-Kras*^{G12D} *Pten*^{L/+} mice) results in more extensive acinar-ductal metaplasia, pancreatic intra-epithelial neoplasia, increased epithelial proliferation, and higher frequencies of invasion and metastasis compared to Pdx1-Cre LSL-*Kras*^{G12D} mice (16). Similar results were observed in the *Kras*^{G12D} mice with conditional *Pten* knockout (*Pten*^{lox/lox}) (17). Pten also functions synergistically with the tumor suppressor, Smad4, during the development and progression of pancreatic cancer in mouse models (18). These data support the notion that PTEN is one of the major tumor suppressors in pancreatic tumorigenesis. Although deletion or loss of function mutations in *PTEN* have not been detected with significant frequency in pancreatic cancer (19, 20), low or no PTEN expression by immunohistochemistry has been reported in up to 70% of human pancreatic ductal adenocarcinoma samples (16). Recently, Feng et al. reported that PTEN expression in pancreatic ductal adenocarcinoma patients with liver metastasis is lower than those patients who had no liver metastasis (21). They also showed that high PTEN expression is associated with a better 5-year survival in pancreatic ductal adenocarcinoma patients with liver metastasis (21). However, the prognostic value of PTEN expression and its clinical impact in patients who underwent surgical resection for pancreatic ductal adenocarcinoma is not clear. In this study, we sought to evaluate by immunohistochemistry the frequency of loss of PTEN expression in 133 patients with stage II pancreatic ductal adenocarcinoma. The results of PTEN expression were correlated with clinicopathologic parameters and patient survival.

MATERIALS AND METHODS

Patient population

Our study population consisted of 133 patients (78 male and 55 female) with stage II pancreatic ductal adenocarcinoma who underwent pancreatectomy as initial treatment for pancreatic ductal adenocarcinoma at our institution from 1990 to 2010. Patient age ranged from 24.9 to 84.8 years (median age: 64.6 years). Five patients with other disease stages (one patient each with stage IA, IB, or III disease and two patients with stage IV disease) were excluded because there were insufficient case numbers to be representative. Patients who received any form of neoadjuvant chemotherapy and/or radiation therapy were excluded. One hundred, thirteen patients (85.0%) underwent pancreaticoduodenectomy, 18 (13.5%) underwent distal pancreatectomy, and 2 (1.5%) underwent total pancreatectomy. One hundred eight patients (81.2%) had an R0 resection (all resection margins negative by histology) and 25 (18.8%) patients had an R1 resection (microscopic disease involving one or more margins). No patients had an R2 resection. Twenty-five (18.8%) patients received adjuvant chemotherapy alone, 73 (54.9%) received combined adjuvant chemoradiation therapy and 35 (26.3%) did not receive adjuvant therapy. The study was approved by the Institutional Review Board of the University of Texas MD Anderson Cancer Center.

Tissue microarray (TMA) construction

Hematoxylin and eosin (H & E) stained slides and their matched formalin-fixed paraffin embedded tissue blocks were retrieved and reviewed for representative areas of tumor and non-neoplastic pancreatic parenchyma. Two 1.0 mm cores from different areas of each tumor and one 1.0 mm core of the matched non-neoplastic pancreatic tissue were used for the TMA construction. The TMA was constructed as previously described using a tissue microarrayer (Beecher Instruments, Sun Prairie, WI)(22).

Immunohistochemistry for PTEN expression

PTEN immunohistochemistry was performed on 4 μ m unstained sections from TMA blocks using a mouse monoclonal antibody against PTEN (DAKO clone 6H2.1, Carpeninteria, CA, 1:100 dilution). The immunohistochemical stained slides were evaluated independently by two pathologists (W.C.F. and H.W.) for percentage of tumor cells with PTEN immunoreactivity and the intensity of PTEN immunoreactivity (0-negative, 1-weak, 2-moderate, and 3-strong). The PTEN expression score was calculated as the product of the percentage of positive tumor cells and the intensity of PTEN staining. We categorized each tumor as either retained PTEN expression (defined as PTEN score >5) or loss of PTEN expression (defined as PTEN score >5). For tumors that retained PTEN expression, we further categorized those tumors as either PTEN-low (PTEN score >5, but 50) or PTEN-high (PTEN score>50). If there was a discrepancy in PTEN expression between two duplicate cores from the same tumor, the tumor was classified based on the result of the core with positive PTEN staining. All cases with a discrepancy in PTEN expression between pathologists were re-reviewed together, and the consensus results for PTEN expression were used.

Statistical analyses

The immunohistochemical staining results were correlated with clinicopathologic features and survival. Chi-square and Fisher's exact tests were used to compare categorical data. Overall survival and disease-free survival curves were constructed using a Kaplan-Meier method, and log-rank tests were used to evaluate the statistical significance of differences. The prognostic significance of clinicopathologic parameters and PTEN expression was determined using a Cox-regression multivariate analysis via a backward stepwise procedure.

Statistical calculations were performed using Statistical Package for Social Sciences software (version 12.0, SPSS Inc., Chicago, IL). A p-value of <0.05 were considered statistically significant.

RESULTS

PTEN expression in pancreatic ductal adenocarcinoma

Of the 133 samples, 34 tumors (25.6%) showed loss of PTEN expression (Figure 1A & 1B) and 99 tumors (74.4%) showed retained PTEN expression. In the cases that had retained expression for PTEN, the tumor showed both cytoplasmic and/or nuclear staining (Figure 1C & 1D). Forty-three (32.3%) cases were PTEN-low and 56 (42.1%) cases were PTEN-high. Heterogenous PTEN expression between the two tissue cores of the tumor was observed in 22 (16.5%) cases; in these cases, the tumor was classified based on the core with positive staining. All 22 tumors were classified as retained PTEN expression. The matched non-neoplastic pancreatic tissue cores retained PTEN expression in all cases examined and served as the internal positive controls for our immunohistochemical staining (Figure 1E and F).

Association of PTEN expression and clinicopathologic features

The correlations between PTEN expression and clinicopathologic features are summarized in Table 1. Of 34 patients whose tumor lost PTEN expression, 30 (88.2%) had local recurrence or distant metastasis compared to 68.7% (68/99) of patients whose tumor retained PTEN expression (p=0.03). Lymph node metastasis was present in 88.2% (30/34) in cases that lost PTEN expression, compared to 71.7% (71/99) in the patients whose tumor retained PTEN expression (p=0.05). We did not observe significant correlation between the loss of PTEN expression and patient gender, age, tumor differentiation, tumor size, resection margin status or post operative chemo/radiation therapies (p>0.05, Table 1). Similarly, we did not observe significant difference in any of the clinicopathologic factors between the group with PTEN-low tumors and the group with PTEN-high tumors (data not shown).

Loss of PTEN expression correlated with poor overall survival

The median follow up time was 18.5 months (range: 4.2 to 82.9 months) for 88 patients who died of disease. For 45 patients who did not die from the disease, the follow-up time ranged from 7.1 months to 214.0 months with the median follow-up time of 40.7 months after pancreatectomy. No patients died in the immediate postoperative period, and none were lost to follow up. The median disease free survival and overall survival in this study population were 12.1 ± 1.9 months and 25.2 ± 3.0 months, respectively. Loss of PTEN expression correlated with poor overall survival. For the patients whose tumor retained PTEN expression, the median overall survival was 32.7 ± 5.0 months compared to the median overall survival of 19.9 ± 3.6 months for patients whose tumor lost PTEN expression (p=0.03, log-rank method, Figure 2A). Patients whose tumor retained PTEN expression also had longer disease free survival (median: 15.4 ± 2.0 months) than those whose tumor lost PTEN expression (median: 9.4 ± 1.7 months) but was not statistically significant (p=0.09, log-rank method, Figure 2B). The univariate analyses of overall survival in correlation with the clinicopathologic parameters and PTEN expression are shown in Table 2. In addition to the loss of PTEN protein expression, overall survival was also associated with tumor size, margin status, and lymph node metastasis (p < 0.05) but not other clinicopathologic factors (p 0.05). No significant difference in either disease free survival or overall survival between the group with PTEN-low tumors and the group with PTEN-high tumors was observed (data not shown). In multivariate analysis, loss of PTEN expression was a prognostic factor for overall survival in patients with stage II disease independent of tumor size, margin status, and lymph nodal metastases (Table 3).

DISCUSSION

In this study, we examined the PTEN expression by immunohistochemistry in 133 patients with stage II pancreatic ductal adenocarcinoma. We found that loss of PTEN protein expression correlated with tumor recurrence/distant metastasis after pancreatectomy and shorter overall survival. Loss of PTEN protein expression was an independent prognostic factor in our patient population. Our data supports the notion that PTEN is an important tumor suppressor in the tumorigenesis and progression of pancreatic ductal adenocarcinoma.

PTEN has been implicated in the tumorigenesis and progression of pancreatic ductal adenocarcinoma. It has been demonstrated that heterozygous Pten deletion in the pancreas accelerates the development of pancreatic intraepithelial neoplasia and pancreatic ductal adenocarcinoma in Kras^{G12D} pancreatic cancer mouse model(17) and that Pten haploinsufficiency was sufficient to induce development of pancreatic ductal adenocarcinoma in murine models. Moreover, mouse survival was dictated in a Pten dosedependent fashion(16). To test this hypothesis in human pancreatic ductal adenocarcinoma samples, we compared the clinicopathologic parameters and survival between a group of patients' tumors with retained PTEN expression and another with loss of PTEN expression. In this study, low or no PTEN expression was observed in 57.9% of stage II pancreatic ductal adenocarcinoma samples. Our findings were consistent with the previous reports, which identified 38.8% and 70% of pancreatic ductal adenocarcinoma samples have low or no PTEN expression (16, 21). We found loss of PTEN expression in 25.6% of our cohort, which was similar to results reported by others using immunohistochemistry performed on both whole tissue and TMA sections (12, 23–25). Given that our study included tumor samples from 133 patients, our results in conjunction with earlier studies show that loss of PTEN staining is not an infrequent occurrence in pancreatic ductal adenocarcinoma.

PTEN functions as an important tumor suppressor in the tumorigenesis of many human malignancies. Loss of PTEN protein expression is reported to be associated with worse survival in advanced endometrial cancer, lung, breast, and colon cancers (26-28), (29, 30). The prognostic significance of loss of PTEN expression in patients with pancreatic ductal adenocarcinoma was unclear. Feng et al reported high levels of PTEN expression in 61.2% pancreatic ductal adenocarcinoma patients without liver metastasis compared to 29.9% in pancreatic ductal adenocarcinoma patients who had liver metastasis (21). Their data suggest that loss of PTEN plays a role in liver metastasis in patients with pancreatic ductal adenocarcinoma. Consistent with this notion, we found that loss of PTEN expression was associated with higher frequency of recurrence/distant metastasis in patients with stage II pancreatic ductal adenocarcinoma compared to those patients whose tumor retained PTEN expression. In addition, we also found that loss of PTEN expression was associated with poor overall survival and was an independent prognostic factor in our patient population with stage II pancreatic ductal adenocarcinoma by multivariate analysis. Loss of nuclear staining for PTEN has been reported to be associated with poor survival in colon cancer (29, 30). In this study, we did not observe a significant correlation in the loss of nuclear staining for PTEN with either disease free survival or overall survival in our study population. Our data did not support the previous report by Pham and colleagues, who found a higher percentage of stage I and II tumors had low levels of PTEN protein expression than stage III and stage IV tumors (25). The limitation of our study was the inclusion of only patients with stage II disease as there were only a small number of patients with other stages of disease. However, our findings are corroborated by a recent report demonstrating high levels of PTEN expression associated with a better 5-year survival rate compared to those with no or low PTEN expression in patients with pancreatic ductal adenocarcinoma metastatic to the liver (21).

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The mechanisms of loss PTEN expression in pancreatic ductal adenocarcinoma are unclear. Recently, Ying *et al* demonstrated the deletion of one or two copies of the *PTEN* locus on 10q23 in 15% of the 61 pancreatic ductal adenocarcinoma samples examined using high resolution array comparative genomic hybridization (16). This finding suggested that loss of PTEN expression in pancreatic ductal adenocarcinoma may be due to the deletion of *PTEN* locus, which had been reported in other malignancies. The loss of PTEN expression is accompanied by the gain/amplification in *AKT2* locus and activation of AKT. The aberrant activation of PI3K/AKT pathway by loss of PTEN and/or activation of AKT accelerates mutant KRAS driven malignant progression of pancreatic ductal adenocarcinoma (31). Our findings that loss of PTEN expression correlated with recurrence/distant metastasis and poor overall survival in patients with pancreatic ductal adenocarcinoma provides additional support for the functions of PTEN/PI3K pathway in pancreatic cancer progression. Therefore, targeting PI3K/mTOR pathway in combination with other treatment modality may provide more effective treatment for pancreatic ductal adenocarcinoma.

In summary, loss of PTEN is not an infrequent occurrence in pancreatic ductal adenocarcinoma. Loss of PTEN expression correlates with recurrence/distant metastasis and survival in patients with stage II pancreatic ductal adenocarcinoma. Thus, loss of PTEN expression supports the implication that PTEN function plays an important role in progression of pancreatic ductal adenocarcinoma. Evaluation of PTEN expression by immunohistochemistry has prognostic implications in patients with stage II pancreatic cancer and may have therapeutic implications as well. As such, the utility of targeting the PI3K/Akt pathway via PI3K inhibitors and mTOR inhibitors should be evaluated in pancreatic ductal adenocarcinoma.

Acknowledgments

Supported by the National Institutes of Health grant (1R21CA149544-01A1) and G. S. Hogan Gastrointestinal Cancer Research Fund at The University of Texas M.D. Anderson Cancer Center

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Figure 1.

Representative micrographs show PTEN expression in pancreatic ductal adenocarcinoma samples and benign pancreatic tissue. A and B, a moderately differentiated pancreatic ductal adenocarcinoma with negative staining for PTEN; C and D, strong cytoplasmic and nuclear staining for PTEN in a moderately differentiated pancreatic ductal adenocarcinoma; E and F, Representative benign pancreatic tissue that is strongly positive for PTEN (Original magnification, $40 \times$ for A, C, and E; $400 \times$ for B, D, and F).

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Figure 2.

Kaplan-Meier curves for overall survival (A) and disease-free survival (B) by PTEN expression in patients with stage II pancreatic ductal adenocarcinoma. Patients whose tumors lost PTEN expression had shorter overall survival than patients whose tumors retained PTEN expression.

Table 1

Clinicopathologic features correlated with PTEN expression in stage II pancreatic ductal adenocarcinomas

Characteristics	n	Retained PTEN $(n = 99)$	Loss of PTEN $(n = 34)$	p value
Age (y)				0.90
<60	39	28	11	
60–70	57	43	14	
>70	37	28	9	
Gender				0.67
Male	78	57	21	
Female	55	42	13	
Tumor size (cm)				0.74
<= 2.0	22	17	5	
> 2.0	111	82	29	
Tumor differentiation				0.92
Well	10	8	2	
Moderate	88	65	23	
Poor	35	26	9	
Margin status				0.07
Negative	108	84	24	
Positive	25	15	10	
Lymph node status (stage)				0.05
Negative (IIA)	32	28	4	
Positive (IIB)	101	71	30	
Adjuvant chemo/radiation				0.26
No	35	29	6	
Yes	98	70	28	
Recurrence or distant metastas	is			0.009
No	35	31	4	
Local recurrence	26	14	12	
Distant metastasis	72	54	18	

Table 2

Univariate analysis of overall survival in patients with stage II pancreatic ductal adenocarcinomas

Variables	n	HR (95% CI)	p value			
Age						
<60 years	39	1.00				
60 – 70 years	57	1.07 (0.65 – 1.77)				
>70 years	37	1.45 (0.83 – 2.55)	0.39			
Gender						
Female	55	1.00	0.14			
Male	78	0.73 (0.48 – 1.11)				
Tumor differentiation						
Well	10	1.00				
Moderate	88	2.70 (1.08 - 6.76)				
Poor	35	2.62 (0.99 - 6.96)	0.05			
Tumor size (cm)						
<= 2.0	22	1.00				
> 2.0	111	3.25 (1.56 - 6.77)	0.002			
Margin status						
Negative	108	1.00				
Positive	25	1.75 (1.04 – 2.93)	0.04			
Lymph node status (stage)						
Negative (IIA)	32	1.00				
Positive (IIB)	101	2.05 (1.18 - 3.55)	0.01			
Adjuvant chemo/radiation						
No	35	1.00				
Yes	98	0.69 (0.42 – 1.14)	0.15			
PTEN expression						
Retained	99	1.00				
Loss	34	1.65 (1.06 – 2.58)	0.03			

^aAbbreviations: HR, hazard ratio; CI, confidence interval

Table 3

Multivariate analysis of overall survival in patients with stage II pancreatic ductal adenocarcinomas

Variables	n	HR ^a (95% CI ^a)	p value			
Tumor size (cm)						
<= 2.0	22	1.00				
> 2.0	111	3.47 (1.66 – 7.25)	0.001			
Margin status						
Negative	108	1.00				
Positive	25	1.39 (0.82 – 2.36)	0.228			
Lymph node status (stage)						
Negative (IIA)	32	1.00				
Positive (IIB)	101	1.52 (0.83 – 2.80)	0.178			
PTEN expression						
Retained	99	1.00				
Loss	34	1.79 (1.14 – 2.80)	0.011			

^aAbbreviations: HR, hazard ratio; CI, confidence interval