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The prognostic significance of MMP-8 tissue Immunoexpression in pancreatic ductal adenocarcinoma after neoadjuvant therapy

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Neoadjuvant therapy (NAT) has become increasingly common in pancreatic ductal adenocarcinoma (PDAC). Still, PDAC remains one of the deadliest cancers and clinically useful biomarkers are needed. Matrix metalloproteinase 8 (MMP-8) has previously been identified as a potential biomarker for PDAC patients undergoing up-front surgery. We investigated the prognostic significance of MMP-8 in PDAC patients treated with NAT and evaluated the association of MMP-8 expression to treatment response. We studied MMP-8 expression using immunohistochemistry in a tissue microarray with samples from 115 NAT and 144 up-front surgery patients. We examined NAT response from resection specimens by estimating the amount of residual tumour cells. We analysed the association of MMP-8 immunoexpression with survival and treatment response. High MMP-8 immunoexpression associated with better survival among patients with strong NAT response (HR 0.22, CI95% 0.05–0.86, $p=0.030$). This association was not observed among patients with poor NAT response nor in the overall NAT group. Furthermore, MMP-8 expression did not differ significantly between the NAT and up-front surgery groups. In conclusion, the MMP-8 tissue expression after NAT is a protective biomarker in PDAC patients with strong NAT response but fails to associate with favourable prognosis in patients with poor NAT response.

Keywords PDAC, Neoadjuvant therapy, Matrix metalloproteinase 8

Pancreatic ductal adenocarcinoma (PDAC) encompasses more than 90% of pancreatic cancer cases, and is one of the leading causes of cancer-related deaths¹. Poor prognosis characterizes PDAC with a 5-year survival rate of less than 10%. This is partly due to nonspecific symptoms leading to late diagnosis and very few effective systemic treatments. PDAC is typically diagnosed at advanced stages with only 10–15% of patients qualifying for surgical treatment with curative intent² and many patients presenting with borderline resectable disease³. Neoadjuvant therapy (NAT) can downstage more advanced disease and thus enable radical surgery also for borderline resectable PDAC^{4,5}.

The combination of leucovorin, fluorouracil, irinotecan and oxaliplatin (FOLFIRINOX) or gemcitabine are the most commonly used and effective NAT regimens for PDAC^{6,7}. While NAT has been suggested to benefit patients with up-front resectable PDAC⁸ recent results from ongoing clinical trials have shown no survival benefit from neoadjuvant FOLFIRINOX compared to up-front surgery in resectable disease⁹. Furthermore, it has been

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shown that preoperative chemoradiotherapy with gemcitabine did not offer significant survival benefit among resectable PDAC patients¹⁰. Current European recommendation limits NAT to borderline resectable disease, whereas the National Comprehensive Cancer Network[®] (NCCN[®]) guidelines for Pancreatic Adenocarcinoma include NAT as an option for resectable disease^{11,12}. NAT is considered safe, offering a benefit in borderline resectable or locally advanced PDAC, and is associated with decreased disease progression^{13–15}. However, treatment responses to NAT remains generally weak with only a few patients demonstrating strong treatment responses on final pathology¹⁶. Histological evaluation is considered the most accurate method for grading NAT response. Some histological features, such as necrosis, fibrosis, and tumour cell atypia, however overlap with characteristics observed in untreated PDAC thus making therapeutic response evaluation challenging^{17,18}. There is a lack of useful diagnostic, prognostic, treatment predictive and response biomarkers in PDAC. The gold standard biomarker, CA19-9, is commonly used for NAT patients, holding value both as a prognostic marker and, to some extent as a treatment response marker^{19,20}.

Matrix metalloproteinases (MMP) comprise a group of genetically distinct but structurally related proteolytic enzymes with varying roles in both physiological and pathological processes^{21,22}. MMPs not only degrade connective tissue proteins but act as immunomodulators and process various bioactive non-matrix molecules such as complement compounds, growth factors and insulin receptors²³. Matrix metalloproteinase 8 (MMP-8) is a collagenase primarily produced by neutrophils but expressed by various other cells as well. MMP-8 degrades collagens and has an important physiological role in tissue remodelling in injuries and in wound healing^{24,25}.

MMP-8 has been shown to exhibit contrasting effects in cancers depending on the tissue of origin²⁶. MMP-8, along with other MMPs, has been reported to play a role in cancer metastasis and invasion, but in contrast to most other MMPs it has also been shown to inhibit tumour progression^{27–29}. Tumour promoting role of MMP-8 has been thought to associate with degradation of ECM collagens and thus enabling invasion and metastatic growth³⁰. However, it has become more evident that MMP-8, along with other MMPs, influence cell proliferation and angiogenesis of cancer. The paradoxical role of MMP-8 was shown first with mouse models where knock-out of *MMP-8* gene was increasing the incidence of skin tumours and it was associated with sustained inflammatory response³¹. It has been further shown that MMP-8 suppresses metastatic potential in breast cancer and it could be due to MMP-8 mediated induction of cancer cell adhesion³². All in all, mechanisms of MMP-8 in cancer remain complex and still subject to speculation. The prognostic significance of MMP-8 has been under active investigation in various cancers with varying results. For instance, high serum level of MMP-8 is a marker of poor prognosis in colorectal cancer and hepatocellular carcinoma^{33,34}.

In PDAC MMP-8 expression is shown to be increased on tissue level compared to normal pancreatic tissue³⁵. In treatment-naïve PDAC, increased MMP-8 expression has been shown to correlate with better prognosis³⁶. The role of MMP-8 in pancreatic cancer however remains unclear³⁵. The aim of the present study was to investigate whether MMP-8 tumour expression patterns vary between patients that have undergone up-front surgery compared to those treated with NAT, and whether MMP-8 can serve as a marker for NAT response in PDAC. Furthermore, we aimed to evaluate the correlation of MMP-8 expression in relation to strong and weak NAT responses. Lastly, to explore whether MMP-8 expression is associated with the administered neoadjuvant regimen.

Materials and methods

Patient cohort and tissue microarray (TMA)

A total of 259 surgically treated patients with histologically verified PDAC diagnoses were included in this retrospective study. All patients underwent surgery with curative intent between 2000 and 2018. NAT was administered to 115 of the patients, while the remaining 144 up-front operated patients were included as controls. Representable samples of resected tumours were assembled into a tissue microarray (TMA) by drilling 1.0 mm size cores from the paraffin embedded tumour blocks. TMA blocks were assembled using a semiautomatic tissue microarrayer (TMA Grand Master, TMA Control 3.0, 3D Histech, Hungary). For immunohistochemical staining 4 µm thick slides were cut from the TMA blocks. Histological diagnosis and quality of the tissue samples was confirmed by an experienced pathologist (JH) prior the TMA assembly.

Immunohistochemistry

TMAs were immunohistochemically stained with an MMP-8 antibody according to manufacturer's protocol on a LabVision Autostainer 480 S (Thermo Fisher Scientific). The TMA samples underwent deparaffinization and rehydration using Sakura Tissue-Tek DRS. For antigen retrieval, a pretreatment protocol was completed using EnVision Flex target retrieval solution (Pre-Treatment module, Agilent Dako, Santa Clara, CA, USA) for 15 min in 98°. Additionally, a protein blocking protocol was performed to prevent non-specific binding of antibodies. Protein blocking was carried out on the Autostainer following the manufacturers protocol (EnVision FLEX peroxidase-blocking reagent, Agilent, Santa Clara, CA, USA). The primary antibody used was a non-commercial MMP-8 antibody previously described by Hanemaaijer et al.³⁷ (dilution of 1:400). The primary antibody was manually added, and incubation was performed overnight at +4°C. The secondary antibody was applied using Autostainer protocol. Dako REAL Antibody Diluent S2022 (Agilent, Santa Clara, CA, USA) was used as antibody dilution for both primary and secondary antibodies. Tissue slides were washed and rinsed with a buffer solution. Finally, DAB and magenta were added as chromogens, and haematoxylin was used as a counter stain. Adequate staining of the tissue samples was ensured by including a positive control sample of tonsillar hyperplasia or gingival tissue.

Scoring of immunohistochemical staining of MMP-8

We analysed tissue expression of MMP-8 microscopically by analysing the immunopositivity of TMA samples from NAT ($n=113$) and up-front surgery patients ($n=141$). Five of the patients were missing representative

tissue material in TMA and therefore MMP-8 expression could not be evaluated for them. We graded the immunopositivity in tumour tissue to four categories based on staining intensity in cancer cells. Score 0 represented non-detectable expression, while scores 1–3 represented low, moderate, and high intensity, respectively. Four TMA samples per patients were examined and the maximum intensity score served as the representative score. Two independent researchers who were blinded to the disease outcome graded the immunopositivity (JH and MK for up-front surgery patients, JH and EK for NAT patients). Any disagreements in scoring were solved by discussion and re-evaluation of the sample.

Evaluation of the histological response to neoadjuvant treatment

Tissue samples from 113 NAT patients were collected at surgical resection. Histological response could not be evaluated for two of the patients due to missing samples. We used a six-tier scheme to grade the NAT response by evaluating the percentage of remaining, viable tumour cells. The responses were graded as follows; 0, no viable, residual tumour cells (RTCs) (0%); 1, only some found with large magnification of 200–400 \times (\leq 5% RTCs); 2, easily found with large magnification of 200–400 \times (6–10% RTCs); 3, easily found with small magnification of 20–40 \times (11–50% RTCs); 4, minimal NAT effect (51–90% RTCs); and 5, no NAT effect identified (91–100% RTCs). Several representative samples of tumour tissue for each patient were evaluated to grade the NAT response. Samples were evaluated by an experienced gastrointestinal pathologist (AR) along with authors AE and EK. The method has been previously described in detail by Eurola et al.¹⁶ and the scheme relied on the recommendations of the latest tumour response consensus statements from the Amsterdam International Consensus Meeting³⁸.

Statistical analyses

For statistical analyses, MMP-8 scores 0 and 1 were combined into a group representing low immunopositivity, while scores 2 and 3 represented high immunopositivity. Response to NAT was further categorized into two groups where responses 0 to 2 represented a strong response (\leq 10% RTCs), while responses 3 to 5 represented a weak response (\geq 11% RTCs).

Comparisons between categorical variables was performed using chi square test. Disease specific survival (DSS) was analysed using Kaplan-Meier analysis and the log-rank test while Cox regression was used for univariate and multivariable analyses. All the statistical analyses were performed using SPSS statistics software for Mac, version 29 (IBM Corp. Armonk, NY) or R version (R core team). P-values < 0.05 were considered statistically significant.

Ethical approval

This study was granted an approval by the Helsinki University Hospital Ethics committee (07.06.2023; Dnr HUS/1223/2021). The Finnish National Supervisory Authority for Health and Welfare has granted this study an approval to use archived tissue samples without requiring individual consent from the patients (Dnr 10041/05.01.03.01/2012).

Results

MMP-8 immunostaining pattern and patient characteristics

The NAT group had more patients with low MMP-8 immunopositivity than the up-front surgery group ($p = 0.029$) and only a few patients in both groups appeared negative for MMP-8 (Table 1). Different immunostaining patterns of NAT patients are demonstrated in Fig. 1. In the NAT group, low MMP-8 tissue expression associated with receiving preoperative radiation therapy ($p = 0.043$). Other clinical parameters such as age, sex, disease stage, histological grade, perineural invasion, perivascular invasion, NAT regimen and NAT response did not associate significantly with MMP-8 expression (Table 2).

	Up-front surgery (n=141)	NAT (n=113)	p-value
MMP-8 expression			
0 = non-detectable	6 (4.3)	2 (1.8)	
1 = low	22 (15.6)	34 (30.1)	
2 = moderate	65 (46.1)	33 (29.2)	
3 = high	48 (34.0)	44 (38.9)	0.663*
MMP-8 low vs. high			
Low expression	28 (19.9)	36 (31.9)	
High expression	113 (80.1)	77 (68.1)	0.029
MMP-8 negative vs. positive			
Negative expression	6 (4.3)	2 (1.8)	
Positive expression	135 (95.7)	111 (98.2)	0.260

Table 1. Matrix metalloproteinase 8 (MMP-8) expression patterns in neoadjuvant therapy (NAT) and up-front surgery groups. Linear-by-linear test* was used for ordinal variables and Chi square test was applied for categorical variables. P-value < 0.05 was considered significant.

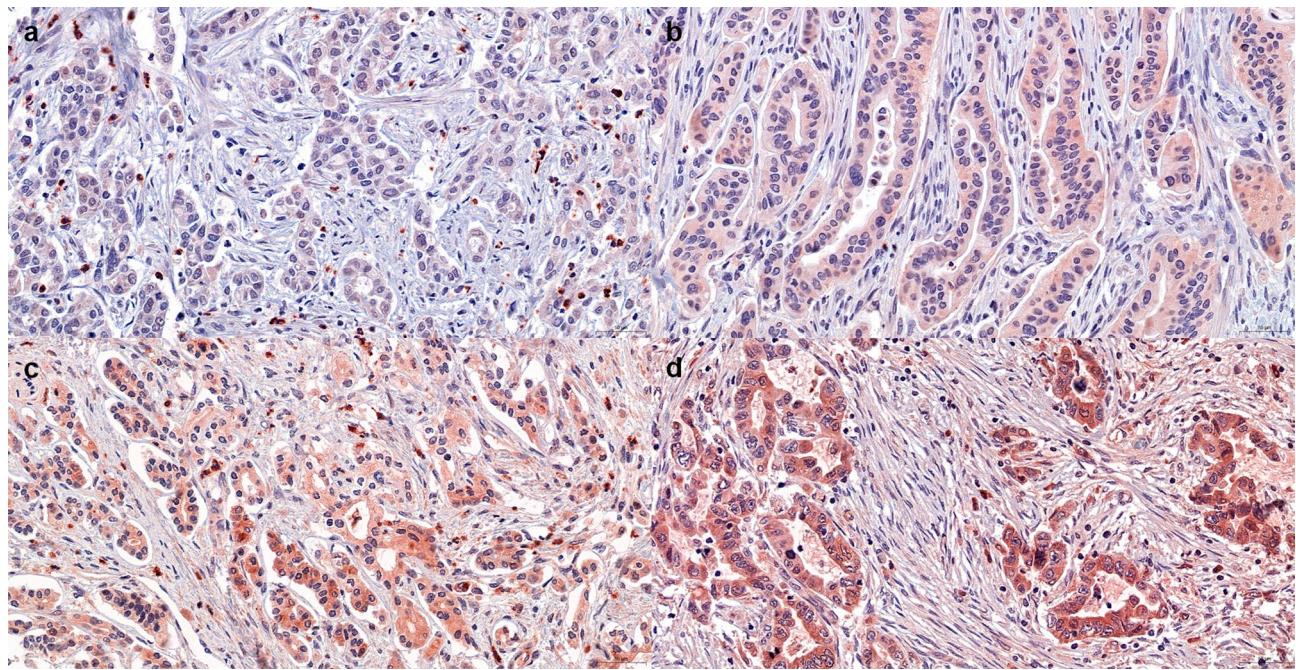


Fig. 1. Immunohistochemical staining patterns demonstrating Matrix metalloproteinase 8 (MMP-8) tissue expression in neoadjuvant therapy (NAT) patients. (A) Non-detectable MMP-8 expression in cytoplasm. (B) Low cytoplasmic immunopositivity. (C) Moderate cytoplasmic immunopositivity. (D) High cytoplasmic immunopositivity. Magnification 30x was used for all images.

Disease specific survival (DSS) and overall survival (OS)

Out of 115 NAT patients, 93 (80.9%) died during follow-up and 88 of them (76.5%) died of PDAC. Thus 94.6% of the deaths were caused by PDAC. In the up-front surgery group, 126 (87.5%) of 144 patients died and of them 120 (83.3%) patients died of PDAC. Consequently, most of the deaths (95.2%) in up-front surgery group were also due to PDAC. The median OS and DSS for NAT patients were 2.57 and 2.24 years, respectively, while for up-front surgery patients, they were 2.20 and 2.26 years. The differences were not statistically significant ($p=0.248$ and 0.211). The median follow-up time was approximately 10 years, and no patients were lost to follow-up.

Neoadjuvant treatment response

Out of 113 NAT patients, 22 (19.5%) had a strong response ($\leq 10\%$ RTCs) to NAT, while 91 (80.5%) patients had a weak NAT response ($\geq 11\%$ RTCs). Only one patient experienced complete pathological response (pCR i.e. no viable RTCs). In univariable Cox regression analysis, a strong NAT response predicted better DSS (HR 0.44, 95% CI 0.24–0.84, $p=0.012$) (Table 3).

MMP-8 expression and survival

We did not find statistically significant difference in DSS between groups with low and high MMP-8 expression among all NAT patients in Kaplan-Meier analyses (Fig. 2A). Univariable Cox regression analyses confirmed that MMP-8 expression was not associated with DSS in the NAT group (HR 0.87, 95% CI 0.56–1.36, $p=0.552$) (Table 3). A similar result was obtained for the up-front surgery group, with survival not being related to MMP-8 expression in Kaplan-Meier analyses (Fig. 3) nor in univariable Cox regression analyses (HR 0.79, 95% CI 0.51–1.23, $p=0.290$, data not shown) when comparing the groups with low and high MMP-8 expression. However, MMP-8 immunopositivity acted as a positive prognostic factor (HR 0.40, 95% CI 0.18–0.92, $p=0.031$) when comparing negative and positive tissue expression in up-front surgery patients (Fig. 3).

As demonstrated in Fig. 2, Kaplan-Meier analyses showed that high MMP-8 tissue expression was associated with better DSS in the subgroup of patients with a strong NAT response, a finding not observed among patients with a weak NAT response. Furthermore, in univariable Cox regression analysis, high MMP-8 expression after NAT indicated better survival in patients with a strong treatment response (HR 0.22, 95% CI 0.05–0.86, $p=0.030$). Among those with a weak NAT response, MMP-8 expression was not related to survival (HR 1.01, 95% CI 0.62–1.67, $p=0.954$) (Table 3).

A multivariable Cox regression analysis for DSS was performed in the entire NAT group including the following parameters: age, sex, stage, grade, preoperative CA19-9 and MMP-8 expression. As demonstrated in Table 4, older age and high preoperative CA19-9 levels were associated with worse survival, but MMP-8 expression was not significantly associated with survival in this model. Due to the small number of patients, multivariable analysis could not be performed for subgroups of strong NAT response ($n=22$) or weak NAT response ($n=91$).

n (%)	Low MMP-8 expression	High MMP-8 expression	p-value
	36 (31.9)	77 (68.1)	
Sex			
Male	15 (41.7)	34 (44.2)	
Female	21 (58.3)	43 (55.8)	0.804
Age			
<65	12 (33.3)	39 (50.6)	
≥65	24 (66.7)	38 (49.4)	0.085
T-stage			
T1	8 (22.2)	18 (23.4)	
T2	19 (52.8)	44 (57.1)	
T3	8 (22.2)	11 (14.3)	
T4	1 (2.8)	4 (5.2)	0.719
N-stage			
N0	13 (36.1)	38 (49.4)	
N1	19 (52.8)	23 (29.9)	
N2	4 (11.1)	16 (20.8)	0.058
Clinical stage			
IA	4 (11.1)	8 (10.4)	
IB	7 (19.4)	23 (29.9)	
IIA	3 (8.3)	5 (6.5)	
IIB	17 (47.2)	22 (28.6)	
III	4 (11.1)	19 (24.7)	
IV	1 (2.8)	0 (0.0)	0.158
Grade			
1	3 (8.3)	10 (13.0)	
2	21 (58.3)	52 (67.5)	
3	7 (19.4)	12 (15.6)	0.681
Missing	8		
Perineural invasion			
Yes	23 (63.8)	54 (70.1)	
No	10 (27.8)	22 (28.6)	0.886
Missing	4		
Perivascular invasion			
Yes	12 (33.3)	24 (31.2)	
No	20 (55.6)	51 (66.2)	0.581
Missing	6		
Preoperative CA19-9			
<37	15 (41.7)	35 (45.5)	
≥37	14 (38.9)	38 (49.4)	
Missing	13		
NAT agent			
Gemcitabine-based	24 (66.7)	57 (74.0)	
FOLFIRINOX	7 (19.4)	17 (22.1)	0.965
Missing	8		
Radiation therapy			
Yes	14 (38.9)	17 (22.1)	
No	20 (55.6)	59 (76.6)	0.043
Missing	3		
NAT response			
Weak	26 (72.2)	65 (84.4)	
Strong	9 (25.0)	11 (14.3)	0.152
Missing	2		

Table 2. Matrix metalloproteinase 8 (MMP-8) and patient characteristics in the neoadjuvant therapy (NAT) group ($n=113$). Chi square test was applied and p-value <0.05 was considered significant.

	All NAT patients			NAT response strong			NAT response weak						
	HR	95% CI	p	HR	95% CI	p	HR	95% CI	p				
Age	1.04	1.01	-1.06	0.005	1.04	0.98	-1.10	0.231	1.03	1.00	-1.06	0.029	
Sex	1.20	0.78	-1.84	0.404	1.45	0.44	-4.84	0.542	1.16	0.73	-1.85	0.518	
Clinical stage (I-IIa vs. IIb-IV)	1.38	0.90	-2.11	0.142	0.59	0.16	-2.23	0.436	1.56	0.97	-2.49	0.065	
Grade	1	Ref.			ref.			Ref.					
	2	0.87	0.47	-1.63	0.663	1.29	0.16	-10.66	0.813	0.88	0.46	-1.70	0.703
	3	0.69	0.32	-1.51	0.354	0.40	0.04	-4.40	0.452	1.05	0.46	-2.39	0.907
Perineural invasion	1.61	0.98	-2.63	0.060	1.10	0.33	-3.61	0.876	1.58	0.90	-2.79	0.114	
Perivascular invasion	1.41	0.90	-2.20	0.138	0.69	0.15	-3.25	0.639	1.56	0.96	-2.52	0.072	
Preoperative CA19-9 (kU/l, log10)	1.82	1.44	-2.30	0.000	1.39	0.69	-2.80	0.362	1.89	1.46	-2.43	0.000	
Gemcitabine vs. FOLFIRINOX	0.64	0.36	-1.12	0.118	1.14	0.24	-5.49	0.873	0.60	0.33	-1.09	0.095	
Preoperative Radiation therapy	1.10	0.69	-1.74	0.695	NA				1.08	0.61	-1.90	0.797	
MMP8 expression High vs. Low	0.87	0.56	-1.36	0.552	0.22	0.05	-0.86	0.030	1.01	0.62	-1.67	0.954	
NAT response Strong vs. Weak	0.44	0.24	-0.84	0.012									

Table 3. Univariable Cox regression analyses of disease specific survival (DSS) in the neoadjuvant therapy (NAT) group and separately in groups of strong or weak response to NAT. p-value 0.05 was considered significant. Statistically significant p-values are bolded in the table.

Treatment response after gemcitabine vs. FOLFIRINOX

Either in combination or as a single agent, gemcitabine was administered to 81 NAT patients, while 24 received the FOLFIRINOX regimen. Information about the specific NAT agent was missing for 8 patients. There was no significant difference in NAT response when comparing gemcitabine with FOLFIRINOX. In the gemcitabine group 16 (19.5%) patients and in FOLFIRINOX group 4 (17.4%) patients experienced a strong NAT response ($p=0.819$). When analysing DSS in subgroups based on NAT agent administered, a strong NAT response predicted better survival in the gemcitabine group while in the FOLFIRINOX group there was no difference in survival when comparing strong or weak NAT response (Fig. 4). Furthermore, patients in FOLFIRINOX group and gemcitabine group expressed MMP-8 similarly in tumour tissue. In the gemcitabine group 57 (70.4%) patients and in FOLFIRINOX group 17 (70.8%) patients expressed high MMP-8 immunopositivity.

Discussion

The aim in this study was to evaluate the prognostic value of MMP-8 tissue expression in PDAC patients receiving NAT prior to surgery. While we hypothesized that higher MMP-8 expression in tumour tissue could predict better prognosis in NAT patients, contradicting results were revealed. We found that the role of MMP-8 varied in the subgroups of NAT-patients as MMP-8 predicted better prognosis in the strong response group but not in the weak response group. In the DSS analysis of all NAT patients, no difference in survival was observed when comparing high or low MMP-8 tissue expression.

In the subgroup of NAT patients with a strong response, high MMP-8 tissue expression predicted better survival. Furthermore, in the group with weak NAT response this protecting effect of MMP-8 expression was not observed. One can speculate, that among those non-responding or only weakly responding to NAT in PDAC, MMP-8 may be either inactive or has adverse effect on survival. Previous studies indicate that in NAT patients, the survival tissue marker profile changes along with NAT response, whereas markers of poor survival may be more pronounced in patients with weak response¹⁶. Moreover it has been previously shown that administration of NAT alters immune microenvironment of the tumour possibly towards more pro-inflammatory profile^{39,40}. This altered inflammatory response could very well affect tumour MMP-8 expression and one can speculate that this is partly reason for better prognosis. However, MMP-8 expression did not differ between the groups with strong or weak NAT response.

In contrast to our previous findings MMP-8 did not act as a prognostic factor in up-front operated patients when analysing DSS in the present analysis. This could be due to differences in grouping of the MMP-8 expression as we here compared low and high expression, while previously there was a comparison between non-detectable and positive MMP-8 expression. Furthermore, we used the strongest tissue immunoexpression score in statistical analyses while previously the median score was used³⁶. Nevertheless, in our additional

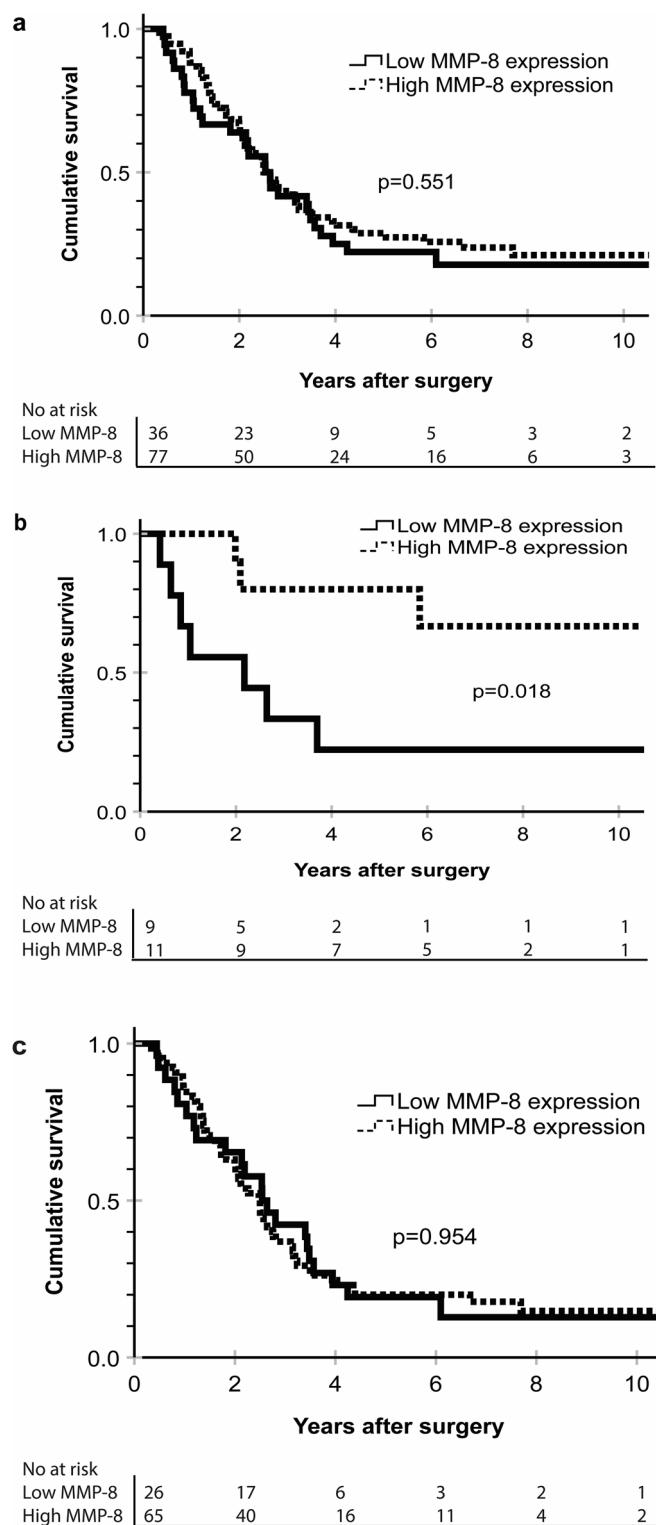


Fig. 2. Kaplan-Meier analyses of disease-specific survival (DSS) in relation to matrix metalloproteinase 8 (MMP-8) tissue expression. (A) Kaplan-Meier analysis showing no correlation between MMP-8 tissue expression and DSS in the whole patient group treated with neoadjuvant therapy (NAT). (B) In the subgroup of patients with strong response to NAT high MMP-8 expression was related to better DSS. (C) MMP-8 expression was not associated with DSS among patients with weak NAT response. Log Rank test was applied and $p < 0.05$ was considered statistically significant.

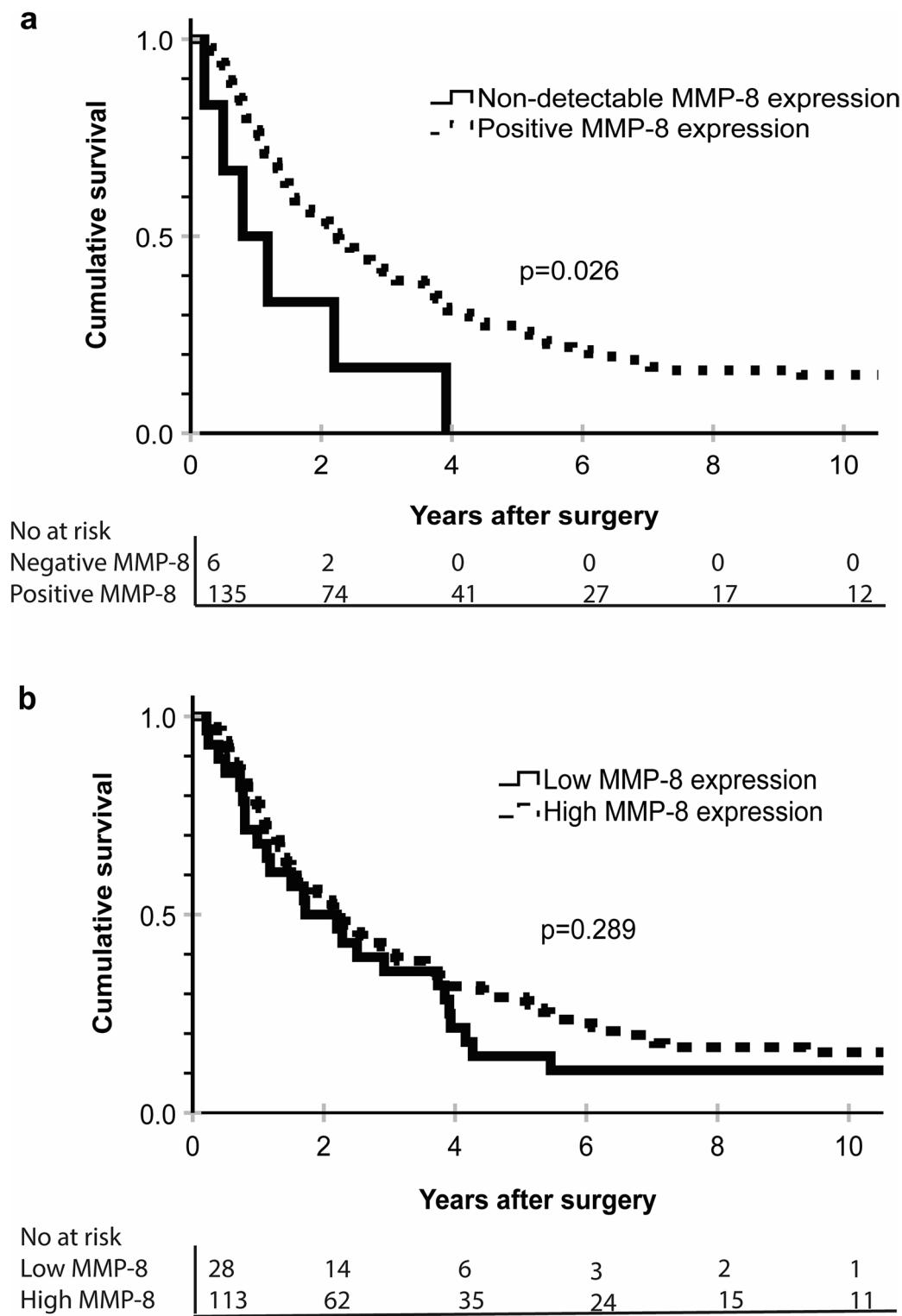


Fig. 3. Kaplan-Meier survival analyses in up-front surgery patients. (A) Immunopositivity of matrix metalloproteinase 8 (MMP-8) predicts better disease-specific survival (DSS) in up-front surgery patients. (B) No correlation was detected between DSS and MMP-8 expression when comparing low and high tissue expression. Log Rank test was applied and $p < 0.05$ was considered statistically significant.

analysis we found that MMP-8 acted as positive prognostic factor also in this patient cohort when comparing non-detectable and positive MMP-8 expression in the up-front surgery group (Fig. 3). However, we noted that very few of the samples had non-detectable MMP-8 expression and this could thus be a source of skewness in statistical analyses.

		HR	95% CI		<i>p</i>
Age		1.04	1.01	-1.07	0.002
Sex		1.06	0.63	-1.78	0.829
Clinical stage (I-IIa vs. IIb-IV)		0.95	0.57	-1.57	0.828
Grade	1	Ref.			
	2	0.78	0.37	-1.66	0.526
	3	1.02	0.42	-2.49	0.970
Preoperative CA19-9 (kU/l, log10)		1.91	1.49	-2.47	0.000
MMP8 expression High vs. Low		0.64	0.37	-1.10	0.106

Table 4. Multivariable Cox regression analysis of disease specific survival (DSS) in neoadjuvant therapy (NAT) group. *p*-value 0.05 was considered significant. Statistically significant *p*-values are bolded in the table.

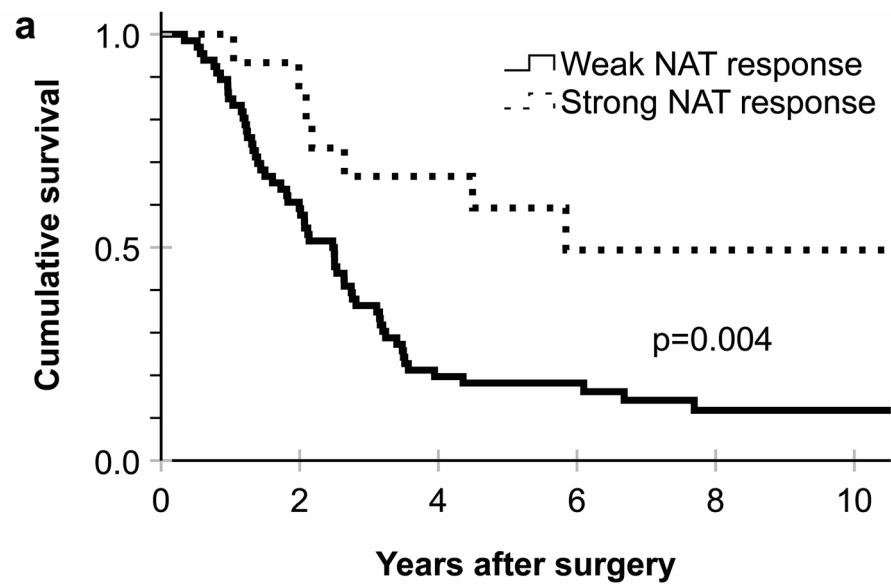
A majority (81%) of NAT patients achieved only a weak treatment response with only one patient out of 113 having a complete pathological response. These results align with previous findings that a strong NAT response in PDAC remains rare¹⁶. In the present cohort, a finding of less than 10% remaining viable tumour cells after NAT is required for a clear survival benefit, thus leaving most of the NAT patients without a statistically significant survival benefit. Some previous studies suggest an even lower, 5% limit of remaining tumour cells for achieving survival benefit⁴¹. Additionally, in our analysis we found that strong NAT response offers survival benefit in patients treated with gemcitabine but not in FOLFIRINOX group. While this finding is interesting, it is likely not clinically true due to small number of patients in the FOLFIRINOX group.

One strength of our study lies in the large and well-described patient cohort. We excluded other tumours of the pancreas or cancers originating from the bile ducts. TMA samples were evaluated by at least two independent investigators. For NAT response analysis the study scheme created for the purpose was previously tested¹⁶.

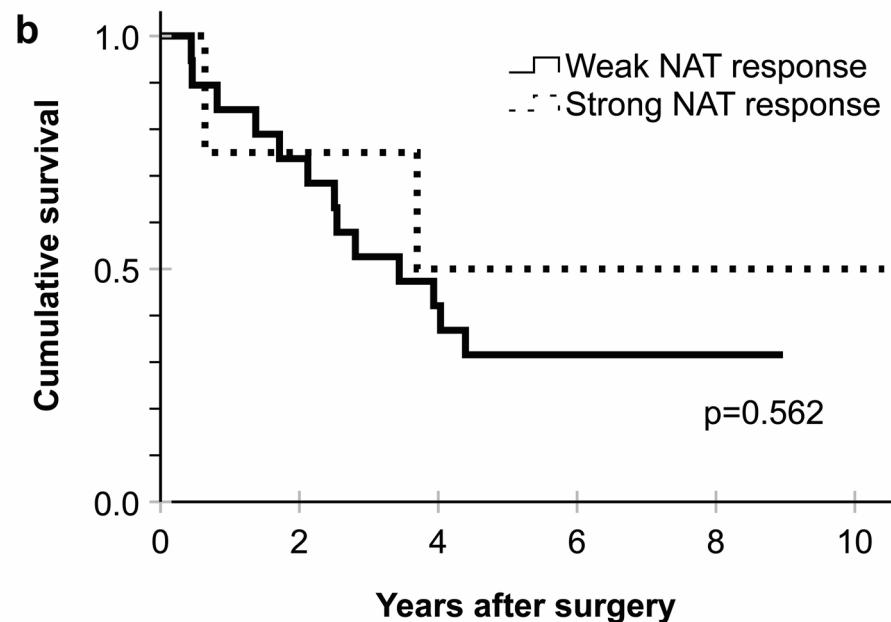
TMAs enable relatively fast microscopic evaluation of multiple tumours, but at the same time limits the visibility of the whole tumour. This was controlled by taking multiple (4) samples of each tumour for TMA. Each TMA sample was evaluated by at least two independent investigators to minimize the impact of subjectivity. NAT-response was determined by an experienced gastrointestinal pathologist. Evaluation of response may be challenging due to some response effects mimicking desmoplasia seen in PDAC. This study featured a limited number of patients in some subgroups, restricting possibilities for conducting multivariable analysis in such settings.

Conclusion

To conclude, the immunoexpression of MMP-8 did not differ significantly between NAT and US patients. Moreover MMP-8 immunoexpression did not correlate with histological response to NAT. However, to our knowledge this is the first study to report MMP-8 as a protective prognostic marker in NAT treated PDAC patients who received a strong treatment response. No prognostic value of MMP-8 was found in patients with a weak NAT response. The underlying reasons for these differences remain unclear and demand further investigation.



No at risk							
Weak response	Strong response	66	39	13	9	5	3
16	13	9	5	3	2		



No at risk							
Weak response	Strong response	19	14	8	5	1	0
4	3	2	2	1	1	1	1

Fig. 4. Kaplan-Meier survival analyses in neoadjuvant therapy (NAT) patients treated with gemcitabine or FOLFIRINOX. (A) Strong NAT response predicted better disease specific survival (DSS) in patients treated with gemcitabine. (B) NAT response did not correlate with DSS in patients treated with FOLFIRINOX regimen. NAT response was considered strong when histological evaluation revealed less than 10% residual tumor cells (RTC). Respectively NAT response was considered weak when more than 10% RTCs. Log Rank test was applied and $p < 0.05$ was considered statistically significant.

Data availability

Data supporting our findings is available on a reasonable request from the corresponding author.

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Author contributions

Data curation, E.K., A.E. and M.K.; Formal analysis, E.K., A.E. and H.M.; Visualization, E.K.; Writing – original draft, E.K. and A.E.; Methodology, A.E. and A.R.; Investigation, J.H. and A.R.; Resources, J.H., T.S. and C.H.; Writing – review and editing, M.K., H.M., J.H., A.R., T.S., H.S., C.H. and M.S.; Conceptualization, H.S., C.H. and M.S.; Supervision, H.S., C.H. and M.S.; Project administration, M.S.

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Declarations

Competing interests

The authors declare no competing interests.

Additional information

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