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# Next-generation sequencing and histological response assessment in peritoneal metastasis from pancreatic cancer treated with PIPAC

Malene Nielsen,<sup>1,2</sup> Martin Graversen ,<sup>3</sup> Signe Bremholm Ellebæk,<sup>3</sup> Thomas Kielsgaard Kristensen,<sup>1</sup> Claus Fristrup,<sup>3</sup> Per Pfeiffer,<sup>2,4</sup> Michael Bau Mortensen,<sup>2,3</sup> Sönke Detlefsen <sup>1,2</sup>

<sup>1</sup>Department of Pathology, Odense Pancreas Center (OPAC) and Odense PIPAC Center (OPC), Odense University Hospital, Odense, Denmark

<sup>2</sup>Department of Clinical Research, Faculty of Health Sciences, University of Southern Denmark, Odense, Denmark

<sup>3</sup>Department of Surgery, Odense Pancreas Center (OPAC) and Odense PIPAC Center (OPC), Odense University Hospital, Odense, Denmark

<sup>4</sup>Department of Oncology, Odense Pancreas Center (OPAC) and Odense PIPAC Center (OPC), Odense University Hospital, Odense, Denmark

## Correspondence to

Dr Sönke Detlefsen, Department of Pathology, Odense University Hospital, Odense 5000, Denmark; sonke.detlefsen@rsyd.dk

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## ABSTRACT

**Background** Peritoneal metastasis from pancreatic cancer (PM-PC) may be treated with repeated pressurised intraperitoneal aerosol chemotherapy (PIPAC). Utility of next-generation sequencing (NGS) to detect cancer-related mutations in peritoneal quadrant biopsies (QBs) and peritoneal fluid (PF) after systemic and PIPAC treatment has not been evaluated. Around 90% of pancreatic cancers (PCs) harbour a *KRAS* mutation, making PC ideal for the evaluation of this aspect.

**Aims** Evaluation of PM-PC in terms of (1) histological response to PIPAC using Peritoneal Regression Grading Score (PRGS), (2) clinical characteristics and (3) frequency of mutations in QBs and PF before and after PIPAC.

**Methods** Peritoneal QBs and PF were obtained prior to each PIPAC. NGS for 22 cancer-related genes was performed on primary tumours, QBs and PFs. Response was assessed by the four-tiered PRGS.

**Results** Sixteen patients treated with a median of three PIPAC procedures were included. The mean PRGS was reduced from 1.91 to 1.58 ( $p=0.02$ ). Fifty-seven specimens (13 primary tumours, 2 metastatic lymph nodes, 16 PFs and 26 QB sets) were analysed with NGS. *KRAS* mutation was found in 14/16 patients (87.50%) and in QBs, primary tumours and PF in 8/12 (66.67%), 8/13 (61.53%) and 6/9 (66.67%). The median overall survival was 9.9 months (SE 1.5, 95% CI 4.9 to 13.9).

**Conclusion** PIPAC induces histological response in the majority of patients with PM-PC. *KRAS* mutation can be found in PM-PC after PIPAC at a frequency similar to the primaries. NGS may be used to detect predictive mutations in PM-PC of various origins, also when only post-PIPAC QBs or PFs are available.

## INTRODUCTION

The incidence of pancreatic cancer (PC) is rising, but the prognosis for the majority of these patients remains poor, especially when the disease has metastasised to the peritoneum. Even in selected patients with peritoneal metastasis from pancreatic cancer (PM-PC), the median survival after systemic combination chemotherapy leads to a median overall survival of only 7–8 months.<sup>1</sup> Pressurised intraperitoneal aerosol chemotherapy (PIPAC) is a treatment alternative to systemic chemotherapy, where studies in patients with peritoneal metastasis (PM) from ovarian, gastric and colorectal cancers and PC show

a high safety profile and promising survival data.<sup>2–6</sup>

Currently, PIPAC is an experimental treatment, and randomised controlled trials are lacking.

The choice of drugs used for PIPAC has depended more on PIPAC tradition and safety rather than disease profile. However, based on the preliminary but promising results of PIPAC-directed treatment in PM-PC, the next obvious step is to look at individual patients with PC that may harbour rare molecular alterations that may be targeted by specific agents.<sup>7,8</sup> The mutational profile of PM of various origin before and after PIPAC may provide important prognostic and predictive information. Studies have shown that chemotherapy treatment in patients with, for example, acute myeloid leukaemia could ‘breed’ new mutations in different genes, so-called subclones.<sup>9,10</sup> It is unknown whether PIPAC can have similar effects regarding the mutational profile of PM. Knowledge regarding if and to what extent there is discordance between the molecular alterations in the primary PC tumour and its PM, and particularly between PM before and after PIPAC, is sparse.

It remains to be elucidated whether treatment may influence the utility of biopsy specimens from the peritoneum for next-generation DNA sequencing (NGS) analyses. As more than 90% of PCs harbour a *KRAS* mutation, this type of cancer is ideal for the evaluation of the latter aspect. The aims of the present study were to evaluate a series of patients with PM-PC treated with PIPAC regarding the following aspects: (1) evaluation of the histological response of PM-PC to PIPAC by means of the Peritoneal Regression Grading Score (PRGS), (2) description of the clinical characteristics and (3) the frequency of mutations in cancer-related genes in peritoneal quadrant biopsies (QBs) and peritoneal fluid (PF) with PM-PC before and after PIPAC.

## MATERIALS AND METHODS

### Clinical data and inclusion of specimens

The patients with PM-PC included in this study are also included in the PIPAC-OPC-1 or PIPAC-OPC-2 studies at Odense PIPAC Center (OPC), Odense University Hospital, Denmark.<sup>11,12</sup> We ensured that patients had not advocated against the use of their tissue in the Danish registry for the use of tissue in research (‘Vævsanvendelsesregisteret’). Patients with PM-PC were included if they had



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biopsy-proven malignancy, clinically proven PM and a maximum of one extraperitoneal metastasis. Five of our 16 patients were included in a previous publication.<sup>13</sup>

### Application of PIPAC and sampling of PF and histological peritoneal biopsies

Using a standard laparoscopic two-trocar approach, a pressure injector and a nebuliser, aerosolised chemotherapeutics (cisplatin and doxorubicin) were distributed within the peritoneal cavity, leading to a deeper and more uniform penetration into the peritoneum compared with traditional chemotherapy.<sup>4 14</sup> The PIPAC procedure was repeated every 4–6 weeks. Prior to each treatment, saline was irrigated into the peritoneal cavity and 150 mL of peritoneal lavage fluid (PF) was collected for further analyses, while the remaining PF was disposed. Quadrant biopsies (QBs) were taken from the peritoneum for assessment of histological response to treatment. The PFs were processed as follows: if a spontaneous coagulum was present, it was fixed in formalin and embedded in paraffin (formalin-fixed paraffin-embedded (FFPE)). Three tubes were filled with 50 mL and centrifuged. From the sediment of the first tube, two smears were produced and stained with Papanicolaou and May-Giemsa Grünwald. From the sediment of the second tube, a cell block was prepared and FFPE. From each of the two FFPE blocks, a thick section of 4–5 µm was cut with a microtome and stained with H&E for microscopic analysis. The sediment of the third tube was stored at –80°C in MagNa Pure LC Lysis Buffer (Roche) for subsequent NGS analysis. For cytological evaluation, a five-tiered score was used: malignant cells, suspicious cells, atypical cells, no malignant cells and other.

### Histology and immunochemistry

The histological QBs were FFPE. Four-micron sections were cut and mounted on FLEX IHC Microscope slides. Three step sections were cut and stained with H&E, followed by a section immunostained for epithelial cell adhesion molecule and a final series of three H&E-stained step sections.<sup>13 15</sup> Sections were dried at room temperature and baked at 60°C for 60 min before immunostaining, which was automated at the Dako Omnis immunostainer using the EnVision FLEX+DAB detection with mouse linker (Dako/Agilent, Glostrup, Denmark). The primary antibody was the clone BS14-Epithelial Specific Antigen/ CD326 (code: BSH-7402–1) (Nordic Biosite, Copenhagen, Denmark). Dilution was 1:600. For retrieval, target retrieval solution buffer with pH 9.0 was used for 30 min at 97°C. Incubation at 32°C was done for 20 min. Nuclear counterstaining was performed using Hematoxylin FLEX at the Dako Omnis platform. Slides were washed, dehydrated and cover slipped using an automated Dako cover slipper (Dako/Agilent, Glostrup, Denmark).

### Peritoneal Regression Grading Score

The four-tiered PRGS for the histological assessment of response to therapy in PM was assessed for each quadrant biopsy and as a mean score for all quadrant biopsies obtained prior to a given PIPAC treatment.<sup>15</sup> The scoring was performed by the same pathologist with special interest for peritoneal pathology (SD) in all cases. It was recently found that the interobserver variability of the PRGS is moderate to good.<sup>16</sup> The PRGS defines four categories, based on the presence of residual tumour cells and the extent of regressive features. Major histological features of regression are fibrosis, inflammation, hyalinosi, acellular mucin pools, ischaemic necrosis, accumulation of macrophages, multinucleated giant cells and granulomas. PRGS 1 corresponds to a

complete regression with absence of tumour cells; PRGS 2 corresponds to a major histological response with regressive features predominant over residual tumour cells; PRGS 3 corresponds to a minor histological response with predominance of residual tumour cells over regressive features; and PRGS 4 corresponds to a lack of histological response to therapy where the tumour cells were not accompanied by any regressive features.<sup>15</sup>

### DNA extraction

For genomic DNA purification from FFPE tissue, GeneRead DNA FFPE Kit (QIAGEN, Hilden, Germany) was used according to the manufacturer's recommendations. When possible, macrodissection was performed to increase the relative amount of tumour DNA, aiming at 20% or more. The GeneRead DNA FFPE procedure was used to remove paraffin and reverses formalin cross-links from the DNA before it was bound to QIAamp MinElute column (QIAGEN). After heating to remove cross-links, the DNA was accessible for removal of deaminated cytosine residues. After binding of the DNA to the spin column, residual contaminants such as salts were washed away with two wash buffers and ethanol. Any residual ethanol was removed by an additional centrifugation step. Finally, DNA was eluted. DNA extraction from PFs was performed using the MagNa pure LC Instrument (Roche Applied Science, Mannheim, Germany). DNA was extracted in a volume of 100 µL of elution buffer using the MagNa Pure LC DNA Isolation Kit I (Roche Applied Science).

### Specimens for NGS

The inclusion criteria for the specimens used for NGS analyses were availability of cancer tissue in biopsy or resection specimen. For histological PM biopsy before and after PIPAC, only QB sets containing cancer cells in at least one of the QBs were included, regardless of the amount of cancer cells. The QB with the highest amount of tumour tissue was chosen. NGS was performed on all PFs with either 'malignant tumour cells' or 'cells suspicious of malignancy'.

### Next-generation DNA sequencing (NGS)

NGS was performed on the Ion Torrent PGM (Life Technologies, Carlsbad, California, USA) instrument, using the Ion AmpliSeq Colon and Lung Cancer NGS Panel v2, containing hotspot regions in 22 target genes: *ALK*, *AKT1*, *BRAF*, *CTNNB1*, *DDR2*, *EGFR*, *ERBB2*, *ERBB4*, *FBXW7*, *FGFR1*, *FGFR2*, *FGFR3*, *KRAS*, *MAP2K1*, *MET*, *NRAS*, *NOTCH1*, *PIK3CA*, *PTEN*, *SMAD4*, *STK11* and *TP53*. Ion Reporter software was used for variant calling. All variants were visualised using Golden Helix Genome-Browse V3.0.0 software (GoldenHelix, Montana, USA). A wild-type control was included in each run. A 1% cut-off was used for known *KRAS* hotspot mutations and a 10% cut-off was used for other variants.<sup>17 18</sup> Only variants with coverage of >2000 were included. Samples with a DNA input of <0.1 ng/µL were discarded.

### Statistics

Values were given as means or percentages. Comparisons were performed using Pearson's  $\chi^2$  test for categorical data and Wilcoxon matched pairs signed-rank test was used where appropriate. P values were two-tailed, and a p value of 0.05 was considered statistically significant. The statistical software STATA V.15.0 was used for statistical analysis.

**Table 1** Baseline characteristics of patients with peritoneal metastasis from pancreatic cancer treated with PIPAC

Demographic variables	Value
Number of patients	16
Age (years), mean (range)	59 (46–72)
Sex, male/female	10/6
ECOG performance status 0, number of patients	5
ECOG performance status 1, number of patients	11
Number of PIPAC procedures, median (range)	3 (1–6)
Previous treatment	
One-line palliative SC, number of patients	12
Two-line palliative SC, number of patients	4

ECOG, Eastern Cooperative Oncology Group; PIPAC, Pressurised IntraPeritoneal Aerosol Chemotherapy; SC, systemic chemotherapy.

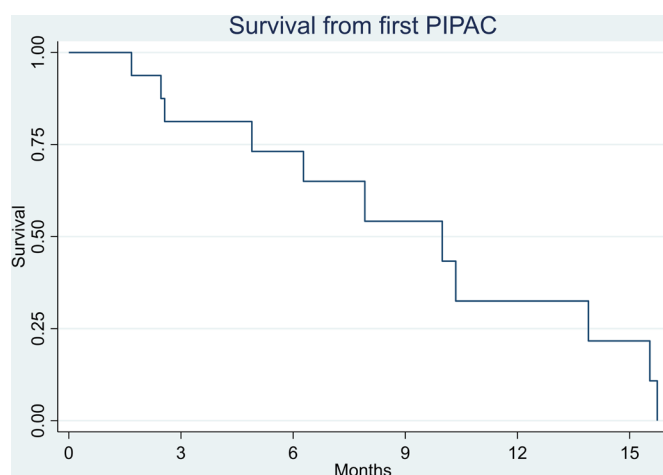
## RESULTS

### Clinical data

Sixteen patients with PM-PC were treated with at least one PIPAC at OPC from January 2016 to October 2019. The median number of PIPAC treatments was 3 (range 1–6). All patients underwent palliative systemic chemotherapy, and six patients underwent synchronous chemotherapy with gemcitabine+S1, gemcitabine+nab-paclitaxel or Folfirinox (folinic acid, fluorouracil, irinotecan and oxaliplatin) (table 1). The median overall survival from the first PIPAC was 9.9 months (SE 1.5, 95% CI 4.9 to 13.9). Kaplan-Meier survival analysis is shown in figure 1.

### Histological response assessment using the PRGS

A total of 44 QB sets were collected, consisting of 111 QBs. Sixteen, 13, 6, 5, 3 and 1 QB sets were taken prior to PIPAC treatments 1, 2, 3, 4, 5 and 6. The remaining biopsies were from the epigastric peritoneum (n=4), the liver capsule (n=1), the right flank (n=3), the falciform ligament and the peritoneum without known precise location (each n=1). Thirteen patients underwent at least two PIPACs and were eligible for response assessment using the PRGS. Seven patients underwent two PIPAC treatments, while six patients underwent three PIPAC treatments or more. Based on the PRGS prior to PIPAC treatment 1 and



**Figure 1** Kaplan-Meier survival analysis of 16 patients with peritoneal metastasis from pancreatic cancer treated with PIPAC. Survival is shown as the time period from the first PIPAC treatment. The median overall survival from first PIPAC was 9.9 months (SE 1.5, 95% CI 4.9 to 13.9). PIPAC, pressurised IntraPeritoneal Aerosol Chemotherapy intraperitoneal aerosol chemotherapy.

PIPAC treatment 2 or 3, 61.50% of patients had a reduction of the mean PRGS. The overall mean PRGS was reduced from 1.91 at baseline to 1.58 prior to PIPAC 2 or 3 ( $p=0.02$ ). The overall findings regarding PRGS are shown in tables 2 and 3.

### Next-generation sequencing

NGS analysis was performed on 57 specimens (13 primary tumours, 2 metastatic lymph nodes, 16 PFs and 26 QBs) (table 4). A mutation was detected in 15 of 16 patients (93.75%), and a *KRAS* mutation (p.Gly12Asp, p.Gly12Val, p.Gly12Ala or p.Gln61Arg) in 14 of 16 patients (87.50%). A *KRAS* mutation was found in QBs, primary tumours and PFs in 8/12 (66.67%), 8/13 (61.53%) and 6/9 (66.67%) patients. A *KRAS* mutation was found in the primary tumour in 8/13 patients (61.53%) and in at least one of the metastatic biopsies/PFs in 10/13 patients (76.92%). A *KRAS* mutation was found in QBs prior to PIPAC 1, in PFs prior to PIPAC 1, QBs after PIPAC and PFs after PIPAC in 8/11 (61.54%), 4/6 (66.67%), 5/8 (62.50%) and 4/7 (57.14%) patients. A *KRAS* mutation was found in both the primary tumour and at least one metastasis in 4/11 patients (36.36%). In three patients, a *KRAS* mutation was detected in a metastasis but not in the primary tumour. In one patient, a *KRAS* mutation was detected in the primary tumour but not in the metastases.

We also found mutations in *TP53* in the primary tumour in two patients (patient 2 and patient 14), a mutation in *MET* in both primary tumour and two metastases (patient 9), a *SMAD4* mutation in the primary tumour and PM (patient 5), a *SMAD4* mutation in QB (patient 10) and a *FGFR2* mutation in the primary tumour (patient 11).

## DISCUSSION

To our knowledge, this is the first study evaluating the utility of NGS for detection of mutations in cancer-related genes in QBs and PFs from PM treated with PIPAC. PIPAC induced a histological response, according to the mean PRGS, in 8/13 patients (61.50%), while 4/13 (30.77%) had stable disease. Using NGS, we found the *KRAS* mutation in peritoneal QBs and PF, even after PIPAC treatment, at a frequency similar to that in the primary tumour biopsies.

A histological tumour response to PIPAC treatment measured by PRGS has been observed in 67% of patients with PM of various origin in a previous study.<sup>13</sup> Our study cohort was more homogenous, and therefore, only patients with PC were included.<sup>2 13 19</sup> In another study on PIPAC treatment of 20 patients with PM-PC, 10 of whom underwent at least two PIPACs or more, a histological tumour response was observed in 35%.<sup>20</sup> PRGS was not used, but instead, tumour regression grade. Horvath *et al* reported a histological regression in 40% of their 12 PIPAC-treated patients with PM from pancreatobiliary cancer (PC (n=6) or cholangiocarcinoma (n=6)), using the PRGS.<sup>21</sup> The histological response was evaluated in 6 of the 12 included patients who had undergone two PIPACs or more. Furthermore, Graversen *et al* demonstrated histological regression, measured by PRGS, in four out of five patients with PM-PC (80%), while one patient had stable disease.<sup>13</sup> Hence, the present study is the to date the largest study providing histological response evaluation in patients with PM-PC treated with PIPAC (n=13). The median survival from first PIPAC that we found, 9.9 months (95% CI 4.9 to 13.9), is similar to the median survival reported by Khosrawipour *et al* and Horvath *et al*.<sup>20 21</sup>

We detected a *KRAS* mutation and other mutations in QBs and PF before and after PIPAC treatment. The *KRAS* gene is mutated in 88% to 95% of patients with PC.<sup>17 18 22</sup> These previous studies



**Table 2** PRGS ranging from 1 to 4 in 16 patients with peritoneal metastasis from pancreatic cancer who underwent at least one PIPAC treatment

	Patient 1	Patient 2	Patient 3	Patient 4	Patient 5	Patient 6
PRGS before PIPAC 1 (highest/mean)	3/2.25	3/2	3/2.25	4/2.75	3/3	1/1
PRGS before PIPAC 2 (highest/mean)	2/1.25	2/2	3/1.75	3/2.25	2/2	
Histological regression	+	U	+	+	+	
PRGS before PIPAC 3 (highest/mean)	2/1.25	3/1.75	2/1.5		3/2	
Histological regression	U	+	+		–	
PRGS before PIPAC 4 (highest/mean)	2/1.75		3/2		3/2.67	
Histological regression	–		–		–	
PRGS before PIPAC 5 (highest/mean)			3/2.5		3/2.33	
Histological regression			–		+	
	Patient 7	Patient 8	Patient 9	Patient 10	Patient 11	Patient 12
PRGS before PIPAC 1 (highest/mean)	1/1	3/3	1/1	3/2.25	2/1.33	1/1
PRGS before PIPAC 2 (highest/mean)	1/1			3/2	1/1	2/1.5
Histological response	U			+	+	–
PRGS before PIPAC 3 (highest/mean)					1/1	1/1
Histological response					U	+
PRGS before PIPAC 4 (highest/mean)					1/1	3/2.66
Histological response					U	–
PRGS before PIPAC 5 (highest/mean)					1/1	
Histological response					U	
PRGS before PIPAC 6 (highest/mean)					1/1	
Histological response					U	
	Patient 13	Patient 14	Patient 15	Patient 16		
PRGS before PIPAC 1 (highest/mean)	1/1	1/1	3/2	3/3		
PRGS before PIPAC 2 (highest/mean)	1/1	1/1	3/2.33	2/2		
Histological response	U	U	–	+		

– indicates histological progression (according to mean PRGS); + indicates histological regression (according to mean PRGS).

PIPAC, Pressurised IntraPeritoneal Aerosol Chemotherapy; PRGS, Peritoneal Regression Grading Score; U, unchanged mean Peritoneal Regression Grading Score.

were mainly based on snap-frozen tissue from pancreatic surgery or FFPE tissue from surgical specimens from chemotherapy-naïve patients, meaning that much more tumour tissue was available than in our QB and PF specimens before and after systemic chemotherapy and PIPAC. In many of our specimens, regressive changes and only few cancer cells were present. In PF specimens before and after PIPAC, we found a mutation in around 60% of the patients. This indicates that NGS for cancer-related genes has a lower sensitivity compared with PCR for messenger RNA (mRNA) of tumour markers in this setting. Graversen *et al* reported a sensitivity of 0.88 and a specificity of 1.00 in patients with PM of various origins, including PC, when performing PCR for a combination of carcinoembryonic antigen (CEA)/EpCAM mRNA on PF specimens.<sup>19</sup> In patient 9, no *KRAS* mutation but

instead a *MET* mutation was found. A previous study has shown that targeting *MET* is possible and can be used for treatments of various cancers.<sup>23</sup> Future studies have to elucidate whether treatment with, for example, crizotinib, targeting *MET*, may be of value in some patients with PM-PC.<sup>24</sup> It has to be emphasised that our data do not support the routine use of NGS in PM treated with PIPAC. However, NGS using post-PIPAC peritoneal biopsies could be considered in the following setting: (1) the NGS panel includes predictive markers (eg, in patients with PM from colorectal cancer, where the *KRAS* mutation has predictive value for treatment with EGF-R antagonists); (2) cancer cells are present in the biopsy (but some histological regression (PRGS 2 or 3) may be present); and (3) an adequate specimen from the primary tumour or from a pre-PIPAC biopsy is not available.

**Table 3** PRGS ranging from 1 to 4 in 13 patients with peritoneal metastasis from pancreatic cancer who underwent at least two PIPAC treatments

	Patient 1	Patient 2	Patient 3	Patient 4*	Patient 5	Patient 7*	Patient 10*	Patient 11	Patient 12
PRGS before PIPAC 1 (highest/mean)	3/2.25	3/2	3/2.25	4/2.75	3/3	1/1	3/2.25	2/1.33	1/1
PRGS before PIPAC 3 (or 2*) (highest/mean)	2/1.75	3/1.75	3/1.5	3/2.25	3/2	1/1	3/2	1/1	1/1
Histological response	+	+	+	+	+	U	+	+	U
	Patient 13*	Patient 14*	Patient 15*	Patient 16*					
PRGS before PIPAC 1 (highest/mean)	1/1	1/1	3/2	3/3					
PRGS before PIPAC 3 (or 2*) (highest/mean)	1/1	1/1	3/2.33	2/2					
Histological response	U	U	–	+					

– indicates histological progression (according to mean PRGS); + indicates histological regression (according to mean PRGS).

\*Did only undergo PIPAC twice, data from PIPAC 2 instead of PIPAC 3.

PIPAC, Pressurised IntraPeritoneal Aerosol Chemotherapy; PRGS, Peritoneal Regression Grading Score; U, unchanged mean PRGS.

**Table 4** Results of NGS analyses of cytological and histological specimens from primary tumour and metastases from 16 patients with PM from pancreatic cancer

	Primary tumour	Histological biopsy prior to PIPAC*	PF before PIPAC	Histological peritoneal biopsy after PIPAC	PF after PIPAC
Patient 1	<i>KRAS</i> (p.Gly12Asp)	<i>KRAS</i> (p.Gly12Asp)	<i>KRAS</i> (p.Gly12Asp)	ND (PIPAC 3 and 4)	NA
Patient 2	<i>KRAS</i> (p.Gly12Val) <i>TP53</i> (p.Cys275Tyr)	<i>KRAS</i> (p.Gly12Val)	NA	<i>KRAS</i> (p.Gly12Val) (PIPAC 2 and 3)	NA
Patient 3	ND	ND	NA	ND (PIPAC 2, 3, 4 and 5)	ND (PIPAC 5)
Patient 4	ND	<i>KRAS</i> (p.Gly12Val)	ND	<i>KRAS</i> (p.Gly12Val) (PIPAC2)	<i>KRAS</i> (p.Gly12Val) (PIPAC 2)
Patient 5	<i>KRAS</i> (p.Gly12Asp) <i>SMAD4</i> (p.Arg135Ter)	NA	NA	<i>KRAS</i> (p.Gly12Asp) (PIPAC 2) <i>SMAD4</i> (p.Arg135Ter) (PIPAC 2) ND (PIPAC 3) <i>KRAS</i> (p.Gly12Asp) (PIPAC 4)	<i>KRAS</i> (p.Gly12Asp) (PIPAC 5)
Patient 6	<i>KRAS</i> (Gly12Asp)	NA	NA	NA	NA
Patient 7	<i>KRAS</i> (p.Gly12Val)	<i>KRAS</i> (p.Gly12Val) (lymph node)	NA	NA	NA
Patient 8	NA	ND	<i>KRAS</i> (p.Gly12Asp)	NA	NA
Patient 9	<i>MET</i> (p.Arg988Cys)	<i>MET</i> (p.Arg988Cys) (found in PM biopsy and lymph node biopsy)	NA	NA	NA
Patient 10	ND	<i>KRAS</i> (p.Gln61Arg) <i>SMAD4</i> (p.Arg361Cys)	<i>KRAS</i> (p.Gln61Arg)	<i>KRAS</i> (p.Gln61Arg) (PIPAC 2)	<i>KRAS</i> (p.Gln61Arg) (PIPAC 2)
Patient 11	<i>FGFR2</i> (p.Asn549Lys)	<i>KRAS</i> (p.Gly12Asp)	NA	NA	NA
Patient 12	<i>KRAS</i> (p.Gly12Asp)	NA	NA	NA	ND (PIPAC 3 and 4)
Patient 13	<i>KRAS</i> (p.Gly12Val)	NA	NA	NA	NA
Patient 14	<i>KRAS</i> (p.Gly12Val) <i>TP53</i> (p.Arg273His)	NA	NA	NA	NA
Patient 15	NA	<i>KRAS</i> (p.Gly12Arg)	<i>KRAS</i> (p.Gly12Arg)	<i>KRAS</i> (p.Gly12Arg)	<i>KRAS</i> (p.Gly12Arg) (PIPAC 2)
Patient 16	NA	<i>KRAS</i> (p.Gly12Ala)	ND	ND	ND (PIPAC 2)

\*NGS of histological biopsy from peritoneum, unless something else is stated.

NA, not available; ND, not detected; NGS, next-generation sequencing; PF, peritoneal fluid; PM, peritoneal metastasis.

However, that being said, a peritoneal post-PIPAC biopsy may be even preferred over a biopsy from the primary tumour, particularly in patients who underwent several PIPACs; as such a biopsy may be more representative of the current mutational profile of the PM compared with a primary tumour biopsy that may have been taken many months (or years) previously.

Recent studies found the histological PRGS useful for evaluation of response to PIPAC.<sup>19 21</sup> The PRGS has a moderate to good interobserver reproducibility and a near-perfect intraobserver reproducibility.<sup>16</sup> The effect of PIPAC has been defined and monitored by different methods, for example, quality of life, median survival, histological tumour regression, PCI score and changes in gene expression.<sup>2</sup> The combined progression index, based on the highest PRGS and cytology of PF, was independently associated with worse prognosis for OS and for PFS in PM.<sup>25</sup>

Four of our patients with PM-PC were treated with a combination of chemotherapy and PIPAC, and all patients underwent systemic chemotherapy. Therefore, it is difficult to assess whether the histological response was based on the systemic chemotherapy or PIPAC or both. This is particularly true for those patients who were found to have a baseline mean PRGS of 1. Both treatments may have led to improvement of overall outcome. The presence of regressive features prior to the first PIPAC indicates a systemic effect. However, an increase in some regressive features from PIPAC 1 to PIPAC 3 could indicate a PIPAC-related effect. The fact that three patients underwent only one PIPAC was the main cause of why some data were missing, which may have had an impact on the results and interpretations. However, we think that it had a very little or maybe no impact on the results regarding the utility of NGS,

because these analyses were independent of how many PIPACs each patient received. A total of five patients underwent more than three PIPACs, but due to the low number of patients, the effect of more than three PIPAC procedures still remains to be investigated.

In conclusion, PIPAC treatment induces objective response according to the PRGS in the majority of patients with PM-PC. The PRGS seems to be a useful tool for the histological assessment of response to PIPAC. PIPAC does not seem to influence

### Take home messages

- Pressurised IntraPeritoneal Aerosol Chemotherapy (PIPAC) can be a treatment alternative to conventional, systemic chemotherapy in patients with peritoneal metastasis from pancreatic cancer (PM-PC).
- PIPAC induces a histological response in peritoneal biopsies taken prior to PIPAC treatment 2 or 3, compared with biopsies taken prior to PIPAC treatment 1, in the majority of patients.
- Next-generation sequencing (NGS) is able to identify the *KRAS* mutation in peritoneal biopsies and peritoneal fluids taken after systemic chemotherapy and PIPAC at a frequency similar to biopsies taken from the primary tumour prior to treatment.
- NGS may in the future become a tool for detection of predictive mutations in patients treated with PIPAC for PM of various origins, also when only peritoneal post-PIPAC biopsies or peritoneal fluids are available.

the utility or sensitivity of NGS because molecular alterations were identified at expected frequencies. Our data indicate that NGS is a useful tool also in peritoneal biopsies and PFs after PIPAC. Hence, it may be hypothesised that NGS in the future may become a tool for detection of predictive mutations in patients treated with PIPAC for PM of various origins, also when only peritoneal post-PIPAC QBs or PFs are available.

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**Competing interests** None declared.

**Patient consent for publication** Not required.

**Ethics approval** The study was conducted according to the Helsinki Declaration and approved by The Regional Committees on Health Research Ethics for Southern Denmark (project-ID S-20180185) and the Danish Data Protection Agency (project-ID 19/10354).

**Provenance and peer review** Not commissioned; internally peer reviewed.

**Data availability statement** Data are available upon reasonable request. Participant data will be retained at the study facility for five years after completion of the trial according to Danish law. After this time period, it will be de-identified and made available for other researchers upon reasonable request to the corresponding author.

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#### ORCID iDs

Martin Gravensen <http://orcid.org/0000-0001-5719-4916>

Sönke Detlefsen <http://orcid.org/0000-0002-9466-2333>

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