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Sulfatase-2: a prognostic biomarker and candidate therapeutic target in patients with pancreatic ductal adenocarcinoma

Sari F Alhasan^{1,6}, Beate Haugk^{2,6}, Laura F Ogle¹, Gary S Beale¹, Anna Long², Alastair D Burt^{2,3}, Dina Tiniakos^{2,4}, Despina Televantou^{1,2}, Fareeda Coxon⁵, David R Newell¹, Richard Charnley⁵ and Helen L Reeves^{*1,5}

¹Northern Institute for Cancer Research, Paul O’Gorman Building, The Medical School, Framlington Place, Newcastle University, Newcastle upon Tyne NE2 4HH, UK; ²Department of Cellular Pathology, Royal Victoria Infirmary, Newcastle upon Tyne Hospitals NHS Foundation Trust, Newcastle upon Tyne NE1 4LP, UK; ³School of Medicine, Eleanor Harrald Building, Frome Road, The University of Adelaide, Adelaide, 5000 South Australia, Australia; ⁴Institute of Cellular Medicine, The Medical School, Framlington Place, Newcastle University, Newcastle upon Tyne NE2 4HH, UK and ⁵Hepatopancreatobiliary multidisciplinary team, Freeman Hospital, Newcastle-upon-Tyne Hospitals NHS Foundation Trust, Newcastle upon Tyne NE7 7DN, UK

Background: Pancreatic ductal adenocarcinoma (PDAC) is the fifth most common cause of cancer death in the UK. Its poor prognosis is attributed to late detection and limited therapeutic options. Expression of SULF2, an endosulfatase that modulates heparan sulfate proteoglycan 6-O-sulfation and is reportedly tumorigenic in different types of cancer, was investigated.

Methods: SULF2 expression was determined immunohistochemically in archival surgical resection tissue sections from 93 patients with a confirmed histological diagnosis of PDAC between 2002 and 2008 followed for a median of 9 years. Relationships with clinico-pathological parameters and patient survival were explored.

Results: The majority of PDACs showed positive SULF2 staining in tumour cells and intratumoural or tumour-adjacent stroma. Greater than 25% SULF2-positive tumour cells was present in 60% of cancers and correlated with tumour stage ($P=0.002$) and perineural invasion ($P=0.024$). SULF2 intensity was scored moderate or strong in 81% of cancers and positively correlated with vascular invasion ($P=0.015$). High SULF2 expression, defined as >50% SULF2-positive tumour cells and strong SULF2 staining, was associated with shorter time to radiological progression ($P=0.018$, HR 1.98, CI 1.13–3.47). Similarly, by multivariate analysis, high SULF2 expression was independently associated with poorer survival ($P=0.004$, HR 2.10, CI 1.26–3.54), with a median survival of 11 months vs 21 months for lower PDAC SULF2.

Conclusions: Elevated SULF2 in PDAC was associated with advanced tumour stage, vascular invasion, shorter interval to radiological progression and shorter overall survival. SULF2 may have roles as a prognostic biomarker and as a therapeutic target for patients with PDAC.

Each year, ~8800 people are diagnosed with pancreatic ductal adenocarcinoma (PDAC) in the UK (Cancer Research UK (Cancer Statistics)). A significant proportion of cancer-related death is

attributed to PDAC (Siegel *et al*, 2012) with a 5-year survival rate in the region of 6% (DeSantis *et al*, 2014). PDAC is asymptomatic in early stages and its poor prognosis is attributed to late detection,

*Correspondence: Dr HL Reeves; E-mail: helen.reeves@ncl.ac.uk

⁶These two authors contributed equally to this work.

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as well as the aggressive nature of the disease (Chrystoja *et al*, 2013). Consequently, only 10–15% of those affected are candidates for surgical resection, with a 5-year survival post resection of just 10–25% owing to frequent regional invasion and distant micro-metastases at the time of resection (Heinemann and Boeck, 2008). The current palliative chemotherapy regimens for those with later stage disease include gemcitabine with or without nab-paclitaxel, or FOLFIRINOX and provide modest benefit with a median overall survival of <1 year (Conroy *et al*, 2011; Thota *et al*, 2014).

Clinically useful biomarkers which inform prognosis or aid treatment stratification for those with PDAC are presently lacking. The contribution of extracellular microenvironment (ECM) to the development and progression of PDAC is recognised (Apte *et al*, 2015), as is a role for the pro-angiogenic vascular endothelial growth factor (VEGF), which is expressed in >90% of cases, and associated with both liver metastases and shorter survival of patients (Seo *et al*, 2000). ECM interactions with angiogenic as well as proliferative signalling pathways are closely regulated by heparan sulfate proteoglycans (HSPGs). VEGF is one of the HSPG-bound proteins, which include other ligands relevant to PDAC progression such as FGF-1 and FGF-2, Wnts and ECM components such as collagens, fibronectin and laminin which are present in PDAC stroma. The heparan sulfate (HS) chains of HSPGs are rich in uronic acid moieties and sulfate groups that are negatively charged, producing binding sites for growth factors, receptors and ECM molecules. Charge, and therefore sequestration *vs* release of growth factors, as well as orientation and binding affinity, are regulated by the degree of sulfation – particularly at the 6-O position of HS chains (Pye *et al*, 1998, 2000; Ai *et al*, 2003). 6-O-Sulfation is in turn regulated by sulfotransferases which add sulfate groups and the extracellular endosulfatases, sulfatase-1 (SULF1) and sulfatase-2 (SULF2), which remove them (Morimoto-Tomita *et al*, 2002; Saad *et al*, 2005; Staples *et al*, 2011).

SULF2 has been reported to be upregulated and to have a candidate oncogenic role in a number of cancers, including glioblastoma, liver, lung, gastric, oesophageal and prostate cancers (Hur *et al*, 2012; Lai *et al*, 2008, 2010; Lemjabbar-Alaoui *et al*, 2010; Lui *et al*, 2012; Phillips *et al*, 2012). In PDAC, Nawroth *et al* (2007) have previously reported an increase in *SULF2* mRNA in cell lines and increased *SULF2* protein expression relative to non-tumour tissues expression in a small number of cases. shRNA-mediated *SULF2* knockdown or transfection with a catalytically inactive dominant form of *SULF2* enzyme in 3 PDAC cell lines reduced cell growth *in vitro* and tumourigenicity *in vivo* (Nawroth *et al*, 2007).

In this study, we have assessed *SULF2* as a candidate immunohistochemical prognostic biomarker in a cohort of 93 patients who underwent PDAC resection. In the majority of cases *SULF2* was overexpressed in PDAC cells, with the percentage of positive cells correlating with tumour stage, and the *SULF2* intensity (SI) correlating with vascular invasion. Combining percentage of positive cells and SI, high-PDAC *SULF2* was associated with vascular invasion, a shorter time to radiological progression and poorer overall survival.

MATERIALS AND METHODS

Patient cohort. As part of a pilot study, *SULF2* expression was assessed in formalin-fixed paraffin-embedded tissues (FFPE) from 19 patients undergoing resection for PDAC and consenting to the use of their tissues for research projects, governed by the Newcastle upon Tyne Hepatopancreatobiliary Research Tissue bank. The data presented include these cases, in combination with data generated subsequently from a larger retrospective series (an additional 74 cases), exploring *SULF2* expression in archived cases from patients with a confirmed histological diagnosis PDAC. Study of

the retrospective case series was approved by the Newcastle and North Tyneside Regional ethics committee (REC approval 11/H0908/02) on 03 March 2011. The selection interval was from 2002 to 2008 and the REC waived the need for informed consent. Patient confidentiality was respected at all times and analyses were on code-linked anonymised data sets. The Newcastle upon Tyne NHS Foundation Trust Research and Development department approved this project.

The study is classed as a retrospective case series, including patients undergoing resection with curative intent, in whom a diagnosis of PDAC was histologically confirmed. Additional inclusion criteria included the availability of FFPE blocks for study. Patients with known distant metastases at the time of resection were excluded, as were those with histologically benign disease or cholangiocarcinoma.

Clinico-pathological information, including Union for International Cancer Control tumour node metastases (TNM) stage, tumour grade, vascular, lymphatic and perineural invasion, lymph node status, resection margin status, time to progression (TTP) and patient survival, was collected from histopathology reports, radiology reports and patient records. Patient demographics and clinico-pathological data are shown in Table 1. The patients were followed-up for a median of 9.4 (range 6.1 to 12.9) years until 31 December 2014.

Immunohistochemistry. *SULF2* immunohistochemistry was performed on a Benchmark Ultra autostainer (Ventana, Tucson, AZ, USA) using anti-*SULF2* antibody (catalogue number MCA5692GA, AbD Serotec, Oxford, UK) at a dilution of 1:150. In brief, the tissue sections were incubated in primary antibody for 32 min following heat-induced epitope retrieval using CC1 buffer (Ventana) for 64 min at 100 °C. Detection and visualisation was achieved using an OptiView IHC DAB Detection Kit and 4 min of OptiView Amplification (Ventana). Control cases without primary antibody confirmed an absence of non-specific staining. Examples are included in Supplementary Figure 1.

Scoring method. An expert pancreatic pathologist (BH) blinded to patient outcome assessed the *SULF2*-immunostained slides at $\times 100$ magnification. The percentage of *SULF2*-positive carcinoma cells was semi-quantitatively assessed with a score from 0–4 (0 = no carcinoma cells positive, 1 = 1–25% carcinoma cells positive, 2 = 26–50% carcinoma cells positive, 3 = 51–75% carcinoma cells positive and 4 = 76–100% carcinoma cells positive). The *SULF2*-specific SI was assessed with a score from 0–3 (0 = no staining, 1 = weak, 2 = moderate, 3 = strong). Representative cases for each score are shown in Figure 1. If a range of SI was noted the predominant score was used. The semi-quantitative score of *SULF2*-positive PDAC cells and the intensity score were added and a summative combined score (range 0–7) was created for each case. In addition, the extent of stromal staining within the tumour and outside the tumour was scored from 0 to 2 (0 = no staining, 1 = focal staining, 2 = extensive staining). Staining of benign and dysplastic epithelium or of tissues unrelated to the tumour was recorded. Presence of endothelial staining was recorded on all cases as an internal positive control. A second experienced pathologist (DT) also blinded to patient outcome, evaluated 10 random *SULF2*-immunostained slides (~10% of cases). Inter-observer agreement assessed by the Kappa measure was very good for percentage of *SULF2*⁺ve tumour cells (0.75) and perfect (kappa statistic 1) for *SULF2* SI.

Statistical analysis. All statistical analyses were conducted using SPSS statistical package (IBM, version 22). Percentage of *SULF2*-positive cells and SI were treated as ordinal categorical variables. To identify relationships between percentage of *SULF2*-positive cells or SI with clinico-pathological features, cross-tabulation and χ^2 -tests were performed. The Monte Carlo correction was applied

to groups with fewer than 5 cases. To explore factors associated with TTP or survival, univariate Cox regression was conducted. Only significantly associated factors were then analysed using multivariate Cox regression. Median TTP and survival times were calculated using Kaplan–Meier survival analysis. Missing data were excluded from statistical studies. Where analyses were performed on <93 cases, (e.g. TTP) numbers in the analysis are stipulated in the text.

RESULTS

Demographic data for the case series studied. Details of the case series of 93 patients undergoing elective curative surgery for PDAC are summarised in Table 1. The median age was 65 years and 53 were male. The vast majority (92/93) underwent a Whipples' pancreatoduodenectomy for tumours in the head or neck of the pancreas. In one case the tumour was in the body and a total pancreatectomy was performed. None of the patients received pre-operative chemotherapy. The series did not include patients with known metastatic disease and 70/93 were classed as having TNM stage 3 disease. Similarly, 61/93 tumours were graded as poorly differentiated, with the majority having vascular, perineural or lymphatic invasion (Table 1). Forty-six patients (50%) received adjuvant chemotherapy. The commonest adjuvant regime (36 patients) was fluorouracil (5FU) and folinic acid, either as the standard of care or as part of the ESPAC III trial. Similarly, six patients received adjuvant gemcitabine. One patient received the DNA methyl transferase inhibitor MG98, another received capecitabine, whereas in two patients, the adjuvant regime was not documented. Thirty two of 88 patients were documented as receiving palliative second-line chemotherapy, most often gemcitabine, whereas two patients entered clinical trials. As a number of patients follow-up palliative care was outside of our tertiary referral centre, TTP was recorded as first documented 'radiological' progression. Consequently the data set was skewed towards those patients fit enough to undergo active monitoring, with analyses on progression being limited to the 67 patients in whom this clearly defined end point was available. Survival data were available in all 93 patients.

Elevated SULF2 expression was common in PDAC and the percentage of cells positive correlated with tumour stage. SULF2 was consistently identified in endothelial cells and relatively weak expression also observed in benign atrophic ductal epithelium. Weak-to-moderately intense cytoplasmic staining was occasionally observed in dysplastic epithelial cells of PanINs, in islet and acinar cells, periductal glands and benign ductal epithelial cells. Examples of these are shown in Supplementary Figure 1, as is variable staining that was present in lymph node sinuses, adipocytes and inflammatory cells. In contrast, expression of SULF2 in PDAC was frequently of greater intensity (Figure 1). In the PDAC cells, SULF2 was observed in the cytoplasm with a granular quality. Focal membranous staining was identified in tumours with clear cell morphology. Sixty per cent (56 out of 93) of PDACs contained >25% of cells positively stained for SULF2. A summary of the immunohistochemical results is shown in Table 1.

The percentage of SULF2-positive tumour cells correlated significantly with tumour stage ($P=0.002$), the majority (16 out of 22; 73%) of cases with >75% tumour cells positive being stage T3, with local invasion. The percentage of SULF2-positive PDAC cells also correlated with perineural invasion ($P=0.024$).

Cytoplasmic SULF2 expression in intratumoural stromal fibroblasts was either focal or marked in 69% (64 out of 93) of PDAC. SULF2 was also evaluated in the stroma outside the tumours and similarly showed either focal or marked staining in

Table 1. Clinico-pathological variables and SULF2 scores in patients with PDAC

| Patient characteristics | Number |
|--|--------|
| Total | 93 |
| Age (years, median) | 65.3 |
| Sex (female:male) | 40:53 |
| Tumour stage | |
| T1: tumour within pancreas <2 cm | 3 |
| T2: tumour within pancreas >2 cm | 19 |
| T3: local invasion | 70 |
| T4: distant or large vessel invasion | 1 |
| Resection (R0:R1) | 28:65 |
| Lymph nodes (-ve: +ve) | 8:85 |
| Tumour grade | |
| Well differentiated | 1 |
| Moderately differentiated | 31 |
| Poorly differentiated | 61 |
| Vascular invasion (no:yes) | 16:77 |
| Perineural invasion (no:yes) | 2:91 |
| Lymphatic invasion (no:yes) | 33:60 |
| Lymph nodes positive | 8:85 |
| Tumour site | |
| Head of pancreas/neck/uncinate | 89/2/1 |
| Neck and body | 1 |
| Adjuvant chemotherapy | |
| None | 47 |
| 5FU regime | 36 |
| Gemcitabine | 6 |
| Other (defined in text) | 4 |
| Palliative second-line chemotherapy | 32/56 |
| Survival (median months) | 16.5 |
| SULF2 scoring | |
| Percentage SULF2-positive tumour cells | |
| 0 = 0% | 3 |
| 1 = 1–25% | 34 |
| 2 = 26–50% | 17 |
| 3 = 51–75% | 17 |
| 4 = 75–100% | 22 |
| SULF2 staining intensity | |
| 0 = None | 3 |
| 1 = Weak | 15 |
| 2 = Moderate | 38 |
| 3 = Strong | 37 |
| SULF2 combined score | |
| 0 | 3 |
| 1 | 0 |
| 2 | 9 |
| 3 | 8 |
| 4 | 35 |
| 5 | 13 |
| 6 | 13 |
| 7 | 12 |
| SULF2 tumour stroma | |
| None | 29 |
| Focal | 54 |
| Marked | 10 |
| SULF2 stroma outside | |
| None | 17 |
| Focal | 48 |
| Marked | 28 |
| Abbreviations: R = resection margins; R0 = negative resection margins; R1 = microscopic tumour infiltration (that is, tumour cells present within 1 mm of the resection margin); SULF = sulfatase. | |

the majority of cases (76 out of 93; 82%). Stromal expression of SULF2 did not correlate with either PDAC grade or stage.

SULF2 intensity in PDAC was positively associated with the presence of vascular invasion. In all, 81% (75 out of 93) of cancers showed moderate or strong intensity of SULF2 expression.

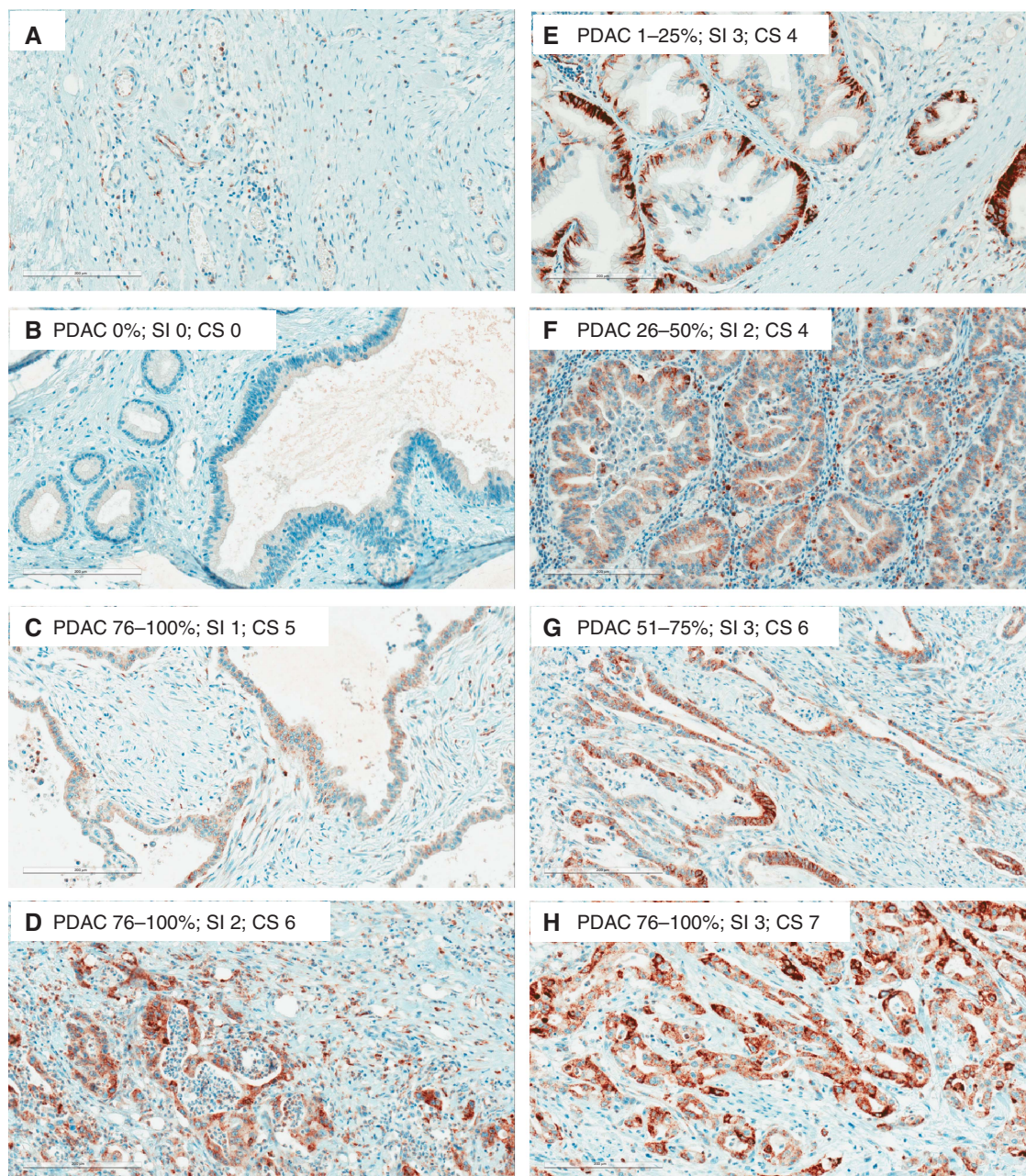


Figure 1. SULF2 expression in pancreatic ductal adenocarcinoma. Representative immunohistochemical images of PDAC cases showing different percentages of SULF2-positive tumour cells (0–100%) and SI of SULF2 in tumour cells. SI 0: none, SI 1: weak, SI 2: moderate, SI 3: strong staining. (A) shows endothelial cell SULF2-positivity used as internal positive control. CS = combined SULF2 score.

The intensity correlated significantly with the presence of vascular invasion ($P = 0.015$).

The SULF2 combined score correlated with the presence of vascular invasion. The combined SULF2 IHC score of percentage cells positive and the level of intensity, as shown in Table 1, correlated significantly with the presence of vascular invasion ($P = 0.026$, Pearson's χ^2 , linear by linear association).

High-PDAC SULF2 was associated with a shorter time to radiological progression. Radiological TTP was documented in 67 out of 93 (62%), with the median TTP being 11.6 months. One factor associated with radiological TTP was lymph node positivity ($P = 0.007$, HR 4.15, CI 1.49–11.58). Age, sex, tumour grade, tumour stage, resection margin status, vascular invasion, lymphatic invasion and perineural invasion had no significant

impact on TTP in this relatively small series. Considering the SULF2 immunohistochemistry (IHC) scores applied, both '>75% PDAC cells SULF2 positive' or SULF2 SI scored as 'strong', were associated with significantly shorter TTP. Combining the IHC scores, as shown in Table 1, identified those cases with a combined score of 6 or 7 as having a poorer outcome. The median TTP was 8 months in this combined group, vs 13 months in combined groups scoring 0–5, as shown in Supplementary Figure 2 ($P = 0.018$, HR 1.98, CI 1.13–3.47). Both lymph node metastasis and the combined SULF2 score (groups 0–5 vs 6–7), were independently associated with a shorter TTP (lymph nodes positive $P = 0.006$, HR 4.28, CI 1.53–12; high SULF2 $P = 0.014$, HR 2.05, CI 1.16–3.63, multivariate Cox regression; Supplementary Table 1).

Of the 67 cases with documented radiological TTP, 36 received adjuvant chemotherapy. Adjuvant chemotherapy had no significant impact on radiological TTP. Standard histopathological

features were not associated with response to chemotherapy. However, the percentage of SULF2-positive PDAC cells trended towards significantly shorter TTP ($P=0.067$, HR 1.28, CI 0.98–1.66), as did SI of PDAC cells ($P=0.062$, HR 1.58, CI 0.98–2.54). High SULF2 using the combined IHC SULF2 score was associated with a median TTP of 10.8 months vs 15.5 months for those with lower SULF2 ($P=0.013$, HR 2.76, CI 1.24–6.17).

High-PDAC SULF2 was significantly associated with shorter overall survival. The overall mortality during the period of

follow-up was 94% (87 out of 93), with a median of 16.5 months post resection until death (range 0.3–148, s.d. 30). Factors significantly associated with survival post resection included age, lymph node metastasis and receipt of adjuvant chemotherapy, as assessed by univariate Cox regression analysis (Table 2). Regarding SULF2, those cases with >75% of PDAC cells positive had a poorer outcome, as did those cases with strong SI compared with cases without SULF2 expression (Table 2). PDAC with a combined SULF2 score 6 or 7 had a significantly poorer overall survival (Table 2). Comparing combined groups 0–5 vs 6–7, the median

Table 2. Factors associated with survival

| Variable | n | Univariate analysis | | | Multivariate analysis | | |
|---|--------------|---------------------|-------------|------------------|-----------------------|-------------|-------------------|
| | | P value | HR | 95% CI | P value | HR | 95% CI |
| Age (median 65.3 years) | 93 | 0.031* | 1.03 | 1.00–1.05 | 0.462 | 1.01 | 0.99–1.04 |
| Sex (male/female) | 53/40 | 0.573 | 1.13 | 0.74–1.73 | Not included | | |
| PDAC TNM stage | | | | | Not included | | |
| 1 = Within pancreas < 2 cm | 3 | 0.595 | | | | | |
| 2 = Within pancreas > 2 cm | 19 | 0.299 | 2.17 | 0.50–9.38 | | | |
| 3 = Local invasion | 70 | 0.217 | 2.44 | 0.59–9.99 | | | |
| 4 = Distant spread | 1 | 0.268 | 3.92 | 0.35–43.9 | | | |
| PDCA grade | | | | | Not included | | |
| Well | 1 | 0.254 | | | | | |
| Moderate | 31 | 0.676 | 1.53 | 0.21–11.3 | | | |
| Poor | 61 | 0.443 | 2.17 | 0.30–15.8 | | | |
| Resection margin (R0/R1) | 28/65 | 0.247 | 1.32 | 0.83–2.10 | Not included | | |
| Vascular invasion (no/yes) | 16/77 | 0.170 | 1.49 | 0.84–2.65 | Not included | | |
| Perineural invasion (no/yes) | 2/91 | 0.128 | 4.66 | 0.65–33.6 | Not included | | |
| Lymphatic invasion (no/yes) | 33/60 | 0.208 | 1.33 | 0.85–2.09 | Not included | | |
| Lymph nodes +ve (no/yes) | 8/85 | 0.011* | 3.28 | 1.31–8.19 | 0.003** | 4.31 | 1.65–11.28 |
| Adjuvant therapy (no/yes) | 47/46 | 0.012* | 0.58 | 0.38–0.89 | 0.009** | 0.50 | 0.30–0.84 |
| Palliative chemotherapy | 32/56 | 0.88 | 0.68 | 0.43–1.06 | Not included | | |
| Percentage of PDAC SULF2 ^{++ve} | | | | | Not included | | |
| 0 | 3 | 0.123 | | | | | |
| 1–25 | 34 | 0.057 | 6.94 | 0.95–51.0 | | | |
| 26–50 | 17 | 0.080 | 6.12 | 0.81–46.5 | | | |
| 51–75 | 17 | 0.084 | 5.99 | 0.79–45.6 | | | |
| 76–100 | 22 | 0.024* | 10.3 | 1.35–76.3 | | | |
| SULF2 intensity | | | | | | | |
| None | 3 | 0.008** | | | (0.066) | | |
| Weak | 15 | 0.045* | 7.96 | 1.04–60.7 | (0.114) | (5.31) | (0.67–42.0) |
| Moderate | 38 | 0.103 | 5.25 | 0.72–38.5 | (0.174) | (4.03) | (0.54–30.1) |
| Strong | 37 | 0.022* | 10.4 | 1.41–76.2 | (0.058) | (7.04) | (0.93–53.0) |
| Combined SULF2 score | | | | | | | |
| 0 | 3 | 0.028* | | | (0.084) | | |
| 2 | 9 | 0.066 | 6.97 | 0.88–55.3 | (0.164) | (4.49) | (0.54–37.3) |
| 3 | 8 | 0.084 | 6.33 | 0.78–51.2 | (0.211) | (3.84) | (0.47–31.7) |
| 4 | 35 | 0.055 | 7.04 | 0.96–51.7 | (0.124) | (4.84) | (0.65–36.3) |
| 5 | 13 | 0.127 | 4.94 | 0.63–38.4 | (0.194) | (3.97) | (0.50–31.9) |
| 6 | 13 | 0.040* | 8.47 | 1.10–65.3 | (0.089) | (6.00) | (0.76–47.3) |
| 7 | 12 | 0.006** | 17.9 | 2.27–142.3 | (0.021)* | (11.65) | (1.44–94.1) |
| Combined SULF2 score 0–5 (lower) vs 6–7 (high) | 68/25 | 0.002** | 2.21 | 1.35–3.61 | 0.004** | 2.10 | 1.26–3.54 |
| Tumour stromal SULF2 | | | | | Not included | | |
| None | 29 | 0.861 | | | | | |
| Focal | 54 | 0.644 | 1.12 | 0.70–1.78 | | | |
| Marked | 10 | 0.647 | 1.19 | 0.57–2.46 | | | |
| Stromal SULF2 (outside) | | | | | | | |
| None | 17 | 0.047* | | | 0.377 | | |
| Focal | 48 | 0.103 | 0.62 | 0.35–1.10 | 0.700 | 1.14 | 0.59–2.17 |
| Marked | 28 | 0.775 | 1.09 | 0.59–2.01 | 0.224 | 1.51 | 0.78–2.91 |

Abbreviations: CI = confidence interval; HR = hazard ratio; PDAC = pancreatic ductal adenocarcinoma; SULF = sulfatase; TNM = tumour node metastases. The multivariate Cox regression analysis included variables (highlighted in bold) with a P value < 0.05 in univariate analyses. The combined bivariate SULF2 score comparing low (0–5) vs high (6–7) was entered into the analysis shown. Data in brackets show similar multivariate analyses entering either SI score or combined SULF2 score with ordinal groups instead. For categorical variables, the comparator was the first listed variable. HR and 95% lower and upper CI are shown for all variables. * P < 0.05; ** P < 0.01.

patient survival was 21 months vs 11 months, respectively ($P=0.002$, HR 2.21, CI 1.35–3.61), as shown in Figure 2A. Multivariate Cox regression analysis, including age, lymph node metastasis, receipt of adjuvant therapy, combined SULF2 IHC score 0–5 vs 6–7 and stromal SULF2 outside the tumour is shown in Table 2. The presence of lymph node metastasis and high SULF2 expression were independently associated with a poorer survival, whereas receipt of adjuvant chemotherapy was associated with improved survival.

In the subgroup of patients who received adjuvant chemotherapy ($n=46$), age, sex and tumour features had no impact on survival. Again, however, higher SULF2-positive PDAC cells or strong SULF2 SI were associated with poorer outcomes. Using the combined IHC score, median survival was 25 months in those with lower SULF2 (score 0–5) vs 12 months in those with a high SULF2 (score 6–7; $P=0.001$, HR 3.23, CI 1.59–6.57) as shown in Figure 2B.

DISCUSSION

The development of personalised or precision medicine for cancer therapy requires the discovery, qualification and clinical use of prognostic and predictive biomarkers. In PDAC the most common driver genes with recurrent mutations; *KRAS*, *SMAD4*, *TP53* and *CDKN2A/p16*, cannot currently be exploited by available targeted therapies. Whole-genome sequencing has identified new candidates which may be druggable, but at low individual prevalence (Waddell *et al*, 2015). Hence predictive biomarkers that can be used to select specific treatments for individual patients have not had a major impact on the management of the disease, with systemic treatment being limited to single agent (e.g., gemcitabine) or combination (e.g., FOLFIRINOX) cytotoxic drug therapy.

The major focus for biomarker discovery in PDAC has therefore been the identification of novel prognostic biomarkers to inform clinical decision making. For example, to identify poor prognosis patients that might be suitable for experimental therapy in the primary setting, and those with a relatively favourable prognosis in whom more aggressive conventional treatments may be appropriate. A large number of conventional clinical, circulating and pathological factors (Giovinazzo *et al*, 2012; Winter *et al*, 2013; Bilici, 2014; Jazieh *et al*, 2014; Lamarca and Feliu, 2014), as well as genomic and proteomic parameters (Marengo and Robotti, 2014), have been the subject of biomarker discovery research in PDAC. However, the glycoprotein CA19–9, the only FDA approved biomarker for PDAC, is the sole factor in widespread routine clinical use. Serum CA19–9 has utility as a diagnostic, prognostic and surrogate response biomarker, but it has well known limitations and its precise value in these settings remains the subject of debate (Bilici, 2014; Jazieh *et al*, 2014).

Thus far, meta-analyses of immunohistochemical prognostic biomarker studies in PDAC have identified VEGF levels (Ansari *et al*, 2011; Smith *et al*, 2011), consistent with a role for the growth factor in tumour angiogenesis and hence progression and prognosis. More recently a meta-analysis assessing expression of the chemokine receptor CXCR4 supported an association between metastatic disease and overall survival in patients with PDAC (Krieg *et al*, 2015). Both of these prognostic biomarkers offer links to the molecular and cellular pathology of PDAC in a way that could ultimately be exploited therapeutically. As yet, however, these hopes have not been realised and there remains a need for additional prognostic biomarkers, preferably ones which may be predictive of responses to novel therapies.

Building on a previous study reporting an increase in SULF2 detected immunohistochemically in 4 of 7 patient PDAC samples relative to non-tumour tissues (Nawroth *et al*, 2007), the current

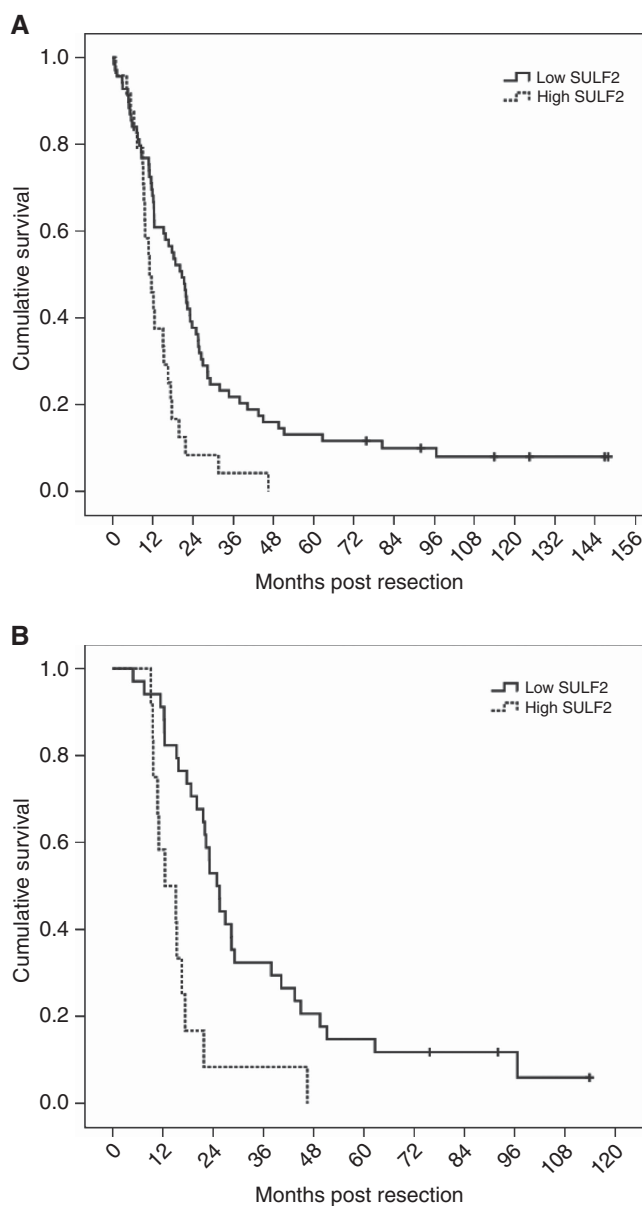


Figure 2. High SULF2 in pancreatic ductal adenocarcinoma is associated with poorer survival. Kaplan–Meier survival plots show (A) cumulative survival of patients with high SULF2 expression compared with patients with lower SULF2 expression. (B) Cumulative survival of a subset of PDAC patients receiving adjuvant chemotherapy. High SULF2 equates to cases with a percentage cells positive and SI combined score of 6–7, vs low SULF2 (scores 0–5).

study has identified expression of the extracellular endosulfatase SULF2 as poor prognostic biomarker in PDAC that is related to tumour stage, vascular invasion, radiological progression and overall survival. By controlling the degree of sulfation of extracellular HSPGs, and hence growth factor binding and signalling, SULF2 can regulate a number of growth factors that have been implicated in PDAC pathobiology, notably VEGF, Wnts and TGF- β (Nawroth *et al*, 2007; Rosen and Lemjabbar-Alaoui, 2010). Importantly, SULF2 is at the interface between the PDAC cell and the tumour microenvironment, and the importance of local tumour-host cell interactions in cancer biology is increasingly recognised. Hence the biological function of SULF2, its involvement in pathways of known importance in PDAC pathogenesis and evidence that SULF2 is a prognostic biomarker, presented here, all identify this enzyme

as a factor that has potential as both a biomarker and a therapeutic target.

Target validation data in PDAC (Nawroth *et al*, 2007), as well as in other tumour types (Rosen and Lemjabbar-Alaoui, 2010), are consistent with SULF2 representing a key signalling node that could be exploited therapeutically. Inhibitors of SULFs are now being described (Schelwies *et al*, 2010; Miller *et al*, 2015) and, once these have been developed to the stage where they are suitable for cell-based and preferably *in vivo* studies, the therapeutic potential of SULF2 inhibition can be explored. The data presented in this study is generated from a relatively small retrospective case series, which is a limitation. However, the pre-clinical data generated using molecular genetic approaches in PDAC models are compelling (Nawroth *et al*, 2007) and, in conjunction with the demonstration in the current study that SULF2 expression is a poor prognostic biomarker, identifies SULF2 as a priority area for further PDAC research.

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CONFLICT OF INTEREST

The authors declare no conflict of interest.

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