Cadherin-1 and cadherin-3 cooperation determines the aggressiveness of pancreatic ductal adenocarcinoma

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Background: Pancreatic ductal adenocarcinoma (PDAC) is characterised by an extensive tissue invasion and an early formation of metastasis. Alterations in the expression of cadherins have been reported in PDAC. Yet, how these changes contribute to tumour progression is poorly understood. Here, we investigated the relationship between cadherins expression and PDAC development.

Methods: Cadherins expression was assessed by immunostaining in both human and murine tissue specimens. We have generated pancreatic cancer cell lines expressing both cadherin-1 and cadherin-3 or only one of these cadherins. Functional implications of such genetic alterations were analysed both in vitro and in vivo.

Results: Cadherin-3 is detected early at the plasma membrane during progression of pancreatic intraepithelial neoplasia 1 (PanIN-1) to PDAC. Despite tumoural cells turn on cadherin-3, a significant amount of cadherin-1 remains expressed at the cell surface during tumourigenesis. We found that cadherin-3 regulates tumour growth, while cadherin-1 drives type I collagen organisation in the tumour. In vitro assays showed that cadherins differentially participate to PDAC aggressiveness. Cadherin-3 regulates cell migration, whereas cadherin-1 takes part in the invadopodia activity.

Conclusions: Our results show differential, but complementary, roles for cadherins during PDAC carcinogenesis and illustrate how their expression conditions the PDAC aggressiveness.

Pancreatic ductal adenocarcinoma (PDAC), the most frequent form of pancreatic cancer, is the fourth leading cause of cancer deaths (Kleeff et al, 2016). During the last decade there has been an alarming worldwide increase in its incidence rate. The absence of early detection tools combined with ineffective treatments are the main reasons of a poor prognosis of pancreatic cancer. The result of these factors combination is that PDAC is projected to become the leading cause of cancer deaths in western countries by 2030 (Rahib et al, 2014).

Pancreatic ductal adenocarcinoma is indeed one of the most lethal cancers with a 5-year survival rate of less than 5% (Kleeff et al, 2016). The low survival rates associated with this human malignancy reflect a diagnosis at later stages characterised by a well-developed, dense, desmoplastic stroma, as well as the presence of metastasis in other organs (Karandish and Mallik, 2016). To overcome this devastating disease, it is essential to develop early detection methods including the identification of new molecular targets.

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Pancreatic ductal adenocarcinoma is thought to evolve through progression of precursor lesions, called pancreatic intraepithelial neoplasia (PanINs) (Singh and Maityra, 2007). PanINs are classified into a four-tier classification (including PanIN-1A, 1B, 2, -3) reflecting a progressive greater architectural distortion of the ducts, loss of the columnar epithelia’s polarity, and nuclear pleomorphism, cumulating in PDAC with the invasion of the basement membrane. Growing evidence from molecular studies indicates that PDAC is genetically very complex with a high diversity of mutations compared with other cancers. However, KRAS (PanIN-1), P16/CDKN2A (PanIN-2), TP53 and MAD4/DPC4 (PanIN-3) are the most commonly altered genes in a stepwise fashion and provide a model for the PDAC development (Hruban et al, 2008). In addition to these commonly recognised genetic drivers of pancreatic carcinogenesis, a recent genomic analysis of PDAC has increased the number of genomic alterations (Waddell et al, 2015; Bailey et al, 2016).

It is well known that tumour progression involves changes in tumour cell capacities to adhere and communicate with neighbouring cells. Among cell–cell adhesion defects, changes in cadherin expression have a critical role in tumour progression and metastasis (Thiery, 2009). Classical cadherins are a family of single-span transmembrane domain glycoproteins that primarily mediate calcium-dependent cell–cell adhesion, which are differentially expressed throughout the body (Nelson, 2008). These glycoproteins establish homophilic adhesion between neighbouring cells through their extracellular cadherin domain and play cohesive organising functions required for tissue integrity (Niesen et al, 2011). The most extensively studied classical cadherins include cadherin-1, -2 and -3, formerly termed E-, N- and P-cadherin (Gul et al, 2017). The intracellular domain of cadherins is associated with the cytoplasmic proteins termed catenins that link cadherins to the cytoskeleton and cell signalling pathways. Therefore, the alterations among cadherins can affect not only cell–cell adhesive properties, but also the signal transduction activity. The latter can modulate a large number of cell functions including tumourigenesis. A cadherin subtype-switching—from cadherin-1 to cadherin-2 was reported in PDAC (Al-Aynati et al, 2004; Nakajima et al, 2004). Moreover, it has been suggested that this phenomenon contributes to metastatic dissemination of tumour cells (Su et al, 2016).

In humans, cadherin-3 is only detected in a few organs (Imai et al, 2008). However, its overexpression is strongly associated with a poor prognosis in some solid tumours including breast, lung and pancreatic cancers (Tanisch et al, 2005; Imai et al, 2008; Ribeiro and Paredes, 2014; Sakamoto et al, 2015). In vitro studies have shown that cadherin-3 induces pancreatic tumour cell motility and invasiveness (Tanisch et al, 2005; Sakamoto et al, 2015), suggesting that blocking cadherin-3 could be a novel therapeutic approach for the treatment of aggressive pancreatic cancer. Although cadherin-3 expression has been reported in PDAC, its expression and role in the early stages of pancreatic carcinogenesis has not been studied (Tanisch et al, 2005; Imai et al, 2008; Sakamoto et al, 2015). Indeed, the cadherin-3 is not yet considered as an early biomarker of PDAC.

Since a cadherin-1 staining was observed at the cell membrane in 50–70% of the PDAC cases (Al-Aynati et al, 2004), it was tempting to speculate that pancreatic cancer cells can express at their surface both cadherin-1 and -3. The aim of this study was therefore to determine whether simultaneous cadherin-1 and -3 expressions could induce a more aggressive biological behaviour in PDAC. To this end, the expression of these two cadherins was immunohistochemically examined through human pancreatic cancer development and in KPC mouse model. Moreover, we have deciphered the role of each cadherin in cell invasiveness both in vitro and in vivo by using orthotopic and ectopic pancreatic tumour mouse models.

**MATERIALS AND METHODS**

**Cell culture and tissues.** BxPC-3 cells were routinely cultured as previously described (Fabre et al, 1993) and authenticated using STR profiling (ATCC). Primary cultures from human tumours (CRCM08 and CRCM110 cells) were cultured in serum-free ductal media as previously published (Gayet et al, 2015). Cells were passaged in the laboratory for no more than 10 passages. Cells were tested for *Mycoplasma* monthly.

A pancreas adenocarcinoma tissue array (#PA484; 24 cases) and a pancreas intraepithelial neoplasia, pancreatitis and cancer tissue array (#BIC14011a; 24 cases) were purchased from Pantomics (Euromedex, Souffelweyersheim, France). The tissue array of 55 PDAC samples from xenografted tumours was obtained either by surgery or endoscopic ultrasound-guided fine-needle aspiration biopsy, as previously described (Duconseil et al, 2015). Some tumour tissue samples were obtained after pancreatic resection (duodeno-pancreatectomy) from patients diagnosed with pancreatic adenocarcinoma (Gastroenterology and Digestive Surgery departments, Timone Hospital, Marseille, France; CRO2 Agreement number DC20131857) between February 2007 and February 2016. All specimens were re-evaluated by an expert pathologist.

PanINs, tumour or control pancreatic tissues from *Pdx1-Cr,Ink4a*/*P16*-LSL-Kras*G12D* mice (Leca et al, 2016) were fixed in 4% (wt/vol) formaldehyde for immunohistochemistry.

**Gene expression microarrays.** Gene expression microarrays from xenografted tumours issued from 55 PDAC tissue samples (see above) were performed as previously described (Duconseil et al, 2015). All array data are available on the Gene Expression Omnibus (http://www.ncbi.nlm.nih.gov.gate2.inist.fr/geo; Accession number GSE55513).

**Cadherin invalidation.** Cadherin-1 or cadherin-3 was stably knocked down in human pancreatic adenocarcinoma BxPC-3 cell line by RNA interference technique. To do so, mission shRNA lentiviral transduction particles (Sigma, St Quentin Fallavier, France) have been used as previously described for melanoma IGR 39 cell line (Siret et al, 2015). Briefly, two lentiviral transduction particles were chosen per cadherin (TRCN000023836 and TRCN0000235419 cadherin-3) and a negative control (MISSION Non-Mammalian shRNA Control Transduction Particles HSC002V) was included. Cells were infected with lentiviral transduction particles according to the manufacturer’s instructions. Small hairpin RNA-containing cells were selected with 5 μg ml⁻¹ puromycin and cloned. Cell surface cadherin extinction was assessed by both immunofluorescence (Figure 3A) and western blot (Supplementary Figure S1).

**Spheroid formation assay.** Spheroid formation assay was used to investigate the effect of cadherins on the cell – cell adhesion properties as previously described (Tercoli et al, 2017). Isolated cells were seeded onto U-bottom untreated tissue culture 96-well plates at a concentration of 5000 cells per well in 100 μl DMEM containing 10% FCS and 0.24% methycellulose. Spheroids were grown for 24 h and pictures were taken using an Olympus microscope (objective × 4). The size of the spheroids was quantified by measuring the area occupied by cells using ImageJ software (rsb.info.nih.gov/ij/).

**3D-invasion assay.** Cells were cultured for 72 h as described above to obtain compact spheroids. The latter were embedded into 1.6 mg mL⁻¹ of bovine collagen I matrix (Advanced Biomatrix) diluted in 0.12% methycellulose and covered with serum-free medium. Spheroids were then incubated for 24 h at a temperature and CO₂ controlled chamber mounted on an inverted microscope Olympus IX83. Images were captured every 60 min using an orca-
flash4 camera (Hamamatsu) with a × 4 objective. The size of the spheroids was quantified by measuring the area occupied by cells using ImageJ software (rsb.info.nih.gov/ij/). In this experiment, cell invasion was independent of cell proliferation since cell treatment with mytomycin C did not alter the results obtained in 3D invasion (data not shown).

**2D-invasion assay.** Owing to their inability to form spheroids, the 3D invasion assay was unachievable for CRCM08 cells. In order to study their invasive properties, in vitro invasion through type I collagen was performed using transwell-based cell culture chamber systems (Millipore-Chemicon). Cells were suspended in DMEM/0.1% BSA and added at a concentration of 20 000 cells per well to the upper chamber containing a polycarbonate membrane filter of 8 μm pore size and a type I collagen layer. The lower chamber was filled with DMEM supplemented with 1% FCS. Plates were incubated under standard culture conditions for 24 h. Cells remaining on the upper side of the filter and type I collagen were removed using a cotton swab. The invasive cells on the lower side of the filter were stained with the crystal violet. Quantification of invasive cells was performed in two independent experiments performed in triplicate. The invasion index was calculated as the mean number of migrated cells counted in 10 microscopic fields per well.

**Wound healing assay.** Cell monolayers were wounded using a sterile tooth-pick, washed twice and incubated in serum-free medium. Plates were incubated for 6 h in a temperature and CO₂ controlled chamber mounted on an inverted microscope Olympus IX83. Images were captured every 5 min using an orca-flash4 camera (Hamamatsu) with a × 40 objective.

**Isolated cell trajectory.** Isolated cells were seeded on a type I collagen covered 12-well plate (12 000 cell cm⁻²) and incubated in serum-free medium for 6 h in a temperature and CO₂ controlled chamber mounted on an Olympus IX83 inverted microscope. Single cell trajectories were analysed by videomicroscopy by capturing images every 5 min using an orca-flash4 camera with a ×40 objective. The cell tracks were obtained using ImageJ software. The values for the assessment of migration speed, velocity and directionality were obtained using ImageJ software. Migration speed was determined by the ratio between the total distance and duration of cell migration. Cell velocity corresponds to the distance between the positions of the cells at the beginning and the end of the experiment divided by time. Cell directionality is the ratio between cell velocity and cell speed.

**Fluorescent-gelatin degradation assay.** Coverslips coated with FITC-conjugated gelatin (Sigma) were prepared as described by (Steffen et al, 2008). Isolated cells were plated on FITC-conjugated gelatin at 70 000 cells cm⁻² for 16 h and then fixed in 4% paraformaldehyde. The areas of degraded matrix were observed with an SP5 Leica confocal microscope equipped with LAS AF Lite software. ImageJ software was used to evaluate the number of invadopodia per cell. At least 150 cells from three different coverslips were chosen randomly and analysed.

**Subcutaneous and orthotopic xenografts of pancreatic cancer cells.** All experimental procedures involving animals were performed in accordance with the French Guidelines and approved by the ethical committee of Marseille (agreement 50-31102012). BxPC-3 cells were harvested by mild trypsinisation, washed twice in PBS, and then suspended in Matrigel at 2 × 10⁶ cells per 100 μl. To induce tumours, cell suspension was injected subcutaneously (s.c.) into the flank of 6–8-week-old female NMRI-Foxn1nu/Foxn1nu mice (Charles River Laboratories, L’Arbresle, France; six mice per group). Injected mice were monitored daily and killed 3 weeks after inoculation. Tumours were removed, weighed, and a digital caliper was used to measure their volumes. Tissue specimens were fixed in 4% formalin, embedded in paraffin and further cut into 4 μm sections, immediately stored at 4 °C. The latter were then subjected to Masson’s trichrome staining (RAL Diagnostics; Prochilab, Lormont, France) or immunohistofluorescence (IHC).

Orthotopic xenografts were performed as described elsewhere (Kim et al, 2009; Chai et al, 2013). BxPC-3 cells in single suspension were mixed with Matrigel (BD Biosciences, Le-Pont-de-Claye, France) at a ratio of 10⁶ cells per 50 μl Matrigel. Mice were anaesthetised with 5% isoflurane in oxygen. An incision (1–1.5 cm) was created in the left abdominal flank under aseptic conditions. The pancreas was carefully exposed, and 50 μl of cell solution diluted in Matrigel was injected into the tail of the pancreas (n = 4 per group). The peritoneum and fascia were closed in separate layers using a continuous 4/0 Prolene suture; the skin wound was closed in a similar way. Mice received s.c. injections of analgesic (Buprenorphine 0.1 mg kg⁻¹) before the operation and 6 h following the suturing.

**Immunofluorescence.** Cells plated on glass coverslips were fixed for 20 min with 2% formaldehyde in PBS. Thereafter, cells were permeabilised with 0.1% saponin in PBS for 20 min, and blocked for 30 min in PBS containing 4% (w/v) BSA. Cadherin-1 and cadherin-3 were sequentially detected by incubation with mouse HECD-1 mAb (Takara, Saint-Germain-en Laye, France), and rabbit Ab (2130s, Cell signalling Technology, Saint-Quentin-en-Yvelines, France), respectively. After three washes, samples were incubated for 1 h with Alexa Fluor 488- and 594-conjugated goat Ig (20 μg ml⁻¹), raised against mouse and rabbit Igs respectively. Samples were then washed and mounted in ProLong Gold (Invitrogen, Illkirch, France). After dewaxing and antigen retrieval, paraffin-embedded 4 μm sections of human pancreatic tumour biopsies were treated and subjected to double immunostaining, as described above. Images were captured and analysed using a SP5 Leica confocal microscope equipped with LAS AF Lite software. The intensity and distribution patterns of specific immunofluorescence staining were evaluated by two blinded, independent observers using a semi-quantitative score (graded as 0 none, 1 weak, 2 moderate, 3 strong staining, and 4 saturated signal). This analysis was performed without any knowledge of anatomopathological evaluation, the diagnosis or the standard performed pathological haematoxylin reaction of each specimen.

**Statistical analysis.** Data are presented as the means ± s.d. for three independent experiments performed in triplicate. Comparison between the two conditions was made using the Mann–Whitney test. P < 0.05 was considered statistically significant in all analyses and is indicated by ‘*’ when P < 0.01, ‘**’ when P < 0.01 and ‘***’ when P < 0.05.

**RESULTS**

**Cadherin-3 expression increases during PDAC progression.** To date, there is no information available concerning cadherin-3 expression in PDAC precursor lesions. We therefore analysed its expression in both human and murine tissue specimens. In healthy human pancreas, cadherin-3 was located in the cell cytoplasm, whereas cadherin-1 was found at the sites of cell–cell contacts (Figure 1A and Supplementary Figure S2). Moreover, cadherin-3 was progressively observed at the plasma membrane during carcinogenesis from PanIN-1 to PDAC. It should be noted that cadherin-1 remained associated with the cell membrane; although a small decrease in the staining intensity was observed between PanIN-2 and PDAC. We scored the expression level of both cadherin-1 and cadherin-3 in all examined tissues. The ratio cadherin-3/cadherin-1 increased from the PanIN-1 to PDAC and was maximal for the grade III of PDAC (Figure 1B). This reflected the concomitant increase of cadherin-3 expression and the partial
cadherin-1 extinction during the tumour progression. Our data were confirmed using the PDX1-Cre, KrasG12D, Ink4a/Arf<sup>flox/flox</sup> mice, a model that develops PDAC from PanIN lesions similar to the human disease (Supplementary Figure S3). In this mouse model, cadherin-1 expression was reduced, while cadherin-3 levels were increased at the cell membrane during progression from PanIN-1 to PDAC.

Taken together these results indicate that cadherin-3 is expressed early during PDAC carcinogenesis, suggesting that this molecule could be an early marker. Despite the appearance of cadherin-3 at the cell membrane, significant amounts of cadherin-1 remained associated with the cell membrane.

**Co-localisation of cadherin-3 with cadherin-1 during PDAC progression.** Double immunostaining experiments were performed to determine whether both cadherin-1 and cadherin-3 are co-expressed in pancreatic tissues. Dual staining showed that once cadherin-3 is expressed, it localises with cadherin-1 in the same cells at cell–cell contact sites (Figure 1A). This observation strongly suggests that cadherin subtype switching is not a general feature in PDAC. To confirm this hypothesis, we analysed both cadherin-1 and cadherin-3 mRNA expression in 55 PDAC samples from patients preserved as xenografts in nude mice. Both cadherin-1 and cadherin-3 transcripts were detected at the same time in a large majority of xenografts (Figure 2A). Quantification of the immunostaining of cadherin-1 and cadherin-3 in those samples confirmed that there is no switch from cadherin-1 to cadherin-3 in PDAC (Figure 2B). It should be noted that cadherin-3 expression level was higher in xenografts issued from metastasis than those obtained from primary tumours (Figure 2C).

Altogether, these results suggest that in PDAC both cadherin-1 and cadherin-3 could elaborate adhesive networks that may regulate tumoural cell behaviour. To decipher the functional effects of cadherin-1 and cadherin-3 co-expression, we established cell models where the expression of these cell–cell adhesion molecules can be manipulated. The human pancreatic cancer cell line BxPC-3 was used as a model system. These cells express indeed high levels of both cadherin-1 and cadherin-3 at cell–cell contacts (Figure 3A). To stably and selectively knockdown the expression of cadherin-1 or cadherin-3, we used shRNA, targeting each one of the two cadherins. We selected among the several clones obtained, one clone with an effective and efficient silencing for cadherin-1 or cadherin-3 (Supplementary Figure S1). The generated stable cell lines were called cadh-1<sup>+</sup>/cadh-3<sup>+</sup> (no cadherin depletion), cadh-1<sup>−/−</sup>/cadh-3<sup>−/−</sup> (cadherin-3 depletion) and cadh-1<sup>−/−</sup>/cadh-3<sup>+</sup> (cadherin-1 depletion). We also used PDAC-derived primary cell cultures derived from patient-derived xenograft (PDX) in nude mice (Figure 3B). CRCM110 cells express both cadherin-1 and cadherin-3, whereas CRCM08 cells express only cadherin-3.

**Figure 1. Cadherin-1 and cadherin-3 expression during PDAC progression.** (A) Sections of human pancreatic tissue samples were sequentially stained with mouse anti-cadherin-1 and rabbit anti-cadherin-3 Abs. Tissues were then incubated with Alexa 488-conjugated goat anti-mouse and Alexa 594-conjugated Abs and were observed with a confocal microscope. Cell nuclei were stained with Draq5. Scale bar: 25 μm. (B) Pancreas adenocarcinoma and pancreatic intraepithelial neoplasia (PanIN) tissue microarrays, as well as cancer tissue array were immunostained and observed as described in (A). Cadherin-1 and cadherin-3 staining were scored as described in Materials and Methods. Results are expressed as cadherin-3 vs cadherin-1 expression ratio. *P<0.05, **P<0.01, ***P<0.001.
Cadherin-1 and cadherin-3 differentially regulate both invasion and cell migration. We next investigated the effect of cadherins on the cell–cell adhesion properties. BxPC-3 cadh-1⁺/cadh-3⁺ cells and CRCM110-cadh-1⁺/cadh-3⁻ cells spontaneously formed compact spheroids when cultured in suspension (Figure 4A), whereas spheroids formed by BxPC-3-cadh-1⁻/cadh-3⁻ cells or CRCM08-cadh-1⁻/cadh-3⁻ cells were less compact. Quantification of the spheroids areas showed that cadherin-3-depleted spheroids were weakly increased compared with BxPC-3-cadh-1⁺/cadh-3⁻ spheroids. These results indicate that cadherin-1 is the major player in cell–cell adhesion, although cadherin-3 regulates this essential biological phenomenon to a small extent.

The invasive capacity of human pancreatic cell lines expressing either both or one of the cadherins of interest was then analysed. BxPC-3-cadh-1⁺/cadh-3⁻ cells organised in spheroids invaded a three-dimensional type I collagen gel (Figure 4B). Videomicroscopy experiments showed that this kind of invasion mainly required cell–cell contacts (Supplementary Figure S4). Cadherin-1 or cadherin-3 silencing inhibited this intercellular contact dependent cell invasion. Since neither cadherin-1 nor cadherin-3 was able to modulate cell proliferation, we concluded that the decrease in the invasive capacity of the spheroids was not due to the cell proliferation inhibition (data not shown). Similar results were obtained with a type I collagen 2D invasion, an assay that did not require cell–cell contacts (Figure 4C). Interestingly, CRCM110 cells were more invasive than CRCM08 cells that express only cadherin-3. All in all, our data indicate that both cadherin-1 and cadherin-3 take part in intercellular-dependent and -independent invasion through type I collagen.

Cell invasion is a complex and multifactorial process that requires cell adhesion, extracellular matrix proteolysis and cell migration. We therefore set out to dissect the role of each cadherin during these crucial events. Collective cell migration (i.e., cell–cell contact dependent cell migration) was assessed by wound healing assay. While BxPC-3-cadh-1⁺/cadh-3⁻ cells resulted in 30% of closed wound at 6 h, there was a significant increase of cell migration for BxPC-3-cadh-1⁻/cadh-3⁻ cells (Figure 5A). Collective migration properties of cells expressing both cadherin-1 and cadherin-3 or only cadherin-1 were quite similar. These findings suggest that collective migration of BxPC-3 cells is enhanced by cadherin-3. We next analysed single cell migration by using time-lapse microscopy. Cell tracking analysis indicated that cadherin-3 expression increased cell speed, but did not impair cell velocity and cell directionality (Figure 5B).

To determine whether cadherins drive invasive structures formations called invadopodia, cells were plated on FITC-labelled gelatin and gelatinolytic activity was detected by loss of FITC-fluorescence. Cadherin-1, but not cadherin-3 depletion, promoted a reduction of the number of degradation areas formed at the ventral surface of cancer cells (Figure 5C and Supplementary Figure S5). Moreover, invadopodia were more abundant in CRCM110 cells compared to CRCM08 cells. These results strongly suggest that cadherin-1 regulates invadopodia formation.

Cadherin-1 and cadherin-3 differentially regulate tumour microenvironment. To assess the differential effect of cadherins on the primary tumour development, the different BxPC-3 cell lines were subcutaneously implanted into the flank of nude mice. Twenty-one days following cell injections, we analysed tumour growth which was expressed as the product of perpendicular diameters. As shown in Figure 6A, cadherin-3 depletion induced the tumour growth delay. However, cadherin-1 expression did not slow down tumoural growth. The measurement of tumour weights confirmed these observations, as shown in Figure 6A. Since
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Figure 3. Cadherin expression in BxPC-3 cell models and in primary cultures from human tumours. (A) Cadherin-1 or cadherin-3 expression was invalidated in BxPC-3 human pancreatic cancer cell line. Generated stable cell lines were called BxPC-3-cadh-1+/cadh-3− (no cadherin depletion), BxPC-3-cadh-1+/cadh-3− (cadherin-3 depletion) and BxPC-3-cadh-1+/cadh-3− (cadherin-1 depletion). Cadherin-1 and cadherin-3 expression was assessed by immunofluorescence. Scale bar: 15 μm. (B) Cadherin-1 and cadherin-3 were sequentially immunostained in CRCM110 and CRCM08 primary cell cultures derived from human tumours. Scale bar: 15 μm.

According to our findings, the role of both cadherin-1 and cadherin-3 in PDAC progression can be summarised as follows: (1) cadherin-3 is progressively expressed at the plasma membrane during progression of PanIN-1 to PDAC. Despite a decrease in cadherin-1 levels, significant amounts remained associated with the cell surface; (2) cadherin-1 localises with cadherin-3 in pancreatic cancer cells suggesting that these two adhesion molecules could elaborate adhesive networks. The latter may regulate pancreatic cancer cell behaviour; (3) both cadherin-1 and cadherin-3 are involved in the cancer cell invasion through distinct processes: cadherin-3 regulates cell migration, whereas cadherin-1 takes part in the invadopodia activity; (4) cadherin-3 regulates tumour growth. However, when associated with cadherin-3, cadherin-1 expression does not slow down tumoural growth; (5) cadherin-1 drives type I collagen fibres organisation in the tumour.

Several previous studies have shown that cadherin-3 upregulation is associated with aggressive pancreatic tumour behaviour (Taniuchi et al, 2005; Imai et al, 2008; Sakamoto et al, 2015). However, its detection at the early stages of pancreatic cancer development has not been studied. Here, we demonstrated that cadherin-3 seems to be upregulated at an early stage of PDAC. Indeed, by using a human tissue microarray and a mouse model that develops PanIN lesions similar to human (PDX1-Cre, KrasG12D, Ink4a/Arflox/flox), we have shown that cadherin-3 is detected at the cell membrane since PanIN-1 development. Moreover, cadherin-3 expression increased at the cell membrane during progression from PanIN to PDAC and was maximal in metastatic pancreatic cancers. This early cadherin-3 expression is not unique to pancreatic cancer, since an aberrant cadherin-3 expression was also described from the earliest morphologically identifiable stage of colon and breast cells transformation (Vieira and Paredes, 2015). Our results have shown that cadherin-3 is expressed early during the time course of PDAC. Since its expression is associated with poor prognosis in patients (Sakamoto et al, 2015), it is tempting to speculate that cadherin-3 could be used as a diagnostic marker for PDAC. This adhesion molecule therefore represents a promising molecular target in the development of a novel anticancer therapy. It would be essential to test the impact of cadherin-3 antagonising molecules during progression from PanIN to PDAC in order to design effective therapies for patients with this lethal disease.

In several types of cancers, cadherins undergo cleavages from cell membrane by ectodomain shedding (De Wever et al, 2007; Ribeiro et al, 2010). The releasing soluble cadherin (s-cad) may still have functional activity as paracrine/autocrine competitor of cadherin function. Significantly elevated s-cad levels, including s-cadherin-1, s-cadherin-2 and s-cadherin-3, have been observed in the sera and urine of cancer patients diagnosed with a variety of cancers. Moreover, in several types of cancers, increased circulating levels of s-cad are indicative of histopathological grade, metastasis recurrence and poor prognosis (Ribeiro et al, 2010). Further studies are required to determine whether circulating s-cadherin-3 is a potential biomarker for pancreatic cancer. According to this, s-cadherin-3 was identified in proteomic analysis of secretomes
from PDAC samples from 55 patients preserved as xenografts in nude mice (our preliminary results).

The role of cadherin-3 in tumour cell invasion is subject to tumour type and cell context (tissue architecture and microenvironment). There are controversial reports regarding the invasive role of cadherin-3 in different cancer models. Although some studies demonstrated an anti-invasive function of cadherin-3 in melanoma, colon cancer and hepatocarcinoma (Van Marck et al, 2005, 2011; Bauer et al, 2014); other studies have revealed pro-invasive function in bladder, breast, ovarian, intra-hepatic collangioma and pancreatic cancer (Paredes et al, 2004; Taniuchi et al, 2005; Cheung et al, 2010; Van Mark et al, 2011; Sakamoto et al, 2015). In the current study, we showed that cadherin-3 downregulation partially reduced tumour growth in mice. Moreover, cadherin-3 depletion significantly decreased cancer cell invasion through type I collagen in 2D (cell–cell contacts independent) and 3D (cell–cell contacts dependent) invasion assays. The latter appears more physiologically relevant, since instead of using matrigel, we have chosen to work with type I collagen, the major component of the PDAC stroma (Sakamoto et al, 2015). Furthermore, we showed that cadherin-3 is responsible of collective cancer cell migration. According to this, cadherin-3 overexpression in cadherin-3-deficient pancreatic cancer cells induced an increase of cell migration (Taniuchi et al, 2005). It was recently shown that cadherin-3 expression in mesenchymal myoblasts regulates planar cell polarisation and collective cell

Figure 4. Both cadherin-1 and cadherin-3 regulate cancer cell invasion. (A) The impact of cadherins expression on cell–cell adhesion properties was assessed by a spheroid formation assay. BxPC-3-cadh-1+/cadh-3+ (no cadherin depletion), BxPC-3-cadh1− (cadherin-3 depletion) BxPC-3-cadh-3+ (cadherin-1 depletion) cells, as well as primary cultures from human tumours CRCM110 and CRCM08 were allowed to form spheroids in suspension for 24 h. The spheroid area was measured by phase contrast microscopy and analysed by ImageJ. Data represent the mean ± s.d. of three separate experiments performed in triplicates. (B) BxPC-3 cell models were allowed to form spheroids for 72 h. Spheroids were then embedded in type I collagen. After the embedding followed by a 24 h incubation period, the spheroid area was measured by phase contrast microscopy and analysed by ImageJ. Data represent the mean ± s.d. of three independent experiments performed in triplicates. (C) Isolated BxPC-3 cells and primary cultures from human tumours CRCM110 and CRCM08 were allowed to invade a layer of type I collagen for 24 h. The invasion index was calculated as the mean number of migrated cells counted in 10 microscopic fields. Data represent the mean ± s.d. of three independent experiments performed in triplicates. **P<0.01, ***P<0.001.
migration (Plutoni et al, 2016). Altogether, these data confirm that cadherin-3 overexpression in PDAC is associated with the invasive phenotype and malignant behaviour.

It is well established that cadherin-1 plays an important role as an invasion suppressor (Rodriguez et al, 2012). However, recent studies suggested that this molecule may also have a promoting role in ovarian, breast and glioma cancer progression (Rodriguez et al, 2012). Nonetheless, much remains to be learned about its role in tumour progression. In the present study, we have demonstrated that cadherin-1 participates in PDAC progression. Indeed, by using BxPC-3 cell models and primary culture from human tumours, we showed that cadherin-1 (1) regulates cell invasion through type I collagen 3D and 2D invasion assays; (2) increases motility of isolated cells; (3) does not affect tumour growth in mice.

It is now clear that a switching from cadherin-1 to cadherin-2 and/or cadherin-3 is a hallmark of epithelial–mesenchymal transition (EMT) in several types of cancer including prostate, ovarian and bladder cancers. By using double immunostaining experiments on human and murine tissue specimens, we have shown that cells co-expressed at their surface both cadherin-1 and cadherin-3 from early PanIN lesions to PDAC. These data indicate that in PDAC cadherin switching refers to a situation in which cadherin-1 levels do not change significantly, but cells turn on cadherin-3 expression. Moreover, the co-expression of cadherin-1 and cadherin-2 was also described in a mouse model of K-ras-
induced PanIN (Su et al., 2016). Taken together these findings suggest that cadherin switching in PDAC is not required for the EMT process.

Herein, we have observed that simultaneous expression of cadherin-1 and cadherin-3 is the most frequent situation in PDAC. Indeed, double immunostaining analysis on a human tissue microarray revealed that these adhesion molecules are jointly expressed at the cell surface of tumoural cells in a large proportion of PDAC (41%). These results were confirmed by using gene transcription and protein expression analysis on 55 samples of PDAC from patient preserved as xenografts in nude mice. According to this, same simultaneous expression of cadherins was described in a quarter of breast carcinomas (Ribeiro et al., 2013). Based on these data, we propose that the co-expression of cadherin-1 and cadherin-3 could be a signature of an increased invasive and tumourigenic potential in PDAC.

PDAC is characterised by a dense desmoplastic stroma with type I collagen as the major component (Feig et al., 2012). However, the process of driving collagen deposition and organisation in PDAC remains unknown. Our results suggest that different type I collagen organisation could be generated depending on the type of cadherin expressed by tumoural cells. Indeed, orthotopic and ectopic injection of BxPC-3 cells in nude mice revealed that malignant cells expressing cadherin-1 drive type I collagen deposition. Moreover, Masson’s trichrome staining of PDAC samples from xenografted tumours of patients confirmed that cadherin-1-positive cells promoted a strong collagen deposition. Therefore, rather than collagen abundance, the architecture and mechanics of collagen fibres adjacent to epithelial lesion may be an indicator of PDAC progression (Laklai et al., 2016). Further studies are required to determine whether the nature of the cadherin(s) expressed by tumour cells alters the organisation of collagen bundles.

Extracellular matrix is a key physical barrier to cancer cell invasion. It is now well known that focal extracellular matrix degradation and remodelling are required to open space and create tracks along which metastatic tumour cells can migrate through tissue. Invadopodia, specialised proteolytically active plasma membrane protrusions, are crucial for this mechanism (Gimona et al., 2008). Extracellular matrix proteolysis mediated by invadopodia is mainly executed by matrix-degrading proteases including membrane type-1 matrix metalloproteinase (MT1-MMP), as well as by matrix metalloproteinase-2 (MMP-2) and -9 (MMP-9) (Castro-Castro et al., 2016). These proteases are overexpressed in metastatic PDAC lesions compared with primary tumours (Ottaviano et al., 2006; Jakubowska et al., 2016). Our
studies revealed that pancreatic cancer cell line BxPC-3 and primary culture from human tumours are able to develop invadopodia to degrade extracellular matrix. Indeed in this assay cadherin-1-deficient cells (BxPC-3-cadh-1+/cadh-3þ and CRCM08-cadh-1+/cadh-3þ) have a reduced focal gelatinolytic activity. We have therefore concluded that the formation of invadopodia involves cadherin-1 expression. The mechanism by which cadherin-1 drives invadopodia formation remains unknown, but needs to be further elucidated. Indeed, clarifying how cadherin-1 signalling regulates extracellular matrix degradation should improve our understanding of invadopodia lifecycle in order to target them and prevent cancer cell invasion and metastasis.

In summary, our data demonstrate that pro-tumoural signalling environments can be generated according to the type of cadherin expressed by cancer cells, which can lead to differential cell responses including invasion, migration or regulation of extracellular matrix deposition and degradation. Importantly, these findings open new avenues towards uncovering innovative options for earlier diagnosis and anti-invasive therapy of pancreatic cancer.

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Figure 7. Cadherins alter ECM deposition. Mice were orthotopically implanted with either BxPC-3-cadh-1+/cadh-3+, BxPC-3-cadh1+ or BxPC-3-cadh-3+ cells resuspended in Matrigel or with Matrigel alone (n = 3 for each condition). (A) Mice were sacrificed 21 days after injection, pancreas were collected and the weight was measured. (B) Sections of graft tissue specimens were subjected to Masson’s trichrome staining. (C) The stromal area of each tumour was measured by microscopy (10 × objective) and analysed by ImageJ. ***P < 0.001.

CONFLICT OF INTEREST

The authors declare no conflict of interest.

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