



GSK3 β as a novel promising target to overcome chemoresistance in pancreatic cancer

Camilla Pecoraro^{a,b,1}, Beatrice Faggion^{a,1}, Beatrice Balboni^{a,c}, Daniela Carbone^b, Godefridus J. Peters^{a,d}, Patrizia Diana^b, Yehuda G. Assaraf^e, Elisa Giovannetti^{a,f,*}

^a Department of Medical Oncology, Amsterdam University Medical Center, VU University, 1081 HV Amsterdam, the Netherlands

^b Department of Biological, Chemical and Pharmaceutical Sciences and Technologies (STEBICEF), University of Palermo, Palermo, Italy

^c Computational and Chemical Biology, Istituto Italiano di Tecnologia, Via Morego 30, 16163 Genoa, Italy, and Department of Pharmacy and Biotechnology, University of Bologna, Via Belmeloro 6, 40126 Bologna, Italy

^d Department of Biochemistry, Medical University of Gdansk, Poland

^e The Fred Wyszowski Cancer Research Laboratory, Department of Biology, Technion-Israel Institute of Technology, Haifa, 3200003, Israel

^f Cancer Pharmacology Lab, Fondazione Pisana per la Scienza, Via Ferruccio Giovannini 13, 56017 San Giuliano Terme (Pisa), Italy

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ABSTRACT

Pancreatic cancer is an aggressive malignancy with increasing incidence and poor prognosis due to its late diagnosis and intrinsic chemoresistance. Most pancreatic cancer patients present with locally advanced or metastatic disease characterized by inherent resistance to chemotherapy. These features pose a series of therapeutic challenges and new targets are urgently needed.

Glycogen synthase kinase 3 beta (GSK3 β) is a conserved serine/threonine kinase, which regulates key cellular processes including cell proliferation, DNA repair, cell cycle progression, signaling and metabolic pathways. GSK3 β is implicated in non-malignant and malignant diseases including inflammation, neurodegenerative diseases, diabetes and cancer. GSK3 β recently emerged among the key factors involved in the onset and progression of pancreatic cancer, as well as in the acquisition of chemoresistance. Intensive research has been conducted on key oncogenic functions of GSK3 β and its potential as a druggable target; currently developed GSK3 β inhibitors display promising results in preclinical models of distinct tumor types, including pancreatic cancer.

Here, we review the latest findings about GSK-3 β biology and its role in the development and progression of pancreatic cancer. Moreover, we discuss therapeutic agents targeting GSK3 β that could be administered as monotherapy or in combination with other drugs to surmount chemoresistance. Several studies are also defining potential gene signatures to identify patients who might benefit from GSK3 β -based therapeutic intervention. This detailed overview emphasizes the urgent need of additional molecular studies on the impact of GSK3 β inhibition as well as structural analysis of novel compounds and omics studies of predictive biomarkers.

1. Introduction

Pancreatic cancer is currently the third most common cancer in Western countries, with an increasing incidence and poor outcome and constitutes one of the most lethal of the common malignancies with a poor five-year survival rate below 10 % (Hill and Chung, 2020; Siegel et al., 2021; Rahib et al., 2014; Coppola et al., 2017; Binenbaum et al., 2015)

Pancreatic ductal adenocarcinoma (PDAC) accounts for approximately 90 % of pancreatic tumors (Sarantis et al., 2020). This

malignancy is among one of most inadequately understood human disorders, posing a significant diagnostic and therapeutic challenge. The lack of specific symptoms and reliable biomarkers for early detection screening of asymptomatic PDAC patients, results in a most dismal prognosis (Kaur et al., 2017; Zhou et al., 2017). In this respect, approximately 52 % of PDAC patients are diagnosed with an advanced-stage or metastatic disease, for which the 5-year survival trend is as low as 3% (Giovannetti et al., 2017; Supadmanaba et al., 2021).

The aggressive nature and the early metastatic behavior of PDAC are

* Corresponding author at: Department of Medical Oncology, Amsterdam University Medical Center, VU University, 1081HV Amsterdam, the Netherlands.

E-mail address: elisa.giovannetti@gmail.com (E. Giovannetti).

¹ Co-first authors.

not the only responsible factors for the poor prognosis of this disorder, but also for its insensitivity to most therapies, such as chemotherapy, radiotherapy and immunotherapy. Among all clinical intervention, surgical resection remains the mainstay chance for cure. However, less than 20 % of patients are good candidates for pancreatectomy due to the usually diagnosed metastatic state of PDAC (Giovannetti et al., 2017). Additionally, chemotherapeutic and radiotherapeutic regimens are often palliative and their high toxicity provides a very marginal improvement in the survival rate of patients with advanced disease (Zeng et al., 2019). This poor treatment efficacy is accompanied by either intrinsic resistance or rapid acquisition of chemoresistance (Arora et al., 2013; Caparello et al., 2016). Thus, despite the advances in the use of combination chemotherapeutic regimens, survival remains dismal, highlighting the tremendous urgency for the design and development of novel therapeutic strategies to overcome the chemoresistant nature of this lethal disease.

In recent years, GSK3 β has emerged as a new potential target in PDAC due to its involvement in promoting neoplastic transformation, tumor cell survival and chemoresistance (Cormier and Woodgett, 2017; Ding and Billadeau, 2020; Uehara et al., 2020). Clinical trials are currently testing several GSK3 β inhibitors either as monotherapy or in combination with chemotherapeutic agents with the aim of developing promising PDAC therapeutic interventions that suppress PDAC growth and prevent disease progression (Garcia-Sampedro et al., 2021; Abrams et al., 2021).

The purpose of the current review is to determine whether or not GSK3 β might be considered a good therapeutic target in advanced PDAC and which patient signatures might be prognostic of good therapy response. Furthermore, it will focus on GSK3 β inhibitors that are currently approved or are undergoing clinical trials. We also discuss possible drug combinations that might prevent tumor recurrence and therapy resistance.

2. Glycogen synthase kinase 3 (GSK3)

2.1. GSK3 β biology in normal cells

Glycogen synthase kinase 3 (GSK3) is a family of serine-threonine kinases which encompasses two highly conserved isoforms, GSK3 α and GSK3 β , sharing approximately 85 % overall sequence homology (Woodgett, 1990). Even if functional redundancy has been observed within the two isoforms, most studies in the oncology field focused on GSK3 β activity, mainly due to its known enigmatic effects on many physiological and pathological processes. By phosphorylating serine and threonine residues of a broad range of functional and structural proteins, GSK3 β regulates many fundamental biological processes in cells such as glycogen metabolism, Wnt/ β -catenin signaling, G-protein-coupled

signal transduction and maintenance of stem cell identity (Cormier and Woodgett, 2017; Doble and Woodgett, 2003; Gao et al., 2013; Xu et al., 2009; Wu and Pan, 2010; Riobó et al., 2006). The large number of GSK3 β substrates explains its emblematic function as tumor promoter or tumor suppressor. Those roles have already been extensively summarized in many recent review articles (Sutherland, 2011; McCubrey et al., 2016; Duda et al., 2020; Xie and Wang, 2017).

The most common targets of GSK3 β are primed substrates harboring a pre-phosphorylated sequence S/T-X-X-S/T(P). Specifically, this provides the binding site for GSK3 β , inducing a functional conformation change that assists the target positioning in the active catalytic domain of the kinase (Fig. 1) (ter Haar et al., 2001; Dajani et al., 2003). Hence, the kinase activity of GSK3 β leads to either suppression and proteasomal degradation or enhanced activation and protein stabilization of target substrates.

Notably, GSK3 β has the unconventional characteristic for a kinase of being normally active in cells under resting conditions. This is mainly correlated with phosphorylation of its tyrosine (Y)216 residue, which induces a conformational change that allows the interaction and phosphorylation of protein targets (Hughes et al., 1993; Kaidanovich-Beilin and Woodgett, 2011). On the other hand, extracellular signals, negatively regulate GSK3 β kinase activity via N-terminal phosphorylation of the serine (S)9 residue which is required for the maintenance of normal cell homeostasis (Sutherland, 2011; Frame et al., 2001). Crystal structure analysis revealed that phosphorylation of the inhibitory serine-9 residue causes the self-association of the GSK3 β N-terminal tail to its substrate binding pocket, thus hampering the interaction with target substrates (Frame et al., 2001; Stamos et al., 2014).

While there are multiple mechanisms that modulate GSK3 β activity, they have not been yet completely elucidated due to the highly complex interconnections with several molecular signaling cascades. Interestingly, consistent experimental evidence reported that various regulatory protein kinases such as Akt, cyclic adenosine monophosphate (cAMP)-dependent, protein kinase A (PKA), p70 S6 kinase (p70S6K), p90RSK and Notch3, increase the inhibitory GSK3 β serine-9 phosphorylation in response to extracellular signals (Fig. 1) (Kaidanovich-Beilin and Woodgett, 2011; Fang et al., 2000; Foltz et al., 2002). In addition, growth factors such as EGF, PDGF and insulin inhibit GSK3 activity via induction of the phosphatidylinositol-3-kinase (PI3K)/MAPK pathway. Other mechanisms that alter the inhibitory phosphorylation status of GSK3 β are represented by elevated intracellular cAMP levels mediated by PKA and amino acid deprivation caused by mTOR signaling (Fig. 1) (Krause et al., 2002; Li et al., 2000).

Next to the phosphorylation status of GSK3 β , which dynamically oscillates in response to extracellular signals and substrate availability, other varying and controversial mechanisms regulate GSK3 β kinase activity such as GSK3 β localization and protein-complex formation

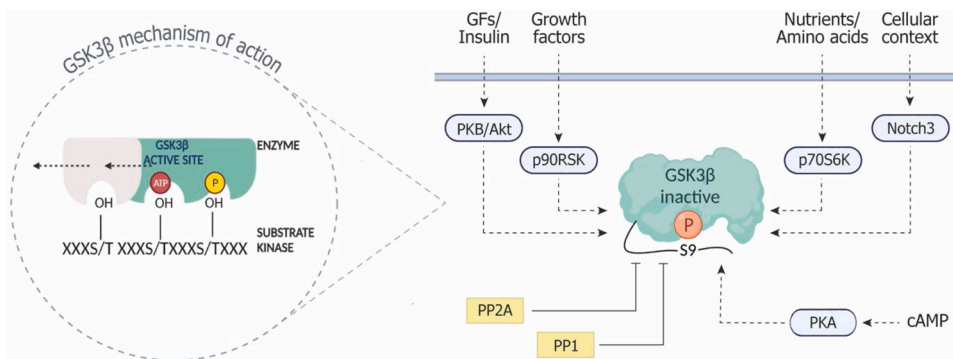


Fig. 1. Mechanism of action of GSK3 β with a special focus on mechanisms regulating GSK3 β activity by phosphorylation of N-terminal Serine-9. A. GSK3 β recognizes a specific amino acid sequence motif S / T-XXX-S / T (P), in which S represents a serine, T a threonine, X a generic amino acid and P indicates the presence of a phosphate group previously bound by another protein kinase which is called kinase primer. The presence of the phosphorylated residue in the recognition sequence allows the substrate to position itself at the active site, thus placing the S/T residue of the target sequence near the kinase site, allowing its phosphorylation. B. Extracellular signals lead to the activation of transduction cascades that result in the phosphorylation of the serine-9 (S9) residue

which blocks the target substrate binding and inactivate the kinase activity of GSK3 β . Kinase phosphorylating S9 residues are represented with blue ovals and dashed lines. Protein phosphatases PP2A and PP1 restore GSK3 β catalytic activity.

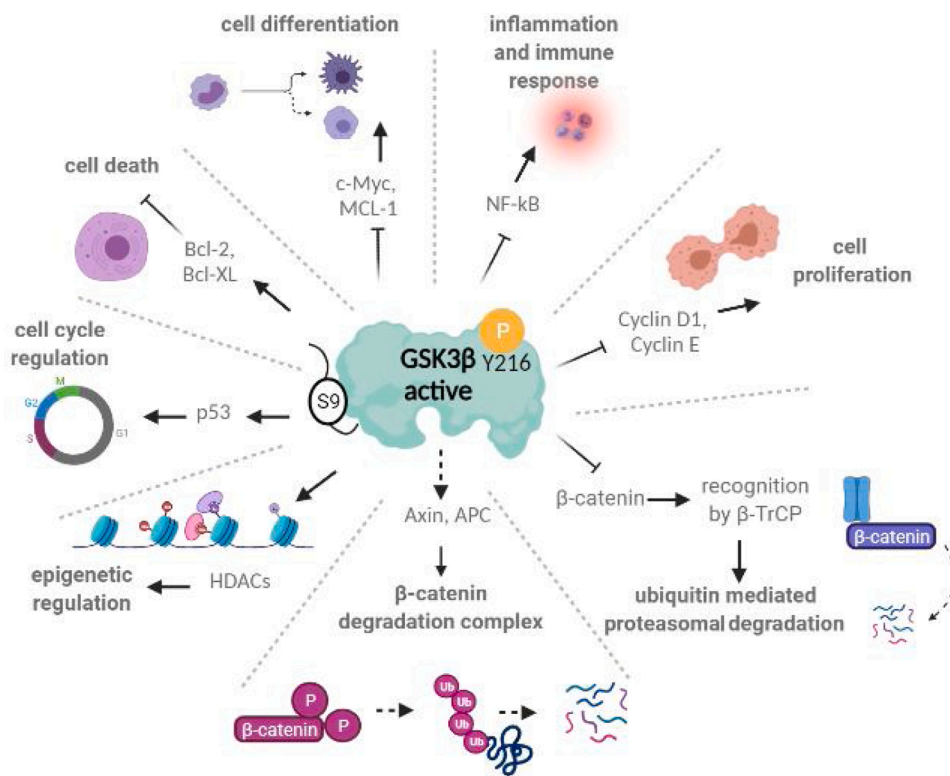


Fig. 3. Multiple roles of GSK3 β and its target substrates in key biological processes for cancer cells. GSK3 β modulates the activity of many cellular substrates involved in cell cycle progression, cell proliferation and differentiation (p53, c-Myc, MCL-1, Cyclin D1, Cyclin E). GSK3 β regulates NF- κ B and CREB (not shown) transcription factors affecting inflammatory and immune responses. In the absence of Wnt ligands, the Axin-APC-CK1-GSK3 β - β -catenin destruction complex allows GSK3 β to phosphorylate β -catenin (residues 41, 37, and 33). This phosphorylation leads to the release of β -catenin from the complex and targets it for proteasomal degradation. GSK3 β phosphorylates several histone deacetylases (HDACs) modulating their regulatory epigenetic functions. GSK3 β -mediated phosphorylation of some molecules of the anti-apoptotic Bcl-2 family results in their stabilization and increased anti-apoptotic effects.

Additionally, recent findings showed that GSK3 cooperates with mTOR to regulate the activity of p70 ribosomal protein S6 kinase 1 (S6K1), which is a pivotal regulator of extracellular signals supporting cell growth (Shin et al., 2011). Other outstanding mechanisms sustaining cancer cell survival and proliferation involve modulation of c-Myc signaling, Hedgehog (Hh) signaling and STAT3 cascade, but further research is required in the context of PDAC (Baumgart et al., 2016; Singh et al., 1995).

Moreover, the WNT/ β -catenin signaling pathway plays an important role in the modulation of apoptosis, differentiation, invasion and epithelial to mesenchymal transition, all critical hallmarks for cancer metastasis (Doble and Woodgett, 2007). GSK3 β is a well-established regulator of β -catenin subcellular localization and deregulated GSK3 β activity may severely impact the tumor-suppressive and tumor-promoting roles of the WNT/ β -catenin signaling cascade (Domoto et al., 2016). Consistently, pharmacologic inhibition of GSK3 β was shown to upregulate β -catenin and c-Myc levels, as well as suppress tumor growth in KRas-mutant PDAC and non-small lung cancer models (Kazi et al., 2018).

Interestingly, it has been recently observed that GSK3 β plays a key role in modulating cell cycle progression at different regulatory levels. In pancreatic cancer models, GSK3 β was observed to directly support the phosphorylation status of many cell cycle modulators such as cyclin D1, p53 and various transcription factors (Kitano et al., 2013; McCubrey et al., 2016). The growth promoting function of GSK3 β in this tumor type is supported by both *in vitro* and in tumor xenograft experiments *in vivo*: specifically, pharmacological inhibition of GSK3 β activity was reported to promote apoptosis by suppressing Cyclin D1 expression, as well as impairing the transcriptional activity of E2F transcription factor and consequent phosphorylation of the Rb protein (Kitano et al., 2013). Furthermore, Yoshino and colleagues observed that apoptosis-resistant PDAC cells treated with GSK3 β inhibitors, exhibited biodynamic cell mechanisms typical of mitotic catastrophe (Yoshino and Ishioka, 2015). This study provided the first proof of fundamental role of GSK3 β in controlling mitotic processes in PDAC cells. Similar observations were also recently made in colorectal cancer cells (Yoshino and Ishioka, 2015;

Dewi et al., 2018).

Overall, within the complexity of the GSK3 β signaling cascades in pancreatic tumorigenesis and tumor progression, cumulative evidence renders GSK3 β a promising therapeutic target in order to improve the survival of PDAC patient and enhance the therapeutic responses.

2.3. Chemoresistance in advanced or metastatic pancreatic cancer

Surgical resection of PDAC remains the curative treatment choice for achieving long-term survival. However, more than 85 % of PDAC patients are diagnosed with advanced-stage or metastatic disease which is not generally amenable to pancreatectomy (Meijer et al., 2020).

The standard-of-care intervention in those tumors with aggressive-stages mainly include chemotherapeutic agents, such as gemcitabine, 5-fluorouracil, oxaliplatin, irinotecan and nab-paclitaxel (El Hassouni et al., 2019). However, after a good initial response of sensitive tumors, overt chemoresistance eventually develops within weeks, thus severely limiting the effectiveness of those therapeutic interventions (Zeng et al., 2019).

Particularly, PDAC cells showed stronger intrinsic or acquired insensitivity to gemcitabine. Recent phase III clinical trials reported superior overall survival in patients with advanced or metastatic PDAC receiving nab-paclitaxel plus gemcitabine or FOLFIRINOX compared to gemcitabine alone (Caparello et al., 2016). However, although these treatments achieved a longer survival rate than gemcitabine use alone, combinations of chemotherapeutic agents generally show higher frequency of toxicity and patients still become resistant after short times (Pusceddu et al., 2019).

Therefore, gaining insights into tumor intrinsic or acquired strategies responsible for chemotherapy resistance are of urgent need in order to develop novel targeted therapeutic approaches that improve patient overall survival, lower toxicity profile and overcome chemoresistance in advanced or metastatic PDAC (Zeng et al., 2019).

2.3.1. GSK3 β and chemoresistance

Even though the underlying mechanisms which trigger

chemoresistance remain controversial, defective pharmacodynamics of biochemical mechanisms, together with perturbations on several cellular signaling cascades, were found in gemcitabine-resistant PDAC cells. These mainly involve nucleoside transport and metabolism, reactivation of EMT and developmental pathways, such as WNT/ β -catenin, Hh and Notch, and growth factor signaling (El Hassouni et al., 2019; Randazzo et al., 2020; Saiki et al., 2012; Ireland et al., 2016; Shukla et al., 2017).

As previously described, mounting evidences report that aberrant activation of NF- κ B plays a crucial role in uncontrolled cell proliferation, tumorigenesis, metastasis, angiogenesis, inflammation and chemotherapy resistance in PDAC (Arlt et al., 2003; Okamoto et al., 2007; Liptay et al., 2003; Holcomb et al., 2008; Mamaghani et al., 2009). Specifically, a strong correlation was observed between basal level of NF- κ B activity and gemcitabine resistance. Indeed, in resistant PDAC cell lines (PancTu-1, Capan-1 and BxPc-3 cells) a strong activation of NF- κ B was detected when compared to sensitive T3M4 and PT45-P1 PDAC cell lines (Arlt et al., 2003).

NF- κ B contribution to chemoresistance was further confirmed by pharmacologic inhibition and by targeting I κ B α super-repressor or GSK3 β , which resulted in increased sensitivity to gemcitabine in non-responsive PDAC cell lines (Arlt et al., 2003). Mechanistically, GSK3 β has been shown to positively regulate NF- κ B maintaining high NF- κ B activity, thus evidencing GSK3 β contribution to chemoresistance onset in PDAC (Ougolkov et al., 2006, 2005; Walz et al., 2017).

PI3K/Akt-mediated signal transduction is another important anti-apoptotic signaling cascade that has been related to chemoresistance of PDAC. Akt is the primary mediator of PI3k-initiated signaling cascade and is specifically involved in the phosphorylation and subsequent inactivation of pro-apoptotic molecules such as Bad, I κ B (I κ B kinase), caspase-9, the forehand family of transcription factors (FKHR/AFX/FOX), CREB, Raf, p21, as well as GSK3 (Massihnia et al., 2017). A recent study reported that activation of Akt and inhibition of GSK3 β through Akt-mediated Serine 9 phosphorylation resulted in the upregulation of Snail1 expression through increased protein stability, promoting EMT-like phenotype and gemcitabine tolerance (Namba et al., 2015).

In order to define the stepwise processes triggering gemcitabine resistance in the clinic, Uehara and colleagues developed a gemcitabine-resistant systemic model derived from the gemcitabine-sensitive human PDAC cell line BxPC-3 (Uehara et al., 2020). Through their work, they demonstrated that GSK3 β facilitates the capacity of PDAC to tolerate chemotherapy by interfering with Rb protein function and E2F transcription factor (E2F)1-mediated transcription. Specifically, gemcitabine-resistant clones were characterized by high expression of ribonucleotide reductase M1 (RRM1), a well-known transcriptional target of the pro-oncogenic E2F1 (Uehara et al., 2020; Yoneyama et al., 2015). Additionally, pharmacological inhibition of GSK3 β was proved to re-sensitize resistant cells to gemcitabine by restoring the functional Rb-mediated regulation of E2F1, attenuating E2F1 transcriptional activity and consequently decreasing RRM1 expression (Uehara et al., 2020; Kitano et al., 2013).

In summary, GSK3 β seems to be involved in many different processes that promote resistance of PDAC cells to gemcitabine and other drugs by sustaining the invasive capacity and stemness phenotype of pancreatic malignant cells.

2.4. Targeting GSK3 β in pancreatic cancer

GSK3 β regulation of several molecular promoters of neoplastic transformation, together with the shorter survival of PDAC patients harboring high GSK3 β expression, strongly sustain the hypothesis of GSK3 β central involvement in PDAC onset and progression (McCubrey et al., 2016). Thus, accumulating evidence on GSK3 β functions has proven the rationale for the clinical development of novel therapeutic strategies targeting GSK3 β in advanced PDAC (Cormier and Woodgett, 2017; Baumgart et al., 2016; Walz et al., 2017; Hoeflich et al., 2000).

The therapeutic and antitumor effects achieved by GSK3 β inhibition have been described in different cancer types, as reviewed recently by Domoto and colleagues (Domoto et al., 2020). Notably, some evidence reveals that GSK3 β inhibition leads to induction of apoptosis in PDAC cells, whereas normal pancreatic epithelial cells seem to be protected from the inhibitory effects associated with the targeted treatment (Marchand et al., 2012). This might find an explanation in the intrinsic biological nature of PDAC cells which present high levels of active GSK3 β aberrantly accumulated in the nucleus (Ougolkov et al., 2006; Walz et al., 2017). Thus, PDAC may become more sensitive to the proteasome-dependent GSK3 β loss from the nucleus that is induced by the treatment with GSK3 inhibitors (Marchand et al., 2012, 2015).

Inhibition of GSK3 β may therefore be a promising precision medicine strategy in PDAC treatment (Domoto et al., 2020; Baudino, 2015). Through cancer cell death promoting effects, this approach may restrict tumor recurrence and metastasis as well as spare harmful consequences on healthy cells and tissues, frequently associated with conventional cytotoxic therapies.

2.4.1. GSK3 β inhibitors

Multiple GSK3 β inhibitors have been developed and many others are now under investigation (Table 1), as previously reviewed by Saraswati AP et al., and Eldar-Finkelman H et al. (Saraswati et al., 2018; Eldar-Finkelman and Martinez, 2011). In the early 1980s, the cation lithium was the first GSK3 β inhibitor being described and then approved by the Food and Drug Administration (FDA) for the treatment of human bipolar depression (Johnson and Amdisen, 1983; Freland and Beaulieu, 2012; Bowden, 2000).

Studies on the mechanism of action of lithium showed that it disrupts the catalytic function of GSK3 β by competing for the binding of magnesium (Mg^{2+}) cofactor, but not for ATP or the substrate (Ryves and Harwood, 2001; Phiel and Klein, 2001; Pasquali et al., 2010). Additionally, lithium indirectly increased the N-terminal inhibitory phosphorylation of GSK3 β either by enhancing the activity of Akt, through the regulation/dissociation of the Akt: β -arrestin 2 (β Arr2): protein phosphatase 2A (PP2A) signaling complex, or by blocking the dephosphorylation of PKB (Pasquali et al., 2010; Zhang et al., 2003; Mora et al., 2002; De Sarno et al., 2002).

Experimental studies on PDAC cells reported reduced tumorigenic potential and cell proliferation, as well as increased apoptosis among lithium treatment (Peng et al., 2013). This outcome was associated with the enhanced ubiquitin-dependent proteasome degradation of the glioma-associated oncogene-1 (GLI1), a crucial downstream component of the Hh signaling pathway, following GSK3 β inhibition (Peng et al., 2013; Zhu and Lo, 2010).

The ATP-binding pocket is an essential site for the catalytic action of GSK3 β ; thus, various GSK3 β inhibitors competing with ATP molecules have been developed in order to block the kinase activation. However, one has to note that, whereas those agents might selectively target GSK3 β , they can also exert inhibitory effects towards cyclin-dependent kinases (CDKs) since some of them including CDK2, share very similar ATP-binding pockets with GSK3 β (Vulpetti et al., 2005). Therefore, the dual inhibitory function of ATP-competitive GSK3 β inhibitors may directly impact cell cycle regulation and enhance the antiproliferative outcome.

Gaisina and colleagues designed a preliminary library of benzofuran-3-yl-(indol-3-yl) maleimides, including some selective and potent ATP competitive inhibitors of GSK3 β . *In vitro* evaluation of the therapeutic potency of these maleimides showed that compounds 1a-e exhibited promising antiproliferative effects against a panel of PDAC cell lines (MiaPaCa-2, HupT3 and BXPC2) (Table 1) (Gaisina et al., 2009). Among all, treatment with compounds 1a and 1e resulted in pronounced inhibition of GSK3 β activity, which correlated with reduced NF- κ B-mediated expression of the antiapoptotic X-linked inhibitor of apoptosis protein (XIAP).

Recently, 1e (9-ING-41), maleimide-based ATP-competitive GSK3 β

Table 1
GSK3 β Inhibitors in pancreatic cancer preclinical and clinical studies.

Category	Inhibitor	Structure	Pharmacologic activity	GSK3 β inhibitory effects	Ref/Clinical trial
ATP competitive	benzofuran-3-yl-(indol-3-yl) maleimides		Potent ATP competitor	Apoptosis induction, reduced NF- κ B-mediated expression of XIAP	Gaisina IN et al., 2009 Ding L et al., 2017
	9-ING-41		Potent ATP competitor	Cell cycle arrest, reduced expression of anti-apoptotic molecules Bcl-2 and XIAP	NCT03678883, 1801 phase 1/2 study
	SB-732881-H		Potent ATP competitor	Apoptosis induction, increased expression of the Bcl-2 protein family	Marchand B et al., 2012
	AR-A014418		Potent ATP competitor	Lowers cytoplasmic β -catenin levels and abrogates NF- κ B transcriptional activation	Bhat et al. (2003) Mamaghani S et al., 2009
	BIO		Reduced inhibitory S9 phosphorylation	Enhanced apoptosis via JNK-dependent mechanism	Meijer L. et al., 2003 Kazi A et al., 2018
	CHIR99021		Potent ATP competitor	Apoptosis induction	Marchand B et al., 2012 NCT01287520, 2018
	LY2090314		Increased inhibitory phosphorylation	Suppressed pro-survival signals	NCT01632306, 2019
	AZD-1080		Potent ATP competitor	Reduced cell cycle progression related genes	Kazi A et al., 2018
	Library of synthetic topoisomerase II inhibitors		Docked in the ATP binding site	Pro-apoptotic signals induction, reduced expression of EMT markers	Carbone D et al., 2021
	Non-ATP competitive	Tideglusib		Binding site not yet defined	Cell cycle arrest, impaired phosphorylation of β -catenin and c-Myc
Lithium		Li^+	Compete for Mg^{2+} cofactor binding	Suppressed hedgehog signaling pathway	Peng Z et al., 2013
Dual inhibitor	Metavert		Inhibition of GSK3 β and HDAC-2	Cell cycle arrest, reduced expression of EMT markers	Edderkaoui M et al., 2018

inhibitor, showed cytostatic effects in PDAC models and is now under investigation in a phase I/II clinical trial (NCT03678883) in patients with advanced solid tumors ([Ding and Billadeau, 2020](#); [NCT03678883, 2021](#); [Carneiro et al., 2020](#)). Even if available results reported a good

9-ING-41 monotherapy tolerance and antitumor efficiency more promising therapeutic effects were obtained under combination chemotherapeutic regimens in patients with refractory PDAC ([Carneiro et al., 2020](#); [Ding et al., 2019](#)).

Increased sensitivity to chemotherapy or chemoresistance regression upon treatment with 9-ING-41 have been newly discovered to be related to GSK3 β regulation of the gemcitabine-induced TopBP1/ATR/Chk1 DNA damage response pathway, as will be discussed later (Ding et al., 2019).

SB-732881-H (SB), a dual inhibitor of both GSK3 isoforms, selectively suppressed the viability of mutant KRas-dependent tumor cells (Kazi et al., 2018; Fleming et al., 2005). *In vitro* studies on human PDAC cells MiaPaCa2 harboring aberrant KRas, revealed high sensitivity to SB treatment (IC₅₀ of 0.4 μ M), resulting in caspase-3 activation and induction of PARP cleavage (Kazi et al., 2018; Demarchi et al., 2003). Furthermore, additional reproducible data suggested that the antitumor effect of SB on KRas-mutant PDAC is supported by apoptosis induction in a c-Myc- and β -catenin-dependent manner (Kazi et al., 2018). In contrast, pharmacologic activity of SB did not affect the viability of non-malignant pancreatic epithelial cells. Overall, these findings are consistent with the pro-survival function of GSK3 in PDAC and further support the assumption that mutant KRas tumors are dependent on GSK3 α/β signalling for cancer cell survival and tumor growth (Kazi et al., 2018; Saiki et al., 2012; Fleming et al., 2005; Demarchi et al., 2003; Bang et al., 2013).

AR-A014418, another dual GSK3 inhibitor, induced a strong dose-dependent growth reduction in various pancreatic tumor models (Bhat et al., 2003; Kunnimalaiyaan et al., 2015). By selectively competing for the ATP-binding pocket of GSK3 β , AR-A014418 lowered the cytoplasmic β -catenin levels and abrogated NF- κ B activation, thus reducing the expression of NF- κ B target genes cyclin D1, XIAP and Bcl-XL (Mamaghani et al., 2009; Bhat et al., 2003). Moreover, a recent *in vitro* study reported that AR-A014418 inhibition of GSK3 α phosphorylation decreased the expression of Notch pathway members, thus attenuating tumor cell survival (Kunnimalaiyaan et al., 2015).

6-bromindirubin-3'-oxime (BIO), a synthetic analog of natural indirubins, has been studied in the context of drug resistance in many different cancer types for its indirubin-related property to concomitantly inhibit CDKs and GSK3 β (Zhang et al., 2017; Liu et al., 2017; Li et al., 2018;), *via* interaction with the ATP-binding pocket of both kinases (Meijer et al., 2003).

Studies on beneficial effects of BIO have been conducted in the context of anti-aging properties, where treatment with 6-bromindirubin-3'-oxime reported ameliorated lipid metabolism and positive modulation of autophagy, inflammation and oxidative stress (Guo et al., 2019).

In PDAC cells, treatment with BIO directly inhibited GSK3 β *via* suppression of Tyr216 phosphorylation and enhanced apoptosis through JNK-dependent mechanisms (Marchand et al., 2015). However, Marchand and colleagues observed that BIO-GSK3 β inhibition augmented the activation of the autophagy/lysosomal network which was elicited through enhanced nuclear localization of the transcription factor EB (TFEB), a master regulator of autophagy and lysosomal biogenesis (Marchand et al., 2015; Zhitomirsky and Assaraf, 2016; Zhitomirsky et al., 2018; Zhitomirsky and Assaraf, 2015). Although the autophagy effects on cancer cells are not well defined, it has been reported that sustained autophagy in PDAC cells elicits cell proliferation by overcoming death signals and favoring oxidative phosphorylation (Yang and Kimmelman, 2011). This metabolic switch provides a proper bioenergetic metabolism and pivotal survival signals to malignant cells under restrictive growth conditions, thus supporting cancer cell growth and tumor progression. Overall, these undesired modulatory effects exerted by BIO, hampered its progression into clinical trials.

Comparably to BIO, the aminopyrimidine derived GSK3 inhibitor CHIR99021 exhibited both apoptosis induction and concomitant increased autophagic response *via* LC3B II expression in PDAC cell models (Tran and Zheng, 2017). Pharmacologic depletion of vacuolar H⁺ ATPase with bafilomycin A1, prevented autophagy by disrupting lysosomal acidification, thus forcing PDAC cells to preferentially respond to the death signals mediated by CHIR99021. Indeed, this study

supported the hypothesis of addressing autophagy induction as a promising mechanism of escape to the antiproliferative effects of GSK3 inhibitors in the setting of PDAC treatment.

Conversely, LY2090314, an ATP-competitive and highly selective GSK3 inhibitor, is currently under clinical evaluation for cancer treatment (clinical trials: NCT01632306 Phase I/II, NCT01287520 Phase I). *In vitro* and *in vivo* studies showed that treatment with LY2090314 in PDAC models increased the inhibitory phosphorylation of GSK3 and significantly suppressed the expression and pro-survival activity of TGF β -activated kinase 1 (TAK1), a crucial mediator of cellular signals that sustain PDAC aggressiveness and chemoresistance (Bang et al., 2013; Melisi et al., 2011; Giovannetti et al., 2014). Moreover, Santoro and co-workers demonstrated that reduced TAK1 expression induced by pharmacologic inhibition of GSK3 impacts the YAP/TAZ functions in PDAC cells, thus affecting their contribution to the progression and drug insensitivity of this malignancy (Santoro et al., 2020; Lin et al., 2015).

The GSK3 inhibitors AZD-1080 and Tideglusib were first designed and tested for the treatment of Alzheimer's disease (AD). Notably, they are currently under investigation in the context of cancer therapy (Lovestone et al., 2015). Ovarian carcinoma cells exposed to AZD-1080 showed a significant downregulation of GSK3 β , as well as cell cycle progression related genes at both the transcript and protein levels (Chen et al., 2016). The high selectivity and remarkable suppression of cancer cell proliferation following AZD-1080 treatment has been recently confirmed in PDAC cell lines.

As BIO and AZD-1080, Tideglusib showed selective inhibitory activity on GSK3 β , impairing the phosphorylation of many GSK3 targets, including β -catenin and c-Myc in refractory PDAC cells and models harboring mutated KRas (Kazi et al., 2018; Domínguez et al., 2012). Mechanistically, Tideglusib differs from the ATP-competitive hallmark of AZD-1080, eliciting an irreversible inhibition of GSK3 β *via* a non-competitive mode, although the exact binding site of this molecule has not been elucidated yet (Domínguez et al., 2012).

Interestingly, Edderkaoui and colleagues observed that even though the treatment with Tideglusib in PDAC cells (HPDE6, Bx-PC3, MIA PaCa-2 and HPAF-II) promoted the expression of the EMT marker vimentin, the combinatorial inhibition of histone deacetylases (HDAC) class I-II and GSK3 β reduced cancer cell survival and the levels of EMT markers (Edderkaoui et al., 2018).

Based on this observational study, they developed Metavert, a novel synthetic molecule designed by combining Tideglusib and suberoylanilide hydroxamic acid (SAHA) active pharmacophores. SAHA (Vorinostat) is an FDA approved inhibitor of HDAC class I-II currently used for the treatment of cutaneous T cell lymphoma. By inhibiting both GSK3 β and HDAC-2, Metavert synergistically impaired *in vitro* PDAC cell proliferation and prevented drug resistance as well as the expression of migration-, EMT- and stemness-associated markers. Furthermore, it significantly reduced tumor cell growth, preventing metastasis and improving overall survival in aggressive PDAC mouse models (Edderkaoui et al., 2018).

Recently, a new library of synthetic topoisomerase II inhibitors with a central replaced 1,2,4-oxadiazole ring reported promising 50 % growth inhibition values against a panel of different human cancer cell lines [49]. Specifically, five of these newly synthesized agents (2a-e) effectively reduced cancer cell viability in PDAC cells (Panc-1, SUIT-2 and Capan-1), with compound 2a displaying the highest cytotoxic activity (IC₅₀ range 0.40–1.19 μ M) (Supplemental Table 1) (Carbone et al., 2021). Compounds 2a and 2e significantly reduce GSK3 β phosphorylation in Panc-1 cells, potentially impacting cancer cell survival and tumor progression. Overall, *in vitro* studies revealed that the antiproliferative effects of these novel topoisomerase II inhibitors correlated with apoptosis induction, reduced cell migration and expression of the EMT markers SNAIL-2 and metalloproteinase-9, thus paving the way for new promising studies for the treatment of PDAC with GSK3 inhibitors (Carbone et al., 2021).

2.4.2. Combined therapy with GSK-3 β inhibitors

Limited second line therapy approaches are currently available for the management of refractory PDAC. Multiple studies in cancer models reported that certain GSK3 β inhibitors enhance tumor sensitivity to chemotherapeutic agents (Abrams et al., 2021; Miyashita et al., 2009; Shimasaki et al., 2012). Therefore, in the past decade various GSK3 β inhibitors have been experimentally and clinically tested in the context of refractory and advanced metastatic PDAC with the purpose of developing effective therapeutic strategies that could prevent or overcome drug resistance, while lowering chemotherapy-associated untoward toxicity (Table 2). *In vitro* and *in vivo* research showed that, as gemcitabine cytotoxicity is dependent on cell cycle regulatory processes, pharmacologic inhibition of GSK3 β can prevent DNA damage repair inflicted by gemcitabine and induced apoptosis in chemoresistant PDAC cells (Ding et al., 2019; Shimasaki et al., 2012). For example, 9-ING-41 ameliorates the antitumor effects of gemcitabine through modulating the ATR-Chk1 DNA damage response (Ding et al., 2019). In fact, observational studies reported that pharmacologic inhibition of GSK3 β in PDAC cells triggers topoisomerase II β binding protein (TopBP1) degradation and impairs ATR activation, consequently reducing gemcitabine-mediated Chk1 phosphorylation (Ding et al., 2019). Additionally, in contrast to results obtained by Mamaghani and colleagues, novel experimental evidence reported that AR-A014418 not only suppressed proliferation of PDAC cells and impaired tumor growth, but also synergistically sensitized tumor cells to gemcitabine treatment. Indeed, transcriptome profiling revealed that inhibition of GSK3 β counteracts the gemcitabine-induced expression of DNA repair, cell death and autophagy-related genes, such as the tumor protein 53-induced nuclear protein 1 (TP53INP1) (Mamaghani et al., 2009; Shimasaki et al., 2012).

Further research revealed that LY2090314 synergistically interacts with clinically relevant chemotherapeutic agents gemcitabine, oxaliplatin, nab-paclitaxel and SN-38, the active metabolite of irinotecan (Santoro et al., 2020; Lin et al., 2015), by modulating the intrinsic chemoresistance of PDAC cells. Interestingly, while treatment with drug combinations decreased PDAC cell viability, mice treated with LY2090314 and nab-paclitaxel exerted improved overall survival with reduced cytotoxic effects on non-malignant pancreatic cells (Santoro et al., 2020; Zamek-Gliszczynski et al., 2013).

Moreover, the pioneer GSK3 β inhibitor lithium has been observed to synergistically improve the antitumor effect of gemcitabine mainly by perturbing the Hh-GLI cascade and enhancing the proteasome-degradation of GLI (Peng et al., 2013). Next to that, Elmaci and Altinoz suggested that the triple-agent regimen comprising already-in-use drugs metformin, pioglitazone and lithium may synergistically target cancer cell metabolism by activating AMPK and PPAR- γ and perturbing GSK3 β , respectively (Elmaci and Altinoz, 2016). In *in vitro* as well as in animal model studies, this triple drug combination increased the intrinsic sensitivity of PDAC cells to apoptosis, potentially providing a novel beneficial adjuvant therapy for refractory PDAC.

Table 2
Drug combination: with GSK3 β inhibitors in PDAC.

Drug Combinations with GSK3 β inhibitors in PDAC				
Treatment	GSK3 β effect	Pharmacological interaction	Tumor stage	Ref/Clinical trial ID
AR-A014418 and Gemcitabine	Impaired DNA repair gene regulation and expression. Inhibition of Notch1 expression.	Synergistic	Preclinical models	Kunnimalaiyaan et al., 2015
9-ING-41 and Gemcitabine	Modulation of ATR-Chk1 DNA damage response	Synergistic	Refractory	NCT03678883 Ding et al., 2019 NCT01632306, 2019
LY2090314 and Gemcitabine, FOLFOX or Gemcitabine + Nab-paclitaxel	CDK-dependent RRM1/2 downregulation and increased DNA damage	Synergistic	Advanced or metastatic	Phase I/II NCT01287520, 2018 Edderkaoui et al., 2018
Metavert and Gemcitabine	Impaired metabolic profile, cell migration capability and cancer stemness. Altered tumor microenvironment.	Synergistic	Locally advanced	
Lithium and Gemcitabine	Impaired Hh-GLI signaling	Additive	Preclinical models	Peng et al., 2013

Furthermore, Metavert, the dual GSK3 β /HDAC inhibitor, when used in combination with irradiation and chemotherapeutic agents paclitaxel or gemcitabine, it reduced tumor growth compared to monotherapy and significantly prolonged the survival rate of mice harboring drug-resistant PDAC (Edderkaoui et al., 2018).

Overall, the combination of GSK3 β inhibitors and other chemotherapeutic agents appears to synergistically reduce tumor cell growth and increase survival in different models, revealing encouraging therapeutic effects towards the overcoming of chemoresistance in refractory PDAC and paving the way for future clinical studies.

2.4.3. Clinical studies on drug combination regimens including GSK-3 β inhibitors

At present, different drug combinations have been explored in pre-clinical and early-phase clinical studies in order to meet the urgent demand for efficient strategies in the therapeutic management of advanced and metastatic PDAC. However, none of them has already been approved.

Cumulative preclinical evidence provided the rationale to clinically test 9-ING-41 (phase I/II, NCT03678883) in combination with the standard chemotherapeutic agents gemcitabine, nab-paclitaxel, carboplatin, paclitaxel, doxorubicin, lomustine or irinotecan in patients with advanced or refractory solid tumors, including PDAC (NCT03678883, 2021; Carneiro et al., 2020). Furthermore, the promising preclinical outcome of LY2090314 and platinum combination in xenograft models have prompted the clinical evaluation of LY2090314 plus carboplatin and pemetrexed in patients with advanced or metastatic cancer (phase I, NCT01287520) (NCT01287520, 2018). Although establishing the efficiency of LY2090314 combined with carboplatin and pemetrexed requires further interventional confirmations, LY2090314 safety profile, pharmacokinetic parameters and optimal drug doses were established (NCT01287520, 2018; Gray et al., 2015). A parallel study conducted on patients with acute leukemia showed that LY2090314 was well tolerated and reported good antitumor activity when combined with chemotherapeutics, while minimal clinical benefits were observed if administered as monotherapy (Gray et al., 2015; Rizzieri et al., 2016). Additionally, phase I/II trial (NCT01632306, 2019) assessing the combination regimen of LY2090314 and different chemotherapies (FOLFOX, gemcitabine and nab-paclitaxel) in patients with metastatic PDAC was recently terminated due to slow enrollment procedure (NCT01632306, 2019).

Currently, combination treatment with lithium, cimetidine, olanzapine and valproate regimen (CLOVA cocktail) is under clinical investigation with simultaneous usage of gemcitabine in advanced PDAC patients (UMIN00005095), but no data are available yet.

Various phase II clinical trials on diverse pathologies are actually evaluating possible therapeutic regimens with the GSK3 β inhibitor tidesglusib (NCT01350362, NCT02586935, NCT02858908), providing potential curative possibilities to be tested in the near future also in

patients with PDAC (Lovestone et al., 2015; Horrigan et al., 2020). Nevertheless, it is important to consider that currently available clinical data provide limited information regarding the overall administration-related adverse events of these combination therapeutic regimens. Indeed, the complexity of cellular GSK3 β interconnections, the small number of ongoing clinical trial and the fact that none of the GSK3 β inhibitors has been approved for clinical use to date, excluding lithium, further complicate the prediction of beneficial or adverse effects in patients. Therefore, additional studies are required before proposing GSK3 β inhibitors-based interventions as therapeutic alternatives in the clinical management of advanced and metastatic PDAC.

2.5. Resistance to GSK-3 inhibitors

Experimental studies revealed that various gene programs are activated upon GSK3 β inhibition, mainly involving metabolic reprogramming, compensatory pro-survival signaling cascades, hyper-activation of NF- κ B and WNT signaling, as well as increased autophagy/lysosomal network activity (Ougolkov et al., 2005; Marchand et al., 2015; Seino et al., 2018; Webster et al., 2000; Sun et al., 2016; Bruton et al., 2020).

Despite the general consensus regarding the antineoplastic activity of GSK3 β inhibitors, a more comprehensive analysis on the downstream effects of GSK3 β inhibition uncovered the potential induction of escape signals mediated by increased autophagic response (Marchand et al., 2015). Although PDAC cells harbor elevated levels of basal autophagy, whether autophagy displays a tumor suppressor role or a potential resistance mechanism to anticancer therapy, remains elusive (Yang et al., 2011; Galluzzi et al., 2015).

A recent study demonstrated that pharmacologic inhibition of GSK3 β in PDAC cells enhanced the transcriptional activity of TFEB, thus positively modulating the autophagic flux (Marchand et al., 2015). Moreover, experimental inhibition of the autophagy cascade with bafilomycin A1 and/or CHIR99021 ameliorated the sensitivity of human PDAC cells to apoptosis triggered upon concomitant treatment with GSK3 inhibitor (Marchand et al., 2015). Similar outcomes were observed in prostate and bladder cancer cells, further sustaining the hypothesis that autophagy exerts a potential mechanism of resistance to GSK3 β inhibition and that a combination drug treatment targeting both GSK3 β and the autophagy/lysosomal network might prevent this issue (Marchand et al., 2015; Kuroki et al., 2019). Furthermore, autonomous production of stromal WNT ligands, sustaining constitutive Wnt signaling, as well as hyper-activated NF- κ B transcriptional activity, have been described in various subsets of PDAC cells (Ougolkov et al., 2005; Seino et al., 2018; Webster et al., 2000; Bruton et al., 2020). Therefore, in light of the molecular interconnections between GSK3 β and these signaling pathways, it would not be surprising if these processes might constitute intrinsic mechanisms of resistance to GSK3 β inhibition in pancreatic tumor cells.

In this regard, several analyses distinguished two PDAC subtypes, the classical and squamous lineages, which are characterized by distinct transcriptomic and proteomic profiles, as well as prognosis (Collisson et al., 2019; Bailey et al., 2016; Le Large et al., 2020). Specifically, while the classical subtype expresses differentiated endoderm cell markers and mostly experiences favorable clinical outcome, the squamous phenotype harbors altered epigenetic landscape, affecting the expression of duct cell markers and leading to a rapid metabolic reprogramming often leading to a worse prognosis (Bailey et al., 2016; Lomberk et al., 2018; Le Large et al., 2017). This was corroborated by Brunton and colleagues, who reported that a subset of squamous pancreatic cell lines rapidly acquired drug resistance upon treatment with GSK3 β inhibitors (Brunton et al., 2020). Specifically, these cancer cells encountered a metabolic adaptation under persistent suppression of glycolysis mediated by pharmacologic inhibition of GSK3 β . Furthermore, this was accompanied by an increased dependency on autophagy and activation of a unique gene transcription program resulting in the self-production of WNT ligands, and ultimately leading to drug resistance.

Overall, sustained autophagy flux and activation of compensatory cascades appear to rapidly induce acquired resistance to GSK3 β inhibition in PDAC cells. Therefore, further studies are warranted to unravel the complexity of autophagy: potential therapeutic applications in PDAC (Gomez Mellado et al., 2015). However, in-depth analysis of unique chromatin landscape signatures and mutation profiles of PDAC cells might improve our understanding of the dynamics of emergence of drug resistance mechanisms, thus supporting the design of more effective therapeutic approaches.

2.6. Tumor chromatin profiling may predict patients with pancreatic tumors sensitive to GSK3B-targeted therapy

The lack of defined biomarkers and the high disease heterogeneity characterizing PDAC are representative of the difficulties in predicting and determining which patients might respond to targeted therapies.

To date observational data highlighted the link between mutant KRas PDAC and GSK3 β overexpression (Zhang et al., 2011; Fleming et al., 2015; Fitzgerald et al., 2015). In fact, oncogenic KRas signaling, positively regulates GSK3 β expression and activity, thus favoring cell proliferation, survival and tumor dedifferentiation. Pharmacologic inhibition of GSK3 β in KRas-dependent tumors was found to impair cancer cell growth and induce apoptosis, partly mediated by c-Myc- and β -catenin-dependent mechanisms (Kazi et al., 2018; Kim et al., 2000). However, the intrinsic plasticity and aggressiveness of KRas-dependent tumors may rapidly lead to the acquisition of drug resistance conclusively discouraging clinical intervention with targeted therapies such as GSK3 β inhibitors (Marchand et al., 2015; Downward, 2015; Cox et al., 2014).

Novel epigenetic and transcriptomic studies allowed the classification of PDAC into two main subtypes according to their gene expression profiles, hence providing a prediction of chemoresistance as well as the prognosis (Chan-Seng-Yue et al., 2020). As mentioned above, concomitant inhibition of HDAC and GSK3 β may prevent the emergence of drug resistance (Edderkaoui et al., 2018). More recently, new analyses were performed to define chromatin accessibility regions to identify epigenetic hallmarks of tumors sensitive to GSK3 inhibition; they reported that increased access to intronic and distal promoters regulating WNT cascade genes, as well as enrichment in transcription factor motifs, may result in WNT cascade amplification and drug resistance (Bruton et al., 2020). These findings, combined with early results from the prospective COMPASS study (NCT02750657), confirmed that chromatin profiling in advanced PDAC may help define tumors that could benefit from target therapies (Chan-Seng-Yue et al., 2020; Aung et al., 2018).

Specifically, Bruton and colleagues suggested that pancreatic tumors with high mutational burden and chromatin instability are more prone to develop drug resistance. Moreover, those tumors harboring loss of hepatocyte nuclear factor 4- α (HNF4 α), an important regulator of endodermal lineage differentiation, more likely maintain sensitivity to GSK3 β inhibitors, due to GSK3 β upregulation and consequent increased tumor dependency (Bruton et al., 2020). Furthermore, ATAC-seq analysis on sensitive pancreatic tumors revealed mutations on chromatin modulators, possibly KFM6A, SETD2, MLL3, ARID1A and SETBP1, that favor distal promoter usage and alterations in either the AMPK signaling activator LKB1, or in WNT canonical pathway transducer LRP6. This proves their crucial role in the maintenance of cancer cell response to GSK3 β -targeted monotherapy.

Many studies clearly showed the complex heterogeneity of pancreatic tumors and their microenvironment aiming to link the tumor genetic mapping with the prediction of patients' response to tailored treatment approaches (Bailey et al., 2016; Boyd et al., 2021). Determining the constellation of tumor genetic and transcriptional alterations not only helps the definition of specific cancer subtypes, but might also have implications to the development of targeted therapeutic strategies specifically designed to address the patient tumor profiles as well as to circumvent or surmount drug resistance.

3. Discussion and conclusions

Pancreatic cancer is a growing global health concern with an increasing incidence-to-mortality ratio. To date, it is the third most common cancer and, due to its frequent dismal prognosis, it will probably become the second-leading cause of cancer-related death in Western countries by 2030 (Rahib et al., 2014). The high genomic complexity and heterogeneity of pancreatic cancer, as well as its intrinsic metastatic behavior, represent an important barrier for the successful treatment of this lethal disease (Boyd et al., 2021). Surgical resection constitutes the only modest chance of cure, while actually standard-of-care therapeutic options are often palliative and offer an average of 5-years survival with most of patients developing drug resistance during the course of the treatment (Zeng et al., 2019). The poor outcomes are mainly related to late diagnosis and the strong aggressive nature of this malignant disease, highlighting the demand for discovering novel tumor vulnerabilities and effective therapies.

Despite dismal statistics, significant progress has been made to elucidate the molecular mechanisms involved in pancreatic cancer progression and chemoresistance. Among all, the glycogen synthase kinase-3 β , a highly conserved isoform of serine-threonine kinase GSK-3 family, has been recently found as an important determinant of PDAC onset and progression (Ougolkov et al., 2006; Zhang et al., 2011;). Initially described as a crucial modulator of glycogen synthesis, GSK3 β is now confirmed to be involved in many fundamental cellular processes. Indeed, aberrant GSK3 β activity has been implicated in different human disorders including bipolar depression, neurodegenerative disorders, acute myeloid leukemia, as well as many other malignancies (Walz et al., 2017; Hooper et al., 2008; Martelli et al., 2021).

Pre-clinical studies showed that pharmacologic inhibition or genetic depletion of GSK3 β drastically reduced cell proliferation and cell survival of multiple human tumor types, further highlighting GSK3 β as an attractive pharmacological target for therapeutic interventions against cancer (Rizzieri et al., 2016; Kotliarova et al., 2008; Korur et al., 2009; Song et al., 2010; Cohen et al., 1998; Carter et al., 2014; Kroon et al., 2014). Moreover, several studies on drug combination in patients with refractory solid tumors have shown that inhibition of GSK3 β sensitizes resistant cancer cells to standard chemotherapeutic agents such as gemcitabine, nab-paclitaxel, doxorubicin, and FOLFIRINOX (Bhat et al., 2003; Kunimallaiyaan et al., 2015; Shimasaki et al., 2012;). Thus, multiple evidence suggested that targeting GSK3 β may reverse chemoresistance and highlighted its key role in many intracellular signaling pathways. These brought GSK3 β to the attention of many researchers which are currently attempting to better understand the role of this this enigmatic kinase in the cellular dynamics of pancreas tumorigenesis and drug resistance.

Although GSK3 β has been previously described as a tumor suppressor regulating the activity of numerous pro-oncogenic molecules such as c-Myc, β -catenin, cyclin D and c-Jun, a series of consistent observational studies reported that GSK3 β is strongly upregulated in PDAC cells and could sustain pancreatic tumorigenesis (Kockeritz et al., 2006; Nagini et al., 2019; Walz et al., 2017). Notably, mutant KRas pancreatic cancers, accounting for 91 % of overall PDAC patients, present the over-expression and nuclear accumulation of active GSK3 β which often correlate with poorly differentiated tumor state and poor outcomes (Christenson et al., 2016).

Tumors expressing mutated Ras usually harbor enhanced activation of mitogenic PI3k signaling and perturbed PTEN phosphatase activity, providing crucial signals driving tumor formation and maintenance (McCubrey et al., 2012a, 2012b; Fitzgerald et al., 2015; Waters and Channing, 2018). However, the intricate role of KRas in sustaining multiple mitogenic signaling pathways, such as activation of Akt, HER2 and EGFR, may explain the lack of success in developing KRas targeted therapies, despite decades of intense research efforts. More intriguing, GSK3 β seems to be fundamental for the survival and growth of KRas-driven PDAC. Pharmacologic inhibition of GSK3 β with SB,

Tideglusib, AZD1080 and BIO, selectively reduced the proliferation of PDAC with dependency on mutant KRas, further evidencing the pro-survival effect of GSK3 β in these tumors.

The controversial anti-tumorigenic or pro-tumorigenic role of GSK3 β in PDAC is finely regulated by diverse mechanisms, including post-translational modifications, cellular localization and trafficking, formation of protein complexes, and substrate priming. All these processes have been extensively studied in order to understand the dynamics governing GSK3 β activation and disruption.

Among all, phosphorylation of tyrosine 216, located within the conserved activation loop, is responsible for the full activation and kinase function of GSK3 β , while serine 9 residue in the N-terminal lobe inhibits GSK3 β activity when phosphorylated by other kinases (Cormier and Woodgett, 2017; Sutherland, 2011; Frame et al., 2001). However, considering pGSK3 β -S9 as the inactive and pGSK3 β -Y216 as the active form is probably over simplistic. In fact, GSK3 β undergoes a dynamic equilibrium within those and others recently identified phosphorylation status in concert with stimulatory signaling molecules and primed substrate concentration. Additionally, serine 9 phosphorylation does not completely abrogate the catalytic activity of this kinase, as proven by the pGSK3 β -S9-mediated phosphorylation of Gli3 within the Hh signaling pathway (Fitzgerald et al., 2015).

These observations might explain why many ATP-competitive GSK3 β inhibitors interacting with the N-terminal lobe exert high concentration IC₅₀ values and low kinase selectivity when compared to covalent inhibitors or non-ATP competitors. In fact, the ATP-binding domain is structurally conserved among most of kinases and, therefore, it is not surprising that some ATP competitive GSK3 β inhibitors target also CDKs and other kinases which share a good degree of homology. Moreover, ATP-competitors increasing the inhibitory phosphorylation on serine-9 might not be optimal to abrogate GSK3 β activity since phosphorylation of this residue, located within the binding pocket for primed-substrates recognition, might still result in GSK3 β -mediated regulation of non-pre-phosphorylated targets.

On the other hand, covalent- or non-ATP-competitive inhibitors of GSK3 β such as tideglusib, display moderate-to-weak binding but improved selectivity and low drug concentrations are required to attain therapeutic effects. Overall, these factors might determine the choice of using these classes of GSK3 β inhibitors in clinical practice, but further studies on GSK-3 protein-substrate are required for future development of potent GSK3 β inhibitors.

The promising therapeutic results expected from GSK3 β targeting in PDAC found a solid base in the progressive increase of GSK3 β expression which, in turn, strongly regulates NF- κ B transcriptional activity (Ougolkov et al., 2006; Demarchi et al., 2003; Ben-Josef et al., 2015). This consequently triggered the stimulation of pathways involved in cell survival, proliferation and a pro-invasive transformation of pancreatic cancer cells, as well as in the promotion of Bim family proteins expression (Marchand et al., 2012).

Moreover, GSK3 β appears to have a negative regulatory role on apoptosis, through phosphorylation and subsequent inactivation of pro-apoptotic molecules such as Bad, Ikk β and caspase 9, further sustaining a pro-survival phenotype (Cervello et al., 2012; Meng et al., 2018). GSK3 β also modulates Wnt/ β -catenin signaling: active GSK3 β phosphorylates β -catenin, targeting it for ubiquitin-mediated proteasomal degradation (McCubrey et al., 2016) mediating Wnt cascade activation. This leads to inhibition of GSK3 β activity, β -catenin accumulation and translocation into the nucleus, causing the expression of proto-oncogenes such as c-Myc and cyclin D1, along with genes promoting cell invasion and migration. In this regard, whether GSK3 β action is dependent or not on β -catenin perturbation is still controversial, sustained WNT cascade activity was observed to drive drug resistance in PDAC models treated with GSK3 β inhibitors (Freland and Beaulieu, 2012; Zhang et al., 2003; Yu et al., 2012).

To date, further mechanisms of drug resistance have been identified to be provoked by treatment with GSK3 β inhibitors, mainly involving

the activation of autophagy/lysosomal network (Marchand et al., 2015). Although coexistence of both apoptosis and autophagic responses have been observed in many *in vitro* studies testing different GSK3 β inhibitors, additional molecular studies reported that the transcription factor EB provided pro-survival autophagic signals by enhancing autophagy/lysosomal activity. Concomitantly, TFEB-depleted PDAC cells exhibited enhanced sensitivity to cell death upon GSK3 β inhibition, further proving the protective role of autophagy in pancreatic cancer cells under GSK3 β disruption. Similar results were observed in prostate cancer models, where GSK3 β depletion enhanced AMP/ATP ratio, eliciting AMPK signals and autophagy activation (Sun et al., 2016). Even if combinatorial inhibition of GSK3 β and TFEB has not been investigated yet, assessing GSK3 β -mediated regulation of autophagic responses might be a future achievement in the development of PDAC treatment.

To prevent the onset of drug resistance, novel advances in understanding escape pathways and chromatin landscape in GSK3 β inhibition have been elucidated in experimental settings. Epigenetic analysis revealed that histone deacetylases are highly expressed in PDAC cells, coordinating cell cycle progression, differentiation and apoptosis. Enhanced HDACs activity, resulting from GSK3 β -mediated activation of Zeb1 transcription factors, consequently induced the repression of E-cadherin expression and lead to a poorer prognosis (Aghdassi et al., 2012). Indeed, concomitant inhibition of GSK3 β and HDACs with the synthetic agent Metavert caused synergistic anti-proliferative effects on PDAC cells and prevented EMT-associated gene expression (Edderkaoui et al., 2018).

Furthermore, chromatin profiling among advanced pancreatic cancer suggests that tumors harboring high mutational burden and chromatin instability are more prone to present drug resistance, and patients with loss of the hepatocyte nuclear factor 4- α and enhanced GSK3 β expression were more likely to respond to treatment with GSK3 β inhibitors (Bruton et al., 2020).

A general optimism is increasing nowadays when combining chemotherapy with novel agents, targeting specific features of different tumors. Indeed, it is overall admitted that drugs targeting various tumor-survival signaling cascades might exert therapeutic advantages compared to single agent treatment approaches. Therefore, considerable advances have been achieved in the development of targeted drugs with reduced toxicities, improved survival benefits and potentiality to overcome or prevent chemoresistance.

However, approved targeted therapies for pancreatic cancer treatment include only olaparib (Lynparza), and erlotinib (Tarceva). Recent preclinical studies underlined the potential of new inhibitors of the focal adhesion kinase (Le Large et al., 2021) or c-Met (Firuzi et al., 2019). Similarly, a few studies showed that administration of GSK3 β inhibitors reduced pancreatic cancer cell proliferation, but more interestingly, it resensitized drug resistant cells to standard of care chemotherapy within combinatorial regimens. Indeed, disruption of GSK3 β has been described to modulate the TAK1-YAP/TAZ axis and the ATR-mediated DNA damage response pathway, thus driving the restoration of effective cytotoxic response (Ding et al., 2019; Lin et al., 2015).

On the other hand, GSK3 β is a fundamental crossroad for multiple anti-oncogenic pathways, by promoting tumor suppressor signaling cascades and this feature has raised a general mistrust in approving such agents for therapeutic interventions against cancer. Thus, the evaluation of GSK3 β inhibitors in clinical trials has been hampered by the concern that inhibition of GSK3 β may stimulate malignant transformation. However, future promising perspectives for GSK3 β inhibitors clinical management of cancer has been recently achieved from observational studies reporting that long-term use of the only approved GSK3 inhibitor, lithium, is not associated with increased risk of cancer in patients with bipolar disorder (Martinsson et al., 2016)

Overall, developing the medical treatment of choice for tumors, such as PDAC, that manifests high recurrence and persistent invasion capacity, metastasis and development of drug tolerance, remains a challenge for current clinical interventions. Chemotherapeutic

interventions, radiation and immunotherapy have indeed minimal effect on patient's survival, highlighting the burning need for additional mechanistic studies exploiting cellular vulnerabilities of advanced and metastatic PDAC. In this respect, additional studies are required to extensively understand the consequence and dynamics regulating aberrant GSK3 β activity. Furthermore, a specific focus should also be directed to GSK-3 α , the other isoform of GSK3, which presents distinctive cellular functions. Since these kinases are differentially expressed within tissues and the majority of cancer studies has focused on GSK3 β , it is questionable if targeting GSK3 α together with GSK3 β has a major effect than single GSK-3 β inhibition.

In conclusion, the combination of GSK3 β inhibitors with chemotherapy is strategically poised to be a promising approach to overcome the emergence of early drug resistance or to overcome chemoresistance in advanced and metastatic pancreatic tumors. Further understanding of the dynamics governing PDAC tumorigenesis and cancer progression involving GSK3 β , might help the achievement of clinical strategies aimed at ameliorating survival benefits and to convert this deadly tumor into a more manageable chronic malignancy.

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Appendix A. Supplementary data

Supplementary material related to this article can be found, in the online version, at doi:<https://doi.org/10.1016/j.drup.2021.100779>.

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