The lipid kinase PI3Kα is required for the cohesion and survival of cancer

cells disseminated in serous cavities.

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Running Title

PI3Kα controls tumor cell cohesion

ABSTRACT (150 WORDS) – MAX 150 WORDS

Breast, ovarian, digestive and lung adenocarcinomas are often associated with the accumulation

of malignant cells in serous cavities. As PI3K is one of the most mutated pathways in cancer, we

investigated the importance of oncogenic PI3Kα in this process. We analyzed tumor cell

organization in ascites from carcinomas at diagnosis. In some malignant ascites, tumor cells grew

as adhesive coherent masses. Ex-vivo patient-derived cell cultures with the addition of

mesenchymal stem cells, as a model of tumoral stroma, favored the compaction of tumorospheres.

Ascites-derived ovarian cancer cell lines frequently harbored *PIK3CA* mutations coexisting with

other mutations. PI3Kα promoted the formation and maintenance of multicellular adhesive

PIK3CA-mutant spheroids, promoting cell survival. Cultures in 3D conditions as opposed to

cultures in cell monolayers increased chemotherapy resistance, which was overcome by PI3Kα

inhibition. We identified a signaling pathway of interest for the treatment of cancer cells

disseminated in serous cavities, limiting cancer progression.

Key words

PI3K signaling, tumor cell aggregation, resistance to treatment, signal-targeted therapies,

malignant serous fluid

(24599 CHARACTERS) - MAX 25000 CHARACTERS (UP TO FIGURE LEGEND –

EXCLUDING STAR METHODS & SUPPL & REFERENCES)

INTRODUCTION

Tumor cell resistance to chemotherapy is now largely considered to be enhanced in

multicellular 3D structures. For example, EMT-6 mammary tumor cells are more resistant to

alkylating agents (cyclophosphamide, cisplatin, thiotepa) when grown in 3D spheroids

(Kobayashi et al., 1993; Tan et al., 2006). Single or clustered cells detach from primary breast or

prostate tumors and are then found in the bloodstream (Aceto et al., 2014). In the peritoneal,

pleural or pericardial cavity, the formation and properties of these tumoral 3D aggregates, which

occur in breast, ovarian, gastrointestinal, pancreatic and lung cancers, remain poorly understood.

Their specific mechanisms of resistance to treatment are largely understudied. However, in

peritoneal carcinomatosis of ovarian cancer patients, it is largely accepted that the peritoneal

microenvironment contributes to chemotherapy resistance (Castells et al., 2012), in particular via

mesenchymal stem cells (MSCs) (Roodhart et al., 2011). They have been shown to contribute to

tumor cells' ability to form spheroids called tumorospheres under anchorage-independent culture

conditions, increasing their capacity to induce tumor formation in vivo after mouse xenografts

(Kabashima-Niibe et al., 2013; Lee et al., 2018).

The class I PI3K (phosphoinositide 3-kinase)/Akt axis is one of the most commonly

deregulated pathways in cancers, including in ovarian, breast and lung cancers. There are four

heterodimeric isoforms of class I PI3K in vertebrates: α, β, γ and δ. PI3K activation results in the

phosphorylation of phosphatidylinositol-4,5-biphosphate (PIP2) into phosphatidylinositol-3,4,5-

triphosphate (PIP3), which acts as a second messenger in the cell. Downstream signals of PI3K,

including the members of the Akt/mTOR (mammalian target of rapamycin) pathway, are

numerous and involved in many cellular processes, including cell-cycle progression and actin

cytoskeleton rearrangement (Baer et al., 2014; Tsujita and Itoh, 2015; Vanhaesebroeck et al.,

2010; Zhang et al., 2017).

One PI3Kα-selective inhibitor, BYL719 (Alpelisib), was just approved for use in breast

cancer with *PIK3CA* mutation (André et al., 2019; Mayer et al., 2017), and BYL719 in

combination with estrogen receptor inhibitors show lower toxicity compared to the use of pan-

PI3K inhibitors. PI3Kα/Akt/mTOR signaling pathway alterations and more specifically

alterations in the frequency of *PIK3CA* mutations (*PIK3CA* encodes for the PI3Kα catalytic

subunit) varied in non-small cell lung cancer (NSCLC) in published studies (Arcucci et al., 2021; Wu et al., 2016). In ovarian cancer, the PI3K signaling pathway is described as a predictor of the invasive and migratory potential of ovarian tumor cells (Bai et al., 2015). Mutations are found in the *PIK3CA* gene in 6%–33% of patients, and amplification or copy gain of this gene is found in 13%–39.8% of cases (Levine, 2005); these alterations are non-exclusive. Regarding the distributions of these alterations in different ovarian carcinoma subtypes, *PIK3CA* mutations are less common in high-grade serous carcinomas, while amplifications are frequently observed. The genetic landscape of ovarian carcinomas suggests that PI3K inhibitors, and in particular those targeting PI3Kα, could be tested in this pathology.

Twenty years ago, peritoneal carcinomatosis was seen as an incurable pathology requiring palliative care only. However, the introduction of loco-regional therapies combining cytoreductive surgery with hyperthermic intraperitoneal chemotherapy (HIPEC) based on platinum salts or taxanes changed the management of this disease (Wright et al., 2015). Signal-targeted therapies, such as those targeting the PI3K/Akt pathway, are currently being tested with the goal of improving disease-free survival rates (Mabuchi et al., 2015).

The importance of PI3Kα activity in tumoral aggregates from adenocarcinomas disseminated in serous cavity is unknown. Our objective is to demonstrate the involvement of PI3Kα during tumorosphere formation and maintenance in serous cavities and their role in chemotherapy resistance.

RESULTS

Peritoneal tumor cells from ascites punctures form adhesive aggregates whose growth is delayed by a high number of initial cells

Ascites of 12 ovarian cancer patients were punctured and analyzed to detect the presence

of adenocarcinomatous cells. We first observed tumor cell aggregates found in ascites using

cytospin or an embedded preparation analyzed with standard immunohistochemistry techniques,

Papanicolaou (PAP), May-Grünwald-Giemsa (MGG), periodic acid–Schiff (PAS) and epithelial

cell adhesion molecule staining (Ber-EP4) (Fig. 1A). We observed heterogeneity in terms of the

distribution of cellular aggregate density (Suppl. Table S1). Most ascites (11/12 patients) were

composed of limited numbers of isolated tumoral cells and numerous aggregated tumoral cells

(representative image, Fig. 1A left). Only one case revealed numerous isolated cells alongside

aggregates (representative image, Fig. 1A right). When grown in 3D-favorable conditions, ex-vivo

cultures of ascites cells formed dense and cohesive spheroids (Fig. 1B). Interestingly, ovarian cell

lines derived from ascites were enriched in *PIK3CA* mutations and/or p53 alterations compared to

primary tumor cell lines (Suppl. Table S2). To reproduce the carcinomatosis development of

cohesive aggregates (Fig. 1C), we first developed a fluorescent 3D culture system of peritoneal

tumor cells with *PIK3CA*-mutant ovarian cancer cell line SKOV-3 stably expressing GFP (Fig.

1D). After 14 days, we analyzed the influence of cell numbers in aggregates and observed an

increase of 75% and 13% in the tumorosphere area when 5,000 and 10,000 cells were seeded,

respectively (Fig. 1D, E). A higher number of initial cells appears to slow tumorosphere growth.

Interestingly, area evolution (+24% in 7 days) (Fig. 1D, E) was associated with the constant

expression of fluorescent GFP as a readout of cellular activity (Fig. 1F). As expected, fluorescence

tends to decrease in the core of tumorospheres (Fig. 1D). Ascites is a mix of different cell

populations (Fig. 1A). To reproduce this heterogeneity, we added a stromal compartment, adding

MSC, known to be recruited at inflammatory tumoral sites. We tested their capacity to aggregate with

tumor cells. We carried out 3D cultures with 10,000 MSC alone and observed a completely

different aggregate evolution profile compared to SKOV-3 cells; there was very fast cell cohesion

the day after seeding, and within 14 days, the aggregate area had decreased by more than 60%

compared to the initial area (Suppl. Fig. S1A and B). We also observed the formation of fibrous

structures within MSC aggregates, the nature of which remains to be determined (Suppl. Fig

S1C). Those structures were also observed in aggregates developed from patient-derived tumor

cells (Fig. 1B, top right). We then co-cultured 5000 GFP-expressing SKOV-3 cells with different

quantities of Qdot-labelled MSCs (Fig. 1G-H) (Latifi et al., 2012). The heterotypic tumorosphere

area decreased proportionally to the number of MSCs, with an apparent increased cohesion (Fig.

1G, H). GFP fluorescence corresponding to tumor cells did not vary regardless of MSC number,

indicating that these cells induced stronger tumorosphere cohesion without modifying tumor cell

proliferation (Fig. 1H, I). At a high ratio of MSCs to tumor cells, MSCs clustered in the center of

tumorospheres and colocalized with tumor cells (yellow staining, Suppl. Fig. S1D).

These results reveal that small-sized tumorospheres exhibit cellular growth and that the

addition of MSCs to tumor cell spheroids results in increased cohesion of tumorospheres,

reminiscent of the observed formation of tumorospheres from patients' ascites cells.

PI3Kα inhibition prevents cohesion of ovarian cancer cell tumorospheres.

The PI3K/Akt pathway plays a role in cell proliferation and survival but is also involved

in actin cytoskeleton regulation, suggesting a role in tumorosphere formation (Baer et al., 2014;

Tsujita and Itoh, 2015; Vanhaesebroeck et al., 2010). We confirmed a concentration-dependent

inhibition of Akt phosphorylation on S473 and T308 after A66 and BYL719 treatment (Fig. 2A-

C), indicating 1) that the PI3K pathway is constitutively activated in SKOV-3 cells, 2) that this

activation is highly dependent on the PI3Kα isoform and 3) that the pharmacological agents used

prevent its activity.

To determine the action of this pathway in the formation of SKOV-3-GFP multicellular

aggregates, we treated them at seeding and 3 days later with the PI3Kα-selective inhibitors A66

and BYL719 (Fig. 2D). We next compared the efficiency of a low (1 µM) and medium (10 µM)

dose of PI3K inhibitors and observed the morphology of the 3D multicellular structures. After 4

and 7 days of treatment with 10 µM of BYL719, we observed a complete reduction of the

tumorosphere area (Fig. 2E–F). Aggregates treated with 10 µM of A66 or 1 µM of BYL719 had

comparable evolution profiles (Fig. 2E), with a decrease in tumorosphere area of about 30%

compared to the control at day 4 (Fig. 2F). This effect was maintained over time.

To measure PI3Kα inhibitor effects on tumorosphere maintenance, we treated already

established SKOV-3-GFP cell aggregates (Fig. 2G). PI3Kα inhibitors caused a concentration-

dependent decrease in tumorosphere area and a tumorosphere morphology change with maximum

effect at day 7 (Fig. 2H-J). Used at 1 µM, treatment with BYL719 was more effective than A66;

there was a significant decrease in tumorosphere area of 86% and 43%, respectively, compared to

the vehicle at day 7 (Fig. 2I and J). This corresponded to a decrease in the initial tumorosphere

size.

Finally, adding MSCs, which compacted the tumorospheres, to BYL-719 at 10 µM

reduced tumorosphere cohesion (Fig. 2K). The absence of a direct effect of all PI3K inhibitors on

MSCs was verified (Suppl. Fig. S1E, F).

PI3Kα inhibitors prevent cohesive tumorosphere formation and maintenance in an α

isoform-specific manner.

PI3Kα inhibition impacts multicellular tumor cell viability in 3D cultures.

To extend our findings, we used 2 other cell lines derived from serous cavities presenting

PIK3CA mutations: the lung cancer cell line H460 and the breast cancer cell line MCF-7. Using a

survival MTT assay, we first verified that they were sensitive in 2D to PI3Kα inhibitors (Fig. 3A).

Heterogeneous responses to PI3K inhibitors were observed (Fig. 3A); the H460 cell line appeared

to be resistant to A66 and to low concentrations of BYL719 (Fig. 3A, middle).

174    We next aimed to determine the impact of the 3D culture on PI3Kα inhibitor efficiency. In

175 particular, we were interested in analyzing the organization and distribution of living and dead

176 cells within tumorospheres under treatment. PI3K inhibitor treatment altered tumorosphere

177 cohesion in SKOV-3 and MCF-7 cells (Fig. 3B, Suppl. Fig. S2A). We quantified the living cell

178 areas using calcein staining and demonstrated that BYL719 significantly decreased this marker in

179 all tested cell lines (Fig. 3B-C, Suppl. Fig S2A). SKOV-3 and H460 cell treatment with PI3Kα

180 inhibitors tended to increase the number of dead cells (Suppl. Fig. S2B). Just 10 µM BYL719

181 caused a significant decrease in the living cell area of all cell lines (Fig. 3B, C). PI3Kα inhibitors

182 induced a rapid decrease in tumorosphere area and viability.

184 **PI3K inhibitors decrease tumorosphere survival in the presence of cisplatin.**

185    If used in the clinic to treat ovarian cancer, PI3K inhibitors will be tested in combination

186 (Pons-Tostivint et al., 2017). A combination of a low concentration (0.05 or 0.5 µM) of cisplatin

187 with PI3Kα inhibitors appeared to alter more efficiently the SKOV-3-GFP tumorosphere area and

188 living cells compared to the control (cisplatin alone), but not significantly more than the

189 corresponding inhibitor alone (Fig. 4A–D). At low doses of cisplatin, both PI3K inhibitors

190 significantly decreased the 3D structure area (Fig. 4C). The addition of 1 µM PI3Kα inhibitors

191 (A66 or BYL719) to a high concentration of cisplatin (5 µM) triggered a huge decrease in living

192 cells compared to each agent alone, suggesting an additive effect for both drugs (Fig. 4D).

193 Differences in the effects observed with these different assays show that the key combined action

194 of PI3Kα inactivation and cisplatin treatment consists in decreasing cell survival and

195 tumorosphere area.

196    Compared to cisplatin treatment, treatment with PI3Kα inhibitors results in decreased

197 tumorosphere survival.

DISCUSSION

Solid cancers associated with peritoneal, pleural or pericardial carcinomatosis frequently develop fluid accumulations called ascites in the case of the peritoneum. This early step in the metastatic process involves tumor cell aggregation within the peritoneal cavity. This 3D structure enhances the ability of tumor cells to survive conventional chemotherapies (Liao et al., 2014; Tan et al., 2006), and we found that PI3Kα activation through its oncogenic mutations contributes to this process.

PI3Kα inhibitors are promising therapeutic tools to prevent local dissemination in serous cavities. The activating mutations of *PIK3CA* and the disabling mutation of *PTEN* have been identified with p53 mutation as the three most frequent point mutations among 21 types of cancers. Several studies using genetically engineered mouse models have demonstrated the role of mutant *PIK3CA* in tumor initiation, progression and maintenance (Arcucci et al., 2021). Despite these data, given the mild effect of PI3Kα inhibitors as clinical monotherapy, the oncogenic potential of mutated PI3Kα is challenged. Analyzing the action of mutant PI3Kα in combination with other genetic alterations in patients and preclinical models is of utmost importance to better understand the cooperation mechanisms between different oncogenic processes and provide novel therapeutic solutions (Pons-Tostivint et al., 2017). p53 mutations are also known to increase *PIK3CA* promoter activity (Gaikwad et al., 2013), contributing to global PI3K/Akt pathway activation in cell lines coming from ascites. Globally, it seems that compared to primary tumors, cell lines coming from ascites are enriched in either mutant PIK3CA and/or genetic p53 inactivation (Table S2).

PI3K, and more particularly the α isoform, plays a role in actin cytoskeleton rearrangement (Baer et al., 2014; Campa et al., 2015). Actin involvement in supporting adhesive connections is well established and, according to our hypothesis, PI3Kα activity would facilitate ovarian tumor cell aggregate formation (Tsujita and Itoh, 2015; Vanhaesebroeck et al., 2010).

To validate this hypothesis, we treated SKOV-3 cell tumorospheres during seeding with different

concentrations of the PI3Kα inhibitors A66 (Jamieson et al., 2011) and BYL719 (Fritsch et al.,

2014). After treatment of these aggregates, microscopy revealed sphere disorganization. Complete

disintegration was seen following treatment with 10 µM BYL719. PI3Kα seems to play a role in

both ovarian tumor cell aggregate formation and maintenance.

PI3Kα action on the actin cytoskeleton could be the source of our demonstrated effect of

PI3K inhibitors on spheroids (Lien et al., 2017). This hypothesis is reinforced by the knowledge

of an existing link between polymerized actin and adhesive molecules like E- or P-cadherins. P-

cadherin expression promotes the aggregation of circulating ovarian cancer cells (Usui et al.,

2014). These elements highlighting the role of actin could in part explain PI3Kα involvement in

cell clustering (van Baal et al., 2018; Saias et al., 2015). Therapies targeting adhesion between

cancer cells of the same type, such as PI3Kα inhibitors, are worth consideration for improving

peritoneal carcinomatosis management.

It has been repeatedly demonstrated that 2D cytotoxicity studies do not account for

complete tissue physiology and overestimate the cytotoxic potential of tested molecules

(Kobayashi et al., 1993; Shield et al., 2009). In a 3D culture, these cells are 18, 15 or 58 times

more resistant to cisplatin at 25, 50 and 100 µM, respectively (Kobayashi et al., 1993). In

addition, proliferation within the spheres decreases in comparison with cells cultured in 2D,

partially explaining why these cells can survive therapies that induce death in replicating cells

(Burleson et al., 2004; Kobayashi et al., 1993). Here, all cells cultured in 3D remain sensitive to

PI3K inhibition, suggesting that PI3K inhibition impacts cells with reduced proliferation.

Tumor aggregate disorganization induced by PI3Kα inhibitors could allow better cisplatin

bioavailability within tumorospheres by breaking the decreasing diffusion gradient from the

periphery to the center of these 3D structures (Sant and Johnston, 2017). Tumorosphere

dissociation could also decrease the solid stress that builds up in 3D culture (Delarue et al.,

2014), which was shown to decrease cell proliferation and chemotherapy efficiency (Rizzuti et

al., 2020). Other parameters we could have analyzed are the response to increased liquid pressure

(Klymenko et al., 2018) or to shear stress caused by ascites fluid motion in the peritoneal cavity

(Rizvi et al., 2013). They were shown by others to increase tumor cell proliferation rate and they

could allow cisplatin to function.

We tested the role of oncogenic PI3Kα in different types of adenocarcinomas with

dissemination in serous fluid, such as ovary, lung and breast cancers. Our findings are potentially

of therapeutic importance in ovarian cancer, which at diagnosis most often presents an advanced

stage associated with peritoneal dissemination. The current therapeutic arsenal lacks treatments

targeting the more advanced disease, which represents a major public health issue due to its

morbidity and mortality. To date, HIPEC is of therapeutic interest for the treatment of peritoneal

carcinomatosis, but its use remains infrequent because of potential toxicities (van Driel et al.,

2018; Goéré et al., 2013). Our data are in favor of adding PI3Kα inhibitors to this therapeutic

arsenal.

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Author contributions

Benoît Thibault: conceptualization; data curation; formal analysis; investigation; methodology; validation; visualization; writing – review & editing. **Adrien Thole:** data curation; formal analysis; investigation; methodology; visualization; writing – original draft; review & editing. **Céline Basset:** data curation; formal analysis; methodology; resources; visualization; writing – review & editing. **Julie Guillermet-Guibert:** conceptualization; data curation; formal analysis; funding acquisition; investigation; methodology; project administration; supervision; validation; visualization; writing – original draft; writing – review & editing.

Conflicts of interest

The authors have no conflicts of interest to declare.

 Figure legends:

Figure 1: Modeling tumor aggregates derived from ovarian cancer ascites. (A)

Immunohistochemistry of high-grade ovarian serous carcinoma from ascites fluid, in which

tumorospheres are indicated by arrows (100x magnification). PAP = Papanicolaou, MGG =

May-Grünwald-Giemsa, PAS = Periodic acid–Schiff. (B) Evolution of tumorosphere

morphology 1, 4 or 7 days after seeding. (C) Model of dissemination in serous cavities from

the primary tumor facilitated by the flow of liquid and formation of tumorospheres. (D)

Morphology and (E) area of SKOV-3-GFP (green fluorescent protein) tumorospheres.

Measurements with ZEN® software (x5 magnification, fluorescence, $n \geq 2$). Comparison of

the tumorosphere area with 5000 or 10000 SKOV-3-GFP cells on day 14 by Student t-test p

<0.01 (**). (F) GFP fluorescence intensity was evaluated at D1, D4 and D7 using ZEN®

software (in fluorescence, x5 magnification, n = 3). (G) Morphology of heterotypic

tumorospheres comprising 5000 SKOV-3-GFP cells (green) cultured with or without different

numbers of mesenchymal stem cells (MSCs) (5, 50, 500, 5000) 1, 4 or 7 days after seeding.

(H) Area and (I) GFP fluorescence intensity were evaluated at D1, D4 and D7 using ZEN®

software (in fluorescence, x5 magnification, n = 3). Mean +/- SEM (ANOVA test comparison

with the condition without MSCs at the corresponding observation day, $p < 0.001$ (***), p

< 0.01 (**), $p < 0.05$ (*)).

Figure 2: PI3Kα inhibition prevents formation and maintenance of ovarian cancer cell

tumorospheres. (A) SKOV-3 cells were serum-deprived overnight, pre-treated for 1 h with

specific PI3Kα inhibitors and then serum-activated for 15 minutes. Actin-standardized

quantification of the amount of (B) phospho-Akt (S473) and (C) phospho-Akt (T308) relative

to the vehicle condition (without treatment). (D) 5,000 SKOV-3-GFP cells that will form

tumoropheres at D1 were treated or not at D0 and D4 with A66 and BYL719 (1 and 10 μM).

(E, F) SKOV-3-GFP tumorospheres were treated according to the schedule in (D) and

morphology was observed 1, 4 or 7 days after seeding. (F) Area was evaluated at D1, D4 and

D7 using ZEN® software (x5 magnification), n ≥ 3, mean +/- SEM (ANOVA test comparison

with the vehicle condition of the corresponding observation day, $p < 0.001$ (***), $p < 0.01$ (**),

$p < 0.05$ (*)). (G) Tumorospheres formed of 5,000 SKOV-3-GFP cells were treated or not at

D1 and D4 with A66 or BYL719 (0.01, 0.1, 1 and 10 μM). (H) Tumorosphere morphology

was evaluated on D1, D4 and D7. (I, J) Tumorosphere area was evaluated on D1, D4 and D7

using ZEN® software (x5 magnification), n ≥ 3, mean +/- SEM (ANOVA test comparison

with the control condition of the corresponding observation day, $p < 0.001$ (***), $p < 0.01$ (**),

$p < 0.05$ (*)). (K) Tumorospheres formed of 5,000 SKOV-3-GFP cells (green) associated with

different ratios of MSCs (labeled in red using Quantum Dots) were produced and treated with

10 μM BYL719. Tumorosphere morphology was evaluated using ZEN® software (in

fluorescence, x5 magnification) on D7.

Figure 3: PI3Kα inhibition impacts multicellular cell viability. (A) SKOV-3, H460 and

MCF-7 cell monolayers were treated or not with A66 or BYL791 and living cells were

quantified after 3 days with a MTT colorimetric assay. (B-C) Tumorospheres formed of 5,000

SKOV-3-GFP, H460 and MCF-7 cells were treated or not at D1 and D4 with A66 or

BYL719. (B) Tumorosphere morphology was evaluated on D7 using ZEN® software (in

fluorescence, x5 magnification); living cells were labeled with calcein (white), n = 3. (C) The

area of living cells after treatment was quantified using ImageJ® software, n = 3. Mean +/-

SEM (ANOVA test comparison with the control condition of the corresponding observation

day, $p < 0.001$ (***), $p < 0.01$ (**), $p < 0.05$ (*)).

Figure 4: PI3K inhibitors decrease tumorosphere survival in the presence of cisplatin. Tumorospheres formed of 5,000 SKOV-3 cells were treated or not on D1 and D4 after seeding with a fixed concentration of 1 µM A66, BYL719, TGX-221 or AZD8186, in combination or not with increasing concentrations of cisplatin (0.05, 0.5, 5 µM). (A) Tumorosphere morphology and (C) area (n ≥ 3) were evaluated using ZEN® software (in fluorescence, x5 magnification). (B) Living cells were marked with calcein and (D) the living cell intensity after treatment was quantified using ImageJ® software (n = 2), mean +/- SEM (ANOVA test comparison with cisplatin concentration and the control of the corresponding observation day, $p < 0.001$ (***), $p < 0.01$ (**), $p < 0.05$ (*)).

Graphical abstract: Schematic representation of PI3Kα involvement in tumor cell aggregates from ascites. 1) Known involvement of PI3Kα in primary ovarian tumors. 2) PI3Kα participates in tumorosphere formation within the peritoneum (treatment with PI3Kα inhibitors causes a delay in the formation of clusters). 3) PI3Kα participates in the maintenance of tumorospheres and in resistance to conventional treatment for peritoneal carcinomatosis. PI3Kα is a target to prevent transcoelomic dissemination and maintenance of tumorospheres in patients with ovarian cancer.

MATERIAL AND METHODS

Patient ascites-derived cell tumorospheres

Ascites fluids were taken from 12 patients with ovarian cancer after consent (CRB Cancer des Hôpitaux de Toulouse, IUCT-O, Toulouse – BB-0033-00014, DC-2008-463, AC-2013-1955). Supplementary Table S1 details patient information; clinical and biological annotations of the samples have been declared to the CNIL (Comité National Informatique et Libertés).

pOVCA-1 and pOVCA-2 were ovarian tumor cells isolated from patients' ascites and cultivated in DMEM supplemented with 10% FCS (fetal calf serum), 1% L-glutamine, 1% penicillin-streptomycin and 0.01% plasmocin (Invitrogen).

Cell lines, 3D culture, treatments and phenotypic analysis

SKOV-3-GFP cells correspond to genetically modified SKOV-3 wild-type (WT) cells expressing GFP. MSCs were derived from healthy donors who underwent hip surgery (Etablissement Français du Sang). Cells were cultured at 37°C, 5% CO_2 in a humid atmosphere, in RPMI 1640 medium (Sigma) for SKOV-3 and H460 cells, or DMEM (Sigma) for MSCs and MCF-7 cells; 50% of each medium was used when MSCs and SKOV-3 cells were mixed. The culture medium was supplemented with 10% FCS (fetal calf serum), 1% L-glutamine, 1% penicillin-streptomycin and 0.01% plasmocin (Invitrogen).

One thousand patient-derived tumoral cells or 5000 SKOV-3-GFP, SKOV-3 WT cells, H460 or MCF-7 cells with or without increasing numbers of MSCs were seeded in round-bottom 96-well plates (Nunclon Sphera, ThermoFisher Scientific). Living cell numbers and metabolic activity were measured with an MTT assay or labeled for 1 h using the LIVE/DEAD® kit (ThermoFisher Scientific) containing 2 µM calcein and 4 µM EthD-1 (Ethidium homodimer-1). The cell clusters obtained were observed and imaged with an AxioVert microscope on days 1, 4,

380   7, 10 and 14 using ZEN® software (Carl Zeiss). The same software measured their diameter and

381   surface. For each condition, triplicates were performed and the average area was compared to the

382   day 1 average of the corresponding experiment (representing 100% of the area). Western blotting

383   was performed as described in our previous work (Baer et al., 2014) with the following

384   antibodies:

Primary antibody	Origin	Dilution	Provider	Size	Secondary antibody
pAkt (S473)	Rabbit	1/1000	Cell signaling #4060	60kDa	1/3300
pAkt (T308)	Rabbit	1/1000	Cell signaling #4056	60kDa	1/3300
Akt total	Rabbit	1/1000	Cell signaling #4691	60kDa	1/20 000
β-Actin	Mouse	1/2000	Sigma Aldrich AC74	42kDa	1/10 000
β-Actin	Mouse	1/2000	Sigma Aldrich AC74	42kDa	1/10 000

Pharmacological inhibitors

Treatments used in this study were as follows: Cisplatin (kindly supplied by Institut

Universitaire du Cancer – Oncopole (IUCT-O)), A66 (Axon Medchem) PI3Kα inhibitor (in vitro

IC50 in nM: p110α: 32; β: >12500; δ: >1250; γ: 3480) (Jamieson et al., 2011); BYL719 (Apex

Bio) PI3Kα inhibitor (in vitro IC50 in nM: p110α: 4.6; β: 1156; δ: 290; γ: 250) (Fritsch et al.,

2014).

Qdot® cell labeling of MSCs

Labeling was performed according to the supplier's recommendations, with the equimolar

mix from the labeling kit at dilution 1/100 incubated for 1 h at room temperature applied directly

to the cell cultures for 45 minutes.

Statistics

Quantitative variables are presented by means and qualitative data by percentages. Group comparisons were made using the ANOVA test (matched data) and Student's t-test for parametric tests. For all experiments, a statistical difference is reached for $p < 0.05$.

Supplementary Figure S1: Development of heterotypic tumorospheres. (A) Area and (B) morphology of MSC tumorospheres. Mean +/- SEM measurements with ZEN® software (x5 magnification, $n \geq 2$). (C) Observation of the formation of fibrous structures within MSC aggregates. (D) Observation of heterotypic tumorospheres of SKOV-3 cells (green) and MSCs (red). 5000 SKOV-3-GFP cells associated with different MSC ratios (500 or 5000) were observed at D1 and D7. (E, F) Spheres formed of 5000 MSCs were produced and treated or not with A66, BYL719 or cisplatin 24 h after seeding. (E) Morphology and (F) area of MSC tumorospheres were measured using ZEN® software (x5 magnification, $n = 3$).

Supplementary Figure S2: Effects of PI3K inhibitors on SKOV-3, H460 and MCF-7 tumorospheres. Tumorospheres formed of 5000 SKOV-3, H460 and MCF-7 cells were produced and treated or not with A66, BYL719 at D1 and D4 after seeding. (A) At D7, living cells were labeled with calcein (green) and dead cells with EthD-1 (red) using the LIVE/DEAD® kit. (B) The ratio of dead cells' fluorescence on living cells' fluorescence and compared to the vehicle condition. Mean +/- SEM, $n = 3$ (Student's t-test, comparison with the vehicle condition, $p < 0.01$ (**)).

Supplementary Table S1: Patient data. Data corresponding to the ascites samples isolated

from patients. Patient number, age (in years), histology, aggregate numbers (per microscopy

field) are represented.

Supplementary Table S2: Frequency of *PIK3CA* or p53 mutations in cell lines derived

from primary tumors versus serous fluid cavities, example of ovarian carcinoma. (A)

PIK3CA and p53 mutational status in ovarian cancer cell lines coming from a primary tumor

(tumor) or ascites fluid (ascites). The exact mutation is indicated when known. WT = wild

type. (B) Frequency of ovarian cancer cell lines harboring a PIK3CA and/or p53 mutation

depending on their origin (primary tumor or ascites). Mainly identified from Cellosaurus or

according to the literature (Beaufort et al., 2014).

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Figure 1

Figure 2

Figure 3

Figure 4

Primary site

1)

PI3Kα

Parietal layer

2)

Dissemination
through the flow

PI3Kα

3)

PI3Kα

Cohesive Tumorosphere

Parietal layer

Resistance to chemotherapies
Sensitive to PI3Kα inhibitors

Pre-metastatic stage in serous cavity

Graphical Abstract