Simultaneous Targeting Tumor Cells and Cancer-Associated Fibroblasts with a Paclitaxel-Hyaluronan Bioconjugate: *in vitro* Evaluation in Non-Melanoma Skin Cancer

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Abstract

Background: Cancers are complex organs which encompass not only the tumor cells but also cells within the surrounding stroma. Such cells, mainly cancer associated fibroblasts (CAFs) and immune infiltrating cells, facilitate many aspects of cancer development providing a structural and supportive framework rich of bioactive compounds. So far, there are emerging studies proposing the combination of conventional anti-cancer therapy, directed against neoplastic cells, to molecules targeting tumor microenvironment.

Methods: The study was designed to evaluated the pharmacological properties of the anti-tumor agent paclitaxel conjugated to hyaluronic acid (HA) on non-melanoma skin cancer (NMSC) and on the surrounding dermal fibroblasts. This molecule, named Oncod-P20 (Onco-P20), targets preferentially cells expressing high level of CD44, the natural ligand of HA.

Results: Consistent with paclitaxel's mechanism of action involving interferences with the normal breakdown of microtubules during cell division, highly sensible carcinoma cells underwent rapidly to apoptotic cell death. Interestingly, less sensible cells such as dermal fibroblasts resisted to the Onco-P20 treatment and experienced a senescent-like status characterized by morphological change, decreased proliferation, and significant modification of the gene expression profile. Onco-P20-treated fibroblasts lowered growth factors production, down-modulate Wnt signal pathway and acquired a pro-inflammatory profile. Independently to the direct exposure to taxol, in presence of Onco-P20-treated fibroblasts or their conditioned medium, carcinoma cells reduced the proliferation rate. Similar to NHF, fibroblasts isolated from skin cancer tumor lesion or from tissue adjacent to the tumor acquired an anti-neoplastic activity under Oncod-P20 treatment.

Conclusion: Collectively, our data demonstrated that Onco-P20, exerting both a direct and a NHF-mediated indirect paracrine effect on carcinoma cells, is a good candidate for an innovative therapy alternative to surgery for treatment of NMSC.

Background

Keratinocytes-derived non-melanoma skin cancer (NMSC), predominantly basal and squamous cell carcinoma, represents the most frequent malignant tumor worldwide [1]. Although they rarely metastasized, their locally aggressive invasion frequently mutilates patients. New results from investigation indicate the importance of tumor microenvironment on the biological properties of tumor, such as the aggressive growth and local recurrence. During malignant transformation, cancer cells acquire invasive properties, violate the basement membrane and invade the underlying dermis. Consequently, dermal stroma and epidermal cancer cells establish unusual heterotypic cell-cell contact and a dynamic cross-talk that leads to alterations of the host tissue [2, 3]. The parenchymal injury associated with a nascent and growing tumor constantly activated stromal cells driving the acquisition of cancer associated phenotype, that neither spontaneously revert to a normal phenotype nor undergoes cell death [4]. A particular feature of cancer stroma is the increased number of mesenchymal or fibroblastic cell type, pathologically activated referred as cancer associated fibroblast (CAFs). CAFs are embedded within an extracellular matrix (ECM) profoundly remodeled compared to the physiological one and rich of immune cell infiltrates and blood/lymphatic vessels. The ECM not only provides structural support for the cells but also participate to biochemical signal transduction through the binding of angiogenic and growth factors whose availability depends also on ECM remodeling [5-7]. CAFs are characterized by a distinct activated phenotype and by expression of a variety of markers, such as fibroblast activated protein-α (FAP-α), platelet-derived growth factor receptor-α and β (PDGFR-α and β), α-smooth muscle actin (α-SMA) and fibroblast-specific protein-1 (FSP1) [8-11]. CAFs in cancer tissue are morphologically similar to myofibroblasts, which are large spindle-shaped cells activated during the wound healing process and found in sites of chronic inflammation [12]. Based on similarity between tumor initiation and wound inflammation linked to myofibroblasts activation tumor has been defined as "wound that never heal" [13]. This notion implies the reversibility of fibroblasts activation and the possibility to target CAFs for cancer therapy [14]. Interestingly, not only fibroblasts within or directly in contact to tumor, but also those from cancer free tissue adjacent to tumor exhibit CAF phenotype [15]. This phenomenon could be explained by the intense release of bioactive molecules by tumor tissue and the consequent bi-directional dynamic cross-talk among cancer cells and surrounding cells. It has been suggested that the high secretory phenotype of CAFs play a key role in tumor progression, but increasing data also argue for their antitumor actions [16, 17]. In mouse model, reduced stromal content accelerates tumor growth and angiogenesis and full depletion of CAFs induces immunosuppression [18, 19]. The capacity to sustain cancer depends largely by altered secretome, augmented pro-mitogenic peptide, regulation of inflammation and ECM whereas the capacity to inhibit cancer is supposed to predominantly depends on the interaction with the immune system. An increased production of a vast repertoire of growth factors has been reported overlapping amongst distinct tumor types including NMSC [4, 20-23]. Among these, VEGF, PDGFs, EGF, FGFs, and Wnts drive tumor growth and vasculature [24]. CAFs also secrete cytokines and chemokines that recruit and modulate the function of immune cells in the tumor microenvironment [7, 25]. The peculiarity of the inflammation associated to tumorigenesis processes demonstrates chronic, non-resolving characteristics [25]. However, similar to mesenchymal cells, inflammatory cells operate in conflicting modalities: both tumor-antagonizing and tumor-promoting [26]. The balance between pro-inflammatory and anti-inflammatory response is deeply implicated in patient's prognosis [27, 28]. It is not fully clear if the dual nature of cancer microenvironment reflects the contemporary presence of heterogeneous populations or if differences reside in disease evolution. Thus, since CAFs co-evolve with genetically altered tumorigenic cells it is possible that an early anti-tumor phenotype is replaced by a pro-tumorigenic one during disease progression. In line with the idea that CAFs co-evolve with tumor cells, it has been demonstrated that normal dermal fibroblasts could be "educated" to assume CAFs gene signature by tumorigenic cells both in vivo and in vitro [29]. On the other hand, in the skin, chronic and long-term exposure to UV radiation, the major environmental risk factor for skin cancer [30], occurs on whole tissue impacting on pre-cancerogenous and resident stromal cells at the same time. Thus, differentially to internal organs, skin fibroblasts undergoes to a continuous extrinsic stimulation. Therefore, in the skin, alteration in the stroma can precede (or act independently) to epithelial cell alterations moving as a driver of the tumorigenic process. Thus, activated fibroblasts may play a relevant role in both initiating and progression of skin carcinogenesis. Consequently, in order to control cancer, we need to focus not only on the malignant cancer cells, but also on the benign microenvironment. In this study, we investigated the effect of the anti-tumor agent paclitaxel (PTX) conjugated to hyaluronic acid (HA), the CD44 ligand, a molecule named Oncod-P20 (Onco-P20) on skin carcinoma cells, dermal fibroblasts and CAFs. PTX, a microtubule-targeting drug, enhances the polymerization of tubulin and also interacts directly with microtubules stabilizing them against depolymerization and enabling cells to form a normal mitotic apparatus [31]. Penetration-enhancing strategies including liposomes,
nanoparticles, implants and albumin-bound PTX have been proposed to bypass the poor water solubility of taxols. Improved skin penetration is desirable due to the proved efficacy of PTX against melanoma and non-melanoma skin cancer [32-34]. HA-conjugate is a highly biocompatible strategy to increase bioavailability and pharmacokinetic of active molecules especially in tissue presenting high hyaluronan receptor density such as the skin. Moreover, in many cancers of epithelial origin has been observed the up-regulation of CD44 [35]. In addition, CD44 is abundantly expressed by CAFs particularly in the hypoxic contest [36]. It has been previously demonstrated that dermal fibroblasts primed with PTX are capable release functional active PTX reducing melanoma cells [37]. In line with this set of preliminary data, using Onco-P20, we analyzed the impact of PTX treatment in the context of NMSC. We demonstrated, Onco-P20 combines in a single molecule two synergetic activities: direct anti-tumor efficacy on keratinocyte-derived skin cancer and an indirect anti-tumor activity dependent on the modulation of CAFs secretory profile. Based on HA-CD44 interaction, Onco-P20 targets fibroblasts more efficiently than free PTX representing a valid alternative to surgery for treatment of non-melanoma skin carcinoma.

Methods

Ethic statement. The Declaration of Helsinki Principles followed and patients gave written informed consent to collect samples of human material for research. Further, the Institutional Research Ethics Committee (Istituti Regina Elena e San Gallicano) approved all research activities involving human subjects.

Preparation of hyaluronan-paclitaxel bioconjugate. The preparation of HA-paclitaxel bioconjugate (Oncofide-P20) with ~20% w/w of paclitaxel (taxol) loading has been previously described [38]. Oncofide-P20 and paclitaxel (donated by Fidia Farmaceutici, Abano Terme, Italy) were diluted with DMSO. Solutions were further diluted at each experimental day in order to achieve a 0.05% final DMSO concentration.

Cell cultures. Primary cultures of normal human keratinocytes (NHK) and fibroblasts (NHF) were isolated from human skin fragments obtained from surgery. NHK were maintained in M154 medium with Human Keratinocytes Growth Supplement (HKGS) supplements (Cascade Biologics Inc., Portland, OR USA), Ca$^{2+}$ (0.07 mM) and antibiotics; NHF were maintained in DMEM (EuroClone S.p.A., Milan Italy) supplemented with 10% FBS and antibiotics (Hyclone Laboratories, South Logan, UT, USA). Cancer-associated fibroblasts (lesional and perilesional) and paired normal fibroblasts were isolated from 5 tumor samples of patients, 4 SCC and 2 BBC using the same methods. Carcinoma cells were washed three times with PBS, then minced into approximately 1-2 mm$^2$ sized pieces and digested using 5 mL of a 0.1% trypsin solution (Gibco, CA, USA). After variable digestion time (2-6 hours), the homogenate was collected and passed through a 70µm strainer and cultured in M154 plus HKGS. A431 and SCC1300-UC cell lines were cultured in DMEM containing 10% FBS or in M154 plus supplements where indicated. Images were recorded using an Axiosvert 25 inverted microscope (Carl Zeiss, Oberkochen, Germany) and a Power Shot G5 digital camera (Canon, Inc., Tokyo, Japan).

Proliferation assays. Briefly, 2.5 x 10$^4$ keratinocyte or 0.8 x 10$^4$ fibroblasts were seeded into the 24-well plates for 24 hours to adhere. Then, growth medium was changed with fresh medium containing treatments (or not for control cells) at the appropriate concentrations. Culture medium and drugs were refreshed twice a week. At the experimental end point cells were incubated with 3-(4,5 dimethylthiazol-2)-5-diphenyl tetrazolium bromide (MTT) for 2 hours. After this time, the medium was removed and the resulting crystals were solubilized in DMSO. The absorbance was measured at 570 nm with a reference wavelength of 690 nm. Absorbance readings were subtracted from the value of blank wells, and results were calculated as a percentage of absorbance respect to control samples. Experiments were performed in duplicates.

Preparation of CAF conditioned medium (CM). Human skin CAFs were plated into a 10 cm$^2$ dish and a conventional culture was carried out for 24 hours before Onco-P20 treatment. Subsequently, fresh DMEM + 10% FBS containing treatment (or not for control cells) was replaced twice a week. After 2 weeks and an additional 48 h period without treatment, medium was replaced with DMEM without FBS before CM harvesting. The supernatant was collected, centrifuged at 1,000 rpm, filtered with a 0.22-µm membrane for sterilization. CM of untreated fibroblasts was used as control medium. Experiments were performed in duplicates.

Trans-well co-culture. Briefly, 12 mm trans-well with 3.0 mm pore membrane insert were used with CAFs and cancers cells alternatively in the lower or upper compartment. 2.0 x 10$^4$ fibroblasts per well were seeded in the lower compartment or alternatively 0.8 x 10$^4$ fibroblasts were seeded in upper compartment and incubated at 37°C 5% CO2 overnight. Then, fresh DMEM + 10% FBS containing treatment (or not) was replaced twice a week. After 2 weeks DMEM was replaced with M154 without treatment. After 48 hours, trans-well inserts containing 2.0 x 10$^4$ carcinoma cells were added. Otherwise, 2.0 x 10$^4$ carcinoma cells were seeded in lower compartment before insert addition. After 96 hours MTT assay was developed. Experiments were performed in duplicates.

Western blot analysis: Cell extracts were prepared with RIPA buffer containing proteases and phosphatases inhibitors. Proteins were separated on SDS-polyacrylamide gels, transferred to nitrocellulose membranes and then treated with the following primary antibodies: mouse monoclonal anti-b-catenin, anti-cyclinE (1:1000) (Santa Cruz Biotechnology Inc., Santa Cruz, CA, USA), anti-bFGF (1:500) (Upstate Biotechnology, Inc., USA), anti-p53, anti-cyclinD1 (1:1000) (Dako, Agilent Technology Italia, Milan, Italy), rabbit polyclonal anti-p21, anti-p27 (1:500), anti-cyclinB1 (1:1000) (Cell Signaling Biotechnology), and goat anti-KGF (1:200), (Santa Cruz Biotechnology) antibodies. Anti-β-actin mouse monoclonal antibody (1:5000) (Sigma Aldrich, Merck KGaA, Darmstadt, Germany) was used to normalize protein content. Horseradish peroxide-conjugated goat anti-mouse, goat anti-rabbit or bovine anti-goat secondary antibody complexes were detected by chemiluminescence (Cell Signalling Technology, MA, USA). Imaging and densitometric analysis were performed with UVITEC Mini HD9 acquisition system (Alliance UVITec Ltd, Cambridge, UK).

Elisa assay. Growth factors and bioactive molecules concentration released in culture medium were measured after 2 weeks treatment using commercially available enzyme-linked immunosorbent assay (ELISA) kits according with manufacturer's instructions: HGF (Cusabio Technology LLC, Baltimore, MD, USA), IGFBP4 and 6 (Aviva Systems, Biology, CA, USA) and VEGF-A (eBioscience, Inc. San Diego, CA, USA). Cells were incubated for 48 hours with DMEM without FBS before medium collection. Results were normalized against protein concentration.
Immunofluorescence analysis. Cells on coverslips were fixed with 4% paraformaldehyde for 20 minutes at room temperature followed by 0.1% Triton X-100 to allow cell permeabilization. Cells were then incubated with anti-PML (1:500), for 1 hour. Primary antibodies were visualized using anti-rabbit IgG Alexa Fluor 488 (BD Biosciences, Milan, Italy). Nuclei were visualized with 4',6'-diamino-2-phenylindole (DAPI). Fluorescence signals were recorded using a CCD camera (Zeiss, Oberkochen, Germany).

Flow cytometry analysis. Cell death and apoptosis were analyzed by annexin-V FITC/propidium iodide (PI) double staining method after 48 hours of treatment. Cells were harvested by trypsinization, suspended in the staining buffer (10 mm HEPES / NaOH, pH 7.4, 140 mm NaCl, 2.5 mm CaCl₂), stained with FITC-labeled annexin V and PI for 15 min at RT in the dark and then kept on ice until be analyzed. For cell cycle distribution, after cells harvesting by trypsinization, cells were fixed in cold 70% ethanol for 10 minutes and then stained with propidium iodide (PI) solution (1 µg/µL PI and 0.125% RNaseA; Sigma Aldrich, St. Louis, MO) at room temperature for 15 minutes. 20,000/sample cells were analyzed using a FACS Calibur instrument (BD) equipped with a 488 nm argon ion laser. The percentage of cells in each phase of the cell cycle was determined using FlowJo software v8.0. The amount of membrane bound CD44 was measured incubating unfixed cells with a CD44-PE antibody (BD Biosciences) before cells wash and cytofluorimetric analysis.

Cytokines protein array. The expression of 20 human cytokines was analyzed using a commercially available antibody array system (RayBio® C-Series Human Inflammation Array C1 Map, RayBiotech, Inc) that uses membrane-bound cytokine-specific antibodies to capture cytokines in biological fluids. The procedure was performed as per the manufacturer's instructions. Cells were seeded in 10-cm culture dishes and treated (or not) with Onco-P20 0.15 mg/mL for 2 weeks. After removing drug cells were maintained in serum-free medium for 48 hours. The cytokine array membranes were blocked in 2 ml 1 x blocking buffer for 30 min at room temperature (RT) and then were incubated with 1 ml of conditioned medium at 4°C overnight. The medium was then decanted from each container, and the membranes were washed three times with 2 ml 1 x wash buffer I, followed by two washes with 2 ml 1 x wash buffer II at room temperature. Next, the membranes were incubated in biotin-conjugated primary antibodies for 2 hours at RT and then washed as described above before incubation in 1:1000-diluted horseradish peroxidase-conjugated streptavidin for 2 hours. The membranes were then washed thoroughly and incubated with a chemiluminescent ECL substrate at RT for 5 min. Imaging and densitometric analysis were performed with UVITEC Mini HD9 acquisition system (Alliance UVItc Ltd; Cambridge, UK).

Statistical Analysis: Results in the figures are representative of several experiments we performed with at least five cell lines from different donors. Quantitative data were reported as mean ± standard deviation (SD). Student t-test was used to assess statistical significance with thresholds of *p ≤0.05 and **p ≤0.01.

Result

Effect of Oncod-P20 treatment on cell proliferation and viability

A431 human squamous carcinoma cells, a commonly model used for studies of skin cancer and normal keratinocytes (NHK) from healthy subjects were treated with a broad panel of Onco-P20 concentration (0.05-1 µg/mL) for 72 hours before MTT assay and cells count. MTT assay showed a strong reduction of NAD(P)H-dependent cellular mitochondrial activity in Onco-P20-treated carcinoma cell cultures that correlated to a significant reduction in the number of cancer cell at all the concentrations tested (Fig.1a and b). In contrast, at the experimental end point in NHK Onco-P20 treatment mildly impacted on metabolic function and on the number of cells resulted modestly reduced (Fig.1c and d). Hyaluronic acid, alone did nor exerted any significant modification of cell viability and proliferation indicating that this portion of the molecule is not implicated in the biological activity observed. Since in addition to cell count, microscopic observation of cell cultures highlighted marked differences comparing normal and tumor cells (Fig. 1e), we additionally evaluated cell cycle distribution demonstrating G2/M accumulation in both cell types and extensive cell death only in cancer cells (Fig. 1f). Significant increase in the number of apoptotic cells was confirmed by AnnexinV/PI staining proving the selective cytotoxic effect on tumor cells (Fig. 1g). Thus, while in carcinoma cultures metabolic activity measurement and the number of apoptotic cells correlated indicating that cell death is the main mechanism of Onco-P20 function, in normal keratinocyte, all the concentrations tested, the treatment reduced the metabolic activity without affecting cell viability. However, high doses of Onco-P20 (>0.25 µg/mL) impacted on keratinocyte proliferation capability as evidenced by cell accumulation in G2/M cell cycle phases.

Since established cell lines maintained in culture for a long period frequently present evidence of deviations from the phenotype of the originating tumor, we used a collection of specimens from patients well characterized for disease stage and follow up to isolate low passage tumor cell lines and patient-matched normal keratinocytes cultures. In this case, cytotoxic effect against short term carcinoma cell cultures showed an overall lighter intensity compared to A431 cells and the MTT assay, that does not discriminate between cytotoxic and cytostatic effect, displayed modest difference between normal and carcinoma cells (Fig. 2a and b). By contrast, AnnexinV/PI staining confirmed that Onco-P20 preferentially reduced viability of neoplastic cells (Fig. 2b). Since A431 cell cultures and fresh isolated carcinoma cells were cultured in non-overlapping growth condition (DMEM supplemented with FBS and chemically-defined medium M154 supplemented with HKGS respectively; see material and method section), we addressed the question if the observed differences could be attributed to culture medium composition or to distinctive biological properties. In this case, we used A431 and SCC1300-UC, one of the low passage SCC cell lines isolated in our laboratory capable to proliferate both in defined media (M154 plus HKGS) as well as in high calcium FBS-containing DMEM medium, to evaluate the impact of cell medium composition on Onco-P20 treatment. As shown in Suppl. Fig. 1, the sensibility to Onco-P20 treatment resulted attenuated in M154 defined medium compared to DMEM plus serum (Fig. S1a) and inversely correlated to the proliferation rate (Fig. S1b). In fact, cell culture composition influenced cell growth demonstrating a significant slower growth curve kinetic in chemically defined medium that correlated with lower pharmacological activity. However, responsiveness to Onco-P20 is also an intrinsic characteristic of different cell lines related to the CD44 level of expression that is not necessarily over-expressed in skin carcinoma compared to normal cells [39]. In fact, FACS analysis confirmed that patient-derived carcinoma specimens present a moderate level of CD44 expression comprised between those of normal cells and one of A431 cells (Fig. 3a). The low presence of CD44 on the membrane could partially explain the higher response of A431 that could be still appreciated using identical growth condition (Fig. 3a). Accordingly, with the idea that the relative abundance of CD44 is not particularly elevated in the keratinocyte lineage, hyaluronan-conjugate formulation failed to demonstrate any significant gain of
function in comparison to conventional free paclitaxel (Fig. S2).

CD44 overexpressing cells comprise not only tumor cells and cancer stem cells but also CAFs [40-42]. Strongly CD44-positive CAFs are particularly frequent in tumor hypoxic areas [36]. Since also dermal fibroblasts physiologically express high level of this receptor (Fig 3b) we evaluated the effect of Oncop-P20 on normal human fibroblasts (NHF). For this purpose, and in line with patient’s characteristics, we used adult fibroblasts isolated from photo-exposed area of geriatric donors. NHF treated with Oncop-P20 (0.05-1.0 µg/mL) for 72 hours displayed attenuation of mitochondrial activity and a significant abatement in cellular proliferation compared to control cells (Fig.4a and b). Similar to normal keratinocytes, fibroblasts did not show acute cytotoxicity as measured by AnnexinV/PI double staining (Fig. 4c). Cell cycle analysis evidenced a marked accumulation in the G2/M phase (Fig. 4d). In addition, evidences of cell hypertrophy (a large cell morphology), a universal marker of in vitro cell senescence [43] appeared at days 3-5 (Fig.4e). Increase expression of p53 and p21, as well as, the presence of several promyelocytic leukemia protein (PML)-nuclear bodies, typical nuclear matrix structures implicated in the induction of the senescence process [44], confirmed diminished proliferative propensity of Oncop-P20-treated cells (Fig. 3f and g). In contrast to p21, another important cell cycle checkpoint CDK inhibitor, p27 resulted decreased. Cyclin E, that is required and rate limiting for S phase entry, is lightly downregulated. We further analyzed the protein level of cyclins in Oncofd-P20-treated cells. The level of cyclin D1 which promotes progression through G1/S phases was elevated, while cyclin B1, promoting G2/M transition, was barely present [45, 46]. According with previous studies [47, 48] the cyclins profile was consistent with G2-arrest. However, fibroblasts seem to be long-term arrested rather than irreversible senescence since after treatment removal cells resumed proliferation even in case of prolonged exposure to Oncop-P20 (Fig. S4).

Finally, Oncop-P20 activity was compared with that of equal amount of free paclitaxel. Following 72 hours drugs exposure, unconjugated paclitaxel exhibited a significant lower effect on NHF (Fig.5a). Thus, due to high level of CD44 expression on the membrane surface of mesenchymal cells, Oncop-P20 exerts a stronger pharmacological activity then free paclitaxel. Moreover, to simulate drug availability in the case of topical application on patient’s skin, we exposed NHF to treatments (Oncop-P20 or paclitaxel) for a defined short period (8 hours) before wash out and subsequent cell culture in drug-free medium. In this case, using the same end point of previous experiments we observed reduced proliferation exclusively in Oncop-P20-treated cells, whereas paclitaxel failed to modified fibroblast proliferation (Fig.5b). By contrast, in carcinoma cells no differences were observed comparing Oncop-P20 and paclitaxel using short period treatment (Fig. 5c). However, it is interesting to note that in this case both molecules exerted a mild effect suggesting or an augmented sensibility to taxol of carcinoma cells compared to fibroblasts or an accelerated trigger of drugs effect.

**Oncop-P20 deeply modifies fibroblasts gene and protein expression profile**

Given that fibroblasts could impact keratinocytes homeostasis through diffusible molecules, we assessed the impact of Oncop-P20 on the expression of secreted proteins involved in tumor progression. Under these circumstances, several growth factors were differentially expressed in Oncop-P20-treated fibroblasts versus control samples (Table 1). Among these, HGF was strongly down-regulated both at mRNA and protein levels (Fig. 6a). Additionally, the high molecular weight of bFGF, frequently associated with a poor prognosis in various human cancer [49], was reduced by Oncop-P20 (Fig. 6b), whereas its mRNA resulted unmodified. However, since high molecular weight isoforms of bFGF are associated to highly proliferative phenotype of fibroblast [50], it is possible that the observed difference reflects an autocrine control of cell growth. Similarly, KGF resulted decrease mostly at protein level. EGF transcript was near to the detection limit in most of the fibroblasts cultures and lightly reduced by the treatment. The IGF was unchanged whereas, members of insulin-like growth factor binding proteins (IGFBPs) superfamily, a group of secreted proteins capable to binding IGFs and to modulate the mitogenic, anti-apoptotic and metabolic actions of IGFs, were found strongly increased. In particular, IGFBP3, IGFBP4, IGFBP5, IGFBP6 and IGFBP7 mRNA were significantly up-modulated. Immunoenzymatic quantification of IGFBP4 and 6 confirmed gene expression analysis (Fig. 6a). Possible repression of neo-angiogenesis is supported by reduced expression of vascular endothelial growth factors (VEGF) at mRNA and protein level (Fig. 6b).

The Wnt/β-catenin signaling pathway, known to be activated in CAFs [51, 52], resulted down-modulated in Oncop-P20-treated fibroblasts as demonstrated by decreased of β-catenin, the pivotal molecule of the Wnt signaling pathway (Fig. 6a). Accordingly, the expression of some negative regulators of the pathway, Wnt5a, DKK1 and SFRP2, were increased suggesting an autocrine mechanism of Wnt signaling regulation. By contrast, TGFβ expression resulted unchanged. Among CAF’s markers, PDGFα and b were reduced whereas the expression of FAP1 and αSMA resulted increased and unmodified respectively. Strong increase of IL1α, IL1b, IL6 and IL8 argues for the activation of inflammatory pathways. To further investigate signal transduction pathways involved in cancer-fibroblasts cross-talk, we investigated a wide range of cytokines and chemokines using gene expression array cards and an antibody membrane array. Among the 93 mRNA studied, 38 were in most or all the sample undetectable and were discarded from the analysis, 13 resulted unmodified by the treatment, 2 lightly decreased and 40 enhanced (≥1.5 fold-increase). Of these highly expressed transcripts 17 reached the statistical significance (p≤0.05 or p≤0.01) (Table 2). Most of the protein up-regulated, including KLK14 and members of MAP kinase family, such as MAPK8 (JNK1) and MAPK14 (p38α), are implicated in the IL1 pathway. The expression of IL1R1 in tumor microenvironment, the main receptor of IL1α and b, due to the interconnection with NF-KB and MAP kinase pathways plays contrasting roles in different tumor stage [53]. Sustained PTGS2 (cox-2) expression confirmed pro-inflammatory profile of NHF Oncop-P20-induced phenotype. Annexin1A (ANXA1), increased by Oncop-P20, has been proposed as a tumor suppressor in head and neck squamous cell carcinoma [54]. However, in prostatic cancer, ANXA1 exacerbatated expression of this marker has been correlated to high amount of cancer stem cells [55]. Interestingly, CD40 an antigen frequently loss by basal and squamous cell carcinoma during tumor escape from activated T cells, was augmented by Oncop-P20 treatment. Intercellular adhesion molecule-1 (ICAM1, CD54), a receptor that support leukocytes accumulation in inflamed tissue, resulted strongly enhanced whereas vascular cell adhesion molecule-1 (VCAM), another mediator of leukocytes trafficking, was lightly increased. Additionally, high LTA4H that catalyzes the hydrolysis of epoxide LTA4 to LTB4, which mainly functions as a neutrophil, macrophage, and T lymphocyte chemoattractant, confirmed the inflammation-enhancing effect of Oncop-P20. The antibody array used to quantified secreted inflammatory factors confirmed a significant rise in IL6 and ILB production whereas IL1α and ILb appeared moderately increased at protein level. Additionally, the secretion of CCL11 (Eotaxin-1), a chemokine implicated in eosinophils recruitment [56], was up-regulated by Oncop-P20 (Table 3). Lastly, CCL1, whose release by fibroblast has been linked to bladder cancer cells invasion [57], significantly diminished.
Secretome of Onco-P20-treated fibroblasts modulates carcinoma cells growth

Cancer and stromal cells communicate mainly by a complex bidirectional cross-talk that evolve during disease progression and the diffusion of soluble factors through basement membrane. To study the effect of Onco-P20-induced fibroblast phenotype on cancer cells simulating the in vivo situation, we used two different experimental systems: fibroblast pre-treated conditioned medium (CM) and the trans-well permeable support (TW) to growth tumor and stromal cells on two different but connecting monolayer. Taking in consideration that previous studies evidenced that mesenchymal cells are capable to release into the microenvironment functional active paclitaxel for 24 hours [37], in both cases we replaced Onco-P20 containing medium with fresh starved medium 48 hours before collect CM or start co-culture experiments. In either way, due to the absence of paclitaxel and serum, cell proliferation depends exclusively by autocrine and paracrine activity of both cell populations. At the experimental end point (72 hours), both systems clearly demonstrated that carcinoma cells dose-dependently slowed the proliferation rate in presence of Onco-P20-pretreated fibroblasts o their CM (Fig.7a-and b). The anti-proliferative effects observed depends on the dose of drug used to treat NHFs and on the relative fibroblasts-carcinoma ratio as demonstrated by the stronger reduction of A431 cells growth in the trans-well upper (smaller) compartment compared to the lower one. Further, co-culture data were confirmed using a patient derived model assembled with NAF, CAF and carcinoma cells from the same donor (Fig.7c) or NAF, CAF and A431 cell line for cases presenting unsuccessful isolation of tumor cells in vitro (Fig.7d). Overall, data demonstrated that the sensibility to Onco-P20 of CAFs is similar to normal fibroblasts.

Discussion

Skin diseases represent a significant health burden. Although epithelial skin cancers rarely metastasize, they grow invasively and their removal often leads to severe functional and aesthetic impairments [58]. Therefore, there is a strong need for the development of cost effective and well tolerated therapies. The use of topical pharmacologic intervention aims to treat superficial form of NMSC decreasing the rate of disease progression and to offer an effective nonsurgical prospective. Increasing studies demonstrated that in tumors, the stroma is not simply a collection of enclosing cells without malignant function [7, 59]. CAFs have been indicated as a key component of the cross-talk between cancer cells and their microenvironment. A dichotomist role has been attributed to CAFs. It is generally accepted that CAFs stimulate cancer cell growth and metastatic behavior [18, 60]. Substantially, cancer cells utilize the CAF-secreted growth factors to facilitate their own survival and proliferation. However, depletion of CAFs led to invasive, undifferentiated tumors with enhanced hypoxia, and cancer stem cells, with reduced animal survival [18, 19]. Thus, based on the idea that fibroblasts have an exceptional phenotypic plasticity, including its secretory phenotype, it has been proposed that CAFs could be re-oriented to an anti-tumor phenotype for therapeutically purpose. However, to efficiently direct CAFs activity against tumor we need to promote exclusively antineoplastic properties. Recent studies demonstrated that CAFs are important modulators of the anti-tumor immune response [61, 62]. Consequently, among the potential approaches targeting the regulation of fibroblast-immune cells cross-talk represents and emerging strategy. Here, we demonstrated that Paclitaxel-Hyaluronan conjugate interacts with CD44, enters cells through a receptor mediated mechanism, and exerts a concentration-dependent inhibitory effect against tumor cells [38, 63, 64]. In preclinical studies, HA-taxol has demonstrated reduced toxicity, enhanced tumor accumulation, a greater anti-tumor efficacy compared to free taxol in ovarian and bladder carcinoma [63, 65]. However, in line with previous studies we observed that during the carcinogenic process, keratinocytes lightly increase the level of CD44 on the membrane and consequently Onco-P20 failed to demonstrate any significant advantage in the pharmacological activity compared to free paclitaxel in this cell type. Thus, we postulated that the modest augmented receptor-mediated uptake is balanced by smaller size and lipophilic nature of free paclitaxel that easily diffuses across cell membrane in vitro [66-68]. Although, the absence of any advantage in receptor-mediated uptake, the use of a hydrophilic HA backbone is still of interest in vivo to overcome the limited aqueous solubility of paclitaxel. Independently to the kinetic of drug internalization, tumor cells under tubulin polymerization/depolymerization anti-cancer agent treatment undertake the apoptotic cell death, whereas normal cells resist adopting a non-proliferative state [69, 70]. The presence of functional p53 plays a crucial role in the level of cytotoxicity by paclitaxel. Cells lacking wild-type p53 respond through a p53-independent apoptosis, whereas cells expressing active p53 resist paclitaxel toxicity by growth arrest [71, 72]. Consistently, we observed a divergent destiny comparing normal and transformed keratinocytes. NHK resist to paclitaxel exposure increasing G2/M arrest. In cells of mesenchymal origin, the switch to a quiescence state following paclitaxel exposure has been already described [73]. However, studies mainly focused on mesenchymal stem cells and on the possibility to use paclitaxel resistant cells as a reservoir of active compound (as a Trojan horse) for local-regional drug delivery [74]. Very recently, Coccè and co-workers, demonstrated that gingival mesenchymal stem cells treated with paclitaxel release vesicle containing an anti-cancer secreteme in addition to free paclitaxel [75].

Here, we demonstrated that Onco-P20-treated dermal fibroblasts present a consistent reorganization of several intracellular signaling. Most of the detected markers are referred to the block of cell cycle progression leading to a stress-induced non-proliferating status resembling a senescent-like phenotype. The observed G2 phase accumulation is a reversible state that at least in vitro does not necessarily fully corresponds to senescence, since after taxol wash out cells re-started to proliferate. This is an important point in consideration of the impact of treatment on tissue integrity and its long-term persistence. In fact, rapidly replaced cells, such as keratinocytes are naturally protected to accumulation of dysfunctional cells, due to normal turnover by new differentiating cells whereas, dermal fibroblasts are long-lived elements capable of proliferation under appropriate stimuli [76, 77].

The presence of senescent cells in premalignant lesions in various mouse tumor models and human patients [78, 79] is consistent with cellular senescence acting as a brake to the development of cancer. Hence, tumor cells must bypass the mechanisms that impose cellular senescence response in order to proliferate [14, 80]. Senescent-associated phenotype sensitizes surrounding cells to senesce. Thus, it is possible that cancer cells senescence, the first line of cancer protection, rapidly spread out to adjacent stromal cells. Accordingly, fibroblast senescence is a very early event in the carcinogenic process [81]. Whilst senescence acts as a tumor suppressor when activated in the epithelial compartment, the senescence of fibroblasts is frequently referred as a characteristic of CAFs suggesting that the associated secretome could play a role in cancer progression. In apparent contradiction to the concept that fibroblasts in the tumor contest resemble senescent fibroblasts, CAF is reported as a hyper activated state associated to an augmented proliferation rate [82-84]. Moreover, CAFs are considered similar to myofibroblasts activated after injury, a situation that promote mesenchymal cells expansion rather than proliferative arrest as in the case
of Onco-P20 treated cells. In addition, reprogramming activated fibroblasts or CAFs back into their dormant state has been proposed strategy for impairing tumorigenesis [16]. Thus, even if Onco-P20-treated fibroblasts display some biomarkers of senescence the observed changes in gene expression do not fully overlap with those of senescence phenotype of CAFs. One of the detrimental aspect of CAFs is the over-production and secretion of growth factors which can induce proliferative signals within cancer cells. By contrast, dermal fibroblasts under continuous treatment with Onco-P20 decrease the production of some growth factors. Notably, we observed down-regulation of HGF and simultaneous increase of most of the IGFBPs that suggests a reduced bioavailability of IGF. Both HGF and IGF stimulate directly and indirectly (supporting angiogenesis) tumor growth [77, 85]. HGF facilitates transformed epithelial cells migration and protect cancer cells from chemotherapeutic agents [77, 86, 87]. Moreover, IGF1 and IGF2 mediate cancer cells resistance to paclitaxel in murine model [88] suggesting that lowered IGF biodisponibility could heightens chemotheraphy. Interestingly, IGF1 diminishes in senescent and physiological aged dermal fibroblasts [89, 90] whereas its production is augmented in CAFs [91] confirming that CAFs are not simply senescent fibroblasts. Also KGF, whose overexpression by fibroblasts enhances proliferation of basal keratinocytes and suppress epidermal cells terminal differentiation [92-94], decrease in presence of Onco-P20. Minor production of growth factors may represent an autocrine control of proliferation when cell cycle progression is not possible, a situation that in the contest of carcinogenesis could play an important role in paracrine regulation of tumor growth. Reduced fibroblasts mitogenic activity could also counteract the increased number of CAFs in the reactive stroma. This idea is supported by a very recent study reporting fewer stromal CAFs in pancreatic cancer patients after treatment with nanoparticles containing paclitaxel [95]. Wnt signaling deregulation in human SCCs has been evidenced not only in cancer cells but also in stromal fibroblasts [96], suggesting that Wnt pathway might act in a paracrine fashion to promote skin carcinogenesis. Fibroblasts treated with Onco-P20 reduced the level of b-catenin expression and increased the production of soluble factors capable to diffuse similar Wnt pathway modulation to neighboring cells. 

In line with the idea that when the cell cycle is blocked downstream, pro-inflammatory pathway is over-activated [97], the most prominent feature of Onco-P20-growth arrested fibroblasts is a consistent pro-inflammatory secretory profile. The pro-inflammatory gene signature has been evidenced in early hyperplastic lesions as well as in end-stage squamous cell carcinomas [29]. Immuno-modulatory cytokines secreted by fibroblasts have a double edged sword effect in cancer biology. Some cytokines overexpressed in Onco-P20 treated fibroblasts influence the mobilization of local immune system playing a relevant role in defense against cancer [98, 99]. For example, CCL11 implicated in eosinophils recruitment and tissue necrosis [100]. IL1, IL6 and IL8 have a bimodal and dynamic behavior. Fibroblast in healthy tissue secrete basal level of IL6 and IL8 that act to maintain tissue homeostasis [101]. IL6 participates in the recruitment and polarization of macrophages, natural killer cells and T cells, promoting immune control of cancer cells [4, 102]. By contrast, high level of IL6 and IL8 favor macrophage immunosuppressive phenotype [103]. Augmented expression of IL6 in the peritumoural skin seems to be related to chronic UV-exposure rather than to the presence of neoplastic cells, since IL6 is almost absent within BCC [20]. IL-1α in combination with EGFR inhibitor can induce a T cell-dependent anti-tumor immune response in head and neck squamous cell carcinoma [104]. Acute inflammation, mainly extensive production of IL1b by CAFs significantly contributes to the efficacy of photodynamic therapy in cutaneous SCC [105]. Thus, inflammation may not always be 'adverse' in the context of cancer and the balance of individual members of the cytokine network can also contribute to therapeutic response.

CAFs influence another major stromal component within tumors by directing tumor-associated macrophages (TAMs) polarization (M1, type I anti-tumor or M2, type II immunosuppressive phenotype) [106]. Even if this study does not evidence any predictive specific effect on macrophages polarization, an interesting patient-based study demonstrated that the antitumor effect of paclitaxel may occur in part via reactivation of the immune response against cancer, guiding tumor-associated macrophages toward the M1-like anti-tumor phenotype [107]. In line with our study, the effect on TAM indicates that paclitaxel-based anti-tumor therapy does not target only cancer cells but also heterotypic interactions. Moreover, in line with our data strong IL1b occurs also in Onco-P20 treated macrophages. The advantage is that targeting tumor cells and normal stromal cells with the same molecule could offer a more effective and durable response since normal cells appear to have a relatively stable genetic constitution in contrast to the genetically instable genomes of cancer cells.

**Conclusion**

Overall our experiments clearly demonstrated that CAF-cancer cells interaction, at least those mediated by secreted factors, could be re-direct by Onco-P20 onto a tumor suppressive function. The enhanced Onco-P20 up-take by fibroblasts compared to standard free paclitaxel also encourage the use of lower therapeutic active doses limiting the toxic effect of uninvolved cells.

**Abbreviations**

BBC: basal cell carcinoma

CAFs: Cancer associated fibroblasts

ECM: Extracellular matrix

EGF: Epidermal growth factor

FAP-α: Fibroblast activated protein-α

FGF: Fibroblast growth factor

FSP1: Fibroblast-specific protein-1

HA: Hyaluronic acid

HKGS: Human Keratinocytes Growth Supplement
NHF: Normal human fibroblasts
NHK: Normal human keratinocytes
NMSC: Non-melanoma skin cancer
PDGF: Platelet-derived growth factor
PDGFR-α and β: Platelet-derived growth factor receptor-α and β
PML: promyelocytic leukemia protein
PTX: Paclitaxel
SCC: Squamous cell carcinoma
VEGF: Vascular endothelia growth factor,
Wnts: wingless-type MMTV integration site family members
α-SMA: α-smooth muscle actin

Declarations

Ethics approval: The Institutional Research Ethics Committee (Istituti Regina Elena e San Gallicano) approved all research activities involving human subjects. The Declaration of Helsinki Principles was followed and patients gave written informed consent to collect samples of human material for research.

Consent for publication: All authors have approved the manuscript and agree with its submission to Journal of Experimental & Clinical Cancer Research.

Availability of data and materials: The datasets used or analysed during the current study are available from the corresponding author on reasonable request.

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Author Contributors: BB designed the study, conducted the experiments, analyzed data, and wrote the manuscript; MP contributed to the study design and critically revised the paper. SC, GL and VM performed part of the experiments. EM selected patients and collected tissue sample. All authors discussed the results and approved the final version of the manuscript.

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Tables
Table 1. Gene expression analysis of growth factors and CAF’s markers in Onco-P20-treated fibroblasts after 2 weeks of continuous treatment. Data represent x-fold increase or decrease respect to untreated cells. Results are mean±SD of seven different experiments performed with fibroblasts isolated from independent donors. *p<0.05 and **p<0.001.

| mRNA      | Onco-P20 (x fold change) |
|-----------|---------------------------|
| HGF       | 0.44±0.36**               |
| IGF       | 0.72±0.97                 |
| IGF1      | 1.31±0.51                 |
| KGF       | 1.41±1.82                 |
| EGF       | 0.06±0.33                 |
| VEGF      | 0.69±0.59                 |
| IGF1P3    | 2.0±±2.05**               |
| IGF1P4    | 2.48±0.59**               |
| IGF1P5    | 2.14±1.9**                |
| IGF1P6    | 2.1±±0.54*                |
| IGF1P7    | 1.62±0.68*                |
| Vimentin  | 2.48±2.77**               |
| CDK1      | 4.21±±2.38**              |
| SFRP2     | 7.60±2.31*                |
| TGFβ      | 0.87±0.24                 |
| PDGFA     | 0.05±0.71                 |
| PDGFB     | 0.57±0.25**               |
| αSMA      | 1.14±0.66                 |
| FAP1      | 4.06±5.67**               |
| IL1α      | 18.8±15.9**               |
| IL1β      | 34.0±43.8**               |
| IL6       | 5.84±1.2**                |
| IL8       | 109.2±105.6**             |

Table 2. Results of human inflammation genes array after 2 weeks treatment with Onco-P20 (or not ctrl cells). Results are mean±SD of seven different experiments performed with fibroblasts isolated from independent donors. *p<0.05 and **p<0.001. Not detected (ND)
Table 3. Semi-quantitative analysis of multiple cytokines secreted by Onco-P20-treated fibroblasts after 2 weeks of continuous treatment. Data represent x-fold increase or decrease respect to untreated cells. Results are mean±SD of four different experiments performed with fibroblasts isolated from independent donors. *p<0.05 and **p<0.001.

| Target | Onco-P20 (x-fold change) |
|--------|--------------------------|
| CCL11  | 2.65±1.0*                |
| CCL24  | 1.16±0.87                |
| GCSF   | 0.57±0.24                |
| GROα   | 1.03±0.44                |
| IFNγ   | 1.49±0.61                |
| IL5a   | 1.55±0.09                |
| IL8    | 2.13±1.58                |
| IL2    | 1.35±0.80                |
| IL3    | 1.00±0.59                |
| IL4    | 3.97±3.12                |
| IL6    | 0.62±4.84*               |
| IL7    | 0.85±0.61                |
| IL18   | 4.00±1.64**              |
| IL50   | 1.22±0.61                |
| IL1    | 0.90±0.43                |
| IL2p83   | 0.86±0.71               |
| IL2pM  | 2.13±2.20                |
| IL10   | 0.73±0.57                |
| CCL1   | 0.26±0.31                |
| TNF-2  | 0.91±0.02                |

Figures
Figure 1

MTT assay (a) and cell count (b) of A431 cells treated (72h) with increasing concentration of Onco-P20 (range 0.05-1.0 μg/mL) or HA 1.0 μg/mL equivalent to the HA concentration at the Onco-P20 higher doses. Data represent the mean±SD of three independent experiments were performed in duplicate. MTT assay (c) and cell count (d) of NHKs treated (72h) with increasing concentration of Onco-P20. Histograms represent the mean±SD of three NHK cell lines isolated from different donors. (e) Microscopic images of normal and carcinoma cells after 72h treatment with Onco-P20. (f) Cell cycle distribution of Onco-P20-treated of normal and carcinoma cells. (g) Apoptosis was evaluated by FACS analysis using annexin V/iodide propidium staining. Dot plots show one representative experiment performed 24 h after Onco-P20 treatment. Statistical significance versus untreated control is reported as *p<0.05 and **p<0.001.
Figure 2

(a) MTT assay of several different NMSC cell lines and patient-matched NHKs after 72 h treatment with Onco-P20 (range 0.05-1.0 μg/mL). (b) Data are also reported as mean value ±SD. Experiments were performed in duplicate. (c) Annexin V/iodide propidium staining evidenced apoptosis cell death only in carcinoma cells. Dot plots show one representative experiment performed 24 h after Onco-P20 treatment.
Figure 3

(a) One representative immunostaining of plasma membrane CD44 expression. Data demonstrated lower intensity in fresh isolated carcinoma cells (SCC138-CI) compared to A431 carcinoma cell line. NHKs isolated from healthy skin of the same patient (NHK138-CI) presented the lowest quantity of CD44. Histogram plot reports fluorescent intensity. (b) Similar to (a) semi-quantitative analysis of CD44 staining showed high expression in all NHF cell culture analyzed compared to A431 and in fresh isolated carcinoma cells (SCC1300-UC).
Figure 4

(a) MTT assay and cell count (b) of several different NHF cell lines treated (72h) with increasing concentration of Onco-P20 (range 0.05-1.0 µg/mL) or HA 1.0 µg/mL equivalent to the HA concentration at the Onco-P20 higher doses. (c) FACS analysis using annexin V/iodide propidium staining excluded apoptosis. Dot plots show one representative experiment performed 24 h after Onco-P20 treatment. Statistical significance versus untreated control is reported as *p<0.05 and **p<0.001.
MTT assay of NHFs treated for 8 h with Onco-P20 or PTX before drug wash out (a) or treated continuously for 72 h with identical treatments (b). Data represent the mean±SD of three independent experiments were performed in duplicate. Statistical significance expressing Onco-P20 versus PTX-treated cells is reported as *p<0.05.
Figure 6

(a) One representative western blot analysis demonstrating the reduction of growth factors and β-catenin expression in Onco-P20-treated (0.15 μg/mL) NHFs. Protein were extracted after 2 weeks treatment. (b) Immunoenzymatic quantification of IGFBP4, IGFBP6 and VEGF released in cell culture medium. After 2 weeks treatment, cells were incubated for additional 48 h with DMEM without FBS before medium collection. Results were normalized against protein concentration.
Figure 7

(a) MTT assay of A431 carcinoma cells grown in presence of CM collected from Onco-P20-treated fibroblast. After 2 weeks and an additional 48 h period without treatment, medium was replaced with M154 without supplements before CM harvesting. CM was diluted 1:5 in fresh M154. CM of untreated fibroblasts was used as control medium. Histograms represent mean±SD of three independent experiments performed with three different NHF cell lines at the experimental end point (72h). (b) Schematic representation of trans-well systems used for co-culture experiments. MTT assay after 72 h of A431 cells in co-culture with fibroblasts demonstrated dose-dependent anti-cancer activity of Onco-P20-treated normal fibroblasts. (c) MTT assay after 72 h of SCC1300-UC carcinoma cells in co-culture with fibroblasts isolated from uninvolved, perilesional and lesional area of carcinoma patients. Results report data relative three independent experiments performed with three different donors. (d) MTT assay after 72 h of SCC316-BA carcinoma cells in co-culture with matched fibroblasts isolated from uninvolved, perilesional and lesional of the same donor (SCC316-BA). Statistical significance versus untreated control cells is reported as *p<0.05 and **p<0.001.

Supplementary Files

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