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Analysis of human renal angiomyolipoma cells: pharmacological modulation of growth and migration in vitro

Dott.ssa Francesca Bertolini

Tutor: Prof.ssa Silvia Anna Racca

Co-tutor: Dott.ssa Barbara Mognetti

PhD Coordinator: Prof. Giuseppe Saglio

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ABSTRACT

Introduction: Renal angiomyolipomas (AMLs) are highly vascular masses variously composed by differentiated cells such as smooth muscle, adipose, and vascular endothelial cells. Pulmonary lymphangioleiomyomatosis (LAM) is an age and gender disease, which hits at first young women, characterized by smooth muscle infiltration of the lung alveolar walls, leading to cystic degeneration. While both can occur clinically as isolated conditions, AML and LAM are also common manifestations of tuberous sclerosis complex (TSC), an autosomal dominant disorder characterized by hamartomas involving the central nervous system, skin, liver, heart, and eyes. Individuals diagnosed with TSC develop AMLs by adulthood in 70-90% of cases, whereas LAM can occur in up to 30-40% of affected women. TSC arises from inactivating mutations of either TSC1 (chromosome locus 9q34.3) or TSC2 (16p13.3) genes that encode for hamartin and tuberin, respectively. These proteins regulate cell proliferation and differentiation, and are involved in the mTOR (mammalian target of rapamycin) signaling pathway. Mutations in TSC2 often result in more severe clinical profiles, including higher frequency and severity of renal angiomyolipomas. No pharmacological treatment for AMLs is available and the most common approach is still the surgical ablation in the case of symptomatic disease or for lesions greater than 4 cm. Although the molecular events responsible for AML development are not fully understood, mTOR inhibitor therapy with sirolimus and everolimus may delay the growth and progression of AML in patients with TSC.

Trials with sirolimus suggest its efficacy in slowing the evolution of the pulmonary disease. In cases of severe functional respiratory or rapid evolution of the disease, the treatment of election is the pulmonary transplantation.

TSC-associated and sporadic AMLs and LAM are more common in female patients. Thus, some Authors have suggested a key role for the estrogen in their development. Nevertheless, it has not been fully clarified yet whether and how hormonal modulation is involved in the different occurrence in male and female patients. The smooth muscle cells of patients affected by both AML and LAM are characterized by structural cytologic analogy and the same somatic mutations; furthermore, AML cells have been detected in donor lungs of patients who had lung transplantation. These observations support the hypothesis that LAM might occur because of the migration of cells from other sites, such as the kidney, lymphatic system, or uterus. Therefore a common cell progenitor originating in the kidney and a metastatic behavior have been suggested. Despite several publications describing AML and LAM, a very few is known about their triggering agents and an eventual correlation between the two pathologies. To date, only a small number of hypotheses have been formulated regarding a link between AML and LAM might be of paramount importance in foreseeing and preventing their evolution, especially in fertile-age women. It must be considered that AML and LAM have, at now, a very dramatic evolution and no definitive cure has been established yet.

The overall objectives of this study are: 1) to ascertain the migratory proprieties of AML cells, 2) to identify common characteristics that might bind AML and LAM, and 3) to define innovative potential pharmacological approaches able to interfere with proliferation and migration of AML cells.

Material and Methods: The study has been performed on two different models: 1) ex vivo experiments on cells isolated from AML, and 2) in vitro experiments on three stabilized AML cell lines. Human primary AMLs cells were isolated from two spontaneous angiomyolipomas, surgically excised from a male (AML3) and a female (AML4) patient, which had neither clinical signs and symptoms nor a family history of tuberous sclerosis. On the other hand, two of the stabilized cell lines were TSC mutated: 4004 male cells carry a mutation in the exon 33 of TSC2 gene (4083-4087 del AGTCG) and 621-101 female cells have biallelic mutations in the exon 16 of TSC2 gene (G1832A). The third cell line (621-103) derives from the 621-101 cell line by restoration of the TSC2 gene.

We tested, in vitro, different concentrations and exposure times of drugs employed for the treatment of LAM and AML patients (sirolimus and everolimus) as well as of alternative selected drugs (simvastatin, zoledronic acid, estradiol, and tamoxifen) to evaluate whether they can modulate AML cells proliferation.

By means of 2D and 3D migration assays, we assessed if AML cells can migrate and if hormonal stimuli and drugs can modulate this property. Moreover, we evaluated the production of metalloproteases principally involved in malignant phenotype acquisition (MMP-2 and MMP-9) by zymography. Finally, by quantitative RT-PCR, we studied the expression of some genes crucial for the progression of these diseases.

Results: In the first part of the study, we isolated and grew cells from primary human spontaneous angiomyolipomas derived from male and female patients. To better understand which type of cells we isolated from fresh masses, we characterized the cellular composition of our cultures by means of immunofluorescence staining against HMB45, Melan-A, S-100, α -Smooth-Muscle Actin, keratin 8/18 and vimentin. The different cellular positivity to the antibodies confirmed the heterogeneous nature of both AMLs.

Subsequently, we examined the expression of estrogen receptor genes; the analysis demonstrated that both primary cell cultures express significantly level of mRNA for GPR30 and ER α , but not for ER β . Moreover, no significant differences in ER gene expressions have been detected between the male and female AMLs.

We evaluated if the treatments with 17- β -estradiol, tamoxifen or the combination of both modulated the proliferation and migration of cells. Concentrations comprised between 0.1 and 100 nM of 17- β -estradiol or between 0.2 and 20 μ M of tamoxifen did not modify cell proliferation. Through the analyses of 2D and 3D migration assays, we observed a different cell behavior. Treatment with estradiol (1 nM) for 2 hours induced a significant increase in 2D motility for both primary cell cultures; in 3D migration assay, only the female cells significantly responded to estrogen after 4 hours. Tamoxifen (2 μ M) had no influence on cell motility, but it was able to erase the effects of estradiol. Moreover, we performed a western blot analysis of Erk1/2 phosphorylation, demonstrating that estradiol incubation activates Erk1/2 pathways.

Finally, we evaluated if the migration of our cells was modulated by SDF-1 α , a chemokine associated with invasion and homing to specific organs. Both cells expressed, in basal condition, similar levels of mRNA for CXCR4, the receptor for SDF-1 α . The addition of SDF-1 α (100 ng/mL) to the culture medium stimulated the migration of both primary cells already after 4 hours of incubation; the stimulation was coherently abolished by the SDF-1 α -receptor antagonist plerixafor (100 nM). At last, we analyzed by zymography the activities of MMP2 and MMP9 on the supernatant collected after 3D migration assay; in both cells treatment with SDF-1 α significantly increased MMP-9 activity in both primary cultures.

Considering the limited number of primary AML cells available, we strengthened our results by studying the stabilized AML cell lines.

In these cells, we evaluated the expression level of specific genes involved in proliferation and migration in either basal condition or under specific drugs treatment. In basal condition, we observed significant expression level of antiapoptotic factors, enzymes involved in the metastatic process, and elements involved in dedifferentiation processes in all cell lines.

At the same concentrations and time tested for primary cells, the treatment with 17- β -estradiol did not induce any significant modification on cell proliferation. Unexpectedly, the proliferation of cell lines was inhibited by 10 μ M tamoxifen already after 24 hours. In three-dimensional migration assay, incubation with estradiol, tamoxifen or both did not modify the migration ratio for male 4004 cells. On the other hand, 1 nM 17- β -estradiol significantly increased the rate of female 621-101 and 621-103 cells migration; 2 μ M tamoxifen increased the migration of these cells also, even if less than compared with 17- β -estradiol (P<0.05). The different behavior of male and female cells was also confirmed, after 4 hours of incubation with the same concentration of 17- β -estradiol, by the increments in the expression level of several genes (such as MMPs, cMET, VEGF, FN1, and IGFR1) involved in motility and cellular adhesion.

Treatment with everolimus (1-100 nM), zoledronic acid, and simvastatin (1-100 μ M) had a direct toxic effect on AML cells, though at different concentrations and incubation time. Moreover, we

tested these drugs in association or alone in a 3D migration assay. For all cell lines, the combination of everolimus and zoledronic acid strongly inhibited the migration at low concentration. We also observed a decrease of migratory capacity induced by 10 μ M simvastatin, that inhibits cell migration already after 4 hours (P<0.05).

Conclusion: Our results suggest that significant differences between the analyzed AMLs are present for what concerns the response to the tested molecules. Both primary and stabilized cells have basic migratory properties. The migration of primary cells is likely modulated by agonists such as SDF-1 α and 17- β -estradiol, as confirmed by the quenching effect of their blockers. Primary cells increase their migration under estrogen influences independently from their gender. This confirms the hypothesis of a potential involvement in propagation and colonization to lung or other organs. In stabilized cells, the female ones are more responsive to estrogenic stimulation in terms of three-dimensional and genetic modulation than the male cells. Drugs such as zoledronic acid and everolimus (alone or in association), can significantly inhibit the growth and migration of the stabilized cells. The AML cells present, in basal condition, expression patterns attributable to invasive behavior, modulation of apoptosis, modification of the microenvironment, and angiogenesis. This feature supports the plasticity and abnormal cell proliferation typical of AML disease. Furthermore, the selected drugs analyzed do not induce a statistically significant modification of the level expression genes so far analyzed. In conclusion, these results represent a starting point for the development of valid pharmacological therapies for AML pathology.

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Appendix

1. Introduction

1.1 Renal angiomyolipoma

Renal angiomyolipoma (AML) is a mesenchymal highly vascular mass composed by three different elements present in variable proportion: smooth muscle, fat, and thick-walled blood vessels [1].

Angiomyolipoma, together with lymphangioleiomyomatosis (LAM) and other clear cell tumor of different sites, belongs to a family of neoplasms characterized by the presence of perivascular epithelioid cell (PEC) and therefore named PEComas [2].

The perivascular epithelioid cell have morphologic, immunohistochemical, ultrastructural and genetic distinctive features such as an epithelioid appearance [3]. Different reports indicate that these cells are immunoreactive for melanocytic, such as HMB-45 and melan-A, and smooth muscle, such as actin, markers [2,4,5].

PEComas of the kidney include different type of angiomyolipoma: classic, cystic, epithelioid, oncocytoma-like AML and lymphangiomyomatosis of the renal sinus [2].

In general AML is a benign mass and represents the 2-6.4% of the renal tumors [6,7].

Renal AMLs can occur sporadically (80-90% of cases) or as part of tuberous sclerosis (TSC), an autosomal dominant syndrome characterized by hamartomas; and less commonly, can occur in association with sporadic lymphangioleiomyomatosis (LAM) [8,9].

Two main types of renal AML are recognized: the sporadic type, usually a unilateral and solitary neoplasm typically seen in middle-aged women, and the multifocal neoplasms, usually small and bilateral, diagnosed in patients with TSC [1]. The sporadic AMLs are asymptomatic and are incidentally discovered during radiographic procedures; in contrast, TSC-associated AML lesions are often symptomatic at presentation and have frequent involvement of both kidneys [10].

TSC-associated and sporadic AMLs are more common and occur at an earlier age than in male in female patients than male (ratio of 2:1) [11].

The most common approach for remove AMLs at present is still the surgical ablation in the case of symptomatic disease or for lesions greater than 4 cm. Although the molecular events responsible for

AML development are not fully understood, mTOR inhibitor therapy with sirolimus and everolimus may delay the growth and progression of AML in patients with TSC. Everolimus has been approved by the Food and Drug Administration (FDA) for the treatment of TSC-associated angiomyolipoma in adult patients [12,13].

The heterogeneity of renal angiomyolipoma cells make it difficult their isolation, identification and phenotype characterization [14]. In the past, several studies have been conducted to analyze AML and LAM behavior an in vivo murine model (Eker rat with TSC2 germline mutation occurring in one of its alleles) or in vitro on ELT3 cell line derived from uterine leiomyomas [15,16,17]. To better understand the nature and characteristics of AML cells different authors tried to isolate cells from human angiomyolipoma masses to create in vitro and in vivo models to study human AML pathogenesis.

In 2004 (Yu et al.) and in the 2005 (Lesma et al.) two human cell lines were successfully isolated from tissue obtained from patients with TSC associated angiomyolipoma [18,19]. These cells were positive for melanocytic and smooth muscle markers and the smooth muscle cells revealed TSC2 LOH and a lack of tuberin expression.

In 2009, Clements and colleagues isolated two cell line populations derived from angiomyolipoma: spindle-shaped cells and epithelioid-like cells. The cultured cells showed constitutive activation of S6K1 protein and strong expression of α SMA, mRNA of gp100, and MART-1 [20].

These in vitro researches highlights the fact that the phenotype of AML cells depends on their microenvironment, exposure to cytokines, growth factors and interactions with other cells Moreover, these studies allowed uncovering new molecules involved in processes of proliferation, vascularization, angiogenesis and differentiation such as VEGF, EGF, IGF-1, TGFB1 and TGFB1 that can become a new therapeutic targets for the AML cells [14].

3

1.2 The "Benign Metastasis" hypothesis

Although the histological characteristics of AML cells are benign, a hypothesis was formulated in which smooth muscle cells with mutations in TSC1 or TSC2 gene could travel to the lungs from renal angiomyolipomas [21,22].

Identical somatic mutations of TSC2 have been described in abnormal lung and kidney cells, but not in healthy cells, in the 60% of women with sporadic LAM and renal AML [23].

Spread of angiomyolipoma smooth muscle cells to the lungs could explain the occasional recurrences of LAM in the donor lung after lung transplantation [23].

This hypothesis is supported by the finding that TSC2-deficient smooth muscle cells have higher migration potential than normal cells in vitro, and different patients with tuberous-sclerosis-associated pulmonary lymphangioleiomyomatosis have large, potentially metastatogenic renal angiomyolipomas [22,24]. The genetic analyses reveal that the metastatic mechanism was related to the presence of mutations in TSC1 or TSC2 gene [25].

The molecular mechanisms underlying the presumed metastatic properties of AML have not still clarified, but the identification, reported in several studies in the literature [26,27,28], of molecules involved in migration, invasion, proliferation and angiogenesis in the cells and tissues of LAM and AML patients supports this unusual disease mechanism.

A recent in vivo and in vitro study discovered that PPARy (peroxisome proliferator-activated receptor gamma) plays a central role in the initiation and propagation of sporadic and TSC-related AML cells [28]. The microarray gene expression analysis revealed strong activation of PPARy in an AML xenograft model, generated by an injection of 10⁶ UMB cells (cells derived by TSC-related AML) into NOD/SCID mice and in primary human AML cells. Consequently, Pleniceanu et al., show that the inhibition of this molecule, induced by one of its antagonists (GW9662), significantly and specifically halts the in vitro growth of cells and strongly limited their tumor-initiation capacity in the in vivo model.

Another in vivo study, conducted on a murine model of TSC, analyzed the mTOR and YAP (Hippo– Yes-associated protein 1) pathways that is involved in tumorigenesis. YAP is up-regulated by mTOR in mouse and human PEComas and it inhibition blunts abnormal proliferation and induce apoptosis of TSC1-2 deficient cells both in in vivo (PEComas of mosaic Tsc1 mutant mice) and in vitro models (HEK293 and MEFs, mouse embryonic fibroblasts, cell lines) [29].

Fibronectin is involved in cell adhesion, cell motility, host defense and metastasis [26]. Evans and colleagues suggested that the aberrant proliferation of LAM cells might be associated with overexpression of fibronectin [30], a glycoprotein present in a soluble and dimeric form in the plasma, and a dimeric or multimeric form on the cell surface and in extracellular matrix (ECM). Altered fibronectin levels have been demonstrated in various diseases such as interstitial lung diseases as well as LAM. Concordant with these findings, they proposed that the deposition of fibronectin might result in tissue remodeling and perhaps fibronectin might be involved in LAM pathogenesis [30]. Other factors that are involved in the metastatic and tumor mechanisms are the matrix metalloproteinases (MMPs).

Physiologically matrix metalloproteinases are a group of enzymes able to degrade components of the ECM; they are secreted as inactive proenzymes which become active when cleaved [31]. MMPs play a major role in cell proliferation, migration (adhesion/dispersion), differentiation, angiogenesis, apoptosis, and host defense [32]. The gelatinases, MMP-2 and MMP-9 can degrade native type IV collagen, denatured type I collagen (gelatin) and elastic fibers [33].

Immunohistochemical studies have found that MMP-2 and MMP-9 are predominantly expressed, in comparison to other MMPs, in LAM cells, suggesting their involvement in the destructive cystic formation and in the pathogenesis of LAM [33,34,35].

MMP-2 is a zync-dependent enzyme capable of remodeling the vasculature, involved in angiogenesis, tissue repair, tumor invasion and inflammation. MMP-2 is predominantly expressed by mesenchymal cells, smooth muscle cells and fibroblast [33]. An increased level of MMP-2 was found in AML cells derived from a LAM patient and in serum of patients with pulmonary LAM [31,32,35].

MMP-9 gene encodes for the enzyme matrix metalloproteinase 9; it is significantly up-regulated in many human diseases (arthritis, diabetes and cancer), and this aberrant activity is thought to contribute to pathological processes such as tissue remodeling, invasion and cell migration [36]. The levels of metalloproteinase 9 in circulating blood are elevated in various pathological states such as asthma, pulmonary emphysema, lung cancer and ischemic heart disease [33].

Cell metastasis to specific organs may depend on cytokines and from the interaction between soluble factors produced by the microenvironment with metastatic cell receptors. In this regard, it has been hypothesized that ligands and receptors of the chemokines can participate in the migration of AML cells to other organs [37]. Stromal cell-derived factor-1 α (SDF-1 α or CXCL12) and its unique receptor CXC chemokine receptor-4 (CXCR4) have prominent roles in invasion and metastasis of a diverse number of cancers [37,38,39,40].

The interaction between SDF-1 α and CXCR4 has been shown to guide tumor cells to organ sites with high levels of SDF-1 α expression, which suggests a key role in chemotaxis and homing of metastatic cells [27]. Therefore, the CXCL12-CXCR4 axis is essential for the regulation of cell migration [37].

1.3 Lymphangioleiomyomatosis

Another disease associated with TSC is lymphangioleyomiomatosis (LAM), which can, nevertheless, occur also in a sporadic form [41].

LAM is an age and gender related disease primarily affecting young women and which, in the TSC form, develops with different degrees of severity and disability [42]. It is characterized, histologically, by smooth muscle infiltration of the lung alveolar walls, leading to cystic degeneration.

Lung function abnormalities consist of decreased expiratory flow expressed as a decreased lung diffusion capacity, leading to a reduction in breathing capacity and hypoxemia during exercise or at rest [43]. LAM presents with dyspnea, recurrent pneumothoraxes, pleural effusions, ascites, and extrapulmonary features that include bleeding renal angiomyolipomas and lymphangioleiomyomas

[8].

In some cases, lung disease progresses slowly; in others, typically in younger women, LAM tends to run a more rapid evolution. In both cases, it occurs a decline in lung function leading to respiratory failure. In cases of severe functional respiratory or rapid evolution of the disease, the treatment of election is the pulmonary transplantation [44].

The median transplant free survival time is of approximately 29 years from the onset of symptoms. There are no efficacious drugs for the treatment of LAM; based on current evidence, it is recommended that LAM patients in whom lung function is declining rapidly or having symptomatic lymphangioleiomyomas, pleural effusions or ascites are treated with sirolimus or everolimus [45, 46]. Trials with sirolimus suggest its efficacy in slowing the evolution of the pulmonary disease [45].

1.4 Tuberous sclerosis

TSC is a rare autosomal dominant multisystem disorder that can cause circumscribed, benign, noninvasive lesions in multiple organs including brain, skin, liver, lung, kidney, heart and eyes [47]. The birth incidence of TSC is estimated to be approximately 1 in 6,000 to 1 in 11,000 [48,49,50]. Renal lesions, collectively occurring in 50–80% of patients with TSC, include angiomyolipomas, renal cysts, renal cell carcinoma, and oncocytomas [50].

TSC arises from inactivating heterozygous or mosaic mutations in either TSC1 (~21%) or TSC2 (~79%) with subsequent deregulation of the Rheb/mTOR/p70S6K pathway. The TSC1 gene is located on chromosome 9q34, and consists of 21 exons encoding the 1164 amino acid protein hamartin. The TSC2 gene on chromosome 16p13 contains 41 exons and encodes the 1807 amino acid protein tuberin [14]. Hamartin and tuberin are likely tumor suppressors regulating cellular proliferation [51,52].

There are different types of gene mutations in TSC1 and TSC2 genes, with an equal distribution in family cases while in the sporadic cases mutation occur more frequently in the TSC2 gene [51]. Mutations in TSC1 are often small insertion or deletions that result in truncated protein and patients

with these mutations have generally a mild clinical phenotype [51]; whereas TSC2 mutations include large deletions, nonsense, and missense mutations [53]. These mutations in these genes are believed to prevent the formation of the complex between TSC1 and TSC2, culminating in loss of inhibition of the mTOR pathway [53]. Mutations in TSC2 gene often result in more severe clinical profiles, including higher frequency and severity of renal angiomyolipomas.

The different nature of hamartomas observed in patients with TSC suggests that these cancers develop according to "second hit" theory (Knudson, 1971). In hereditary cases, a first genetic alteration ("first hit") is inherited in the germ line by a parent affection, while the second mutation occurs in a cell that already has the first mutation. The second mutation can cause the complete inactivation of either of the two genes of TSC1 or TSC2 and, of consequence, loss of heterozygosity (LOH) [22].

LOH in the TSC1 or TSC2 region occurs in most angiomyolipomas, rhabdomyomas, and astrocytomas from TSC patients. TSC2 LOH in renal lesions occurs in 10% of sporadic angiomyolipomas and in 60% of TSC-associated angiomyolipomas [25, 54].

1.5 mTOR pathway

The TSC1/TSC2 complex is the principal cellular regulator of mTOR (mammalian target of rapamycin), a serine/threonine protein kinase in the PI3K-related kinase family that controls different important biological processes such as cellular growth, proliferation, transduction, transcription, autophagy and metabolism [55]. A few of the metabolically active tissues in which mTOR plays an important role include the heart, pancreas [56] and kidney [57].

mTOR is a component of two functional complexes: TORC1 and TORC2; TORC1 is the rapamycinsensitive mTOR complex responsible for the regulation of protein translation initiation and efficiency, and is activated by the presence of growth factors, amino acids and energy status [58]. TORC2 responds to growth factors and is not directly inhibited by rapamycin and induces cell cycle exit and differentiation [58]. The role of the proteins encoded by TSC1 and TSC2 (hamartin and tuberin respectively) in this pathway is to form a heterodimer that activates a GTPase enzyme called Rho, which by reducing Rheb activity, indirectly inhibits mTOR. The production of tuberin and hamartin is also physiologically regulated by different factors such as PI3K (phosphoinositide 3-kinase) and PDK1 (phosphoinositide-dependent kinase-1), which induce the phosphorylation of Akt. Activated Akt, as well as extracellular signal-regulated kinase (ERK), phosphorylates TSC2 resulting in inhibition of its activity as a GTP-ase [14].

In tuberous sclerosis, mutations in one of the TSC genes result in the abnormal formation of the hamartin/tuberin complex and so the subsequent inhibition of Rheb is lost and this induces an increase of cell proliferation [14].

Therefore, deficiency of hamartin or tuberin due to mutations of TSC1 or TSC2 in patients with TSC gives rise to hyperactivation of the mTOR pathway, resulting in a downstream kinase signaling cascade that can consequently lead to abnormalities in numerous cell processes, including cell cycle progression, transcription, translation, and metabolic control [46].

1.5.1 The PI3K/Akt and MAPK pathways

mTOR can be activated either through the PI3K/Akt or MAPK (mitogen-activated protein kinase) ERK1/2 axes [59]. The PI3K/Akt and MAPK pathways play important roles in modulating cellular function in response to extracellular signals, including growth factors and hormones. These pathways are activated in certain kinds of malignancies for example bone and soft tissue tumors and certain sarcomas, and in renal angiomyolipoma [60,61,62].

Akt is a serine/threonine kinase activated by phosphoinositide 3-kinase (PI3K), regulates many cellular processes (proliferation, survival and cell growth) and when deregulated can contribute to the progression of cancer [63].

Akt has also been shown to contribute to tumor invasion and metastasis by promoting the secretion of MMP-9 [64] and the induction of epithelial–mesenchymal transition [65]. The higher secretion of MMP-9, associated with an overexpression of AKT, facilitates the movement and thus invasion of tumor cells [63,66]. The PI3K/AKT/mTOR pathway is regulated by a wide-range of upstream signaling proteins and it regulates many downstream effectors by collaborating with various compensatory signaling pathways, primarily with RAF/MEK/ERK pathway [67].

The Ras/Raf/MEK/extracellular signal regulated kinase (ERK) pathway, also known as the MAPK pathway, also regulates a variety of cell functions such as proliferation, growth, and survival. The two forms of ERK, ERK1 and ERK 2, belong to the family of MAP kinases, are ubiquitously expressed and are involved in the regulation of meiosis, mitosis, post-mitotic functions in differentiated cells and in cell proliferation and survival [68,69].

The activation of the ERK1/2 cascade mainly depends on the membrane receptors, such as G protein– coupled receptors (GPCRs), ion channels, and others; ERK1/2 were demonstrated to regulate several members in the subgroup of nuclear receptors, including estrogen receptor and in different type of cells, the rapid activation of ERK1/2 pathway can be induced by estradiol [69].

1.6 mTOR inhibitor: Everolimus

Since the TSC gene products form a tumor suppressor complex that inhibits mTORC1 activity, this latter complex can be targeted for therapy in patients with TSC renal, lung and subependymal giant cell astrocytoma disease. Sirolimus and everolimus are administrated as mTOR inhibitors; sirolimus is a macrolide compound that is used to prevent organ transplant rejection and to treat a rare lung disease (LAM). Everolimus is an orally bioavailable, structurally similar derivative of sirolimus that exhibits antiprofilerative and immunosuppressive effects.

Everolimus was designed to improve the pharmacokinetics of the sirolimus; in fact, it has greater stability, solubility, rapid and consistent absorption and lower nephrotoxicity, and reaches its peak concentration after 1.3–1.8 hours [70].

Everolimus forms a complex with the intracellular binding protein FKBP12 which directly interacts with mTORC1; this complex which block the constitutively upregulated downstream PI3K/Akt/mTOR signaling by reducing the phosphorylation of downstream mTOR effectors and the downstream signaling events responsible for the many manifestations of TSC [13,71].

At now the pharmacological treatment of AML and LAM associated with TSC is limited to mTOR inhibitors. Everolimus is administrated to reduce the tumor progression but, unfortunately, not all patients respond adequately; moreover, it has very important side effects such as ulcers, fatigue, rash, mucositis, anorexia, diarrhea, nausea, arthralgia, thrombocytopenia, and effects on lipid metabolism [72]. The recent literature suggests that everolimus is efficacious and safe in controlling AML tumor burden in patients with TSC, while preserving the renal parenchyma. However, tumor responses after treatment with only everolimus are usually only partial, and regrowth occurs after drug withdrawal [73]. For example, Yang and colleagues tested the antitumor efficacy of everolimus in combination with sorafenib, a kinase inhibitor, on renal lesions in patients with TSC. They suggested that this combination might improve therapeutic efficacy, compared to everolimus alone, for TSC-associated solid tumors [73].

Questions regarding the durability of responses, period of treatment and impact of toxicity from chronic therapy remain, and the role of mTOR inhibitors in the treatment of patients with non-TSC associated AMLs is still to be determined [9, 74].

1.7 The role of estrogen in AML

The higher frequency of LAM and AML in women, suggests the hypothesis that the growth and migration of TSC2-deficient cells can be influence by estrogens [24].

The majority of female patients with these diseases are at child-bearing age at the time of diagnosis; pregnancy and use of oral contraception, in LAM, are associated with a higher frequency of exacerbations and a more aggressive disease course [75]. Boorjian and colleges observed a positivity of both estrogen receptors on smooth muscle cells of sporadic and TSC-associated renal AML.

Moreover, the carboxy terminal of TSC2 interacts with the estrogen receptor alpha functioning as transcriptional corepressor of estrogen receptor. Sex steroids might play a role in the pathogenesis of renal AMLs, hormone receptors could be potential therapeutic targets [76].

17-β-estradiol (estrogen or E2) is a typically gender hormone; through the estrogen receptors (ER) it acts a variety of biological processes including reproduction, differentiation, cell proliferation [77]. This hormone exerts its activity in different tissues through three major receptors: the soluble nuclear receptors, ER α and ER β , and GPR30.

The two nuclear receptors, ER α and ER β , are highly homologous in their DNA- and ligand-binding domains [78]. Although ER α and ER β are coexpressed in certain target tissues, they also exhibit different tissue/cell expression patterns and are functionally distinct. ER α is a more potent transcriptional activator than ER β , and in tissues where both ERs are expressed, ER β has been suggested to have a role as an attenuator of ER α [79].

GPR30 is a seven-transmembrane spanning G protein-coupled receptor [80] that binds 17-β-estradiol with high affinity and lead to rapid and transient activation of numerous intracellular signaling pathways, for example calcium mobilization, cAMP production, PI3K activation and ERK1/2 activation in a G-protein dependent manner [81].

These receptors can mediate gene transcription events, either directly (genomic), as in the case of ER α and ER β [Edwards D.P. et al., 2005; Marino M. et al., 2006], or indirectly (non-genomic), as in the case of GPR30 in which the MAPK, PI3K, Scr kinase and related pathways are activated [78,83,84].

The most known antagonist of the 17-β-estradiol is tamoxifen (TAM); TAM is a non-steroidal antiestrogen, extensively used in the treatment of hormone responsive breast cancer [85]. The principal mechanism of action of TAM is the inhibition of estrogen receptor, inducing a conformational change in the receptor with a consecutive modification in the expression of estrogen dependent genes [86]. Tamoxifen acts as an anti-estrogen (inhibiting agent) in the mammary tissue, but as an estrogen (stimulating agent) in cholesterol metabolism, bone density, and cell proliferation in the endometrium [85]. Tamoxifen is a partial agonist of ER α , a selective agonist for GPR30 and a pure antagonist for ER β [86]. The different effects, agonism or antagonism, mediated by TAM in various tissues may depend on the different expression of alpha and beta estrogen receptors and on the type of receptor-ligand interaction [86,87].

Recent experimental studies have revealed, other than estrogen receptors, new tamoxifen targets, such as protein kinase C, phospholipase C and antiangiogenic agents, which are key mediators of signaling pathways activating additional non-ER-mechanisms [88].

Tamoxifen has been used, besides the breast and endometrial carcinomas [89], in the treatment of malignancies such as renal carcinoma [86]. In this ER-negative cancer, the therapeutic efficacy of TAM has been obtained at doses at 4 to 8 folds above those used in ER-positive tumors [86]. The anti-tumor effect of tamoxifen is believed to be due to a combination of ER and non-ER-mediated mechanisms [86].

Since sex steroids might play a role in the pathogenesis of renal AMLs, a possible strategy for the treatment of AML and LAM might correspond to the inhibition of the effect of the estrogen [68].

1.8 Other approaches in AML treatment

Despite publication of several observational papers [1,90], little is known about efficient pharmacological treatments on AML mass. Aside from mTOR inhibitors, no other drugs are known for the therapy of this disease; therefore, an effort has to be done to identify molecules able to slow down its development. Since very few recent in vitro studies on AML cells are published in the literature, we based our literature researches on other cell models with a similar origin and behavior of AML.

Several in vitro experiments conducted on renal cancer cells and other cancer cell models suggest that drugs such as simvastatin and zoledronic acid (ZOL) might be used also for their anti-proliferative effect [91,92,93].

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1.8.1 Zoledronic acid

Bisphosphonates (BPs) are simple chemical compounds and are similar to endogenous pyrophosphates [94].

Some in vitro cell culture experiments have shown that BPs induce apoptosis and inhibit cancer cell invasion and angiogenesis in several human tumor cell lines such as breast, prostate, lung and kidney; [92,94]. Data from the literature report that caspase-dependent apoptosis appears to be the major mechanism responsible for BP-induced tumor cell apoptosis, and caspase-3 is certainly the major player in the antitumor activity [92,95].

BPs exert anticancer activity by stimulating the expansion of $\gamma\sigma$ T-cells (a subset of human T-cells that have antitumor activity) [96] by mimicking phosphoantigens and/or rising circulating phosphoantigen levels, and increasing sensitivity of cancer cells to the cytotoxic effects of $\gamma\sigma$ T-cells [97].

Zoledronic acid (ZA) belongs to the third generation of BPs; it inhibits the activity of farnesyl pyrophosphate synthase (FPPS) in the mevalonate pathway. The inhibition of FPPS reduces the activity of some small GTPases, such as Rho and Ras, which are among the key signal pathways that promote cell proliferation, vesicular traffic, cell adhesion and induction of phosphorylation of mTOR [98].

The possible mechanism of action through which zoledronic acid induces a decrease in tumor growth includes the inhibition of angiogenetic factors such as VEGF, and modification of the tumor microenvironment (i.e. reduction of vascularization and macrophage infiltration) [99,100].

ZA inhibits the activity of some metalloproteinases (such as MMP-2 and MMP-9) essential for cell migration and adhesion [101,102], and modulates the expression of adhesion molecules (such as cadherin and integrin) which regulate the cellular invasion [103]. Therefore, in vitro studies on prostate and breast carcinoma cells hypothesized the use of zoledronic acid as a growth and cell migration inhibitor.

1.7.2 Simvastatin

Statins are inhibitors of the first committed enzyme of mevalonate pathway, 3-hydroxy-methylglutary (HMG) CoA reductase. In general, statins decrease LDL levels, and are effective in preventing cardiovascular diseases [104].

At now, several in vitro experiments have demonstrated that liphophilic statins, such as simvastatin and atorvastatin, exert antiproliferative effects on different cells such as renal cancer [91] and prostate cancer [105]. Antiproliferative effects of statin include growth inhibition caused by cell cycle arrest, induction of apoptosis, and reduction of metastatic potential. These effects are regulated by Akt/mTOR axis through a Rho dependent mechanism [104,106,107].

Fang and colleagues found that the treatment of human renal carcinoma cell lines with simvastatin, significantly suppresses the phosphorylation/activation of Akt and inhibits the phosphorylation of mTOR. This suggests that simvastatin may exert its anti-cancer effects by inhibiting the Akt/mTOR axis [91].

Moreover, Atochina-Vasserman et al. compared the effect of simvastatin and atorvastatin on in vitro and in vivo mouse model of LAM, and showed that simvastatin, through AKT/PKB axis, inhibits the growth of these cells better than atorvastatin. Additionally, simvastatin, but not atorvastatin, also induces concentration-dependent inhibition of phosphorylation of ERK1/2 and mTORC1 [108].

2. Aim of the study

The overall objective of this study is the improvement of the knowledge of AML cells biology and, in particular, the ascertainment of their migratory properties. Despite being classified as a benign tumor, hypothesis, never confirmed, have been put forward concerning their metastatic potential. Our study will try to clarify if and how AML cells can migrate to lung or other organs, and if their behavior is influenced by hormonal milieu.

Moreover, our study may clarify the high diversity between the incidence of AML in female and male patients and the possible link between LAM and AML diseases.

In conclusion, we want to identify molecules capable to modify AML cell proliferation, migration which might be an outstanding information for considering new treatment protocols in AML patients, in order to replace as much as possible surgical removal of AML.

3. Material and methods

3.1 Angiomyolipoma cells, tissue and ethical approvals

Human primary AML cells have been obtained from patients that underwent surgical nephron-sparing AML ablation for therapeutic purposes at the Urology Unit of the San Luigi Gonzaga Hospital. The study was approved by the Ethical Committee of the San Luigi Gonzaga Hospital, (Protocol 0006771, approved on April 18, 2016). All patients provided written informed consent in accordance with the Declaration of Helsinki.

AML3 derives from a male patient, AML4 from a female patient. None of the patients had any clinical signs or symptoms or a family history of tuberous sclerosis.

Primary cells were isolated from excess material not required for diagnostic use, which was divided into small fragments and treated with type II collagenase. Resulting cell suspensions were plated into T25 tissue culture flasks in AML medium (adapted from Lesma et al.), composed of phenol red DMEM medium, ferrous sulphate 1.6 μ M, 20% fetal bovine serum (FBS) and 10 μ g/mL epidermal growth factor.

3.2 Stabilized cell culture

Human 621-101 cells derived from renal angiomyolipoma of LAM female patients, kindly provided by Prof. Henske, (Harvard Medical School, USA) and human 4004 cells derived from renal angiomyolipoma cells derived from TSC male patients were purchased from ATCC (Rockville, MD, USA). 621-103 cell lines, also kindly provided by Prof. Henske, derive from the 621-101 cell line by restoration of the TSC2 gene.

621-101 cells carry biallelic mutations in exon 16 of the TSC2 gene (G1832A) [18], 4004 cells carry mutations in exon 33 of TSC2 gene (4083-4087 del AGTCG) [109].

Cells were maintained at 37°C in a humidified 5% CO₂ atmosphere in phenol red DMEM medium containing 1% penicillin and streptomycin solution and 10% Fetal Bovine Serum (FBS).

3.3 Immunofluorescence

Primary cells were grown on round sterilized coverslips under standard conditions for 2 days, and then washed with PBS followed by fixation in 4% paraformaldehyde (PAF) for 15 minutes. After washing in PBS, cells were treated with PBS containing 1% normal goat serum (NGS), 0.1% Triton X-100 at room temperature (RT) for 1 hour. After incubation time cells were incubated overnight at 4°C with the following primary antibodies (diluted in PBS) against: S-100 (rabbit, 1: 800; Dako, Glostrup, Denmark), α-Smooth-Muscle Actin (mouse, 1: 100; NeoMarkers, Fremont, CA), HMB45 (mouse,1:100, Dako, Glostrup, Denmark), Keratin 8/18 (mouse, 1:100 Menarini, Florence, Italy) Vimentin (mouse, 1:70; Novocastra Lab, Newcastle, UK) and Melan-a (mouse; 1:100; NeoMarkers, Fremont, CA). After washing, cells were incubated for 1 hour at RT with the appropriate secondary antibodies: goat anti-mouse IgG Alexa-Fluor-488-conjugated (1: 200, Molecular Probes, Eugene, Oregon) and CY3-conjugated anti-rabbit IgG (dilution 1:400, Dako, Milan, Italy). Finally, washed slides 3 times in PBS, added DAPI solution (dilution 1:1000) and after the last washing mounted with antifade mounting media [110]. The immunostained coverslips were analyzed on a Zeiss fluorescence microscope and images were captured with an Axiovision Imaging System.

3.4 Proliferation assay

Angiomiolipoma cells, primary or stabilized, were seeded into flat-bottomed 96-well microplates $(1,000/100 \ \mu L \ culture \ medium/well)$ and allowed to attach overnight in complete medium before drugs addition.

Drugs were added to culture medium testing various concentrations from 24 to 72 hours:

- From 0.1 to 100 nM and 17-β-estradiol [18] and from 0.2 to 20 μM tamoxifen [18] on primary and stabilized cells.
- From 0.001 to 100 μM zoledronic acid [111] and everolimus [111], and from 1 to 100 μM simvastatin [91] for stabilized cells.

The MTT ((3-(4,5-Dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide assay was performed according to routine protocols. Briefly, 10 μ l of MTT prepared at a concentration of 5 mg/ml in PBS, was added to each well. Cell culture was placed in the incubator for another 3 hours at 37°C; after discarding supernatant, 100 μ l of DMSO were then added to each well and the absorbance was measured using a microculture plate reader (Microplate 450, Bio-Rad, Hercules, CA, USA) with a wavelength of 595 nm. Data (mean \pm standard errors) were the average values of 8 replicates. Each experiment was repeated thrice. Cell viability was expressed as percentage of living cells with respect to controls.

3.5 Two-dimensional migration assay-wound healing

Primary AML cells were seeded in a 12-well plate at 300,000 cells/well. When they were confluent, a cross "wound" was made in each well with a p200 tip, then wells were washed thrice with PBS, and cultured in AML medium supplemented either with SDF-1 α (Peprotech, London, UK) 13 nM, plerixafor 100 nM, 17- β -estradiol 1 nM, or tamoxifen 2 μ M. We photographed "wounds" on time-laps every hour to highlight migration, until a maximum of 8 hours. Experiments were repeated three times, and every time five different spots for each experimental condition were considered. Images were analysed using ImageJ software (Wayne Rasband, NIH, USA): the healing percentage was quantified comparing the wound area at t = 0 to the following time-points for each treatment.

3.6 Three-dimensional migration assay

To measure the three-dimensional (3D) movements of primary and stabilized cells in response to different drugs, migration assays were performed trough transwell (BD Falcon cell culture inserts incorporating polyethylene terephthalate membrane with 8.0 μ M pores, $6\pm2x10^4$ pores /cm²). DMEM was placed on the bottom of a 24-well microplates, and the transwell was inserted within the same: the multiwell plate was placed in the incubator for 30 minutes for medium equilibration. In the meanwhile the angiomyolipoma cells were suspended in 200 μ L; cell suspension was carefully placed

in the transwell and multiwell was placed in incubator. At the end of migration time, each transwell was taken off from its well and gently washed twice with PBS Ca²⁺/Mg²⁺ to prevent the detachment of cells from the membrane of the transwell and then with use of cotton swabs the cells located in the internal part of the transwell taken off. The transwell was immersed in a solution of glutaraldehyde 2% for 20 minutes to fix cells and then washed five times in deionized water and allowed to dry. Each transwell was immersed in crystal violet 0.1% for 20 minutes to stain cells and then washed again five times in deionized water and allowed to dry. Each transwell was indeionized water and allowed to dry. Wells were photographed using a BRESSER MikroCam 3 Mpx camera, with an optical microscope (Leica DC 100) at 100x. Five pictures were randomly chosen per well, and used to count the migrated cells with ImageJ software using cell-counter plug-in. Results from different experiments (performed at least three times in duplicate) were expressed as mean ± standard error.

The experimental scheme for the different performed tests was schematically described in the following tables (Table 1-3).

The concentration of drugs used in transwell assay represent non-toxic concentration at 24 hours as per the described viability assay.

Experimental time	Upper chamber (200 µL)	Lower chamber (800 µL)
1 and 8 hours	Drimory colls (5×10^4)	Fresh culture medium (CTRL)
4 and 8 nours	Finnary cens (5x10)	17-β-estradiol (1 nM)
		Fresh culture medium (CTRL)
4 1	Stabilized cells (8x10 ⁴)	17-β-estradiol (1 nM)
4 nours		tamoxifen (2 μM)
	Stabilized cells $(8x10^4)$ + tamoxifen $(2 \mu M)$	17-β-estradiol (1 nM)

Table 1: Three-dimensional migration assay with estrogen incubation

Experimental time	Upper chamber (200 µL)	Lower chamber (800 µL)
	Stabilized cells (8x10 ⁴)	Fresh culture medium (CTRL)
	Stabilized cells $(8x10^4)$ + simvastatin $(10 \mu M)$	
4 hours	Stabilized cells $(8x10^4)$ + zoledronic acid $(10 \mu M)$	
	Stabilized cells $(8x10^4)$ + everolimus (10 nM)	Fresh culture medium
	Stabilized cells $(8x10^4)$ + zoledronic acid $(10 \mu M)$	
	+ everolimus (10 nM)	

Table 2: Three-dimensional migration assay with drug incubation

Experimental time	Upper chamber (200 µL)	Lower chamber (800 µL)
	Primary cells (5x10 ⁴)	Fresh culture medium (CTRL)
4		SDF-1α (13 nM)
4 and 8 nours		plerixafor (100 nM)
	Primary cells $(5x10^4)$ + plerixafor* (100 nM)	SDF-1 α (13 nM)

Table 3: Three-dimensional migration assay in presence of SDF-1 α and plerixafor. * For this test, primary cells were pre-incubated for 30 min at 37°C with plerixafor.

3.7 Western blot

Cells were seeded in 10 cm-diameter Petri dishes, cultured until sub-confluence and treated with drugs (17- β -estradiol 1nM, tamoxifen 2 μ M, simvastatin 10 μ M, zoledronic acid 10 μ M and everolimus 10nM). Following incubation period (5 min -1 hours for estradiol and 4 hours for others treatment), the medium was removed and the cell monolayer was first washed with PBS, then covered with ice-cold PBS and incubated for 5 minutes to facilitate detachment. Subsequently, adherent cells were gently scraped, collected and centrifuged at 1000 rpm for 5 minutes. Then the pellet was resuspended in 50 μ l of RIPA buffer, placed in ice for one hour and gently shuffled every 20 minutes to facilitate the membrane breakup. The mixture was then centrifuged at 13200 rpm for 30 minutes at 4°C, the supernatant collected and protein content quantified with the Bradford assay (1 μ l RIPA suspension/999 μ l Bradford solution 1:5) (Bio-Rad): sample were read with a spectrophotometer (Beckman DU® 640 Spectrophotometer – U.S.A.) at 595 nm wavelength.

Thirty µg of proteins were resolved in the Biorad system by SDS-PAGE gels, denaturing conditions, and then transferred onto nitrocellulose membranes (GE Healthcare), and immunoblotted with the following antibodies (Table 4) according to Racca et al 2012 [112].

Primary antibody	Protein molecular weight	Diluition	Antibody manufacturer
Anti-vinculin (mouse)	116 kDa	1:2000	
Anti-estrogen receptor α (rabbit)	67 kDa	1:600	Sigma Aldrich
Anti-estrogen receptor β (rabbit)	59 kDa	1:600	Sigilia Alurich
Anti-GPR30 (rabbit)	42 kDa	1:600	
Anti-p42/44 Erk 1/2 (mouse)	42/44 kDa	1:2000	
Anti-phospho p42/44 Erk 1/2 (mouse)	42/44 kDa	1:2000	Cell Signaling
Anti-Akt (rabbit)	60 kDa	1:1000	Technology
Anti-phospho Akt (rabbt)	60 kDa	1:1000	

Table 4: Antibodies used for performing western blot.

The anti-vinculin, anti-estrogen alpha and beta were diluted in PBS Tween 0.1%; the other antibodies were suspended in 5% w/v nonfat dry milk + PBS tween 0.1%.

HRP-conjugated anti-mouse and anti-rabbit (Amersham-GE Healthcare, Buckinghamshire, UK) were diluted 1:6000 in PBS Tween 0.1%. Bands were quantified using the ImageJ software.

Phosphorylation levels of $ERK_{1/2}$ and Akt were expressed as ratio $pERK_{1/2}/ERK_{1/2}$ and pAkt/Akt respectively. Each experiment was repeated thrice and all data were expressed as percentage modification relative to control conditions.

3.8 Gelatin zymography

MMP-2 and MMP-9 activities in medium samples were assayed by gel zymography. Proteins (100 µg) were separated by electrophoresis in 8% SDS-PAGE gel containing gelatin (0.8 mg/mL) under non-reducing conditions. The gel was washed with Tris buffer (2.5% Triton X-100 in 50 mM Tris-HCl, pH 7.5, final solution) for 1 hour, then incubated overnight at 37°C in a proteolysis buffer (40 mM Tris-HCl, 200 mM NaCl, 10 mM CaCl₂, 0.02% NaN₃, pH 7.5, final solution). The gel was stained for 3 hours with Coomassie Blue solution (0.05% Coomassie Brilliant Blue R-250, 50% methanol, 10% acetic acid, final solution) and finally destained with 5% methanol and 7% acetic acid (final solution). Reagents and chemicals were obtained from VWR International (Milan, Italy). MMPs activity was detected as a clear band on a blue background and estimated by densitometric analysis using ImageJ Software. The results were expressed as percentages of control values.

3.9 Quantitative real-time PCR (qRT-PCR)

In order to perform quantitative real-time PCR (qRT-PCR), total RNA was extracted from treated cells by using Trizol (Invitrogen Life Technologies Italy). After its purification and treatment with DNAse I (Fermentas, St. Leon-Rot, Germany), 1 µg was retrotranscribed in cDNA with the RevertAidTM H Minus First Strand cDNA Synthesis Kit (Fermentas) using oligo (dT) primers. Gene assays were performed in triplicate for each treatment in a 20 µL reaction volume containing 1 µL of RT products, 10 µL Sso-Fast EVA Green SMX (Bio-Rad), 500 nM each forward and reverse primers. Gene expression was normalized on the housekeeping gene ribosomal 18S rRNA. Table 5 resumes the primer sequences that were adopted. Automated CFX96 real-time thermocycler (Bio-Rad) was used and the reaction conditions were 95°C for 1 minute, followed by 45 cycles 98°C for 5 seconds and anneal–extend step for 5 seconds at 60°C, with data collection. At the end of these cycles, a melting curve (65°C to 95°C, with plate read every 0.5°C) was performed in order to assess the specificity of the amplification product by single peak melting temperature verification. Results were analysed with Bio-Rad CFX Manager. Calculations and statistical analyses were performed using GraphPad Prism version 5.00 for Windows (GraphPad Software, San Diego California USA).

GENE	SEQUENCE	AMPL. SIZE	NCBI REF. SEQ.	
EDa	Fw: 5'-TGGAGTCTGGTCCTGTGAGG-3'	172 hr	NT 025741 16	
EKű	Rev: 5'-CCCACCTTTCATCATTCCCACT-3'	172 op	N1_025/41.10	
Erß	Fw: 5'-GAGCAAAGATGAGCTTGCCG-3'	142 hn	NM_001437.2	
пр	Rev: 5'-AGCTGGGCCAAGAAGATTCC-3'	142 Up		
CDD 20	Fw: 5'-AGTCGGATGTGAGGTTCAG-3'	240 hp	NM_001505.2	
UFK50	Rev: 5'-TCTGTGTGAGGAGTGCAAG-3'	240 Op		
CYCPA	Fw: 5'-TGACGGACAAGTACAGGCTGC-3'	406 bp	NM_001348056.1	
CACR4	Rev: 5'-CCAGAAGGGAAGCGTGATGA-3'			
	Fw: 5'-GGCCCTGTCACTCCTGAGAT-3'	474 bp	NIM 001202510 1	
IVIIVIT-2	Rev: 5'-GGCATCCAGGTTATCGGGGA-3'		NWI_001302310.1	
MMD 0	Fw: 5'-CAACATCACCTATTGGATCC-3'	480 bp	NM 004004 2	
IVIIVII - 9	Rev: 5'-CGGGTGTAGAGTCTCTCGCT-3'		INIM_004994.2	
EN1	Fw: 5'-AGACCCCAGGCACCTATCAC-3'	263 bp	NM 017002605 1	
1/1/1	Rev: 5'-TCGGTCACTTCCACAAACTG-3'	Ĩ	NWI_017003095.1	
SDE 1a	Fw; 5'-GGTGGAGCTGGAGAAGACAGA-3'	72 hn	NIM 000600 6	
SDF-IU	Rev: 5'-CAGCCGGGCTACAATCTGAA-3'	73 Up	11/11/1_000009.0	
18S	Fw: 5'-GTGGAGCGATTTGTCTGGTT-3'	201 hr	V02205 1	
rRNA	Rev: 5'-ACGCTGAGCCAGTCAGTGTA-3'	201 bp	A03203.1	

Table 5: Primers sequences, size of the amplification product and NCBI Reference Sequence. Fw=forward, Rev= reverse

3.10 Statistical analysis

All the data in this study were shown as mean \pm standard error (SE). Two group means were compared using the unpaired t-test, and more than two group means were analyzed by one-way analysis of variance (ANOVA), where P <0.05 was considered statistically significant. For gene expression level comparison One-way ANOVA with Dunnett's post tests were performed using GraphPad Prism version 5.00 for Windows (GraphPad Software, San Diego California USA).

Coefficient of Drug Interaction (CDI) was used to define the type of interactions between the employed drugs. CDI was calculated by means of the equation: $CDI = AB/(A \times B)$, where AB is the relative cell migration of the combination; A or B, relative cell migration of the single agent. CDI < 1 indicates a synergistic effect; CDI = 1 indicates an additive effect; CDI > 1 indicates an antagonistic effect.

4. Results
Part I: Primary cells

4.1 Immunohistochemistry to confirm the AML diagnoses

AML diagnoses were confirmed by standard histological examination including specific immunostaining for alpha-smooth muscle Actin, HMB-45 and Pancytokeratin antigens conduct by the Patholgy Unit of San Luigi Gonzaga Hospital.

4.2 Cells characterization by immunofluorence

To better characterize isolated AML primary cells, immunofluorescence was performed with several antibodies. Overall, both cell lines showed similar immunophenotype and partly elongated, partly rounded shapes. Both primary culture cells were strongly and totally positive for smooth muscle actin antibody (in both elongated and rounded cells), with a diffused stain throughout the cytoplasm (Figure 1a and la'). Furthermore, in both cell lines there were single rounded element positive for keratin 8/18 and elongated cells strongly positive for vimentin (Figure 1d, 1e, 1d', and 1e') together with single negative ones for both the antigens. Finally, a strong nuclear and cytoplasmic positivity was found for S100 in both cells (Figure 1f and 1f'). Overall this phenotype confirmed the mixed (either muscle, epithelioid, lipomatous and mesenchimal) nature of the cultures, according to the heterogeneous nature of AML. As a matter of fact, even if scattered, some AML3 cells were focally positive for Melan-A antigen (Figure 1c and 1c'), both of which being consistent with the AML phenotype.



Figure 1: Primary angiomyolipoma cells characterization by immunofluorescence.

Cells isolated from the two AML were stained with different antibodies to detect their immunophenotype. Fields were chosen to clearly show both the morphological aspect and the specific marker expression.

4.3 Estrogen and SDF-1a receptor gene expression

Early passages AML cells underwent qRT-PCR for CXCR4, GPR30, ER α and ER β mRNA expression analysis, which demonstrated a significant presence of mRNA for CXCR4 (SDF-1 α receptor), GPR30, ER α , but not for ER β (Figure 2). The level of each gene was similar in the two cultures, with no significant difference.



Figure 2: Receptors gene expression.

Early passages AML cells underwent qRT-PCR for CXCR4, GPR30, ER α and ER β mRNA expression analysis. Data are shown as the absolute mRNA expression normalized by the housekeeping 18S rRNA.

4.4 Influence of estrogen on cell proliferation

Treatment of primary cells for 24-72 hours with 17- β -estradiol from 0.1 nM to 100 nM (Figure 3A) or with tamoxifen from 0.2 to 20 μ M (Figure 3B) did not modify AML cells proliferation, nor did the combination of 17- β -estradiol (1 nM) and tamoxifen (2 μ M) (Figure 4).



Figure 3: Proliferation assay

Effect of increasing concentration of 17- β -estradiol (A) or tamoxifen (B) after 24, 48 and 72 hours culture.



Figure 4: Effect of 17-\beta-estradiol alone and of its combination with tamoxifen on AML cells growth *Proliferation assay after 72 hours culture in presence of 17-\beta-estradiol (1 nM), tamoxifen (2 \muM) or the combination of both.*

4.5 Influence of estrogen on two- dimensional migration (wound healing)

The in vitro wound healing assay was performed on male and female cells to observe whether estrogen modifies the motility of AML cells. Figure 5A shows a scratch wound generated after cell confluence; figure 5B shows the same wounded area after 8 hours of assay. Wound healing was quantified, and data are displayed graphically as healing percentage (Figure 5 D-E). Panel C compares basal migration of AML3 and AML4: the early migration rate of AML3 was higher, a significant difference being demonstrated at 4 hours. This difference was promptly quenched since after 8 hours the movement rate of the two AMLs was similar. Two-dimensional migration was significantly modulated by 17- β -estradiol (1 nM), although with some differences between the two AMLs. In fact, AML3 (D) of male origin, showed a significant motility increase in the first 4 hours of incubation with 17- β -estradiol, and a prevalent logarithmic pattern. On the other hand, migration of AML4 cells, of female origin (E), significantly increased, following an exponential pattern, at any of the time points considered. The treatments with the ER-antagonist tamoxifen alone (2 μ M) had no influence on two-dimensional motility of any cell type, while it was able to abolish the effects of estradiol.



Figure 5: Analyses of primary AML cells migration by *in vitro* wound healing assay.

Scratch wound healing assay in a representative experiment before (A) and after (B) the incubation period. Panel C shows bi-dimensional migration of the two AMLs in basal conditions. Migration in presence of 17- β estradiol (1 nM) or its antagonist tamoxifen (2 μ M), or a combination of both, is shown in panel D (AML3) and E (AML4). Migration is expressed in arbitrary units. Data are expressed as the percentage of migration vs t = 0 h. *=P<0.05 vs control; #=P<0.05 vs 17- β -estradiol.

4.6 Influence of estrogen on three-dimensional migration

The transwell migration experiments performed in presence of 17- β -estradiol demonstrated a different behavior between cells originating from the two AMLs. Migration of AML3 cells of male origin was not modified all along the stimulation period (Figure 6A). AML4 cells (Figure 6B) responded to the stimulation by increasing the migration rate already at 4 hours while after 8 hours the stimulating effect was lost.



Figure 6. Three-dimensional migration.

Cells were incubated with 17- β -estradiol (1 nM) for 4 and 8 hours. Migration is expressed in arbitrary units. *=P < 0.05 vs control.

4.7 ERK phosphorylation

Treatment of AML3 cells with 17-β-estradiol increased ERK phosphorylation at 4 hours, while pERK was significantly augmented in AML4 cells already at 5 minutes and was stable until 4 hours (Figure 7).



Figure 7: ERK phosphorylation

Effect of 17- β -estradiol (1 nM) on pERK/ERK in AML3 and AML4 cells after 5 minutes and 4 hours of incubation. Vinculin as internal control. *=P<0.05 vs control.

4.8 Influence of SDF-1α on two and three-dimensional migration

SDF-1 α treatment, in wound healing assay, induced statistically significant motility increase already after 2 hours treatment in cells from both AMLs (Figure 8 A-B) and the difference persisted all along the experimental period (up until 8 hours). The results, for the transwell assay, show that the number of migrating cells significantly increased in response to SDF-1 α both after 4 and 8 hours incubation (Figure 8 C-D). In both assays, the effects induced by SDF-1 α were abolished by the SDF-1 α -receptor antagonist plerixafor, while no significant effects were induced by plerixafor alone.



Figure 8: Two and three-dimensional migration (SDF-1α)

Bi-dimensional migration in presence of SDF-1 α 13 nM or its receptor antagonist plerixafor 100 nM, or a combination of both (A-B). Three-dimensional migration with SDF-1 α (13 nM) and its receptor blocker plerixafor (100 nM) or a combination of both (C-D). *=P<0.05 vs control; #=P<0.05 vs SDF-1 α .

4.9 Metalloproteases activity in supernatant derived from 3D-migration test

The activity of two metalloproteases involved in malignant phenotype acquisition, MMP-2 and MMP-9, was measured by zymography in the supernatant collected at the end of the 3D-migration test performed in presence of SDF-1 α , plerixafor and 17- β -estradiol (4 hours) (Figure 9). While the diverse treatments induced no significant difference in MMP-2 activity, SDF-1 α enhanced MMP-9 enzymatic activity. Coherently, plerixafor inhibited MMP-9 activity increase provoked by SDF-1 α . No difference was induced by incubation with 17- β -estradiol.



Figure 9: Metalloproteases activity

MMP-2 and MMP-9 activity in presence of SDF-1 α , *plerixafor and 17-* β *-estradiol. Data are expressed as the relative activity calculated by densitometric analyses.* *=P<0.05 vs control; #=P<0.05 vs SDF-1 α .

Part II: Stabilized cell lines

4.10 Estrogen receptor expression

The qRT-PCR analysis for estrogen gene receptor demonstrated a significant presence of mRNA for GPR30, ER α and ER β for 4004 and 621-103; the level of GPR30 expression appeared to be lower in 621-101 cells than the other cells (Figure 10A). Instead, by western blot we detected the protein expression for all the estrogen receptor considered. In this case, high levels of expression for ER β were observed in 4004 and 621-103 cells (Figure 10B). We observed a significant expression of ER α only in female 621-101 cell lines by immunofluorescence, The receptor was localized in the cell cytoplasm (Figure 10C). Unfortunately, due to technical problems, the expression for ER β and GPR30 were not available by immunofluorescence.



Figure 10: Receptors gene expression

(A) 4004, 621-101 and 61-103 cells underwent qRT-PCR for ER α , ER β and GPR30 mRNA expression analysis. Data are shown as the absolute mRNA expression normalized by the housekeeping 18S rRNA. (B) Expression of ER α , ER β and GPR30 protein underwent western blot. (C) Expression of ER α in cell lines by immunofluorence. Figures show representative fields of 4004, 621-101 and 621-103 cells immune-stained with anti-ER α antibody.

4.11 Influence of 17-β-estradiol, tamoxifen and combination of both on cell proliferation

Treatment for up to 72 hours with increasing concentration of 17-β-estradiol (0.1-100 nM) did not modify 4004, 621-101 and 621-103 cell proliferation *in vitro* (Figure 11A).

Unexpectedly, the growth of all angiomyolipoma cell lines was strongly inhibited by 10 μ M tamoxifen already after 24 hours of treatment (Figure 11B).



Figure 11: Influence of 17-β-estradiol and tamoxifen on cell proliferation

Effect of 17- β -estradiol (A) or tamoxifen (B) on 4004, 621-101 and 621-103 cell growth. *=P<0.05 vs control.

The treatment of all cell lines with 17- β -estradiol (10 μ M) did not modify the cell proliferation. Treatment for 24 and 48 hours demonstrated that only in female 621-101 cells the combination of tamoxifen (10 μ M) and 17- β -estradiol (10 μ M) provoked an unusual increase of cell proliferation (Figure 12 B). In the male 4004 and female 621-103 cells the combination of these drugs did not modify the proliferation (Figure 12 A, C).



Figure 12: Influence of 17- β -estradiol and tamoxifen and combination of both on cell proliferation *Effect of 17-* β -estradiol (10 μ M) and tamoxifen (10 μ M) on 4004 (A), 621-101 (B) and 621-103 (C) cells. *=P<0.05 vs control; #=P<0.05 vs 17- β -estradiol; •= P<0.05 vs tamoxifen.

4.12 Influence of 17-β-estradiol and tamoxifen on three-dimensional migration

The migration of all cell lines was evaluated after 4 hours of incubation time in basal conditions, or in response to 17- β -estradiol (1 nM), tamoxifen (2 μ M) and the combination of both. None of the tested molecules modified the migration of male 4004 cells. On the other hand, 17- β -estradiol (1 nM) significantly increased the rate of female 621-101 and 621-103 cell migration (50% and 37%, respectively). The treatment with tamoxifen (2 μ M) and its combination with 17- β -estradiol also increased the number of migrating female cells (P<0.05) (Figure 13).



Figure 13: Three-dimensional migration assay

Cells were incubated with 17- β -estradiol (1 nM), tamoxifen (2 μ M) or a combination of both for 4 hours. Data are expressed as the percentage of migration vs control. *=P<0.05 vs control; #=P<0.05 vs 17- β -estradiol.

4.13 ERK and AKT phosphorylation

No significant modification in phosphorylation rate of ERK was induced on 4004 and 621-103 cells by 17- β -estradiol or tamoxifen incubation. Furthermore, for the female 621-101 cells the treatment with estradiol and tamoxifen for 4 hours induced a lower increase in pERK (Figure 14).

Neither of the two tested drugs induced a significant change in pAKT/AKT ratio in all cell lines (Figure 15).



Figure 14: ERK phosphorylation

Effect of 17-8-estradiol (1 nM) and tamoxifen (2 μ M) on pERK/ERK after 4 hours of incubation on all cell lines. *=P<0.05 vs control.



Figure 15: AKT phosphorylation

Effect of 17-6-estradiol (1 nM) and tamoxifen (2 μ M) on pAKT/AKT after 4 hours of incubation on all cell lines.

4.14 Effect of 17-β-estradiol and tamoxifen on adhesion/invasiveness gene expression

Cell lines were treated with $17-\beta$ -estradiol (1 nM) and tamoxifen (2 μ M) for 4 hours.

In 4004 cells (Figure 16) treatment with estradiol provokes a low decrease of the expression of FN1, GPER30 and MMP-2 genes; with tamoxifen treatment the diminution of the expression of these genes were more evident for MMP-2 especially. Both the treatments induced an increase, without evident differences, of the expression of SDF-1 α .



Figure 16: Effects of 17-β-estradiol (1 nM) and tamoxifen (2 μM) on gene expression.

Data from qRT-PCR are expressed as fold variation of gene expression in 4004 cells. Data are expressed as mean \pm SD of at least three experiments. *=P<0.05 vs control.

In 621-101 cells, the treatment with estradiol induce a diminution in the MMP-2 gene expression, whereas tamoxifen treatment provoked a significant reduction of FN-1 gene expression. For the other analyzed genes, the treatment did not induce noteworthy variation in their expression (Figure 17).



Figure 17: Effects of 17-β-estradiol (1 nM) and tamoxifen (2 μM) on gene expression.

Data from qRT-PCR are expressed as fold variation of gene expression in 621-101 cells. Data are expressed as mean \pm SD of at least three experiments. *=P<0.05 vs control.

Finally, the treatment with estradiol on 621-103 female cells (Figure 18) induced an increase in the expression of GPR30 gene, and after treatment with tamoxifen, the expression of MMP-2 gene was strongly decreased as observed in 4004 male cells.





Data from qRT-PCR are expressed as fold variation of gene expression in 621-103 cells. Data are expressed as mean \pm SD of at least three experiments. *=P<0.05 vs control.

4.15 Influence of simvastatin, zoledronic acid and everolimus on cell proliferation

To evaluate the effect of different drugs on the proliferation of AML cells; 4004, 621-101 and 621-103 cells were exposed to different concentrations of simvastatin, zoledronic acid and everolimus for 24, 48 and 72 hours. Simvastatin showed its inhibitory effect on cell growth at 10 μ M after 48 hours incubation (Figure 19).

The 621-101 cells were more sensitive to simvastatin at 72 hours respect to 4004 and 621-103 cells; the inhibition growth of 621-101 cells, at 1 μ M, was 10% more than the other cells.



Figure 19: Influence of simvastatin on cell proliferation

Effect of increasing concentration of simvastatin on 4004, 621-101 and 621-103 cell growth. *=P<0.05 vs control.

Zoledronic acid (0.001-100 μ M) strongly inhibited cell proliferation; 4004 cells growth was inhibited by concentrations as low as 1 μ M after 48 hours and even more strikingly after 72 hours culture (Figure 20A). Growth inhibition on 621-103 cells was less striking at 48 hours but comparable at 72 hours (Figure 20B, C).



Figure 20: Influence of zoledronic acid on cell proliferation

Effect of increasing concentration of zoledronic acid on 4004 (A), 621-101 (B) and 621-103 (C) cell growth after 24, 48 and 72 hours of incubation. *=P<0.05 vs control.

Everolimus inhibited cell proliferation at 1 μ M for 4004, 10 μ M for 621-101 and 100 μ M for 621-103 after 24 hours of treatment (Figure 21A, B, C). After 48 hours of treatment 4004 and 621-101 cell growth were inhibited after already at 0.001 μ M.



Figure 21: Influence of everolimus on cell proliferation

Effect of increasing concentration of everolimus on 4004 (A), 621-101 (B) and 621-103 (C) cell growth. *=P < 0.05 vs control.

To evaluate if everolimus and zoledronic acid have a synergic effect we also tested the combination of these two drugs on all cell lines. We observed that the simultaneous presence of these drugs influenced proliferation less than zoledronic acid alone. At 48 hours the combination of these drugs, caused a 40% (in 4004 and 621-101 cells) or a 20% (in 621-103 cells) growth inhibition, a

significantly greater effect on cell growth compared with the treatment with everolimus alone (Figure 22). The table 6 reports the coefficient of drug interaction (CDI) after 24 hours of treatment.





hours of incubation.

Table 6: Drug interactions after 24 hours of treatment

	АВ	Α	В	CDI
4004	zoledronic acid + everolimus (24 hours)	zoledronic acid (10µM)	everolimus (1 nM)	0,98
			everolimu (10 nM)	0,86
621-101	zoledronic acid + everolimus (24 hours)	zoledronic acid (10µM)	everolimus (1 nM)	0,87
			everolimu (10 nM)	0,81
621-103	zoledronic acid + everolimus (24 hours)	zoledronic acid (10µM)	everolimus (1 nM)	0,95
			everolimu (10 nM)	0,91

CDI < 1 indicates a synergistic effect; CD=1 indicates an additive effect

4.16 Three dimensional migration assay in presence of zoledronic acid, everolimus and

simvastatin

Treatment with zoledronic acid, everolimus and the combination of both significantly decreased the migration rate of all cell lines if compared to the control conditions (Figure 23). In this test, the simultaneous presence of zoledronic acid and everolimus influenced migration less than each single drug.



Figure 23: Effect of zoledronic acid, averolimus and combination of both on three-dimensional migration

Effect of 4 hours treatment with zoledronic acid (10 μ M), everolimus (10 nM) or a combination of both on 3D migration. Data are expressed as the percentage of migration vs control. *=P<0.05 vs control; #=P<0.05 vs zoledronic acid; •=P<0.05 vs everolimus.

Simvastatin decreased migration of all cell lines. Likewise the proliferation test, we observed in 3D migration assay a significant difference between cells: 621-101 cells were much more inhibited than 4004 and 621-103 (50% vs 34 and 30%, respectively) (Figure 24).



Figure 24: Effect of simvastatin on three-dimensional migration

Effect of 4 hours treatment with simvastatin (10 μ M). Data are expressed as percentage of migration vs control. *=P<0.05 vs control.

4.17 ERK and AKT phosphorylation

Treatment with zoledronic acid (10 μ M) and everolimus (10 nM) for 4 hours did not modify significantly the pERK/ERK ratio for any cell line. pERK/ERK ratio was similarly decreased by simvastatin (10 μ M) in 4004 and 621-101 cells; in 621-103 cells the treatment did not change the ratio of ERK (Figure 25).



Figure 25: ERK phosphorylation

Effect of zoledronic acid (10 μ M), everolimus (10 nM) and simvastatin (2 μ M) on pERK/ERK after 4 hours of incubation on all cell lines. *=P<0.05 vs control.

Exposure of 4004 and 621-103 cells to everolimus and simvastatin for 4 hours significantly decreased pAKT/AKT, while zoledronic acid after 4 hours had no effect compared to the untreated control. On the other hand, in 621-101 cells the treatments with zoledronic acid and everolimus induced a significant inhibition of pAKT/AKT, in this case the treatment with simvastatin did not significantly change this ratio (Figure 26).



Figure 26: AKT phosphorylation

Effect of zoledronic acid (10 μ M), everolimus (10 nM) and simvastatin (2 μ M) on pAKT/AKT after 4 hours of incubation on all cell lines. *=P<0.05 vs control

4.18 Effect of zoledronic acid, everolimus and simvastatin on adhesion/invasiveness gene expression

The treatment with zoledronic acid, simvastatin and everolimus on 4004 cells induced change in the level gene expression. Zoledronic acid induced only a reduction in MMP-2 level gene, simvastatin induced a significant increase in FN-1 and SDF-1 α expression and a significant decrease in GPR30 and MMP-2 gene expression (Figure 27).



Figure 27: Effects of zoledronic acid (10 μ M), everolimus (10 nM) and simvastatin (10 μ M) on gene expression.

Data from qRT-PCR are expressed as fold variation of gene expression in 4004 cells. Data are expressed as mean \pm SD of at least three experiments. *=P<0.05 vs control.

Treatment of 621-101 cells with zoledronic acid and simvastatin induced a significant increase in the expression level of MMP-2 gene, while the treatment with everolimus induced a decrease in the expression of FN-1 and SDF-1 α genes.

The treatments did not induce any noteworthy variation in the expression of the other genes analyzed (Figure 28).





Figure 28: Effects of zoledronic acid (10 μ M), everolimus (10 nM) and simvastatin (10 μ M) on gene expression.

Data from qRT-PCR are expressed as fold variation of gene expression in 621-101 cells. Data are expressed as mean \pm SD of at least three experiments. *=P<0.05 vs control.

Finally, in 621-103 cells zoledronic acid and simvastatin induced a strong increase of SDF-1 α expression, furthermore only the zoledronic acid provoked a reduction in MMP-2 gene expression. Everolimus did not induce any significant modification in the expression of the genes analyzed (Figure 29).



Figure 29: Effects of zoledronic acid (10 μ M), everolimus (10 nM) and simvastatin (10 μ M) on gene expression.

Data from qRT-PCR are expressed as fold variation of gene expression in 621-103 cells. Data are expressed as mean \pm SD of at least three experiments. *=P<0.05 vs control.

5. Discussion

Renal angiomyolipoma is a rare disease that can occur sporadically or in association with TSC.

Despite different observational studies, at now AML cell biology is still poorly defined and the underlying mechanisms of this disease are not yet known.

The heterogeneous nature of this benign mass adds more complexity in understanding the behavior and the possible therapeutic approaches.

In the first part of this study, for the first time, we describe the behavior of primary AML cells, originating from male and female patients, in terms of proliferation and migration in vitro, both in basal condition and in response to environmental stimuli.

Our findings clearly demonstrated that primary AML cells were able to migrate in vitro. We isolated a great number of cells by male (AML3) and female (AML4) angiomyolipoma, without a clinical history of TSC, on which we made different in vitro experiments.

First, we characterized the isolated cells: the two cultures were similar to each other regarding a mixed cell composition, according to the heterogeneous nature of AML. None of the cell types (smooth muscle, epithelioid, lipomatous and mesenchimal) in the cultures seemed to be predominant and this characteristic is an advantage in a pharmacology study, since the complexity of the tumor should be taken into account.

Independently from their TSC mutational status, we decided to ascertain whether these cells were able to migrate and if this process could be modulated pharmacologically.

In particular, taking advantage from the fact that our cells originated from male and female patients, we focused our attention on estrogen to investigate the high difference between the incidences of AML in patients with opposite gender. Moreover, we evaluated if the hypothesis that renal AML cells might migrate to the lung or other organs is true.

Despite the presence of the estrogen receptors, we have observed that there was no estrogenic influence on proliferation of both AMLs.

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Although some common properties between the two AMLs have been described, we reported that other aspects seemed to differentiate the cells originating from male and female patients. We showed that the basal unstimulated migration of male AML3 had more rapid onset, respect to female AML4. This difference in spreading was already striking 4 hours after seeding, and also evident in the ability to invade, as demonstrated by three-dimensional assays. Together with this observation, we also reported that the higher basal migration of AML3 is less influenced by estrogen, while spreading of AML4 cells is significantly increased already after 2 hours incubation with 17- β -estradiol (1 nM). In order to explain the response to estrogen, we investigated the expression of estrogen receptor: ER α , ER β , and GPR30.

In our primary cells, we detected an abundant mRNA expression of GPR30, as described by Marino et al., respect to the nuclear receptor; this is coherent with the rapid non-genomic response to $17-\beta$ -estradiol that we observed. The existence of a rapid, non-genomic response has also been confirmed by the ability of estradiol to quickly increase ERK1/2 phosphorylation in female AML4 (after 5 minutes of treatment with 17- β -estradiol 1 nM) [113, 82].

Moreover, to verify whether the SDF-1 α is indispensable for migration also in our cells, we analyzed how they cells responded to it in this process. Our results are in accordance with the well-known role of SDF-1 α in cell migration; in fact, we have shown its significant effect in increasing the bi- and three-dimensional migration of these cells. The specificity of this observation was confirmed by the effect of the treatment with the selective SDF-1 α receptor antagonist plerixafor, which completely abolished SDF-1 α stimulation.

Analysis of the culture media collected from three-dimensional migration experiments revealed that the stimulation of migration induced by SDF-1 α was accompanied by an augmented release of MMPs, in particular of MMP-2. Also in this case, these effects were consistently reverted by the selective receptor antagonist plerixafor. Strikingly, such an increase in MMPs production was not observed in estradiol-stimulated conditions, despite a significant increase in cell migration. We therefore speculated that, while SDF-1 α -induced migration occurred through metalloproteases production, the migratory response to estrogen most likely did not involve MMPs activity. Our results therefore might support the hypothesis that the host microenvironment tissue *stimuli* could exert a specific chemotactic signal to promote homing of AML cells as already seen in other study [37]. The primary culture approach, has inherent strengths and unfortunately, some limitations.

The restrictions are clear: since AML is a rare disease, it is difficult to obtain fresh AML tissues, indeed we could analyze two samples only. Moreover, the use of primary cultures restricted the number of studies that we could perform before losing them. For these reasons, we decided to strengthen our results by studying three immortalized AML cell lines: 4004 from male and 621-101, 621-103 from female patients with TSC.

In the second part of the study, to improve the results obtained on the primary cells, the same experiments were repeated on stabilized cells; then, we evaluated the effects of some drugs, belonging to different therapeutic classes, on the proliferation and migration of AML cells and on the expression of some genes crucial for these mechanisms.

At first, we decided to evaluate the effect of estrogen on the proliferation and migration of male and female cells. Renal AML masses are known to express estrogen receptor [76]; the analysis by means of western blot allowed us to affirm that even in all our cells great protein level of these receptors were expressed. However, through a qRT-PCR we observed an abundant mRNA expression of transmembrane estrogen receptor (GPR30) in 621-103 and 4004 cells with no significant differences between them. The discrepancy between the absence of signal in qRT-PCR and protein expression that we observed in 621-101 cells is astonishing, and hence we supposed that the most realistic way to explain it is to ascribe to a technical problem occurred during cell processing.

Although the presence of estrogen receptor, we reported also for these stabilized cells, that treatment with an increasing concentration of 17- β -estradiol did not modify their proliferation.

The treatment of our cells with estrogen antagonist, tamoxifen, emphasized its dual action; already after 24 hours, low concentrations of tamoxifen provoked an increase of growth (statistically not significant), while over 10 μ M it induced a strong inhibition of proliferation in all cell lines. This

behavior confirmed the hypothesis of Prossnitz and colleagues according to tamoxifen promotes or inhibits cell proliferation based on the estrogen receptor it binds to [78].

Tamoxifen is able to bind ER α and ER β and to inhibit proliferative stimulation, but it can also bind the transmembrane estrogenic receptors on which it acts like an agonist stimulating the proliferation [78]. Increased cell growth at low tamoxifen concentrations may indicate a greater affinity of the drug to transmembrane receptors rather than to nuclear estrogenic receptors.

About migration, we observed that also these stabilized cells were able to migrate in vitro in basal conditions. After 4 hours, the female cells were more responsive to the estrogen *stimuli* respect to the male cells.

Tamoxifen induced an increase in migration only for the female cells; this rapid increase might be associated to the agonist effect of tamoxifen on GPR30 receptor [82]. Moreover, in female 621-101 cells, this rapid increase in migration, induced both by $17-\beta$ -estradiol and tamoxifen, can be associated with the phosphorylation of p42/44 MAPK that we observed through western blot analysis; an analogous situation has been described by Liu et al. in non-small cell lung cancer [114].

In 4004 and 621-103 cells the treatment with these molecules did not induce any modification in phosphorylation of ERK1/2; the ratio of pAkt/Akt was not modified in any cell line after these treatments. Therefore, we can suggest that the phosphorylation of Akt did not correlate with the migration under hormonal *stimuli*. Analogous results were previously observed by Clements and colleagues in a xenograft AML in which phosphorylation of Akt and p42/44 MAPK were not affected by oestrogen treatment [20].

The effect on the MMP-2 gene expression on human renal angiomyolipoma cell lines treated with 17- β -estradiol and tamoxifen for 4 hours underlined that TSC status was involved. The decrease of MMP-2 gene was similar in 4004 and 621-101 (TSC2^{-/-}) cells under the effect of 17- β -estradiol; on the contrary, after the treatment with tamoxifen, the inhibition on MMP-2 was similar in the 4004 and in 621-103 (TSC2^{+/+}).

Although the molecular events responsible for angiomyolipomas development are not fully understood, mutations in TSC1 or TSC2 result in a hyper-activation of the mammalian target of rapamycin (mTOR) signaling pathway [115,116].

Rapamycin and its derivatives, such as everolimus, have been suggested as therapeutic approaches for AML, since they block mTOR pathway, and some clinical trials demonstrated that these drugs reduce AML size. [46; 117].

The first study demonstrating an anti-proliferative effect of mTOR inhibitors in angiomyolipoma, was published in 2008 [118], since then, clinical data suggest that alternative or adjuvant drugs are still needed because of the side effects of the mTOR inhibitor.

The treatment with everolimus reduced the growth already after 24 hours of treatment at different drug concentrations for our cell lines. For TSC2^{-/-} cells the inhibition occurred at 1 and 10 μ M (for 4004 and 621-101 cells respectively); while for 621-103 the inhibition occurred after at 100 μ M. This different sensitivity to everolimus could be caused by mutation in TSC2: when TSC2 is missing mTOR is active, when TSC2 is present (as in 621-103) mTOR is inactive. We may suppose that everolimus exert a greater cytotoxic effect in cells in which mTOR is hyperactive; such as on 4004 and 621-101 cells.

The treatment of AML with bisphosphonates such as like zoledronic acid provoked a decrease in cell proliferation probably by inhibiting angiogenetic factors or inducing microenvironment changes that decreased tumor growth [97, 99,119,100]. In all cell lines, the association of these drugs caused a inhibition of cell growth at any incubation time.

Moreover, the simultaneous incubation of zoledronic acid and everolimus potentiate their inhibitory migration properties; however the combination of these drugs had major effect on female 621-101 cells compared to the male ones. The association of everolimus and zoledronic acid led to a double action on the mTOR pathway: everolimus acted directly on mTOR, zoledronic acid acted further upstream, inhibiting the action of GTPase Rho and leading to a consequent inactivation of mTOR.

Our encouraging results on AML cells allow us to affirm that combination of these drugs may be used to limit the side effects of high everolimus doses. Additionally, we have studied drugs with different mechanism of action, such as simvastatin, that have been proposed to inhibit AML cell growth and migration through the signal transduction Akt-mTOR axis. In our study, the treatment with simvastatin inhibited both proliferation and three dimensional migration of our cells, so it may be a potential therapeutic agent for the treatment of AML.

Mevalonate is not only a precursor of cholesterol synthesis but also a precursor of nonsteroidal isoprenoid compounds. In a precedent study, Martin and colleagues described that statins inhibit the activity of PPAR γ [120]. PPAR γ has been classically regarded as a master regulator of adipogenesis [121], but more recent studies have demonstrated its involvement in various cellular processes, including proliferation, apoptosis, angiogenesis, and cancer [28]. On these bases, we can speculate that the action of statin observed on our cells might depend on PPAR γ , as recently observed on in vivo xenograft AML model by Pleniceanu and colleagues. Therefore, also in our cells PPAR γ might be a new potential therapeutic target for AML.

To clarify the signaling pathways underlying everolimus, zoledronic acid and simvastatin mediated responses in AML cells we further examined the effect of these drugs on the activation of the AKT and ERK pathways. We found that the treatment for 4 hours with zoledronic acid and everolimus significantly suppressed the phosphorylation of Akt in 621-101 cells; in the other cell lines the reduction of this phosphorylation was observed after treatment with everolimus or simvastatin. Treatment with simvastatin induced a reduction in ERK phosphorylation only in 621-101 and 4004 cells only.

A striking difference was observed between cell lines in terms of Akt and Erk phosphorylation and the modulation of the expression of genes involved in adhesion/invasiveness despite the similar response to drugs in terms of cell migration.

It is important to underline that all drugs exerted important effects in regulation of MMP-2 and FN-1 genes, both implicated in neoplastic aggressiveness, in 621-101 female cells compared to 4004 cells

and also, in 4004 and 621-103 cells. The differences observed in the expression of these important genes involved in migration, suggest that it is necessary a different therapeutic approach in female and male patients with TSC-associated or sporadic AML.

In conclusion, the results obtained from primary and stabilized cells, for what concerns the estradiol treatment, confirmed the different behavior of female and male cells with or without mutations in TSC2 gene. In either models, estrogen did not modify the proliferation of AML cells, but increased the migration only for the female cells. Moreover, analyzing the different experiments performed on stabilized cells, we can conclude that, in most of our tests, the male 4004 TSC2^{-/-} cells and the female 621-103 TSC^{+/+} cells had a similar behavior, such as that the restoration of the mutation was responsible for masculinization of the cellular phenotype.

These studies have provided valuable results to support the hypotheses seen in the literature of the involvement of estrogens in the migration of AML toward other sites such as lung and potential development of LAM. This might be of great importance under a diagnostic and prognostic point of view, since patients with AML might be monitored more closely to identify LAM as soon as possible. Furthermore, our results could lead to the definition of an effective drug therapy that almost completely replaces, or delays, AML mass removal surgery or nephrectomy in order to offer a higher quality of life to the patient.

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7. PUBLICATIONS

- Francesca Bertolini, Giulia Casarotti, Luisella Righi, Enrico Bollito, Carlo Albera, Silvia Anna Racca, Donato Colangelo, Barbara Mognetti. Human renal angiomyolipoma cells of male and female origin can migrate and are influenced by microenvironmental factors. Submitted for publication to Plos One.
- Giuseppe La Montagna, Francesca Bertolini, Barbara Mognetti. Zoledronic acid and leuprorelide acetate affect DU-145 migration towards stem cell conditioned medium. Submitted for publication to The Prostate.

Orbassano, October 16th, 2017

Dear Editor, dear Reviewers We are grateful for the determination of the Academic Editor.

We hope we fulfilled your request, we thank you for having positively reconsidered our manuscript for publication and we are waiting for a your final communication on PLOSONE decision. Please do not hesitate to contact us for any further information you might need.

Best Regards Bonbaro Mognetti Barbara Mognetti

1 2	Human renal angiomyolipoma cells of male and female origin can migrate and are influenced by
3	microenvironmental factors
4	
5	Running title: In vitro migration of cells derived from human renal angiomyolipoma
6	
7	<u>Authors:</u>
8	Francesca Bertolini ¹ , Giulia Casarotti ² , Luisella Righi ³ , Enrico Bollito ⁴ , Carlo Albera ¹ , Silvia Anna Racca ¹ ,
9	Donato Colangelo ² , Barbara Mognetti ^{1*} .
10	
11	¹ Department of Clinical and Biological Science, University of Turin, Regione Gonzole 10, 10043, Orbassano,
12	Italy
13	² Department of Health Sciences, Università del Piemonte Orientale, Via Solaroli 17, 28100 Novara, Italy
14	³ Pathology Unit, Department of Oncology, University of Turin, Regione Gonzole 10, 10043, Orbassano, Italy
15	⁴ Pathology Unit, San Luigi Gonzaga Hospital, Regione Gonzole 10, 10043, Orbassano, Torino, Italy
16	
17	* <u>barbara.mognetti@unito.it (BM)</u>
18	
19	
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21	Angiomyolipoma; Lymphangioleiomyomatosis; Cell Migration; Estrogen Effect; In Vitro Techniques
22	

23 Abstract

Improving the knowledge of angiomyolipoma physiopathology might help in understanding the link with 24 25 pulmonary lymphangioleiomyomatosis lesions. We investigated if angiomyolipoma cells have migratory 26 properties, how their growth and motility can be influenced by the hormonal milieu, and if this can be related 27 to a specific gender. Primary cells were isolated from angiomyolipomas surgically resected for therapeutical 28 reasons in a female and in a male patient. Bi- (wound healing) and three-dimensional (transwell assay) 29 migration were analyzed in vitro in basal conditions and under the influence of 17- β -estradiol and SDF-1 α . Treatment up to 72 hours with 17-β-estradiol (0.1-100 nM), tamoxifen (0.2-20 μM) or with both, does not 30 modify angiomyolipoma cells proliferation. On the other hand, SDF-1 α and 17- β -estradiol treatment induce a 31 significant motility increase (both bi- and three-dimensional) which becomes evident already after 2 hours of 32

both the metalloproteases principally involved in malignant phenotype acquisition, i.e. MMP-2 and MMP-9.
Angiomyolipoma cells behave similarly, despite their different source.

incubation. Angiomyolipoma cells express mRNA coding for SDF-1 α and 17- β -estradiol receptors and secrete

Primary angiomyolipoma cells migrate in response to hormonal milieu and soluble factors, and produce active metalloproteases, both aspects being consistent with the theory claiming they can migrate to the lungs (and/or other organs) and colonizing them. No main feature, among the aspects we analyzed, seems to be referable to the gender of origin.

40

41 Introduction

Widespread use of cross-sectional imaging of kidneys has resulted in a significant increase in incidentally 42 43 diagnosed small masses. The prognosis is usually favourable since they rarely progress to metastases [1]. Angiomyolipomas (AMLs) most commonly occur in the kidneys as small masses and, although often 44 45 asymptomatic, may enlarge and bleed leading to haemorrhage and renal impairment [2]. These mesenchymal lesions are characterized by proliferation of spindle cells, epithelioid cells and adipocytic cells in concert with 46 many thick-walled blood vessels [3]. A normal tissue counterpart has not been identified and genetic analyses 47 indicate that all three tissue components derive from a common progenitor cell [4,5]. In case of intractable 48 49 pain, large mass size (>4 cm) and risk of bleeding, surgical intervention is needed [6]. The preferred treatment for AML is nephron-sparing surgery or selective renal artery embolization, since both methods preserve 50 residual renal function in comparison to radical nephrectomy [7]. On the other hand, asymptomatic patients 51 are managed conservatively with long-term surveillance (mainly by imaging). 52

Although most AMLs are clinically insignificant benign tumors, an uncommon subtype, the epithelioid AML,
can behave more aggressively and develop distant metastases [8,9,10].

55 AMLs are twice as common in females, and can occur sporadically or in association with other disorders, such 56 the autosomal dominant condition Tuberous Sclerosis Complex (TSC) and sporadic 57 lymphangioleiomyomatosis (LAM). In particular, LAM is a progressive disease of the lung histologically characterized by a diffuse proliferation of atypical smooth muscle cells (LAM cells) in the alveoli and cystic 58 59 degeneration of the normal lung parenchyma [11].

AML and LAM share the same origin from mesenchymal perivascular epithelioid cell (PEC) and therefore both are considered as belonging to the PEComas lesion family [12]. The smooth muscle–like LAM cells that diffusely infiltrate the lungs and the lymphatic vessels have a low proliferation index and little or no evidence of cellular atypia. In the handful of patients who have had multiple tissues available for sequencing, identical inactivating mutations of TSC1 or TSC2, with subsequent deregulation of the Rheb/mTOR/p70S6K pathway, were demonstrated in AML, in lymph nodes, and in pulmonary LAM cells, but not in normal lung of the samepatient [13].

67 It has been also shown that both AML and LAM cells share immune-expression of HMB-45 antigen [14,15].

Furthermore, both LAM and AML cells express estrogen receptor α [16], and estrogen is thought to cause clinical worsening in women with LAM [17]. Even more strikingly than AML, LAM preferentially affects women, especially at childbearing age, more often than AML. To date the underlying reasons for this behaviour are not known.

These, and other data [18,19], support a model in which both LAM and AML pathogenesis share some genetic and biological mechanisms, and are consistent with the hypothesis that pulmonary LAM might result from the metastatic spread of AML smooth muscle cells [20], possibly influenced by the hormonal milieu.

Therefore, considering the diffuse approach of delaying AML ablation in asymptomatic patients to preserve renal function, and that there is no reliable imaging technique able to differentiate a benign AML from one undergoing malignant change, we deem of paramount importance the study of migratory properties of AML cells. A clear demonstration that AML cells migration is involved in pathogenesis of lesions in several different organs might suggest important hints on new preventive drug therapies.

For this reason, we undertook a study on the proliferative and migratory properties of primary cells isolated from two different surgically removed AMLs, respectively from a male and a female patient. We evaluated if 17- β -estradiol could modulate their growth and their two- and three-dimensional migration. Furthermore, we compared the hormone-dependent effects to those induced by the stromal cell-derived factor 1 α (SDF-1 α), a soluble factor known to induce cellular migration [21]. We also investigated metalloproteases 2 and 9 (MMP-2 and MMP-9) activities of AML cells, both in basic and stimulated conditions, since these MMPs play a pivotal role in the pathogenesis of cystic lung destruction in LAM [22,23].

MMPs modification of the extracellular matrix usually contributes to cell migration as well as to tissue
invasion and metastasis. Similar modifications may facilitate AML cell migration and pulmonary colonization
[20]. In fact, MMPs imbalance, together with other factors like a strong expression of cathepsin K, bcl-2 and
HMB-45, characterize this pathology [19].

We focused our attention on estrogens in order to investigate if there are some differences in AML cells
behaviour according to the hormonal milieu, as suggested by the hypothesis that pulmonary LAM in women
might derive from the metastatic spread of AML abnormal smooth muscle cells.

95 Materials and Methods

96 Materials

All reagents were purchased from Sigma (St. Louis, MO, USA) unless otherwise stated. Tissue culture
plasticware was from Falcon (Franklin Lakes, NJ, USA).

99

Angiomyolipoma cells, tissue and ethical approvals

Human primary AML cells have been obtained from patients that underwent surgical nephron-sparing AML ablation for therapeutic purposes at the Urology Unit of the San Luigi Gonzaga Hospital. The study was approved by the Ethical Committee of the San Luigi Gonzaga Hospital, University of Turin, Orbassano, Turin, Italy (Protocol 0006771, approved on April 18, 2016). All patients provided written informed consent in accordance with the Declaration of Helsinki.

AML3 derives from a male patient, AML4 from a female patient. None of the patients had any clinical signs

07 or symptoms or a family history of tuberous sclerosis. No genetic analysis of TSC mutation were performed.

.08 AML diagnoses were confirmed by standard histological examination including specific immunostaining for

alpha-smooth muscle Actin, HMB-45 and Pancytokeratin antigens.

Primary cells were isolated from excess material not required for diagnostic use, which was divided into small fragments and treated with type II collagenase. Resulting cell suspensions were plated into T25 tissue culture flasks in AML medium (adapted from Lesma et al. [24]), composed of phenol red DMEM medium, ferrous

sulphate 1.6 μ M, 20% foetal calf serum (FCS) and 10 μ g/mL epidermal growth factor. Experiments were

14 performed on cells at passage 3-6.

15

16 Immunofluorescence

AML cells were fixed in 4% paraformaldehyde (PAF) for 15 minutes. After washing in PBS cells were treated with PBS containing 1% normal goat serum (NGS), 0.1% Triton X-100 at room temperature (RT) for 1 hour.

.19	Cells were incubated overnight at 4°C with the following primary antibodies against: S-100 (rabbit, 1: 800;
20	Dako, Glostrup, Denmark), α-Smooth-Muscle Actin (mouse, 1: 100; NeoMarkers, Fremont, CA), HMB45
21	(mouse,1:100, Dako, Glostrup, Denmark), Keratin 8/18 (mouse, 1:100 Menarini, Florence, Italy) Vimentin
22	(mouse, 1:70; Novocastra Lab, Newcastle, UK) and Melan-a (mouse; 1:100; NeoMarkers, Fremont, CA).
23	After washing, cells were incubated for 1 hour at RT with the appropriate secondary antibodies: goat anti-
24	mouse IgG Alexa-Fluor-488-conjugated (1:200, Molecular Probes, Eugene, Oregon) and CY3-conjugated
25	anti-rabbit IgG (dilution 1 :400, Dako, Milan, Italy) [25]. The immunostained coverslips were analyzed on a
26	Zeiss fluorescence microscope and images were captured with an Axiovision Imaging System.

.27

28 2D migration assay-wound healing

Two-dimensional migration assays were performed as described in Mognetti et al. [26]. Briefly, primary AML cells were seeded in a 12-well plate at 300,000 cells/well. When they were confluent, a cross "wound" was made in each well with a p1000 tip [27], then wells were washed thrice with PBS, and cultured in AML medium supplemented either with SDF-1 α (Peprotech, London, UK) 100 ng/mL, plerixafor 100 nM, 17- β estradiol 1 nM, or tamoxifen 2 μ M. We photographed "wounds" on time-laps every hour to highlight migration, until a maximum of 8 hours.

Experiments were repeated three times, and every time five different spots for each experimental condition were considered. Images were analysed using ImageJ software (Wayne Rasband, NIH, USA): the healing percentage was quantified comparing the wound area at t = 0 to the following time-points for each treatment.

.38

3D migration assay

The transwell migration assay, performed as previously described in Mognetti et al. [28], was used to measure the three-dimensional movements of cells. Migration assays were performed in transwells (BD Falcon cell culture inserts incorporating polyethylene terephthalate membrane with 8.0 μ M pores, $6\pm 2x10^4$ pores /cm²) in 24-well plates. When tests were performed in the presence of SDF-1α blocker, cells were preincubated for 30 min at 37°C in
100 nM plerixafor conditioned medium.

Cells $(5x10^4)$ were suspended in 200 µL of culture medium and seeded in the upper chamber of a transwell. .46 The lower chamber was filled with fresh culture medium with or without 100 ng/mL SDF-1 α or 17- β -estradiol .47 1 nM, and placed in the incubator. After 4 or 8 hours, cells were treated as detailed by Gambarotta et al. [27]. .48 .49 Wells were photographed using a BRESSER MikroCam 3 Mpx camera, with an optical microscope (Leica DC 100) at 100x. Five pictures were randomly chosen per well and used to count the migrated cells with ImageJ 50 software using cell-counter plug-in. Results from different experiments (performed at least three times in .51 duplicate) were expressed as mean ± standard errors. In order to avoid any cytotoxic effect potentially .52 53 confounding migration results, we performed a cytotoxicity test at the same time and same conditions of every migration test. .54

55

56 **Proliferation assay**

Primary AML cells were seeded into flat-bottomed 96-well microplates at a density of 1,000 cells in 100 μ L culture medium *per* well and allowed to attach overnight in complete medium. Drugs were then added to culture medium at concentrations ranging from 0.1 to 100 nM 17-β-estradiol, and from 0.2 to 20 μ M tamoxifen [29] for 24–72 hours, according to protocols. The MTT (3-(4,5-dimethylthiazol-2-yl)-2,5diphenyltetrazolium bromide) assay was performed as described in Mognetti et al. [28]. Data (mean ± standard errors) were calculated as the mean values of 8 replicates. Each experiment was repeated thrice. Cell viability was expressed as the percentage of living cells *versus* the untreated controls.

.64

65 Quantitative real-time PCR (qPCR)

In order to perform quantitative real-time PCR (qPCR), total RNA was extracted from treated cells by using
Trizol (Invitrogen Life Technologies, Italy). After RNA purification and treatment with DNAse I (Fermentas,
St. Leon-Rot, Germany), 1 µg was retrotranscribed in cDNA with the RevertAid[™] H Minus First Strand

cDNA Synthesis Kit (Fermentas) using oligo(dT) primers. Gene assays were performed in triplicate for each .69 treatment in a 20 µL reaction volume containing 1 µL of RT products, 10 µL Sso-Fast EVA Green SMX (Bio-.70 Rad, Hercules, CA, USA), 500 nM each forward and reverse primers. Gene expression was normalized on the .71 housekeeping gene ribosomal 18S rRNA. Table 1 resumes the primer sequences that were used. Automated .72 .73 CFX96 real-time thermocycler (Bio-Rad) was used and the reaction conditions were 95°C for 1 minute, followed by 45 cycles 98°C for 5 seconds and anneal-extend step for 5 seconds at 60°C, with data collection. .74 At the end of these cycles, a melting curve (65°C to 95°C, with plate read every 0.5°C) was performed in .75 .76 order to assess the specificity of the amplification product by single peak melting temperature verification. Results were analysed with Bio-Rad CFX Manager. Calculations and statistical analyses were performed .77 using GraphPad Prism version 5.00 for Windows (GraphPad Software, San Diego California, USA). .78

.79

.80	Table 1. Primers seq	uences, size of the	e amplification	product and NCBI	Reference Sequence
		,			

GENE		SEQUENCE	AMPL. SIZE	NCBI REF. SEQ.
Ene	Fw:	5'-TGGAGTCTGGTCCTGTGAGG-3'	172 h	NT 025741 16
Erα	Rev:	5'-CCCACCTTTCATCATTCCCACT-3'	172 бр	N1_025741.16
E-0	Fw:	5'-GAGCAAAGATGAGCTTGCCG-3'	142 hr	NIM 001427.2
Егр	Rev:	5'-AGCTGGGCCAAGAAGATTCC-3'	142 bp	NM_001437.2
CDD20	Fw:	5'-AGTCGGATGTGAGGTTCAG-3'	240 hz	NIM 001505 2
GPK30	Rev:	5'-TCTGTGTGAGGAGTGCAAG-3'	240 бр	NIVI_001505.2
	Fw:	5'-GGCCCTGTCACTCCTGAGAT-3'	474 ha	NIM 001202510.1
MIMP-2	Rev:	5'-GGCATCCAGGTTATCGGGGA-3'	474 bp	NM_001302310.1
	Fw:	5'-CAACATCACCTATTGGATCC-3'	490 h.c	NIM 004004 2
MIMP-9	Rev:	5'-CGGGTGTAGAGTCTCTCGCT-3'	480 bp	NM_004994.2
195 "DNA	Fw:	5'-GTGGAGCGATTTGTCTGGTT-3'	201 hn	X02205 1
105 fKNA	Rev:	5'-ACGCTGAGCCAGTCAGTGTA-3'	201 op	A03203.1

Fw=forward, Rev= reverse

.81 .82

.83

.84

85 Western blotting

Cells were seeded in 10 cm diameter Petri dishes, cultured until sub-confluence, then 17-β-estradiol (1 nM)
was added. After 5 minutes and 4 hours incubation, cells were collected and treated as detailed by Mognetti et
al. [21].

Blots were probed with primary monoclonal antibody anti-vinculin (mouse; 1: 2000 Sigma) resuspended in PBS Tween 0.1% and with anti-ERK1/2 (mouse; 1:2000), anti-pERK1/2 (mouse; 1:2000) (Cell Signaling Technology, Danvers, MA, USA) resuspended in 5% w/v nonfat dry milk + PBS tween 0.1%. Vinculin was used as an internal control. HRP-conjugated anti-mouse (Amersham-GE Healthcare, Buckinghamshire, UK) was diluted 1:6000 (ERK1/2 and pERK1/2) and 1:8000 (vinculin) in PBS Tween 0.1%. Densitometric analysis was performed by ImageJ software.

The ratio pERK1/2/ ERK1/2 was expressed as percentage optical density modification relative to control conditions. Experiments were repeated three times.

.97 .98

.99 Gelatin zymography

200 MMP-2 and MMP-9 activities in medium samples were assessed by gel zymography. Proteins (100 µg) were separated by electrophoresis in 8% SDS-PAGE gel containing gelatin (0.8 mg/mL) under non-reducing 201 202 conditions. The gel was washed with Tris buffer (2.5% Triton X-100 in 50 mM Tris-HCl, pH 7.5, final solution) for 1 hour, then incubated overnight at 37°C in a proteolysis buffer (40 mM Tris-HCl, 200 mM 203 NaCl, 10 mM CaCl₂, 0.02% NaN₃, pH 7.5, final solution). The gel was stained for 3 hours with Coomassie 204 Blue solution (0.05% Coomassie Brilliant Blue R-250, 50% methanol, 10% acetic acid, final solution) and 205 finally destained with 5% methanol and 7% acetic acid (final solution). Reagents and chemicals were obtained 206 from VWR International (Milan, Italy). MMPs activity was detected as a clear band on a blue background and 207 estimated by densitometric analysis using ImageJ Software. The results were expressed as percentages of 208 209 control values.

211 Statistical analysis

- All the data in this study are shown as the mean \pm SE. Two group means were compared using the unpaired t-
- test, and more than two group means were analyzed by one-way analysis of variance (ANOVA), where P
- 214 <0.05 was considered statistically significant [30].</p>
- For gene expression level comparison One-way ANOVA with Dunnett's post tests were performed using
- GraphPad Prism version 5.00 for Windows.
- 217

Results

Cell characterization by immunofluorescence

To better characterize isolated AML primary cells, immunofluorescence was performed with specific 220 221 antibodies. Overall, both cell lines showed similar immunophenotype and partly elongated or rounded shapes. Both primary culture cells were strongly and totally positive for smooth muscle actin antibody (in both 222 223 elongated and rounded cells), with a diffused stain throughout the cytoplasm (Fig 1a and 1a'). Furthermore, in 224 both cell lines there were single rounded element positive for keratin 8/18 and for elongated cells strongly positive for vimentin (Fig 1d, 1e, 1d' and 1e'), together with single negative ones for both the antigens. Finally 225 226 a strong nuclear and cytoplasmic positivity was found for \$100 in both cells (Fig 1f and 1f'). As a matter of 227 fact, even if scattered, some AML3 cells were focally positive for intracytoplasmic HMB45 (Figure 1b and 228 1b'), and AML4 cells were focally positive for Melan-A antigen (Fig 1c and 1c'), which is consistent with the 229 AML phenotype.

230

Figure 1: Primary angiomyolipoma cells characterization by immunofluorescence.

Cells isolated from the two angiomyolipomas were challenged with specific antibodies to reveal their immunocytochemical characteristics: α -actin antibody (a and a') specific for smooth muscle cells; HMB45 (b and b') and Melan-A (c and c'), both typical of AML; keratin 8/18 (d and d') labeling the epithelial-like cells; vimentin (e and e'), a marker of fibroblasts and S100 (f and f'), a marker of lipid-containing cells. Fields were chosen to show both the morphological aspect and the specific marker expression.

237

Estrogen and SDF-1α receptors gene expression

The analyses of gene expression of both AMLs demonstrated the presence of mRNA for CXCR4 (SDF-1 α receptor), GPR30, ER α , but not for ER β (Fig 2). The level of each gene was similar in both primary AML cells, with no significant difference.

Fig 2. Receptors gene expression.

Early passages AML cells underwent qRT-PCR for CXCR4, GPR30, ER α and ER β mRNA expression analysis. Data are shown as the absolute mRNA expression normalized by the housekeeping 18S rRNA.

246

247 ERK phosphorylation

Treatment of AML3 cells with $17-\beta$ -estradiol increased ERK phosphorylation at 4 h, while pERK was significantly augmented in AML4 cells already at 5 minutes and was stable until 4 hours (Fig 3).

250

251 Fig 3. ERK phosphorylation

Effect of 17-β-estradiol (1 nM) on pERK/ERK in AML3 and AML4 cells after 5 minutes and 4 hours of incubation. Vinculin as internal control. *=P<0.05 vs control.

254

Influence of 17-β-estradiol or tamoxifen on AML cell proliferation

The treatment for up to 72 hours with concentrations of $17-\beta$ -estradiol ranging from 0.1 nM to 100 nM with or without tamoxifen, or with tamoxifen alone (0.2-20 μ M, two representative experiments are shown in Fig 4) did not induce any modification on AML cells proliferation regardless of their gender. No significant difference was detected at any time point or culture condition (data not shown). Figure 4 reports two typical experiments as an example.

261

Fig 4. Effect of 17-\beta-estradiol alone and of its combination with tamoxifen on AML cells growth. Proliferation assay after 72 hours culture in presence of increasing concentration of 17- β -estradiol (A) or with 17- β -estradiol 1 nM, tamoxifen 2 μ M, or both (B). Anova and Dunnett's post test analyses demonstrated that no significant modification in cell growth respect to untreated controls was induced by the molecules at any of

the concentrations tested.

Two-dimensional Motility Assay (wound healing)

The two-dimensional motility was quantified, and data are displayed graphically as healing percentage (Fig 5, panels C, D and E). Panel C compares basal migration: the early migration rate of AML3 was higher, a significant difference being demonstrated at 4 hours. This difference is promptly quenched since after 8 hours the migration rate of the two AMLs was similar.

273

Fig 5. Analyses of primary AML cells migration by *in vitro* wound healing assay.

Wounded area in a representative experiment of *in vitro* wound healing assay is shown before (A) and after (B) the incubation period. Bi-dimensional migration was then quantified in basal conditions for both AMLs (C), and for AML3 (D) and AML4 (E) in presence of SDF-1 α 100 ng/mL or its receptor antagonist plerixafor 100 nM, or a combination of both. Migration is expressed in arbitrary units. *=P<0.05 vs control; #=P<0.05 vs SDF-1 α .

280

SDF-1 α treatment induced a statistically significant motility increase already after 2 hours treatment in cells from both AMLs. Significant difference persisted all along the experimental period (up until 8 hours). The effects induced by SDF-1 α were abolished by the SDF-1 α -receptor antagonist plerixafor, while no significant effects were induced by plerixafor alone.

Two-dimensional migration was significantly modulated by 17- β -estradiol (Fig 6), although some differences in migration patterns were evident. In fact, AML3 (A) of male origin, showed a significant motility increase in the first 4 hours of incubation with 17- β -estradiol, and a prevalent logarithmic pattern. AML4 cells, of female origin (B), showed an exponential pattern and a significant increase in estradiol induced migration respect to the untreated control, evident at any of the time point considered. The treatments with the ER-antagonist tamoxifen had no influence on two-dimensional motility of both cell types, while it was able to abolish the effects of estradiol.

Fig 6. Wound healing assay in presence of 17-β-estradiol 1 nM and/or tamoxifen 2 μM.

Data are expressed as the percentage of migration vs t = 0 h. *=P<0.05 vs control; [#]=P<0.05 vs 17- β -estradiol.

296 Three-dimensional Migration Assays

Cells were seeded in the upper chamber of a transwell filter and allowed to migrate for 4 or 8 hours under either basal conditions, or in response to stimuli added in the lower chamber. Fig 7, panel A and B, shows that the number of migrating cells significantly increased in response to SDF-1 α .

800

Fig 7. Three-dimensional migration (SDF-1α).

Cells were incubated with SDF-1 α (100 ng/mL) and its receptor blocker plerixafor (100 nM) or a combination of both for 4 or 8 hours. In each experimental condition cells were counted in 5 fields *per* insert. Data are expressed as the percentage of migration *vs* control. *=P<0.05 vs control; #=P<0.05 vs SDF-1 α .

305

Data were similar for cells originated from the two AMLs and were significant after 4 hours of continuous exposure. SDF-1 α stimulation was completely abolished by the SDF-1 α -receptor antagonist plerixafor, while no differences were shown for longer exposures.

The transwell migration experiments performed in presence of 17-β-estradiol indicated a different behavior

between cells originating from the two AML. AML3 cells of male origin needed at least 8 hours of stimulation

before a significant difference in migration could be appreciated (Fig 8A). AML4 cells (Fig 8B) responded to

the stimulation at 4 hours, while after 8 hours no differences respect to the control were evident.

313

Fig 8. Three-dimensional migration (17-β-estradiol).

Cells were incubated with 17- β -estradiol (1 nM) for 4 and 8 hours. Migration is expressed in arbitrary units. *=P<0.05 vs control.

818	Metalloproteases activity in supernatant derived from 3D-migration test
819	The activity of two metalloproteases involved in malignant phenotype acquisition, MMP-2 and MMP-9, was
820	measured by zymography in the supernatant collected at the end of the 3D-migration test (4 hours) (Fig 9).
821	
822	Fig 9. Metalloproteases activity in supernatant collected from 3D-migration test.
323	Data are expressed as the relative activity calculated by densitometric analyses. *=P<0.05 vs control;
824	$^{\#}=P<0.05 \text{ vs SDF-1}\alpha.$
825	
826	While the diverse treatments induced no significant difference in MMP-2 activity, SDF-1a enhanced MMP-9
827	enzymatic activity. This data was in accordance with the increased migration induced by SDF-1 α . Coherently,
828	plerixafor inhibited MMP-9 activity increase provoked by SDF-1a. No difference was induced by incubation
829	with 17-β-estradiol.
30	Figure 10, panel A, shows absolute MMPs activity (not normalized) and demonstrates that MMP-2 activity, in
31	basal conditions, is higher than MMP-9 activity for both AMLs. Consistently, Fig 10B, shows coherent
32	mRNA expression for both enzymes.
333	
34	Fig 10. Absolute MMP-2 and MMP-9 activity (A) and their corresponding mRNA expression (B) in
35	basal conditions.
836	Data are expressed as the activity calculated by densitometric analyses. The gene expression is shown as the
37	level of expression normalized by the housekeeping 18S rRNA. The difference in MMP-2 vs MMP-9 activity
38	or expression is always significant (P<0.001).
39	
840	

Discussion

842 In this paper we describe for the first time the behaviour of primary AML cells, originating from male and 343 female patients, in terms of proliferation and migration in vitro, both in basal conditions and in response to environmental stimuli. Our findings clearly demonstrate that primary AML cells are able to migrate *in vitro*. 344 345 The main limit of this study was, by far, the small sample number. It is worth underlying, though, that AML is 846 a rare disease and it is quite difficult to come across AML of male origin, and to isolate and grow primary 847 cells, what we actually did. The cellular composition of the cultures were similar, as demonstrated by 848 immunofluorescence and shown in Fig. 1. Overall, this phenotype confirmed the mixed (either muscle, 849 epithelioid, lipomatous and mesenchimal) nature of the cultures, respectively, according to the heterogeneous nature of AML [24]. None of the cell type seemed to be predominant; we consider the miscellaneousness of 850 the two cultures an advantage in a pharmacological study, since the complexity of the tumor should be taken 851 352 into account.

353 The original aim of this study was to understand if AMLs, independently from their TSC mutational status, 354 were able to migrate and if there was the possibility to modulate this process pharmacologically. In particular, we focused on estrogen because of the underlying hypothesis that in females during childbearing age AML 355 cells might migrate to the lungs and generate LAM. We also investigated SDF-1 α since this factor is usually 856 857 associated to migration and homing of several cell types [31,21]. We have shown that both AML3 (male origin) and AML4 (female origin) have analogous significant expression levels of estrogen and SDF-1a 858 receptor mRNAs. Despite the presence of the estrogen receptors, we have observed that there was no 359 estrogenic influence on proliferation of both AMLs. Therefore, the hormone therapy might not be optimal 860 pharmacological choice to treat this pathology in female patients, and other options should be taken into 861 862 consideration. Another aspect that we demonstrated in common for both AMLs was their migratory response to SDF-1a. In fact, we have shown its significant effect in increasing the bi- and three-dimensional migration 863 of these cells. The specificity of this observation was confirmed by the effect of the treatment with the 864 865 selective SDF-1 α receptor antagonist plerixafor, which completely abolished SDF-1 α migratory stimulation.

Our results are in accordance with the well known role of SDF-1 α in cell migration, and might support the hypothesis that the host microenvironment tissue stimuli could exert a specific chemotactic signal to promote homing of AML cells [32].

Although some common properties between the two AMLs have been described so far, we report that other aspects seem to differentiate the cells originating from male from those deriving from female patients. We showed that the basal unstimulated migration of male AML3 had a more rapid onset, respect to female AML4. This difference in spreading was striking already 4 hours after seeding, and was particularly evident in the ability to invade, as demonstrated by three-dimensional assays. Alongside with this observation, we also report that the higher basal migration of AML3 is less influenced by estrogen, while spreading of AML4 cells is significantly increased already after 2 hours incubation with estradiol 1 nM.

In order to clarify the short-term effects of estrogen, we investigated the expression of ER α , ER β and GPR30. It is worth to underline an abundant mRNA expression of GPR30 [33], which might likely mediate the early response to estradiol that we observed.

In this work we present some data on the response of AML cells to hormonal stimuli. While many Authors 379 have thoroughly described the long term effects of these stimuli [34, 35], their influence on the colonization 880 881 potential and short term effects needs further research. In particular, our attention focused on GPR30 role in these processes, and we have chosen the timing in order to better analyze the responses referred to this 882 receptor pathway. The multiple effects of estrogens can be explained by different modulation of transcription 883 884 and by rapid signaling events that are not associated with altered gene transcription [34]. GPR30 hormone stimulation is able to induce rapid MAPK activation pathway and ERK1/2 phosphorylation via MMP-EGFR 885 and is responsible for several cell responses and signaling. 886

We have included to this paper some Western blot experiments that demonstrate that E2 stimulation is able to increase ERK1/2 phosphorylation in AML4. Noteworthy, both AML3 and AML4 constitutively express significant levels of ERK1/2 and pERK. In particular, AML3 has a basal level of pERK higher than AML4, and this might explain the reduced effects of E2 on pERK/ERK observed in AML3. These data are in accordance with the 2D and 3D migration data shown in Figure 6 and Figure 8 and justify why the early response to estradiol in migration of AML4 cells is not related to a higher mRNA expression compared to AML3.

ERK protein is activated when tuberin function is lost [36], and estrogen-mediated non-genomic ERK signaling activated by GPER is involved in cell viability and motility of TNBC cells [37].

An interesting aspect of GPR30-E2 stimulation is the activation of the PI3K-Akt signaling pathway via mTOR and S6K that leads to different activity on DNA transcription and on proliferation. This pathway is activated from both E2 and tamoxifen, and it could explain, at least in part, the data shown in Figure 6.

Analysis of the culture media collected from three-dimensional migration experiments revealed that the stimulation of migration induced by SDF-1 α was accompanied by an augmented release of MMPs. These effects were consistently reverted by the selective receptor antagonist plerixafor. Strikingly, such an increase in MMPs production was not observed in estradiol-stimulated conditions, despite a significant increase in cell migration. We therefore speculate that, while SDF-1 α -induced migration occurs through metalloproteases production, the migratory response to estrogen most likely does not involve MMPs activity.

Noteworthy we have shown that tamoxifen induced unexpected effects on migration of the two cell types. The peculiar mechanism of action of this drug, which can bind both intracellular and membrane estrogen receptors, might give us some hints in the interpretation of our results. Further characterization of the cells may be of value, but it is outside of the scope of the present study. We are currently conducting more exhaustive

investigations on this topic, also trying to correlate the different aspects to TSC mutational status, but since

they go further the aim of the present study, the data will be reported in the next future.

Our work demonstrates that primary AML cells migrate and produce active metalloproteases, both aspects being consistent with the theory claiming these cells can migrate and invade other tissues, and for some yet unknown reasons colonize the lung.

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Zoledronic acid and leuprorelide acetate affect DU-145 migration towards stem cell conditioned medium

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Key Words:	prostate cancer, bisphosphonate, cytotoxicity, metastasis, GnRH analogues



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11	6	10043 Orbassano Italy
12	0	10045, Oloassailo, Italy
13		
14	7	
15		
16	8	Correspondence to: Barbara Mognetti, Department of Clinical and Biological Science,
17	9	University of Turin Regione Gonzole 10, 10043, Orbassano (TO)
10	10	
18	10	
19		
20	11	
21	12	Corresponding author: Barbara Mognetti
22	13	Tel.: +39 0116705439
23	14	Fax: +39.0119038639
24	14	
24	15	E-mail: <u>barbara.mognetti(@unito.it</u>
25		
26	16	First author: Giuseppe La Montagna
27	17	Tel.: +39 0116705439
28	18	Fax: +39 0119038639
29	10	E mail: 285103@edu unito it
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35 Abstract

 Background: To study the effect of GnRH analogues (leuprolide acetate, LA) and
bisphosphonates (zoledronic acid, ZA), alone or in combination, on human prostate cancer
(PCa) cells proliferation and migration, in vitro.

Methods: Cell proliferation (MTT assay) and three-dimensional (transwell assay) migration were analyzed on human PCa cell line (DU-145) in basal condition, under drug treatment or in MSC-CM (mesenchymal stem cells conditioned medium), whose cytokines content was previously measured by Milliplex bead immunoassay; finally, we observed by western blot analysis how these LA and ZA modify Akt phosphorylation.

Results: ZA (5 μ M) cytotoxicity appears after 48-hours incubation, while LA cytotoxicity only after 72-hours at 100 μ M. Both subcytotoxic ZA and LA concentrations decrease 3D PCa cell migration rate. pAkt/Akt ratio is diminished by LA and, though less strikingly, by ZA, in agreement with the respective inhibition migration ratios. MSC-CM significantly increases PCa cell migration (210%± 2.2; P<0.05), but this phenomenon is quenched both by ZA and LA.

Conclusions: Our results suggest that LA and, mostly, ZA have a direct toxic effect on cancer
cells. Furthermore, they inhibit cellular migration even under attractive stimuli exerted by
MSC; this might contribute to explain their effect in limiting metastatization.

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Introduction

Prostate cancer is the most frequent genitourinary tumor, representing 11% of the male malignancies in Europe, and is one of the leading causes of morbidity and mortality in the world [1].

Administration of GnRH agonist or other analogues (such as leuprolide acetate, buserelin, deslorelin, goserelin and istrerelin) is a well-established treatment of prostate cancer inducing a pharmacological castration [2] and resulting in cancer regression. Despite pharmacological treatment, the onset of metastatic dissemination represents, together with the development of androgen-independent growth, a critical progression step of human prostate cancer that largely determines the clinical course of the disease and survival of the patients [3]. Bone is a preferential site of metastases [4], which produce a crucial impact on patients' functional status and quality of life due to significant pain and high risk of skeletal-related events, including pathologic bone fractures (both vertebral and non-vertebral), spinal cord compression, surgery and radiotherapy to bone [5]. The burden of metastatic disease can be treated by administering a potent inhibitor of osteoclast activity such as zoledronic acid (ZA), a bisphosphonate widely used to treat skeletal complications of malignancy and considered the drug of choice for both the prevention and the treatment of bone mass loss. In vivo, it inhibits the release of growth factors from osteoblasts and bone marrow stromal cells [6]. Moreover, bisphosphonates modulate many other cellular and physiologic processes relevant to bone metabolism and tumor initiation and progression [7].

Prostate cancer cells tropism for the bone is the result of a sequential series of molecular
events: bone metastases arise as a result of a crosstalk between metastatic cells, bone matrix,
osteoblasts and osteoclasts, and cellular components of the bone marrow microenvironment.
Among these, bone marrow mesenchymal stem cells (BM-MSCs) play a paramount role in

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85	the so-called metastatic niche [8, 9]. Prostate cancer cells migration can be influenced by
86	MSC-CM soluble factors [8], and drugs normally used for prostate cancer treatment, such as
87	zoledronic acid and leuprolide acetate, interact with this process [10]. We aimed at examining
88	the effect of such molecules on the behavior of the human prostate cancer cell line DU-145 in
89	an in vitro cell co-culture model of invasion assay. Cells were exposed to the drugs, alone or
90	in combination, prior and/or during the migration test. We previously demonstrated that an
91	up-regulation of Akt phosphorylation may exert a crucial role in DU-145 cell migration under
92	conditioned medium stimulus [9]. Knowing that PI3-K/Akt signaling pathway plays a critical
93	role in cell invasion and in modulation of cell migration [11], we also examined if zoledronic
94	acid and leuprolide acetate modulate AKT level.
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96	Materials and Methods
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98 99	Materials All reagents were purchased from Sigma (St. Louis, MO, USA) unless otherwise stated.
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110 *Prostate cancer cell culture*

Human androgen independent DU-145 prostate cancer cells were purchased from ATCC
(Rockville, MD, USA). Cells were maintained at 37°C in a humidified 5% CO₂ atmosphere
in RPMI 1640 containing 10 mL/L penicillin and streptomycin solution, NaHCO₃ 2 g/L
(7.5% w/v), 10% Fetal Bovine Serum (FBS).

115

116 **Proliferation assay**

DU-145 cells were seeded into flat-bottomed 96-well microplates $(1,000/100 \ \mu L \ culture$ 117 118 medium/well) and allowed to attach overnight in complete medium before drugs addition. 119 Drugs were added to culture medium, alone or in combination, testing various concentrations 120 from 2.5 μ M to 50 μ M (ZA) [12] and from 0.5 μ M to 100 μ M (LA) [13] for 24–96 hours, according to protocols. The MTT (3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyltetrazolium 121 122 bromide) assay was performed as previously described [14]. Data (mean ± standard 123 deviation) were the average values of 8 replicates. Each experiment was repeated thrice. Cell 124 viability was expressed as percentage of living cells with respect to controls.

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126 Mesenchymal stem cells isolation and MSC-CM collection

Bone marrow cells were obtained from femurs of adult rats as described in Mognetti et al, 2013. They were grown in complete α MEM containing 10% FBS, 2 mM L-glutamine, 100 U/ml penicillin and 100 µg /mL streptomycin at 37 °C and 5% CO₂ for 3 days as previously described [9].

For migration assay, conditioned medium was collected after three days of culture,
centrifuged at 4000 rpm for 5 minutes at 4°C in order to eliminate cells and cellular debris,
and used for migration assays or frozen.

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Three-dimensional migration assay

Three-dimensional (3D) migration assay was used to measure the invasiveness of DU-145 cells in response of various stimuli. Migration assays were performed in transwells (BD Falcon cell culture inserts incorporating polyethylene terephthalate – PET – membrane with 8.0 μ M pores, $6\pm2*10^4$ pores /cm²) as previously described [9].

Briefly, 10^5 cells were resuspended in 200 µL of RPMI containing 2% FBS with or without drugs (zoledronic acid 20 µM, leuprolide acetate 100 µM or both, representing non-toxic concentration at 24 hours as per the described viability assay) and then seeded in the upper chamber of a transwell; in the lower chamber we added RPMI or MSC-CM as detailed in Table I. In some cases, before seeding cells were grown for 18 hours in presence of zoledronic acid 20 µM, leuprolide acetate 100 µM or both (Table I).

147 Transwells were placed in the incubator at 37 °C and 5% CO₂ for 6 hours and finally treated
148 as detailed by Mognetti et al. [9].

Wells were photographed using a BRESSER MikroCam 3 Mpx camera, with an optical microscope (Leica DC 100) at 100x. Five pictures were randomly chosen per well, and used to count the migrated cells with ImageJ software using cell-counter plug-in. Results from different experiments (performed at least three times in duplicate) were expressed as mean \pm standard deviation. In order to avoid any cytotoxic effect of potentially confounding migration results, we performed a cytotoxicity test at the same time and same conditions of every migration test.

157 Immunoassay to detect cytokine content in MSC-CM

Cytokine profiles in MSC-CM were determined using the Human Cytokine/ Chemokine
Magnetic Bead Panel protocol from the "Milliplex[®] Human Cytokine 5 Plex" kit (Billerica,
MA). The procedure was conducted according to the manufacturer's protocol.

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Briefly, the assay plates were washed with washing buffer, and shaken on an orbital plate shaker for 10 minutes at room temperature. The washing buffer was decanted and the standards, assay buffer, or samples/controls were mixed with serum matrix in each well, incubated overnight at 4°C on an orbital shaker with specific antibody to detect GM-CSF, MCP1/CCL2, IL-10, IFN γ , TNF α ; well contents were then removed and wells washed. Biotinylated detection antibodies were then added and incubated for 1 hour at room temperature while shaking. After incubation, well contents were removed and streptavidinphycoerythrin was added and incubated for 30 minutes at room temperature, then washed and resuspended in Sheath Fluid. Plates were read on the Luminex MagPix[®] machine and data were collected using the Luminex xPONENT[®] software (v. 4.2); data analysis was performed using the Milliplex[®] Analyst software (v. 5.1).

173 Western blotting

174 Cells were seeded in 10 cm diameter Petri dishes, cultured until sub-confluence and then 175 drugs were added (ZA 20 μ M and LA 100 μ M, to reproduce the same conditions of migration 176 assay). After 6 hours incubation, cells were collected, treated and immunoblotted as detailed 177 according to Mognetti et al. [9].

Blots were probed with primary polyclonal antibody (Cell Signaling Technology, Danvers,
MA, USA) suspended in TBS Tween 0.1% as follows: anti-Akt (mouse, 1:800), anti-pAkt
(Ser473, rabbit, 1:500), and anti-vinculin (developed in rabbit, Sigma). Vinculin was used as
an internal control.

182 HRP-conjugated anti-mouse (Amersham-GE Healthcare, Buckinghamshire, UK) and anti183 rabbit (Santa Cruz Biotechnology) were diluted (1:6000 and 1:8000, resp.) in TBS Tween
184 0.025%. Bands were quantified using the ImageJ software.

185	Phosphorylation levels of Akt were expressed as ratio pAkt/Akt. All data were expressed as
186	percentage modification relative to control conditions.
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188	Statistical analysis
189	All the data in this study were shown as the mean \pm standard deviation. Statistical analyses
190	were performed by One-way ANOVA, with Dunnett's post tests, or two-way ANOVA using
191	GraphPad Prism version 5.00 for Windows (GraphPad Software, San Diego California USA).
192	Coefficient of Drug Interaction (CDI) was used to define the type of interactions between the
193	employed drugs. CDI was calculated by means of the equation: $CDI = AB/(A \times B)$, where AB
194	is the relative cell migration of the combination; A or B, relative cell migration of the single
195	agent. CDI < 1 indicates a synergistic effect; CDI = 1 indicates an additive effect; CDI > 1
196	indicates an antagonistic effect.
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200	Results
201	Effect of zoledronic acid, leuprolide acetate and combination of both on cell proliferation
202 203	Incubation up to 24 hours with 20 and 40 μ M zoledronic acid induced a weak cytotoxic effect
204	on DU-145 (Figure 1A). A statistically significant cytotoxicity appears after 48 hours of
205	incubation, already at the 5 µM concentration.
206	Leuprolide acetate displayed no cytotoxicity on DU-145 cells (Figure 1B), within the
207	concentration range tested, after 24 and 48 hours of incubation.
208	A significant toxicity is appreciable only after 72 hours of incubation from 5 μ M on.

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209	Addition of leuprolide acetate 100 μ M did not increase zoledronic acid cytotoxicity at any
210	concentration after 48 hours incubation (Figure 1C).
211	
212	3D migration assay
213	
214	Both drugs (ZA and LA) decreased migration rate of DU-145 cells when compared to control
215	conditions (Figure 2). At these conditions, leuprolide acetate was significantly more effective
216	than ZA in inhibiting cell migration. The simultaneous presence of the two drugs influenced
217	migration less than each single drug; association decreased the efficiency of leuprolide
218	acetate alone and did not seem to differ from incubation with zoledronic acid alone.
219	Single drug pre-incubation did not potentiate the effect of simple incubation, in any case
220	(Figure 2). The effects of any single drug were not significantly potentiated by any pre-
221	incubation, even, in some cases, abolished (pre-incubation with ZA or LA and pre-incubation
222	with ZA following by migration with ZA).
223	When cells were pre-incubated with the single drugs before migration test performed in
224	control medium, migration was not inhibited at all. On the other hand, when cells were pre-
225	incubated with both drugs, migration, even if occurring in RPMI, was inhibited as much as
226	when cells were not pretreated but incubated with both drugs simultaneously. MSC-CM
227	significantly increases the rate of migration towards control medium, and the addition of both
228	drugs quenched the attractive effect of conditioned medium (Figure 2).
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230	Drug interaction

CDI results suggest an antagonistic effect in the incubation LA + ZA, slightly additive for
Pre-incubation ZA + incubation LA and synergistic for Pre-incubation LA + incubation ZA
as reported in Table II.

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235	Cytokines content in MSC- CM
236	Cytokines were detected in MSC-CM as follows:
237	• GM-CSF: 93.40 pg/ml
238	• MCP1/CCL2: 12.95 µg/ml
239	• IL-10: 1.81 pg/ml
240	• IFNγ: 30.75 pg/ml
241	• TNFα: 40.34 pg/ml
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243	pAkt/Akt ratio after drugs incubation
244	Dath drugs inhibit Alst absorbarylation command to control (Figure 2) after 2 hours
245	Both drugs minor Akt phosphorylation compared to control (Figure 5) after 5 hours
246	incubation. This effect was completely lost after 6 hours of treatment.
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249	Discussion
250	In this work, we have demonstrated that ZA and LA inhibits, in vitro, the proliferation of
251	human prostate cancer cells: this is particularly true for ZA, whose cytotoxicity appears
252	already after 48- hours incubation and at lower concentrations than those necessary to
253	observe a LA-induced cytotoxic effect. Since patients often receive both drugs, we searched
254	for eventual interactions between the two drugs in vitro. Therefore, we incubated DU-145
255	cells with both drugs simultaneously (at variable concentrations of ZA). No synergistic effect
256	was induced by the simultaneous incubation with ZA and LA on cells.
257	Besides cytotoxicity, we wondered if ZA and LA could have other effects on prostate cancer
258	cells in vitro. Therefore, because of its relevance on metastasis phenomenon, we decided to

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investigate migration by means of the transwell assay and the evaluation of Aktphosphorylation level.

Both ZA and LA provoke a significant decrease in Akt phosphorylation in 3 hours, but after 6-hours it returns to the control levels; this observation suggests that their inhibitory effect occurs early, the timing being in agreement with that observed in inhibition of cell migration (Figure 2). Furthermore, no synergistic effect is provoked by ZA and LA co-incubation (Table II); even, LA effect seems to be significantly reduced by the presence of ZA. On the other hand, pre-incubation with LA (but not with ZA) significantly potentiate inhibitory migration properties of both LA and ZA in a synergistic manner. Nevertheless, the simultaneous presence of the drugs has to induce some effect on cell migratory properties, since when cells were pre-incubated with the single drugs their migration in culture medium was not inhibited at all, while 18-hours pre-incubation with ZA and LA significantly decreased their migration, even in RPMI. This may suggest that incubation with each drug singularly can cause not relevant/reversible changes, which cells are able to repair, while specific pre-incubation and/or specific drugs sequences (in particularly pre-incubation with LA followed by incubation with ZA) could lead to hard reversible changes affecting cell migration. We do not know, at present, how to explain this phenomenon, but a durable effect of the simultaneous presence of ZA and LA is probably worth further investigations.

About the ability to migrate more towards MSC-CM, it has been shown that migration and formation of metastases by prostate cancer cells in vivo is largely influenced by several factors produced by bone cells [15-19]. We observed a massive migratory increase under MSC-CM stimulus. It is interesting to note that both ZA and LA inhibit DU-145 3Dmigration at the same magnitude also when cells undergo MSC-CM stimuli (Figure 2). According to the literature, we found that the MSC-CM contains several factors involved in cancer cell survival and migration, such as TNF α , MCP1/CCL2 [15, 16, 17] and GM-CSF

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[20]. Aggressive cancer cell lines such as DU-145 and PC3 express a higher amount of 284 285 CCL2-specific receptor CCR2 compared with the less aggressive cancer cells such as LNCaP 286 or non-neoplastic PrEC and RWPE-1 cells [21]; a positive correlation has also been 287 established between CCR2 expression and prostate cancer progression [17]. Furthermore, Rivas and coll. [20] shown that prostate cancer cells express functional high-affinity GM-288 289 CSF receptors and therefore this hematopoietic growth factor may have an effect on prostate carcinoma cells. The increased expression of GM-CSF receptors in prostatic hypertrophy and 290 291 neoplastic prostate ephitelium suggests a relationship between prostatic epithelial cell growth 292 and GM-CSF [20]. At least, as shown by Gao and coll. [22], TNF α factor in endothelial cells 293 may increase the activation and ligation of $\alpha\nu\beta3$ integrins [22] to facilitate cell migration, and 294 regarding prostate cancer activation have a central role in prostate cancer metastatization 295 [23].

As previously reported in literature, most of these factors are affected by action of ZA and LA [24-29]. In particular, a direct correlation between GM-CSF and MCP-1 decreased levels and cell invasiveness has been demonstrated [24, 27]. On the basis of these observations, we could speculate that, as demonstrated in other systems, it can be confirmed in our own that the action of ZA and LA in decreasing migration could be both directed to the tumor cells and indirectly on potential metastatic niche.

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304 Conclusion

Our results suggest that: (i) drugs in use in vivo for prostate cancer treatment have a direct effect on prostate cancer cells proliferation, and this could contribute to justify the results obtained in vivo; (ii) effect of drugs overlapping is difficult to predict a priori; we are unable, with data at our disposal, to identify a regimen that clearly enhances the effect of the two

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309	drugs. However, we have shown that specific co-incubation prolongs inhibitory effect on
310	prostate cancer cells migration; (iii) drugs affect migratory ability of prostate cancer cells,
311	and this could contribute to justify their limiting effect of in vivo metastases (mostly LA);
312	(iv) in the same way ZA and LA diminish the chemoattractive effect of the bone marrow
313	mesenchymal stem cells and, hence, the potential role that they can play in the phenomenon
314	of metastasis; (v) both ZA and LA are also able to decrease cell migration, and this probably
315	by acting on the PI3K/Akt pathway [30].
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318	Acknowledgments
319	This study was supported by University of Turin (ex 60%). Authors thank Prof.ssa Biasi for
320	sharing her experience in cytochines quantification.
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22	155	Figure 1 Effect of zoledronic acid and leunrolide acetate on DU 145 growth
23	455	Figure 1. Effect of zoleutonic actu and leuptonide acetate on DO-145 growth.
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25	456	(A) Proliferation assay after 24, 48 and 72 hours culture in presence of increasing
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20	457	concentration of zoledronic acid $(2.5-50 \text{ \mu M})$ (B) Proliferation assay after 24 48 and 72
2/	107	concentration of zeroatome acta (2.0 co part). (2) Frontenation accup atter 21, 70 and 72
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29	458	nours culture in presence of increasing concentration of leuprolide acetate (0.5-100 μ M). (C)
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31	459	Effect of LA 100 µM and ZA (0-40 µM), alone or in combination, on DU-145 cell after 48
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33	460	hours incubation
34	400	nours incubation.
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36	461	*=P<0.05 vs control.
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40	463	Figure 2. Three-dimensional migration
41	105	rgure 2. Three anneholonal inigration.
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43	464	Migration rate after incubation or pre-incubation with drugs and/or in conditioned media
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44 45	465	(CM) compared to control. Migration is expressed in arbitrary units.
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46	100	*-D<0.05 via control: ***- D<0.001 via control: $^{\bullet}$ -D<0.001 via ZA, $^{\#}$ -D<0.001 via LA - ZA
47	400	-P<0.03 vs control, $-P<0.001$ vs control, $-P<0.001$ vs ZA, $-P<0.001$ vs LA+ZA.
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51	468	Figure 3 Western blotting
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57	469	Drugs effect on pAkt/Akt in DU-145 cell line after 3 and 6 hours of incubation with LA 100
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55	470	μ M or ZA 20 μ M. Vinculin as internal control.
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5	470	Table I. Experimental conditions for migration assay. I.A = loweralide contate 100 uM:
6	472	Table 1. Experimental conditions for migration assay. LA- reupronde acetate 100 µM,
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8	473	$ZA=$ zoledronic acid 20 μ M; MSC-CM= Mesenchymal stem cells conditioned medium.
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13	476	Table II. Drug interactions. CDI < 1 indicates a synergistic effect; CDI = 1 indicates
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15	477	additive effect: CDI > 1 indicates an antagonistic effect
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13x10mm (600 x 600 DPI)







12x6mm (600 x 600 DPI)

Pre-incubation 18 hours	Migration test 6 hours
	RPMI
none	ZA
re-incubation 18 hours none LA ZA ZA+LA none	LA
	RPMI
LA	LA
	ZA
	RPMI
ZA	ZA
	LA
ZA + LA	RPMI
	MSC-CM
none	MSC-CM + ZA
	MSC-CM + LA

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3	AB	Α	В	CDI
4 5	Incubation LA + ZA	Incubation ZA	Incubation LA	2,02
6 7 8	Pre-incubation ZA + incubation LA	Pre- incubation ZA	Incubation LA	1,02
9 10	Pre-incubation LA + incubation ZA	Pre-incubation LA	Incubation ZA	0,58
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