



LIPOXIN A4 ANALOG SELECTIVELY ALTERS THE TUMOR-ASSOCIATED MACROPHAGE PROFILE LEADING TO CONTROL OF TUMOR PROGRESSION

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3 **LIPOXIN A₄ ANALOG SELECTIVELY ALTERS THE TUMOR-ASSOCIATED**
4 **MACROPHAGE PROFILE LEADING TO CONTROL OF TUMOR**
5 **PROGRESSION**
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32 *Our study demonstrates, for the first time, that lipoxin, an important anti-inflammatory*
33 *mediator, selectively down-modulates the tumor progression stimulated by tumor-*
34 *associated macrophages, inducing the shift of macrophage profile and leading to tumor*
35 *cell apoptosis. These unexpected results reveal important differences in M2*
36 *macrophage subsets particularly those associated with tumors. This work suggests that*
37 *this lipid mediator may become a new tool for reducing the development and*
38 *progression of cancer.*
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3 Abbreviations used in this article:

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5 ALX/FPR2; formyl-peptide receptor type 2;

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7 ATL, aspirin-triggered lipoxin – 15-epi-LXA₄;

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9 ATL-1, stable analog of ATL

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11 BOC-1,N-t-butoxycarbonyl-methionine-leucine-phenylalanine;

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13 DPI, Diphenyleneiodonium;

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15 EC, endothelial cells;

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17 FITC, fluorescein isothiocyanate;

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19 GPCR, G-protein-coupled receptor;

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21 HMEC-1, human dermal microvascular endothelial cell line;

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23 iNOS, inducible nitric oxide synthase;

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25 LX, lipoxin;

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27 LXA₄, one of the lipoxins derived from arachidonic acid

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29 Mφ, macrophages;

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31 MRC1, mannose receptor, M2 macrophage marker

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33 MV3, melanoma cell lineage;

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35 NO, nitric oxide;

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37 PE, phycoerythrin;

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39 Per-CP, peridinin chlorophyll

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41 PMA, phorbolmyristate acetate;

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43 ROS, reactive oxygen species;

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45 SPMs, specialized pro-resolving lipid mediators;

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47 TAM, tumor-associated macrophages;

Abstract

In tumor microenvironments, pro-inflammatory macrophages (M1) acquire anti-inflammatory and pro-tumor characteristics. These tumor-associated macrophages (TAMs), often referred to as myeloid suppressors, exhibit an M2-like profile, with low cytotoxic properties and a deficient modulation of NO and ROS production. Lipoxins (LX) and 15-epi-lipoxins are lipid mediators inducing anti-inflammatory and pro-resolution activities in mononuclear cells, but their effects on TAMs remain to be elucidated. This study tested the hypothesis that ATL-1, a synthetic analogue of 15-epi-lipoxin A₄, could modulate TAM activity profile. We show that human macrophages (MΦ) differentiated into TAMs after incubation with conditioned medium from MV3, a human melanoma lineage cell. In contrast with the effects observed in the other M2 subset and M1 profile macrophages, ATL-1 selectively decreased M2 surface markers in these TAM, suggesting unique behaviour of the M2d subset. The effect was dependent on VEGF signaling and importantly, reproduced by the natural lipoxins, LXA and 15-epi-LXA₄. In parallel, ATL-1 stimulated TAM to produce NO by increasing the iNOS/arginase ratio and activated NADPH oxidase, triggering ROS production. These alterations in TAM profile induced by ATL-1 led to loss of the anti-apoptotic effects of TAMs on melanoma cells and increased their cytotoxic properties. Finally, ATL-1 was found to inhibit tumor progression *in vivo*, which was accompanied by alterations in TAM profile and diminished angiogenesis. Together, the results suggest unexpectedly that ATL-1 down-modulates tumor progression stimulated by TAM probably by inducing a change in the TAMs from an M2- to an M1-like profile thereby triggering tumor cell apoptosis.

1. Introduction

The relationship between the inflammatory response and cancer has been extensively investigated. In some types of cancer, the microenvironment is similar to that observed in inflammatory processes and tissue repair, with the presence of inflammatory cells and mediators, including chemokines, cytokines, and growth factors, as well as changes in the processes of tissue remodeling and angiogenesis¹. An early event in tumor progression is the recruitment of monocytes to the tumor site, where they differentiate into macrophages (M Φ)^{2,3}. These cells, as important effectors of the inflammatory response and its resolution, can provide an immediate defense against neoplastic elements but may promote tumor development through action at different stages⁴⁻⁵. This paradoxical role of M Φ in cancer could be explained by their functional plasticity, which may result in the polarized expression of either pro- or anti-tumoral functions. During the acute inflammatory response, M Φ with an M1 profile acquire a potent cytotoxic profile, with increasing production of nitric oxide (NO) and reactive oxygen species (ROS), which mediate the elimination of pathogens or neoplastic cells⁴. However, evidence has shown that tumor cells can suppress these immune functions, primarily through down-regulation of M Φ activity, allowing growth and spread of the cancer throughout the body, resulting in death of the host⁶⁻⁷. Tumor cells create a microenvironment that signals the pro-inflammatory and anti-tumoral M1 M Φ to acquire an M2-like profile. These M Φ , designated tumor-associated macrophages (TAMs), possess anti-inflammatory and pro-tumor characteristics, and, in contrast to M1 M Φ , exhibit low cytotoxic properties with defective production of NO and ROS, which allow tumor progression⁸⁻⁹. A number of studies have reported a positive correlation between high TAM density and poor prognosis in several human tumors¹⁰⁻¹².

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3 The dual role of MΦ in both endogenous tumor defense and tumor progression suggests
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5 that a better understanding of the molecular and cellular mechanisms involved in
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7 macrophage–tumor interactions is critical to developing new tools to control cancer
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9 progression.
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12 The balance between pro- and anti-inflammatory mediators may modulate the
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14 outcome of the inflammatory response and consequent tumor development. Lipoxins
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16 (LX) are specialized pro-resolving lipid mediators (SPMs), produced by lipoxygenase
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18 interactions under a variety of conditions, including infection and inflammation. They
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20 can be produced by different biosynthetic routes in particular biological settings¹³⁻¹⁴
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22 and can exert selective effects on monocytes and Mφs, polarizing these cells towards
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24 an anti-inflammatory profile, *in vitro* and *in vivo*¹⁵⁻¹⁹ with potential for regulation of
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26 tumor growth. Although some studies reveal down-regulation of lipoxins in certain
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28 tumors²⁰⁻²¹, the anti-inflammatory actions of lipoxins indicate that they play unique
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30 roles as mediators in their resolution of inflammation and could be important targets for
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32 the therapeutic control of cancer, although at present little is known about their effects
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34 on TAM behavior in the tumor microenvironment.
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39 In this study, we investigated the role of lipoxins in the activity of TAMs,
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41 using ATL-1(15-epi-16-(para-fluoro)phenoxy-LXA₄), a more stable analog of 15-
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43 epi- lipoxinA₄, an aspirin-inducible lipoxin. We show that this LXA₄ analog down-
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45 modulates the pro-tumor activity of TAM, inducing the shift from the M2 to the M1
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47 profile, promoting tumor cell apoptosis and controlling tumor growth *in vivo*.
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2. Methods

2.1. Reagents

Dulbecco's modified Eagle's medium (DMEM), RPMI 1640, and penicillin/streptomycin/amphotericin B solution were purchased from Life Technologies-Invitrogen (Carlsbad, CA). SDS and type IV horseradish peroxidase (HRP) was purchased from Sigma (St Louis, MO). Ficoll-Hypaque and Percoll were purchased from Amersham Biosciences (Piscataway, NJ). Antibodies were purchased from ABCAM (Cambridge, MA) and Biolegend (San Diego, CA). All other reagents and chemicals were purchased from Sigma (St Louis, MO). ATL-1, the stable 15-epi-LXA₄ analog, was a generous gift from Dr. John F. Parkinson, Bayer Healthcare Pharmaceuticals.

2.2. Melanoma cell culture

MV3 human melanoma cells were obtained from the Temple University Center for Neurovirology and Cancer Biology (PA, USA). B16F10 murine melanoma cells were obtained from American Type Culture Collection (ATCC). MV3 cells were maintained in DMEM and B16F10 were maintained in RPMI, both enriched with 10 % fetal bovine serum), 3.7 g/L sodium bicarbonate, 5.2 g/L HEPES, 0.5 U/mL penicillin, and 0.5 mg/mL streptomycin at 37°C in a humidified atmosphere of 5 % CO₂. Cells were grown to 80-90 % confluence in 75-cm² culture flasks and were then cultured for 3 days to generate conditioned medium.

2.3. Murine melanoma model

C57BL/6 mice at 6–7 wk of age were provided by the Department of Pharmacology and Psychobiology and all experimental procedures were performed according to guidelines

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3 of the Committee on Ethical Use of Laboratory Animals of the State University of Rio
4 de Janeiro, Rio de Janeiro, Brazil (CEUA/077/2012/, UERJ, Rio de Janeiro, Brazil).
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6 Mice were caged with free access to food and fresh water in a temperature-controlled
7 room (22–24°C) on a 12-hr light/dark cycle. For the induction of tumors, mice (n = 6-8)
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9 were administered B16 melanoma cells (2×10^5 cells/mouse suspended in 30 μ l of
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11 saline) by the subcutaneous (s.c.) route. ATL-1 (1 μ g/mouse) or ethanol (Vehicle) were
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13 injected intravenously 14 days after melanoma cell injection. For experimental analysis,
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15 mice were sacrificed on day 14 or 21 after melanoma induction and tumor mass was
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17 removed for histological analysis, morphometry, immunohistochemistry, and cellular
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19 profile. Lung and liver were analyzed to detect metastasis.
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27 ***2.4. Macrophage isolation and TAM differentiation***

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29 Isolated PBMCs were obtained from EDTA (0.5 %)-treated venous blood of healthy
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31 volunteers by Ficoll-Hypaque density gradient, as described previously¹⁸. Cells were
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33 plated for 1 hr to isolate lymphocytes and monocytes were cultured for 7 days for M Φ
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35 maturation. To obtain TAMs, the M Φ were cultured for an additional 3 days with RPMI
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37 1640 medium plus conditioned medium from MV3 cultures (MV3-CM; 1:1), as
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39 described above.
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44 ***2.5. Measurement of TAM survival***

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46 A modified MTT (3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide) assay
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48 was used to quantify the effect of lipoxins on TAM survival. This method measures
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50 mitochondrial function as described previously¹⁸. Briefly, M Φ and TAM (2×10^5
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52 cells/well) were incubated with or without ATL-1 (10-100 nM), LXA₄ (10nm) and 15-
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3 epi LXA₄ (10nM) for 48 hr. Survival was assessed and expressed as % of live cells
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5 above the negative control.
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10 **2.6. FACS analysis**

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13 Fluorescence-activated cell sorting (FACS) analysis was conducted with an ACCURI
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15 C6 flow cytometer with CFLOW software (Becton Dickinson, Heidelberg, Germany).
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17 MΦ and TAMs (1x10⁶ cells/well) cultured in 24-well plates were treated with ATL-1
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19 (1-100 nM), LXA₄ (10nM), and 15-epi-LXA₄ (10nM) for 48 hr. Following incubation,
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21 the cells were stained with conjugated monoclonal antibodies MRC1-phycoerythrin
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23 (PE), CD68-fluorescein (FITC) or CD86-PerCP-Cy5.5(Biolegend) with which they
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25 were incubated for 60 min at RT in the presence of blocking solution (2%bovine serum
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27 albumin,, 5%fetal bovine serum, and 0.1%sodium azide). An isotype control antibody
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29 was used as the negative control.
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36 **2.7. RNA isolation and RT-PCR**

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38 MΦ and TAMs (3x10⁶ cells/well) were incubated with or without ATL-1 (10 nM) for 6
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40 hr. Total RNA was isolated from TAMs using RNeasy Mini kits (Quiagen®, Hilden,
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42 Germany) . After DNase treatment (RQ1 RNase-Free DNase), the mRNA was reverse
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44 transcribed using high-capacity cDNA reverse transcription kits as described
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46 previously²². Primers were designed based on the sequence of human IL-10 (GeneBank
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48 accession n° NM_000265 - IL-10: sense (5'-AATAAGGTTTCTCAAGGGGCT-3');
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50 anti-sense (5'- AGAACCAAGACCCAGACATCAA-3'), GAPDH sense (5'-
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52 CAGTCAGCCGCATCTTCTTT-3'); anti-sense (5'-AATTCCGTTGACTCCGACCTT-
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54 3')). PCR was performed with the following program: 1 cycle of 95°C for 5 min and 32
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3 cycles of denaturation at 95°C for 45 sec, annealing at 58°C for 30 sec, and elongation
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5 at 72°C for 30 sec. Primers were used to validate the cDNA in each reaction). PCR
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7 products were separated by 2% agarose gel electrophoresis and visualized by UV
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9 exposure on a transilluminator. Relative band densities were determined by
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11 densitometric analysis using National Institutes of Health Image Software, and the
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13 ratios calculated.
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16 17 18 **2.8. NO production** 19

20 Nitrite release by TAM was determined by the accumulation of a stable nitrite end-
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22 product in cell-free culture supernatants using a modified Griess reaction method, as
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24 performed by Silva *et al.*(22). Briefly, TAMs (2×10^6 cell/mL) were incubated with
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26 ATL-1 (1-100 nM) for 8 hr, and the samples incubated with the Griess reagent (1 %
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28 sulfanilamide in 5 % phosphoric acid and 0.1 % naphthylethylenediamine). The nitrite
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30 concentration was then determined as previously described²².
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36 37 **2.9. ROS production assays** 38

39 MΦ and TAM were suspended in Hank's balanced salt solution (HBSS) and placed in a
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41 white, 96-well plate (2×10^5 cells/well, final volume 200 μ L). Following pretreatment
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43 with or without DPI (10 μ M) or BOC-1 (10 μ M) for 15 min, cells were loaded with
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45 Luminol (50 μ M) and hydroxyphenyl-fluorescein (HPF, 10 μ M), and stimulated with
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47 ATL-1 (10 nM) or PMA (10 ng/mL), a known NADPHox inducer, as a positive control.
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49 Cells that remained unstimulated were considered the control group. ROS production
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51 was assessed as described before²³. The area under the curve is expressed as the mean \pm
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53 standard deviation (S.D.).
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2.10. MV3 apoptosis

MV3 apoptosis was assessed via cell cycle analysis by flow cytometry. MV3 cells were distributed into 24-well plates at a density of 5×10^5 cells/well in 10 % FBS medium. MΦ and TAM (1×10^6 cells/mL) were cultured in inserts (0.4 μm) of transwell plates and the TAMs incubated for 15 min with or without DPI (10 μM), aminoguanidine (AG, 50 μM), or PEG-catalase (30 U/mL), followed by treatment with ATL-1 (10 nM) for 15 min. The analog and inhibitors were then removed, and TAM were co-cultured with MV3 cells for 72 hr. MV3 cells treated with cycloheximide (5 μM) was used as the positive control. Thereafter, the MV3 cells were detached with 5 mM EDTA in HBSS. DNA content was accessed by propidium iodide (PI) staining (50 μg/ml PI; 0.1 mg/ml RNase A; 0.05% Triton X-100) and analyzed by flow cytometry with Accuri C6® (Becton Dickinson), as previously described²³.

2.11. Preparation of cell extracts

MΦ and TAM (1×10^6 /mL) were incubated with different concentrations of ATL-1 (1-100 nM) and then frozen in an ice bath to stop the reaction. To obtain whole cell extracts, cells were resuspended in lysis buffer as previously described¹⁸. The total protein contents of the cell extracts were determined by the Bradford method²⁶.

2.12. Immunoblotting

The membranes were blocked with 5 % BSA in Tris-buffered saline (TBS) containing 0.1% Tween-20, and then probed with anti-iNOS (1:500), anti-ALX/FPR2, anti-arginase-1, anti-tubulin or anti-actin (1:1000) antibodies for 2 hr and developed as described previously¹⁸. Bands were visualized via an enhanced chemiluminescence

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3 system (ECL, Amersham, Arlington Heights, IL). In all instances, band density values
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5 were corrected by subtracting the background values.
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8 9 **2.13. Immunohistochemistry**

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11 Twenty-one days after administration of B16F10 cells, mice were killed, the tumor
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13 mass removed and immersed in the fixative solution for 24 hr. It was then embedded in
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15 optimum cutting temperature (O.C.T., Tissue-tek®) and sliced (10 µm) perpendicular to
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17 the tumor base (apical-basal axis) and stained with H&E. Sections were hydrated and
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19 the slides washed in TBS with 0.05% Tween 20 (T-PBS, Sigma-Aldrich), blocked with
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21 block solution (BSA 5% in T-TBS), and immunostained with antibodies to endothelial
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23 cells (CD105), MΦ (F480) and TAM (MRC1), used at a 1/100 dilution in PBS/Tween
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25 buffer overnight at 4°C. Immunofluorescent images were developed with antibodies
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27 conjugated with fluorescein or phycoerythrin and counterstained with DAPI for nuclear
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29 staining. An irrelevant isotype IgG was used as negative control.
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40 **2.14. Statistical analysis**

41 The data are representative of different experiments and expressed as the mean ± S.D.
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43 The means from different treatments were compared by Student's *t*-test and ANOVA,
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45 followed by the *Bonferroni t*-test for unpaired values. Statistical significance was set at
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47 $P < 0.05$.
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3. Results

3.1. Characterization of a TAM phenotype induced by a simulated tumor microenvironment *in vitro*

Characteristic TAM markers are the macrophage mannose receptor (MRC1), and CD163, which are usually expressed on M2 cell membranes²⁷. For induction of a TAM phenotype by a simulated tumor microenvironment *in vitro*, MΦ derived by maturation of peripheral blood monocytes were incubated with conditioned medium obtained from cultures of melanoma cells. MΦ were converted to a TAM profile expressing higher levels of MRC1 and CD163 than do naïve MΦ (Fig. 1A and B).

One key feature of TAM is the high level of expression of IL-10, an immunosuppressive cytokine produced by tumor cells and TAMs that allows these cells to assume M2-related functions². Indeed, under our experimental conditions, the TAM also presented high levels of IL-10 mRNA (Fig. 1C).

LX and their analogs exert their anti-inflammatory activities via activation of ALX, a GPCR expressed in a variety of cellular types¹⁶. We demonstrated that TAM express ALX and the treatment of the cells with ATL-1 (10 nM) did not change this expression (Fig. 1D). Furthermore, in the presence of ATL-1, the cells maintained viability even after 48 hr of incubation, as demonstrated by MTT assays (Fig. 1E).

3.2. ATL-1 selectively decreases M2 marker expression by TAM

To investigate the effects of ATL-1 on the TAM phenotype, we treated the cells with the analog for 72 hr. As demonstrated in Figure 2A, ATL-1 treatment decreased MRC1 expression to levels similar to those observed in MΦ. Furthermore, ATL-1 inhibited IL-10 release by TAMs (Fig. 2B). The natural lipoxin, LXA₄ and its aspirin-induced analogue, 15-epi-LXA₄ induced similar effects to ATL-1 (Fig 2C). Intriguing

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3 contrast, ATL-1 had no effect on the M2 MRC1 phenotype induced by other M2
4 stimuli, such as IL-4, TGF- β and VEGF, and even induced an M2 profile in M1 cells
5 (Fig 2D). This suggests that TAM represent a subset of M2 macrophages which are
6 selectively down-modulated by ATL-1 to an M1-like profile *in vitro*. A further response
7 of TAM was the significant reversal of MRC1 expression by an M1 stimulus.
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16 **3.3. ATL-1 effect on TAM is vascular endothelial growth factor (VEGF)-** 17 **dependent.** 18 19

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21 In the tumor microenvironment, VEGF is the most important angiogenic factor
22 and is involved in several steps of tumor progression²⁸. Recently, our group
23 demonstrated that ATL-1 impairs VEGF actions both *in vitro* and *in vivo*²⁹⁻³⁰. On this
24 evidence, we investigated the involvement of VEGF in the ATL-1-induced TAM
25 effects. First, we found that conditioned medium from MV3 (CM-MV3) induced an M2
26 profile independent of VEGF, since treatment with a neutralizing anti-VEGF antibody
27 did not affect MRC1 expression on TAM (Fig. 2E). On the other hand, when M2
28 polarization triggered by CM-MV3 was performed in the absence of VEGF, the ability
29 of ATL-1 to alter the TAM profile was impaired (Fig.2E). These results suggest that the
30 effects of ATL-1 on the TAM profile are dependent of VEGF actions.
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45 **3.4. ATL-1 induces NO production by increasing the iNOS/Arginase-1 ratio in** 46 **TAM** 47 48

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50 In the tumor microenvironment, TAMs, in contrast to M1 cells, exhibit deficient
51 production of NO due to inhibition of iNOS activity by elevated levels of arginase-1^{8,31}.
52 Figure 3B shows that TAM induced in the present study express high levels of arginase-
53 1, in accord with these observations. Treatment of TAM with ATL-1 decreased
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3 arginase-1 and enhanced iNOS expression, thus increasing the iNOS/arginase-1 ratio
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5 (Fig.3A,B and C). Corroborating these results, ATL-1 (10 nM) also induced significant
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7 NO production by TAM, an effect that was inhibited by BOC-1, a selective inhibitor of
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9 ALX/FPR2 (Fig. 3D).
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11 12 13 14 **3.4. ATL-1 induces ROS production by TAMs**

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16 In addition to their limited ability to produce NO⁸, TAM also produce low
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18 amounts of ROS (Fig.4A and B). After treatment with ATL-1 (10nM), TAM generated
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20 significantly larger amounts of ROS, an effect impaired by both BOC-1 and DPI, an
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22 NADPH oxidase inhibitor (Fig. 4A and B). Furthermore, ATL-1 had no effect on ROS
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24 production by MΦ. Additionally, we show that ATL-1, as a possible consequence of an
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26 increased production of ROS and NO³², induces peroxynitrite formation, an effect
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28 inhibited by BOC-1 (Fig 4C).
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34 **3.6. ATL-1 induces MV3 apoptosis in a ROS- and NO-dependent manner**

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36 Since the activation of NO and ROS could lead to the recovery of the cytotoxic
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38 properties of TAM², we analyzed the melanoma cell cycle after 72 h of co-incubation
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40 with ATL-1-treated TAM. Compared to MΦ-co-cultured tumor cells, MV3 melanoma
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42 cells co-cultured with TAM were protected from apoptosis (Fig.5A). By contrast, the
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44 treatment of TAM with ATL-1 (10nM) reversed this anti-apoptotic activity of TAM on
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46 MV3 to levels comparable to those seen with MΦ and after an M1 stimulus (Fig. 5A).
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48 The incubation of tumor cells with supernatant from the ATL-1-treated TAM culture
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50 did not interfere with MV3 apoptosis (Fig. 5A), suggesting that the pro-apoptotic effect
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52 depends on the presence of the TAM.
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3 To investigate the contribution of the NADPH oxidase-derived ROS to the pro-
4 apoptotic effects of ATL-1-treated TAM, cells were pretreated with DPI (10 μ M) before
5 incubation with the analog (10nM). DPI impaired ATL-1-induced MV3 apoptosis,
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7 indicating that the analog induces tumor cell death via activation of the NADPH oxidase
8 system in TAM (Fig. 5B).
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14 A simultaneous increase in the production of superoxide/H₂O₂ and NO leads to
15 potent cytotoxic properties displayed by ATL-1-stimulated TAM. To test this
16 hypothesis, TAM were treated with aminoguanidine (AG, 50 μ M) and PEG-catalase
17 (30U/mL), inhibitors of iNOS and H₂O₂, respectively, followed by the treatment with
18 ATL-1 (10 nM). Treatment with the inhibitors completely blocked the pro-apoptotic
19 effects of ATL-1, but did not potentiate the inhibitory effect of either one alone,
20 suggesting that NO and H₂O₂ act through a common pathway (Fig. 5C).
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31 32 **3.7. ATL-1 inhibits tumor progression *in vivo*.**

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34 In view of these *in vitro* results, we investigated the effects of ATL-1 on tumor
35 progression in a murine model *in vivo*. We observed that treatment with a unique dose
36 of ATL-1 at the 14th day after implantation inhibits tumor growth, as observed in
37 Figures 6A and B. This potent inhibition of ATL-1 was accompanied by the impaired
38 angiogenic process and a decrease in the TAM marker, as shown in Figures 6C and D.
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4. Discussion

The relationship between the inflammatory response and cancer has been widely investigated. In spite of the strong evidence for their anti-inflammatory and pro-resolution actions, reports on the effects of lipoxins on tumor progression are limited. In this study, we present novel and promising data on the effects of ATL-1 on the behavior of TAM in the tumor microenvironment. We show that the treatment with ATL-1 selectively alters TAM to an M1-like profile, increasing their tumoricidal activity and leading to control of tumor progression *in vivo*. These effects emphasize the important role of lipoxins in modulation of the macrophage profile in the tumor microenvironment with consequent induction of pro-resolutive properties. As described before it has been demonstrated by a number of groups, including ours, that lipoxins and other lipid mediators can induce anti-inflammatory properties in mononuclear cells¹⁷⁻¹⁹. Indeed, Li et al, screening the effects of lipoxins on naïve and inflammatory monocytes and macrophages derived from the U937 cell line, demonstrated that LXA₄ induces the M2 anti-inflammatory phenotypes³³. Vasconcelos *et al* showed that lipoxins allowed *in vivo* the control of macrophage phenotypic profile, displaying a characteristic M2 phenotype³⁴.

Depending on the inducing stimulus, M2 macrophages can be classified into four subsets³⁵⁻³⁶. M2a and M2b, activated by IL4/IL13 or immune complexes/LPS, respectively, exert immune regulatory functions, whereas M2c macrophages activated by IL-10 and TGF- β have a predominant role in suppressing immune responses and in promoting tissue remodeling. TAM have been reported as a novel M2 subset, called M2d, that are apt to promote tumor growth, tissue remodeling, angiogenesis, and suppression of adaptive immunity^{1,3,4,36}. Although there is a paucity of evidence for the lipoxin effects on macrophages present in the conditions of the tumor

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3 microenvironment, we have demonstrated for the first time that the stable derivative of
4 aspirin-induced LXA₄, ATL-1, alters the TAM profile to an M1-like phenotype. In
5 contrast, we observed that ATL-1 induces an M2 profile in Mφs activated by LPS and
6 IFNγ that exhibit an M1 profile, and maintains M2 polarization in Mφ treated with
7 IL4, a classical M2 stimulus, indicating a specificity of the LX effect on TAM
8 polarization.
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12 We demonstrated that the effect of lipoxins on TAM polarization appears to be
13 dependent on VEGF released in the tumor microenvironment. Our data strongly
14 suggest that prior signaling triggered by VEGF during the differentiation of Mφs in
15 TAMs would be an essential step for the specific effect of lipoxins on TAM. We
16 propose that VEGF, in synergy with other factors released into the tumor
17 microenvironment, such as TGF-β, VEGF, IL4 and IL10, may trigger distinct
18 intracellular signaling, which induces the unique M2-like profile, which differentiates
19 TAM from other M2-phenotypes. Corroborating this hypothesis, Linde *et al* showed
20 that, beyond its chemotactic effect, VEGF induces the production of IL- 4 by a
21 macrophage subset and of IL10 by tumor cells, so potentiating M2 polarization³⁰.
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39 The recruitment, maturation, and differentiation of infiltrating leukocytes in
40 tumors are dependent upon the local conditions in the tumor microenvironment,
41 which play a central role in the activation of specific pathways expressed by TAM³⁷.
42 In this study, we observed that ATL-1, besides decreasing the expression of the
43 mannose receptor MRC1, a characteristic surface antigen marker of TAM²⁷, also
44 drastically inhibited IL-10 gene expression. IL-10, a known anti-inflammatory and
45 immunosuppressive cytokine, reported to be present in the microenvironment of
46 different tumors, inhibits the expression of pro-inflammatory mediators, including
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3 IL-12, inducible cyclooxygenase, and nitric oxide synthase, by phagocytic cells^{7,38-}
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5⁴⁰. This inhibitory effects of the ATL analog contrast clearly with other reports
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7 showing that lipoxins can induce IL-10 synthesis by macrophages in a non-tumor
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9 environment⁴¹. Moreover, this inhibition of IL-10 expression by ATL-1 suggests that
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11 the alteration in macrophage profile could facilitate the restoration of their potent
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13 cytotoxic activities, thereby increasing NO and ROS production.
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17 Indeed, we demonstrated that ATL-1, in changing the TAM
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19 phenotype, increases the production of NO and induces ROS generation by
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21 activating the NADPHox system. These pro-oxidative effects triggered in TAM
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23 contrast, again, with those obtained in other M ϕ phenotypes, where inhibitory effects
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25 of lipoxins and other ALX/FPR2 agonists on reactive species production were
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27 demonstrated⁴²⁻⁴⁴. Corroborating the hypothesis of a selective effect of ATL-1 on
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29 TAM, we demonstrated that it had no effect on ROS production by peripheral blood-
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31 derived M ϕ s. The NO production is consequence of a balance between iNOS and
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33 arginase-1 activity⁹. The arginase pathway limits arginine availability for iNOS action
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35 and NO synthesis. Interestingly, NO production by TAM triggered by ATL-1 seems to
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37 be more closely related to a decline in arginase-1 expression than to an
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39 augmentation in iNOS expression, resulting in an increase in the iNOS/arginase-1
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41 ratio.
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46 Several studies have demonstrated that malignant cells are able to grow
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48 indefinitely because of unregulated proliferative capacity and acquired resistance to
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50 apoptosis, allowing these cells to become invasive and metastatic⁴⁵⁻⁴⁶. Alterations in
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52 the TAM profile and the recovery of the cytotoxic properties of M ϕ s have been
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54 proposed as part of an efficient strategy to control tumor progression⁹. Our data
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56 indicate that ATL-1, by triggering ROS and NO production, enables TAM to induce
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3 apoptosis of MV3 melanoma cells. Inhibition of either NADPH oxidase or iNOS
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5 systems alone impaired the pro-apoptotic effect of ATL-1-stimulated TAMs
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7 on the tumor cells, suggesting that both reactive species, ROS and NO, share a
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9 common pathway. An increased production of superoxide/H₂O₂ and NO leads to the
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11 production of the highly cytotoxic mediator peroxynitrite³² consistent with our
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13 observation that ATL-1 induces peroxynitrite production by TAM. Conceivably this
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15 metabolite could be the main factor responsible for the potent cytotoxic properties
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17 displayed by ATL-1- stimulated TAM. The data would appear to indicate the
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19 involvement of different M2 subsets under the same umbrella of the M2 markers used
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21 to define the M2 phenotype in this study.
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25 Tumor cells need to avoid apoptosis, which is also induced by limited supplies
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27 of oxygen, nutrients, and growth factors⁴⁷. ATL-1, in addition to its ability to alter
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29 TAMs to a tumor pro-apoptotic phenotype, also inhibited human endothelial cell
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31 tubulogenesis stimulated by TAM *in vitro* (Supplemental Figure S1). Furthermore, the
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33 treatment with ATL-1 decreased, in parallel with M2 cell population, angiogenic
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35 markers in the tumor microenvironment, *in vivo*. The data corroborate early studies
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37 that suggested ATL-1 as a potent inhibitor of angiogenesis by down-regulating
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39 VEGFR activation, and inhibiting migration and proliferation of endothelial cells³¹⁻³².
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41 Taken together, our data support the idea that the shift in macrophage profile can also
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43 contribute to the anti-angiogenic effects of ATL-1 *in vivo* .
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48 The production of ATL by aspirin *in vivo* results from the inhibition of COX-2
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50 through its acetylation¹⁴. Non-steroidal anti-inflammatory drugs (NSAIDs), including
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52 aspirin, were reported to impair tumor progression through the suppression of
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54 proliferation, the induction of apoptosis, and the inhibition of the invasiveness of
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56 neoplastic cells⁴⁸⁻⁴⁹. Interestingly, Claria *et al* showed that ATL was produced by
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3 aspirin-activated neutrophils co-cultured with tumor cells⁵⁰, and proposed that lipoxins
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5 may have a function in the control of cancer by affecting leukocyte activity.
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7 Corroborating this suggestion, we demonstrate for the first time that lipoxins, by
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9 inducing a shift from the M2-like (TAM) to the M1-like profile, promote inhibitory
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11 effects on tumor cell proliferation and survival, leading to tumor cell apoptosis and
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13 control of tumor progression *in vivo*. This work highlights an important role for lipoxins
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15 in tumor progression and suggests that these lipid mediators may become new targets
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17 for reducing the development and progression of cancer.
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Or Peer Review

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For Peer Review

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Figures Legends

Figure 1 - Characterization of a TAM phenotype induced by simulated tumor

microenvironment *in vitro*. A-B) MΦ were cultured with or without conditioned medium from melanoma cells for 72hr to TAM differentiation. Expression of the M2 markers, MRC1(**A**) and CD163 (**B**) was assessed by flow cytometry using the anti-MRC1-PE and CD163-APC antibodies. CD68-FITC and CD80-PerCP-Cy5 were used as the MΦ marker. The figure is representative of three experiments performed separately. **C)** MΦ and TAM (3×10^6 cells) were treated with ATL-1 (10 nM) for 6 hours. Cells were lysed in TRIZOL® and IL-10 gene expression was investigated by RT-PCR analysis. GAPDH was used as load control. Figure is representative of three experiments performed separately. **D)** MΦ and TAM (1×10^6 cells) were treated with ATL-1 (10 nM) for 48 hr. ALX/FPR2 expression was investigated by Western blot. The figure is representative of three experiments performed separately. **E)** MΦ and TAM (2×10^5 cells) were treated with ATL-1 (10-100 nM) for 48 hr. Cell viability was assessed by the MTT assay. Results are expressed as the mean \pm S.D of three independent experiments.

Figure 2 - ATL-1 selectively decreases M2 markers expression by TAMs. **A)** MΦ and

TAMs (1×10^6 cells) were incubated with or without ATL-1 (1-100nM) for 72 hours. MRC1 expression was assessed by flow cytometry using the antibody anti-MRC1-PE. The results are expressed as the mean \pm S.D of three independent experiments. *, $P < 0.05$ when compared to MΦ; **, $P < 0.05$ when compared to TAMs. **B)** MΦ and TAM (1×10^6 cells) were treated with ATL-1 (10 nM) for 24 hours. IL-10 release was assessed by ELISA using an anti-IL10 antibody. The results are expressed as the mean \pm S.D of

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3 three independent experiments. *, P < 0.05 when compared to TAM alone. C) MΦ and
4 TAM (1x10⁶ cells) were incubated with or without ATL-1(10nM), 15-epi-LXA₄(10nM),
5 LXA₄ (10nM) and in the latter case, with M1 stimulus (LPS1μg/mL + IFNγ 40ng/mL)
6 for 72 hr. MRC1 expression was assessed by flow cytometry using the antibody anti-
7 MRC1-PE. The results are expressed as the mean ± S.D of three independent
8 experiments. *, P < 0.05 when compared to control MΦ; **, P < 0.05 when compared to
9 TAM. D) MΦ (1x10⁶ cells) were incubated with medium alone, MV3 conditioned
10 medium (TAM), M1 stimulus (LPS1μg /mL + IFNγ 40ng/mL), M2 stimuli: IL-4
11 (100ng/mL), TGF-β (10ng/mL) or VEGF (10ng/mL) for 3 days. After washing cells
12 were then incubated in the presence or absence of ATL-1 (10 nM) for 72 hr. MRC1
13 expression was assessed by flow cytometry using the anti-MRC1-PE antibody. The
14 results are expressed as the mean ± S.E. of three independent experiments. *, P < 0.05
15 when compared to control MΦ; **, P < 0.05 when compared to TAM. #, P < 0.05 when
16 compared to M1. E) MΦ were incubated with MV3 conditioned-medium (CM-MV3) in
17 the presence or absence of monoclonal anti-VEGF (0.1μg/mL) for 72 hours. The results
18 of treating MΦ and TAMs (1x10⁶ cells) with or without ATL-1 (10 nM) for 48 hr are
19 shown for comparison. MRC1 expression was assessed by flow cytometry using the
20 anti-MRC1-PE antibody. The results are expressed as the mean ± S.D of three
21 independent experiments. *, P < 0.05 when compared to untreated MΦ; **, P < 0.05
22 when compared to untreated TAM.

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50 **Figure 3. ATL-1 induces NO production by increasing the iNOS/Arginase-1 ratio**
51 **in TAMs. A-C) TAMs (1x10⁶ cells) were incubated with ATL-1 (10-100 nM) for 2 hr.**
52 **The cells were lysed in a specific buffer and analysed by SDS-PAGE. A) iNOS/actin**
53 **ratio. B) Arginase-1/actin ratio. C) iNOS/Arginase-1 ratio. Protein expression was**
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3 evaluated by immunoblotting with specific antibodies. The results are expressed as the
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5 mean \pm S.D. The data shown are representative of three independent experiments. *, $P <$
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7 0.05 compared with M Φ .**, $P <$ 0.05 compared with TAMs alone. **D)** TAM (2×10^5 /mL)
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9 were cultured in 96-well plates, pretreated with BOC-1 (10 μ M), and treated with ATL-
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11 1 (1-100 nM) for 8 hr. NO production was assessed by the Griess Reaction and
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13 analyzed on an Envision® plate reader, as described in the Methods. *, $P <$ 0.05 when
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15 compared with TAMs alone. **, $P <$ 0.05 when compared with TAMs treated with
16
17 ATL-1.
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22 **Figure 4. ATL-1 induces ROS production by TAMs.** **A)** M Φ and TAMs (2×10^5 /mL)
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24 were pre-treated with BOC-1 (10 μ M) or DPI (10 μ M) and then incubated with or
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26 without ATL-1 (10 nM) for 1 hr in the presence of luminol as a chemiluminogenic
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28 probe (50 μ M). BOC-1 alone had no effect. PMA was used as a positive control. The
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30 figure is representative of 4 experiments. **B)** ROS production. Mean \pm S.D. of the area
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32 under the curve. **C)** M Φ and TAMs (2×10^5 /mL) were pre-treated with BOC-1 (10 μ M)
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34 and then incubated with or without ATL-1 (10 nM) for 30 min in the presence of HPF
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36 as a chemiluminogenic probe (10 μ M). The results are expressed as the mean \pm S.D of
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38 four independent experiments. *, $P <$ 0.05 compared with M Φ . **, $P <$ 0.05 compared
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40 to cells treated with ATL-1.
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47 **Figure 5. ATL-1 induces MV3 apoptosis in a ROS and NO dependent manner.** **A)**
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49 M Φ and TAM (1×10^6 cells) were cultured in *transwells* and the TAM treated with or
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51 without ATL-1 (10 nM) or an M1 stimulus (LPS 1 μ g /mL + IFN γ 40ng/mL) for 15 min.
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53 The analog was then removed, and the cells were co-cultured with MV3 for 72 hours.
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55 MV3 apoptosis was assessed by cell cycle analysis via flow cytometry. Cycloheximide
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3 (CHX)-treated MV3 cells were used as the positive control. SN TAM+ATL-1;
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5 supernatant from the ATL-1-treated TAM. Dashed line, MV3 cells alone. **B-C**) MΦ
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7 and TAM (1×10^6 cells) cultured in *transwells* were pretreated for 15 min as shown in
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9 the Figure with or without **B**)DPI (10 μ M); or **C**) aminoguanidine (AG, 50 μ M) and
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11 polyethyleneglycol-catalase (PEG-Cat) (30 U/mL), followed by treatment with ATL-1
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13 (10 nM) for 15 min. The analog and inhibitors were then removed, and the cells co-
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15 cultured with MV3 cells 72 hr. MV3 apoptosis was assessed by cell cycle analysis via
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17 flow cytometry. The results are expressed as the mean \pm S.D of three independent
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19 experiments. *, $P < 0.05$ when compared to MΦ; **, $P < 0.05$ when compared to TAMs;
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21 #, $P < 0.05$ when compared to TAM plus ATL-1.
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27 **Figure 6. ATL-1 inhibits tumor progression *in vivo*.** Groups of 6-8 mice were
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29 inoculated with B16F10 tumor cells. The tumor mass was removed from the animals on
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31 the 14th and 21st day after B16F10 injection. ATL-1 (1 μ g/mouse) or ethanol (vehicle
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33 alone) was injected on day 14, as described in Methods section 2.3. **A**) Tumor volume
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35 and **B**) weight were analyzed. The excised tumors were immunostained with antibodies
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37 for: **C**) blood vessels (CD105-PE) and **D**) whole macrophage (F4/80-FITC) or M2
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39 macrophages (MRC1-PE). DAPI was used for nuclear staining shown in the upper
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41 right-hand square. An isotype IgG was used as negative control. Pictures are
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43 representative of each group. *, $P < 0.05$. **, $P < 0.01$.
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Figure 1 – Simões R.L. et al - LIPOXIN A₄ ANALOG SELECTIVELY ALTERS THE TUMOR-ASSOCIATED MACROPHAGE PROFILE LEADING TO CONTROL OF TUMOR PROGRESSION

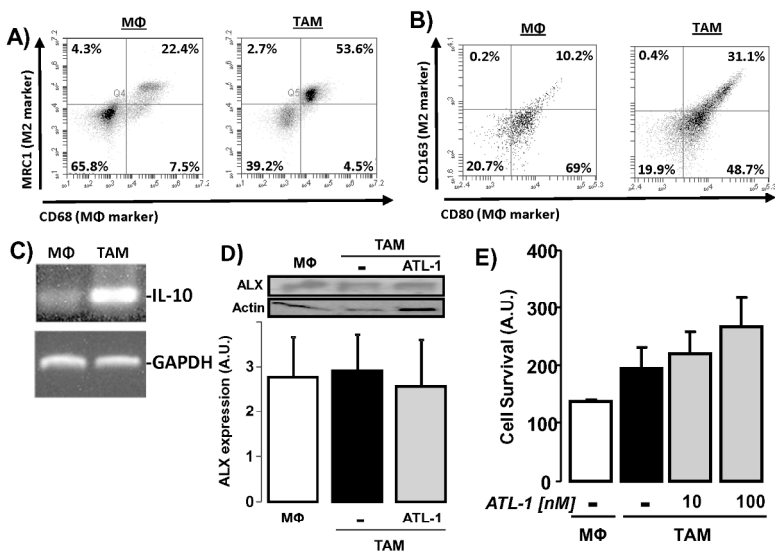


Figure 1
190x254mm (300 x 300 DPI)

Figure 2 – Simões R.L. et al - LIPOXIN A₄ ANALOG SELECTIVELY ALTERS THE TUMOR-ASSOCIATED MACROPHAGE PROFILE LEADING TO CONTROL OF TUMOR PROGRESSION

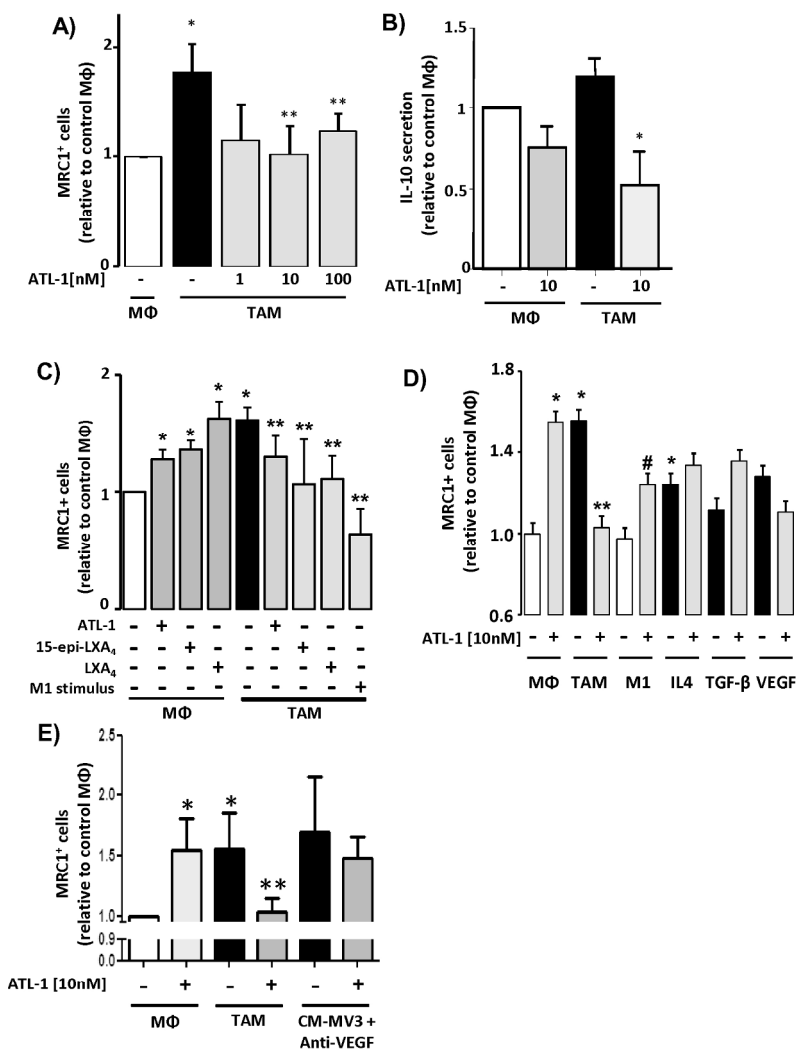


Figure 2

190x254mm (300 x 300 DPI)

Figure 3 – Simões R.L. et al - LIPOXIN A₄ ANALOG SELECTIVELY ALTERS THE TUMOR-ASSOCIATED MACROPHAGE PROFILE LEADING TO CONTROL OF TUMOR PROGRESSION

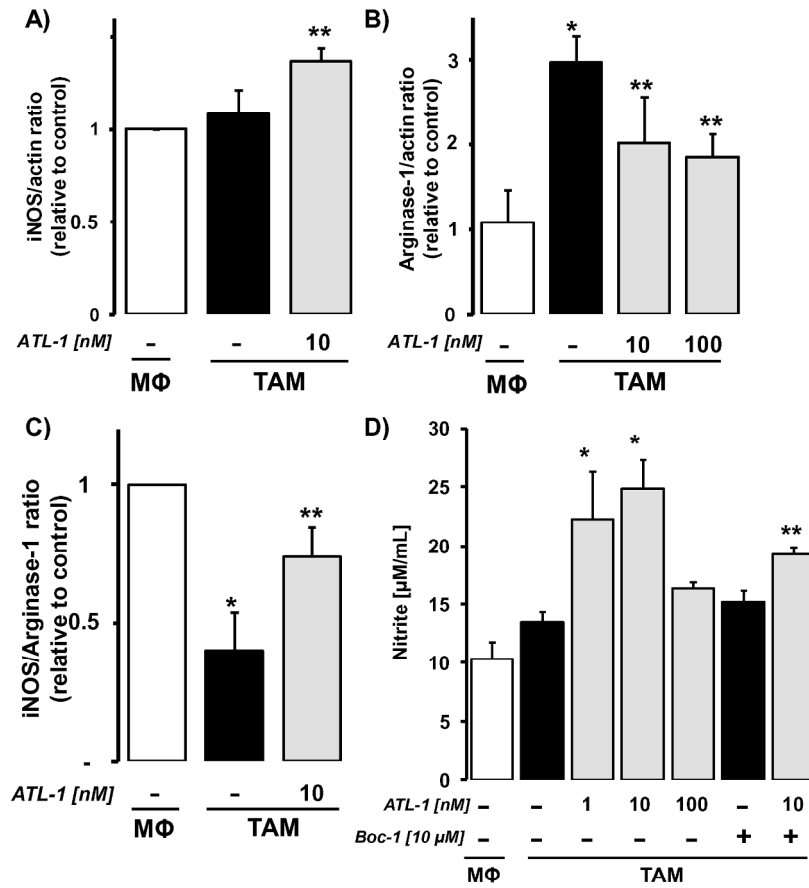


Figure 3
190x254mm (300 x 300 DPI)

Figure 4 – Simões R.L. et al - LIPOXIN A₄ ANALOG SELECTIVELY ALTERS THE TUMOR-ASSOCIATED MACROPHAGE PROFILE LEADING TO CONTROL OF TUMOR PROGRESSION

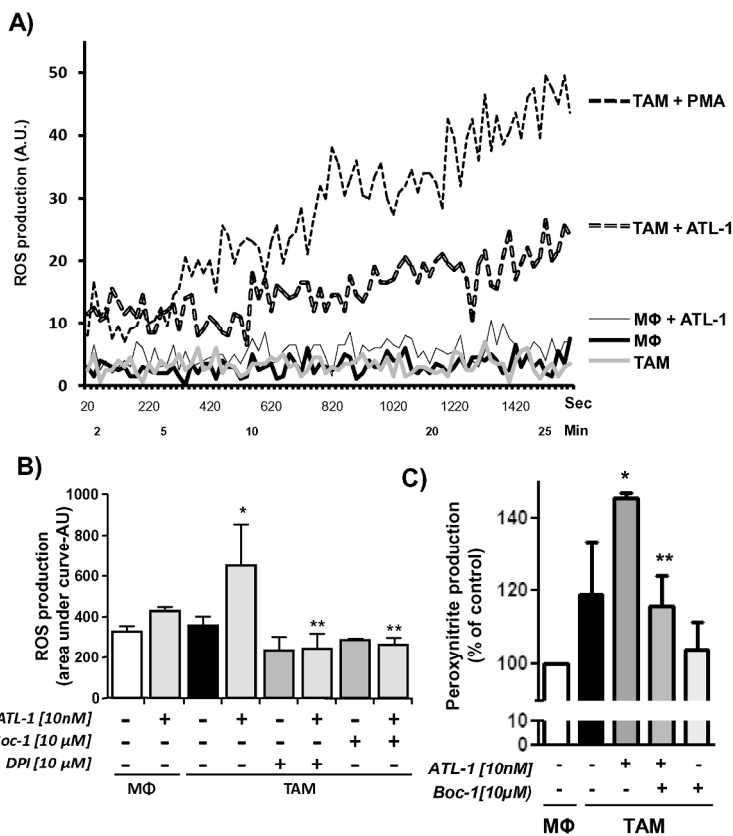


Figure 4
190x254mm (300 x 300 DPI)

Figure 5 – Simões R.L. et al - LIPOXIN A₄ ANALOG SELECTIVELY ALTERS THE TUMOR-ASSOCIATED MACROPHAGE PROFILE LEADING TO CONTROL OF TUMOR PROGRESSION

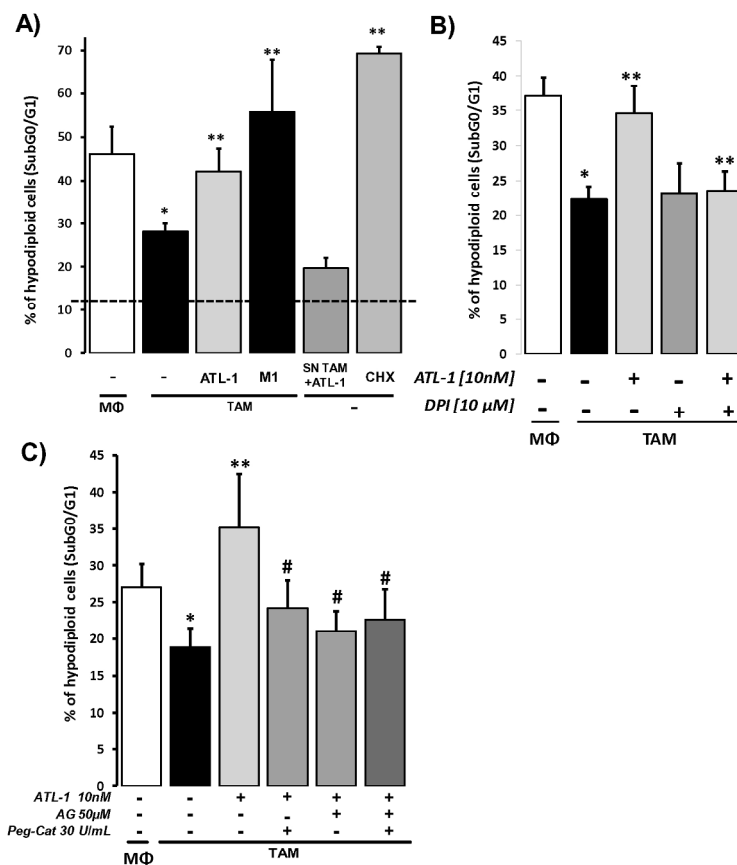


Figure 5
190x254mm (300 x 300 DPI)

Figure 6 – Simões R.L. et al - LIPOXIN A₄ ANALOG SELECTIVELY ALTERS THE TUMOR-ASSOCIATED MACROPHAGE PROFILE LEADING TO CONTROL OF TUMOR PROGRESSION

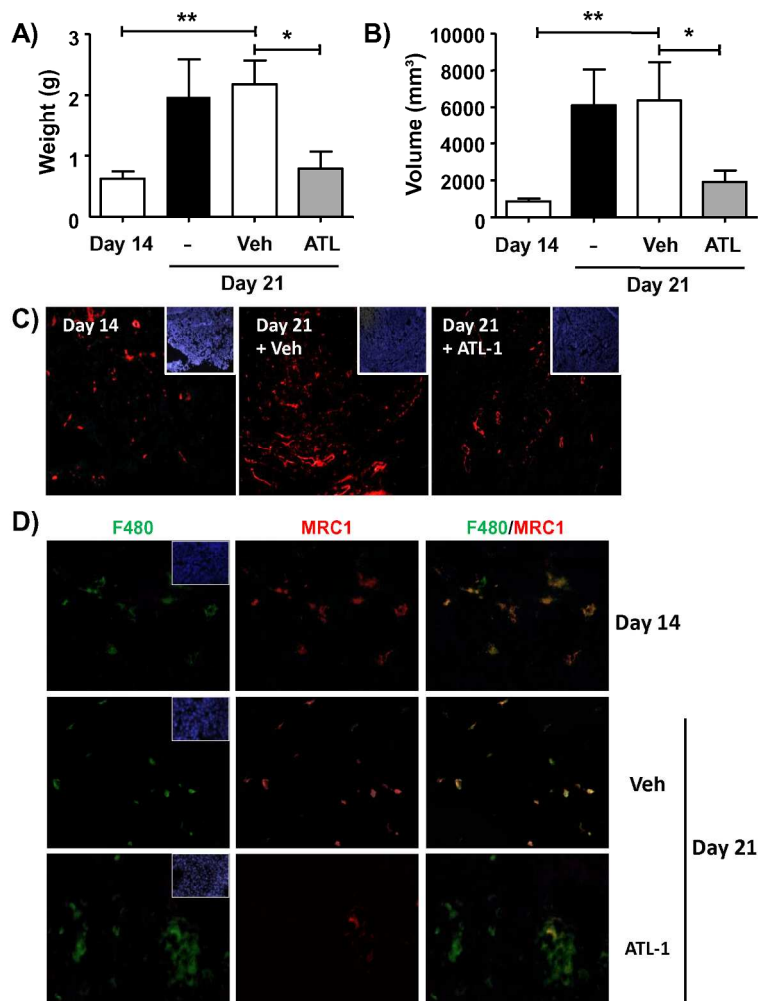


Figure 6
190x254mm (300 x 300 DPI)