

LYMPHOCYTE SUBSETS, DENDRITIC CELLS AND CYTOKINE PROFILES IN MICE WITH MELANOMA TREATED WITH *Uncaria tomentosa*

Ivan Lozada-Requena^{1,2,a}, Cesar Nuñez^{1,2,b}, Yubell Alvarez^{1,c}, Laura Kahn^{1,d}, Jose Aguilar^{1,e}

ABSTRACT

Objectives. To evaluate the immunomodulatory effect on lymphocyte subsets, dendritic cells (DC), Th1 / Th2 / Th17 and inflammatory cytokines on systemic level and/or in the tumor microenvironment of mice with or without melanoma. **Materials and methods:** Peripheral blood and/or primary tumors samples were obtained of mice with B16 melanoma treated or not with a hydroalcoholic extract of *Uncaria tomentosa* (UT) with 5.03% of pentacyclic oxindole alkaloids (UT-POA) obtained from the bark of the plant. All cell assays and cytokine measurements were performed by flow cytometry. **Results.** UT-POA systemically increased CD4/CD8a relation while cell activation was inversely proportional; increased the proportion of DCm; induced a pro-inflammatory Th1 profile and reduced Th17 response. TNF- α and IL-17A positively and negatively correlated with CD4/CD8a relation. **Conclusions.** The increase of Th1 (TNF- α) may result in the increase of CD4 or M1 macrophage activation. Although UT-POA shows increased DCm, is not dose-dependent. Th17(IL-17A) decreased can support the function of CD8a lymphocytes. UT-POA shows better systemic immunomodulatory effects than intratumoral.

Key words: *Uncaria tomentosa*; Lymphocyte activation; Dendritic cells; Melanoma (source: MeSH NLM).

INTRODUCTION

Malignant melanoma, a tumor derived from skin melanocytes, is the deadliest form of skin cancer⁽¹⁾. The increased incidence of human melanoma is related to hereditary or ethnic factors, environmental factors such as UV exposure, and nutritional and lifestyle factors⁽²⁾. Murine models such as melanoma B16 cells in BALB/c and C57BL/6 mice are accepted among scientists because of the understanding of the genetics and because they reproduce characteristics of human cancer⁽³⁾. Therefore, murine models are used to evaluate drugs, immunotherapies, and medicinal plants.

Medicinal plants have few side effects, low cytotoxicity, and the ability to act on target cells without affecting

normal cells, and they are also a low-cost therapeutic option. Research on medicinal plants has yielded encouraging results. Park W.-B. *et al.* evaluated the effect of *Viscum album* (mistletoe) lectins on mice with melanoma, and observed proapoptotic, antiangiogenic, antimetastatic, and survival-enhancing effects⁽⁴⁾. Currier N.L. *et al.* showed upregulation of natural killer (NK) cells in mice with leukemia that were treated with *Echinacea purpurea* (Echinacea)⁽⁵⁾. Epigallocate polyphenol (EG) derived from *Camelia sinensis* (green tea) has an antimetastatic activity. Likewise, epigallocatechin-3-gallate (EGCG) and dacarbazine reduce metastases and tumor growth in mice with melanoma⁽⁶⁾.

Uncaria tomentosa (UT; Cat's claw) is a vine from the Peruvian Amazon jungle that has anti-inflammatory,

¹ Immunology laboratory. Department of cellular and molecular sciences. Faculty of sciences and philosophy. Universidad Peruana Cayetano Heredia. Lima, Peru.

² EMINDES SAC (*Empresa de Investigación y Desarrollo en Cáncer*). Lima, Perú.

^a Master in Science, ^b physician; ^c biologist; ^d degree of bachelor in biology; ^e rheumatologist specializing in immunology
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antioxidant, immunomodulatory, and antiproliferative properties^(7,8). Using the same extract as in our present study, Dreifuss *et al.* showed antitumor and antioxidant effects in Walker-256 rats with carcinosarcoma⁽⁹⁾. The antiproliferative and antitumor activity of UT has also been shown in cell lines of breast cancer (MT-3) and Ewing's sarcoma (MHH-ES-a)⁽¹⁰⁾. Although there are not many studies on humans, one in particular stands out, which shows an improvement in the quality of life of patients with solid tumors⁽¹¹⁾ and another one that shows alleviation of neutropenia and repair of the DNA damage related to chemotherapy⁽¹²⁾. UT contains quinovic acid glycosides, steroids, polyphenols, polyhydric triterpenes, proanthocyanidins, catechins, saponins, tannins, sterols, flavonoids, and alkaloids. The most clinically relevant effects occur when these compounds act synergistically⁽¹³⁾. Sheng *et al.* showed that C-MED100 (a UT extract with less than 0.05% of oxindole alkaloids) increases the numbers of T lymphocytes (TL) in rats with doxorubicin-induced leukopenia⁽¹⁴⁾. It also promotes repair of splenocyte DNA in UV-irradiated rats⁽¹⁵⁾. Aguilar *et al.* and Aquino *et al.* showed that ethanolic and methanolic UT extracts reduce inflammation in plantar edema^(7,16). Sandoval *et al.* showed that inhibition of inflammation is associated with downregulation of TNF- α and is mediated by inhibition of NF- κ B activity⁽¹⁷⁾. Fazio *et al.* showed that UT reduces melanoma tumors, lung metastases, and downregulates TNF- α and IL-6⁽⁸⁾.

Regarding inflammation and cancer, it is still unknown whether it is the state of chronic inflammation that leads to the development of neoplastic cells or vice versa⁽¹⁸⁾. Coussens *et al.* demonstrated the presence of growth factors, TNF- α , and reactive oxygen species (ROS) in the tumor microenvironment (TME); ROS damage DNA and initiate neoplasms⁽¹⁹⁾. Tumor immunotherapy induces cell-mediated immunity (for example, dendritic cell [DC] vaccines or activation of CD4+ TLs involved in the direct protection against the development of tumors and in tumor regression)⁽²⁰⁾. Likewise, upregulation of IL-2 because of tumors causes a dose-dependent decrease in tumor mass⁽²¹⁾, and it has been observed that PBI-1393 is capable of reducing the size of tumors and can stimulate lymphocytes to produce antitumor interferon (IFN) and IL-2⁽²²⁾. Everything indicates that TLs play a crucial role in the defense of the host against the disease. CD4+ TLs differentiate into the following subsets: Th1, Th2, Th17, Th22, T_{reg}, Th9, and Th follicular helper cells⁽²³⁾. Nonetheless, the Th1/Th2 system has been an immunological paradox for a long time. Th1 (IFN- γ and IL-2) or proinflammatory cytokines promote a response to intracellular pathogens. Th1 lymphocytes recognize tumor antigens, whether directly (some melanomas express on-site MHC II) or via

DC cross-presentation mechanisms⁽²⁴⁾. It is possible that Th1 immune cells, through IFN- γ generate an antitumor response in which CD8+ TLs do not participate directly because they activate NKs, macrophages, and monocytes involved in the defense against neoplasms⁽²⁵⁾, while Th2 anti-inflammatory cytokines (IL4, IL-5, and IL-10) stimulate elimination of extracellular parasites and induce polarization of tumor-associated macrophages (TAMs) or M2d that tend to promote progression of cancer⁽²⁶⁾. On the other hand, TAMs or M2d are also reported to play an anticancer role because they can indirectly activate the phagocytic system to release ROS and nitric oxide synthase (NOS) that are cytotoxic to tumor cells⁽²⁷⁾.

There are *in vivo* and *in vitro* melanoma studies that show Th2 directionality⁽²⁸⁾. The other newly discovered subset is Th17, which is generated in the presence of TGF- β and IL-6 and proliferates under the influence of IL-23⁽²⁹⁾. During the development of tumors, Th17 cells were found in prostate cancer and in the TME⁽³⁰⁾. The produced cytokines, including IL17, have controversial functions. On the one hand, they impair the function of CD8+ TLs and promote carcinogenesis and neovascularization of tumors via STAT3; on the other hand, they have been reported to perform antitumor functions. Because they may cause inflammation and destruction of tissue, they are potentially useful for cancer treatment. IL-17 induces inflammatory cytokines and chemokines close to the TME, which can facilitate the recruitment of DCs specialized in presentation of tumor antigens as well as cytotoxic effector cells such as CD8+ TLs, NKs, and NKT cells that can attack the tumor⁽³¹⁾. DCs have several subsets, the most important of which are myeloid DCs (mDCs) and plasmacytoid dendritic cells (pDCs)⁽³²⁾. They are both important to an antitumor response⁽³³⁾.

All these data suggest that immunocompetent cell populations, the Th1/Th2/Th17 balance, and inflammatory cytokines have a strong influence on the development of cancer. All the evidence indicates a correlation, but the specific role of cytokines and the complexity of their interactions have not been identified yet. Therefore, in this study, we evaluated the modulatory effect of a UT extract on lymphocyte populations and DCs and the profile of inflammatory and Th1/Th2/Th17 cytokines systemically and in the TME in a murine melanoma model.

MATERIALS AND METHODS

EXPERIMENTAL DESIGN

Sixty-two mice were used, subdivided into 4 or 5 experimental groups according to the type of evaluation (Table 1). They received a UT hydroalcoholic extract

Table 1. Study groups according to the experimental design

Evaluation	Without melanoma		With melanoma		
	NC (n=2)	PC (n=9)	UG50 (n=6)	UG500 (n=6)	UG1000 (n=9)
Peripheral blood	NC (n=2)	PC (n=9)	UG50 (n=6)	UG500 (n=6)	UG1000 (n=9)
Tumoral microenvironment	---	PC (n=9)	UG100 (n=9)	UG500 (n=7)	UG1000 (n=5)

NC = negative control, PC= positive control, UG50-1000 = UT-POA at 50–1000 mg/ kg of body weight

with 5.03% pentacyclic oxindole alkaloids (UT-POA) or distilled water (negative control, CN) via a gastric tube 7 days prior to the inoculation of B16 cells, and during 22 days in the case of peripheral blood (PB) evaluation or until the tumor reached 4.5 mm for the TME evaluation.

Uncaria tomentosa (UT)

We used a hydroalcoholic extract, which was obtained from the bark of UT, in the form of a fine reddish brown powder. This extract was kindly provided by Peruvian Heritage® and prepared by decoction with ethanol and water in the 70:30 ratio for 1 hour at 20°C; subsequently, it was spray dried. The extract contained 5.03% of pentacyclic oxindole alkaloids, quantified by high-performance liquid chromatography (HPLC), as described by Dreifuss *et al.* (9). A UT-POA stock solution was prepared: 30 g was dissolved in 1 L of double-distilled water, boiled for 30 minutes, decanted, filtered twice through Whatman No. 3 paper and microfilter (pore size 0.22 µm; for sterilization).

ANIMALS

C57BL/6 mice were obtained, randomized as females and males, from the Cayetano Heredia Peruvian University Vivarium. The experiments were performed on groups with 6 mice on average (7 to 9 weeks old and weighing 20–25 g). They were given standard balanced food and water *ad libitum*. The UT-POA extracts and distilled water were administered daily via a gastric tube (Table 1).

B16 CELL CULTURE

The B16/BL6 (murine melanoma) cell line was cultured in a humidified atmosphere containing 5% of CO₂ at 37°C in the RPMI 1640 medium supplemented with 10% fetal bovine serum, penicillin (100 U/mL), streptomycin (100 µg/mL), and 0.01% sodium pyruvate. After the cells reached >70% confluence, they were trypsinized,

washed, counted, and resuspended in PBS pH 7.2 for further use.

INDUCTION OF A TUMOR

The standard dose for subcutaneous injection was 5×10^4 cells per 100 µL of a suspension. The cells were injected into the left flank of the abdominal region. In the case of PB evaluation, the animals were euthanized on day 22, whereas for the TME analysis, they were euthanized when the tumor reached the size of 4.5 mm.

ISOLATION AND CULTURE OF PERIPHERAL BLOOD MONONUCLEAR CELLS (PBMC) AND TUMOR MICROENVIRONMENT MONONUCLEAR CELLS (TMMC).

On day 22, PB samples were obtained via cardiac puncture (approximately 1 mL) and the blood of 2 to 3 animals was pooled by diluting the blood in equal parts with complete RPMI 1640 (Sigma, St. Louis, MO, USA). In the case of a solid tumor (4.5 mm), it was dissected, fragmented mechanically, digested with 1% type I collagenase (Gibco, USA) for 30 minutes at 37°C, and diluted with equal volume of complete RPMI 1640. Density gradient centrifugation was performed on both types of samples with tubes containing 3 or 1 mL of 1.083 g/mL Histopaque (Sigma), respectively. They were centrifuged at 1800 rpm at room temperature (RT) for 30 minutes, without braking. The cell ring was recovered, washed twice with 15 mL of complete RPMI 1640 at RT during the first wash and at 4°C during the second wash at 2500 rpm for 10 minutes. The cells were counted in a Neubauer chamber, 1:1 ratio with Trypan blue (Sigma). Once the PBMCs or TMMCs were isolated, 5×10^4 cells were plated for each 100 µL of a suspension in sterile tubes, and they were incubated at 37°C and 5% CO₂ in complete RPMI 1640 for 24 hours. The culture supernatants (CSs) were collected to quantify the cytokines, and the cells were washed and resuspended

in a cell wash solution (1% fetal bovine serum [Hyclone] in PBS [pH 7.4]).

FLOW CYTOMETRY

Specific fluorochrome-labeled monoclonal antibodies were used: anti-CD3-PerCP, anti-CD4-FITC, anti-CD8a-PE, anti-CD44-APC, anti-CD69-PerCP, anti-CD8a-APC, anti-CD11b-FITC, anti-I-Ad-PE, and anti-CD11c-APC, and their respective isotype controls (Becton Dickinson, San Jose, CA, USA). The titration, compensation, and acquisition of antibody data were performed on a FACSCanto™ II flow cytometer (BD Immunocytometry Systems, USA). The results were analyzed in the Flow Jo software v. X10.0.7r2.

In all cases, the respective combinations of four antibodies, according to the population or populations to be identified, were added to tubes that contained the cells. They were incubated for 30 minutes at 2–8°C and washed twice with 1 mL of the cell wash solution with centrifugation at 1800 rpm for 5 minutes. Finally, the cell pellets were resuspended in 500 µL of the cell fixative (1% paraformaldehyde in pH 7.4 PBS) and analyzed on the flow cytometer.

FLOW CYTOMETRY FOR ANALYSIS OF CYTOKINES VIA CBA (CYTOMETRIC BEAD ARRAY)

Cytokine assay kits for cytometry (BD CBA Mouse Th1/Th2/Th17 and Inflammation kits, USA) were used for quantitative analysis of Th1 cytokines (IFN-γ IL-2), Th2 (IL-4, IL-6, IL-10), Th17 (IL-17A), and proinflammatory cytokines (IL-6, IL-10, MCP-1 [monocyte chemoattractant protein], IFN-γ, TNF-α, and IL-12p70). The CSs were processed in strict accordance with the manufacturer's instructions. Data were acquired on a FACSCanto™ II flow cytometer (BD Immunocytometry Systems, USA) and analyzed in the BD™ Cytometric Bead Array Software, version 1.4.

STATISTICAL ANALYSIS

The results are presented as mean ± SEM (standard error of the mean). We performed the one-tailed *t* test and one-way analysis of variance (ANOVA) followed by Dunnett's *post hoc* mean comparison test and Pearson's one-tailed correlation test in the GraphPad Prism software, version 6.00 for Mac (GraphPad Software, La Jolla California USA). Differences with *p* < 0.05 were considered significant.

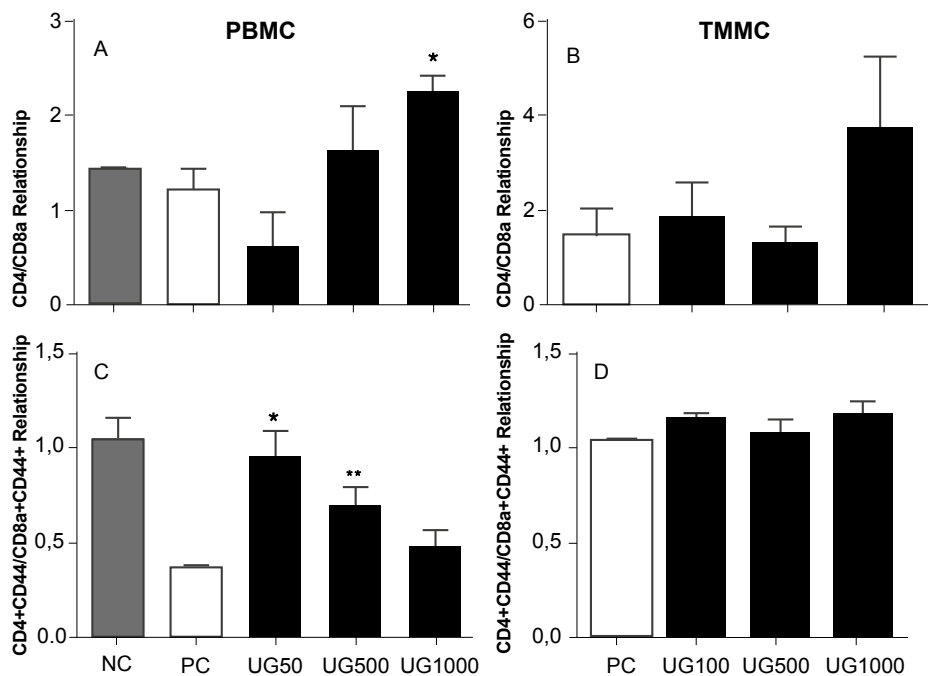


Figure 1. The ratio of systemic T lymphocytes (TLs) and the tumor microenvironment. The percentages of helper T cells (CD3e+CD4+) were determined by flow cytometry, as were percentages of cytotoxic TLs (CD3e+CD8a+) and the activation level (CD44+ or CD69+) for both populations. The following ratios were then determined and plotted: CD4+/CD8a+ and CD4+CD44+/CD8a+CD44+ or CD4+CD69+/CD8a+CD69+ for both peripheral blood mononuclear cells (PBMCs) and tumor microenvironment mononuclear cells (TMMCs), respectively. The data represents mean ± standard error of the mean (SEM). One-way ANOVA and *post hoc* Dunnett's test were performed. Differences with *p* < 0.05 (*, **) were considered significant relative to PC. NC = Negative Control, PBMCs of mice without melanoma/UT-POA; PC = Positive Control, PBMC/TMMC of untreated mice with melanoma; UG50-1000 = PBMCs of mice treated with 50, 500, or 1000 mg/kg UT-POA; UG1001000 = TMMCs of mice treated with 100, 500, or 1000 mg/kg UT-POA.

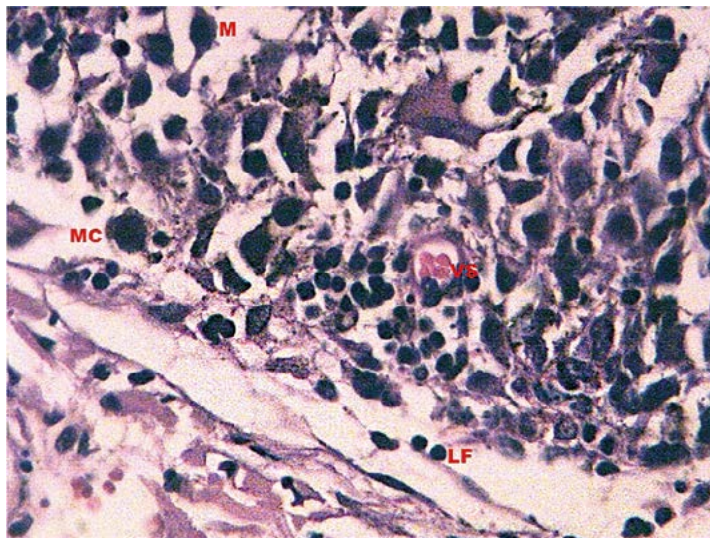
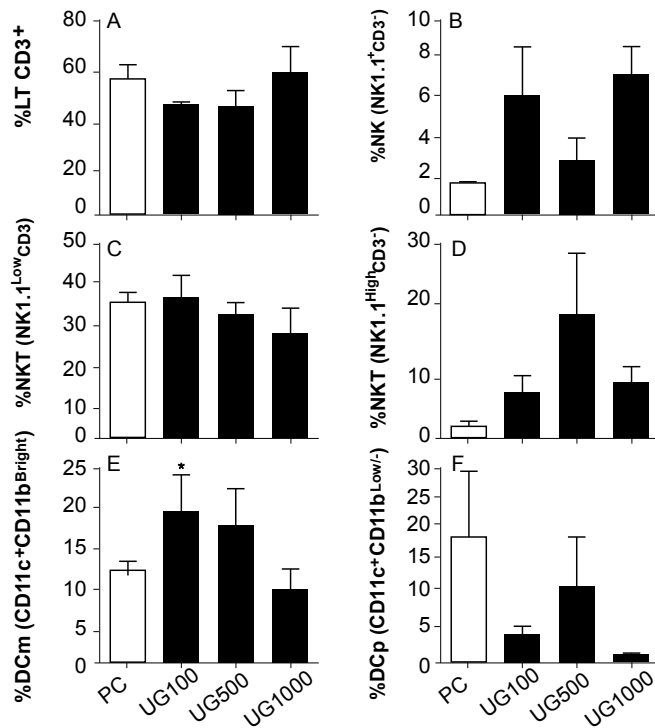


Figure 2. Immunocompetent populations of the tumor microenvironment. The following percentages were determined by flow cytometry A) Total TLs (CD3e+); B) NKs (NK1.1+CD3e-); C) NKT cells (NK1-1LowCD3e+); D) NKT cells (NK1-1HighCD3e+); E) DCs (CD11c+CD11bbright); F) DCs (CD11c+CD11bLow/-) based on the TMMCs, and G) Histological section (hematoxylin and eosin staining) representative of a solid melanoma tumor from a mouse from group UG1000, 40x magnification. The data represent mean ± standard error of the mean (SEM). One-way ANOVA and *post hoc* Dunnett's test were performed. Differences with *p* < 0.05 (*) were considered significant relative to PC. PC = positive control, TMMCs of untreated mice with melanoma; UG100-1000 = TMMCs of mice treated with 100, 500, or 1000 mg/kg UTPOA; M = melanocytes; MC = macrophages; LF = lymphocytes; and VS = blood vessel with red blood cells.

ETHICAL CONSIDERATIONS

All experiments on animals were conducted in accordance with the *Principles of Laboratory Animal Care* (Publication NIH 85-23, revised in 1985). The project was approved by the Institutional Ethics Committee of Cayetano Heredia Peruvian University, registration code 56290 (SIDISI).

RESULTS

CD4+/CD8a+ RATIO AND LYMPHOCYTE ACTIVATION IN PB AND TME

The percentages of helper T cell (CD3⁺CD4⁺) populations and cytotoxic cells (CD3⁺CD8a⁺) were determined, and the CD4/CD8a ratio was calculated. Additionally, the activation level was measured for both TL populations with two markers that may be used interchangeably, CD44 and CD69. The subpopulations in groups that were treated with UT-POA or untreated were analyzed for PBMCs (UT-POA dose: 50, 500, or 1000 mg/kg) and for the TMMCs (UT-POA dose: 100, 500, or 1000 mg/kg). The results indicated that the systemic CD4/CD8a ratio increased significantly in response to 1000 mg/kg UT-POA (Figure 1A), while the systemic cell activation (judging by the CD4⁺CD44⁺/CD8a⁺CD44⁺ ratio) was significant in response to the doses 50 and 500 mg/kg. Although at 1000 mg/kg, we saw stronger activation than in the positive control (PC), the effect was not significant (Figure 1C). In the TME, no significant changes were observed either in the CD4/CD8a or in the activation ratio of CD4⁺CD69⁺/CD8a⁺CD69⁺ (Figure 1B and 1D).

IMMUNOCOMPETENT POPULATIONS IN THE TME

We determined the percentages of total TLs (CD3⁺); NK lymphocytes (NK1.1⁺CD3⁺); NKT lymphocytes (NK1.1⁺CD3⁺) and myeloid dendritic cells (mDC, CD11c⁺CD11b^{bright}); and plasmacytoids (pDC, CD11c⁺CD11b^{low/-}) in the TME of mice either treated or not treated with UT-POA (100, 500, or 1000 mg/kg). The results revealed that none of the TL, NK, or NKT populations showed significant variations in response to the UT-POA treatment (Figures 2A-D and 3). Likewise, we found that UT-POA at 100 mg/kg significantly increased the

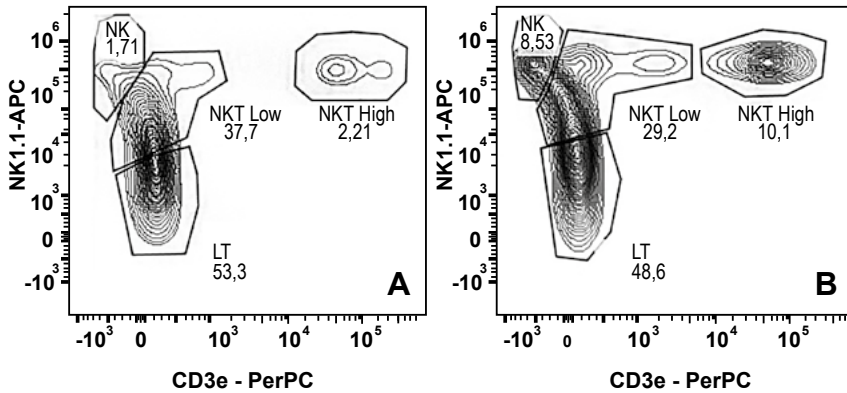


Figure 3. NKT lymphocytes of the tumor microenvironment. The following percentages were determined by flow cytometry: total LTs (CD3e+); NKs (NK1.1+CD3e-), and NKT cells (NK1-1LowCD3e+ and NK1-1HighCD3e+) on the basis of the TMMCs. The results of two separate experiments are presented, A) PC, TMMCs of untreated mice and B) UG100, TMMCs of mice treated with 100 mg/kg UT-POA.

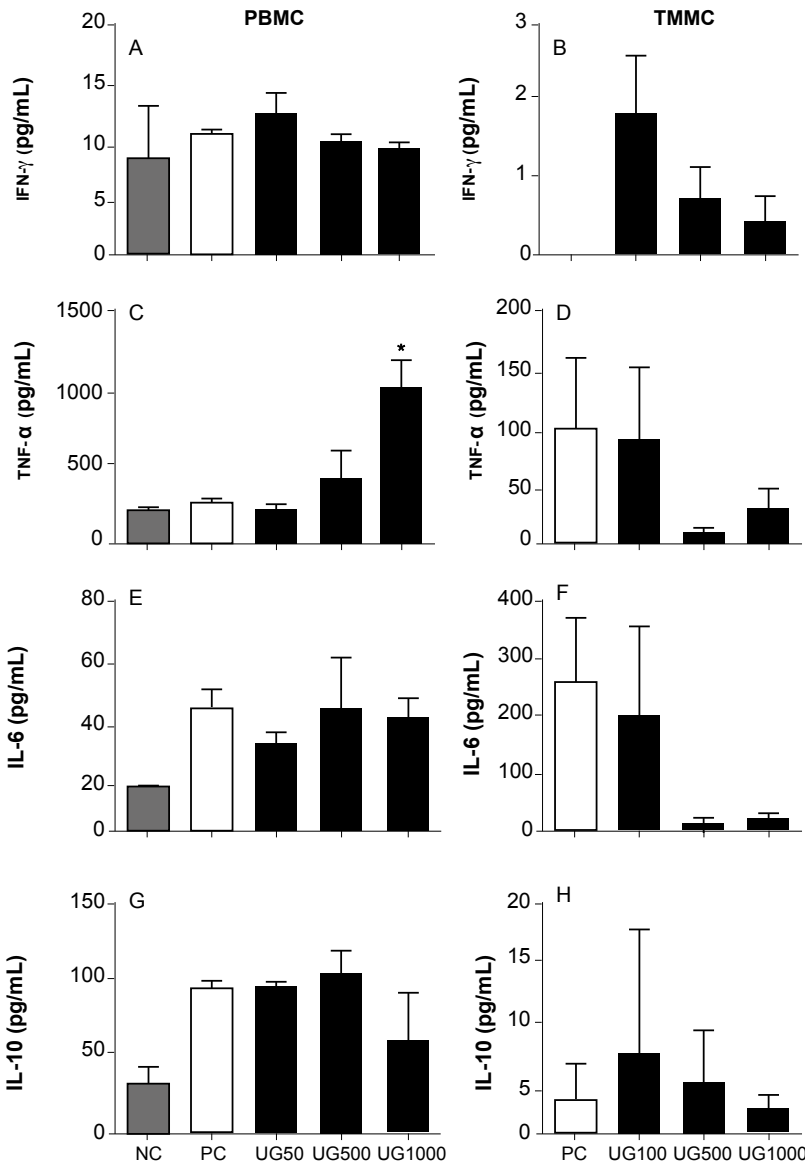


Figure 4. Production of IFN- α , TNF- α , and IL-10 by PBMCs and TMMCs. The concentrations of the cytokines IFN- α and IL10 in culture supernatants of PBMCs and TMMCs were quantified by cytometry (CBA). **NC**, negative control with PBMCs of untreated healthy mice; **PC**, positive control with PBMCs or TMMCs of untreated mice with B16 melanoma; **UG50-1000**, treatment groups (PBMCs of mice with B16 melanoma treated with 50, 500, or 1000 mg/kg UT-POA); **UG100-1000**, treatment groups (TMMCs of mice with B16 melanoma treated with 100, 500, or 1000 mg/kg UT-POA). The data represent mean \pm standard error of the mean (SEM). One-way ANOVA with *post hoc* Dunnett's test and the onetailed *t* test were performed. Differences with $p < 0.05$ (*) were considered significant relative to PC. PBMCs = peripheral blood mononuclear cells, TMMCs = tumor microenvironment mononuclear cells, UT-POA = hydroalcoholic *Uncaria tomentosa* extract with 5.03% pentacyclic oxindole alkaloids.

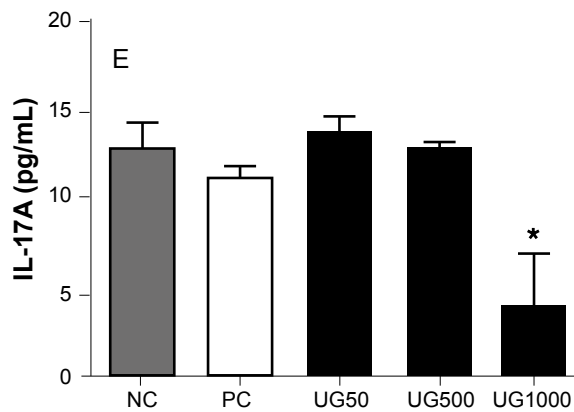


Figure 5. IL-17A production by PBMCs. The concentration of the cytokine IL-17A in the culture supernatants of PBMCs was quantified by cytometry (CBA). **NC**, negative control (PBMCs of untreated healthy mice); **PC**, positive control (PBMCs of untreated mice with B16 melanoma); **UG50-1000**, treatment groups (PBMCs of mice with B16 melanoma treated with 50, 500, or 1000 mg/kg of UT-POA, respectively). The data represent mean ± standard error of the mean (SEM). The one-tailed *t* test was performed. Differences with *p* < 0.05 were considered significant relative to PC. PBMCs = peripheral blood mononuclear cells, UT-POA = hydroalcoholic *Uncaria tomentosa* extract with 5.03% pentacyclic oxindole alkaloids.

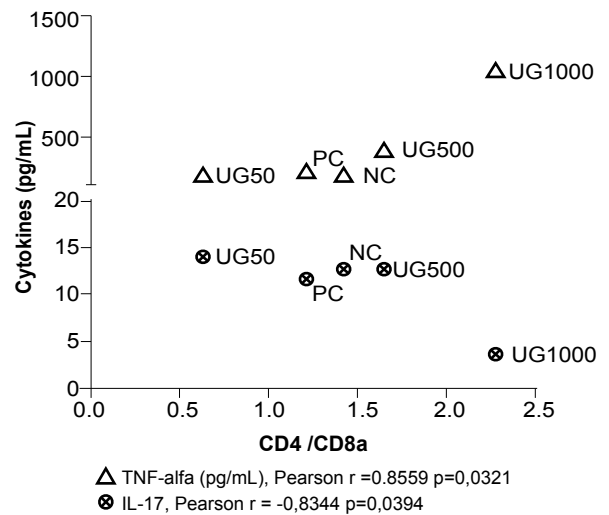


Figure 6. Correlation of the CD4/CD8a values with the concentrations of the cytokines TNF-α and IL-17A. The figure represents the intersection of the average values for each group: **NC**, negative control (PBMCs of untreated healthy mice); **PC**, positive control (PBMCs of untreated mice with B16 melanoma); **UG50-1000**, treatment groups (PBMCs of mice with B16 melanoma treated with 50, 500, or 1000 mg/kg extract). A one-tailed Pearson correlation analysis was performed. Negative or positive *r* values indicate negative or positive correlations; correlations with *p* < 0.05 were considered significant. PBMCs = peripheral blood mononuclear cells.

percentage of mDCs, while the pDCs showed no changes (Figure 2E and F). The presence of immunocompetent cells was detected histologically in the TME (Figure 2G).

SYSTEMIC AND TME CYTOKINES

The concentration of Th1 (IFN-γ, TNF-α) and Th2 (IL-6, IL-10) cytokines were determined via cytometry (CBA) in the PBMC and TMMC supernatant; Th17 (IL-17A) in PBMCs; Th1 (IL-2) and Th2 (IL-4) in PBMCs; IL-12p70 and MCP-1 in TMMCs. Upregulation of TNF-α was detected (Figure 4C) as well as significant downregulation of IL-17A in PBMCs at 1000 mg/kg UT-POA (Figure 5). There were no significant differences in IL-2 and IL-4 (PBMCs) or in IL-12p70 and MCP-1 (TMMC; data not shown).

CORRELATION BETWEEN TLs AND SYSTEMIC CYTOKINES

The correlations among the CD4/CD8a ratio of TLs, TNF-α, and IL-17A were evaluated. The results indicated that there was a positive correlation (*r* = 0.8559) between CD4/CD8a and TNF-α and a negative correlation (*r* = -0.8344) between CD4/CD8a and IL-17A. In both cases, the correlations were significant (*p* < 0.05; Figure 6).

DISCUSSION

This is the first study to evaluate the effects of a UT extract on the TME. There have been previous studies on the TME only to assess the oxidative parameters in tumor homogenates in a model based on Walker-256 rats. Also, UT has been evaluated in terms of B16 melanoma lung metastases (8). Populations of CD4 and CD8a TLs and Th1/Th2 cytokines have been quantified in splenocyte cultures, but in *healthy* BALB/c mice (34). Our results indicate that there is a significant increase in the CD4/CD8a ratio of the TLs from the PBMCs of mice with melanoma treated with UT at a dose of 1000 mg/kg, in contrast to the TME. This increase implies the presence of a higher percentage of CD4+ TLs or helper cells, which are lymphocytes characterized by active participation in the cellular immune response because they are capable of activating NKs, NKT cells, macrophages, cytotoxic TLs, and B lymphocytes.

These results are similar to those of Domingues *et al.*, who found upregulation of CD4+ TLs at doses 125 and 500 mg/kg of an hydroethanolic extract, although as mentioned above, they used healthy mice. Therefore, it is likely that a higher dose of UTPOA is necessary in mice with melanoma in order to achieve similar upregulation as in our study.

The activation level (CD44+) of the CD4+CD44+/CD8a+CD44+ ratio showed the opposite pattern, meaning that there was a significant increase at lower doses (50 and 500 mg/kg UT-POA), whereas at 1000 mg/kg, the ratio was lower but still greater than in the PC. These results suggest that UT-POA is capable of increasing not only the percentage of CD4+ T_H1s but also its activation level. It is possible that the activation level depends on the dose of UT-POA, but it was expected that this difference would remain in the TME. Nevertheless, it did not change significantly relative to PC. Although there are no differences in terms of percentages and activation of these cells in the TME, it is important to point out their presence, as well as the possibility of their activity's being affected by tumor factors released by the melanoma. In this study, it was not possible to evaluate other systemic lymphocyte populations. Nevertheless, they were analyzed in the TME, where these lymphocyte populations (represented by total T_H1s, NKs, and NKT cells) were not affected by the UT-POA treatment. Their presence is also an important finding, as is the possible inhibitory effects of tumor factors.

The ability of mDCs to intervene and induce Th1 antitumor responses was demonstrated here. Our results show that UT-POA at 100 mg/kg induced a significant increase in the % of mDCs, and at this same concentration, the % of pDCs is 7.5 times lower than the percentage of mDCs. This finding could be explained by the studies of Zuniga *et al.*, who reported that after a viral infection, pDCs can turn into mDCs via phenotypic and functional changes, including an improved ability to present antigens. In our case, it is possible that the UT-POA has an effect similar to that observed by Zuniga *et al.* Nevertheless, the mechanisms would have to be identified as well as whether these changes indeed occur or simply represent an effect directed specifically at mDCs while the pDCs are at normal levels.

TNF- α is produced not only by activated macrophages but also by tumor cells, and in cancer, this cytokine is reported to show paradoxical behavior because it is required for proliferation and function of NKs, T_H1s, B₁ cells, macrophages, and DCs, and it is an important molecule to the cell-mediated elimination of certain tumors, at least during the acute stage of the disease. When TNF- α is produced by macrophages, it causes M1 polarization that could be useful at the acute stage. On the other hand, there is evidence that TNF- α is one of the main mediators of cancer-related inflammation and acts as a protumor factor, at least during the chronic stage of the disease⁽³⁵⁾. Our results show significant systemic upregulation of TNF- α at 1000 mg/kg UT-POA, and this increase correlates positively with the CD4/CD8a ratio,

which in our model can be interpreted as a UT-POA-induced effect of an adaptive cellular response during the 29 days of treatment, which we can consider the acute stage. IL-17A also shows paradoxical behavior because it can impair the action of CD8+ T_H1s but can also induce inflammatory cytokines and chemokines that can attract CD8+ T_H1s, NKs, and DCs⁽³¹⁾.

Our results revealed a negative correlation between IL-17A and the CD4/CD8a ratio at 1000 mg/kg UT-POA, which means that downregulation of IL-17A under the influence of UT-POA may be associated with a low systemic proportion of CD8a T_H1s. Although in this study, we could not assess the levels of IL-17A in the TME, it would be useful to determine whether the presence of lymphocytes and DCs correlates with that cytokine. The evaluation of some immunological parameters allows us to conclude that UT-POA has better systemic effects than the effects in the TME, mainly because of the improved CD4/CD8a ratio and induction of Th1, inflammatory, or M1 macrophage polarization responses and a reduction in the Th17 response. Although these effects can be observed at high doses, we should take into account that after the first hydroalcoholic extraction, there was a second aqueous extraction, which may have dissolved some compounds and thus their potency may have been reduced. Nevertheless, UT exerted immunomodulatory effects. We recommend evaluating this extract after the first hydroalcoholic extraction. According to the above observation, the concentrations that we used should serve as a reference. Even though the model under study involves an aggressive type of cancer, our results are a good reference for future studies on the antitumor potential of UT-POA and its use as an adjunctive therapy during conventional treatment of cancer.

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Correspondence: Iván Lozada Requenaz
Address: Av. Honorio Delgado 430 Urb. Ingeniería – San Martín de Porres. Lima, Peru.
Phone number: 511-998674601
E-mail: ivan.lozada@upch.pe



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