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The small-molecule compound AC-73 targeting CD147 inhibits leukemic cell proliferation, induces autophagy and increases the chemotherapeutic sensitivity of acute myeloid leukemia cells

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ABSTRACT

D147 is a transmembrane glycoprotein with multiple functions in human healthy tissues and diseases, in particular in cancer. Overexpression of CD147 correlates with biological functions that promote tumor progression and confers resistance to chemotherapeutic drugs. In contrast to solid tumors, the role of CD147 has not been extensively studied in leukemia. Understanding whether CD147 represents a new hematologic target and whether its inhibitor AC-73 may be used in leukemia therapy may reveal an alternative treatment strategy in patients with acute myeloid leukemia (AML). We analyzed CD147 expression and function in hematopoietic progenitor cells from normal cord blood, in several leukemic cell lines and in primary leukemic blasts obtained from patients with AML. We investigated the effects of AC-73, used alone or in combination with arabinosylcytosine (Ara-C) and arsenic trioxide (ATO), on leukemic cell proliferation. We demonstrated that CD147 overexpression promotes leukemic cell proliferation. We showed that AC-73 exhibits a potent growth inhibitory activity in leukemic cells, by inhibiting the ERK/STAT3 activation pathway and activating autophagy. We demonstrated that AC-73 exerts an anti-proliferative effect additive to chemotherapy by enhancing leukemic cell sensitivity to Ara-C-induced cytotoxicity or to ATO-induced autophagy. We also reported CD147 expression in the fraction of leukemic blasts expressing CD371, a marker of leukemic stem cells. Altogether, our study indicates CD147 as a novel potential target in the treatment of AML and AC-73 as an anti-proliferative drug and an inducer of autophagy in leukemic cells to use in combination with chemotherapeutic agents.

Introduction

Targeted therapy for acute myeloid leukemia (AML) represents an ongoing challenge and in this context, cluster of differentiation 147 (CD147) represents an attractive target for therapeutic intervention in AML and in other hematologic neoplasms.¹⁻³ CD147, also known as basigin or extracellular matrix metalloproteinase inducer (EMMPRIN), is a type-I transmembrane glycoprotein that belongs to the immunoglobulin superfamily. Among the numerous studies that have documented the significance of CD147 in various physiological processes, the best characterized function of CD147 is related to its role in tumor metastasis, angiogenesis and chemoresistance processes.³⁻⁶ Overexpression of CD147 correlates with a number of biological functions that promote tumor progression (e.g. cellular proliferation, angiogenesis, matrix metalloproteinase production) and confers resistance to chemotherapeutic drugs such as adriamycin,^{7,8} cisplatin.⁹ CD147 mediates molecuHaematologica 2019 Volume 104(5):973-985

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lar events by interacting with various binding partners, such as tumor- and inflammation-associated molecules including integrins, monocarboxylate transporters (MCTs), cyclophilins, caveolin-1, and E-selectin, explaining its significant role in the pathogenesis of several diseases.^{3-6,10} CD147 overexpression and more recently its coexpression with MCTs^{11,12} are regarded as unfavorable prognostic factors in cancers associated with hypoxia, a common feature of solid tumors, but also a major component of the bone marrow (BM) microenvironment, crucial in leukemia progression.^{13,14} However, in contrast to solid tumors, the function of CD147 remains poorly defined in leukemia.

Recent studies have shown growing interest in the CD147 molecule in AML^{15,16} and in some hematologic neoplasia, in particular in multiple myeloma (MM), where CD147 expression levels have a prognostic value and are required for the proliferation of MM cells.¹⁷⁻¹⁹ Moreover, CD147 is over-expressed in erythroid cells of myelodysplastic syndrome (MDS) with 5q deletion.¹⁸ Here, we show that CD147 is expressed in normal CD34⁺ hematopoietic progenitor cells (HPCs) and down-regulated during monocytic and granulocytic differentiation of HPCs. We then show that CD147 is over-expressed in blasts pertaining to different subtypes of AML and promotes leukemic cell proliferation. Interestingly, we report that CD147 is expressed at the level of CD34⁺CD371⁺ AML cells, previously described for their leukemia-initiating properties.²⁰

Recently, the small-molecule AC-73 has been proposed as a specific inhibitor for CD147.²¹ First, we checked that the response to AC-73 treatment is not involved in an offtarget mechanism in leukemic cells. Then, we analyzed the effects of CD147 inhibition by AC-73 in AML cell lines and in primary leukemic blasts. We found that AC-73 inhibits leukemic cell proliferation by suppressing the ERK/STAT3 activation pathway, known to play a role in AML cell proliferation and survival,²² but also by activating autophagy, an essential phenomenon for hematopoietic stem cell (HSC) maintenance, resistance to stress, survival and differentiation, the machinery of which might be disrupted in AMLs.²³⁻²⁵ Next, we analyzed whether AC-73 enhanced the sensitivity of leukemic cells to conventional chemotherapeutic agents. We used arabinosylcytosine (Ara-C), one of the most active cytotoxic agents in myeloid leukemia, and arsenic trioxide (ATO), an active anti-proliferative agent used in the treatment of patients with acute promyelocytic leukemia (APL) (AML-M3)^{1,2,26} [although with low efficacy in AML lacking the t(15;17) translocation], and also an inducer of autophagy.25,27 We found that AC-73 used in vitro in combination with Ara-C or ATO, increases the effects of these agents.

Altogether, our data suggest that CD147 plays a key role in leukemic cell proliferation and represents a potential therapeutic target in AML patients, and that AC-73 is a new promising inhibitor that could be used in combination with conventional chemotherapeutic agents as a novel treatment strategy in AML.

Methods

Cell cultures

Human cord blood (CB) was obtained from healthy donors after informed consent. Leukemic blasts were isolated from BM $\,$

obtained from patients with newly diagnosed AML, using Ficoll-Hypaque density gradient. Informed consent was obtained from patients in accordance with the Declaration of Helsinki. This study was approved by the local ethical committees of the Istituto Superiore di Sanità and the University of Tor Vergata, Rome, Italy.

Cord blood CD34⁺ HPC purification, unilineage monocytic (Mo) and granulocytic (G) differentiation and morphological analyses were performed,^{28,29} as described in the *Online Supplementary Methods*.

Human primary AML blasts were maintained in culture in Iscove medium supplemented with 10% FCS, GM-CSF (10 ng/mL), SCF (50 ng/mL), IL-3 (10 ng/mL) (PeproTech Inc., Rocky Hill, NJ, USA), as described.²⁹

Human AML cell lines used in our study were: U937 as a model of AML-M5, NB4 and HL-60 as models of AML-M3 and AML-M4, respectively; NB4-R4 as AML-M3 resistant to all-*trans* retinoic acid (ATRA) treatment; MV4-11 as AML-M2 mutated for FLT3-ITD; Kasumi-1 as AML-M2 with the t(8;21) translocation. All cell lines were grown in RPMI medium supplemented with 10% FCS (Gibco, Carlsbad, CA, USA).

Cell growth, cell cycle profile, viability and apoptosis analysis were performed, as described in the *Online Supplementary Methods*.

Flow cytometry, western blot and quantitative real-time RT-PCR analysis were as described in the *Online Supplementary Methods.*

Knockdown of CD147 expression by RNA interference, clonogenic assays and colony formation assays were performed, as described³⁰ in the *Online Supplementary Methods*.

Cyto-ID autophagy detection was performed using the Cyto-ID assay (Enzo Life Sciences ENZ-51031-K200) as described in the *Online Supplementary Methods*.

AC-73 treatment of leukemic cells used alone or in combination with chemotherapeutic agents

AC-73 (3-{2-[([1,1'-biphenyl]-4-ylmethyl) amino]-1-hydroxyethyl}phenol) (Specs ID number AN-465/42834501, Specs, Zoetermeer, the Netherlands) was dissolved in 20% DMSO (Sigma, St. Louis, MO, USA) and diluted in DMEM, with a final DMSO concentration of no more than 0.2% for all in vitro studies.²¹ In leukemic cell lines, dose-response and time-course analysis were performed using AC-73 at 1.0, 2.5, 5.0 and 10 μ M from 1 to 4 days of treatment; results were compared with 0.2% DMSOtreated leukemic cells, indicated as control leukemic cells. AC-73 was added in cultures every 2 days to maintain its activity. Combinations of treatment were performed using AC-73 (2.5 μ M) alone over 24 hours and then by adding Ara-C (0.01; 0.1 and 1.0 $\mu M)$ or ATO (0.1; 0.01 and 1.0 $\mu M)$ for another 1 (for NB4 and NB4-R4 cells) or 2 (for U937, HL-60, MV4-11 and Kasumi-1 cells) days. Cell viability assays were performed to evaluate the effect of AC-73 alone or in combination on cell growth and viability of these cells.

Analysis of The Cancer Genome Atlas data

Datasets of The Cancer Genome Atlas (TCGA) Research Network 2008, were processed and obtained directly from the public access data portal (*http://tcga-data.nci.nih.gov/*).

Statistical analysis

Student *t*-test was applied to assess statistical significance of differences between multiple/group of experiments. Data were analyzed using GraphPad Prism software. For univariate survival analysis, Kaplan-Meier plots with a log-rank test were presented using the overall survival data of AML patients from the TGCA. Additional information is provided in the *Online Supplementary Appendix.*

Results

CD147 is directly involved in hematopoietic progenitor cell proliferation

We analyzed CD147 expression at both mRNA and protein level in CD34⁺ HPCs and during Mo and G proliferation and differentiation of these cells. CD147 is well expressed in CD34⁺ HPCs and its level of expression decreases during Mo and G differentiation and maturation of these cells (Figure 1A-C and Online Supplementary Figure

S1A and B). Western blot analysis shows that CD147 protein is highly glycosylated (HG-CD147 40-60 kDa) in CD34⁺ HPCs and at all stages of Mo and G cell differentiation (Figure 1B), indicating the presence of CD147 protein in a stable and biologically active conformation mainly translocated to plasma membrane,⁵ as also shown by flow cytometry analysis (Figure 1C and Online *Supplementary Figure S1B*). To analyze the role of CD147 in HPCs, we performed siRNA-mediated CD147 knockdown experiments in CD34⁺ HPCs. We transiently trans-



Figure 1. CD147 is down-regulated during monocytic (Mo) and granulocytic (G) differentiation of CD34⁺ hematopoietic progenitor cells (HPCs) and is involved in HPC proliferation. (A) gRT-PCR analysis of CD147 mRNA expression during selective Mo and G proliferation and differentiation of CD34* HPCs, as compared to U937 and HL-60 leukemic cells. (B) Western blot analysis of CD147 protein expression level in CD34⁺ cells and Mo and G differentiating HPCs; Jurkat and SKBR3 cells are shown as positive controls of CD147 expression; HG- and LG- are indicated for High- and Low- glycosylated CD147 isoforms; actin is shown as an internal control; molecular weights are indicated (kDa). (C) Flow cytometry analysis of CD147 membrane protein expression during Mo and G differentiation and maturation of HPCs. (D) Flow cytometry analysis of CD147 membrane protein expression in Mo and G differentiating transfected (CD147-siRNA)-HPCs, as compared to transfected (c-siRNA)-HPCs of control, at day 2 of Mo and G cultures corresponding to day 3 post transfection. (E) Cell growth inhibition of (CD147-siRNA)-HPCs, as compared to (c-siRNA)-HPCs, grown under both G and Mo liquid culture conditions. (F) Clonogenic assays performed under G and Mo culture conditions with (CD147siRNA)-HPCs, as compared to (c-siRNA)-HPCs. (A, C-F) Mean±Standard Error of Mean of three independent experiments is shown. **P*<0.05; ***P*<0.01; ***P<0.001. (B) One representative experiment out of three is shown. AU: arbitrary units; MFI: mean fluorescence intensity

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fected HPCs with specific-CD147 interfering small RNAs (CD147-siRNA) or with a non-targeting control siRNA (csiRNA). Silencing of CD147 mRNA decreased CD147 protein levels in transfected (CD147-siRNA)-HPCs, as compared to (c-siRNA)-HPCs of control (Figure 1D); then, transfected-HPCs were grown under culture conditions, to allow their selective G or Mo proliferation and differentiation. We observed that the decreased expression of CD147 impaired the proliferation of both G and Mo differentiating (CD147-siRNA)-HPCs, as compared to control (c-siRNA)-HPCs (Figure 1E), without any significant modulation of G or Mo specific antigen expression on these cells (data not shown) and without any significant induction of apoptosis (data not shown). Clonogenic G and Mo progenitor assays performed with transfected-HPCs, showed a significant decrease in the number of CFU-GM and CFU-M colonies obtained from (CD147-siRNA)-HPCs, compared with the number of colonies obtained from (c-siRNA)-HPCs (Figure 1F). Altogether, our data indicate that CD147 plays a role in HPC proliferation and clonogenic activity.

Over-expressed in AML, CD147 is down-regulated during differentiation of leukemic cells

By analyzing CD147 mRNA expression levels in primary leukemic blast cells obtained from 48 patients with different subtypes of AML,²⁹ we found that CD147 is over-expressed in all AML subtypes, in particular in the AML-M3 subtype, as compared to normal CD34⁺ HPCs (Figure 2A, left panel), in line with the data from AML samples generated by the TCGA Research Network (Figure 2A, right panel).

Then, we also examined CD147 expression in several AML cell lines induced to terminal differentiation. All leukemic cell lines analyzed over-expressed the fully glycosylated mature CD147 protein, biologically active (Figure 2B, HG-CD147), like also its mRNA, as compared to normal CD34⁺ HPCs (Figure 2C). We found that CD147 expression is down-regulated during vitamin D3-induced Mo differentiation of U937 cells and ATRA-induced G differentiation of HL-60 and NB4 cell lines, at both mRNA and protein levels during terminal differentiation (Figure 2D-F and Online Supplementary Figure S1C-H), similarly to Mo and G terminal differentiation of normal HPCs (Figure 1A-C). However, CD147 expression is not affected by ATRA treatment of the NB4-R4 cell line (Figure 2G and Online Supplementary Figure S1I and L), indicating that induction of cell differentiation is required for CD147 downregulation.

AC-73 inhibits leukemic cell proliferation by blocking ERK/STAT3 signaling and induces autophagy

To investigate whether a high level of CD147 can promote cell proliferation, we used AC-73 to inhibit CD147 function in normal and leukemic cells.

First, we examined the effect of AC-73 treatment on G and Mo differentiation of normal HPCs. AC-73 used at 5 μ M moderately decreased cell growth without affecting cell cycling or differentiation of HPCs (Figure 3A-C). Morphology analysis of HPCs at different days of culture supported these observations, showing that AC-73-treated cells regularly progress along G and Mo differentiation and do not show any significant morphology abnormality or delay in differentiation (Figure 3D).

We analyzed the effects of chronic AC-73 administra-

tion on leukemic cell growth, apoptosis and viability of AML cell lines. Leukemic cells were treated from 1 to 3 days, with various AC-73 doses (2.5, 5 and 10 μ M) to determine their sensitivity to AC-73 treatment. We observed a time- and dose-dependent effect of AC-73 on leukemic cell growth, with growth inhibition of all leukemic cell lines treated (Figure 4A and Online Supplementary Figure S2B). We could not detect any effect of AC-73 on cell growth and apoptosis of the Chinese Hamster Ovary (CHO) cells (data not shown) used as a CD147-negative cell line (Online Supplementary Figure S2A, left panel).

NB4 and NB4-R4 cell lines were more sensitive to AC-73 treatment than U937 and HL-60 cell lines. Indeed, a higher significant decrease in cell growth (Figure 4A), apoptosis (Figure 4B), and viability (Figure 4C and D) was observed in NB4 and NB4-R4 cells treated with low-dose AC-73 (2.5 μ M) than in U937 and HL-60 lines (2 days) (Figure 4D). MV4-11, Kasumi-1 cells were the most resistant to AC-73 treatment (*Online Supplementary Figure S2B-E*) when compared to other cell lines (Figure 4A-D). Significant cell growth inhibition and sensitivity of MV4-11 and Kasumi-1 cells occurred after treatment with high concentration of AC-73 (5 and 10 μ M) (*Online Supplementary Figure S2B-D*) or when cells were treated for longer (3-4 days) (*Online Supplementary Figure S2E*).

AC-73 treatment had no significant effect on cell cycle distribution in the leukemic cell lines tested as compared to control cells (*data not shown*), suggesting that AC-73 decreases the cell growth rate but does not inhibit the progression of cells in the cell cycle. Analysis of CD11b, CD14 and CD15 expression levels in leukemic cell lines treated for 3 days with 5 μ M AC-73 showed no significant effect of AC-73 on leukemic cell differentiation (*data not shown*). By analyzing the effects of AC-73 (5 μ M) on clonogenic growth of U937 and NB-4 cells we found that AC-73 decreases the *in vitro* colony formation of both U937 and NB-4 cell lines, indicating an inhibitory effect of AC-73 on the clonogenetic capacity of leukemic cells (*Online Supplementary Figure S2F*).

Altogether, despite the different sensitivities to AC-73 manifested by the different leukemic cell lines, probably related to the specific molecular alteration, such as *PML/RARA, FLT3-ITD* or *RUNX1/RUNX1T1*, of these cell lines, AC-73 exhibits potent growth inhibition and cytotoxic activity on leukemic cells only at high doses. Because low doses of AC-73 inhibit leukemic cell proliferation, but do not cause cell death via apoptosis or cell cycle arrest, we investigated the possibility that AC-73 treatment induces autophagy in leukemic cells. First, we assessed the effect of AC-73 on the level of the autophagic indicator LC3 by western blotting in U937 and NB4 cells. Our results showed a dose-dependent effect of AC-73 on the increase of LC3-II/LC3-I ratio in leukemic cells as compared to control (-) cells (Figure 4E), indicating induction of autophagy.³¹ Then we monitored autophagy flux by flow cytometry analysis in live leukemic cells treated for 72 hours (h) by AC-73 compared to control cells. Our data demonstrated that AC-73 induces dose-dependent autophagy (Figure 4F) in CD147-expressing leukemic cells (Figure 4G), but not in CHO cells (Online Supplementary) *Figure S2A, right panels*), again indicating that the biological effects induced by AC-73 require CD147 expression on target cells. Considering that AC-73 suppresses CD147/ERK1/2/STAT3/MMP-2 pathways in hepatocellular carcinoma cells, and that ERK/STAT3 signaling plays a significant role in promoting AML cell proliferation, survival and autophagy,^{22,23,32,33} we also investigated the effect of AC-73 on ERK1/2 and STAT3 activation in leukemic cells. In line with previous studies,^{31,32} western blot analysis showed that ERK and STAT3 are constitutively phosphorylated in all leukemic cell lines [*Online Supplementary Figure S3*, lanes (-)]. We then found that AC-73 (5 μ M, for 3 days) notably decreases both pERK and pSTAT3 (S727) levels without affecting the levels of total ERK and STAT3, in leukemic cells (lanes 5 μ M) (*Online Supplementary Figure S3*, lower panels).

Altogether, our data demonstrate that AC-73 inhibits leukemic cell proliferation, in part by suppressing the ERK/STAT3 activation pathway in these cells, in part by activating a non-apoptotic, autophagic form of cell death, which may account for the efficacy of AC-73 in leukemic cells.

AC-73 has an additive anti-proliferative effect on chemotherapeutic treatment of leukemic cells and increases ATO-induced autophagy in leukemic cells

To evaluate the effect of AC-73 in combination with standard drugs used for AML treatment, AML cell lines

d5

d2

ATRA

ATRA

Figure 2. Over-expressed in acute myeloid leukemia (AML), CD147 expression is down-regulated during monocytic and granulocytic differentiation of leukemic cells. (A) qRT-PCR analysis of CD147 mRNA expression in primary leukemic cells of AMLs pertaining from M0 to M5 subtypes of the French-American-British (FAB) classification, as compared to normal CD34* hematopoietic progenitor cells (HPCs) (left panel); CD147 mRNA expression data from AML samples generated by TCGA Research Network (right panel). (B) Western blot analysis of CD147 protein expression level in AML cell lines; densitometry analysis [arbitrary units (AU)] of CD147 protein expression levels compared with actin levels is indicated. (C) qRT-PCR analysis of CD147 mRNA expression in leukemic cell lines, as compared to normal CD34⁺ HPCs. (D-G) Western blot (left panels) and flow cytometry (right panels) analysis of CD147 total and membrane protein expression levels during vitamin D3-induced monocytic differentiation of U937 cell (D), ATRAinduced granulocytic differentiation of HL-60 cells (E), ATRA-induced differentiation of NB4 cells (F), and in NB4-R4 cells, resistant to ATRA treatment (G). (A and C) Mean±Standard Error of Mean (SEM) of three independent experiments is shown. *P<0.05; **P<0.01; ***P<0.001. (B, D-G, left panels) One representative western blot experiment out of three is shown. High (HG) and Low (LG) glycosylated CD147 isoforms are indicated. Actin is shown as internal control. (D-G, right panels) Mean±SEM of three independent experiments by flow cytometry analysis is shown. *P<0.05; **P<0.01; ***P<0.001. ns: not significant; RPKM: Reads Per Kilobase Million; MFI: mean fluorescence intensity.



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and primary AML blasts were initially treated for 24 h with a low concentration (2.5 μ M) of AC-73 and then cotreated for 48 h with Ara-C or ATO at different concentrations (0.01-1 μ M). Cell viability assays were performed to assess leukemic cell survival and sensitivity to AC-73 treatment used in combination with Ara-C or ATO, as compared to single AC-73, Ara-C, or ATO treatment (Figure 5A and B and Online Supplementary Figure S4). We observed a significant decrease in leukemic cell viability of all leukemic cell lines after AC-73 treatment in combination with Ara-C or ATO, as compared to AC-73, Ara-C or ATO used alone (Figure 5A and B and Online Supplementary Figure S4). Furthermore, we identified three subgroups of leukemic cells according to their Ara-C drugsensitivity: high-sensitivity lines (U937 and HL-60); intermediate-sensitivity lines (NB4 and NB4-R4); low-sensitivity lines (MV4-11 and Kasumi-1). AC-73 potentiates the sensitivity to Ara-C treatment of all leukemic cells, even those belonging to the low-sensitivity subgroup (Figure 5A and B and *Online Supplementary Figure S4*). The antiproliferative effect of AC-73 enhances sensitivity to ATO treatment of both M3 leukemic cells, such as NB4 and NB4-R4 (Figure 5B and *Online Supplementary Figure S4B*), and non-M3 leukemic cells, such as U937, HL-60, Kasumi-1 and MV4-11 cells (Figure 5A and *Online Supplementary Figure S4A*, *C and D*).

Then, we showed that when used in combination with Ara-C and ATO, AC-73 treatment inhibits ERK and STAT3 activation in leukemic cells (Figure 5C).

Because ATO is also an inducer of autophagy,²⁵ we analyzed the autophagy flux in both M3 and non-M3 leukemic cells treated by ATO or AC-73 used alone or in combination, as compared to control cells. Our data showed that AC-73 and ATO are both inducers of autophagy and the autophagy flux is significantly increased in M3 (NB4) and non-M3 (U937) leukemic cells



treated by ATO and AC-73 used in combination, as compared to single agent treatment or control cells (C) (Figure 5D and E).

AC-73 increases the sensitivity of primary acute myeloid leukemia blasts co-expressing CD147 and CD371 to chemotherapy

Altogether, our data demonstrate that AC-73 is a potent, novel anti-proliferative molecule that enhances the sensitivity of leukemic cells to conventional chemotherapeutic agents, but also increases ATO-induced autophagy in M3 and non-M3 leukemic cells. We evaluated the effects of AC-73 used alone and in combination with Ara-C or ATO on primary AML blasts. First, we analyzed CD147 protein expression in samples obtained from different subtypes of AML and we controlled the expression of CD34, CD38 and CD371 mark-





ers previously identified in LSCs.³⁴ We confirmed that CD147 is expressed at high levels in all AML subtypes (M1, M2, M3, M5 and M5a) analyzed (Table 1). We also report for the first time that CD147 is co-expressed with CD34, and particularly with CD371, in these cells (Table 1 and Figure 6A). It is interesting to note that in 8 non-M3 AMLs analyzed for CD147 and CD371 expression, there was a strong positive correlation between CD147 and CD371 expression (P<0.001) (Table 1). These data suggest

the existence of CD34⁺/CD147⁺ and CD371⁺/CD147⁺ subpopulations in AMLs that may have functional properties of leukemic stem cells (LSCs).

Then, we investigated the effects of AC-73 used alone or combined with Ara-C and ATO at low concentration in primary AML blasts, and particularly in M2, M3 and M5 AML cells. We found that AC-73 used alone has no significant effect on AML cell apoptosis, unless used at high concentration (10 μ M) for AML(7)-M3 and AML(8)-M5



Figure 5. AC-73 potentiates the sensitivity of leukemic cells to chemotherapeutic treatment. (A and B) Cell viability assays on U937 and NB4 leukemic cells, treated 24 hours (h) with AC-73 used alone (2.5 µM), then in combination with Ara-C (AC-73 + Ara-C, left panels) or ATO (AC-73 + ATO, right panels) for 48 h, as compared to treatment with single drug, AC-73 (AC-73 + 0.00 µM Ara-C; AC-73 + 0.00 μM ATO), Ara-C and ATO. (C) Western blot analysis of phospho-STAT3 (ser727) (pSTAT3(S727), STAT3, phospho-ERK1/2 (pERK) and ERK1/2 (ERK) protein expression in U937, NB4 and MV4-11 leukemic cell lines. U937, NB4 and MV4-11 cells were treated for 3 days with AC-73 in combination with ATO (AC-73 + ATO 1 μM), or Ara-C (AC-73 + Ara-C 1 μM), as compared to AC-73 (2.5 µM), ATO (1 µM) or Ara-C (1 µM) used alone or to control (C) leukemic cells. Actin was used as an internal control. Quantification of total and phosphorylated ERK1/2 and STAT3 proteins by densitometry analysis is indicated. (D and E) Analysis of the autophagy flux induces by AT0 (1 μM) and AC-73 (5 μM) used alone or in combination in U937 and NB4 leukemic cells compared to control cells (C). (A. B and E) Mean±Standard Error of Mean of three independent experiments is shown. *P<0.05: **P<0.01: ***P<0.001. ns: not significant. (C and D) One representative experiment out of three is shown. (C) A.U.: arbitrary units. pSTAT3/STAT3 Ratio and pERK/ERK of three independent experiments is shown. kDa: molecular weights.

(Figure 6B). However, AC-73 induces autophagy by increasing the LC3-II/LC3-I ratio in leukemic blasts, as compared to control (-) leukemic cells, even at 2.5 or 5 μ M (Figure 6C). However, combination of low-dose AC-73 (2.5 μ M) and ATO or Ara-C (0.1 and 1 μ M) more efficiently decreased leukemic blast viability, compared with single AC-73, ATO or Ara treatment (Figure 6D). Overall, our data indicate that AC-73 activates a non-apoptotic autophagic form of cell death in AML blasts and increases the sensitivity of these cells to Ara-C or ATO agents.

Analysis of CD147 expression in acute myeloid leukemia based on TCGA dataset

By providing selected genetic and clinical data from 200 AML patients, useful for prognosis and diagnosis, the TCGA dataset offered a unique opportunity to explore a possible relationship between the recurrent gene mutations observed in AML and the level of CD147 expression. TCGA data set analysis indicates that CD147 level is particularly elevated in AML-M3, those AML subtypes bearing *PML-RARA* fusion gene^{1,26} (Figure 7A, RARA), in





line with the data observed from fresh samples obtained from AML-M3 patients shown in Figure 2A, while significantly lower levels were observed in AMLs blasts mutated for *NPM1*, *DNMT3A* or *FLT3* (Figure 7A). By analyzing CD147 levels in all AMLs that we have stratified into three risk groups according to the European LeukemiaNet (ENL) risk classification, we found that: 1) the lower CD147 levels were observed in poor- and intermediaterisk AMLs; 2) surprisingly, the highest CD147 levels were observed in AMLs with favorable risk profile, mainly due to the presence in this group of AML-M3 (Figure 7B). However, it is important to point out that approximately 20% of poor- and intermediate-risk AMLs also display elevated CD147 expression (Figure 7B). Because AML-M3 have the best prognosis among all AML subtypes for the efficiency of target therapy based on ATRA and ATO,²⁶ those patients with favorable risk and high CD147 levels may represent a bias for our analysis. Therefore, we evaluated a possible correlation between CD147 expression level and overall survival of AML patients, excluding AML-M3 patients, and according to CD147 levels classified as: low (CD147<70); medium (CD147; range, 70-120); high (CD147 >120) (Figure 7C). In these non-M3 patients who died within 60 months after diagnosis, earlier death (within 20 months) was observed more frequently in cases with high CD147 levels, as compared with (non-M3)-AML subgroups with low-medium CD147 levels (P=0.003 and P=0.0003, respectively) (Figure 7D). These three (non-M3)-AML subgroups had comparable age and white blood count (WBC) at diagnosis (Figure 7D, right panels). Our data indicate that, excluding AML-M3, high CD147 levels are correlated with early death of AML patients.

Discussion

CD147 represents a prognostic marker in several solid tumors and in hematologic malignancies.^{3,6-9,15-19} However, the biological function of CD147 and its potential role as a marker in leukemia remain poorly defined. Apart from the erythrocyte lineage, the expression, regulation and function of CD147 in normal and leukemic hematopoietic cells have not been extensively studied.³⁵ CD147, previ-

ously identified as a carrier molecule for the blood group antigen OKa,36 has been involved in the recirculation of mature erythrocytes from the spleen into the general circulation.³⁷ In our study, we show that CD147 is important for the proliferation of normal CD34⁺ HPCs and its expression is down-regulated during Mo and G differentiation. Notably, we found that CD147 is over-expressed in all leukemic cell lines and in the large majority of primary leukemic blasts analyzed, as compared to normal CD34⁺ HPCs. However, CD147 expression significantly decreases during Mo and G differentiation of leukemic cells, thus mimicking what occurs in Mo and G lineage cells. By using AC-73 to inhibit CD147 function in leukemic cell lines and in primary AML blasts, we demonstrated that CD147 overexpression promotes leukemic cell proliferation. Interestingly, we observed that AC-73 treatment inhibits leukemic cell proliferation by suppressing the activation of the ERK/STAT3 signaling pathway, in line with previous studies.^{5,21} As constitutive STAT3 activity has been described to promote AML cell proliferation and survival,^{22,38} targeting key tyrosine kinases upstream of STAT3 has been proposed as a strategy to treat AML.³⁸⁻⁴⁰ However, due to its role in stem cell renewal, the potential risk of systemic STAT3 inhibition could be a deregulation of hematopoiesis.⁴¹ Here, we found that CD147 is expressed in normal HPCs and also showed that a low dose of AC-73 induces only a slight inhibition of cell proliferation, without affecting differentiation of CD34⁺ HPCs. Importantly, we found that AC-73 does not cause cell death via apoptosis or cell cycle arrest, but induces autophagy in leukemic cells, as described in previous studies carried out in AML cells treated with ATO as an inducer of autophagy,²³ or with other novel targeted drugs.^{23,27,42-} ⁴⁴ In some cases, autophagy accounted for a non-apoptotic decrease in cell viability.44

To further validate its potential therapeutic activity, we used AC-73 in combination with conventional antileukemia treatment, and showed that the anti-proliferative effect of AC-73 on leukemic cells enhanced the sensitivity of leukemic cells to chemotherapeutic treatments such as Ara-C or ATO, which could be consequently used at lower concentration. Our findings are in line with recent studies showing that other compounds, such as chidamide that down-regulates the JAK2/STAT3 signaling,⁴⁵

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Patient n.	FAB subtype	CD147 (%)	CD34 (%)	CD371 (%)	CD11b (%)	CD14 (%)	
1	M1	63.1	21.2	80.3	31	1	
2	M1	93.7	92.4	88.6	ND	ND	
3	M1	44.4	98.6	64.5	28	2.3	
4	M2	95	11.1	88.3	5	1	
5	M2	92.5	39.5	94	78	18.5	
6	M3	63.7	0.2	31	4	ND	
7	M3	97.9	12.5	95.9	3.2	0	
8	M5	71.5	5	90.9	60	21	
9	M5a	96.6	8.3	93.2	33.6	8.8	
10	M5	91.3	73	ND	69	17	

Flow cytometry analysis of CD147, CD34, CD371 and CD11b, CD14 cell surface antigen expression in 10 primary acute myeloid leukemia (AML) samples, obtained from patients with AML pertaining to different French-American-British (FAB) classification subtypes. Results are shown as percentage (%) of expression. ND: not determined.

inhibit the viability of AML cells and synergize with cytarabine in inhibiting AML cell viability.⁴⁶ Importantly, we also showed that AC-73 increases ATO-induced autophagy in both M3 and non-M3 leukemic cells, indicating a possible synergistic anti-tumor interaction of ATO with an inhibitor of CD147, both agents being able to induce autophagy.

The analysis of CD147 expression in AML subsets presented several interesting findings. First, CD147 is overexpressed in all French-American-British classification AML subtypes, and particularly in AML-M3; a finding confirmed through the analysis of primary AML samples. This finding was also corroborated by the analysis of the TCGA data set. Moreover, about 20% of non-M3 AMLs displayed elevated CD147 levels. Although the overall long-term survival of these patients (defined as highlyexpressing CD147) was similar to the other two non-M3 AML groups (defined as middle- and low-expressing CD147), death occurs earlier in patients showing high expression of CD147. This observation supports the negative prognostic role of high CD147 expression levels in AMLs, as observed also in other tumors, including some hematologic neoplasia.^{3,19} Other recurrent AML mutations,^{1,2} such as *NPM1*, *FLT3-ITD* and *DNMT3A*, were not associated to particularly elevated levels of CD147. Therefore, AC-73 used in combination with Ara-C or ATO may have clinical potential implications in treatment of AML patients expressing CD147. Future toxicological and pharmacodynamic studies in suitable animal models will be required for a preclinical evaluation of the possible impact of AC-73 as an anti-leukemic drug.

Our study also reports for the first time that CD147 is expressed in a sub-fraction CD34⁺CD371⁺ of AML cells that is able to engraft immunodeficient mice,²⁰ CD34 being predominantly regarded as a marker of hematopoietic stem cells (HSC) and HPCs, and CD371 as a marker of



Figure 7. CD147 gene expression level does not predict overall survival in acute myeloid leukemia (AML), but is correlated in early death of (non-M3)-AML patients. (A) Relationship between the most recurrent gene mutations observed in all AMLs and the level of CD147 mRNA expression, according to the TCGA dataset. (B) CD147 mRNA levels were analyzed in AMLs stratified into 3 risk groups: poor, intermediate and favorable, according to the European LeukemiaNet (ENL) risk classification. (C) Kaplan-Meier survival analysis in a new group of AML patients that excludes AML-M3 patients, based on CD147 gene expression and stratified according to their CD147 mRNA levels in 3 groups: low (CD147<70); medium (CD147, range, 70-120); high (CD147 >120). (D) Kaplan-Meier survival analysis in the new group of (non-M3)-AML patients deceased within 50-60 months years after diagnosis, indicates that the kinetics of death is more rapid (within 20 months) among CD147 high patients, as compared with those with low (P<0.01) or medium (P<0.001) CD147 levels: P-values calculated by log-rank test. The histograms show that white blood count (WBC) number, age at diagnosis and proportion of patients with poor cytogenetics were comparable in the 3 (non-M3)-AML subgroups subdivided according to CD147 expression level.

LSCs and a novel prognostic predictor in AML.^{20,34,47} Interestingly, in non-M3 AMLs a strong positive correlation was observed between CD147 and CD371 expression. In this context, while a dedicated future study would shed light on the functional properties of LSCs of the fractions of (CD34⁺CD147⁺) and (CD34⁺CD371⁺CD147⁺) AML cells, our data suggest that CD147 is expressed in LSCs and may be a potential new prognostic marker in AML and a potential therapeutic target.

Taken together, our study demonstrates a pivotal role for CD147 in leukemic cell proliferation and indicates CD147 as a potential therapeutic target of AC-73 in AML. Thus, targeting CD147 by using AC-73 or derivate molecules to induce autophagy or to inhibit the growth of leukemic clones, including LSCs, may be an effective option to improve leukemia treatment.

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