

Research Article



Induced Antileukemic Activity after Blast Modulation is Independent from Immune Checkpoint Marker Expression on Patients' Blasts or T Cells

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Abstract

Background: Acute myeloid leukemia is still associated with a poor prognosis. New therapeutic strategies are necessary. Myeloid leukemic blasts can be converted into dendritic cells (DC) of leukemic origin leukemiaderived DC (DC_{len}) using 'DC-generating Picis' or 'DC-generating Kits', resulting in enhanced leukemia-specific antileukemic immune responses.

Methods: DC/DC_{leu} were generated out of leukemic Peripheral Blood Mononuclear Cells (PBMNC) or whole blood (WB) using DC/DC_{leu} generating protocols and used to stimulate T cell enriched immunoreactive cells in mixed-lymphocyte culture (MLC), followed by a cytotoxicity fluorolysis assay (CTX). We evaluated the expression profiles of immune checkpoint molecules (CD279 (PD-1), CD273 (PD-L2) and CD274 (PD-L1)) on uncultured blasts and T cells from AML patients respectively on monocytes and T cells from healthy donors after MLC with Kit pre-treated WB and correlated immune checkpoint (CP) expressions with patients' clinical data and functional cytotoxicity.

Results: We were able to generate DC/DC_{leu} from leukemic (and healthy) PBMNC and WB without induction of blast proliferation. Stimulation of immunoreactive cells after MLC with Kit pre-treated $\mathrm{DC/DC}_{\mathrm{leu}}$ containing WB resulted in downregulated CP expression on blasts and DC_{leu} and in increased anti-leukemic cytotoxicity. CP expressing T cells (T₂₇₀₊) correlated negatively with response to induction therapy and with improved blasts lysis ex vivo (T_{274+}) .

Conclusion: Through this immunomodulatory approach, we demonstrated the potential to induce or enhance leukemia-specific and anti-leukemic activity ex vivo, largely independent of checkpoint inhibitor expression on blasts, T cells, or patients' clinical characteristics. Building on these findings, our data further suggest that DC/DC_{leu} could be generated ex vivo using 'DCgenerating Picis' or 'DC-generating Kits' for subsequent adaptive transfer to patients, or that in vivo treatment with Kit-M could be explored to help stabilize remissions in AML patients.

Keywords: Acute myeloid leukemia; Immune checkpoint molecules; Leukemia derived dendritic cells; Anti-leukemic functionality

Introduction

Classification and Diagnosis of AML

Acute myeloid leukemia (AML) is characterized by uncontrolled proliferation of myeloid progenitor cells ('blasts') in the bone marrow (BM) and peripheral blood (PB), leading to suppression of normal haematopoiesis and resulting in

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anaemia, bleeding, infections and impaired antitumor immune response [1]. Diagnosis is based on cytomorphological, cytogenetic, molecular and immunophenotypic analyses, which allows for subtyping and prognostic classification of cases [2-4]. The French-American-British (FAB) and WHO classifications divide into subtpyes M0 – M7, while the ELN risk classification enables prognostic subgrouping in 'favorable', 'intermediate' and 'adverse' risk [2].

Treatment strategies for AML patients (pts)

Standard therapy consists of high-dose cytarabine plus anthracycline chemotherapy, followed by hematopoietic stem cell transplantation (HSCT) in younger pts or treatment with hypomethylating agents, often combined with venetoclax [2,5-7]. Although remissions reach up to 80%, most pts relapse within two years, resulting in an unsatisfying prognosis with a 5-year-survival rate of 31.9% [8]. Relapse is driven by residual blasts and/or immune evasion through inhibition of effector T cells, natural killer cells (NK cells) and dentritic cells [9].

Novel therapeutic approaches aim to overcome immune evasion and trigger tumor-specific immune response, e.g. by epigenetic modulation, targeting immune checkpoint molecules or employing antibody-based or cellular immunotherapies (CAR-T cells, NK cells, DC-based strategies) [2,10]. DC-based strategies have the competence to prime and enhance (leukemia-specific) immune responses by presenting individual patients' whole leukemic antigen repertoire and inducing immunological memory [11,12].

Dendritic cells (DC) and leukemia-derived DC (DC $_{\rm leu}$)

DC are potent antigen-presenting cells that stimulate and regulate immune responses. They can be generated from patients' monocytes and loaded with leukemic antigens. Moreover leukemic blasts from Peripheral Blood Mononuclear Cells (PBMNC) or whole blood (WB) can be converted into leukemia-derived DC (DC_{leu}), expressing DC antigens as well as leukemia-associated antigens. DC/ DC_{leu} were generated using specific response modifiers in accordance with established DC/DC_{len}-generating protocols. Different compositions of response modifiers were employed during this process, including the so-called 'DC-generating Picis' (Pici1 (GM-CSF, IL-4, Picibanil and PGE1) or Pici2 (GM-CSF, IL-4, Picibanil and PGE2)) or 'DC-generating Kits' (Kit-I (GM-CSF and OK-432), Kit-K (GM-CSF and PGE2) or Kit-M (GM-CSF and PGE1)) [13], without the induction of blast proliferation [14,15]. Experimental data from leukemic rat models and clinical applications in refractory AML patients before or after alloHSCT indicate that in vivo DC/DC_{leu} induction can activate leukemia-specific immune responses and promote disease stabilization [16,17].

DC/DC_{leu} thus enhance 'graft-versus-leukemia' (GvL) effects and establish leukemia-specific immunologic memory [10-13,18].

Immune checkpoint molecules

Immune checkpoint molecules (CP) regulate immune responses by balancing T and NK cell activation [19-21]. While essential for maintaining self-tolerance, they also modulate immune reactions against pathogens or tumors [20,22,23].

CD279 (PD-1), expressed on activated T, B, NK and antigen-presenting cells, interacts with two main ligands CD273 (PD-L2) and CD274 (PD-L1) [24-26]. CD273 and CD274 are found on antigen-presenting, hematopoietic and non-hematopoietic cells and on the surface of tumor cells [27,28]. Engagement of CD279 with these ligands inhibits cytokine production, reduces T cell activation up to T cell exhaustion and promotes regulatory T cell (Tregs) expansion, which further hampers the effector function of CD8 T cells [9,29-31]. Over-expression of CP in AML leads to immune escape through evading detection and elimination by the immune system [32-35].

Immune checkpoint inhibitors (ICI) aim to restore antitumor immunity by blocking CP [36–38], leading to the reactivation of T cells [39,40]. While ICI monotherapy (e.g. nivolumab, pembrolizumab, atezolizumab or durvalumab) has shown success in solid tumors, clinical efficacy in AML remains limited. Initial studies combining ICI with hypomethylating agents suggest potential synergistic effects and acceptable safety profiles [23,29,41–44]. However, immune-related adverse events may occur during ICI treatment [29,45,46], which vary in severity and can potentially affect almost any organ [47,48].

Aim of this work

This study investigates the role of CP CD273, CD274 and CD279 in AML in the context of DC/DC_{leu}-based immunotherapeutic approaches. Specifically, we analysed CP expression on uncultured blasts from AML pts and healthy donor monocytes, as well as on DC/DC_{leu} generated from leukemic and healthy WB using DC/DC_{leu}-generating Pici-methods (Pici1, Pici2) or Kits (Kit-I, Kit-K, Kit-M). To evaluate the immunomodulatory potential, T cell-enriched immunoreactive cells were stimulated in a mixed lymphocyte culture (MLC) with Kit-pretreated WB and the effect of DC/ DC_{leu} stimulation on immune cell composition as well as the induction of antileukemic cytotoxicity was analysed. In addition, CP expression profiles were determined on T cells before and after MLC. Finally, the observed CP expression patterns were correlated with clinical subtypes, response to therapy and the extent of antileukemic activity.



Material and Methods

Sample Collection

WB was collected from AML pts (n=38) and from healthy donors (n=15) using lithium-heparin tubes. Samples were provided by the University Hospitals of Augsburg, Munich, Oldenburg and Tuebingen. Informed consent was obtained and the experiments were performed in accordance with the Helsinki protocol and the local ethics committee of Ludwig-Maximilians-University-Hospital Munich.

Patient characteristics

The median age of AML pts was 59 years (range 21-79) with a male:female ratio of 1.1:1. Healthy donors had a median age of 28 years (range 20-56), male:female ratio 1:1.5. AML was classified according to FAB-classification, subtype, cytogenetic risk and ELN prognostic categories [49]. An overview of the patient characteristics is given in Table 1.

Table 1: Patients' characteristics.

| Patient No. | Age | Sex | FAB | Stage | ELN risk | Response to induction therapy | Blast Phenotype (CD) | IC blasts (%) | Conducted Experiments | |
|-------------|-----|-----|--------|-----------------------|--------------|-------------------------------------|------------------------|------------------|---------------------------|--|
| | | | | | | AML | | | | |
| 1426 | 61 | f | s – M5 | first dgn | adverse | CR | 13, 33, 34, 64, 117 | 40 | DCC (WB, PBMNC), MLC, CTX | |
| 1430 | 79 | m | p – M5 | first dgn | favorable | NCR | 13, 33, 34, 117 | 70 | DCC (WB, PBMNC), MLC, CTX | |
| 1432 | 34 | m | p – M5 | first dgn | intermediate | CR | 13, 33, 34, 64 | 80 | DCC (WB, PBMNC), MLC, CTX | |
| 1434 | 61 | f | s – M? | first dgn | adverse | NCR | 7, 13, 33, 34, 64, 117 | 59 | DCC (WB, PBMNC), MLC, CTX | |
| 1439 | 61 | f | s – M5 | first dgn | favorable | ND | 13, 33, 34, 117 | 15 | DCC (WB, PBMNC) | |
| 1441 | 60 | m | s – M4 | first dgn | favorable | NCR | 13, 33, 64, 65, 117 | 81 | DCC (WB), MLC, CTX | |
| 1442 | 73 | f | s – M4 | first dgn | intermediate | CR | 33, 117 | 14 | DCC (WB, PBMNC), MLC, CTX | |
| 1443 | 64 | m | s – M? | first dgn | adverse | NCR | 13, 33, 34, 117 | 28 | DCC (WB, PBMNC) | |
| 1444 | 35 | f | p – M1 | first dgn | favorable | CR | 15, 33, 34, 65, 117 | 50 | DCC (WB), MLC, CTX | |
| 1447 | 21 | m | p – M5 | first dgn | intermediate | CR | 4, 33, 56 | 65 | DCC (WB), MLC, CTX | |
| 1449 | 78 | m | s – M? | first dgn | favorable | NCR | 15, 64, 65 | 62 | DCC (WB), MLC, CTX | |
| 1452 | 44 | m | p – M? | first dgn | intermediate | NCR | 13, 33, 34, 117 | 55 | DCC (WB), MLC, CTX | |
| 1454 | 60 | f | s – M? | first dgn | intermediate | ND | 33, 34, 117 | 33 | DCC (WB) | |
| 1459 | 54 | m | p – M4 | first dgn | favorable | CR | 33, 56, 64 | 14 | DCC (WB), MLC, CTX | |
| 1460 | 78 | f | p – M4 | first dgn | intermediate | CR | 15, 34, 117 | 68 | DCC (WB), MLC, CTX | |
| 1462 | 49 | f | p – M5 | first dgn | favorable | CR | 13, 33, 34, 56, 64 | 60 | DCC (WB) | |
| 1466 | 47 | f | p – M5 | first dgn | adverse | CR | 13, 15, 33, 34, 117 | 15 | DCC (WB, PBMNC), MLC, CTX | |
| 1471 | 39 | m | p – M1 | first dgn | adverse | NCR | 13, 33, 34, 117 | 69 | DCC (WB, PBMNC), MLC, CTX | |
| 1472 | 33 | f | p – M2 | first dgn | favorable | CR | 13, 15, 34, 117 | 30 | DCC (WB), MLC, CTX | |
| 1473 | 73 | m | p – M2 | first dgn | adverse | NCR | 13, 33, 34, 117 | 55 | DCC (WB, PBMNC), MLC, CTX | |
| 1480 | 66 | m | s – M? | first dgn | adverse | NCR | 13, 33, 117 | 38 | DCC (WB), MLC, CTX | |
| 1481 | 62 | f | p – M4 | first dgn | favorable | CR | 13, 56, 64, 117 | 29 | DCC (WB, PBMNC), MLC, CTX | |
| 1483 | 77 | m | p – M5 | first dgn | adverse | ND | 13, 33, 34, 64 | 93 | DCC (WB), MLC, CTX | |
| 1489 | 55 | f | p – M0 | first dgn | favorable | NCR | 13, 33, 34, 65, 117 | 58 | DCC (WB), MLC, CTX | |
| 1492 | 52 | f | s – M2 | first dgn | nd | NCR | 13, 33, 34, 117 | 42 | DCC (WB), MLC, CTX | |
| 1494 | 55 | f | p – M5 | first dgn | adverse | NCR | 13, 34, 117 | 88 | DCC (WB), MLC, CTX | |
| 1464 | 72 | m | s – M? | persisting disease | - | - | 20, 33, 34, 117 | 44 | DCC (WB), MLC, CTX | |
| 1467 | 59 | f | s – M? | persisting disease | - | - | 13, 33, 34, 117 | 30 | DCC (WB, PBMNC) | |
| 1468 | 66 | m | p – M? | persisting disease | - | - | 33, 34, 56, 117 | 75 | DCC (WB) | |
| 1463 | 60 | f | s – M? | rel | - | - | 13, 33, 34, 56 | 30 | DCC (WB), MLC, CTX | |
| 1469 | 49 | m | p – M4 | rel | - | - | 13, 33, 34, 117 | 94 | DCC (WB) | |
| 1475 | 77 | m | s – M? | rel | - | - | 13, 33, 34, 117 | 20 | DCC (WB, PBMNC), MLC, CTX | |
| 1486 | 77 | m | s – M1 | rel | - | - | 33, 34, 56 | 45 | DCC (WB) | |
| 1424 | 37 | f | s – M4 | rel after HSCT | - | - | 13, 14, 33, 117 | 30 | DCC (WB), MLC, CTX | |
| 1470 | 67 | m | p – M? | rel after HSCT | - | - | 33, 34, 56, 117 | 9 | DCC (WB, PBMNC), MLC, CTX | |
| 1474 | 70 | m | p – M? | rel after HSCT | - | - | 33, 34, 56, 117 | 80 | DCC (WB, PBMNC) | |
| 1476 | 63 | f | s – M? | rel after HSCT | - | - | 13, 33, 34, 65 | 20 | DCC (WB) | |
| 1482 | 75 | m | s - M? | rel after HSCT | - | - | 33, 64, 117 | 12 | DCC (WB), MLC, CTX | |

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| | Healthy | | | | | | |
|------|---------|---|--|----------------------|--|--|--|
| 1421 | 27 | f | | DCC (WB, PBMNC), MLC | | | |
| 1425 | 27 | m | | DCC (WB, PBMNC), MLC | | | |
| 1428 | 56 | f | | DCC (WB, PBMNC), MLC | | | |
| 1429 | 22 | f | | DCC (WB, PBMNC), MLC | | | |
| 1431 | 22 | m | | DCC (WB, PBMNC), MLC | | | |
| 1436 | 25 | m | | DCC (WB, PBMNC), MLC | | | |
| 1438 | 31 | f | | DCC (WB, PBMNC), MLC | | | |
| 1440 | 20 | f | | DCC (WB), MLC | | | |
| 1446 | 25 | m | | DCC (WB, PBMNC), MLC | | | |
| 1448 | 27 | f | | DCC (WB), MLC | | | |
| 1458 | 21 | f | | DCC (WB), MLC | | | |
| 1478 | 31 | m | | DCC (WB), MLC | | | |
| 1479 | 35 | f | | DCC (WB, PBMNC), MLC | | | |
| 1485 | 21 | f | | DCC (WB), MLC | | | |
| 1493 | 25 | m | | DCC (WB), MLC | | | |

f female; m male; p primary AML; s secondary AML; M? FAB type not classified; first dgn first diagnosis; rel relapse before or after HSCT (hematopoietic stem cell transplantation); nd no data; CR complete remission; NCR blast persistence; IC blasts immunocytologically determined blasts (Bold letters: blast markers used for quantification of blasts and DCleu); WB whole blood; PBMNC peripheral blood mononuclear cells; DCC dendritic cell culture; MLC mixed lymphocyte culture; CTX cytotoxicity fluorolysis assay.

Cell Culture with whole blood (WB) Isolation of PBMNCs and CD3⁺ T cells

PBMNCs were isolated from WB by density gradient centrifugation, washed with phosphate-buffered saline (PBS, Biochrom, Berlin, Germany) and used for CD3⁺T cell isolation via magnetic bead separation due to MACS-technology (Milteney Biotech, Bergisch Gladbach, Germany), reaching a T cell purity of Ø 84.99% (range 54.24 – 99.56%). Cells were cryopreserved in RPMI-1640-medium (Biochrom) with

dimethyl sulfoxide (DMSO, Sigma Aldrich Chemie GmbH, Steinheim, Germany) and fetal calf-serum (FCS, Biochrom, Berlin, Germany) at -80°C and thawed as needed [12].

Generation of DC/DC $_{\rm leu}$ from PBMNCs

DC/DC_{leu} were generated from PBMNCs of AML pts or healthy donors using DC/DC_{leu}-generating protocols for 'Pici1' and 'Pici2' containing specific combinations of response modifiers (further referred to as PBMNC^{DC(Pici1)}, PBMNC^{DC(Pici2)}) as given in Table 2 [13]. A culture without

Table 2: DC/DC_{leu}-generating protocol.

| DC/DC _{leu} Protocol | DC/D _{cleu} Source | Compo | osition | Time of addition (day (d)) | Time of Culture (days) | Reference | | |
|-------------------------------|-----------------------------|----------------------------|----------|----------------------------|---------------------------|--|--|--|
| Pici1 | | GM-CSF | 500 U/ml | d0 | | | | |
| | DDMMC | IL-4 | 250 U/ml | d0 | | [13,50] | | |
| | PBMNC | Picibanil | 10 µg/ml | d7 | 9-10 | | | |
| | | PGE₁ | 1 µg/ml | d7 | | | | |
| Pici2 | | GM-CSF | 500 U/ml | d0 | | | | |
| | PBMNC | IL-4 | 250 U/ml | d0 | | | | |
| | PBIVING | Picibanil PGE ₂ | 10 µg/ml | d7 | | | | |
| | | | 1 µg/ml | d7 | | | | |
| Kit-I | WB | GM-CSF | 800 U/ml | | | [13,50] European Patent No. 15 801 987.7 - 1118 US Patent 15-517627 MODIBLAST GmBH | | |
| | VVD | Picibanil | 10 µg/ml | | | | | |
| Kit-K | WB | GM-CSF | 800 U/ml | 40 40 3 | 7-8 | | | |
| | VVD | PGE ₂ | 1 µg/ml | d0, d2-3 | | | | |
| Kit-M | MD | GM-CSF | 800 U/ml | | | | | |
| | WB | PGE₁ | 1 μg/ml | | | | | |

Mode of Action:

GM-CSF: Induction of myeloid (DC-) differentiation; IL-4: Induction of DC-differentiation; Picibanil: Danger signaling, DC maturation; PGE, / PGE2: Danger signaling, DC maturation

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added response modifiers served as control (PBMNC^{DC(Control)}). Therefore, 3-4x10⁶ isolated PBMNCs were pipetted in 12-multiwell-plates and diluted with 2ml X-Vivo-15-medium (Lonza, Basel, Switzerland).

Generation of DC/DC_{leu} from WB

DC/DC_{leu} were also generated directly from WB using DC/DC_{leu}-generating protocols 'Kit-I', 'Kit-K' and 'Kit-M' [50] given in Table 2. 500 μ l WB (corresponding to 5.0 – 30.3 x 10⁶ PBMNC) were cultured in 24-multiwell culture plates (ThermoFisher Scientific) and diluted with 500 μ l X-Vivo-15-medium. A culture without added response modifiers was served as a control (WB^{DC(Control)}).

All cell culture experiments were conducted at standard laboratory conditions comprising 37°C, 21% O₂ and 5% CO₂.

Cell-characterization by flow cytometry

Frequencies, subsets and phenotypes of leukemic blasts, T cells, B cells, monocytes and DC/DC_{leu} were quantified by flow cytometric analyses were done as previously described [12]. Therefore cells were stained with monoclonal antibodies (moAbs) labelled with Fluorescein isothiocyanat (FITC), phycoerythrin (PE), tandem Cy7-PE conjugation (Cy7-PE) or allophycocyanin (APC). Dead cells were excluded using 7AAD^b. Erythrocytes in WB were lysed prior to staining. Staining was performed in PBS containing FCS and the corresponding moAbs for 15 min in the dark.

Cell-Evaluation and quantification was done via fluorescence-activated cell sorting Flow-Cytometer (FACSCaliburTM) and the CellQuestPro-acquisition and analysis software (Becton Dickson, Heidelberg, Germany) with appropriate isotype controls.

Leukemic blasts, DC and DC $_{leu}$ were analysed before and after cell culture using a refined gating strategy. DC $_{leu}$ were defined by co-expression of at least one blast marker (CD15, CD34, CD65, CD117) including lineage-aberrant markers (CD56) and one or two DC markers (CD80, CD206) absent on naïve blasts. DC/DC $_{leu}$ maturation was assessed by CCR7 expression. Subgroup analyses required $\geq 10\%$ DC in the total cell fraction.

Mixed-lymphocyte-culture (MLC) of T cell-enriched immune-reactive cells with Pici/Kit-treated-cell-suspensions

Autologous T cells were stimulated with DC/DC_{leu} generated from WB (WB^{DC}) generating T cell-enriched immunoreactive cells. Therefore, 1x10⁶ autologous CD3⁺ T cells ('effector cells') were co-cultured with a stimulator cell suspension containing 2.5 x 10⁵ DC/DC_{leu} in 24-mulitwell-tissue-culture plates (ThermoFisher Scientific, Darmstadt, Germany) and diluted in RPMI-1640 medium containing 100 U/ml Penicillin (Biochrom, Berlin, Germany) and 15%

human serum. 50 U/ml Interleukin 2 (IL-2, PeproTech, Berlin, Germany) was added on day 0 and day 2-3. Cells were harvested after 6-8 days (further referred to as WB^{DC-MLC}) and used for the cytotoxicity-fluorolysis-assay. T cell subsets were quantified by flow cytometry before and after the MLC [12].

Cytotoxicity fluorolysis assay

A fluorolysis assay was done to analyse the lytic activity of T cell-enriched immunoreactive cells against leukemic blasts cells [11,50]. Therefore, effector cells (WB^{DC-MLC}) containing 1 x 10⁶ T cells and 1 x 10⁶ thawed PBMNCs (blast containing target cells) were co-cultured in RMPI-1640 medium with 100 U/ml penicillin and 15% human serum for 3 and 24 hours. Controls consisted of effector- and target-cells cultured separately and combined only prior to analysis.

Target cells were pre-stained with blast specific moAbs. After incubation, cells were resuspended in PBS containing 7 AAD and a defined number of Fluorospheres Beads (Beckman coulter, Krefeld, Germany) to quantify viability and cytotoxicity. Data were acquired using flow cytometry with a refined gating strategy [11].

Statistical Methods

Data is presented as mean \pm standard-deviation-values. Statistical comparisons of two groups were performed using the two-tailed t-test in cases with normal distribution or with the Mann-Whitney-Wilcoxon-Test in cases with no normal distribution. Statistical analyses and figures were performed with Microsoft Excel 2010 ® (Microsoft, Redmond, Washington, USA) and GraphPad Prism8© (GraphPad Software, California, USA).

Differences were considered as 'not significant' in cases with p values > 0.1, as 'borderline significant' with p values between 0.1 and 0.05 and as 'significant' with p values < 0.05.

Results

Prolog

In the first part of this study, we generated DC/DC_{leu} with the DC/DC_{leu}-generating Kits for Kit-I, Kit-K and Kit-M from leukemic WB or blood from healthy probands as we did in analogy for PBMNC with Pici-generating methods (Pici1, Pici2). We also cultivated WB or PBMNC without addition of any response modifiers as a control. Afterwards we stimulated T cell enriched immunoreactive cells with Pici/Kit treated cells and carried out a cytotoxicity fluorolysis assay (CTX).

In the second part of this study, we analysed the expression of CP on blasts, DC or DC_{leu} before and after DC/DC_{leu}-generation. In addition, we analysed the expression of CP on T cells after MLC. Conclusively, we correlated the CP expression on immunoreactive cells with the anti-leukemic

| Group | Subgroup | Surface marker | Refers to | Abbreviation | References |
|---------------|---|--|-------------|--|------------|
| Blast cells | blasts | BLA ⁺ e.g. CD34 ⁺ , CD117 ⁺ | WB or PBMNC | BLA/WB or /PBMNC | [13] |
| | dendritic cells | DC+e.g. CD80+, CD206+ | WB or PBMNC | DC/WB or /PBMNC | [13] |
| DC | leukemia derived DC | DC+BLA+ | WB or PBMNC | DC _{leu} /WB or /PBMNC | [13] |
| | mature DC | DC⁺CCR7⁺ | WB or PBMNC | DC _{mat} /WB or /PBMNC | [50] |
| Monocytes | CD14⁺ monocytes | CD14⁺ | WB or PBMNC | mo ₁₄₊ /WB | [50] |
| Taalla | CD3 ⁺ T cells | CD3⁺ | WB or PBMNC | T/WB or /PBMNC | [11] |
| T cells | non-naive T cells | CD3⁺CD45RO⁺ | CD3 | T _{non-naive} / T | [11] |
| | CD279⁺ expressing blasts | Bla⁺CD279⁺ | Bla | Bla ₂₇₉₊ /Bla | [51] |
| | CD279 ⁺ expressing monocytes | CD14+ CD279+ | CD14 | mo ₁₄₊₂₇₉₊ /mo ₁₄₊ | [52] |
| | CD279⁺ expressing DC _{leu} | DC⁺Bla⁺CD279⁺ | Bla | DC _{leu279+} /Bla | |
| | CD279 ⁺ expressing T cells | CD3⁺CD279⁺ | CD3 | T ₂₇₉₊ / T | [53] |
| | CD274⁺ expressing blasts | Bla⁺CD274⁺ | Bla | Bla ₂₇₄₊ /Bla | [51] |
| CP expressing | CD274 ⁺ expressing monocytes | CD14⁺ CD274⁺ | CD14 | mo ₁₄₊₂₇₄₊ /mo ₁₄₊ | [54] |
| cells | CD274 ⁺ expressing DC _{leu} | DC+Bla+CD274+ | Bla | DC _{leu274+} /Bla | |
| | CD274⁺ expressing T cells | CD3⁺CD274⁺ | CD3 | T ₂₇₄₊ / T | [53,55] |
| | CD273⁺ expressing blasts | Bla⁺CD273⁺ | Bla | Bla ₂₇₃₊ /Bla | [51] |
| | CD273 ⁺ expressing monocytes | CD14+ CD273+ | CD14 | mo ₁₄₊₂₇₃₊ /mo ₁₄₊ | [54] |
| | CD273⁺ expressing DC _{leu} | DC+Bla+CD273+ | Bla | DC _{leu273+} /Bla | |
| | CD273 ⁺ expressing T cells | CD3 ⁺ CD273 ⁺ | CD3 | T ₂₇₃₊ /T | [53] |

Table 3: Cells and cell subsets as evaluated by flow cytometry.

cytotoxic activity, the patients' age, sex, ELN risk groups and patients' response to induction chemotherapy. Abbreviations of all cell types are given in Table 3.

Expression of CP on uncultured AML patients' blasts or T cells or on healthy donors' monocytes or T cells

We determined the expression of several CP (CD274, CD279 and CD273) on leukemic blasts in uncultured WB of AML pts as well as frequencies of CP expressing monocytes in uncultured healthy WB. In addition, we quantified CP expressing T cells in leukemic WB and in WB of healthy probands (Figure 1).

CP expressing blasts and healthy monocytes

We quantified CD274, CD279 and CD273 expressing leukemic blasts in uncultured WB: We found low frequencies of CD273⁺ leukemic blasts (most times < 10% CD273⁺ blasts). In only 3 out of 37 cases we found more than 20% CD273 expressing blasts. We found higher frequencies of CD279 expressing leukemic blasts (on approximately 50% of blasts). In 25 of 37 cases, we observed more than 20% CD279-expressing blasts. We found also high frequencies of CD274 expressing blasts in uncultured WB (Figure 1A).

We found low frequencies of CP (CD273, CD279 and CD274) expressing uncultured monocytes in healthy samples (Figure 1B).

CP expressing uncultured T cells from AML patients and healthy probands

We found low frequencies of CD273 and CD274 expressing uncultured T cells in leukemic WB as well as in WB of healthy probands. In only 1 out of 24 cases we noticed more than 20% CD273 respectively CD274 expressing T cells in AML patients' WB. In contrast, higher frequencies of CD279 expressing T cells were found in healthy (Ø 45%) as well as in AML patients' samples (Ø 58%) (Figure 1C, Figure 1D).

DC/DC_{leu} generation from healthy and leukemic PBMNC

Significantly higher frequencies of DC were generated with Pici1 and Pici2 compared to control from leukemic as well as from healthy PBMNC. We could generate DC or DC_{leu} in PBMNC from healthy and leukemic samples with both Pici1 and Pici2 vs. untreated control cultures, thereby confirming data shown before [13] (Figure 2A, B).

DC/DC_{leu} generation from healthy and leukemic WB

We evaluated the effect of the several Kits on the generation of DC/DC_{len} for leukemic as well as healthy WB.

WB samples from AML patients (C) and healthy probands (D) were cultured with or without Kits for 7 days. In parallel PBMNC samples from AML patients (A) and



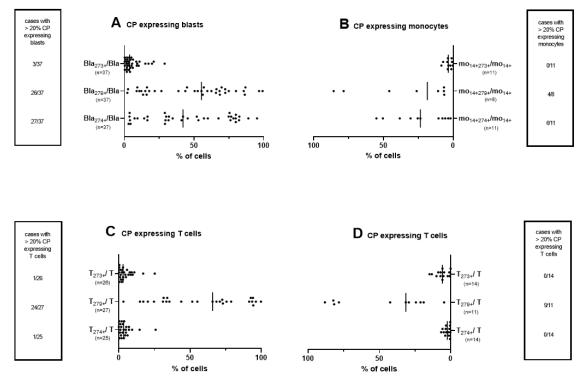


Figure 1: Expression of CP (CD274, CD279 and CD273) on uncultured blasts from AML patients (left side), monocytes from healthy probands (right side) or on T cells from AML patients and healthy probands.

Given are median frequencies of CP (CD274, CD279 and CD273) expressing uncultured leukemic blasts from AML pts (**A**) or monocytes from healthy probands (**B**) as well as T cells from pts with AML (**C**) or from healthy probands (**D**) in uncultured WB coexpressing CP markers. Each dot plot represents one individual sample. Abbreviations of all cell types are given in Table 3.

healthy volunteers (**B**) were cultured with or without Picis for 9 days. Given are mean frequencies \pm standard deviation of generated DC- and DC subgroups in WB/PBMNC. Statistical analyses were conducted using multiple t-test: Differences were considered as highly significant with p values <0.005 (**) and as significant with p values <0.05 (*). Abbreviations of all cell types are given in Table 3.

Significantly higher frequencies of DC were generated with Kit-I, Kit-K and Kit-M compared to control from leukemic as well as from healthy WB. We observed significantly higher frequencies of DC_{leu} with Kit-I, Kit-K and Kit-M compared to Kit-Control for leukemic WB. In summary, we were able to generate significantly higher frequencies of DC and DC_{leu} with immunomodulatory Kits from healthy and leukemic WB as compared to controls thereby confirming data shown before [13,50] (Figure 2C, D). The proliferation of blasts was not induced under Kit treatment (data not shown).

Expression of CP on AML patients' blasts and DC_{leu} after DC/DC_{leu} -culture

We evaluated CP (CD274, CD279 and CD273) expressing blasts and DC_{leu} of AML patients' WB after DC/DC_{leu}-generation with 'Kits' (WB^{DC(I)}, WB^{DC(K)}, WB^{DC(M)}) compared to control (WB^{DC(control)}) (Figure 3A). In addition, we give

the relative changes of CP (CD274, CD279 and CD273) expressing leukemic blasts and DC $_{\rm leu}$ as percentual differences ('delta' (Δ %)) for Kit-I, Kit-K and Kit-M compared to control (Figure 3B). Overall, we found lower frequencies of CP expressing blasts after DC/DC $_{\rm leu}$ -culture for all three Kits and significantly lower frequencies of CD279 expressing blasts for WB^{DC(K)} compared to control.

Lower frequencies of CP (CD279 and CD274) expressing DC_{leu} were seen after the influence of Kit-I, Kit-K and Kit-M. Borderline significantly lower frequencies of Bla_{279+}/DC_{leu} were found for Kit-K compared to control.

Relative changes of CP (CD274, CD279 and CD273) expressing leukemic blasts and DC_{leu} compared to control are given in Figure 3B. Overall, we show in almost all cases a decrease in CP expression on blasts as well as on DC_{leu} compared to control.

Stimulatory impact of Kit pre-treated WB on T cells in MLC

To evaluate the potential stimulating effect of generated DC/DC_{leu} on the composition of immunoreactive cells, T cell compositions were evaluated before (uncultured MLC) and after stimulation with Kits-treated WB (WB^{MLC(I)}, WB^{MLC(K)},



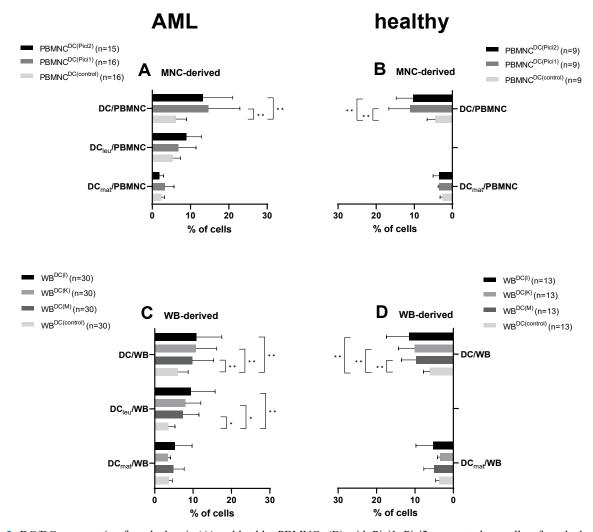


Figure 2: DC/DC_{leu}-generation from leukemic (A) and healthy PBMNCs (B) with Pici1, Pici2 vs. control as well as from leukemic (C) and healthy WB (D) with Kit-I, Kit-K, Kit-M vs. control.

WB samples from AML patients (\mathbf{C}) and healthy probands (\mathbf{D}) were cultured with or without Kits for 7 days. In parallel PBMNC samples from AML patients (\mathbf{A}) and healthy volunteers (\mathbf{B}) were cultured with or without Picis for 9 days. Given are mean frequencies \pm standard deviation of generated DC- and DC subgroups in WB/PBMNC. Statistical analyses were conducted using multiple t-test: Differences were considered as highly significant with p values <0.005 (**) and as significant with p values <0.005 (*). Abbreviations of all cell types are given in Table 3.

 $WB^{MLC(M)}\ WB^{MLC(control)}$). Significantly higher frequencies of $T_{non-naive}$ cells after Kit-M treatment could be found in comparison to control. We could also find borderline significantly increased $T_{non-naive}$ cells in Kit-K pre-treated samples (Figure 4A).

Expression of CP on AML patients' T cells after stimulation in MLC with Kit pre-treated WB

Frequencies of CP expressing T cells were analysed after stimulation in MLC with Kit pre-treated WB of AML patients and also of healthy probands (Figure 4A). We quantified frequencies of CP (CD274, CD279, CD273) expressing T cells in AML patients' WB^{MLC(I)}, WB^{MLC(K)}, WB^{MLC(M)} compared to WB^{MLC(control)}. We observed comparable frequencies of T₂₇₃₊/T

and T₂₇₄₊/T in WB^{MLC(I)}, WB^{MLC(K)} and WB^{MLC(M)} compared to WB^{MLC(control)} for leukemic and healthy WB. We could detect lower frequencies for T₂₇₉₊/T in Kit pre-treated WB of healthy probands compared to WB^{MLC(control)}. Although we observed slightly lower frequencies of CP expressing T cells in Kit pre-treated WB compared to WB^{MLC(control)} of AML patients as well as healthy probands, results did not differ significantly (Figure 4A).

In addition, we compared the relative changes of CP (CD274, CD279 and CD273) expressing T cells (Figure 4B) Results are given as percentual differences (Δ%) of CP expressing T cells in WB^{MLC(K)}, WB^{MLC(K)}, WB^{MLC(K)} in proportion to WB^{MLC(control)}. We could not find any significant differences between the individual values.

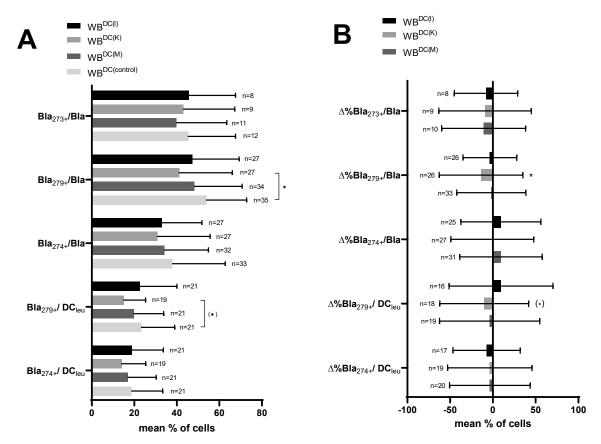


Figure 3: CP (CD274, CD279 and CD273) expressing blasts and DC_{leu} from AML patients after DC/DC_{leu}-generation.

Given are mean frequencies \pm standard deviation of activated non-naive T cells and CP (CD274, CD279 and CD273) expressing leukemic or healthy T cells after stimulation in MLC with Kit pre-treated WB (WB^{MLC(I)}, WB^{MLC(K)}, WB^{MLC(K)}) compared to untreated WB (WB^{MLC(control)}) (A). Given are the relative changes of non-naive T cells and CP (CD274, CD279 and CD273) expressing T cells as percentual differences ('delta' (Δ %)) for Kit-I, Kit-K and Kit-M compared to control (B). Statistical analyses were conducted using multiple t-test: Differences were considered as significant with p values <0.05 (*) and as borderline significant ((*)) with p values 0.10 to 0.05. Abbreviations of all cell types are given in Table 3.

To evaluate the anti-leukemic cytotoxicity of DC/DC_{leu}stimulated immunoreactive cells, we analysed the lytic activity of WBMLC(I), WBMLC(K), WBMLC(M) and WBMLC(control) through CTX after 3 h and after 24 h of incubation of effector and leukemic target cells. After 3 h we could see a lysis of target cells in WBMLC(I) in about 30.8% of cases, whereas we could observe a lysis of target cells in $WB^{\text{MLC}(K)}$ and $WB^{\text{MLC}(M)}$ in more than 60% of cases. In the control group we observed a blast lysis in 40% of cases. After 24 h, more cases of WBMLC(I), WBMLC(K), WBMLC(M) and WBMLC(control) attained lysis. in which WB^{MLC(M)} showed the highest percentage of cases with lysis (94.7%) compared to WBMLC(control) (75% of cases) (Figure 5A). Selecting the best achieved result after 3h or 24h, we discovered blast lysis in 100% of cases for WBMLC(K) and $WB^{\text{MLC(M)}}$. $WB^{\text{MLC(I)}}$ and $WB^{\text{MLC(control)}}$ achieved lysis in about approximately 80% of cases (Figure 5A). Improved lysis, defined as the relative improvement of blast lysis of Kit treated vs. not pre-treated cases (WBMLC(control)), was observed after 3h in 38.5% of cases for WBMLC(I) and in more than 60%

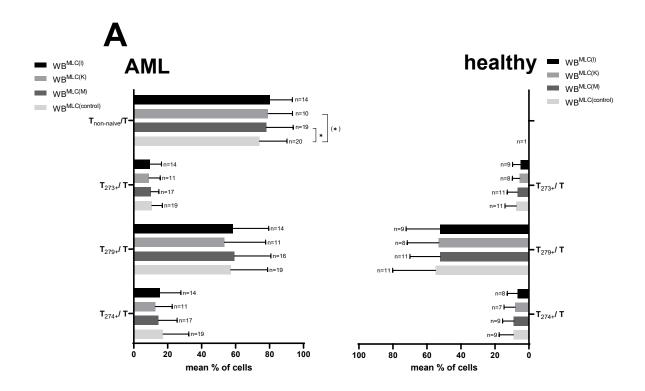
of cases for WB^{MLC(K)} and WB^{MLC(M)}. After 24h more than 50% of cases for WB^{MLC(I)} and WB^{MLC(M)} showed an improved lysis. Selecting the best achieved result after 3h or 24h, we found improved blast lysis in almost 90% of cases for WB^{MLC(M)}. WB^{MLC(I)} and WB^{MLC(K)} showed improved lysis in more than 60% of cases (Figure 5B).

We observed highest frequencies of lysed blasts especially for Kit-K and Kit-M after 24h leading to best improved lysis compared to the control thereby confirming data shown before (Figure 5C) [13,50].

Correlation analyses

Correlation of frequencies of CP expressing uncultured cells with patients' clinical data

We correlated frequencies of CP expressing uncultured blasts or T cells in samples from AML patients with patients' clinical data. We did not find correlations between frequencies of uncultured CP expressing blasts or T cells with patients' sex, ELN risk or response vs. no response to



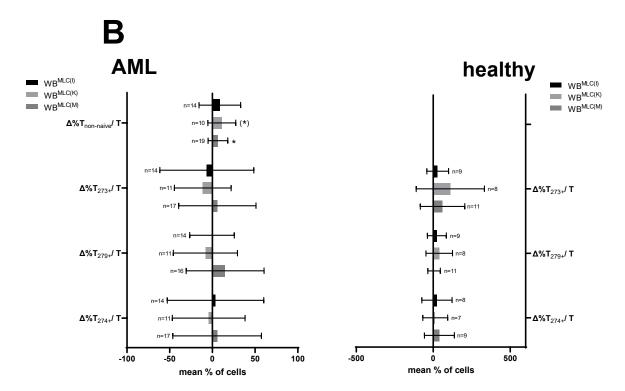


Figure 4: Activated non-naive T cells and CP (CD274, CD279 and CD273) expressing T cells after MLC with Kit pre-treated vs. untreated WB. Given are mean frequencies ± standard deviation of activated non-naive T cells and CP (CD274, CD279 and CD273) expressing leukemic or healthy T cells after stimulation in MLC with Kit pre-treated WB (WB^{MLC(I)}, WB^{MLC(K)}, WB^{MLC(M)}) compared to untreated WB (WB^{MLC(control)}) (A). Given are the relative changes of non-naive T cells and CP (CD274, CD279 and CD273) expressing T cells as percentual differences ('delta' (Δ%)) for Kit-I, Kit-K and Kit-M compared to control (B). Statistical analyses were conducted using multiple t-test: Differences were considered as significant with p values <0.05 (*) and as borderline significant ((*)) with p values 0.10 to 0.05. Abbreviations of all cell types are given in Table 3.

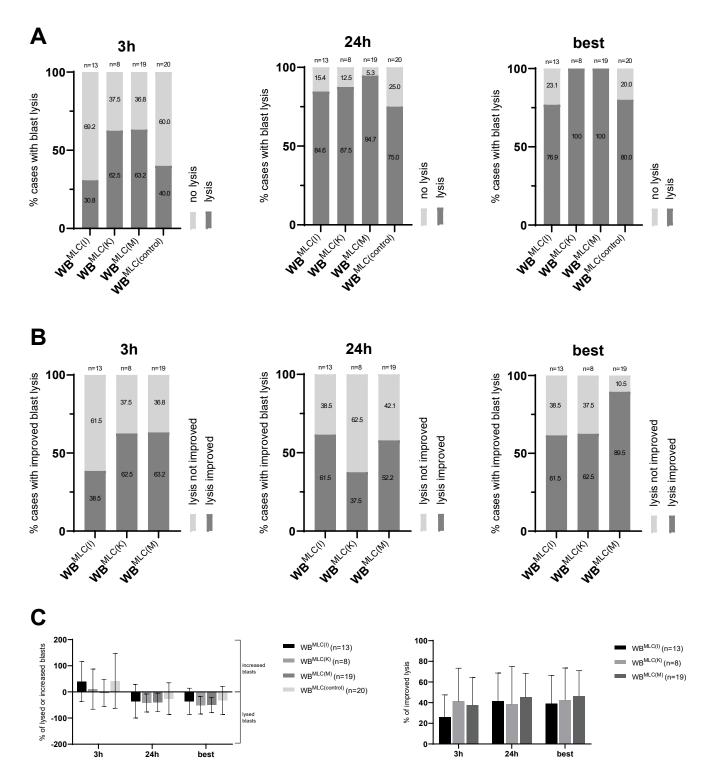


Figure 5: Stimulatory effect of DC/DC_{leu} generated leukemic WB on the cytotoxic activity of immunoreactive cells after MLC as measured by CTX.

Given are the proportions of cases with blast lysis (**A**) and the mean \pm range of lysed or increased blasts in WB^{MLC(K)}, WB^{MLC(K)}, WB^{MLC(K)} and WB^{MLC(Control)} after 3 h, 24 h and the "best" achieved blast lysis after 3 h or 24 h (**C**); the proportions of cases with an improved blast lysis (**B**) in WB^{MLC(K)} and WB^{MLC(K)} in relation to WB^{MLC(control)} after 3 h, 24 h and the "best" achieved improved blast lysis after 3 h or 24 h. Abbreviations of all cell types are given in Table 3.



induction chemotherapy, although frequencies of T_{279+}/T were (significantly) higher in cases with favourable ELN risk as well as in responders to induction chemotherapy (Figure 6).

Correlation of frequencies of CP expressing uncultured blasts or T cells with improved blast lysis after MLC with Kit pre-treated WB

We correlated frequencies of CP expressing blats and T cells in uncultured WB samples from AML patients with achieved improved blast lysis after Kit pre-treated vs. not pre-treated MLC. We found no correlations between frequencies of CP expressing blasts or T cells and achieved improved blast lysis with Kit-I or Kit-M pre-treated WB after MLC. Kit-K results could not be correlated due to low case numbers (Figure 8).

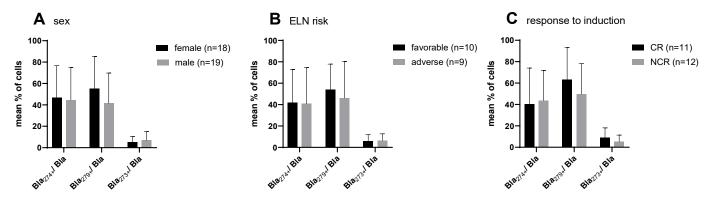
Correlation of frequencies of CP expressing blasts or T cells after DC/DC₁₋₁-generation with improved blast lysis

after MLC with Kit pre-treated WB

We observed (non-significant) negative correlations between improved blast lysis and relatively changed frequencies of CP expressing blasts after MLC with Kit-I or Kit-M pre-treated WB vs. control (Figure 9). Noteworthy, we found significant positive correlations between frequencies of CP expressing T cells after MLC and improved blast lysis via CTX for Kit-M pre-treated samples (Figure 9). An examination of T cells after MLC for Kit-I and Kit-K was not possible due to low case numbers.

Correlation of relative changed frequencies of CP (CD274, CD279 and CD273) expressing leukemic blasts and T cells from AML pts with the relative improvement of blast lysis (= improved blast lysis) in Kit-I and Kit-M treated WB after MLC. Correlation coefficients (r) and p-values (one-tailed) are given, evaluated by Pearson correlation analyses. Abbreviations of all cell types are given in Table 3.

CP expressing blasts



CP expressing T cells

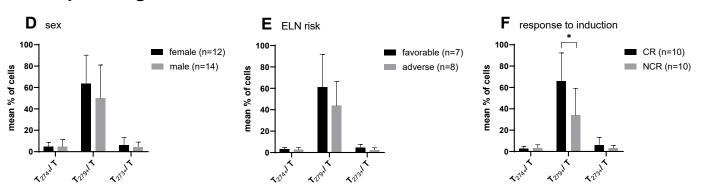
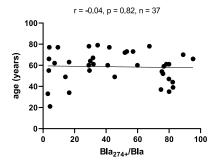


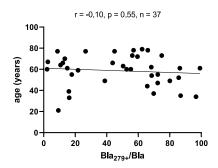
Figure 6: Correlation of frequencies of CP expressing uncultured blasts or T cells with patients' sex, ELN risk group & response to induction chemotherapy.

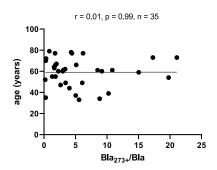
Given are mean frequencies \pm standard deviation of CP (CD274, CD279 and CD273) expressing uncultured leukemic blasts and T cells from AML pts subdivided into sex (female vs. male) (**A**, **D**), ELN risk groups (favorable vs. adverse) (**B**, **E**) and patients' response to induction chemotherapy (complete remission vs. blast persistence) (**C**, **F**). Differences were considered as significant with p values <0.05 (*). Abbreviations of all cell types are given in Table 3.



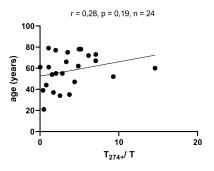
CP expressing blasts

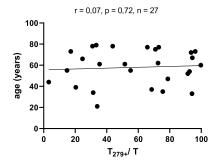






CP expressing T cells





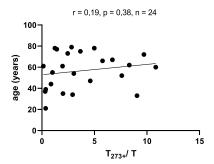


Figure 7: Correlation of frequencies of CP expressing uncultured blasts or T cells with patients' age.

Correlation of CP (CD274, CD279 and CD273) expressing uncultured leukemic blasts and T cells from AML pts with the patients' age. Correlation coefficients (r) and p-values (one-tailed) are given, evaluated by Pearson correlation analyses. Abbreviations of all cell types are given in Table 3.

Discussion

Immunotherapy of AML

Recent immunotherapeutic strategies for AML target leukemic blasts either through passive approaches, such as antibody- or cell-based therapies, or through immune (re)activating approaches including vaccines or immune checkpoint blockade [56].

 $\mathrm{DC}_{\mathrm{leu}}$ -based immunotherapies mediate leukemia-specific immune responses using individual patients' ex vivo generated or in vivo produced $\mathrm{DC/DC}_{\mathrm{leu}}$ (e.g. [12,57,58]). We confirm, that $\mathrm{DC/DC}_{\mathrm{leu}}$ generation from leukemic PBMNC or WB is successful using Pici-methods (Pici1, Pici2) or blastmodulatory Kits (Kit-I, Kit-K, Kit-M) [12,13,18]. No statistically significant differences were detected between the individual Kits or Pici protocols. However, a trend favoring Kit-M was observed, showing slightly higher efficiency in $\mathrm{DC/DC}_{\mathrm{leu}}$ generation compared to Kit-I and Kit-K (Figure 1).

DC/DC_{leu} generation and Antileukemic cytotoxicity

We could also confirm, that Pici or Kit pre-treated

PBMNC or WB leads to improved T/immune cell activation (Figure 4A), resulting in (improved) blast lysis, as shown by CTX assay after MLC [13,50,57] (Figure 5). These cytotoxic effects appear to involve distinct mechanisms: a rapid perforin/granzyme-mediated pathway, likely responsible for blast lysis after 3 hours of incubation, and a delayed Fas/FasL-dependent pathway becoming effective after 24 hours [11,12]. Ongoing studies aim to determine whether DC/DC_{leu} generation out of leukemic blasts can be achieved using Kits in vivo [58].

CP expressing blasts or monocytes and T cells in AML

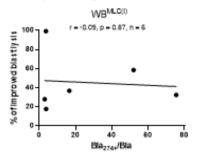
The expression of CD274, CD279 and CD273 was analysed on uncultured leukemic blasts and on monocytes from healthy probands. In addition, we quantified CP expressing T cells in leukemic WB and in WB of healthy probands (Figure 1).

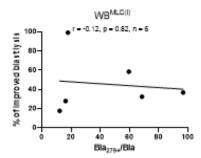
CP are known for their immunoregulatory properties. We found higher frequencies of CP expressing blasts for all three CP markers (CD274, CD279 and CD273) compared

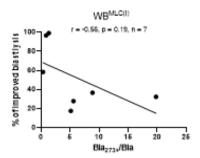


Kit-I

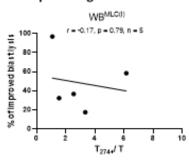


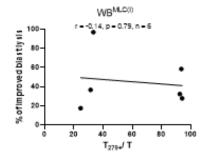


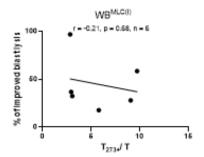




CP expressing T cells

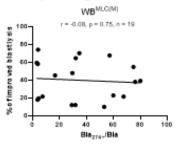


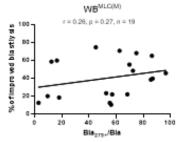


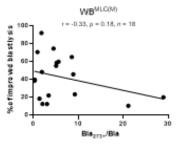


Kit-M

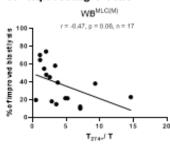
CP expressing blasts

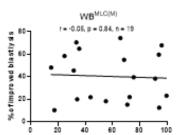






CP expressing T cells





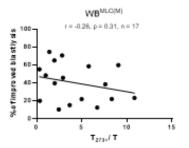


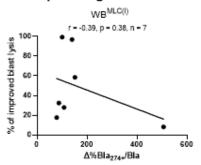
Figure 8: Correlation of achieved 'improved blast lysis' with frequencies of uncultured CP expressing blasts or T cells

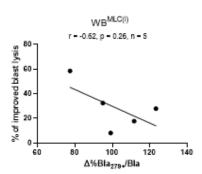
Correlation of CP (CD274, CD279 and CD273) expressing uncultured leukemic blasts and T cells from AML pts with the relative improvement of blast lysis (= improved blast lysis) in Kit-I and Kit-M treated WB after MLC. Correlation coefficients (r) and p-values (one-tailed) are given, evaluated by Pearson correlation analyses. Abbreviations of all cell types are given in Table 3.



Kit-I

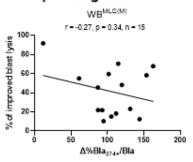
CP expressing blasts

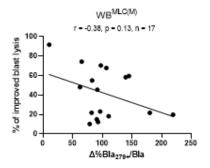




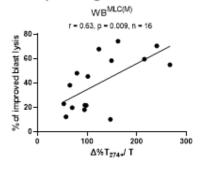
Kit-M

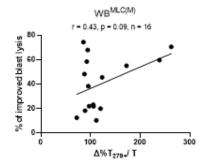
CP expressing blasts





CP expressing T cells





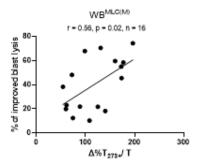


Figure 9: Correlation of achieved 'improved blast lysis' with frequencies of CP expressing blasts or T cells after MLC

Correlation of relative changed frequencies of CP (CD274, CD279 and CD273) expressing leukemic blasts and T cells from AML pts with the relative improvement of blast lysis (= improved blast lysis) in Kit-I and Kit-M treated WB after MLC. Correlation coefficients (r) and p-values (one-tailed) are given, evaluated by Pearson correlation analyses. Abbreviations of all cell types are given in Table 3.

to frequencies of CP expressing monocytes of healthy probands (Figure 1). It is known, that CD279 expression is higher on blast and T cells in newly diagnosed and relapsed AML patients compared to healthy controls [35,56,59-61]. Frequencies of CD279 expressing monocytes seems to be significantly higher in renal cell carcinoma patients comparing to healthy probands [52]. Royer et al. [54] showed varying and modified expression rates of CD279 expressing monocytes under stress condition (e.g. in vitro stimulation, hip fracture or sepsis) although CP expression between young and old patients did not differ in the basal state. Previous studies showed high frequencies of CD279 expressing monocytes in patients with hepatocellular carcinoma, but did not show CD279 expression in monocyte subsets of healthy probands [62].



Expression of CD279 on T cells was shown to increase with disease progression [56]. Confirming data shown before, we found higher frequencies of CD279 expressing T cells in AML patients' samples compared to healthy controls (Figure 1), thereby confirming data shown before [56,63,64]. Previous studies also showed variable, but approximately 60% CD279 coexpressing healthy memory CD8 T cells [65]. In patients with lung cancer low frequencies of naïve T cells show low (~1%) whereas high frequencies of central memory and effector memory T cells (40–60%) coexpressing CD279 were found [66-68]. With aging, expression of CD279 is increased on CD8 T cells as well as on CD4 T cells [69].

CD274 is expressed by hematopoietic cells such as DC or monocytes and non-hematopoietic cells [20,70]. The expression of CD274 changes over time during an AML disease: it is only minimally expressed by leukemic cells at the time of initial diagnosis but significantly upregulated as the disease progresses [71]. The varying expression rates over time on blasts could be explained by varying frequencies of CD274 expressing uncultured blasts. We found rather low frequencies of CD274 expressing monocytes in healthy probands, thereby confirming previous results [72,73] (Figure 1).

We found low expression levels of CD273 on leukemic blasts or monocytes as well as T cells from healthy or leukemic blood donors (Figure 1), thereby confirming previous data, showing low expression on DC, macrophages or activated T cells [74].

The rather low frequencies of CP expressing monocytes, found by us, could be explained by differential expression of CP within monocyte subtypes [54]. The functionality of CP within monocytes is currently still poorly understood.

An interesting approach could be to reactivate patients' immune system against blasts by modulating CP expression or functionality. To investigate the influence of Kit-treatment on CP expression, we compared frequencies of CP expressing blasts after DC/DC_{leu}-generation as well as T cells' CP expression after MLC. Overall, we found (significantly) lower expression levels of CD274, CD279 and CD273 on blasts after Kit-treatment compared to control (Figure 3).

We confirm with our data, that Pici or Kit pre-treated AML (MNC or WB) samples lead to increased frequencies of mature/leukemic derived DC vs. control and moreover to increased frequencies of activated T cells leading to improved blasts lysis vs. control (Figure 2, Figure 4, Figure 5) [12,13].

We found comparable frequencies of CP expressing T cells from AML patients' after MLC with Kit pre-treated compared to untreated WB (Figure 4), demonstrating that Kits pre-treatment of leukemic WB does not increase frequencies of CP expressing T cells after MLC compared to control. This

could also mean, that CP expression on T cells is not centrally involved in (Kit mediated) antileukemic reactions.

Current studies have indicated the potential clinical benefits of ICI against solid tumors as well as AML [38]. Also a combination with intensive chemotherapy, hypomethylating agents or other targeted therapies is possible and lead to improved outcome compared to monotherapy with ICI [35,36]. Moreover, several new co-inhibitory pathways are currently studied for their potential impacts on improving anti-tumor immune responses [34].

Correlations

We confirm data, that Kit pre-treatment of leukemic blood increases blast lysis after MLC (Figure 5). That is why we correlated frequencies of CP expressing T cells/blasts before and after DC or MLC-culture with several clinical and functional parameters. Although some studies showed that high frequencies of CD279 as well as CD274 expressing blasts in AML patients are associated with a poor prognosis [23,64,75], we did not observe significantly different frequencies of CD279 or of CD274 expressing blasts for AML patients concerning to their ELN risk group (Figure 6), thereby conforming data of Zajac et al. [75], who did not identify significant differences in overall survival between groups with high or low CD274 expression [75].

In contrast to data from Zajac et al. [75], we did not show positive correlations between age and CP expression [75]. We can add in addition, that frequencies of CP expressing leukemic blasts or T cells were independent of patients' age, sex or ELN risk group (Figure 6, 7). In contrast to data published before [76], we did not find a correlation between frequencies of CP expressing blasts with patients' response to induction therapy, but in contrast lower frequencies of T_{279+}/T in AML patients with no response to induction chemotherapy (Figure 6).

Whereas we did not find significant correlations between frequencies of uncultured blasts or T cells coexpressing CP with later on achieved improved blasts lysis (mediated by Kit pre-treated WB) (Figure 8), we found a negative correlation of CP expressing blasts in Kit treated samples (in relative to control) with improved blasts lysis and in Kit-M pre-treated samples (in relative to control) with improved blast lysis (Figure 9). These findings could be explained by the inhibitory effects of CP expressing blasts leading to a decreased T cell activation [74,77]. Especially CD279 is a known as a T cell inhibitor [78,79]. Inhibition of immune cells like DC_{len} through CD279 significantly impairs antigen presentation required to stimulate adaptive immune responses [52]. Previous results suggest that both CD279 and CD274 DC are capable to inhibit T cell response by regulating effector functions of T cells and antitumor immunity [80]. Tumor growth can be effectively suppressed following the transfer of CD279-deficient DC [80].



Surprisingly, there was a significant positive correlation found between improved blast lysis and the frequencies of CP expressing T cells after MLC for Kit-M (Figure 9). These data point to a gained immune activating antileukemic role of CP expressing T cells achieved after the influence of Kit-M, as CD279-expressing DC modulate T cell responses directly in the tumor microenvironment [80]. The enhancement of antileukemic activity, particularly observed ex vivo with Kit-M, could also occur via a CP-independent mechanism.

Future directions

Our results show increased frequencies of CP expressing blasts and T cells in the blood of AML patients. Kit pretreatment of patient WB, particularly with Kit-M, followed by MLC, resulted in enhanced blast lysis without increasing the frequency of CP expressing blasts, DC and T cells. High, relatively (compared to control) decreased frequencies of blasts coexpressing CP markers and high relatively (compared to control) increased frequencies of CP expressing T cells correlated with improved blast lysis. These data has to be interpreted as a change of functionality of CP expressing cells under the influence of Kit-M - independent of patients' sex, age or clinical/haematological data and more or less from frequencies of CP expressing blast or T cells.

Targeted modulation of immune escape mechanisms, particularly through the regulation of CP expression on blasts and effector cells, could improve antileukemic immunity – an approach that has been insufficiently investigated so far, especially in the context of DC_{leu}-based strategies. Therefore the combination of ICI with DC/DC_{leu}-based immunotherapies could represent a promising strategy for improving T-cell activation and restoring immune regulation in AML. However, this needs to be confirmed by further studies. Ongoing research and additional studies will be essential to clarify the efficacy, safety and optimal use of DC-based immunotherapies, ICI and their potential combination in AML and thus improve clinical outcomes for AML patients.

Author Contributions: C.A. conducted DCC (WB, PBMNC), MLC and CTX experiments and all flow cytometric and statistical analyses. M.W., D.C.A., C.S. (Christoph Schwepcke), F.D.-G. performed additional DCC, MLC, CTX, and CSA experiments, which were analysed by C.A. D.M.K., A.R., C.S. (Christoph Schmid) provided leukemic whole blood samples and corresponding diagnostic reports. H.M.S. designed the study. C.A. and H.M.S. drafted the manuscript.

Statement of Ethics: Sample collection was conducted after obtaining written informed consent of the blood donor and in accordance with the World Medical Association Declaration of Helsinki and the ethic committee of the Ludwig-Maximilians-University Hospital Munich (vote No. 33905).

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Conflicts of Interest:

H.M.S. is involved with Modiblast Pharma GmbH (Oberhaching, Germany), which holds the European Patent 15 801 987.7-1118 and US Patent 15-517627, 'Use of immunomodulatory effective compositions for the immunotherapeutic treatment of patients suffering from myeloid leukemias'. For all other authors, there are no conflicts of interest to declare.

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