

Communication

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Communication

Circulating CD34+/CD38-/CD26+ Leukemia Stem Cells along Chronic Myeloid Leukemia Progression: Differences between Chronic, Accelerated and Blast Phase

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Abstract: In Chronic Myeloid Leukemia (CML) patients, CD34*/CD38*/CD26* cell population represents a "CML specific" Leukemia Stem Cell (LSC) compartment. Indeed, preliminary studies showed that the expression of CD26 discriminates bone marrow CML Leukemic Stem Cells (LSCs) from normal Hematopoietic Stem Cells (HSCs) or from LSCs of other myeloid neoplasms. We were first to demonstrate that at diagnosis CD34*/CD38*/CD26* cells are easily measurable also in Peripheral Blood (PB) and that residual circulating CD26*LSCs persist, with a fluctuating trend, in most patients in optimal response during treatment with Tyrosine Kinase Inhibitors (TKIs) and even after successful TKI discontinuation. These data corroborate and confirm the possibility of using flow-cytometry CD26* evaluation as an important diagnostic tool that, combined with molecular biology and cytogenetic, could provide a rapid diagnosis of Chronic Phase (CP) CML starting from a simple PB sample. Yet, few data are available regarding the behavior of CD26*LSCs during Accelerated Phase (AP) or Blast Phase (BP) CML and the role, if any, this peculiar staminal cell compartment may play in disease progression. In the present study we compared the presence and phenotypic characteristics of circulating CD26*LSCs in CP CML patients at diagnosis, during AP and in cases of progression to lymphoid BP, inquiring a possible role of these cells during disease evolution.

Keywords: CP-CML; Accelerated Phase (AP); Blast Phase (BP); CD26+LSCs

1. Introduction

Chronic Myeloid Leukemia (CML) is a tri-phasic myeloproliferative disorder, characterized by a specific cytogenetic and molecular marker (Ph chromosome, BCR::ABL1 fusion gene) and consisting of a chronic, accelerated and blastic phases [1]. At diagnosis, CML patients present most frequently in Chronic Phase (CP), whereas 4-5% of patients are firstly diagnosed in Accelerated Phase (AP) and only 1-2% of patients present in Blastic Phase (BP, or blastic crisis) [1,2]. From a physiopathology point of view, CP is the latent phase, in which often the patients are mild symptomatic or experience symptoms such as fatigue, weight loss and sweating correlated to marked leukocytosis, anemia and splenomegaly [1,2]. AP phase is characterized by additional genetic alterations, cytopenia and increased blasts, while BP phase that may present with myeloid, lymphoid or mixed phenotype, leads to a fatal outcome in nearly all CML patients due to resistance to current treatment strategies. Tyrosine Kinase Inhibitors (TKIs) have significantly improved the outcome of

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CP-CML patients by inducing a stable molecular response in the great majority of patients on continuous treatment and offering overall in about 15-20% of them the possibility of discontinuing TKI without disease recurrence. [3]. However, the risk of progression to AP or BP phase still remains, mainly in those patients not achieving an optimal response with available first, second and third generation TKIs.

CML LSCs have been identified in the fraction CD34*/CD38*/Lin* and specifically co-express CD26, which expression discriminates CML LSCs from normal HSCs and from LSCs of other myeloproliferative disorders [4]. In our previous studies, we demonstrated that CD26*LSCs are easily measurable in all Peripheral Blood (PB) samples of CP-CML patients at diagnosis and they are detectable during first line TKI treatment (imatinib, nilotinib, dasatinib) and during TFR, albeit with fluctuating values [5–7]. A confirmation that CD26 represent a marker of stemness, was obtained from a study performed with a confocal laser microscope in which it has been showed that CD26 antigen is co-expressed with the protein Polycomb BMI1. Usually, BMI1 is highly expressed in CML despite a lower transcript level of BCR::ABL1; in this study it has been demonstrated that the fraction CD34*/CD38*/CD26* represent the reservoir of the BMI1 protein [8].

Thanks to the "know how" acquired by performing these studies on CML LSCs, we developed a rapid custom-made flow cytometry tool for the measurement of CD34*/CD38-/CD26* population in PB samples of CP-CML patients and divulgated a standardized method useful for the diagnosis of CP-CML. However, few data are available regarding the behavior of CD26*LSCs during AP or BP-CML and the role, if any, this peculiar staminal cell compartment may play in disease progression. In the present study we compared the presence and phenotypic characteristics of circulating CD26*LSCs at diagnosis in CP-CML patients, in an AP-CML case and in 4 cases of progression to lymphoid BP.

2. Materials and Methods

The expression of CD34*/CD38-/CD26*LSCs was evaluated by flow cytometry using lyophilized reagents (CD34-FITC, CD26-PE, CD38-APC, and CD45-V500 BD, Biosciences) on EDTA PB samples of patients with leukocytosis suspected of CML. Data acquisition has been done using BD FACSCantoTM II or a BD FACSLyricTM cytometers (BD, Biosciences). Data analysis has been performed using BD FACSDivaTM or BD FACSuiteTM softwares (BD, Biosciences) using a strict gating strategy as previously described [6]. All CML diagnosis have been confirmed with the measurement of BCR::ABL1 transcript levels by RT-PCR assays, according to European Leukemia Net (ELN) criteria.

3. Results

Newly diagnosed CP-CML patients

In the contest of a prospective multicenter Italian study (Prospective Flowers Study) coordinated by Siena Hematology Unit, 270 newly diagnosed CP-CML patients have been studied. Flow cytometry evaluation on PB samples have been performed, according to our previous standardized procedure, and in 100% of the patients we detected the hallmark CD34+/CD38-/CD26+ population, characterized by a low expression of CD45 (CD45dim). More in details, flow cytometry measurements performed on PB CP-CML samples at diagnosis showed that the median value of CD34+ circulating cells, accounted for 0.36% (range 0.001-13%) of CD45+ cells; the CD34+/CD38- subset, represented 17% (range 0.3-71%) of total CD34+ cells. Within the CD34+/CD38- stem cell fraction, the median percentage of CD26+ cells were 35% (range 0.07–98%). Figure 1 shows a representative CP-CML case at diagnosis in which is evident that LSCs express CD45dim.

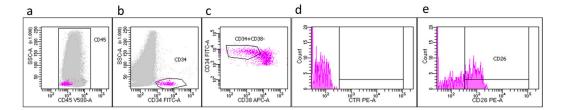


Figure 1. PB immunophenotypic analysis of a representative CP-CML case at diagnosis. **a.** CD45/SSC dot-plot. **b.** Expression of CD34⁺ cells. **c.** Gate CD34⁺/CD38⁻ in CD34⁺ cells. **d.** Isotype control. **e.** CD26 expression on CD34⁺/CD38⁻ gated cells.

Accelerated Phase CML case

On February 2022, PB sample of a 49-years-old male, suspected new diagnosis for CML, referred to our laboratory for the search of CD26⁺ LSCs. Cell blood count showed leukocytosis (WBC 68.405 x10⁹/L), anemia (Hb 9.4 g/dl) and thrombocytosis (PLT 1058 x10⁹/L). Surprisingly, we found a high total amount of CD34⁺ cells (27% of WBCs) that was in part CD45⁻ (54%) and in part CD45⁻dim (46%). We found that 70% of the CD45⁻dim/CD34⁺/CD38⁻ fraction co-expressed CD26, thus accounting for the CD26⁺LSCs population. However, when we analyzed the CD26 expression within the CD45⁻/CD34⁺/CD38⁻ cell fraction (supposedly, a blastic population) we found still a detectable, yet lower expression, of CD26⁺ cells (8%) (Figure 2). The patient underwent first line dasatinib therapy and was monitored for the presence of CD26⁺LSCs at 3- and 6-months during treatment. Along with the achievement of a molecular response, both flow cytometry evaluations showed the absence of CD26⁺LSCs and of the blastic cell population.

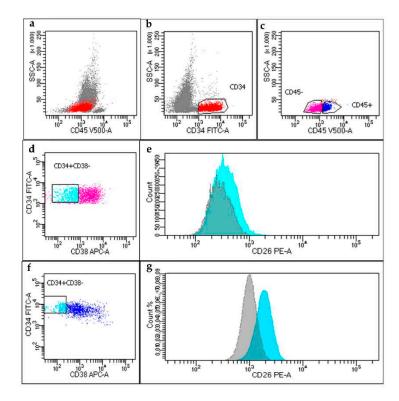


Figure 2. Immunophenotype of PB blasts during the accelerated phase of disease. **a.** CD45/SSC dotplot. **b.** Expression of CD34⁺ cells (red). **c.** CD34⁺ cells are distinguished into CD45⁻ fraction (pink) and CD45⁻dim fraction (blue). **d.** Gate CD34⁺/CD38⁻ in CD45⁻/CD34⁺ cells. **e.** CD26 partial expression on CD45⁻/CD34⁺/ CD38⁻ gated cells (blue peak) compared to isotype control (grey peak). **f.** Gate CD34⁺/CD38⁻ in CD45⁻dim/CD34⁺ cells. **g.** CD26 expression on CD45⁻dim/CD34⁻/ CD38⁻ gated cells (blue peak) compared to isotype control (grey peak).

Blastic Phase CML cases

We studied 4 cases of BP-CML patients. As following reported, we documented different cellular phenotypes.

Case #1

The first BP-CML case was a 67-years-old male, with cardiovascular co-morbidities, diagnosed in August 2016 with CP-CML. Patient started treatment with second generation TKI dasatinib (100mg/die) until complete morphological and molecular remission (June 2017). However, therapy was then discontinued for onset of dyspnea and pleural effusion and, after 6 months of discontinuation, patient showed a molecular relapse (BCR::ABL1 0.2%, MR 2), and started treatment with imatinib (400mg/die). On June 2021, he presented leukocytosis at PB (WBCs 18.610x10°/L), Hb 10.4g/dL and PLT 72x10°/L with a suspect of a BP-CML evolution. Blastic phase was confirmed by the presence of 72% of blast cells at the PB smear, FISH and molecular assays (85% of nuclei and 62% of BCR::ABL1 transcript, respectively). Flow cytometry analysis showed a cell population CD45-/CD34*/CD19*/CD10*/TdT* confirming the morphological evaluation. All CD34* cells were CD38- and CD26- at both PB and BM samples (Figure 3A). Patient was admitted to Hyper-CVAD + Ponatinib treatment.

Case #2

This case refers to an 83-years-old male who received diagnosis of CP-CML in February 2019. At that time, FISH confirmed CML diagnosis in 90% of cell studied and RT-PCR detected a p210 transcript type b2a2. The patient started treatment with first line imatinib (400 mg/die) achieving a complete cytogenetic response. A PB sample was then sent to Siena Hematology lab, in February 2021, to investigate the presence of CD26*LSCs during TKI treatment. Unexpectedly, cell blood count was found 2x10°/L. Immunophenotypic analysis showed the presence of CD34*/CD38* cells (9,3% of total cellularity in the sample), that were surprisingly negative for both CD45 and CD26 (Figure 3B). The latter evidence suggested an evolution to BP, that was shortly confirmed by a BM analysis, performed at local center, that documented a blast quote of 36%.

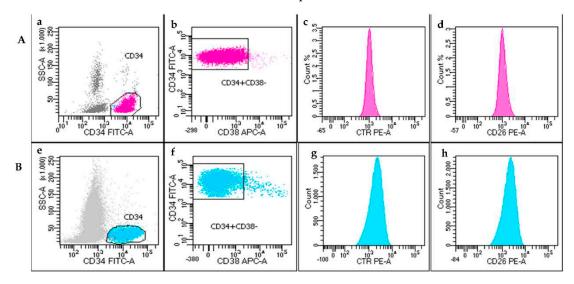


Figure 3. Immunophenotype of PB cells during Blastic Phase in Case #1 (**Panel A**) and Case #2 (**Panel B**). **a** and **e**. Expression of CD34⁺ cells. **b** and **f**. Gate CD34⁺/CD38⁻ in CD34⁺ cells. **c** and **g**. Histogram of control isotype. **d** and **h**. Histogram of CD26 expression.

Case #3

The present case involves a 34-years-old female patient that was diagnosed with a de novo lymphoid BP-CML. Cell blood count at diagnosis was the following: WBC 62.880 x10°/L, Hb 9.4 g/dl

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and PLT 51 x10°/L. Morphological examination of blood smear showed a consistent quote of myeloid immature cells and 23% of blasts with increased nuclear-cytoplasmatic ratio and prominent nucleoli. PB flow cytometry test showed the presence of lymphoid blasts CD45·/CD34*/CD19*/CD10*/TdT* and RT-PCR detected the presence of p210 BCR::ABL1 transcript (b3a2). Bone marrow aspirate confirmed a B-Lymphoid BP-CML. Circulating PB CD26*LSCs were not detected at that time (Figure 4A). The patient underwent induction treatment with Hyper-CVAD + Ponatinib and at recovery, concomitantly with the response to treatment and PB lymphoblasts disappearance, we documented a slight, but evident population of CD34* (0.091% of CD45* cells, with CD45dim) and CD34*/CD38* (12% of CD34*) cells. CD26* LSCs were 24% of CD34*/CD38* population (Figure 4B). Yet clearly detectable, the absolute percentage of CD26*LSCS accounted for 0.002% of CD45+ cells.

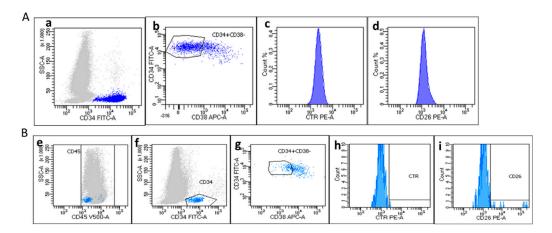


Figure 4. Immunophenotype of case #3 at the blastic phase (**Panel A**) and at recovery after induction treatment (**Panel B**). **Panel A** shows the following gates: **a** Expression of CD34⁺ cells. **b.** Gate CD34⁺/CD38⁻ in CD34⁺ cells. **c.** Histogram of control isotype. **d.** Histogram of CD26 expression. **Panel B** shows the same gating strategy of Panel A (gates **e-g**). A slight, but evident population of CD34⁺/CD38⁻/CD26⁺LSCs is present after induction therapy (gate **i**) compared it to the isotype control (gate **h**).

Case #4

The present case is referred to a 68-years old male, diagnosed in a peripheral hematology center in November 2021 with CP-CML. Diagnosis of CML was confirmed by CD26+LSCs evaluation, molecular and cytogenetic analysis in our laboratory. He started treatment with Imatinib (400 mg/die) but, due to GI toxicity, he discontinued therapy for two weeks and then reduced Imatinib dosage to 200 mg/die for intolerance. Few months later, patient's BM and PB samples were sent to Flow Cytometry Lab in Siena due to leukocytosis (WBCs 15.903x10°/L) and suspicion of lymphoid blast crisis. Surprisingly, the immunophenotypic analysis showed the presence of a population CD45dim/CD34+/CD38-, similar to the phenotype of a CP-CML, co-expressing CD19 and CD10 antigens, as for a typical lymphoid blast crisis. Consequently, we searched for the expression of CD26 and, indeed, the analysis confirmed that the CD45dim/CD34+/CD38-/CD19+/CD10+ cells co-expressed this antigen.

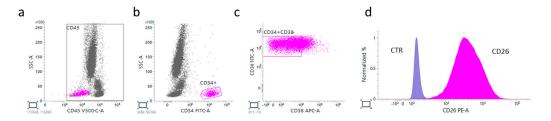


Figure 5. Immunophenotype of a lymphoid BP-CML case (#4) showing a population CD45^{dim}/CD34⁺/CD38⁻/CD10⁺ (gates **a**, **b**, **c**) positive for CD26 expression (gate **d**).

4. Discussion

Based on these results, we can confirm that the search of PB CD26⁺ LSCs cells is a powerful tool for the diagnosis of CML in CP, since these cells symbolize a "signature" of CP-CML. Indeed, as demonstrated in our previous studies, and documented in our prospective study, CD26 represents a specific marker of CML LSCs, expressed on 100% of PB samples of CP-CML at diagnosis. However, we also observed an interesting and misleading/controversial immunophenotypic panel in the AP and BP-CML samples studied.

In the AP-CML case, we identified the presence of two distinct PB CD34⁺/CD38⁻ populations: the first subset included CD45^{dim} cells in which CD26 antigen was expressed (70% of CD34⁺/CD38⁻ fraction) while, simultaneously, a second subset CD45⁻/CD34⁺/CD38⁻ was present in which blast cells were predominant, but in which we documented that the CD26 was still expressed on the CD34⁺/CD38⁻ population, yet in a lower percentage of cells (8%).

Regarding the four lymphoid BP-CML cases, we documented a homogeneous phenotype in 3 patients, where the blasts were negative for CD45 expression and consequently, also for CD26 antigen. Surprisingly, the case #4 was countertrend compared with others, showing a blast phenotype CD45dim/CD34+/CD38-(CD19+/CD10+ and the expression of CD26+ (99% of CD34+/CD38- cells). In this latter case lymphoid blasts displayed a phenotype CD45dim/CD34+/CD38- comparable to that of LSCs found in CP-CML at diagnosis. Even if we studied only few patients, the different CD26 expression documented in our AP- and BP-CML cases, induced to ask new questions and to formulate possible hypothesis. We confirmed, as reported by others, that CD26+LSCs are not detectable (or "disappear"?) during a classical CD45- lymphoid BP-CML. However, we documented also that a "blastic" CD45dim/CD26+LSCs population may coexist.

This preliminary, yet unique, evidence arise the question regarding the role of CD26⁺LSCs during blastic transformation, suggesting two possible scenarios: CD26 LSCs may "shift" phenotype, first losing CD45 expression and subsequently CD26 expression and then acquiring a blastic phenotype as it would suggest the case studied during AP and to some extent the BP case #4; alternatively, CD34⁺ blasts arise from a different subset and CD26⁺ LSCs cells may coexist during BP albeit surmounted by the bulk of CD34⁺ blasts and thus not detectable.

In conclusion, the present study reports for the first time, additional information about CD26⁺ expression in AP and BP CML cases, never investigated until now. However, more studies are needed to better enlighten if the CD26 antigen concur or not to the progression of CML from Chronic to Accelerated and Blast Phase.

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Informed Consent Statement: Informed consent was obtained from all subjects involved in the study.

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