

CHRONIC MYELOID LEUKEMIA

CLINICAL ROLE OF PERIPHERAL BLOOD LEUKEMIA STEM CELLS AT DIAGNOSIS IN CHRONIC MYELOID LEUKEMIA: FINAL RESULTS OF THE PROSPECTIVE FLOWERS STUDY

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Background: We previously demonstrated that CD26+ leukemia stem cells (LSCs) are detectable by flow cytometry in the peripheral blood (PB) of chronic myeloid leukemia (CM-L) patients (pts) at diagnosis, during tyrosine kinase inhibitor (TKI) therapy, and in treatment-free remission (TFR). However, prospective data on the dynamics of PB CD26+LSCs from diagnosis and their potential correlation with molecular response (MR) are lacking.

Methods: We report final results of a prospective, multicenter Italian study enrolling newly diagnosed chronic phase (CP) CML. Pts were centrally monitored by flow cytometry to quantify PB CD26+LSCs from diagnosis up to 24 months (mos) of therapy.

Results: 242 consecutive CP-CML pts were included (132 treated with imatinib, 72 with nilotinib, and 38 with dasatinib). The CD26+LSC count at diagnosis varied among pts, with a median of 7.14 cells/ μ L (IQR 2.18-33.26 cells/ μ L). During TKI, we observed a rapid and significant reduction of CD26+LSCs, with median values decreasing to 0.013 cells/ μ L (IQR 0-0.034), 0.011 cells/ μ L (IQR 0-0.031), and 0.007 cells/ μ L (IQR 0-0.025) at 3, 12, and 24 mos, respectively. No significant differences in LSC log-reduction were observed according to the type of TKI employed. CD26+LSCs were significantly higher in the high Sokal risk group compared to intermediate or low risk groups (22.65 cells/ μ L vs 5.60 cells/ μ L vs 6.16 cells/ μ L) ($p=0.018$). A significant association was found between low CD26+LSCs count at diagnosis and opti-

mal MR (BCR::ABL1 <10% at 3 mos and <0.1% at 12 and 24 mos). Pts achieving optimal MR at 3 mos had a median CD26+LSCs at diagnosis of 6.21 cells/ μ L (IQR 1.79-31.50), whereas suboptimal responders had a median of 19.87 cells/ μ L (IQR 5.37-39.81) ($p=0.03$). Similarly, optimal responders at 12 and 24 mos had significantly lower median CD26+LSCs at diagnosis compared to suboptimal responders (5.50 vs. 16.87 cells/ μ L, $p=0.004$ and 6.05 vs. 20.52 cells/ μ L, $p=0.009$, respectively). Furthermore, pts who switched TKI for failure had higher baseline CD26+LSCs counts (median 14.59 cells/ μ L; IQR 3.76-46) than those who did not switch (median 5.82 cells/ μ L; IQR 2.35-26.70; $p=0.034$). Three tertiles of CD26+LSCs at diagnosis were defined (<3.21, 3.21-19.21, and >19.21 cells/ μ L), showing significant correlation with MR rates: at 3 mos, the incidence of BCR::ABL1 <10% was 93.5% in the lowest tertile vs. 78.8% in the highest ($p=0.027$); at 12 mos, optimal response was 78.5% vs. 62.8% ($p=0.015$); at 24 mos, response rates were 90.8% vs. 77.9% ($p=0.079$).

Conclusions: This prospective study demonstrated a rapid rate of reduction of CD26+LSCs during TKI, confirming their long-lasting persistence even at very low levels. For the first time, a correlation between the amount of CD26+LSCs at diagnosis and the response to TKI treatment was documented. Given these results, the bulk of CD26+LSCs at diagnosis could represent an easily and rapidly measurable, new prognostic tool for predicting TKI response.