Ibrutinib-induced lymphocytosis in patients with chronic lymphocytic leukemia: correlative analyses from a phase II study

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Abstract

Ibrutinib and other targeted inhibitors of B-cell receptor signaling achieve impressive clinical results for patients with chronic lymphocytic leukemia (CLL). A treatment-induced rise in absolute lymphocyte count (ALC) has emerged as a class effect of kinase inhibitors in CLL and warrants further investigation. We here report correlative studies in 64 patients with CLL treated with ibrutinib. We quantified tumor burden in blood, lymph nodes, spleen, and bone marrow, assessed phenotypic changes of circulating cells, and measured whole blood viscosity. With just one dose of ibrutinib the average increase in ALC was 66%, and in over 40% of patients the ALC peaked within 24 hours of initiating treatment. Circulating CLL cells on day 2 showed increased Ki67 and CD38 expression, indicating an efflux of tumor cells from the tissue compartments into the blood. The kinetics and degree of the treatment-induced lymphocytosis was highly variable; interestingly in patients with a high baseline ALC the relative increase was mild and resolution rapid. After two cycles of treatment the disease burden in lymph node, bone marrow, and spleen...
decreased irrespective of the relative change in ALC. Whole blood viscosity was dependent on both ALC and hemoglobin. No adverse events were attributed to the lymphocytosis.

**Keywords**
Ibrutinib; lymphocytosis; safety

**Introduction**

Chronic lymphocytic leukemia (CLL), a malignancy of mature B-cells that involves blood, bone marrow and lymphoid tissues, is the most common leukemia in western countries.\(^1\) In the US alone ~120,000 people live with CLL\(^2\) while an estimated 15,680 people will be newly diagnosed with, and 4,580 will die of CLL in 2013.\(^3\) Median survival with early stage disease is 10.7 years; however, the clinical course is quite heterogeneous, depending on prognostic markers like Immunoglobulin Heavy Chain Variable (IGHV) gene mutation status, ZAP70 expression, deletion of the short arm of chromosome 17 (del(17p)) or presence of SF3B1 and NOTCH1 mutations.\(^4,\(^5\)\)

B-cell receptor (BCR) signaling has emerged as a key pathway in the pathogenesis of CLL.\(^6\)–\(^8\) A role for antigenic stimulation of the clonal cells was first indicated by the observation that CLL cells use a restricted repertoire of IGHV genes that encode the antigen interacting interface of the BCR.\(^9,\(^10\)\) More recently, the discovery of BCR stereotypes shared by a substantial proportion of CLL cases, suggests that CLL cells may arise from B-cells with defined antigen specificities.\(^11,\(^12\)\) BCR signaling and activation of the NF-κB pathway occur primarily in the lymph node microenvironment promoting cell growth, proliferation, and survival.\(^13,\(^14\)\) Thus, antigenic stimulation emerges as a driving pathway in the pathogenesis of CLL; a mechanism that is also implicated in an increasing number of mature B-cell malignancies.\(^15,\(^16\)\)

Bruton’s tyrosine kinase (BTK), a cytoplasmic non-receptor tyrosine kinase, is recruited early in the BCR signaling cascade in conjunction with SYK and PI3K\(^\delta\).\(^16,\(^17\)\) BTK couples BCR activation to intracellular calcium release and activation of NF-κB and is essential for normal B-cell development and response of B-cells to antigenic stimulation.\(^18\) Knockdown of BTK is lethal to select lymphoma cell lines derived from activated B-cell like diffuse large B-cell lymphoma\(^19\) and decreases the viability of primary CLL cells.\(^20\) In addition, genetic ablation of BTK inhibits disease progression in mouse models of CLL.\(^20,\(^21\)\) Ibrutinib, an orally active agent, covalently binds to Cys-481 of BTK thereby irreversibly inactivating the kinase.\(^22\) In the phase I study, ibrutinib was well tolerated and active across a spectrum of mature B-cell malignancies, with the highest response rates in CLL and mantle cell lymphoma (MCL).\(^23,\(^24\)\) More recently, overall response rates of >70%, and an estimated 26 month progression free survival rate of 75% for previously treated patients with CLL was reported.\(^25\) In vitro ibrutinib has been shown to inhibit proliferation, adhesion and migration of CLL cells.\(^26\)–\(^29\) Further, murine CLL models suggest that ibrutinib inhibits homing of CLL cells to tissue sites.\(^28,\(^30\)\)

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In addition to ibrutinib, several other inhibitors of kinases in the BCR pathway are in clinical development.\textsuperscript{7, 8, 16, 31} Initial clinical experience with such BCR inhibitors raised concerns due to a sometimes dramatic worsening of peripheral lymphocytosis,\textsuperscript{32, 33} which is now recognized as a class effect. Concerns about patient safety due to the treatment-induced lymphocytosis have been somewhat alleviated by the increasing experience with these agents in clinical trials. However, many questions remain. Here we focused on the kinetics and inter-individual variability in treatment-induced lymphocytosis, characterized changes in the immune-phenotype of circulating CLL cells on treatment, assessed concomitant changes in disease distribution in different anatomic compartments, and sequentially determined whole blood viscosity during the period of peak lymphocytosis. We found that the ibrutinib induced lymphocytosis developed almost immediately after the first dose of drug, peaked within 24 hours in many patients, and showed pronounced inter-patient variability. We provide direct \textit{in vivo} evidence that the initial rise in lymphocytosis is in large part due to the release of previously activated cells from the lymph node. Furthermore, substantial reductions in tumor burden in lymph node, bone marrow, and spleen independent of changes in the number of circulating CLL cells underscores the notion that in CLL patients treated with BCR inhibitors, the ALC is not a valid surrogate of overall disease burden or activity.\textsuperscript{32}

\section*{Materials and Methods}

\subsection*{Patients, blood counts, and whole blood viscosity}

We report correlative analyses on 64 CLL patients enrolled between January 2012 and October 2013 in our ongoing, investigator-initiated phase II study of ibrutinib (NCT01500733). The study was approved by the local ethics committee; informed consent was obtained from all patients in accordance with the Declaration of Helsinki. Briefly, both treatment naïve and relapsed/refractory patients with either del(17p) or age \textgeq 65 years were eligible (Table 1) and treated with ibrutinib 420 mg orally once daily until disease progression or the occurrence of intolerable side effects. Mutation status of the immunoglobulin heavy chain variable (\textit{IGHV}) gene was assessed as described and CLL with <2\% mutations was classified as unmutated.\textsuperscript{13} Complete blood counts were determined in the NIH central laboratory. Groups of patients having distinct patterns of treatment-induced lymphocytosis (in regards to degree of change in ALC, and the kinetics of onset and resolution) were identified using hierarchical clustering (Eisen Laboratory, Stanford University, CA). Whole blood viscosity was determined by quantitative viscometry of whole blood (ARUP Laboratories, University of Utah, Salt Lake City, UT). Adverse events were graded according to CTCAE v 4.03.

\subsection*{Modeling CLL tumor burden and tissue redistribution}

Tumor volume in spleen and lymph node was computed from CT scans using syngovia software (Siemens, Cary, NC). Of the spleen volume, 70\% was considered due to CLL cells.\textsuperscript{34} Bone marrow volume was calculated as previously published,\textsuperscript{35} and the content of CLL cells was calculated from the overall marrow cellularity multiplied by the degree of CLL cell infiltration. Cellular density was estimated at 10^{9} cells/ml based on measurements of thymocyte density.\textsuperscript{36} Peripheral blood volume was calculated using the Nadler formula.\textsuperscript{37}
The number of CLL cells was calculated based on measured ALC and the percentage of CD19+ cells among all lymphocytes.

**Flow cytometry**

Peripheral blood mononuclear cells (PBMC) were prepared by density-gradient centrifugation (Lymphocyte Separation Media; ICN Biomedicals, Irvine, CA). Lymph node derived single cell suspensions were obtained from core biopsies. Cell suspensions were stained as previously described (BD Biosciences, Franklin Lakes, NJ). For Ki67 staining, cells were fixed in 4% paraformaldehyde and permeabilized in 70% EtOH. To assess cell death fresh whole blood samples were collected, red cells were removed by ACK lysis, and CLL cell viability was assessed with the LIVE/DEAD® Fixable Violet Dead Cell Stain Kit (Invitrogen, Grand Island, NY). Cells were analyzed on a FACS Canto II flow cytometer (BD Biosciences) using FACS-DIVA 6.1.1 and FlowJo software (Version 8.8.6; TreeStar, Ashland, OR).

**Statistical analysis and mathematical modeling**

When comparing patient groups, a linear model was applied (JMP10 software, SAS Inc, Cary, NC). When comparing specific events among patient groups a Fisher’s exact test was used. All other statistical analyses were by Student’s T-test (Prism5, GraphPad, La Jolla, CA).

To test the hypothesis that the treatment-related increase in ALC is due to efflux of CLL cells from the lymph node, the predicted percentage of CLL cells expressing Ki67 (Ki67_TPB) on day 2 was calculated from the change in ALC and the measured percentage of CLL cells expressing Ki67 pre-treatment in peripheral blood (Ki67_PB pre) and lymph node (Ki67_LN pre):

\[
Ki67_{PB\, Day\, 2} = \left( \frac{ALC_{pre}}{ALC_{pre} \times Day^2} \right) \times Ki67_{PB\, pre} + \left( 1 - \frac{ALC_{pre}}{ALC_{pre} \times Day^2} \right) \times Ki67_{LN\, pre}
\]

**Results**

**Ibrutinib-induced lymphocytosis develops within hours of starting treatment and is highly variable between patients**

Within 24 hours of initiating ibrutinib the mean baseline ALC of 105 000/μL increased on average by 66% (P<0.001) and continued to increase throughout the first cycle before it gradually declined (Figure 1a and b). Within four hours of starting treatment the mean ALC had already increased in most patients (data not shown). Thus, the onset of ibrutinib-induced lymphocytosis was immediate and in many patients reached its peak within days; in 46% of patients on day 2 and in 78% by day 28 (Figure 1c). The magnitude of the treatment-induced lymphocytosis was highly variable between patients (Figure 1d), without any correlation to prior treatment history or the presence of absence of del(17p) (Supplementary Figure S1a–d). However, patients with IGHV mutated CLL showed a more pronounced rise in ALC that resolved more slowly than in patients with IGHV unmutated CLL (P<0.001; Supplementary Figure S1e–f).

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To test for factors that might predict distinct patterns of lymphocytosis, we used an unsupervised clustering analysis of the relative change in ALC over the first 6 months on treatment compared to baseline. Three separate clusters of patients that differ in the degree and the kinetics of the lymphocytosis were identified (Figure 2a–c). Notably, the pre-treatment ALC was significantly different between the three clusters ($P<0.002$, Figure 2d). The mean pre-treatment ALC was highest for patients in cluster 1 and in this group of patients the treatment-induced lymphocytosis peaked within days and resolved rapidly. Conversely, patients in cluster 3 tended to have lower pre-treatment ALC and showed a greater relative increase in lymphocytosis that developed and resolved more slowly. Compared to the other clusters, cluster 3 was enriched for patients having $IGHV$ mutated CLL and bulky lymphadenopathy (at least one node with largest diameter >5 cm; Table 2). Conversely, patients in cluster 1 frequently had advanced Rai stage (80%). Thus, it appears that CLL patients with moderate lymphadenopathy may have a shorter time to peak ALC and a faster resolution of lymphocytosis, while bulky disease was here associated with more prolonged lymphocytosis. Notably, there was no significant difference in the degree of nodal response at 6 months among the three clusters (Table 2), suggesting that a prolonged lymphocytosis does not predict for inferior response; in agreement with a recent report by Woyach et al.$^{38}$

**Concurrent reduction in total tumor burden irrespective of treatment induced lymphocytosis**

We estimated a patient’s total disease burden from the ALC, the extent of lymphadenopathy and splenomegaly on CT scans, and the degree of bone marrow infiltration. Figure 3 summarizes data for four representative patients. The distribution of disease among the different anatomic compartments varied greatly between individual patients (Figure 3a, Supplementary Table 1). During treatment, we repeatedly assessed the tumor burden in these sites for four patients chosen to represent cases with both diverse involvement of the four compartments pre-treatment and different kinetics of the treatment-induced lymphocytosis. The aggregate volume of lymphadenopathy as calculated from CT scans is shown for a representative patient in Figure 3b; ibrutinib reduced the pre-treatment lymph node volume of 3 161 mL (equal to an estimated $3.2\times10^{12}$ CLL cells) by >80% within two months, with continued improvement at 6 months. Similarly, ibrutinib also rapidly reduced the tumor burden in the spleen (Figure 3c). The degree of bone marrow infiltration was estimated from anti-CD79a stained bone marrow biopsies (Figure 3d). Irrespective of the initial tumor distribution, all four representative patients showed a pronounced decrease in the number of CLL cells over time, with the most rapid decrease occurring during the first two months (Figure 3e). One patient showed an apparent increase in the total number of CLL cells at 2 months despite reductions in disease burden in all three tissue sites due to a pronounced increase in ALC. This likely reflects release of CLL cells from the liver and intestinal tract, sites known to be significant reservoirs of B-cells, but not included in our model.$^{39-41}$

The substantial reduction in total tumor burden cannot be explained by mere disease redistribution between different anatomic sites and suggests that there is a substantial amount of treatment-induced cell death. For example in three of the four patients shown in
Figure 3e trillions of CLL cells “disappear” within the first two months; and that is only from the four sites amenable to study. However, evidence of cell death has been elusive. For example none of the patients in our study showed signs of tumor lysis syndrome, and in ficoll-ed, viably frozen PBMCs we failed to detect an increase in cell death or apoptosis. When we switched to analyzing fresh whole blood samples we found that pre-treatment a median 2.4% of the circulating tumor cells were dead or dying. On day 28 on ibrutinib the frequency of dead or dying cells in circulation on average more than doubled compared to pre-treatment ($P=0.03$; Figure 3f).

Evidence for efflux of tissue resident CLL cells into the peripheral blood

*In vitro* and murine studies suggest that both inhibition of cell adhesion and homing of circulating cells to tissue sites may contribute to treatment-induced lymphocytosis in CLL. Based on the rapid onset of the treatment-induced lymphocytosis, we hypothesized that efflux of CLL cells from tissue sites fuels the rise in circulating cells. To distinguish CLL cells in circulation from tissue-resident cells we made use of prior observations that CLL cells are activated and proliferate in the tissue microenvironment, and the two populations therefore differ in their immuno-phenotypic characteristics (Figure 4a). Expression of activation markers, including CD38, CD69, and CD86, and the proliferation marker Ki67 have been found to be downregulated by BCR inhibitors. As expected we found downregulation of these markers on circulating tumor cells on day 14 of ibrutinib therapy (data not shown). However, on day 2 concurrent with the increase in ALC the frequency of CD38 and Ki67 positive CLL cells in circulation increased significantly ($P \leq 0.004$, Figure 4b–c, and Supplementary Figure S2a) consistent with an influx of tissue resident cells into the blood. Next we used the relative change in ALC and in the fraction of circulating cells expressing Ki67 on day 2 to estimate to what degree the influx of tissue resident cells can account for the rise in ALC. Figure 4d illustrates these changes in a representative patient. The fraction of Ki67 expressing CLL cells as measured by flow cytometry pre-treatment was 6.4% in the peripheral blood and 12.8% in the lymph node. On day 2 the ALC had increased from 90 000/μl to 128 000/μl. Thus one third of the cells on day 2 were “additional cells”. If all these additional cells are derived from the lymph node, the frequency of Ki67 positive cells is predicted to increase to 8.3%. Indeed the actual frequency measured by flow cytometry was 8.2%. We could assess eight patients in this manner and consistently, actual and predicted frequency of Ki67 expressing cells were virtually identical ($R=0.99, P<0.001$; Figure 4e). Using this same formula, we could also predict the expected rise in ALC based on the pre-treatment and day 2 Ki67 values ($R=0.99, P<0.001$; Figure 4f). Thus, the treatment-induced rise in ALC on day 2 can be accounted for by the influx of lymph-node resident CLL cells into the blood.

Whole blood viscosity is determined by ALC and hemoglobin

Patients with certain types of leukemia may develop leukostasis leading to central nervous system and respiratory impairment. CLL patients, even those with very high lymphocyte counts, are rarely reported to suffer such complications; however, the rapid increase in ALC and the release of more activated cells from tissue sites might contribute an additional risk. Fortunately, no adverse events were attributed to leukostasis or hyperviscosity syndrome in this study, where the highest ALC was 475 000/μl, and 42% of patients had an ALC greater

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than 200,000/μl at least once during the first two months. Prospectively measured whole
cell blood viscosity pre-treatment and on days 2 and 28 correlated with ALC, but did so even
more strongly with hemoglobin (P<0.001, Figure 5a and b). Overall, there was no
significant increase in whole blood viscosity on treatment (Figure 5c). Only in two patients a
slightly increased whole blood viscosity was recorded, in one patient on day 2 with an ALC
of 405,000/μl and in the other on day 28 with an ALC of 388,000/μl. ALC and hemoglobin
were inversely correlated and patients with high ALC tended to be more anemic
(Supplementary Figure S3). Interestingly, in patients with a pre-treatment hemoglobin ≥10
g/dL there was a decrease in mean hemoglobin from 11.7 to 10.9 g/dL during the first 28
days on treatment (P ≤0.02, Figure 5d). Figure 5e illustrates the interaction of hemoglobin
and ALC on whole blood viscosity. Notably, the effect of a high ALC may in part be offset
by a degree of anemia.

Discussion

Treatment-induced lymphocytosis has become a hallmark of BCR inhibitors in
CLL, 23, 25, 32, 33 Here we report clinical and biologic aspects of ibrutinib-induced
lymphocytosis in patients with CLL; we show that the rise in ALC is virtually immediate,
driven by the efflux of cells from tissue compartments, and paralleled by a substantial
decrease in total tumor burden. In prospective measurements of whole blood viscosity we
identified a small subset of patients in whom whole blood viscosity can exceed the normal
range. However, no clinical adverse events were attributed to the lymphocytosis.

Prior studies have reported treatment-induced lymphocytosis with BCR inhibitors within
weeks of initiating treatment, 23, 25, 33 Here we show that the onset of lymphocytosis occurs
within hours of the first dose of ibrutinib and identify a group of patients in whom the
lymphocytosis peaks within the first days and resolves before the end of the first cycle. The
variability in degree and kinetics of the ibrutinib-induced lymphocytosis in our cohort
appears to be somewhat more heterogeneous than seen in other studies. While we included
both treatment naïve and relapsed/refractory patients of all age groups, and a large
proportion of patients having del(17p), none of these characteristics predicted kinetics or
degree of lymphocytosis. Of practical consequence is the observation that patients with a
high pre-treatment ALC tend to have a rapid but relatively modest rise in ALC followed by
fast resolution of the leukemic disease. Conversely, the group of patients showing a slow but
continuous rise in ALC also had slow resolution of leukemic disease, had lower ALC pre-
treatment, a higher proportion of patients with bulky disease, and commonly had IGHV
mutated CLL. Thus, patients in this group would be predicted to have a more indolent
course and extended benefit of treatment, at least with conventional
chemoimmunotherapy. 1, 49 Therefore, as long as there is no evidence of progressive disease
in other sites, it seems appropriate to continue treatment and await the eventual resolution of
the lymphocytosis. Overall, compared to patients with IGHV unmutated CLL patients with
IGHV mutated CLL had a greater relative increase in ALC and slower resolution of
lymphocytosis confirming an earlier report. 25 While IGHV mutated patients may therefore
have lower response rates by IWCLL criteria, Woyach et al. recently showed that patients
with persistent lymphocytosis have a similar PFS compared to those who reach an IWCLL
objective response within the first year. 38 In accordance with this, we did not find a

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significant difference in the degree of nodal response at 6 months among the three clusters, further suggesting that a prolonged lymphocytosis does not indicate an inferior response.

Resolution of lymphadenopathy concurrent with the rise in ALC has been a strong indication that treatment-induced lymphocytosis is not a sign of progressive disease but reflects redistribution of tumor cells between different anatomic compartments. Here we present for the first time direct evidence that the initial rapid rise in ALC is driven by the release of CLL cells from tissues, in particular from the lymph node. This conclusion is based on the increased frequency of CLL cells expressing Ki67 and CD38 in the blood within 24 hours of starting ibrutinib and supported by a mathematical model derived from in vivo measurements in both blood and lymph node resident cells (Figure 4d). At least in the first 24h the bulk of cells released into the blood appear to come from the lymph node but we cannot rule out that some cells also egress from the bone marrow or other tissue sites.

While several markers have been reported to be differentially expressed between CLL cells in the blood and tissues, not all are suitable to assess shifts in cell populations on ibrutinib. For example the activation markers CD69 and CD86 are so rapidly downregulated by ibrutinib that they cannot identify different cell populations once treatment is initiated. In contrast, we observed an increase of CXCR4 expression on CLL cells during treatment with ibrutinib (data not shown), which precludes the identification of the CXCR4$^{\text{dim}}$/CD5$^{\text{bright}}$ population reported by Calissano et al. Similarly, on continued treatment both Ki67 and CD38 are downregulated by BCR inhibitors. Therefore, we could not distinguish different cell populations at later time-points. Thus, while we show that an efflux of cells from the tissue into the blood accounted for the observed rise in ALC by day 2, we cannot assess to what degree reduced homing of CLL cells to the tissue may contribute to the persistence of the lymphocytosis. However, the latter mechanism may contribute to the drop in ALC during treatment interruptions as seen in early studies that cycled ibrutinib 4 weeks on, one week off.

Leukostasis is a rarely reported complication of CLL. However, in some settings the treatment-induced worsening of lymphocytosis has raised concerns about patient safety. We therefore prospectively incorporated whole blood viscosity measurements. Clinically we did not observe any symptoms or signs suggestive of leukostasis. Further, whole blood viscosity was only slightly elevated in two patients. While our experience is limited in patients with ALC greater than 300 000, the positive correlation between ALC and whole blood viscosity suggests that blood flow may be altered at very high ALCs especially in patients with normal hemoglobin (Figure 5e). The fact that most patients with high ALC had hemoglobin $<10$ g/dL could in part account for their normal viscosity readings and may justify withholding administration of blood products in patients with high ALC.

Treatment-induced lymphocytosis has attracted considerable attention. However, in most patients, the circulating CLL cells are only a minor fraction of the total tumor volume. Importantly, the bulk of the disease, located in tissue sites such as bone marrow, lymph node, spleen and likely the intestinal tract, is substantially decreased on ibrutinib (Figure 3). Also, 27% of patients already had a reduction in ALC below baseline by two weeks. Thus, in most patients there is a rapid reduction in total tumor burden indicating an increased rate of cell death on ibrutinib. In fresh whole blood samples obtained pre-
treatment we identified a median 2.4% of the circulating tumor cells as dead or dying, which is within the range of the previously calculated daily death rates in untreated CLL patients using heavy water labeling.42 Our measurements suggest that the frequency of dead or dying cells in circulation more than doubled on ibrutinib (Figure 3f). As dead cells are constantly cleared through the reticuloendothelial system this is most consistent with an increased death rate. An absolute increase in the rate of cell death by 2% per day would result in a >50% reduction in tumor burden within 28 days, which is in good agreement with the initial kinetics of tumor response seen in many patients. In addition, inhibition of tumor proliferation by ibrutinib, as reported elsewhere,46, 47 will prevent or reduce the replacement of the dying cells. A slow but steady decay rate of the tumor is also in accordance with the notable absence of reports of tumor lysis syndrome with single agent ibrutinib,23, 25 and the modest degree of apoptosis induced by ibrutinib in vitro.26, 28 As BCR inhibitors decrease NF-κB signaling and expression of the anti-apoptotic BCL2 family member BCL2A130, 45, 46 cell death may be a consequence of reduced pro-survival signaling rather than of a direct cytotoxic effect. However, this remains to be more fully evaluated.

Ibrutinib is active in several mature B-cell malignancies.23, 52 Notably, patients with MCL also frequently experience treatment-induced lymphocytosis.44, 52 Although the mechanisms leading to redistribution of disease may be similar to those in CLL, patterns of lymphocytosis in MCL and their relation to clinical response and patient safety need to be independently studied. Results presented here support the positive benefit to risk profile of continued ibrutinib treatment for most patients with CLL showing treatment-induced lymphocytosis. In fact, treatment-induced lymphocytosis in CLL may be viewed as an on-target effect of BCR inhibitors, with the caveat that it may be missed as the peak can occur within the first days or even hours of starting ibrutinib. Thus treatment-induced lymphocytosis is neither a sign of progressive disease nor can it be used as a surrogate of response.

Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

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Ibrutinib-induced lymphocytosis develops rapidly, peaks within days, and is highly variable between patients. (a–d) Data on 41 patients with complete counts for the first six months on ibrutinib are depicted. (a) Change in mean ALC over time; vertical lines indicate SEM. (b) Mean relative change of ALC on treatment over baseline; vertical lines indicate SEM. (c) Cumulative proportion of patients reaching their peak ALC at the indicated time point. Note >40% of patients reached the peak ALC on day 2. (d) Box and whisker plots demonstrating distinct inter-patient variability in ALCs.
Figure 2.
Variability in development and resolution of ibrutinib-induced lymphocytosis. (a) Hierarchical clustering of change in ALC normalized to baseline (day 1) in individual patients (n=41) reveals three distinct patterns. The degree of change is color coded according to the legend shown. (b–c) The mean for each of the three patient clusters is shown as (b) ALC or (c) relative change of ALC compared to pre-treatment. (d) The pre-treatment ALC for the three patient clusters are significantly different by Student’s T-test.
Figure 3.
Ibrutinib rapidly decreased total tumor burden and increases the rate of cell death (a) A graphic representation of the estimated tumor burden in different anatomic compartments pre-treatment is shown for four representative patients (identified by study code). (b–d) Change in tumor burden from pre-treatment (Pre) to 2 and 6 months (MO) on treatment in different tissues. (b) Graphic representation of total lymph node volume computed from whole body CT scans. (c) Graphic representation of spleen volume computed from CT scan. (d) CLL cell infiltration of bone marrow visualized by CD79a staining. (e) Changes in disease burden on treatment is shown for the four representative patients. Also see Supplementary table S1. (f) The viability of circulating CLL cells was measured in fresh whole blood samples using the LIVE/DEAD stain (n=12, each symbol represents a different patient). Ibrutinib doubles the rate of cell death (P=0.03 by paired Student’s T-test.)
Figure 4.
Ibrutinib induced lymphocytosis is driven by the release of cells from the lymph node. (a) More CLL cells in the lymph node (LN) express Ki67 (proliferation marker) and CD38 (activation marker) compared to CLL cells in the peripheral blood (PB). All data are from untreated patients, red symbols represent ibrutinib study patients (n=5). (b) Shown is the relative change in the frequency of Ki67 positive CLL cells in the peripheral blood on day 2 (after one dose of ibrutinib) compared to pre-treatment (n=28). (c) Change in the frequency of CD38 positive CLL cells on day 2 compared to pre-treatment (n=11, CD38+ patients only). (d) Efflux of CLL cells from the lymph node increases the fraction of Ki67 positive cells in the blood. Shown is the data for one representative patient. The frequency of Ki67 expressing CLL cells pre-treatment in lymph node (measured in single cell suspension by flow cytometry) and in the peripheral blood is shown. On day 2, 30% of the ALC consists of “additional cells” and the frequency of Ki67 positive CLL cells has increased to 8.2%. If all the additional cells derive from the lymph node (with Ki67 expression in 12.8%) the predicted frequency of Ki67 positive cells is 8.3%. (e–f) Regression analysis of predicted vs. actual percent Ki67 expression (e) and ALC values (f). Dashed lines indicate 95% confidence intervals (n=8).
Whole blood viscosity is rarely elevated in CLL and is influenced by ALC and hemoglobin. (a) Correlation of whole blood viscosity (normal range 3.6–6 cP, shaded) with ALC (n=105; R=0.41; P<0.001). (b) Correlation of whole blood viscosity with hemoglobin concentration (n=105; R=0.59; P<0.001). (c) Whole blood viscosity pre-treatment (pre) and on treatment days 2 and 28. (d) Change in hemoglobin in patients with an initial value ≥10 g/dL (n=26). (e) Contour plot of whole blood viscosity as a function of both ALC and
hemoglobin. Whole blood viscosity measurements are color coded from low (white) to high (red). Correlations by Pearson’s test.
Table 1

|                   | n=  | Sex (% M) | Rai Stage (% 3,4) | % IGHV Unmutated | % CD38 | Median Pre ALC |
|-------------------|-----|-----------|-------------------|------------------|--------|----------------|
| >65 years old     |     |           |                   |                  |        |                |
| TN                | 14  | 57%       | 64%               | 45%              | 29%    | 142            |
| RR *              | 15  | 47%       | 80%               | 71%              | 64%    | 96             |
| del(17p)          |     |           |                   |                  |        |                |
| TN *              | 24  | 54%       | 63%               | 53%              | 22%    | 74             |
| RR                | 11  | 55%       | 64%               | 73%              | 36%    | 94             |
| All *             | 64  | 58%       | 70%               | 61%              | 37%    | 93             |

1 Unmutated indicates < 2% change in IGHV gene sequence compared to germline.

2 CD38 positive (+) indicates ≥30% of CLL cells express CD38 above isotype control.

* One patient in the >65 RR group and six patients in the del(17p) group do not have available IGHV mutational information or CD38 expression data. Shown are the percentages of the patients where information was available.
Table 2

Characteristics of patients by cluster\(^1\).

|        | n= | Treatment history (% TN) | Sex (% Male) | Rai stage (% 3,4) | % IGHV unmutated\(^2\) | % del(17p) | % CD38\(^3\) | % Bulky disease\(^4\) | Median Age (y) | Median Pre ALC (10^9/L) | Median % LN reduction (6 MO) |
|--------|----|--------------------------|--------------|-------------------|-------------------------|------------|--------------|----------------------|----------------|-------------------------|-----------------------------|
| Cluster 1 | 20 | 40%| 60%| 80% | 70% | 45% | 30% | 15% | 69 | 139 | 69% |
| Cluster 2 | 11 | 55%| 64%| 73% | 55% | 55% | 45% | 55% | 67 | 68 | 78% |
| Cluster 3 | 10 | 40%| 30%| 40% | 30% | 50% | 50% | 80% | 66 | 36 | 77% |

\(^1\) Clusters were determined by unsupervised hierarchical clustering as depicted in Figure 2a.

\(^2\) Unmutated indicates < 2% change in *IGHV* gene sequence compared to germline.

\(^3\) CD38 positive (+) indicates ≥30% of CLL cells express CD38 above isotype control.

\(^4\) Bulky disease was defined as patients with at least one lymph node measurement >5 cm.