Lenalidomide-induced upregulation of CD80 on tumor cells correlates with T-cell activation, the rapid onset of a cytokine release syndrome and leukemic cell clearance in chronic lymphocytic leukemia

Georg Aue, Ndegwa Njuguna, Xin Tian, Susan Soto, Thomas Hughes, Berengere Vire, Keyvan Keyvanfar, Federica Gibellini, Janet Valdez, Carol Boss, Leigh Samsel, J. Philip McCoy Jr, Wyndham H. Wilson, Stefania Pittaluga, and Adrian Wiestner

1Hematology Branch, NHLBI; 2Office of Biostatistics Research, NHLBI; 3Department of Pharmacy, Clinical Center; 4Flow Cytometry Core Facility, NHLBI; 5Metabolism Branch, CCR, NCI, and 6Laboratory of Pathology, NCI, of the National Institutes of Health, Bethesda, MD, USA

ABSTRACT

Background
In chronic lymphocytic leukemia lenalidomide causes striking immune activation, possibly leading to clearance of tumor cells. We conducted this study to investigate the mechanism of action of lenalidomide and the basis for its unique toxicities in chronic lymphocytic leukemia.

Design and Methods
Patients with relapsed chronic lymphocytic leukemia were treated with lenalidomide 20 mg (n=10) or 10 mg (n=8) daily for 3 weeks on a 6-week cycle. Correlative studies assessed expression of co-stimulatory molecules on tumor cells, T-cell activation, cytokine levels, and changes in lymphocyte subsets.

Results
Lenalidomide upregulated the co-stimulatory molecule CD80 on chronic lymphocytic leukemia and mantle cell lymphoma cells but not on normal peripheral blood B cells in vitro. T-cell activation was apparent in chronic lymphocytic leukemia, weak in mantle cell lymphoma, but absent in normal peripheral blood mononuclear cells and correlated with the upregulation of CD80 on B cells. Strong CD80 upregulation and T-cell activation predicted more severe side effects, manifesting in 83% of patients as a cytokine release syndrome within 8-72 h after the first dose of lenalidomide. Serum levels of various cytokines, including tumor necrosis factor-α, increased during treatment. CD80 upregulation on tumor cells correlated with rapid clearance of leukemic cells from the peripheral blood. In contrast, neither the severity of the cytokine release syndrome nor the degree of T-cell activation in vitro correlated with clinical response.

Conclusions
Upregulation of CD80 on tumor cells and T-cell activation correlate with unique toxicities of lenalidomide in chronic lymphocytic leukemia. However, T-cell activation appears to be dispensable for the drug’s anti-tumor effects. This provides a rationale for combinations of lenalidomide with fludarabine or alemtuzumab.

Key words: chronic lymphocytic leukemia, lenalidomide, tumor flare, CD80, cytokine release syndrome.


©2009 Ferrata Storti Foundation. This is an open-access paper.
Introduction

The immunomodulating drug lenalidomide is active in chronic lymphocytic leukemia (CLL). Response rates of 47% and 32% have been reported. However, several side effects are more severe in CLL than in other diseases and have hindered the widespread use of lenalidomide in CLL. Most important among the CLL-specific side effects is the tumor flare reaction, characterized by tender lymph node swelling, low-grade fever, pain and rash. Significant morbidity and fatal adverse events have been attributed to tumor flare reaction, often in conjunction with tumor lysis syndrome. Based on these observations, clinical trials have been modified to include lower starting doses and close monitoring. In contrast, lenalidomide 80 mg daily in multiple myeloma and 25 mg daily in relapsed or refractory aggressive non-Hodgkin’s lymphoma has not caused tumor flare reaction or tumor lysis syndrome. The basis for these differences in the side effect profile of lenalidomide is unclear.

Multiple biological effects of lenalidomide have been described; however, the relative contributions of individual effects to this drug’s anti-cancer activity remain to be defined. Despite the rapid decrease in leukemic cell count in patients at the start of treatment, lenalidomide has no apparent pro-apoptotic effect on CLL cells in vitro. Lenalidomide has been thought to modulate the tumor microenvironment by downregulating cytokines including tumor necrosis factor alpha (TNFα), interleukin (IL)-6, and vascular endothelial growth factor (VEGF). However, clinical studies in CLL have reported increased serum cytokine levels during lenalidomide treatment. A postulated inhibitory effect of lenalidomide and related immunomodulating drugs. Lenalidomide can activate T cells, NK cells, or both and in some situations can lead to an expansion of immune effector cells in vivo. Lenalidomide has been found to enhance immunological synapse formation, which could increase the anti-tumor effect of immune effector cells. Upregulation of co-stimulatory molecules and/or surface antigens on CLL cells, which could render the leukemic cells more immunogenic, has been described. A partially overlapping mechanism of action has been attributed to DNA oligonucleotides containing CpG motifs that are also known to increase expression of co-stimulatory molecules and to cause immune activation. However, evidence linking immune activation and clinical response to lenalidomide is missing.

Design and Methods

Patients, clinical samples, flow cytometry and in vitro assays

CLL patients treated with lenalidomide were enrolled on the institutional review board-approved NHLBI protocol 07-H-0104 (ClinicalTrials.gov Identifier: NCT00465127). The patients’ baseline characteristics are given in Table 1. In this study lenalidomide was given in cycles: 3 weeks on, 3 weeks off. The starting dose for the first 10 patients was 20 mg, whereas that for subsequent patients was 10 mg. Allopurinol was given for 14 days in cycles 1 - 3. Side effects were classified according to the NCI common toxicity criteria (version 3.0). Control peripheral blood mononuclear cells (PBMC) from healthy volunteers (n=12) were obtained through the Department of Transfusion Medicine, NIH. Cells from patients with mantle cell lymphoma (MCL) (n=12, Online Supplementary Table S1) were collected under NHLBI protocol 04-H-0012 (ClinicalTrials.gov Identifier: NCT0071045). PBMC isolated by gradient centrifugation (Lymphocyte Separation Media, MP Biomedicals, Irvine, CA, USA) were adjusted to 5×10^6/mL and cultured in AIM V serum-free media (Invitrogen, Carlsbad, CA, USA). Lenalidomide (Sequoia Research Products, Berkshire, UK) and CpG oligonucleotides (Integrated DNA Technologies, Coralville, IA, USA) were used at doses of 2 μM and 1 μM, respectively. The mean fluorescence intensity (MFI) of surface antigens was measured by flow cytometry on BD FACSCanto II equipment and analyzed with BD FACS Diva software (BD BioSciences, San Jose, CA, USA). Mouse anti-human antibodies were: fluorescein isothiocyanate (FITC)-conjugated CD2 and CD19, phycoerythrin (PE)-conjugated CD25, CD69, CD80, CD86, and CD95, CD5 alkaline phosphatase (APC), mouse isotype G1 FITC, and mouse isotype G2a PE (BD BioScience, San Jose, CA, USA).

<table>
<thead>
<tr>
<th>Patient</th>
<th>Sex</th>
<th>Age (years)</th>
<th>Rai stage</th>
<th>WBC (x10^9/L)</th>
<th>Bulky disease</th>
<th>Prior therapies</th>
<th>GFR (mL/min)</th>
</tr>
</thead>
<tbody>
<tr>
<td>1</td>
<td>Male</td>
<td>69</td>
<td>I</td>
<td>4.6</td>
<td>yes</td>
<td>3</td>
<td>59</td>
</tr>
<tr>
<td>2</td>
<td>Male</td>
<td>78</td>
<td>III</td>
<td>23.5</td>
<td>yes</td>
<td>1</td>
<td>63</td>
</tr>
<tr>
<td>3</td>
<td>Male</td>
<td>60</td>
<td>IV</td>
<td>106</td>
<td>yes</td>
<td>1</td>
<td>74</td>
</tr>
<tr>
<td>4</td>
<td>Female</td>
<td>72</td>
<td>I</td>
<td>153</td>
<td>no</td>
<td>1</td>
<td>74</td>
</tr>
<tr>
<td>5</td>
<td>Male</td>
<td>69</td>
<td>IV</td>
<td>8.3</td>
<td>yes</td>
<td>3</td>
<td>59</td>
</tr>
<tr>
<td>6</td>
<td>Male</td>
<td>45</td>
<td>IV</td>
<td>25.5</td>
<td>yes</td>
<td>6</td>
<td>88</td>
</tr>
<tr>
<td>7</td>
<td>Male</td>
<td>77</td>
<td>I</td>
<td>77.6</td>
<td>yes</td>
<td>7</td>
<td>76</td>
</tr>
<tr>
<td>8</td>
<td>Male</td>
<td>68</td>
<td>IV</td>
<td>61.7</td>
<td>yes</td>
<td>3</td>
<td>87</td>
</tr>
<tr>
<td>9</td>
<td>Female</td>
<td>69</td>
<td>III</td>
<td>59</td>
<td>yes</td>
<td>3</td>
<td>62</td>
</tr>
<tr>
<td>10</td>
<td>Female</td>
<td>63</td>
<td>III</td>
<td>111</td>
<td>no</td>
<td>3</td>
<td>77</td>
</tr>
<tr>
<td>11</td>
<td>Male</td>
<td>53</td>
<td>II</td>
<td>11.4</td>
<td>yes</td>
<td>1</td>
<td>105</td>
</tr>
<tr>
<td>12</td>
<td>Male</td>
<td>76</td>
<td>II</td>
<td>95</td>
<td>no</td>
<td>1</td>
<td>49</td>
</tr>
<tr>
<td>13</td>
<td>Male</td>
<td>53</td>
<td>II</td>
<td>56.4</td>
<td>no</td>
<td>4</td>
<td>87</td>
</tr>
<tr>
<td>14</td>
<td>Male</td>
<td>71</td>
<td>IV</td>
<td>146</td>
<td>yes</td>
<td>5</td>
<td>73</td>
</tr>
<tr>
<td>15</td>
<td>Female</td>
<td>63</td>
<td>II</td>
<td>70</td>
<td>yes</td>
<td>1</td>
<td>56</td>
</tr>
<tr>
<td>16</td>
<td>Male</td>
<td>71</td>
<td>II</td>
<td>38</td>
<td>yes</td>
<td>5</td>
<td>64</td>
</tr>
<tr>
<td>17</td>
<td>Male</td>
<td>75</td>
<td>IV</td>
<td>139</td>
<td>yes</td>
<td>4</td>
<td>36</td>
</tr>
<tr>
<td>18</td>
<td>Male</td>
<td>62</td>
<td>I</td>
<td>6.1</td>
<td>no</td>
<td>1</td>
<td>126</td>
</tr>
</tbody>
</table>

Table 1. Patients’ characteristics at the start of lenalidomide treatment.

Patients 1 and 18 had SLL and visceral lymphadenopathy detected by computed tomography, all other patients had CLL. At least one nodal area >5 cm in largest diameter; estimated glomerular filtration rate (Cockcroft-Gault formula).
Serum cytokines

Serum samples obtained from study subjects prior to treatment (n=15), on day 4 (n=4) and on day 8 (n=13) of the first cycle of treatment were stored at -80°C. Samples from patient 15 were collected on days 1, 2, 3, 4, and 5. The cytokines and chemokines IFN-γ, TNF-α, IL-10, IL-1β, IL-1Ra, IL-2, IL-4, IL-5, IL-6, IL-10, IL-17, CXCL5, CXCL8/IL-8, CCL2, CCL3, and CCL4 were measured using Fluorokine MultiAnalyte Profiling (F-MAP) kits from R&D systems (Minneapolis, MN, USA) on a Lumines 15100 instrument (Lumex Corp. Austin, TX, USA) and analyzed using Masterplex software (Hitachi Software Engineering America, South San Francisco, CA, USA) (Online Supplementary Table S2).

Peripheral blood lymphocyte subsets and lymph node analysis

T, B and NK cells stained with CD4 APC, CD3 FITC, CD8 PE, CD45 PerCP, CD16, and CD56 PE and CD 19 APC using the BD FACS Sample Prep Assistant IVD (Becton Dickinson, Franklin Lakes, NJ, USA) were quantified at the indicated times on a FACS Caliber using FACS multiSET software (Becton Dickinson, Franklin Lakes, NJ, USA). Core biopsies from superficial lymph nodes obtained prior to lenalidomide treatment and on day 8 of cycle 1 of treatment in 11 patients were stained for CD3, CD68 and CD56. The number of CD3+ cells was scored in five representative high-power fields. Images were captured at 400x fold magnification on an Olympus Bx41 microscope (Center Valley, PA, USA).

Statistics

Changes in surface antigen expression and lymphocyte subsets were examined by paired t tests. MHF measurements of surface antigens were compared using an unpaired t test, assuming unequal variances. Pearson’s correlation coefficient was used to describe correlations. All tests were two-tailed and a p value of less than 0.05 was used as the criterion for statistical significance.

Results

Lenalidomide induces expression of co-stimulatory molecules preferentially on malignant B cells but responses are weaker than those induced by immune stimulatory CpG oligonucleotides

Lenalidomide and CpG oligonucleotides can increase the expression of co-stimulatory molecules on tumor cells. The effect of CpG differs between various B-cell malignancies. Whether the response to lenalidomide also differs between different types of B cells has not been determined. We measured the effect of lenalidomide on the expression of CD80, CD86 and CD95 on tumor cells in vitro by flow cytometry. Seventeen of the 18 patients with CLL enrolled on a phase II clinical trial using lenalidomide were studied. For comparison, we included peripheral blood B cells from healthy volunteers (n=12), and from patients with leukemic MCL (n=12). After 48 h of in vitro exposure to lenalidomide, CD80 or CD95 expression on normal B cells was unchanged. In contrast, average expression of CD80 and CD95 increased almost 2-fold on CLL B cells (Figure 1A). CD86 was upregulated on normal B cells (average increase 85±30%) and more strongly on CLL cells (average increase 125±81%). Leukemic MCL cells upregulated all three markers in the same way as CLL cells (Figure 1A). For comparison we determined the effect of CpG oligonucleotides on the same samples (Figure 1B). Upregulation of CD80 was again stronger on CLL and MCL cells than on normal B cells, but CpG increased expression of CD86 and CD95 on all three types of B cells equally and was at least twice as potent as lenalidomide in inducing CD80 and CD86. Thus, while lenalidomide preferentially upregulated co-stimulatory molecules on malignant B cells, CpG oligonucleotides were less selective but more potent stimulators.

In the presence of chronic lymphocytic leukemia cells, lenalidomide induces T-cell activation

To assess T-cell activation in response to lenalidomide, we measured expression of CD69 on T cells in the same PBMC samples used above for B cells (Figure 1C). In normal PBMC, we found only minimal T-cell activation (average increase in CD69 8±22%, p=0.1). In contrast, lenalidomide induced T-cell activation in CLL (average increase in CD69 95±95%, p<0.01). T cells from most patients with MCL (n=12) showed a weak response (average increase in CD69 25±27%, p=0.03). Consistent with the stronger effect of CpG oligonucleotides on B cells, we also found more pronounced T-cell activation in response to CpG than to lenalidomide (Figure 1D). T-cell activation in response to lenalidomide correlated strongly with CD80 upregulation on B cells (r=0.57, p=0.0001, Figure 1E) but not with upregulation of CD86 (r=0.1, p=0.53, data not shown).

Clinical presentation of immune stimulation in chronic lymphocytic leukemia: a potentially life-threatening cytokine release syndrome

The unique side effects of lenalidomide in CLL appear to be related to immune activation. This is exemplified by patient 13, who 16 h after his first dose of 10 mg developed fatigue and back pain. On hospital admission he was febrile to 40.1°C, slightly hypotensive, without a rash or change in lymphadenopathy. Lenalidomide was withheld and intravenous hydration was started. Within hours the fever rose to 41°C and the patient developed hypotensive shock that was controlled with 8 L of intravenous fluids and methylprednisolone. There was no increase in white cell count, no evidence of tumor lysis syndrome and a work-up for infection was negative. All symptoms resolved over 3 days. This clinical presentation is reminiscent of the acute cytokine release syndrome experienced by CLL patients within hours of the first administration of rituximab or with the administration of oblimersen, a CpG antisense nucleotide. We, therefore, measured the cytokine serum levels in patient 13 during the acute presentation, finding a striking increase in inflammatory cytokines, most notably TNFα and IL6 (Figure 2A). We then measured cytokine levels in prospectively collected serum from day 8, cycle 1 in an additional 12 patients (Figure 2B). The levels of the cytokines and chemokines, IL8/CXCL8, TNFα, CCL2, CCL3, and CCL4 and the soluble receptor IL-1Ra...
showed statistically significant increases on day 8 compared to pretreatment levels \((p < 0.05, \text{Figure 2B})\). IFN\(_\gamma\), IL-1\(\alpha\), IL-1\(\beta\), IL-2, IL-4, IL-5, and IL-17 were below the limit of detection in all samples (Online Supplementary Table S2). An increase in cytokines was already apparent on day 4 in the four patients available for analysis, suggesting that cytokine release within the first few days of treatment is a general reaction to lenalidomide. This interpretation is consistent with the rapid onset of clinical symptoms (see below).

While patient 13 had an exceptionally severe presentation, similar symptoms were noted in the majority of patients. To standardize the clinical presentation of the cytokine release syndrome, we uniformly recorded and graded the main symptoms and findings in all patients (Table 2). Among the 18 patients enrolled in the study, the following symptoms were observed within 8 to 72 h (average 38 h) of starting lenalidomide: fatigue (72%), pain (61%), a greater than 25% increase in lymph node size and/or in absolute lymphocyte count (ALC) over baseline (50%), fever exceeding 38\(^\circ\)C in the absence of signs of infection (44%), dehydration (39%), chills (38%) and a rise in creatinine (33%). Three out of 18 patients (17%) did not experience any of these side effects. In order to correlate the clinical presentation with laboratory and biological features we summarized...
the number and clinical grade of the individual manifestations into a score (Table 2). This score captures the degree of an inflammatory state induced by lenalidomide well, as shown by its close correlation with C-reactive protein (CRP) levels, a laboratory marker of inflammation that was rapidly induced in most patients (Figure 2C and D). Both the clinical score and CRP levels illustrate the considerable variability in lenalidomide-induced immune activation between different patients.

**In vivo effects of lenalidomide on lymphocyte subsets in chronic lymphocytic leukemia**

Recently, Ferrajoli and colleagues described that overall CD3 T-cell numbers remained stable at 3 and 6 months in 19 patients receiving continuous administration of lenalidomide. However, there are no reports on early changes during the time of the most pronounced clinical side effects and an analysis of lymphocyte subsets is not available. We, therefore, monitored B, T, and NK cell numbers during the first cycle of lenalidomide treatment (Figure 3). In the first week, almost all patients had a rapid reduction in circulating lymphocytes averaging 30-40% for B, T, and NK cells (p<0.01, Figure 3A). By the end of week 3 the normal lymphocyte subsets started to recover and after the 3-week period off drug had reached pre-treatment levels (Figure 3B). In contrast, the CLL cell count continued to decrease and by day 21, ten of 12 patients who completed 3 weeks of treatment had an average reduction of 62% (p<0.001). After the 3-week drug-free period the leukemic cell count exceeded pre-treatment levels (Figure 3C). CD56 and CD68 stained few immune effector cells into tumor sites or cell proliferation in vivo.

Table 2. Adverse events during cycle 1 of lenalidomide therapy define a clinical cytokine release syndrome.

<table>
<thead>
<tr>
<th>Patient</th>
<th>1</th>
<th>2</th>
<th>3</th>
<th>4</th>
<th>5</th>
<th>6</th>
<th>7</th>
<th>8</th>
<th>9</th>
<th>10</th>
<th>11</th>
<th>12</th>
<th>13</th>
<th>14</th>
<th>15</th>
<th>16</th>
<th>17</th>
<th>18</th>
</tr>
</thead>
<tbody>
<tr>
<td>Lenalidomide (mg)</td>
<td>20</td>
<td>20</td>
<td>20</td>
<td>20</td>
<td>20</td>
<td>20</td>
<td>20</td>
<td>20</td>
<td>20</td>
<td>10</td>
<td>10</td>
<td>10</td>
<td>10</td>
<td>5</td>
<td>10</td>
<td>10</td>
<td>5</td>
<td>10</td>
</tr>
<tr>
<td>Onset (hours)</td>
<td>24</td>
<td>40</td>
<td>48</td>
<td>48</td>
<td>8</td>
<td>48</td>
<td>24</td>
<td>24</td>
<td>48</td>
<td>72</td>
<td>16</td>
<td>12</td>
<td>72</td>
<td>72</td>
<td>24</td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Lymph node size ± ALC</td>
<td>I</td>
<td>I</td>
<td>I</td>
<td>I</td>
<td>I</td>
<td>I</td>
<td>I</td>
<td>III</td>
<td>I</td>
<td>I</td>
<td>I</td>
<td>I</td>
<td>I</td>
<td>I</td>
<td>I</td>
<td>I</td>
<td>I</td>
<td></td>
</tr>
<tr>
<td>Fever</td>
<td>I</td>
<td>I</td>
<td>I</td>
<td>I</td>
<td>I</td>
<td>I</td>
<td>I</td>
<td>I</td>
<td>I</td>
<td>I</td>
<td>I</td>
<td>I</td>
<td>I</td>
<td>I</td>
<td>I</td>
<td>I</td>
<td>I</td>
<td>I</td>
</tr>
<tr>
<td>Pain</td>
<td>I</td>
<td>I</td>
<td>III</td>
<td>IV</td>
<td>I</td>
<td>II</td>
<td>II</td>
<td>I</td>
<td>I</td>
<td>III</td>
<td>I</td>
<td>I</td>
<td>I</td>
<td>I</td>
<td>I</td>
<td>I</td>
<td>I</td>
<td>I</td>
</tr>
<tr>
<td>Fatigue</td>
<td>I</td>
<td>II</td>
<td>I</td>
<td>I</td>
<td>I</td>
<td>III</td>
<td>I</td>
<td>I</td>
<td>I</td>
<td>I</td>
<td>I</td>
<td>I</td>
<td>I</td>
<td>I</td>
<td>I</td>
<td>I</td>
<td>I</td>
<td>I</td>
</tr>
<tr>
<td>Chills</td>
<td>I</td>
<td>I</td>
<td>I</td>
<td>I</td>
<td>I</td>
<td>I</td>
<td>I</td>
<td>I</td>
<td>I</td>
<td>I</td>
<td>I</td>
<td>I</td>
<td>I</td>
<td>I</td>
<td>I</td>
<td>I</td>
<td>I</td>
<td>I</td>
</tr>
<tr>
<td>Hypotension/ dehydration</td>
<td>I</td>
<td>I</td>
<td>II</td>
<td>II</td>
<td>I</td>
<td>II</td>
<td>II</td>
<td>I</td>
<td>I</td>
<td>I</td>
<td>I</td>
<td>I</td>
<td>I</td>
<td>I</td>
<td>I</td>
<td>I</td>
<td>I</td>
<td>I</td>
</tr>
<tr>
<td>Creatinine</td>
<td>I</td>
<td>I</td>
<td>I</td>
<td>I</td>
<td>I</td>
<td>I</td>
<td>I</td>
<td>I</td>
<td>I</td>
<td>I</td>
<td>I</td>
<td>I</td>
<td>I</td>
<td>I</td>
<td>I</td>
<td>I</td>
<td>I</td>
<td>I</td>
</tr>
<tr>
<td>Hospitalization</td>
<td>Yes</td>
<td>Yes</td>
<td>Yes</td>
<td>Yes</td>
<td>Yes</td>
<td>Yes</td>
<td>Yes</td>
<td>Yes</td>
<td>Yes</td>
<td>Yes</td>
<td>Yes</td>
<td>Yes</td>
<td>Yes</td>
<td>Yes</td>
<td>Yes</td>
<td>Yes</td>
<td>Yes</td>
<td>Yes</td>
</tr>
<tr>
<td>Score</td>
<td>2.5</td>
<td>2.5</td>
<td>6</td>
<td>2</td>
<td>5.5</td>
<td>0</td>
<td>3</td>
<td>2</td>
<td>1</td>
<td>5</td>
<td>5.5</td>
<td>0</td>
<td>9</td>
<td>0</td>
<td>4</td>
<td>1</td>
<td>1.5</td>
<td>1.5</td>
</tr>
</tbody>
</table>

*Patients 6, 12 and 14 did not experience a cytokine release reaction. After patient 10, the starting dose was lowered to 10 mg daily for the first two cycles and lenalidomide dose was adjusted according to renal function (patients 12 and 17). Grading of each adverse event is given in Roman numerals according to CTCAE v3.0. The cytokine release score was calculated as follows: 0.5 point for each adverse event, plus 3 points for grade III, 4 points for grade IV.*

---

**Figure 2.** Cytokine release in CLL patients treated with lenalidomide: (A) daily after the initiation of lenalidomide in patient L13 and (B) in patients pre-treatment (squares, n=13), on day 4 (triangle, n=4) and on day 8 (diamonds, n=13) of lenalidomide treatment. (C) CRP serum levels. (D) Correlation of CRP (maximal value week 1) and cytokine release score.
NK cells and macrophages, respectively, but the numbers of these cells did not change in response to lenalidomide (data not shown). Collectively, our analysis of paired lymph node biopsies provides little evidence that tumor-infiltrating T cells are responsible for lymph node swelling in the context of flare reactions.

**T-cell activation and leukemic cell clearance are distinct effects of lenalidomide**

The severity of clinical side effects summarized in the cytokine release score correlated with the *in vitro* response to lenalidomide (Figure 4A) measured by upregulation of CD80 on CLL cells (*r*=0.71, *p*=0.001), and CD69 on T cells (*r*=0.75, *p*=0.001), and inversely with treatment-induced changes in T-cell numbers (*r*=-0.64; *p*=0.006, data not shown). There was no correlation between the clinical score and the decrease in leukemic cell count, age, RAI stage, number of prior therapies, renal impairment, or bulky disease. The average score in all patients was 2.89 (range, 0-9) with no difference between the 20 mg and the 10 mg cohort (mean scores of 2.95 and 2.81, respectively, *p*=0.9).

Next we tested the association of clinical and biological characteristics with leukemic cell clearance. We found a strong correlation between the *in vitro* response of CLL B cells to lenalidomide, measured by CD80 upregulation, and the reduction in leukemic cells during the first cycle of treatment (Figure 4B). In contrast, upregulation of CD86 and CD95 was not correlated with treatment response. Metrics of immune activation, including CD69 upregulation on T cells (Figure 4B), the cytokine release score, individual cytokine serum levels, or CRP did not correlate with leukemic cell clearance *in vivo*, suggesting that T-cell activation and anti-leukemic activity of lenalidomide are separate effects.

**Discussion**

The clinical use of lenalidomide in CLL has been com-
The cytokine release syndrome in CLL appears to be determined by tumor characteristics, which raises the possibility of prospectively identifying patients at risk of severe reactions. The main predictor of leukemic cell clearance, which reached an average of 62% by day 21, was lenalidomide-induced upregulation of CD80 on CLL cells (Figure 4B). Response data are currently not mature enough to test whether CD80 could serve as a predictor of overall response, and this should be addressed.
Lenalidomide-induced immune activation in CLL

prospectively. It will also be interesting to test whether CD80 upregulation on tumor cells correlates with clinical responses in other B-cell malignancies. Whether CD80 expression is just a marker of unidentified biological effects of lenalidomide on CLL cell biology or whether it has a functional role in mediating leukemic cell clearance remains to be determined. In this regard it is interesting to note that CD80 is a signaling molecule that has been reported to transmit growth inhibitory and pro-apoptotic signals in B-cell lymphoma cells.11,17

While we cannot exclude that lenalidomide could have different mechanisms of action in different diseases, we favor the hypothesis that T-cell activation in CLL causes increased toxicity independently of antitumor effects. This provides a rationale to test the combination of lenalidomide with immunosuppressive agents such as glucocorticoids, purine analogs or alemutzumab.

Authorship and Disclosures

This work was presented in part at the annual American Society of Hematology meeting December 6th to 9th 2008 in San Francisco, CA, USA.

GA, NN, XT, SS, TH, JV, CB, WHW and AW were investigators of the study. GA, NN, BV, KK, FG, LS, JPM, and SP performed the laboratory work for this study. GA, NN, XT and AW participated in the statistical analysis. GA, JPM, and AW coordinated the research. GA, NN, XT, SP and AW wrote the manuscript. The authors report no potential conflicts of interest.

References


