Tumor-Associated Macrophages in Pediatric Classical Hodgkin Lymphoma: Association with Epstein-Barr Virus, Lymphocyte Subsets, and Prognostic Impact

Mário Henrique M. Barros1, Rocio Hassan2, and Gerald Niedobitek1

Abstract

Purpose: Tumor-infiltrating macrophages are associated with adverse outcome in adult classical Hodgkin lymphoma (cHL). We have previously shown age-related changes in the lymphocyte composition of pediatric cHL. We therefore hypothesized that the number, function, and prognostic impact of macrophages in pediatric cHL would be different from adult cases.

Experimental Design: We analyzed the number of macrophages and dendritic cells (DC) in the tumor microenvironment of pediatric cHL by immunohistochemistry. Results were analyzed in context of age, histologic characteristics, Epstein-Barr virus (EBV) status, clinical follow-up, and our previous study of T-cell populations in these cases.

Results: One hundred cHL cases were studied, including 69% nodular sclerosis and 23% mixed cellularity cases. A total of 44.8% of cases were EBV-positive. Patients ≤10 years displayed more CD14+ cells (P = 0.025). In comparison with nodular sclerosis, mixed cellularity was characterized by higher numbers of CD14+ (P = 0.003) and CD163+ cells (P = 0.027). EBV+ cases exhibited higher numbers of CD14+ (P < 0.0005), CD68+ (P = 0.005), and CD163+ cells (P = 0.02). CD68-positive cells did not display an effect on outcome. Worse overall survival was observed in cases with CD163/CD8 ratio ≥2 (P = 0.007). High numbers of CD163+ cells were associated with worse progression-free survival (PFS; P = 0.015). Furthermore, high numbers of CD163+ and granzyme B+ cells were associated with worse PFS in EBV-negative (P = 0.005) but not in EBV-positive cases.

Conclusion: Our results suggest that macrophage composition in pediatric cHL is distinct from adults. Functional status of macrophages and their value as prognostic indicators in pediatric cHL may depend on EBV status. Clin Cancer Res; 18(14); 3762–71. ©2012 AACR.

Introduction

Classical Hodgkin lymphoma (cHL) microenvironment is modulated by cytokines and chemokines produced by Hodgkin/Reed–Sternberg (HRS) cells and is thought to contribute to tumor cell growth and escape from immune-surveillance (1). We have shown recently that in Epstein-Barr virus (EBV)+ pediatric cHL the tumor microenvironment is characterized by a cytotoxic/T-helper cell 1 (Th1) phenotype (2). Furthermore, cases with a small number FOXP3+ regulatory T cells (Treg) and a FOXP3/CD8 ratio less than 1 in the tumor microenvironment were associated with an improved outcome. We, therefore, hypothesized that in pediatric EBV+ cHL an effective cytotoxic immune response directed against viral or tumor antigens may be triggered in the tumor microenvironment (2–3).

Recent studies have consistently reported that high numbers of tumor-associated macrophages (TAM) are associated with adverse outcome in adult cHL (4–6). As yet, the specific role of TAM in pediatric cHL has not been studied.

Macrophages can be polarized toward a proinflammatory phenotype (M1 macrophages) with tumoricidal activity or toward a regulatory function (M2 macrophages), which are characterized by functions in tissue repair, remodeling and promotion of Th2 immune response (7–9). It is generally thought, that M2 macrophages are predominantly found in the microenvironment of malignant tumors. In agreement with this notion, macrophages expressing CD163 (considered to be a marker of M2 macrophages) were associated with adverse outcome in a study of adult cHL (5).

Dendritic cells (DC) are critically important in the initiation and regulation of immune responses, driving Th1 or...
Translational Relevance

Tumor-infiltrating macrophages are potential independent indicators of poor outcome in classical Hodgkin lymphoma (cHL) in adults and represent possible therapeutic targets. In this study, we investigated macrophages and dendritic cells in pediatric cHL taking into account Epstein-Barr virus (EBV) status and lymphocyte composition of the tumor microenvironment. Our results show that the prognostic role of macrophages in pediatric cHL is affected by EBV status of the Hodgkin and Reed–Sternberg cells. Furthermore, we suggest that use of microenvironment variable as prognostic indicators in cHL may require the combined analysis of cell populations from the innate and adaptive immune systems. Our results emphasize differences in the microenvironment between adult and pediatric cHL and suggest that it may be necessary to reevaluate studies of adult patients taking EBV status into account.

Materials and Methods

Patients

One hundred HIV-negative children and adolescents (up to 18 years old) diagnosed with cHL at the Instituto Nacional do Câncer (INCA, Brazil) between 1999 and 2006 were included in this study. The clinical features of these cases have been previously described (2). All patients were treated according to standard pediatric protocols as previously described (2). For 5 cases, there was not sufficient clinical data and these were excluded from the survival analysis. Children were classified in 2 age groups (≤10 years vs. >10 years) to investigate the characteristics of macrophages and DCs in younger versus older children (17–21). This study was approved by the INCA Ethics Committee.

Histology

Diagnosis of cHL was established by morphologic criteria according to the World Health Organization classification (22). All cases were independently reviewed by 2 pathologists (M.H.M. Barros and G. Niedobitek) as previously described (2).

Tissue microarray design and immunohistochemistry

Construction of tissue microarray (TMA) blocks has been described previously (2). From each case, 2 1-mm-diameter cores, selected from 2 different tumor areas rich in neoplastic cells, were included. TMA blocks were sectioned at a thickness of 3 mm. Antigen retrieval was carried out by heat treatment in a pressure cooker for 1 minute. Buffers used for antigen retrieval and primary antibodies are listed in the Supplementary Material (Supplementary Table S1). Briefly, the antibodies used were CD14 (clone 7, Novocastra), CD68 (clone PGM1, Dako), CD163 (clone ID6, Novocastra), CD83 (clone 1H4b, Novocastra), and CD207 (Langerin; clone 12D6, Abcam). After incubation with primary antibody (30 minutes), immunodetection was conducted using Zytocach Plus HRP Polymer Kit (Zytomed Systems; Supplementary Material, Supplementary Table S1), using 3,3′-diaminobenzidine (DAB) chromogen as substrate. Sections were counterstained with hematoxylin. External and internal controls included in the TMA were taken into consideration to interpret staining results.

EBV detection

Latent EBV infection was determined in all cases by in situ hybridization (ISH) with fluorescein-conjugated probes for EBV-encoded RNAs (EBER-ISH) and by immunohistochemistry against latent membrane protein 1 (LMP1) as described (23).

Computer-assisted microscopic analysis

For cell subset quantitative evaluation, each core was photographed using AxioCam MRc camera (Zeiss) at a ×200 magnification. The numbers of labeled macrophages and DCs were determined per 1 mm² using the image analysis software HISTO (Biomas).

Thresholds

For each cell subset, the 25th and 50th percentiles were used to categorize the intensity of the cell infiltration. Cases displaying no labeled cell in both cores were considered not evaluable for technical reasons.

Statistical analysis

Pearson χ² and Fisher exact test were used to test association between dichotomous variables. Mann–Whitney U test was used to compare medians, whereas Spearman rank correlation was used to test association between continuous variables. First-order partial correlations were used to correlate 2 or more variables while keeping constant 1 or more additional variables and were computed as rank correlations (continuous variables) or point biserial rank correlations (categorical vs. continuous variables). Differences were considered significant at P < 0.05 in 2-tailed tests. Hierarchical cluster analysis using average linkage and binary simple matching measure allowed to explore the structure of association among variables of
the tumor microenvironment and EBV status. Primary treatment was defined as a failure if the lymphoma had progressed at any time after the initiation of therapy; treatment success was defined as the absence of progression or relapse. Progression-free survival (PFS) was the interval (in months) from diagnosis to progression at any time, relapse from complete response, or initiation of new, previously unplanned treatment or to the last follow-up in the patients with treatment success. Overall survival (OS) refers to the interval (in months) from the diagnosis to death or last follow-up. Survival distributions were estimated by the Kaplan–Meier method and differences were compared using log-rank test. Multivariate Cox proportional hazard regression method was used to determine the independent prognostic value of statistically significant variables in univariate analyses. Data were analyzed using Statistical Package for the Social Sciences 13.0 (SPSS).

Results
Clinicopathologic data of this pediatric group have been described previously (2). In brief, the age at diagnosis ranged from 3 to 18 years (median, 14 years) and 27% of children were ≤10 years. The majority of patients had stage I/II disease (59 of 95, 62.1%) and favorable clinical presentation (50 of 95, 52.6%). Extranodal disease was observed in 11 children (11 of 95, 11.6%). Nodular sclerosis was the predominant subtype (69 of 100, 69%), followed by mixed cellularity (23 of 100, 23%). Granulomas were observed in 37% of cases (37 of 100 cases; ref. 2). As described previously, EBV-positive HRS cells were detected in 44.8% of cases and no association with age groups was observed (3). A detailed description of EBV association in relation to histologic features has been published previously (2, 3). Distribution of lymphocyte subsets in the tumor microenvironment in relation to age group, histology, EBV status, and their prognostic impact has been reported previously (2, 3) and are summarized in Supplementary Table S3. The main clinical and histologic characteristics of the patients are described in the Supplementary Material (Supplementary Table S2).

Macrophages and DCs in pediatric cHL
CD14+ monocyte/precursor cells, as well as CD163+ macrophages, CD83+ and CD207+ DCs were distributed uniformly in the tumor microenvironment, without specific distribution pattern (Fig. 1). CD68+ cells also showed uniform distribution in the majority of cases, but in 5 cases a rosette-like arrangement around HRS cells was observed.

The results of the quantitative analysis of CD68+, CD163+, CD14+, CD207+, and CD83+ cells are summarized in Table 1. In general, the number of CD68+ cells was higher than that of CD163+ cells and of CD14+ cells. As expected, a strong correlation between CD68+ and CD163+ cells was observed (P < 0.001), as well as between CD68+ and CD14+ cells (P < 0.001).

The numbers of CD83+ cells and CD14+ cells were similar. In contrast, Langerin+ cells were infrequently detected in the tumor microenvironment (Supplementary Material, Supplementary Fig. S1). No significant bivariate correlation was observed between CD83+, CD207+, and CD14+ cells.

TAMs are associated with clinical and histologic features
The young age group (≤10 years) was characterized by a higher number of CD14+ cells (median, 70 vs. 44 cells/mm² for the old age group, P = 0.025, Mann–Whitney U test). No other age-related differences were observed.

The numbers of CD14+, CD68+, CD163+, CD83+, and CD207+ cells were not associated with clinical stage, favorable or unfavorable clinical presentation, B symptoms, number of involved anatomic sites, lymphopenia, or anemia.

In comparison with nodular sclerosis subtype, mixed cellularity subtype was characterized by higher numbers of CD14+ (median, 105 vs. 49 cells/mm²) to nodular sclerosis,

Figure 1. Examples of the diversity of immunostains for antigen-presenting cells markers. Some cases displayed high numbers of CD68+ cells (A), whereas in others, only few CD68+ cells were seen (D; original magnification, ×10). Similarly, there were cases with high numbers of CD163+ cells (B), whereas others exhibited few number of these cells (E; original magnification, ×10). As a typical example of CD83 expression by DCs (original magnification, ×40), CD207+ cells were infrequent in the tumor microenvironment of pediatric cHL (F; original magnification, ×40).


Cases with granulomas exhibited a higher number of CD14\(^+\) cells (median, 64 vs. 49 cells/mm\(^2\) without granulomas, \(P = 0.04\), Mann–Whitney \(U\) test) and CD163\(^+\) cells (median, 3 vs. 1 cell/mm\(^2\) without granulomas, \(P = 0.03\), Mann–Whitney \(U\) test). Although not statistically significant, the cases with granulomas also exhibited higher numbers of CD68\(^+\) cells (median, 310 vs. 269 cells/mm\(^2\) without granulomas, \(P = 0.1\) and CD163\(^+\) cells (median, 167 vs. 126 cells/mm\(^2\), \(P = 0.2\)).

### Table 1. Description of the macrophage and DC subsets in the tumor microenvironment of pediatric cHL and their association with PFS and OS

<table>
<thead>
<tr>
<th>Variable</th>
<th>Labeled cells/mm(^2)</th>
<th>Cases analyzed (%)</th>
<th>5-Year PFS rate (%)</th>
<th>Univariate analysis ((P)) for PFS</th>
<th>5-Year OS rate (%)</th>
<th>Univariate analysis ((P)) for OS</th>
</tr>
</thead>
<tbody>
<tr>
<td>CD14(^+)</td>
<td>Range 0 to 524 (83.1/53)</td>
<td>20/78 (25.6)</td>
<td>78.9</td>
<td>0.8</td>
<td>85</td>
<td>0.6</td>
</tr>
<tr>
<td>CD68(^+)</td>
<td>Range 44 to 601 (290.8/284)</td>
<td>20/81 (24.7)</td>
<td>78.9</td>
<td>0.6</td>
<td>84.2</td>
<td>0.8</td>
</tr>
<tr>
<td>CD163(^+)</td>
<td>Range 0 to 560 (164.1/143)</td>
<td>20/79 (25.3)</td>
<td>84.2</td>
<td>0.2</td>
<td>85</td>
<td>0.8</td>
</tr>
<tr>
<td>CD207(^+)</td>
<td>Range 0 to 32 (4.3/2)</td>
<td>49/76 (64.5)</td>
<td>74.4</td>
<td>0.6</td>
<td>86.7</td>
<td>0.9</td>
</tr>
<tr>
<td>CD83(^+)</td>
<td>Range 11 to 410 (66.9/55)</td>
<td>21/79 (26.6)</td>
<td>80</td>
<td>0.6</td>
<td>90</td>
<td>0.6</td>
</tr>
</tbody>
</table>

\(P = 0.003\), Mann–Whitney \(U\) test) and CD163\(^+\) cells (median, 191 vs. 117 cells/mm\(^2\) to nodular sclerosis, \(P = 0.027\), Mann–Whitney \(U\) test; Supplementary Material, Supplementary Table S3).

Cases with granulomas exhibited a higher number of CD14\(^+\) (median, 64 vs. 49 cells/mm\(^2\) without granulomas, \(P = 0.04\), Mann–Whitney \(U\) test) and CD207\(^+\) cells (median 3 vs. 1 cell/mm\(^2\) without granulomas, \(P = 0.03\), Mann–Whitney \(U\) test). Although not statistically significant, the cases with granulomas also exhibited higher numbers of CD68\(^+\) cells (median, 310 vs. 269 cells/mm\(^2\) without granulomas, \(P = 0.1\) and CD163\(^+\) cells (median, 167 vs. 126 cells/mm\(^2\), \(P = 0.2\)).

### TAM and DCs are associated with lymphocyte subsets in tumor microenvironment

The analysis of lymphocyte subset distribution in this pediatric cHL series has been published previously, and all data about the quantitative assessment of lymphocytes have been taken from those studies (refs. 2, 3; Supplementary Material, Supplementary Table S3).

Numbers of macrophages and DCs were lower than the total number of CD3\(^+\) T cells. However, the number of CD68\(^+\) cells was higher than that of CD20\(^+\) B cells and the number of CD163\(^+\) cells was similar to that of CD8\(^+\) cells (Supplementary Material, Supplementary Fig. S1).
As previously shown, the size of these lymphocyte subpopulations differed according to age, with a more cytotoxic composition observed in the ≤10 years group compared with a more suppressive profile in children older than 10 years (ref. 2; Supplementary Material, Supplementary Table S3). To investigate whether macrophages and DCs are associated with lymphocyte subpopulations a partial rank correlation analysis was carried out, controlled by age.

The details of this analysis are provided in the Supplementary Material (Supplementary Table S4). In general, the numbers of CD14⁺ cells showed a direct correlation with the numbers of TBET⁺ (P < 0.0005), CD8⁺ (P = 0.019), TIA1⁺ (P = 0.024), and granzyme B⁺ (P = 0.009) cells.

The numbers of CD68⁺ cells were directly correlated with the numbers of FOXP3⁺ (P < 0.0005), TIA1⁺ (P = 0.001), and granzyme B⁺ (P = 0.001) cells (Supplementary Material, Supplementary Table S4).

Numbers of CD163⁺ cells were directly correlated with those of TBET⁺ (P = 0.04), CD8⁺ (P = 0.025), TIA1⁺ (P = 0.001), and granzyme B⁺ (P < 0.0005) cells (Supplementary Material, Supplementary Table S4).

Finally, numbers of CD83⁺ cells were directly correlated with the numbers of CD4⁺ (P = 0.01), FOXP3⁺ cells (P = 0.003), and CD20⁺ lymphocytes (P = 0.018; Supplementary Material, Supplementary Table S4).

**EBV status of HRS cells is associated with the number of TAM and DCs in pediatric cHL.**

Because EBV has been shown to modulate the tumor microenvironment composition in adult and childhood cHL (2, 24), a preliminary data classification strategy using hierarchical cluster analysis was conducted to identify underlying patterns of tumor microenvironment cell subsets according to EBV status (Fig. 2). In this analysis, 2 distinct clusters emerged: cluster I included the EBV-associated cases and cases with high numbers of CD8⁺, TIA1⁺, TBET⁺, CD14⁺, CD68⁺, CD163⁺, and CD207⁺ cells; cluster II was mainly composed of EBV-negative cases and those with high numbers of FOXP3⁺, CMAF⁺, and CD83⁺ cells.

Statistical associations between the numbers of macrophages/DCs and EBV status of HRS cells were investigated with the Mann–Whitney U test. EBV-associated cases displayed higher numbers of CD14⁺ (median, 83 vs. 38 cells/mm² in EBV-negative cases, P < 0.0005), CD68⁺ (median, 331 vs. 234 cells/mm² in EBV-negative cases, P = 0.005), and CD163⁺ cells (median, 146 vs. 111 cells/mm² in EBV-negative cases, P = 0.02). No association between EBV status of HRS cells and numbers of DCs was observed. Specifically, we did not observe any association of EBV status of HRS cell with the number of tumor-infiltrating CD207⁺ cells.

![Dendrogram using average linkage obtained by hierarchical cluster analysis. Two main identified clusters (I and II) are indentified by brackets.](image-url)

**Figure 2.** Dendrogram using average linkage obtained by hierarchical cluster analysis. Two main identified clusters (I and II) are indentified by brackets. Num, order of variable input. CD14 high, high number of CD14⁺ cells/mm² (>50th percentile). CD163 high, high number of CD163⁺ cells/mm² (>50th percentile). TBET high, high number of TBET⁺ cells/mm² (>50th percentile). CD8 high, high number of CD8⁺ cells/mm² (>50th percentile). EBV positive, EBV-associated cHL. CD68 high, high number of CD68⁺ cells/mm² (>50th percentile). CD20 high, high number of CD20⁺ cells/mm² (>50th percentile). CMAF high, high number of CMAF⁺ cells/mm² (>50th percentile). Langerin high, high number of Langerin (CD207⁺) cells/mm² (>50th percentile). EBV-negative, absence of EBV association in cHL. CMAF high, high number of CMAF⁺ cells/mm² (>50th percentile). Granzyme B high, high number of granzyme B⁺ cells/mm² (>50th percentile). FOXP3 high, high number of FOXP3⁺ cells/mm² (>50th percentile). CD83 high, high number of CD83⁺ cells/mm² (>50th percentile).
To assess the correlations between the numbers of macrophages, DCs, and lymphocytes subpopulations, independently of the EBV status in HRS cells, point biserial rank correlations were carried out, using EBV status as controlling variable (Table 2). CD14<sup>+</sup> cells were directly correlated with the number of TBET<sup>+</sup> cells independently of EBV status. The same was observed for CD68<sup>+</sup> and granzyme B<sup>+</sup> cells, CD163<sup>+</sup> and granzyme B<sup>+</sup> cells, and CD83<sup>+</sup> and FOXP3<sup>+</sup> cells.

On the other hand, a positive correlation was observed between CD68<sup>+</sup> and FOXP3<sup>+</sup> cell numbers only in EBV-negative cases, indicating that this is the source of the overall association observed between these 2 subpopulations in the microenvironment (see earlier). These analyses in EBV-negative cases also disclosed a relationship of CD163<sup>+</sup> cells with both TBET<sup>+</sup> cells and TIA1<sup>+</sup> cells (Table 2).

### TAM and DCs are associated with outcome

The OS at 60 months was 89.4%; stage and type of clinical presentation did not have impact on the OS (2). No association was observed between single macrophage or DC populations and OS. A worse 5-year OS rate was observed in cases with CD163/CD8 ratio >2 (64.7% vs. 91.4% for ratio <2, P = 0.009, log-rank test) and with CD83/CD20 ratio >2 (50% vs. 89.7% for ratio <2, P = 0.001, log-rank test; Fig. 3). The other ratios between macrophages, DCs, and lymphocyte populations were not associated with survival. In the multivariate Cox regression, CD83/CD20 ratio ≥2 maintained statistical significance [P = 0.008; HR, 7.3; 95% confidence interval (CI), 1.6–31.5], whereas CD163/CD8 ratio showed a borderline significance (P = 0.06; HR, 3.3; 95% CI, 0.9–11.6).

PFS at 60 months was 78.6% (2). High numbers of CD163<sup>+</sup> cells (≥50th percentile) were associated with worse 5-year PFS rate (P = 0.015, log-rank test), whereas CD68<sup>+</sup> cells showed no influence on survival (Fig. 4). Extranodal disease (54.5% vs. 81% for absence of extranodal disease, P = 0.028, log-rank test), leukopenia (50% vs. 80% for absence of leukopenia, P = 0.034, log-rank test), and presence of granulomas (63.6% vs. 84.5% for absence of granulomas, P = 0.026, log-rank test) were also associated with worse 5-year PFS rate (2). Stage and type of clinical presentation did not impact on PFS (2). Considering the variables statistically significant in the log-rank test, only extranodal disease maintained statistical significance in the Cox regression (P = 0.025; HR, 3.4; 95% CI, 1.1–10.2).

We have reported previously that high numbers of granzyme B<sup>+</sup> cells were also associated with worse 5-year PFS rate (69.6% vs. 90.9%, P = 0.045, log-rank test; ref. 2). Next, we combined both microenvironment variables significantly associated with PFS: CD163<sup>+</sup> and granzyme B<sup>+</sup> cells. This allowed the distinction of 3 patient groups: group 1, high numbers of granzyme B<sup>+</sup> cells and CD163<sup>+</sup> cells; group 2, high numbers of granzyme B<sup>+</sup> cells or high number of CD163<sup>+</sup> cells; and group 3, low numbers of granzyme B<sup>+</sup> cells and CD163<sup>+</sup> cells. On the basis of this approach, the worst PFS was observed in cases with high numbers of both, CD163<sup>+</sup> and granzyme B<sup>+</sup> cells (P = 0.031, log-rank test).
In the Cox regression, extranodal disease (P = 0.01; HR, 4.1; 95% CI, 1.4–12.1) and the combination of CD163+ and granzyme B+ cells (P = 0.038; HR, 2.2; 95% CI, 1.0–4.9) maintained the statistical significance.

EBV infection of HRS cells may influence the prognostic significance of CD163+ cells in the tumor microenvironment

Given that EBV appears to be able to modulate the tumor microenvironment composition and local EBV-specific immunity (2, 24, 25), the impact of the numbers of CD14+, CD68+, CD163+, CD207+, and CD83+ cells on PFS was investigated, according to EBV-status. In the EBV-negative group, a worse PFS rate was observed for cases with high numbers of CD163+ cells (>50th percentile; 44.4%, P = 0.008, log-rank test), whereas no influence of a high number of CD163+ macrophages was observed in the EBV-positive group, although this was the group that generally exhibited higher numbers of CD163+ macrophages. The other cell populations showed no influence on PFS when cases were stratified by EBV status.

Concerning the combined effect of granzyme B+ and CD163+ cells, again, the impact on PFS was observed only in the EBV-negative group, with the worst PFS rate occurring in cases with high numbers of both CD163+ and granzyme B+ cells (35.7%, P = 0.005, log-rank test; Fig. 4).

The effects of microenvironment variable on 5-year OS rate were also affected by EBV status. CD163/CD8 ratio ≥2 and CD83/CD20 >2 were associated with worse prognosis only in EBV-negative cases (50%, P = 0.005 and 40%, P = 0.002, respectively; log-rank test; Fig. 3).

Discussion

In comparison with adults, children display distinct physiologic immunologic features (17–21). In pediatric cHL, these distinct features may contribute to a different immune response against HRS cells reflected by qualitative and quantitative differences in lymphocyte subset distribution in the tumor microenvironment (2). Adaptive and innate immune systems are integrated and cross-talk between these systems contributes to a robust and balanced
To understand the nature of the local immune responses in the tumor microenvironment, it is essential to analyze the joint analysis of TAM, DCs, and lymphocytes. Therefore, in the context of solid tumors (7, 9), we did not observe a direct immune response (15–16). Hence, the CD163 expression of CD8 T cells is also more prevalent in chL cases of the younger age group. It is unknown if age-related changes exist in the frequencies of CD14+ cells in peripheral blood or tissues; therefore, at present it is not possible to say if our result is secondary to a particular immune response against HRS cells in the young age group or merely reflects an age-related physiologic phenomenon.

CD14 is expressed on cells of the myelomonocytic lineage, including monocytes, a proportion of macrophages and Langerhans cells (26). Though our study did not observe age-related changes in the numbers of macrophages or Langerhans cells, it is possible that the majority of the CD14+ cells represent monocytes (macrophages precursors). However, this issue requires further analysis.

As described previously in adult cHL (5, 27), we observed higher numbers of TAM in EBV-positive cases, independently of their maturation/activation status, as evidenced by CD14, CD68, and CD163 immunohistochemistry. It is well recognized that the immune axis shaped by T11+ cells, that is, CD8+ cells, macrophages and DCs, is essential for the control of virus infection (16, 28). In conjunction with our previous observations (2), these results raise the possibility that in EBV-associated pediatric chL, an antiviral immune response may be triggered against viral proteins present on HRS cells, resulting in an increased recruitment of macrophages and cytotoxic/T11+ cells to the tumor microenvironment (2). In addition, a putative cross-talk of T11+ and CD8+ cells with macrophages (15), might explain the direct correlation observed between the number of macrophages and the number of T11+ cells, seen in this study. At present, it is not understood how monocyte/macrophages are recruited to the microenvironment of HRS cells and how these and other tumoral cells contribute to the final phenotype.

In contrast to in vitro studies, animal studies or investigations of solid tumors (7, 9), we did not observe a direct correlation between numbers of putative M2 macrophages (CD163+ cells) and the numbers of Tregs. Moreover, unexpectedly, we noticed a direct correlation between the numbers of CD163+ cells and granzyme B+ cells, independently of EBV status. These results suggest that a high number of CD163+ macrophages per se do not reflect in an immunosuppressive tumor microenvironment composition. It is possible that CD163+ macrophages in a cytotoxic environment may either represent part of a “negative-feedback loop” to avoid an excessive tissue destruction by cytotoxic T cells or alternatively, may have different differentiation/activation status, as it has been described in non–small cell lung tumor (29) and in cutaneous squamous cell carcinoma (30).

Moreover, it is of note that in our pediatric series, therapy response was not negatively affected by the presence of TAM in EBV+ cases, despite the high number of these cell subset that characterized the EBV+ cases. This reinforces the idea that, in the EBV+ cases, most TAMs are not of the M2 protumoral subset and that CD163 probably is not a definitive marker of M2 polarization of macrophages.

In addition, we found a correlation between the numbers of CD68+ macrophages and Tregs in the EBV-negative cases, which again points to differences in the macrophage differentiation in the EBV+ and EBV- groups.

We did not observe an independent prognostic impact of CD68+ cells on OS, contrary to what has been reported for adult chL (3). We used the macrophage-specific PG-M1 clone, contrary to many studies of adult chL, which used CD68 KP1 clone to identify macrophages (4, 5, 33). As KP1 is known to cross-react with different cell populations, including large lymphocytes (34, 35), it is possible other cell populations stained with the KP1 antibody contribute to the adverse prognostic effect observed in previous studies. The absence of impact on the OS was also observed for EBV+ cases. It remains unclear in our specific pediatric population, if the absence of impact of macrophages and DC subpopulations (when analyzed separately) on OS was due to the success of rescue therapy or secondary to any other factor. Nevertheless, we observed that CD163/CD8 ratio >2 and CD83/CD20 >2 were associated with worse OS, suggesting that the immune system may contribute to treatment response; and that tumor microenvironment variables can be informative about this.

High numbers of CD163+ cells were associated with worse PFS only at the univariate level, in line with previously published results in adult cases (5). However, when this cell population was analyzed together with granzyme B+ cells, we observed a worse PFS in cases with high numbers of both granzyme B+ cells and CD163+ cells in both univariate and multivariate analysis. Because of the complexity of microenvironment cell interactions, prognostic prediction in chL may require combining microenvironment variables in an index or score, as shown here for the combination of the numbers of CD163+ cells and granzyme B+ cells. A similar strategy was also proposed in a previous study of adult chL (6) and may be fruitful for future clinical studies focused on the modulation of tumor microenvironment.

CD83 (clone 1H4b) has been shown to be a specific marker for mature and activated DCs and does not cross-react with macrophages (26, 36). In support of this notion, in our study, numbers of CD83+ cells did not correlate with those of CD14+, CD68+, or CD163+ cells. Also, we observed that the numbers of CD83+ DCs were directly
correlated with the numbers of FOXP3+ cells. As it has been described that pDCs are able to induce primary Treg cell differentiation (13, 14), it is possible that part of these CD83+ cells were pDC and that they were engaged in Treg cell differentiation. However, the CD83 antibody is not able to differentiate myeloid DCs from pDCs (36, 37) and this issue requires further investigation. The prognostic impact of tumor-associated DCs on survival of patients with cancer is controversial; in some studies a higher number of these cells was associated with a better outcome (38, 39), whereas in others, the specific pDC subset has been associated with worse outcome (40–42). Here, an excess of DCs was associated with worse OS, when analyzed in relation to B cells (CD83/CD20 ratio ≥2). This highlights the necessity to understand better the contribution of DCs to the immune response against HRS cells.

CD207 (Langerin) is a type II membrane-associated C-type lectin exclusively expressed by Langerhans cells, a subtype of DC (43). Recently, it has been suggested that the presence of CD207+ Langerhans cells in the tissue microenvironment may be an exclusive feature of EBV-associated disease, including cHL (44). Our results are different in 2 respects. First, we found CD207+ cells in comparable numbers in both, EBV-positive and EBV-negative pediatric cHL cases. Moreover, in our study, these represented only a very minor cell population. The reasons for these discrepancies are currently unclear but likely are due to technical differences, for example, related to the choice of CD207 antibodies. The Langerin (CD207) antibody (clone 12D6, Abcam) used in our study produced reproducible staining of epidermal Langerhans cells in control experiments (not shown). Thus, our results suggest that CD207+ cells are not specifically recruited to the microenvironment of pediatric cHL and, at least in this setting, the presence of CD207+ cells is not a specific feature of EBV+ cases. We are aware that the number of cases in this study imposes limitation in relation to the analysis of prognostic impact and prospective studies are mandatory to confirm our results. However, our series of sequential cases with similar distribution in relation to EBV status is appropriate for the immunologic evaluation conducted here.

In summary, our results suggest that in pediatric cHL the macrophage composition is numerically and functionally distinct from adults and that the functional status of macrophages and their value as prognostic indicators in pediatric cHL may depend on EBV status of HRS cells. Finally, our results emphasize the complexity of the cHL microenvironment and suggest that for an assessment of the impact on outcome, multiple cell populations of the innate and adaptive immune systems may have to be considered together.

Disclosure of Potential Conflicts of Interest
No potential conflicts of interest were disclosed.

Authors’ Contributions
Conception and design: M.H.M. Barros, G. Niedobitek
Development of methodology: M.H.M. Barros
Acquisition of data (provided animals, acquired and managed patients, provided facilities, etc.): M.H.M. Barros, G. Niedobitek
Analysis and interpretation of data (e.g., statistical analysis, biostatistics, computational analysis): M.H.M. Barros, R. Hassan
Writing, review, and/or revision of the manuscript: M.H.M. Barros, R. Hassan, G. Niedobitek
Administrative, technical, or material support (i.e., reporting data, constructing databases): M.H.M. Barros
Study supervision: R. Hassan, G. Niedobitek

Acknowledgments
The authors thank Dr. Fernando Soares (Pathology Department of A.C. Camargo Hospital, São Paulo, Brazil) for kindly constructing the TMA and Dr. Luizpold Distel for kindly providing the image analysis software.

Grant Support
This work was supported by the Fundação de Amparo à Pesquisa do Estado do Rio de Janeiro [TAPERI E26/110.432/2010] and INCI para Controle do Câncer [grants CNPq 573806/2008-0 and TAPERI E26/170.026/2008; Brazil] and the Berliner Krebsgesellschaft [NFII 201004]. M.H.M. Barros is supported by Alexander von Humboldt Foundation. The costs of publication of this article were defrayed in part by the payment of page charges. This article must therefore be hereby marked advertisement in accordance with 18 U.S.C. Section 1734 solely to indicate this fact.

Received January 17, 2012; revised April 13, 2012; accepted May 11, 2012; published OnlineFirst May 29, 2012.

References
Macrophages in Pediatric Classical Hodgkin Lymphoma

Tumor-Associated Macrophages in Pediatric Classical Hodgkin Lymphoma: Association with Epstein-Barr Virus, Lymphocyte Subsets, and Prognostic Impact

Mário Henrique M. Barros, Rocio Hassan and Gerald Niedobitek


Updated version
Access the most recent version of this article at:
doi:10.1158/1078-0432.CCR-12-0129

Supplementary Material
Access the most recent supplemental material at:
http://clincancerres.aacrjournals.org/content/suppl/2012/05/29/1078-0432.CCR-12-0129.DC1

Cited articles
This article cites 40 articles, 10 of which you can access for free at:
http://clincancerres.aacrjournals.org/content/18/14/3762.full#ref-list-1

Citing articles
This article has been cited by 2 HighWire-hosted articles. Access the articles at:
http://clincancerres.aacrjournals.org/content/18/14/3762.full#related-urls

E-mail alerts
Sign up to receive free email-alerts related to this article or journal.

Reprints and Subscriptions
To order reprints of this article or to subscribe to the journal, contact the AACR Publications Department at pubs@aacr.org.

Permissions
To request permission to re-use all or part of this article, contact the AACR Publications Department at permissions@aacr.org.