Rhesus Macaque Lymph Node PD-1\textsuperscript{hi}CD4\textsuperscript{+} T Cells Express High Levels of CXCR5 and IL-21 and Display a CCR7\textsuperscript{lo}ICOS\textsuperscript{+}Bcl6\textsuperscript{+} T-Follicular Helper (Tfh) Cell Phenotype

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Abstract

CD4 T follicular helper (Tfh) cells play a unique and essential role in the generation of B cell responses in the lymph node microenvironment. Here we sought to determine if differential expression of PD-1 could be used to delineate Tfh cells in rhesus macaque lymph nodes (LN). CD3\textsuperscript{+}CD4\textsuperscript{+} T cells were found to harbor a unique subset of cells that expressed the Program death-1 (PD-1) receptor at significantly high levels that were enriched in the LN compartment as compared to peripheral blood. The LN CD4\textsuperscript{+}PD1\textsuperscript{hi} T cells expressed a predominantly CD28\textsuperscript{-}CD95\textsuperscript{+} central memory phenotype and were CCR7\textsuperscript{lo}ICOS\textsuperscript{+}Bcl6\textsuperscript{+}. Additionally, CD4\textsuperscript{+}PD1\textsuperscript{hi} T cells preferentially expressed high levels of CXCR5 and IL-21 and significantly correlated with Bcl6\textsuperscript{+}Ki-67\textsuperscript{+}IgG\textsuperscript{+} B cells. As Bcl6 is primarily expressed by proliferating B cells within active germinal centers, our results suggest that LN CD4\textsuperscript{+}PD1\textsuperscript{hi} T cells likely localize to active GC regions, a characteristic that is attributable to Tfh cells. Overall, our findings suggest that high levels of PD-1 expression on CD4\textsuperscript{+} T cells in LN of rhesus macaques can serve as a valuable marker to identify Tfh cells and has implications for studying the role of Tfh cells in Human immunodeficiency virus (HIV), Simian immunodeficiency virus (SIV) and other infectious diseases that use the rhesus macaque model.

Introduction

CD4 T cells are a heterogeneous mix of lineages such as T-helper-1 & 2 (Th1 & Th2), T-helper-17 (Th17), T-regulatory cells (Tregs) and T-follicular helper cells (Tfh) that have specialized functions during an immune response [1]. Of these, Tfh cells are thought to play a central role in the generation and differentiation of T and B cell responses and play a key role in the germinal center reaction [2,3,4]. Studies have shown that Tfh cells are essential for the generation of long-lived memory and plasma B cells [5,6].

Tfh cells in human and mouse have been shown to express a unique phenotype characterized by high levels of program death receptor-1 (PD-1), the transcription factor Bcl6, the Inducible T cell Co-stimulator (ICOS), low levels of CCR7 and preferentially express the chemokine receptor CXCR5, that allow these cells to preferentially home to GC rich regions of the LN and provide critical help to B cells [7,8,9,10] and CD8 T cells [11,12]. Tfh cells are a major source of IL-21 in germinal centers [7] where it plays an essential role in increasing Bcl6 expression within B cells and the formation of germinal centers [10,13] and mediating B cell differentiation and maturation [4].

Rhesus macaques have been extensively used as an animal model to study various human infections, and for the development of vaccines against various pathogens. Recent studies [14,15] have shown that Tfh cells were expanded during simian immunodeficiency virus (SIV) infections in rhesus macaques and these expanded Tfh cells likely played a role in B cell dysregulation seen in these macaques. Most studies to date have relied on using multiple markers to identify Tfh cells. We sought to determine if differential expression of PD-1 was sufficient to phenotypically delineate Tfh cells in rhesus macaques. We show that Tfh cells in rhesus macaque LN can be delineated based on the high expression of PD-1 and these CD4\textsuperscript{+}PD-1\textsuperscript{hi} T cells inherently express high levels of CXCR5 and IL-21 and display a phenotype that is similar to Tfh cells in mouse and humans. Additionally, we found a significantly high positive correlation between CD4\textsuperscript{+}PD-1\textsuperscript{hi} T cells and Bcl6\textsuperscript{+}Ki-67\textsuperscript{+}IgG\textsuperscript{+} B cells indicating that both these populations of cells likely co-associate within the GC regions of LN.

Materials and Methods

Ethics Statement

The rhesus macaques (Macaca mulatta) used in this study were housed at Bioqual, Inc., in accordance with the recommendations...
of the Association for Assessment and Accreditation of Laboratory Animal Care International Standards and with the recommendations in the Guide for the Care and Use of Laboratory Animals of the United States – National Institutes of Health. The Institutional Animal Use and Care Committee of BIOQUAL approved these experiments. When immobilization was necessary, the animals were injected intramuscularly with 10 mg/kg of Ketamine HCl (Parke-Davis, Morris Plains N.J.). All efforts were made to minimize suffering. Details of animal welfare and steps taken to ameliorate suffering were in accordance with the recommendations of the Weatherall report, “The use of non-human primates in research”. Animals were housed in an air-conditioned facility with an ambient temperature of 21–25°C, a relative humidity of 40%–60% and a 12 h light/dark cycle. Animals were socially housed when possible or individually housed if no compatible pairing could be found. The animals were housed in suspended stainless steel wire-bottomed cages and provided with a commercial primate diet and fresh fruit twice daily, with water freely available at all times.

Animals and samples

Rhesus macaques of Indian origin (n = 7) were used for this study. The animals were seronegative for simian immunodeficiency virus (SIV), simian retrovirus (SRV) and simian T-cell leukemia virus (STLV) type-1.

Mesenteric lymph nodes (LN) and peripheral blood were obtained after sacrifice. Peripheral blood mononuclear cells (PBMC) were isolated by density gradient centrifugation and mesenteric lymph nodes were processed as previously described [16,17,18,19,20,21]. Samples collected from the 7 healthy animals were used in the various experiments.

Antibodies and flow cytometry

Isolated cells were labeled with different combinations of anti-CD3-Cy7APC, anti-CD4-PB, anti-CD28-Cy5-PE, anti-CD95-FITC, anti-CCR7-Alexafluor700, anti-CD-1-Biotin: anti-Biotin-APC, anti-ICOS-PE and analyzed by flow cytometry. All the antibodies except for anti-CCR7 (R&D Systems) and anti-CD1 (eBiosciences, San Diego, CA), and anti-Biotin-APC (Miltenyi Biotec, Cambridge, MA) were obtained from BD Biosciences (San Diego, CA), and titrated using rhesus macaque PBMC and LN cells. The optimal number of cells required for efficient staining was also titrated using LN cells. Bcl6 (BD Biosciences) expression was determined by Intracellular staining using the eBiosciences Fix/Perm kit.

The capacity of CD4+PD-1hi cells to produce IL-21 was determined in an in vitro stimulation assay using intracellular staining and flow cytometry as described previously [16,17,18,19,20,21]. Isolated cells were stimulated with 10 ng/ml of PMA (Sigma, St Louis, MO) and 500 ng/ml ionomycin (Sigma, St Louis, MO) in the presence of 1 μM Brefeldin A (Golgiplug; BD biosciences) for 4 hours in RPMI 1640 medium containing 10% fetal bovine serum. Cells were harvested and surface labeled with CD4 and PD-1. After fixing, the cells were permeablized (Cytofix/perm kit from BD Biosciences) and labeled with anti-CD3-Cy7APC, anti-IL-21-PE (BD Biosciences) and anti-IFNγ-FITC (BD Bioscience). Labeled cells were fixed with 0.5% Paraformaldehyde and analyzed using a Becton Dickinson LSR II flow cytometer.

To determine the expression of Bcl6 and Ki-67 in B cells, isolated LN cells were surface labeled with anti-CD3-Cy7APC, anti-CD20-PB, and anti-IgG-APC. After fixing, the cells were permeablized with the e-biosciences Fix/Perm kit and labeled

Figure 1. Lymph nodes are enriched for CD4+PD-1hi T cells. (a) Representative dot plot b) Mean fluorescence intensity (MFI) of PD-1 expression on lymph node CD4+ and CD4- T cells subsets. c) Relative proportions of CD4+PD-1hi T cells and CD4- PD-1hi T cells in LN and d) relative proportions of CD3 gated CD4+PD-1hi T cells in LN and peripheral blood mononuclear cells (PBMC). doi:10.1371/journal.pone.0059758.g001
with anti-Bcl6-PE and anti-Ki-67-FITC. All the antibodies except for anti-CD20 (e-biosciences) were obtained from BD Biosciences. Labeled cells were fixed with 0.5% Paraformaldehyde and analyzed using a Becton Dickinson LSR II flow cytometer.

For ex vivo analysis of CXCR5 and IL-21 mRNA expression, viable CD3+CD4+ PD-1-+, CD4+PD-1int, and CD4+PD-1hi T cell subsets were sorted from the LN using a BD FACS Aria sorter. Cells were labeled with anti-CD3, anti-CD4, anti-PD-1 and live/dead amine reactive dye VIVID [22]. VIVID+ dead cells were excluded.

Relative qPCR for IL-21 and CXCR5

The expression of IL-21 and CXCR5 mRNA was determined in purified subsets of CD4 T cells. RNA was isolated using RNeasy Kit (Qiagen Sciences, Gaithersburg, MD) and treated with Ambion Turbo DNase (Applied Biosystems, Austin, TX) to remove contaminating DNA. Purified RNA was reverse transcribed with Superscript III First Strand Synthesis kit (Invitrogen, Carlsbad, CA) to make cDNA that was used to determine the expression of IL-21 and CXCR5 using the ABI 7500 instrument (Applied Biosystems). Taqman qPCR was performed using high fidelity Platinum Taq polymerase (Invitrogen) as described previously [16,17,18,19,20,21] with Macaca mulatta specific (a) IL-21 primers: IL-21-F (TGTGAATGACTTGGACCCTGAA) and IL-21-R (AAACAGGAAATAGCTGACCACTCA), and probe IL-21-P (FAM- TCTGCCAGCTCCAGAAGATGTAGAGA-CAAACT-BHQ1) described previously [23], (b) CXCR5 primers: CXCR5-F (TTCACCTCCCGATTCCTCTA) and CXCR5-R (TGTGCACTACCCCCACTGTA), and probe CXCR5-P (FAM- GGATTCCTGCTGCCCCATGCT-BHQ1) and normalized to Macaca mulatta β-Actin house keeping gene using β-Actin specific primers: β-Actin -F (ATGCTTCTAGGCGGACTGTG), β-Actin-R (AAAGCCATGCGCAATCTC), and probe β-Actin-P (FAM-TGGGGTACACGGCTTCTGTGACAAAACC-BHQ1). Primers and probes for CXCR5 and β-Actin were designed using Primer-3 software [24]. Collected data were analyzed using the ddCT method with the ABI 7500 software.
was determined 

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showing the gating strategy and sort purity of CD4

Data analysis

Flow cytometric data was analyzed using FlowJo version 9.2 (Tree Star, Inc., Ashland, OR). Statistical analysis was performed using Mann-Whitney U test with GraphPad Prism Version 4.0 software (GraphPad Prism Software, Inc. San Diego, CA). A $p<0.05$ was considered significant. Error bars represent standard error. Linear regression analysis was performed to determine line of fit, and correlations were derived using Pearson’s correlation and a $p<0.05$ was considered significant.

Results

CD3$^+$PD-1$^{hi}$ T cells are predominantly CD4$^+$ T cells and are enriched in the lymph nodes of rhesus macaques

Tfh cells in mice and humans have been shown to express PD-1 at high levels [4]. As Tfh are predominantly CD4$^+$ T cells and mediate their function within the organized LN, we sought to determine if PD-1 was differentially expressed on LN CD4$^+$ T cells as compared to other T cells, and if the proportions of PD-1 expressing CD4$^+$ T cells differed from that of peripheral blood.

We first evaluated the expression of PD-1 on CD4$^+$ and CD4$^-$ T cell subsets in lymph nodes (Fig. 1a). Our results showed that PD-1 was differentially expressed on both CD4$^+$ and CD4$^-$ T cells with a majority of both these T cell subsets expressing either no or intermediate levels of PD-1 (Fig. 1b). Only a minor subset of CD4$^+$ and CD4$^-$ T cells were found to harbor a subset that expressed PD-1 at high levels. The proportion of CD4$^+$PD-1$^{hi}$ T cells was significantly higher than CD4$^-$PD-1$^{hi}$ T cell subsets (Fig. 1c) in the LN. When compared to peripheral blood, LN was found to harbor significantly higher proportions of CD4$^+$PD-1$^{hi}$ T cells (Fig. 1d).

Rhesus macaque CD4$^+$PD-1$^{hi}$ T cells display a predominantly ICOS$^+$Bcl6$^+$CCR7$^-$ and central memory phenotype

Tfh cells in mice and humans have been shown to coexpress ICOS and Bcl6 at high levels, and express CCR7 at low levels [1]. To determine if rhesus macaque Tfh cells exhibit a similar phenotype, we first evaluated the expression of CCR7, ICOS and Bcl6 on LN CD4$^+$PD-1$^{hi}$ T cells. As in humans and mice, essentially most CD4$^+$PD-1$^{hi}$ T cells co-expressed ICOS and Bcl6 on their surface whereas CCR7 levels were significantly lower than other PD-1 subsets (Fig. 2a–e). In contrast, CD4$^-$PD-1$^{hi}$ T cells were predominantly ICOS$^-$ (21%±17%) though a higher proportion of them were found to be Bcl6$^+$ (61%±14%).

Next we examined the expression of CD28 and CD95 on CD4$^+$PD-1$^{hi}$ T cells. CD28 and CD95 have been extensively used to delineate naive and memory subsets of CD4 T cells in rhesus macaques [26]. A majority of the CD4$^+$PD-1$^{hi}$ T cells coexpressed CD28 and CD95 indicative of a central memory phenotype (Fig. 2b–c). Previous studies have shown that Tfh cells are predominantly central memory T cells [4].

LN CD4$^+$PD-1$^{hi}$ T cells express high levels of CXCR5 and IL-21

CXCR5 is an essential chemokine receptor expressed on Tfh cells [1,4]. The binding of CXCR5 to CXCL13, its cognate ligand in the GC allows the Tfh cells to migrate into the GC where they directly interact with B cells. On the other hand, IL-21 is a critical effector cytokine produced by Tfh cells during Tfh:B cell interaction, that plays a major role in mediating B cell maturation and differentiation in within the GC [1,4].

We evaluated the expression of CXCR5 and IL-21 mRNA ex vivo in unstimulated sorted subsets of LN CD4$^+$PD-1$^{hi}$ T cells and compared them to LN CD4$^+$PD-1$^-$ T cells (Fig. 3a–b). Our results showed that LN CD4$^+$PD-1$^{hi}$ T cells expressed $\sim$100 fold more CXCR5 mRNA and $\sim$1000 fold more IL-21 mRNA as compared to LN CD4$^+$PD-1$^-$ T cells. Next we determined if LN CD4$^+$PD-1$^{hi}$ T cells preferentially produced IL-21 following mitogenic stimulation with PMA and ionomycin. The capacity of the cells to produce IL-21 was determined by intracellular staining and flow cytometry. LN CD4$^+$PD-1$^{hi}$ T cells produced significantly higher levels of IL-21 as compared to LN CD4$^+$PD-1$^-$ T cells with few LN CD4$^+$PD-1$^{hi}$ T cells producing IFN$\gamma$ (Fig. 3a–b). There was
a significant difference in the ratio of IL-21:IFNγ between the two subsets (Fig. 4c) with LN CD4⁺PD-1⁺ T cells harboring ~7x more IL-21 producing cells than LN CD4⁺PD-1⁻ T cells suggesting the LN CD4⁺PD-1⁺ T cells were primed to preferentially produce IL-21.

LN CD4⁺PD-1⁺ T cells significantly correlates with LN Bcl6⁺Ki-67⁺IgG⁺ B cells

Tfh cells directly interact with B cells in the GC and this interaction is essential for the induction of Bcl6 in GC B cells. Previous studies have shown that B cells that participate in GC reaction co-express high levels of Bcl6 [27,28,29].

To determine if LN CD4⁺PD-1⁺ T cells were associated with GC B cells, we evaluated the expression of Bcl6 and Ki-67 on LN IgG⁺ B cells and correlated them with LN CD4⁺PD-1⁺ T cells (Fig. 5a-d). Our results showed that there was a significantly high positive correlation between LN CD4⁺PD-1⁺ T cells and Bcl6⁺Ki-67⁺IgG⁺ B cells in LN suggesting that LN CD4⁺PD-1⁺ T cells are correlated closely with B cells in the GC. In contrast, we did not find a correlation between LN CD4⁺PD-1⁻ T cells and Bcl6⁺Ki-67⁺IgG⁺ B cells (Fig. 5d).

Discussion

Tfh cells play a critical role in immunity and delineating these cell subsets have relied on using a combination of markers such as Bcl6, ICOS, CCR7 and CXCR5 [1]. The lack of some of these critical reagents such as CXCR5 in the extensively used rhesus macaque model has been a limiting factor in better clarifying Tfh cells in this model. Our studies suggest that rhesus macaque Tfh could be clearly delineated based on the high expression of PD-1, a marker that is expressed at significantly high levels on Tfh cells.

We first characterized the expression of CCR7, Bcl6, and ICOS in LN CD4⁺PD-1⁻ T cells. CCR7 is a homing molecule expressed by T cells that traffic through the LN, whereas cells resident in the LN down regulate the expression of CCR7. LN CD4⁺PD-1⁻ T cells expressed significantly low levels of CCR7 as compared to other CD4 T cell subsets suggesting that these cells were enriched in the LN. In line with this, we found that LN harbored significantly higher proportions of CD4⁺PD-1⁻ T cells as compared to peripheral blood. Petrovas et al [15] showed that rhesus macaque Tfh cells expressed lower levels of CCR7 as compared to other CD4 T cells.

Tfh cells in humans and mice have been shown to express high levels of ICOS and Bcl6. ICOS is a costimulatory molecule expressed on Tfh cells that interact with the ICOS ligand on B cells during the GC reaction and this interaction is thought to play an essential role in influencing the outcome of Tfh mediated B cell maturation [30,31]. On the other hand, Bcl6 is a transcription factor that is an essential determinant of Tfh cells and critical for the differentiation of Tfh cells [1,4]. A majority of the LN CD4⁺PD-1⁻ T cells coexpress both ICOS and Bcl6 suggesting
that these cells display a phenotype similar to Tfh cells which is in line with previous studies showing high levels of Bcl6 and ICOS expression on rhesus macaque Tfh cells [15]. Interestingly, few CD4+ T cell subsets expressed ICOS though a majority of them expressed Bcl6 suggesting that these subsets probably localize to the GC rich regions of the LN. Quigley ME et al [32] showed that CXCR5+CD8 T cells frequently infiltrate B cell follicles whereas others have shown that NKT and γδ T cells were present in GC [33,34,35,36].

Tfh cells preferentially home to the GC rich regions of the LN where they interact with B cells to provide the critical help required for the differentiation and maturation of B cells [1,2,4]. Homing of Tfh to GC rich regions requires the binding of chemokine receptor CXCR5 on Tfh cells to its cognate ligand, CXCL-13 in the GC. As such, CXCR5 has been thought to be an essential marker to delineate Tfh cells [3]. Our results were found to support these observations as LN CD4+PD-1hi T cells expressed ~100 fold more CXCR5 as compared to cells that did not express PD-1 and in line with high levels of CXCR5 expression on rhesus macaque Tfh cells [15].

Like CXCR5, the ability of Tfh cells to produce IL-21 is considered to be an essential functional determinant of Tfh cells. IL-21 is produced primarily by Tfh cells within the GC microenvironment and binds to the IL-21 receptor expressed on B cells and T cells where it is thought to drive isotype class switching [7] and the expression of Bcl6 in B cells [10]. LN CD4+PD-1hi T cells were found to express ~1000 fold more IL-21 ex vivo and preferentially produce IL-21 following mitogenic stimulation as compared to LN CD4+PD-1 T cells suggesting that

Figure 5. CD4+PD-1hi T cells significantly correlates with CD20+Bcl6+Ki-67+IgG+B cells in LN. (a) Representative dot plots showing the gating strategy used for analyzing Bcl6 and Ki-67 expression on IgG+ B cells in the lymph node (LN) and b) relative proportions of IgG+ B cells in LN expressing Bcl6 and Ki-67. Correlation between LN CD4+PD-1hi T cells and c) CD20+Bcl6+Ki-67+IgG+B cells and d) CD20+Bcl6+Ki-67+IgG+B cells in LN. doi:10.1371/journal.pone.0059758.g005
Cells and Bcl6–Ki-67 not in other cells such as the mantle-zone B cells, plasma cells, lymphoid tissues, and inter- and intra-follicular CD4 T cells but not in other cells such as the mantle-zone B cells, plasma cells, dendritic cells, and macrophages. In these studies, we observed no correlation between LN CD4^+PD-1^hi T cells and Bcl6^hi^IgG^+^B cells. The expression of Bcl6 has been shown to be critical for the differentiation and proliferation of B cells in the GC and mice lacking BCL6 fail to form GC. Others have shown that Tfh cells directly interact with B cells within a GC reaction, whereas Hong et al. showed that PD-1^hi^CD4^+^T cells are present within the GC in the B cell zones of rhesus macaques. Good-Jacobson et al. showed that PD-1 regulates germinal center B cell survival and the formation and affinity of long-lived plasma cells.

Taken together, our results suggest that high levels of PD-1 expression on CD4^+^T cells can be an effective marker to discriminate rhesus macaque Tfh cells in LN and can serve as a valuable tool to study the role of Tfh cells in the rhesus macaque model.

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Author Contributions

Conceived and designed the experiments: JJM OO. Performed the experiments: OO JG. Analyzed the data: OO JG. Contributed reagents/materials/analysis tools: ML. Wrote the paper: J JM OO JG ML.

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