Immuno-PET in Inflammatory Bowel Disease: Imaging CD4-Positive T Cells in a Murine Model of Colitis

Amanda C. Freise, Kirstin A. Zettlitz, Felix B. Salazar, Richard Tavaré, Wen-Ting K. Tsai, Arion F. Chatzioannou, Nora Rozengurt, Jonathan Braun, and Anna M. Wu

1Crump Institute for Molecular Imaging, Department of Molecular and Medical Pharmacology, David Geffen School of Medicine at UCLA, Los Angeles, California; and 2Department of Pathology and Laboratory Medicine, David Geffen School of Medicine at UCLA, Los Angeles, California

Inflammatory bowel diseases (IBDs) in humans are characterized in part by aberrant CD4-positive (CD4+) T-cell responses. Currently, identification of foci of inflammation within the gut requires invasive procedures such as colonoscopy and biopsy. Molecular imaging with antibody fragment probes could be used to noninvasively monitor cell subsets causing intestinal inflammation. Here, GK1.5 cystodiabody (cDb), an antiumouse CD4 antibody fragment derived from the GK1.5 hybridoma, was used as a PET probe for CD4+ T cells in the dextran sulfate sodium (DSS) mouse model of IBD.

Methods: The DSS mouse model of IBD was validated by assessing changes in CD4+ T cells in the spleen and mesenteric lymph nodes (MLNs) using flow cytometry. Furthermore, CD4+ T-cell infiltration in the colons of colitic mice was evaluated using immunohistochemistry. 89Zr-labeled GK1.5 cDb was used to image distribution of CD4+ T cells in the abdominal region and lymphoid organs of mice with DSS-induced colitis. Region-of-interest analysis was performed on specific regions of the gut to quantify probe uptake. Colons, ceca, and MLNs were removed and imaged ex vivo by PET. Imaging results were confirmed by ex vivo biodistribution analysis.

Results: An increased number of CD4+ T cells in the colons of colitic mice was confirmed by anti-CD4 immunohistochemistry. Increased uptake of 89Zr-maleimide-deferoxamine (malIDFO)-GK1.5 cDb in the distal colon of colitic mice was visible in vivo in PET scans, and region-of-interest analysis of the distal colon confirmed increased activity in DSS mice. MLNs from colitic mice were enlarged and visible in PET images. Ex vivo scans and biodistribution confirmed higher uptake in DSS-treated colons (DSS, 1.8 ± 0.40; control, 0.45 ± 0.12 percentage injected dose [%ID] per organ, respectively), ceca (DSS, 1.1 ± 0.38; control, 0.35 ± 0.09 %ID per organ), and MLNs (DSS, 1.1 ± 0.58; control, 0.37 ± 0.25 %ID per organ).

Conclusion: 89Zr-malIDFO-GK1.5 cDb detected CD4+ T cells in the colons, ceca, and MLNs of colitic mice and may prove useful for further investigations of CD4+ T cells in preclinical models of IBD, with potential to guide development of antibody-based imaging in human IBD.

Key Words: immuno-PET; CD4; inflammatory bowel disease; 89Zr; diabody

J Nucl Med 2018; 59:980–985
DOI: 10.2967/jnumed.117.199075

The gut is a crucial compartment of the immune system. Lymphocytes reside in multiple locations throughout the gut, including the intestine, gut-associated lymphoid tissue such as Peyer patches and lymphoid follicles, and mesenteric lymph nodes (MLNs) (1). The immune system must balance between tolerating the many food- and microbiome-derived antigens sampled from the gut and mounting an immune response to pathogens. Dysregulation of this delicate balance can lead to inflammatory bowel disease (IBD), a family of disorders characterized in part by abnormal and excessive T-cell responses (2).

CD4-positive (CD4+) T cells are known mediators of inflammation in IBD. Crohn’s disease and ulcerative colitis each involves different subsets of CD4+ T cells, including T helper type 1 (Th1), Th2, and Th17 (2). Mouse models of IBD have helped advance understanding of how certain immune cell subsets, including CD4+ T cells, contribute to gut inflammation. Several models, such as the CD4+ CD45RBhigh T-cell transfer model and the 2,4,6-trinitrobenzenesulfonic acid/oxalozole model, are CD4+ T-cell–dependent and are used to study contributions of the adaptive immune system to IBD (3). The dextran sulfate sodium (DSS) model of colitis is often used to investigate innate immune cell contributions, as it is characterized by an initial influx of neutrophils, followed by infiltration of T cells. Although CD4+ T cells are not required for the induction of DSS colitis (4,5), they do increase in number in the colon as the disease progresses (6). Hall et al. (6) found that the peak number of CD4+ T cells occurred on day 12 after initiation of DSS administration. In addition, changes in the percentage of T cells in spleen and MLNs can occur as early as day 1 (6), indicating that during acute colitis the adaptive immune system is already changing, even though T cells are not required for induction. Given these findings, continued investigation into the role of CD4+ T cells in the DSS colitis model is warranted.

Molecular imaging has been used to assess inflammation in murine models of colitis. 18F-FDG and PET imaging were used in the DSS model to detect inflammation at several time points during progression of acute disease (7) and to correlate uptake in the colon with histologic damage and expression of perforin (8). 18F-FDG PET was used to identify inflammation in models with mild or severe chronic colitis, including the CD4+ CD45RBhigh and G0i2−/− CD3+ T-cell transfer models (9). In humans with IBD, 18F-FDG PET can identify inflammation with estimated sensitivity of 85% and specificity of 87% (10) and provide information complementary to standard methods such as colonoscopy and clinical symptom tracking (11); one caveat is that the highly variable physiologic uptake of 18F-FDG PET in the gut may reduce its utility as a screening tool (12).
Investigating specific cell subsets involved in IBD requires more specific probes. Molecular imaging with antibodies enables imaging of cell-surface biomarkers (13), and antibodies have been used to image immune cells in models of colitis. Kanwar et al. used a non-depleting intact anti-CD4 antibody for SPECT imaging of CD4+ T cells in DSS colitis and showed that uptake in the lower abdomen correlated with disease severity and the number of CD4+ T cells present (14). The integrin $\beta_7$ subunit, expressed on lymphocytes activated in Peyer patches and MLNs, was recently used as a target for SPECT imaging in DSS-treated colitic mice (15). Dearling et al. developed fragments of the antiintegrin $\beta_7$ antibody probe and used them to image lymphocytes in the intestines of colitic mice (16).

We previously described GK1.5 cys-diabody (cDb), an antimouse CD4 antibody fragment, and demonstrated its use in tracking immune system reconstitution posthematopoietic stem cell transplant (17). Subsequent studies investigated the effect of protein dose on image contrast and biologic effect on CD4+ T cells and showed that low-dose GK1.5 cDb produces high-contrast images of lymphoid organs with minimal effects on CD4+ T-cell function in vivo (18). Here, the utility of anti-CD4 GK1.5 cDb as an immuno-PET probe for regions of CD4+ T-cell infiltration in the murine DSS model of IBD has been evaluated and correlated with ex vivo biodistribution analysis, flow cytometry, and immunohistochemistry of CD4+ T cells in the intestines and lymphoid organs.

**MATERIALS AND METHODS**

**Animals**

Female C57BL/6 mice 6–12 wk old were obtained from Jackson Laboratories and housed by the Department of Laboratory Animal Medicine at the University of California, Los Angeles, under specific pathogen-free conditions. Animal studies were conducted under protocols approved by the institutional Chancellor’s Animal Research Committee.

**Induction and Assessment of Colitis**

DSS (40,000 kDa; Sigma) was dissolved in autoclaved tap water, sterile-filtered, and provided ad libitum. On the basis of published literature (6) and dose-finding pilot studies (data not shown), we determined that 4% DSS administered for 5 d induced infiltration of mononuclear cells into the colon by day 12. Regular water was re-stored on day 6. Mice were weighed every day and examined for signs of colitis, which was evaluated by a disease activity index modified from Sha et al. (19): weight loss (0, none; 1, 1%–4%; 2, 5%–10%; 3, 11%–20%; 4, >20%), fecal blood (0, none; 2, blood present in stool; 4, gross bleeding from anus), and stool consistency (0, normal; 2, loose stools; 4, diarrhea). On day 12, the mice were euthanized, and the colon, cecum, and MLNs were removed. MLNs were pooled per individual mouse, weighed, and divided by the number of MLNs collected to get the average weight of 1 MLN for each mouse.

**GK1.5 cDb Administration and PET/CT Imaging**

Antimouse CD4 GK1.5 cDb was produced, purified, and radiolabeled as described previously (17,18).

**FIGURE 1.** Characterization of DSS-induced anatomic changes in colitic mice. (A) Schematic showing days of DSS treatment, injection of $^{98}$Zr-malDFO-GK1.5 cDb, and PET imaging. (B) Change in body weight during and after DSS treatment ($n = 8$). (C) Colons cut just below cecum, laid flat, and measured ($n = 8$). (D) Ratio of colon weight (after removing stool) to length ($n = 8$). (E) MLNs removed, counted, pooled per mouse, and weighed. Each dot represents average weight of 1 MLN from individual mouse ($n = 8$ mice). Image shows 3 representative MLNs from 2 control mice (left) and 2 DSS-treated mice (right). $^*P < 0.05$. $^‡P < 0.0005$. $^†P < 0.0001$.

**FIGURE 2.** Characterization of CD4 cells in colons and MLNs of DSS-treated mice. (A) Representative images of antimouse CD4 immunohistochemical staining in colons from control (top) and DSS-treated (bottom) mice. Left panels: ×4 magnification; scale bar, 500 μm. Right panels: ×10 magnification; scale bar, 200 μm. (B) Number of cells staining positively for CD4 from entire colon section divided by total number of cells. Two sections of colon, 100 μm apart, were analyzed per mouse, with each point representing average percentage of CD4+ T cells in colon from 1 mouse (control, $n = 12$ mice; DSS, $n = 13$). (C) Flow cytometric analysis of CD4+ CD8+ T cells from MLNs (control, $n = 9$; DSS, $n = 10$) of control and DSS-treated mice. $^*P < 0.05$. 

- Anti-CD4 Immuno-PET in Colitic Mice • Freise et al. 981
Flow Cytometry

Lymphocytes in MLNs and spleens were analyzed by flow cytometry. Organ harvesting and preparation were performed as described previously (18).

Changes in CD4+ T Cells in DSS-Treated Mice

Anti-CD4 immunohistochemical staining showed increased infiltration of CD4+ T cells into the colons of DSS-treated mice (Fig. 2A). CD4+ cells were on average 0.66% ± 0.46% of the total cells in each section of colon in control mice, whereas DSS-treated mice had an average of 1.9% ± 1.5% CD4+ cells (P < 0.05; control n = 12, DSS n = 13) (Fig. 2B). Cecal from colitic mice did not show a significant increase in the percentage of CD4+ T cells (Supplemental Fig. 3A). Changes in CD4+ T cells in spleens and MLNs were evaluated by flow cytometry. The percentage of CD45+ CD4+ cells in the spleen increased slightly in DSS mice, though not significantly (control, 20.6% ± 4.4%; DSS, 23.7% ± 4.1%; n = 11 mice) (Supplemental Fig. 3B). The percentage of CD45+ CD4+ cells in MLNs of DSS mice decreased significantly (control, 33.0% ± 9.4%; DSS, 24.6% ± 6.1%; P < 0.05; control n = 9 mice; DSS n = 10 mice) (Fig. 2C).

RESULTS

DSS-Induced Colitis Results in Gross Anatomic Changes

Mice treated with 4% DSS for 5 d (Fig. 1A) showed weight loss beginning on day 6. On day 12, the weight of DSS-treated mice was significantly reduced to 87% ± 0.06% of initial weight, whereas control mice did not display any weight loss (P < 0.0001; n = 8) (Fig. 1B). DSS-treated mice also had significantly higher scores on the disease activity index than control mice as a result of increased weight loss, loose stool, and fecal bleeding (Supplemental Fig. 1A; supplemental materials are available at http://jnm.snmjournals.org).

Colons from DSS-treated mice were significantly shorter (P < 0.05) (Fig. 1C) and heavier (data not shown) than control colons. The colon weight-to-length ratio was significantly higher in colitic mice (P < 0.001) (Fig. 1D). Individual MLNs from DSS-treated mice weighed more on average (6.1 ± 3.0 mg; range, 1.4–11 mg) than MLNs from control mice (2.1 ± 1.1 mg; range, 0.4–4.4 mg; P < 0.0001; n = 8 mice) (Fig. 1E); furthermore, average MLN cellularity was higher in DSS mice (n = 4 mice; Supplemental Fig. 2).

Histopathologic analysis showed that colons from DSS-treated mice had epithelial cell loss, crypt destruction, and infiltration of inflammatory cells into the mucosa (Supplemental Figs. 1B and 1C). These signs were severe but localized in most mice. DSS-treated mice also had significantly more lymphoid aggregates throughout the colon (Supplemental Fig. 1D).

Statistical Analysis

Prism software (GraphPad) was used to perform 2-tailed Student t tests. A P value of less than 0.05 was considered to be statistically significant. For t tests on organs from the ex vivo biodistribution study, the Holm–Sidak correction for multiple comparisons was applied. Values are reported as mean ± SD unless indicated otherwise.
Immu-no-PET Imaging of CD4 in Colitie Mice

GK1.5 cDb was site-specifically conjugated to malDFO and radiolabeled with $^{89}$Zr. Radiolabeling efficiency was more than 99% ($n = 3$). Specific activity was 0.19 ± 0.023 MBq/µg (9.9 × 10$^5$ ± 1.2 × 10$^5$ MBq/mmol), and radiochemical purity was more than 99% ($n = 3$). $^{89}$Zr-malDFO-GK1.5 cDb (2 µg; 0.38 MBq) was injected into groups of mice (2 experiments of 4 mice per group). PET imaging showed tracer uptake in spleens and lymph nodes, consistent with previous results of studies of untreated mice (Fig. 3A) (18). PET scans of colitic mice demonstrated increased signal from MLNs. The colons of colitic mice also showed uptake of $^{89}$Zr-malDFO-GK1.5 cDb, whereas the control mice showed little to no uptake in the colon (Fig. 3A). Oral and intrarectal instillation of CT contrast agent delineated the intestine clearly on the CT scan (Supplemental Figs. 4A and 4B) and enabled 3-dimensional isocontour ROIs of the complete small and large intestines (gut ROI) and of the distal colon only (colon ROI) to be created on the basis of intensity of CT signal after manual removal of the skeleton (Supplemental Fig. 4C; Fig. 3B). DSS mice had significantly increased uptake in the colon ROI ($P < 0.005$) but not the gut ROI compared with control mice ($P = 0.548$) (Fig. 3C); spillover signal from the kidneys into the ROI may have masked actual differences.

Ex vivo scans confirmed the results of the in vivo scans, showing visibly increased uptake in colons, ceca, and MLNs from DSS mice (Fig. 4). Biodistribution studies showed that total activity was higher in DSS-treated colons, ceca, and MLNs (3.1-fold, 3.9-fold, and 3.0-fold, respectively) (Fig. 5A; Table 1). Comparing concentration of activity (percentage injected dose per organ [%ID/g]) in organs known to be affected by DSS-induced colitis (colon, cecum, and MLN) did not result in significant changes in uptake (Fig. 5B; Supplemental Table 1).

**DISCUSSION**

Abrupt activation and infiltration of CD4+ T cells is a hallmark of IBD in humans and in numerous mouse models of IBD. The contribution of CD4+ T cells to DSS colitis is still controversial. Although the DSS model does not require T cells for induction of colitis (4,5), they still infiltrate into the colon (6), and CD4+ T cells appear to play a role in the chronic stage of colitis (21,22). Adoptive transfer of DSS-primed CD4+ T cells into recipient mice treated with DSS resulted in exaggerated colitis (23). Conversely, athymic CD1 nu/nu mice treated with DSS developed more severe clinical signs of colitis than normal mice, suggesting that T cells may be protective in DSS colitis (5). Molecular imaging could be of use to further investigate how CD4+ T cells affect the induction and progression of DSS colitis.

The DSS model and resulting changes in CD4+ T cells were validated using quantitative immunohistochemistry and flow cytometry. Radiolabeled anti-CD4 GK1.15 cDb was used to see if the presence of increased CD4+ T cells could be visualized and quantified with immuno-PET. $^{89}$Zr-malDFO-GK1.5 cDb successfully detected CD4+ T cells in the colons and MLNs of colitic mice, and quantification of the distal colon region with ROI analysis showed that uptake significantly increased in colitic mice. This result is consistent with other studies, which report that the most severe damage in the DSS model occurs in that region (4,24). Quantifying uptake in the gut ROI, which included both the small and the large intestines, was challenging because the probe cleared through the kidneys, resulting in signal spillover into the anterior region of the ROI; differences in the gut ROIs were not significant between control and DSS-treated mice.

Significantly increased uptake of $^{89}$Zr-malDFO-GK1.5 cDb in the distal colon ROI, as well as in MLNs, was supported by the ex vivo organ scans. Interestingly, uptake was also increased in the cecum of colitic mice, although we did not observe increased CD4+ T cells by immunohistochemistry in colitic cecum; this may be due at least in part to enhanced permeability of the cecum due to DSS-induced inflammation. Quantification of uptake via ex vivo biodistribution analysis confirmed a significant increase in %ID per organ for colons, MLNs, and ceca of DSS-treated mice; however, the concentration of activity in these organs (presented as %ID/g) was not significantly higher in DSS mice. In colons of DSS mice, this may be explained by the finding that they were heavier than control colons. In DSS-treated MLNs, although the percentage of CD4+ T cells decreased, MLNs were significantly enlarged, which suggests that both CD4+ T cells and other types of immune cells were present in higher total numbers, resulting in increased size but not increased concentration.

In situ and ex vivo immuno-PET images, colitic MLNs appear to have a higher %ID/g than control MLNs, but ex vivo biodistribution analysis indicates that they have the same %ID/g. This finding is explained by the partial-volume effect, in which the concentration of activity in organs near the resolution of the PET scanner is underestimated (17). Colitic MLNs were dramatically enlarged and more easily detected by immuno-PET than the tiny

---

**FIGURE 4.** Ex vivo immuno-PET images of colons, ceca, and MLNs. After in vivo PET/CT scan mice were euthanized, colons, ceca, and MLNs were dissected and laid out for ex vivo PET/CT maximum-intensity projections (MIPs) ($n = 8$ mice per group). Representative images are shown, displayed as 12-mm maximum-intensity projections.
control MLNs. The partial-volume effect, especially in the context of murine lymph nodes, has been observed and discussed in previous work from our laboratory (17). The observation of signal from enlarged MLNs in colitic mice demonstrates the importance of MLNs and associated lymphatics in the development of DSS colitis and raises the possibility of the prognostic value of inflammatory staging sites such as MLNs, though more work is needed to fully understand the timing and cellular dynamics of MLN enlargement.

These results confirm and expand on the work of Kanwar et al., who used 111In-labeled intact anti-CD4 antibody for SPECT imaging of CD4+ T cells in DSS mice; in that work, a protein dose of 350 µg was used (14). We previously explored protein doses and found that amounts as low as 2 µg of GK1.5 cDb enabled imaging of CD4+ T cells in the lymph nodes and spleen with minimal biologic effects (18). Here, 2 µg of GK1.5 cDb were sufficient to image CD4+ T cells in the DSS model. Furthermore, Kanwar et al. drew an ROI that encompassed the entire lower abdomen and referred to this as “colon uptake” but did not discuss signal from CD4+ T cells in the MLN (14), which were likely included in the ROI. Therefore, while whole-abdomen ROIs may have prognostic value in IBD, they must be interpreted with consideration of the presence of MLNs and associated lymphatic tissue.

Although GK1.5 cDb enables visualization of total CD4+ cells in the gut, it does not provide information about CD4+ T-cell subsets such as Th1, Th2, Th17, or regulatory T cells, which is disadvantageous given the different pro- and antiinflammatory roles of each of these subsets. However, GK1.5 cDb can be used to identify sites of CD4+ T-cell accumulation, followed by detailed analysis with traditional methods such as biopsy to determine the subsets present as needed. Further studies using GK1.5 cDb in

<table>
<thead>
<tr>
<th>Site</th>
<th>Control Mean</th>
<th>Control SD</th>
<th>Control n</th>
<th>DSS Mean</th>
<th>DSS SD</th>
<th>DSS n</th>
</tr>
</thead>
<tbody>
<tr>
<td>MLN</td>
<td>0.37</td>
<td>0.25</td>
<td>8</td>
<td>1.09*</td>
<td>0.58</td>
<td>8</td>
</tr>
<tr>
<td>Colon</td>
<td>0.45</td>
<td>0.12</td>
<td>8</td>
<td>1.75†</td>
<td>0.40</td>
<td>8</td>
</tr>
<tr>
<td>Cecum</td>
<td>0.35</td>
<td>0.09</td>
<td>8</td>
<td>1.09‡</td>
<td>0.38</td>
<td>8</td>
</tr>
<tr>
<td>Small int.</td>
<td>5.12</td>
<td>1.33</td>
<td>8</td>
<td>4.87</td>
<td>1.13</td>
<td>8</td>
</tr>
<tr>
<td>Stomach</td>
<td>0.53</td>
<td>0.13</td>
<td>8</td>
<td>0.37*</td>
<td>0.07</td>
<td>8</td>
</tr>
<tr>
<td>Thymus</td>
<td>0.17</td>
<td>0.14</td>
<td>8</td>
<td>0.16</td>
<td>0.11</td>
<td>8</td>
</tr>
<tr>
<td>Spleen</td>
<td>6.89</td>
<td>0.98</td>
<td>8</td>
<td>7.56</td>
<td>1.38</td>
<td>8</td>
</tr>
<tr>
<td>ALN</td>
<td>0.41</td>
<td>0.14</td>
<td>8</td>
<td>0.45</td>
<td>0.17</td>
<td>8</td>
</tr>
<tr>
<td>ILN</td>
<td>0.31</td>
<td>0.09</td>
<td>8</td>
<td>0.46*</td>
<td>0.07</td>
<td>8</td>
</tr>
<tr>
<td>Blood</td>
<td>0.28</td>
<td>0.08</td>
<td>8</td>
<td>0.22</td>
<td>0.08</td>
<td>8</td>
</tr>
<tr>
<td>Liver</td>
<td>9.84</td>
<td>2.64</td>
<td>8</td>
<td>13.37</td>
<td>1.01</td>
<td>8</td>
</tr>
<tr>
<td>Kidney</td>
<td>23.84</td>
<td>6.26</td>
<td>8</td>
<td>27.30</td>
<td>3.68</td>
<td>8</td>
</tr>
<tr>
<td>Heart</td>
<td>0.15</td>
<td>0.04</td>
<td>8</td>
<td>0.21</td>
<td>0.07</td>
<td>8</td>
</tr>
<tr>
<td>Lung</td>
<td>0.38</td>
<td>0.26</td>
<td>8</td>
<td>0.36</td>
<td>0.03</td>
<td>8</td>
</tr>
<tr>
<td>Muscle</td>
<td>0.04</td>
<td>0.01</td>
<td>8</td>
<td>0.05</td>
<td>0.01</td>
<td>8</td>
</tr>
<tr>
<td>Femur</td>
<td>0.69</td>
<td>0.17</td>
<td>8</td>
<td>0.49</td>
<td>0.10</td>
<td>8</td>
</tr>
<tr>
<td>Tail</td>
<td>4.19</td>
<td>3.63</td>
<td>8</td>
<td>3.25</td>
<td>0.72</td>
<td>8</td>
</tr>
<tr>
<td>Carcass</td>
<td>19.53</td>
<td>3.24</td>
<td>8</td>
<td>16.64</td>
<td>0.96</td>
<td>8</td>
</tr>
</tbody>
</table>

*P < 0.05.  
†P < 0.005.  
‡P < 0.0001.  
Int. = intestine; ALN = axillary LN; ILN = inguinal LN. Data are total activity (%ID) present in blood and organs from DSS-treated mice compared with control mice 26 h after injection of 89Zr-maDFO-GK1.5 cDb.
mouse models of IBD could include imaging of CD4 in a T-cell–dependent model or imaging at several time points during the development of colitis to investigate the course of T-cell infiltration into the intestines and MLNs.

CD4+ T cells play a key role in human IBD, and identifying them throughout the bowel could be useful to monitor disease severity and predict treatment success. Imaging is less intrusive than colonoscopies, which are uncomfortable for patients and carry a risk of bowel perforation; furthermore, inflammation can vary depending on the position within the bowel. Anti-CD4 immuno-PET could provide an overview of inflammatory sites throughout the intestines. Other immune targets, including granulocytes (24), E-selectin (25), and CXCL8 (12), have been imaged in IBD patients. Aarntzen et al. suggest that anti-CXCL8 imaging could be a tool for making decisions about immune-modulating treatment (12); a similar application could apply for anti-CD4 imaging.

CONCLUSION

Anti-CD4 immuno-PET with GK1.5 cDb enabled detection of CD4+ T cells in inflamed mouse colons and in enlarged MLNs and should provide a useful tool for investigating CD4+ T cells in murine models of IBD and other diseases with a CD4 inflammatory component.

DISCLOSURE

Histological analysis was provided by the UCLA Imaging and Stem Cell Core at the CURE Digestive Diseases Research Center (NIH P30DK41301). Flow cytometry was performed in the UCLA Jonsson Comprehensive Cancer Center (JCCC) and the Center for AIDS Research Flow Cytometry Core Facility (NIH P30CA16042 and 5P30AI028697). Small-animal PET scans were performed in the Preclinical Imaging Technology Core at the Crump Institute for Molecular Imaging at UCLA, which is funded by the UCLA JCCC and National Institutes of Health (NIH P30CA16042). Amanda C. Freise was supported by the Dissertation Year Fellowship from UCLA. Richard Tavaré was funded by NIH award R21AI14255 and the UCLA Scholars in Oncologic Molecular Imaging training program (NIH R25ST09010). Anna M. Wu is a member of the JCCC and is a stockholder in and consultant to ImaginAb, Inc. No other potential conflict of interest relevant to this article was reported.

ACKNOWLEDGMENTS

We thank Kha Huynh, Alec Estrada, Waldemar Ladno, Theresa Falls, Jason Lee, Richard Tsachereau, and Clara Magyar for technical assistance.

REFERENCES

Immuno-PET in Inflammatory Bowel Disease: Imaging CD4-Positive T Cells in a Murine Model of Colitis

Amanda C. Freise, Kirstin A. Zettlitz, Felix B. Salazar, Richard Tavaré, Wen-Ting K. Tsai, Arion F. Chatziioannou, Nora Rozengurt, Jonathan Braun and Anna M. Wu

Published online: January 11, 2018.
Doi: 10.2967/jnumed.117.199075

This article and updated information are available at:
http://jnm.snmjournals.org/content/59/6/980

Information about reproducing figures, tables, or other portions of this article can be found online at:
http://jnm.snmjournals.org/site/misc/permission.xhtml

Information about subscriptions to JNM can be found at:
http://jnm.snmjournals.org/site/subscriptions/online.xhtml