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# Disruption of the $\beta_2$ -integrin CD11d ( $\alpha_D\beta_2$ ) gene fails to protect against experimental autoimmune encephalomyelitis

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# Abstract

The fourth member of the  $\beta_2$ -integrin family of adhesion molecules, CD11d ( $\alpha_D\beta_2$ ), is expressed on a wide variety of immune cells, however its function in autoimmune diseases, including EAE remains unknown. We induced EAE in wild type and CD11d<sup>-/-</sup> C57BL/6 mice using myelin oligodendrocyte glycoprotein (MOG<sub>35-55</sub>) peptide. The clinical course and histopathology of EAE was identical in both groups of mice throughout the disease course. There were no significant differences in the infiltration of leukocyte subsets into the central nervous system or in the production of cytokines from T cells isolated from the spleen or spinal cord from both groups of mice. Our data demonstrate that CD11d is not required for the development of EAE and, to date, is the only  $\beta_2$ -integrin molecule whose deletion does not result in attenuated disease.

# Keywords

adhesion molecules;  $\beta_2$ -integrins; EAE; neuroimmunology

# 1. Introduction

A critical requirement for the development of multiple sclerosis (MS) and its animal model experimental autoimmune encephalomyelitis (EAE) is the trafficking of immune effector cells into the central nervous system (CNS). These cells include macrophages and antigen-specific T cells which can participate in lesion formation and, in turn, demyelination (Sospedra and Martin, 2005). There are several families of cellular adhesion molecules known to function in the trafficking of effector cells to sites of inflammation, including the CNS (Archelos et al., 1999; Butcher et al., 1999; Campbell et al., 1998; Engelhardt, 2006; von Andrian and Mackay, 2000). In MS, the most notable of these are the  $\alpha_4$  integrins, specifically  $\alpha_4\beta_7$  and  $\alpha_4\beta_1$ , since treatment with antibody to  $\alpha_4$  reduces the development of new lesions, resulting in fewer clinical relapses (Miller et al., 2003). However inhibition of  $\alpha_4$  integrins, by a variety of approaches, does not completely prevent disease, due most likely to the participation of other cellular adhesion molecules in the disease process (Engelhardt et al., 1998; Kent et al., 1995; Myers et al., 2005; Yednock et al., 1992).

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In addition to the  $\alpha_4$  integrins, members of the  $\beta_2$ -integrin family, in particular CD11a and CD11b have been shown to contribute to EAE, however the results have often been contradictory. For example, treatment with anti-CD11a antibody in EAE produced results ranging from disease inhibition to exacerbation (Cannella et al., 1993; Gordon et al., 1995; Welsh et al., 1993; Willenborg et al., 1996). Similarly, several studies have shown variable levels of protection in antibody-mediated inhibition of CD11b. Those studies demonstrating attenuated disease attributed protection largely to prevention of myelin phagocytosis (Brocke et al., 1999; Bruck, 1997; Bruck and Friede, 1990; Gordon et al., 1995; Huitinga et al., 1993; Smith, 1999). Studies using CD11b-deficient mice, have shown that CD11b expression on T cells is critical for disease development, independent of cell trafficking into the CNS or phagocytosis (Bullard et al., 2005). More recent studies using CD11c-deficient mice have also shown a protective phenotype in EAE compared to control mice (Bullard et al., submitted). In contrast to these three  $\beta_2$ -integrins family members, nothing is known regarding the role of CD11d in EAE.

Previous studies have demonstrated multiple roles for CD11d in immune responses including facilitation of macrophage activation and infiltration into inflamed tissues (Bao et al., 2004; Mabon et al., 2000; Saville et al., 2004), development of IgG complex induced lung injury (Shanley et al., 1998) and T cell proliferation (Wu et al., 2004). In a spinal cord injury model (SCI), treatment with antibody to CD11d markedly reduced neutrophil and macrophage infiltration leading to improved recovery and reduced secondary damage (Gris et al., 2005; Gris et al., 2004; Mabon et al., 2000). These results, combined with those demonstrating reduced cellular infiltration in CD11b-deficient mice during EAE, suggested to us that loss of CD11d-mediated trafficking mechanisms would prove beneficial in EAE. We show here that deletion of CD11d, in contrast to other members of the  $\beta_2$ -integrin family, has no effect on the development and progression of EAE. These data indicate that  $\beta_2$ -integrin functions are not redundant in EAE and suggest differential functions in acute versus chronic inflammation.

# 2. Materials and Methods

#### Mice

CD11d<sup>-/-</sup> mice were generated through the replacement of exons 1 and 2 of the CD11d gene with a neomycin cassette as previously described (Wu et al., 2004). All CD11-deficient mice were backcrossed for at least six generations onto the C57BL/6 background. Inbred C57BL/6 mice were used as controls for all experiments. All studies were performed with approval from the University of Alabama at Birmingham Institutional Animal Care and Use Committee.

### EAE induction

Control and CD11d<sup>-/-</sup> mice were immunized with MOG35–55 peptide as previously described (Szalai et al., 2002). MOG peptide was synthesized by standard 9-fluorenyl-methoxycarbonyl chemistry and was >95% pure as determined by reversed-phase HPLC (Biosynthesis International). Briefly, on day 0 mice were injected subcutaneously with MOG35-55 emulsified in complete Freud's adjuvant. Pertussis toxin was injected intraperitoneally on day 0 and 2. Onset and progression of EAE symptoms were monitored daily using a standard clinical scale ranging from 0 to 6 as follows: 0, asymptomatic; 1, loss of tail tone; 2, flaccid tail; 3, incomplete paralysis of one or two hind limbs; 4, complete hind limb paralysis; 5, moribund; 6, dead. Mice scoring at least a 2 for more than 2 consecutive days were judged to have EAE. For each animal, a cumulative disease index (CDI) was calculated from the sum of the daily clinical scores observed between days 7 and 29.

# **Isolation of Spinal Cord Leukocytes**

Spinal cords were removed from control and CD11d<sup>-/-</sup> mice with active EAE (day 10 and 21) following saline perfusion and were ground through a cell strainer, washed in PBS, resuspended in 40% Percoll, and layered on 70% Percoll. After centrifugation at 2000 rpm (RT, 25 min), cells at the interface were removed, washed in PBS, and stained as described below.

# Flow Cytometry

Cells obtained from spinal cords, draining lymph nodes, or spleens were incubated with anti-CD16/32 (24G2, FcR block) to prevent nonspecific staining. Spinal cord leukocytes were stained with anti-CD45-FITC (30F11), anti-CD4-Sav-PerCP (GK1.5), anti-CD8-APC (53-6.7), anti-TNF- $\alpha$ -PE (MP6-XT22), and anti-IFN- $\gamma$ -FITC (XMG1.2; all from eBioscience). To determine if  $\beta_2$ -integrin expression changed during EAE on T cells or macrophages from CD11d<sup>-/-</sup> mice relative to control mice, cells were isolated from spleens or lymph nodes at day 21 post-immunization, stained with anti-CD16/32 as described above and then stained with anti-CD3-FITC (145-2C11), or F4/80-FITC (BM8) and anti-CD11a-PE (M17/4), CD11b Biotin-Sav-PercP (M1-70) or anti-CD11c-APC (HL3). Stained cells were analyzed using a FACSCalibur (BD Biosciences) and data was analyzed using CellQuest software (BD Biosciences).

### T cell proliferation and cytokine production

Ag-specific T cell proliferation assays were performed as described previously (Szalai et al., 2002). Single-cell suspensions from spleens obtained 21 days after EAE induction were cultured in 96-well plates at  $5 \times 10^5$  cells/well with increasing concentrations of MOG35–55 peptide or 3ug/ml ConA in triplicate. After 48 h, cultures were pulsed with [<sup>3</sup>H] thymidine for an additional 18 h, and incorporation of thymidine was measured. The in-vitro cytokine assays were performed essentially as described for the proliferation assay. Duplicate cultures were stimulated with MOG peptide (5 µg/ml). Culture supernatants were collected at 48 h for use in cytokine ELISAs. ELISA kits for TNF- $\alpha$ , IFN- $\gamma$ , IL-4, IL-2, IL-17 and TGF- $\beta$ 1 were purchased from R&D Systems. Each assay was performed according to the manufacturer's instructions. Cytokine production by cultures of CD11d deficient cells is reported as a percentage of the wild-type cell production. The data are pooled from two separate experiments.

### **Statistical Analysis**

Statistical significance between control and CD11d<sup>-/-</sup> mice for active EAE experiments was calculated using the Mann-Whitney U test. Statistical significance in the proliferation, compensation analysis and cytokine production assays was tested using the student's t test. All statistical analysis was done using Prism software v.4 (GraphPad).

# 3. Results

### CD11d-deficiency does not alter the disease course or cellular infiltration of active EAE

To assess the role of CD11d<sup>-/-</sup> in EAE, we performed active EAE using  $MOG_{35-55}$  peptide. We induced EAE in control and CD11d<sup>-/-</sup> mice and followed the course of disease for 29 days. The CD11d<sup>-/-</sup> mice presented with a phenotype remarkably similar to the wild-type mice as shown in Figure 1. The deletion of CD11d<sup>-/-</sup> does not alter the onset (15 days vs. 14 days; Table 1), incidence (100% for both groups) or the severity of disease as determined by cumulative disease index (CDI) for each group (46.5 vs. 45.6; Table 1). We next determined if the extent of cellular infiltrate was similar between the two groups of mice. For these studies, we induced active EAE and isolated spinal cords 10 and 21 days later. Leukocytes were isolated from spinal cords as described in Materials and Methods and analyzed for total leukocyte infiltration

(CD45<sup>+</sup> cells) and for T cell subsets (CD4<sup>+</sup> and CD8<sup>+</sup>). We observed essentially identical infiltration of CD45<sup>+</sup> cells in both groups of mice at day 10 (Figure 2A) and at later time points (data not shown). Neutrophil (Gr-1<sup>+</sup> cells) and macrophage (CD11b<sup>+</sup>) cells) infiltration was also not significantly different (p>0.05, student's t test) between wild type and CD11d<sup>-/-</sup> mice (data not shown). To determine if CD11d<sup>-/-</sup> T cells infiltrated the CNS to a similar extent as T cells from wild type mice, we isolated cells from spinal cords of control and CD11d<sup>-/-</sup> mice with active EAE and analyzed for the presence of CD4<sup>+</sup> and CD8<sup>+</sup> T cells. At day 10, the composition of the T cell infiltrate in CD11d<sup>-/-</sup> mice was not significantly different from that of wild type mice (Figure 2B, p>0.05, student's t test). On Day 21 post-immunization, the spinal cords of control and CD11d<sup>-/-</sup> mice contained similar numbers of CD4<sup>+</sup> T cells (3.3% and 5.5% respectively), while CD11d<sup>-/-</sup> mice had ~2.5-fold more CD8<sup>+</sup> T cells than control mice (4.4% and 1.7%, respectively; Figure 2B). The difference in the numbers of CD4<sup>+</sup> T cells between the two groups was not statistically significant (p>0.05, student's t test). To further characterize the cellular infiltrate in CD11d<sup>-/-</sup> versus wild type mice, we examined for the production of intracellular TNF- $\alpha$  and IFN- $\gamma$  in the CD4<sup>+</sup> T cell population at days 10 and 21. Wild type and CD11d<sup>-/-</sup> CD4<sup>+</sup> T cells produced comparable amounts of TNF- $\alpha$  at day 10 of active EAE (Figure 3). CD4<sup>+</sup> T cells from CD11d<sup>-/-</sup> mice produce modestly elevated levels of IFN- $\gamma$  early in disease compared to wild type mice, however, by day 21, both wild type and CD11d<sup>-/-</sup> T cells produce equivalent amounts of IFN- $\gamma$  and very little TNF- $\alpha$  (Figure 3). None of the differences in cytokines produced by CD4+ T cells between the two groups of mice was significantly different (p>0.05, student's t test).

# Splenic CD11d<sup>-/-</sup> T cells and macrophages express comparable levels of $\beta_2$ -integrins compared to wild type cells

To determine if the expression of the remaining  $\beta_2$ -integrin family members may have changed such that they might compensate for the loss of CD11d, we isolated T cells and macrophages from spleen and lymph nodes from CD11d<sup>-/-</sup> and wild type mice at day 21 post-immunization. We observed no statistically significant differences in the expression of CD11a-c on T cells from spleen or lymph nodes between CD11d<sup>-/-</sup> and wild type mice (Table 2). In contrast, CD11b and CD11c expression on CD11d<sup>-/-</sup> macrophages was significantly lower than that seen on control macrophages. There were no other significant changes in expression of the  $\beta_2$ -integrins on macrophages isolated from CD11d<sup>-/-</sup> and wild type mice (table 2).

# Splenic CD11d<sup>-/-</sup> T cells show no proliferative deficits or differences in cytokine production compared to wild type T cells

We next looked for differences in the proliferative capacity and cytokine profile of T cells isolated from the spleens of CD11d<sup>-/-</sup> and wild type mice. An *in-vitro* proliferation assay was performed as described previously (Bullard et al., 2005). Proliferation of MOG-sensitized T cells from wild type and CD11d<sup>-/-</sup> mice did not vary significantly (p>0.05, student's t test) when restimulated with various concentrations of MOG peptide (Figure 4). We also examined for cytokine production in MOG-sensitized splenic T cells from wild type and CD11d<sup>-/-</sup> mice. No significant differences were observed in the levels of IFN- $\gamma$ , TNF- $\alpha$ , IL-2, IL-17, IL-4, or TGF- $\beta$ 1 between the two groups of mice (Figure 5).

# 4. Discussion

Members of the  $\beta_2$ -integrin family of adhesion molecules are primarily known for their roles in leukocyte trafficking and, in the case of CD11b and CD11c, phagocytosis (reviewed in (Mayadas and Cullere, 2005; Underhill and Ozinsky, 2002; von Andrian and Mackay, 2000)). Although the results have often been contradictory, inhibition of  $\beta_2$ -integrin functions via treatment with antibody generally results in a better disease outcome regardless of the model system employed. This is also true in the CNS, where anti-CD11a and anti-CD11b treatment

reduced clinical signs of disease, cellular infiltration and phagocytosis of myelin (reviewed in (Barnum, 2001; Engelhardt, 2006)). Surprisingly the results reported in this study indicate that CD11d, unlike the other members of the  $\beta_2$ -integrin family, is not required for development or progression of EAE. In contrast, in a SCI model system employing spinal cord transection, inhibition of CD11d through the use of anti-CD11d monoclonal antibodies, partially prevented the infiltration of neutrophils and macrophages (Mabon et al., 2000). The long-term effectiveness of antibody treatment was not assessed with respect to cellular infiltration, neuronal survival or functional outcome, and the protective effect was attributed largely to disruption of neutrophil interaction with VCAM-1 on the vascular endothelium (Mabon et al., 2000). More recent SCI studies, in which the spinal cord was injured by compression (as opposed to transection), demonstrated not only reduced cellular infiltration but also reduced secondary injury and improved functional outcome for up to 12 weeks post treatment (Gris et al., 2005; Gris et al., 2004). Importantly, anti-CD11d treated mice had reduced myelin loss and more rapid remyelination compared to control mice (Gris et al., 2004). Based on the treatment regimen in these studies (3 injections in 48 hrs) and the long-term improved outcome, it is clear that transient inhibition of leukocyte infiltration with anti-CD11d, during the acute phase response, provides significant neuroprotection.

The effect of blocking or deleting CD11d with respect to T cell infiltration has not been previously examined. In the present study we observed that both CD4<sup>+</sup> and CD8<sup>+</sup> T cells readily infiltrated the spinal cords of CD11d<sup>-/-</sup> mice during EAE (Figure 2B). In fact, our data indicate that greater frequencies of both T cell subsets migrate into the spinal cords of CD11d<sup>-/-</sup> mice during EAE compared to control mice. This data combined with the observation that CD45<sup>+</sup> cells (as representative of the total leukocyte infiltrate) also traffic readily into the spinal cords of CD11d<sup>-/-</sup> mice, indicates that CD11d is not required for CNS infiltration in EAE. Although it is possible that other  $\beta_2$ -integrins or members of other integrin families compensate for the loss of CD11d under these circumstances, we did not observe this by flow cytometry (Table 2) and, deletion of any of the remaining  $\beta_2$ -integrin  $\alpha$ -chains results in significant attenuation of EAE ((Bullard et al., 2005), J. Hu, D. Bullard and S.R. Barnum, unpublished observations).

Aside from cellular trafficking, it is possible that deletion of CD11d could affect T cell proliferation and effector functions in a way that would alter the course of EAE relative to control mice. However, we observed no difference in antigen-specific T cell proliferation in CD11d<sup>-/-</sup> and control mice (Figure 3), indicating that there is no intrinsic proliferative defect in the absence of CD11d. Wu and colleagues demonstrated that CD11d<sup>-/-</sup> T cells proliferate comparably to wild type T cells when the T cell receptor is bypassed on treatment with PMA and ionomycin (Wu et al., 2004). However, in the same study, it was shown that CD11d<sup>-/-</sup> T cells proliferate poorly on treatment with PHA, ConA and several superantigens including SEA, SEB and SEE (Wu et al., 2004). The reason(s) for these differences in proliferative capacity remain unclear. Similar to the proliferation assays using T cells from CD11d<sup>-/-</sup> mice, there were no differences in production of IFN- $\gamma$  and TNF- $\alpha$ , in T cells isolated from the spinal cords, compared to control mice. In contrast, splenic T cells from CD11d<sup>-/-</sup> mice produced two-fold greater levels of IFN- $\gamma$ , but lower levels of TNF- $\alpha$  compared to control mice. The reasons for differential production of cytokines in these two tissue sites is unclear but suggests that evaluation of T cell cytokine production in the spleen, is not necessarily representative of cytokine production at sites of demyelination.

Adhesion molecules and their respective ligands represent an attractive therapeutic target for demyelinating disease. Inhibition of effector lymphocyte and phagocytic cell trafficking into the CNS by inhibiting adhesion molecule interactions provides significant protection from clinical disease as reported in animal models and clinical trials (Bullard et al., 2005; Cannella et al., 1993; Engelhardt, 2006; Gordon et al., 1995; Miller et al., 2003; Nataf et al., 2000; Welsh et al., 1993; Willenborg et al., 1996). Although this strategy has proven effective for many

adhesion molecules, it is not uniformly effective. For example, deletion of E and P selectins and PSGL-1 has no effect on the outcome of EAE (Brocke et al., 1999; Engelhardt, 1997; Engelhardt et al., 2005; Osmers et al., 2005). The failure to protect against demyelinating disease through targeted inhibition of these molecules is likely due to redundancy of function for some family members through the utilization of unexpected receptor/ligand pairs or signaling pathways (Fagerholm et al., 2004; Hynes, 2002; Richter et al., 1990). Which of these possibilities account for the results we report here for CD11d remains to be fully explored. Our results support growing evidence that  $\beta_2$ -integrins participate in unique rather than highly overlapping functions and indicate that successful therapeutic targeting even of closely related family members might be disease-specific.

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#### Figure 1.

The clinical course of MOG-induced EAE in wild-type mice and  $CD11d^{-/-}$ mice is identical. Active EAE was induced and symptoms were scored as described in the materials and methods. Results shown are the daily mean clinical scores for wild type (n=11) and CD11d^{-/-}mice (n=11) from 3 experiments. Adams et al.



### Figure 2.

Leukocyte subsets in spinal cord of control and CD11d<sup>-/-</sup> deficient mice with EAE are identical. Leukocytes isolated from spinal cords of control and CD11d<sup>-/-</sup> mice as described in Materials and Methods were immunostained for CD4, CD8 and CD45. A) The level of CD45<sup>+</sup> cells infiltrating the spinal cords of wild type (n=5) and CD11d<sup>-/-</sup> (n=6) mice 10 days postimmunization are identical. B) The percentage of CD4<sup>+</sup> T cells in the spinal cords of wild type and CD11d<sup>-/-</sup> mice at days 10 (n=5 and 6, respectively) and 21 (n=4 and 5, respectively) postimmunization were nearly identical (upper left quadrant in each panel). The percentage of CD8<sup>+</sup> T cells is modestly elevated in CD11d<sup>-/-</sup> mice at day 21 post-immunization compared

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to wild type mice (lower right quadrant). The results shown are from cells pooled within each group of mice and a representative experiment is shown.





### Figure 3.

Intracellular levels of TNF- $\alpha$  and IFN- $\gamma$  are identical in spinal cord-derived CD4<sup>+</sup> T cells from wild type and CD11d<sup>-/-</sup> mice with active EAE. CD4<sup>+</sup> T cells were isolated from the spinal cords of wild type and CD11d<sup>-/-</sup> mice at 10 (n=5 and 6, respectively) and 21 days (n=4 and 5, respectively) post-immunization and stained for intracellular TNF- $\alpha$  and IFN- $\gamma$  as described in Materials and Methods. IFN- $\gamma$  levels were modestly elevated at day 10 in CD11d<sup>-/-</sup> mice compared to wild type mice. Cytokine levels were similar for both groups at day 21. The results shown are from cells pooled within each group of mice.

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### Figure 4.

 $CD11d^{-/-}$  T cells proliferate comparably to wild-type T cells. Encephalitogenic T cells enriched by nylon wool adherence from the spleens of wild type (n=4) or  $CD11d^{-/-}$  (n=4) mice with active EAE were co-cultured with irradiated splenic APCs plus MOG peptide. Cells were pulsed with [<sup>3</sup>H]thymidine and harvested at 18 h for determination of radioisotope incorporation. The results shown are from two experiments and are expressed as the mean ± SEM of fold induction of T cell proliferation relative to background proliferation.



#### Figure 5.

The cytokine repertoire of CD11d<sup>-/-</sup> T cells is not significantly altered compared to wild-type T cells during EAE. Encephalitogenic T cells enriched by nylon wool adherence from the spleens of wild-type (n=6) or CD11d<sup>-/-</sup> (n=6) mice with active EAE were co-cultured with irradiated splenic APCs and restimulated with MOG peptide. Supernatants were collected 48h after stimulation and assayed by ELISA to quantitate production of each cytokine. The production of each cytokine for both CD11d<sup>-/-</sup> and wild type mice is presented as the mean  $\pm$  S.D. in pg/ml. The data are the mean of two experiments.

# Table 1

# EAE Symptoms in wild type and CD11d-deficient mice.

|                           | CDI <sup>A</sup> | Disease Onset <sup>B</sup> | Disease Incidence <sup>C</sup> |
|---------------------------|------------------|----------------------------|--------------------------------|
| Wild type n=11            | 46.5             | 15d                        | 100                            |
| CD11d <sup>-/-</sup> n=11 | 45.6             | 14d                        | 100                            |

 $^{A}$ Cumulative Disease Index is the mean of the sum of daily clinical scores observed between days 7 and 29.

 $^{B}$ Disease onset is defined as the first day of two consecutive days with a clinical score of two or more.

 $^{C}$  Disease incidence is defined as the percent of mice that displayed any clinical signs of disease.

### Table 2

Changes in expression of  $\beta_2$ -integrins on T cells and macrophages in CD11d-deficient mice during EAE

| T cells                             |                    |                |                |  |  |  |
|-------------------------------------|--------------------|----------------|----------------|--|--|--|
|                                     | CD11a <sup>A</sup> | CD11b          | CD11c          |  |  |  |
| Spleen (WT)                         | $25\pm0.78$        | $6.8 \pm 1.8$  | $1.7 \pm 1$    |  |  |  |
| Spleen (CD11d <sup>-/-</sup> )      | $17\pm5.5$         | $5.4\pm0.6$    | $2.7\pm0.6$    |  |  |  |
| Lymph node (WT)                     | $48\pm5.9$         | $4.9\pm0.57$   | $3.2\pm1.5$    |  |  |  |
| Lymph node (CD11d <sup>-/-</sup> )  | $45\pm9.7$         | $5.6\pm1.6$    | $3.3\pm1.4$    |  |  |  |
| Macrophages                         |                    |                |                |  |  |  |
|                                     | CD11a              | CD11b          | CD11c          |  |  |  |
| Spleen (WT)                         | $15\pm1.9$         | $B_{16\pm0.9}$ | $C_{10\pm1.1}$ |  |  |  |
| Spleen (CD11d <sup>-/-</sup> )      | 13 ± 1.9           | $13\pm0.9$     | $4.4 \pm 2.0$  |  |  |  |
| Lymph nodes (WT)                    | $2.6 \pm 0.7$      | $2.3 \pm 1.0$  | $1.5\pm0.9$    |  |  |  |
| Lymph nodes (CD11d <sup>-/-</sup> ) | 2.4 ± 1.3          | $2.0\pm0.9$    | 2.1 ± 1.7      |  |  |  |

<sup>A</sup>Data are presented as the mean  $\pm$  S.D. from three mice.

 $^{B}$ Significantly different compared to spleens from CD11d<sup>-/-</sup> mice, p=0.02.

 $C_{\text{Significantly different compared to spleens from CD11d}^{-/-}$  mice, p=0.01.