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# Borrelia miyamotoi activates human dendritic cells and elicits T cell responses

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## 16 Abstract

The spirochete Borrelia miyamotoi has recently been shown to cause relapsing fever. Like the 17 18 Lyme disease agent, Borrelia burgdorferi, B. miyamotoi is transmitted through the bite of 19 infected ticks, however, little is known about the response of the immune system upon infection. Dendritic cells (DCs) play a central role in the early immune response against B. 20 burgdorferi. We investigated the response of DCs to two different strains of B. miyamotoi using 21 in vitro and ex vivo models and compared this to the response elicited by B. burgdorferi. Our 22 23 findings show that B. miyamotoi is phagocytosed by monocyte derived-DCs, causing 24 upregulation of activation markers and production of proinflammatory cytokines, in a similar manner to *B. burgdorferi*. Recognition of *B. miyamotoi* was demonstrated to be partially 25 26 mediated by TLR2. DCs migrated out of human skin explants upon inoculation of the skin with B. miyamotoi. Finally, we showed that B. miyamotoi-stimulated DCs induced proliferation of naïve 27 CD4+ and CD8+ T cells, to a larger extent than B. burgdorferi. In conclusion, we show here that 28 29 DCs respond to and mount a immune response against *B. miyamotoi* that is similar to the 30 response to *B. burgdorferi*, and are able to induce T cell proliferation.

#### 32 Introduction

The tick-borne relapsing fever spirochete Borrelia miyamotoi was recently shown to be 33 34 pathogenic (1-3). Similar to the Lyme disease agent, Borrelia burgdorferi sensu lato, B. 35 miyamotoi infects Ixodes ticks in Europe, North America and Asia, however, the clinical presentation is more similar to other relapsing fever Borrelia, as symptoms are more systemic 36 and viral-like and include high fever, headache, myalgia, arthralgia and malaise (3-5). The 37 development of chronic meningoencephalitis has been described in three immunocompromised 38 patients infected with *B. miyamotoi* (1, 2, 6). As yet, it is not clear what the public health burden 39 40 of this disease is (7). Furthermore, as it has only recently become possible to cultivate B. miyamotoi in vitro (8), up to now very few in vitro studies have been performed and little is 41 known about the immunopathogenesis of *B. miyamotoi*. 42

43 In contrast, B. burgdorferi has been studied extensively in vitro and much of the immunopathogenesis has been elucidated. Upon entry into the skin via tick bite, B. burgdorferi 44 encounters dendritic cells (DCs); professional antigen presenting cells which orchestrate 45 adaptive immune responses by presenting phagocytosed antigen to CD4+ T cells which they 46 47 encounter upon migration into draining lymph nodes (9). DCs prime CD4+ T cells to become effector T helper (Th) cells, of which various subsets exist with each a different specialization, 48 49 depending on the type of response needed to clear the pathogen (10). DCs have been shown to phagocytose and respond to B. burgdorferi in vitro, producing cytokines and upregulating 50 activation markers (11). DCs recognize pathogens through pattern recognition receptors (PRR), 51 which can recognize pathogen-associated molecular patterns expressed by pathogens. The 52

central PRR responsible for recognition of *B. burgdorferi* is Toll like receptor 2 (TLR2) (12, 13) that recognizes lipoproteins expressed on the outer surface of the spirochete. We have shown previously that, in an *ex vivo* human skin model, DCs migrated out of the skin in response to inoculation with *B. burgdorferi* in a TLR-2 dependent manner (14). In *in vitro* studies, DCs primed with *B. burgdorferi* were able to induce Th cells, and elicited a protective immune response *in vivo* in mice (15). *B. burgdorferi* induced mixed Th1/Th2 responses in mice infected through syringe inoculation (16).

It is not known whether *B. miyamotoi* is recognized by DCs and whether the spirochete elicits 60 61 DC responses. Furthermore, the adaptive T cell responses that arise from DC-B. miyamotoi 62 interactions have not been previously studied. In this study, using two laboratory strains of B. miyamotoi and in vitro (monocyte derived (mo)DC) and ex vivo (human skin) models, we 63 demonstrated that human DCs phagocytose B. miyamotoi and become activated, migrate out of 64 the skin and produce cytokines in response, in a manner similar to B. burgdorferi. B. miyamotoi-65 stimulated DCs were able to induce T cell proliferation. Unraveling the ways in which the 66 67 immune system responds to and fights B. miyamotoi is crucial to our understanding of the 68 disease and may help to explain why the clinical presentation of infection with *B. miyamotoi* differs from that of Lyme disease. 69

70

#### 71 Materials and methods

### 72 Borrelia culture

73 Borrelia miyamotoi strains HT31 (Asian isolate) and LB-2001 (American isolate) were cultured in 74 MKP-F medium (AMC, the Netherlands) (8) and Borrelia burgdorferi sensu stricto strain B31 75 was cultured in MKP medium (AMC, Amsterdam, The Netherlands) at 33°C until cultures 76 reached the mid to late log growth phrase. Spirochetes were assessed for motility, quantified by dark-field microscopy and recovered by centrifugation at 4000 x g for 8 minutes. Subsequently, 77 78 spirochetes were resuspended in Roswell Park Memorial Institute (RPMI) 1640 medium (Gibco, 79 Paisley, UK) supplemented with 10% foetal calf serum (FCS, Lonza, Verviers, Belgium) and L-80 glutamine (Gibco) (for in vitro experiments) or phosphate buffered saline (PBS) (Fresenius Kabi, 81 Graz, Austria) (for ex vivo experiments). Spirochetes were passaged no more than 5 times prior 82 to experimentation.

### 83 Monocyte derived dendritic cell (moDC) generation and stimulation

MoDCs were generated from peripheral blood mononuclear cells (PBMCs) from fresh 84 heparinized blood donated by consenting, healthy volunteers as described previously (17, 18). A 85 86 total of 1 x  $10^5$  moDCs in 100µl RPMI 1640 medium supplemented with 10% FCS and L-87 glutamine were stimulated in a 96-wells flat-bottom plate (Corning<sup>®</sup>, Sigma-Aldrich) for 48 88 hours at 37°C with 5% CO<sub>2</sub> with various multiplicities of infection (MOIs) of Borrelia strains, 10 89 ng/ml ultrapure Lipopolysaccharide (LPS) of *Escherichia coli* (Invitrogen, Paisley, UK) or 50 ng/ml Pam3CSK4 (Invivogen, San Diego, CA, USA). For lysate stimulation experiments, lysates of LB-90 91 2001 strains expressing vsp1 or vlpC2 were prepared as previously reported (19). In selected 92 experiments 10  $\mu$ g/ml TLR2-blocking  $\alpha$ TLR2.5 (Ebioscience, San Diego, CA, USA) was used. Cells were washed and resuspended in FACS buffer (0.5% bovine serum albumin (BSA), 0.01% NaN<sub>3</sub> 93

and 0.35 mM EDTA in PBS) and stained with fluorophore-conjugated antibodies against HLA-DR,
CD83 or CD86 (BD Biosciences, Franklin Lakes, NJ) and flow cytometry was performed using a
FACS-CANTO II (BD Biosciences). The results were analysed using Flow-Jo (Treestar, Ashland,
OR) and Prism 7 (GraphPad Software, La Jolla, CA).

98 Phagocytosis

Phagocytosis experiments were performed in essence as previously described (17). Viable CFSEstained *B. burgdorferi or B. miyamotoi* were incubated with moDC at 37°C at an MOI of 5 or 50 for various time points. Phagocytosis was stopped by adding ice-cold FACS buffer (0.5% bovine serum albumin (BSA), 0.01% NaN<sub>3</sub> and 0,35 mM EDTA in PBS) and putting the samples at 4°C. The cells were then washed and analysed with flow cytometry. The phagocytosis index was defined as the mean fluorescence intensity × fraction CFSE-positive cells, minus the 4 °C binding control.

106 Lymphocyte proliferation

PBLs were isolated from fresh heparinised blood donated by consenting, healthy volunteers by density centrifugation on FicoII and PercoII (both GE healthcare, Chicago, IL), washed in PBS and labelled with 1  $\mu$ M CFSE in PBS. Labelled PBLs were then co-cultured together with washed allogenic *Borrelia*-stimulated moDCs at DC:PBL ratios of 1:1, 1:2 and 1:5 in 96-wells flat bottom plates (Corning<sup>®</sup>, Sigma-Aldrich) in Iscove's Modified Dulbecco's Medium (IMDM, Gibco) with 10% FCS. After four days, supernatant was harvested and PBLs were stained with fluorophoreconjugated αCD4 and αCD8 antibodies (BD Biosciences) and analyzed by flow cytometry. For each DC:PBL donor combination, the DC:PBL ratio which produced optimal proliferation results was selected and is shown in the results and analysis. *Borrelia*-stimulated moDCs were confirmed to have similar expression of maturation markers and cytokine production to earlier results prior to co-culture. All donors were seronegative, as measured by a commercially available *B. burgdorferi* C6 ELISA (Immunectis, XX, USA).

#### 119 *Ex vivo experiments*

120 Ex vivo skin experiments were conducted as previously described (14, 18). Briefly, human 121 surplus skin from breast or abdomen was obtained from healthy donors undergoing corrective 122 surgery at the Kennemer Gasthuis Haarlem, the Netherlands. A total of 10<sup>4</sup> or 10<sup>6</sup> spirochetes or 100 ng/ml ultrapure LPS of Escherichia coli were injected intradermally in 50 ul PBS. Eight mm 123 biopsies were taken immediately from the injection site and then washed in Iscove's Modified 124 Dulbecco's Medium (IMDM) (Life Technologies, Paisley, UK) with 2% FCS, before being 125 126 transferred into 1 ml IMDM with 10% FCS and L-glutamine - without antibiotics - and incubated 127 for 48 hours at 37°C with 5% CO2. Cells were harvested, washed twice, counted and 128 resuspended in FACS buffer for staining with fluorophore-conjugated antibodies against HLA-129 DR, CD11c, CD83 and CD1a (BD Biosciences). Flow cytometry was performed as for moDCs. The 130 number of DCs migrated per condition was calculated from the percentage of total migrated cells that were HLA-DR+/CD11c+. The supernatant and migrated cells from three separate 131 132 biopsies, taken from random locations of the skin, were pooled together to constitute one 133 replicate of each condition.

134 *Cytokine measurement* 

135 Concentrations of the cytokines IL-1 $\beta$ , IL-6, IL-8, TNF $\alpha$ , IL-12p70 and IL-10 were measured in 136 supernatant of 48h stimulated moDCs (1 x 10<sup>5</sup> in 100 $\mu$ l) or the supernatant of migrated DCs 137 using a human inflammatory cytometric bead array (CBA) kit (BD Biosciences), according to the 138 manufacturer's instructions. Results were analyzed using Flow-Jo and GraphPad Prism 7 139 software.

140 *Statistics* 

Statistical analysis was performed with GraphPad Prism 7. Data was considered significant at p <</li>
0.05, with one asterisk (\*) representing 0.01 
< 0.01 and three asterisks (\*\*\*) representing 0.0001 < p < 0.001. One-way ANOVA and Tukey's</li>
post-hoc tests were used to analyze normally distributed data and Kruskal-Wallis and Dunn's
post-hoc tests were used to analyse data that was not normally distributed.

146 *Ethics statement* 

*In vitro* experiments were performed using surplus human healthy volunteer blood collected for the purpose of a separate study. Written informed consent was obtained from human blood donors in accordance with protocols approved by the Medical Ethics Committee of the Academic Medical Center (Approval number NL34294.018.10) and the declaration of Helsinki. As for the human skin, the Medical Ethical Committee of the Academic Medical Center, Amsterdam, deemed that no approval was necessary for the use of (otherwise) discarded tissue.

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#### 155 Results

#### 156 B. miyamotoi is phagocytosed by DCs

To determine whether DCs are capable of phagocytosing *B. miyamotoi*, monocyte-derived DCs were incubated with CFSE-labelled strains HT31 and LB-2001 for various time points at 37°C. Uptake by DCs was analyzed by flow cytometry and compared to that of *B. burgdorferi* strain B31. The CFSE signal (fig 1a) and phagocytosis index (fig 1b) of DCs incubated with both *B. miyamotoi* strains increased over time, as similarly observed for B31, indicating that *B. miyamotoi* is phagocytosed by DCs, in a comparable manner to *B. burgdorferi*.

#### 163 **B. miyamotoi activates DCs to produce cytokines**

The activation of DCs and production of cytokines are crucial for their function. DCs were 164 165 stimulated with viable B. miyamotoi strains HT31 and LB-2001 and B. burgdorferi strain B31 for 48 hours at an MOI of 5 and 50 and maturation marker expression was measured by flow 166 cytometry (fig 2a,b). Notably, expression of the maturation markers CD83 and CD86 was 167 significantly upregulated by stimulation with *B. miyamotoi* strains at both MOIs, compared to 168 unstimulated DCs, similarly to stimulation with LPS. The *B. burgdorferi* strain B31 induced CD83 169 170 expression at both MOIs, whereas CD86 was only significantly upregulated at the higher MOI. 171 HLA-DR expression was only significantly induced by HT31 at an MOI of 50. Next we measured 172 cytokine production by B. miyamotoi-stimulated DCs (fig 2c). Both B. miyamotoi strains as well 173 as *B. burgdorferi* strain B31 induced IL-8, IL-6 and TNF $\alpha$  production at an MOI of 50. DCs stimulated with *B. miyamotoi* strains also produced IL-12, at significantly higher levels than B31-174 stimulated DCs, and HT31-stimulated DCs additionally produced significantly higher levels of IL-175

1β than unstimulated DCs. These results indicate that *B. miyamotoi* strains are capable of
 inducing DC activation and cytokine production in a similar manner to *B. burgdorferi*.

B. miyamotoi expresses variable major proteins (VMPs) - surface lipoproteins that form the 178 basis of the humoral immune response against the spirochete, and populations of spirochetes 179 180 have different VMP expression by spontaneous full gene switching in an attempt to evade 181 antibody responses. In a prelimary effort to explore whether VMPs affect the activation of DCs by B. miyamotoi strains, we measured the maturation markers of DCs stimulated with lysates of 182 LB-2001 containing different VMPs. We selected an LB-2001 that expressed the surface 183 184 lipoprotein variable small protein 1 (Vsp1), an LB-2001 strain that had switched to express 185 variable large protein C2 (VlpC2) after challenge with  $\alpha$ Vsp1 antibodies, and an LB-2001 strain 186 challenged with  $\alpha$ OVA (i.e. still expressing Vsp1) as a control. Our preliminary data show that all three lysates (at a concentration of 10 ng/ml) induced upregulation of CD83 and CD86 and no 187 differences were observed in the expression of these markers between the different lysates, 188 suggesting that activation of DCs by B. miyamotoi is independent of the VMP expressed (data 189 190 not shown).

### 191 DCs migrate out of the skin in response to inoculation of *B. miyamotoi*

To investigate whether *B. miyamotoi* activates DCs in a more physiological *ex vivo* model, we inoculated human skin with HT31 or LB-2001, took biopsies of the inoculated skin and cultured the biopsies in medium for 48 hours, and we analysed the number of migrated DCs. Migration of DCs from skin inoculated with HT31 or LB-2001 was approximately 10-fold higher than migration of DCs from PBS-inoculated skin, and similar to (i.e. not statistically different from) the numbers of DCs migrated from B31-inoculated skin (fig 3a). Preliminary data indicate that, as we have previously shown for *B. burgdorferi* [REF], that LCs consisted of the minory of migrated cells upon *B. miyamotoi* inoculation, i.e. 0,5-2% (data not shown). In addition, significantly higher levels of the proinflammatory cytokines IL-6 and IL-8, but not IL-1 $\beta$  or IL-10 were measured in the supernatants of *Borrelia*-inoculated biopsies compared to PBS-inoculated biopsies (fig 3b). IL-12 and TNF $\alpha$  levels were under the detection limit. These results strongly suggest that *B. miyamotoi* induces DC migration from the skin.

#### 204 DC responses to *B. miyamotoi* are partially mediated by TLR2

205 To determine whether TLR2, the main PRR responsible for recognition of B. burgdorferi, is 206 involved in the recognition of *B. miyamotoi*, DCs were preincubated with or without a blocking TLR2 antibody and stimulated with HT31 or LB-2001 for 48 hours. Notably, the antibody against 207 TLR2 strongly diminished the expression of CD83 induced by both strains (fig 4b). Moreover, the 208 TLR2 antibody suppressed CD83 upregulation induced by TLR2-agonist Pam3CSK4, but not that 209 210 induced by the TLR4-agonist LPS, indicating that the antibody is specific for TLR2 (fig 4a). CD86 211 upregulation in response to HT31 was also abrogated by the TLR2 antibody, and a trend was 212 observed towards lower CD86 expression in LB-2001 stimulated DCs, although this was not 213 significant. A trend toward lower production of the cytokines IL-12 and IL-6 (fig 4b) as well as IL-8 and TNFα (data not shown), by *B. miyamotoi*-stimulated DCs preincubated with αTLR2 was 214 215 also observed, although not statistically significant. These results demonstrate that TLR2 plays a role in DC recognition of *B. miyamotoi*. 216

# B. miyamotoi-stimulated DCs are better inducers of PBL proliferation than B. burgdorferi stimulated DCs

In order to explore the adaptive responses induced by B. miyamotoi-stimulated DCs, and 219 compare these to DCs stimulated by B. burgdorferi, DCs stimulated with HT31, LB-2001 or B31 220 221 for 48 hours were co-cultured with allogenic CFSE-labelled PBLs for 4 days, after which time the 222 CFSE signal of responder cells was measured by flow cytometry. Proliferating cells had a lower CFSE signal, as CFSE was halved between daughter cells during cell division (fig 5a). Significantly 223 224 higher percentages of total PBLs, CD4+ and CD8+ T cells were induced to proliferate by HT31-225 and LB-2001-stimulated DCs compared to unstimulated DCs (fig 5b,c,d). The percentage of 226 proliferating PBLs induced by B. miyamotoi-stimulated DCs was also significantly higher than 227 that of *B. burgdorferi*-stimulated DCs, which did not induce significant proliferation compared to 228 unstimulated DCs. These results demonstrate that DCs stimulated by *B. miyamotoi* are able to induce T cell proliferation. 229

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#### 231 Discussion

In this study, we investigated the response of DCs to *B. miyamotoi*, from recognition to induction of T cell responses. Uptake of pathogens and activation of DCs is the first step towards orchestration of adaptive immune responses. We demonstrated that DCs were able to phagocytose two different strains of *B. miyamotoi*, HT31 and LB-2001, in a manner similar to the *B. burgdorferi* strain B31. Most probably due to technical reason we were unable to show

phagocytosis in the ex vivo skin model (data not shown). After 48 hours of stimulation, DCs 237 238 became activated, upregulating the costimulatory molecule CD86, necessary for T cell activation, and CD83 which characterizes DC activation. HLA-DR, the major histocompatibility 239 240 complex II molecule which presents antigen to T cells was also upregulated in several experiments although this varied among donors. Proinflammatory cytokines IL-6, IL-8 and TNFa 241 242 were produced by DCs in response to B. miyamotoi strains at high MOIs, as with B31. B. 243 miyamotoi strains elicited a greater cytokine response than B. burgdorferi, as, unlike stimulation 244 with B31, DCs stimulated with HT31 or LB-2001 additionally produced the Th1-inducing cytokine IL-12 and HT31 induced production of significantly higher concentrations of IL-1β, although a 245 246 trend was also seen for LB-2001 and B31 and which may be significant at higher MOIs.

247 B. miyamotoi induced migration of DCs out of human skin explants in a more physiological ex vivo model of early B. miyamotoi infection. This result was comparable to previous work in 248 which we have shown that DCs migrate out of skin in response to the spirochetes B. burgdorferi 249 250 (14), Treponema pallidum and Leptospira interrogans (18). Spirochetes were inoculated 251 intradermally in numbers comparable to those found in biopsies of erythema migrans, in order 252 to mimic a tick bite (20). Due to variations in the skin used for experiments, i.e. thickness, 253 region (breast or abdomen), age of donor we observed variability in the migration of DCs in 254 response to stimulation. This variation is in line with our previous observations. As expected, 255 no differences in the expression of activation markers was observed in DCs migrated from B. 256 miyamotoi and PBS inoculated skin (data not shown) as DCs that have migrated – also in 257 response to PBS – have a mature and activated phenotype. However, skin explants inoculated

with *B. miyamotoi* produced greater amounts of IL-6 and IL-8, whereas IL-1 $\beta$  and IL-10 production was not significantly induced after inoculation by either *B. miyamotoi* or *B. burgdorferi* compared to PBS, as we have observed in previous experiments with *B. burgdorferi* (14). Although it is likely that DCs are the main producers of these cytokines, alongside DCs, other possible sources for the cytokine production observed in inoculated skin are keratinocytes and fibroblasts (21, 22).

264 Our data strongly suggest that TLR-2 is involved in the recognition of *B. miyamotoi* by DCs, as 265 blocking the receptor using an antagonistic TLR2 antibody impaired activation and cytokine 266 production in response to B. miyamotoi stimulation. TLR2 recognises diacylated and triacylated 267 lipopeptides by forming heterodimers with TLR1 and TLR6 (23, 24). Upon recognition, the 268 Myeloid differentiation primary response 88-dependent activation of nuclear factor kB signaling 269 cascade is induced, which triggers transcription of co-stimulatory molecules required for T cell activation, pro-inflammatory and anti-inflammatory cytokines, matrix metalloproteinases and 270 adhesion molecules, which enable the maturation and migration of DCs (9). We did not observe 271 a complete block of B. miyamotoi-induced DC activation and cytokine production upon blocking 272 273 TLR2. This may be due to incomplete blocking by the antagonistic antibody, as we also observed residual cytokine production by DCs stimulated by the TLR2 agonist Pam3CSK4 in the presence 274 of aTLR2. Another explanation could be that other PRR are involved in the recognition of B. 275 276 miyamotoi alongside TLR2. Since other PRRs such as TLR5, 7,8 and 9 have been reported to recognize B. burgdorferi (9), they may be similarly implicated in recognition of B. miyamotoi. 277

278 As with other tick-borne relapsing fever spirochetes, B. miyamotoi expresses variable major 279 proteins (VMPs) including Vlps and Vsps (25, 26), surface proteins that are the basis of the 280 humoral immune response against relapsing fever spirochetes. The B. miyamotoi strain LB-2001 has recently been demonstrated to predominantly express Vsp1, with a minority population 281 expressing VIpC2. After infection of a mammalian host, an IgM response is mounted against the 282 283 predominantly expressed VMP, in this case Vsp1, however the minority populations expressing 284 a different VMP are not affected and are able to evade the antibody response of the host, 285 enabling relapses in spirochetemia to occur (19). Our preliminary data suggest that DCs become activated in response to LB-2001 strains expressing either VMP, indicating that they are capable 286 287 of recognizing either or that other spirochetal proteins are driving the activation.

288 DCs stimulated with *B. miyamotoi* were able to elicit a T cell response, as we demonstrated 289 proliferation of the CD4+ and CD8+ PBL fractions, as well as total PBLs using an allogenic MLR. This response was greater than that generated by *B. burgdorferi*-stimulated DCs. The reason 290 291 remains to be investigated. Regardless, differences in DC-induced adaptive immune responses 292 might explain why B. miyamotoi appears to be readily cleared after human infection - and 293 relapses only seldomly occur - and B. burgdorferi is able to cause an chronic infection. Indeed, preliminary unpublished data indicate that *B. burgdorferi* was able to inhibit expression of genes 294 295 involved in antigen presentation in monocytes to a greater extent than *B. miyamotoi* (personal communication Leo Joosten, Radboudumc). B. miyamotoi-stimulated DCs released IL-12, a Th1 296 297 polarizing cytokine. Of interest, Th1 and Th2 responses have previously been implicated in B. 298 burgdorferi infection; IL-4 and IFN-γ were detected in the lymph nodes of mice infected with B.

*burgdorferi* (16), and Th1 responses have been observed in the joints and skin of Lyme borreliosis patients (27-29). Further studies should determine which subsets of Th cells are primed as part of the immune response to *B. miyamotoi*.

In summary, we have shown that *B. miyamotoi* strains HT31 and LB-2001 are phagocytosed by, and activate DCs, leading to upregulation of activation markers and production of proinflammatory cytokines in a manner similar to that of the *B. burgdorferi* strain B31. In addition, *B. miyamotoi* activated DCs to migrate out of human skin explants. Activation of DCs was found only to be partially dependent on TLR2. Interestingly, *B. miyamotoi*-stimulated DCs induced significantly more proliferation of CD4+ and/or CD8+ T cells compared to *B. burgdorferi*stimulated DCs.

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- 402
- 403 Figure legends

## 404 Fig. 1: DCs phagocytose B. miyamotoi

MoDCs were incubated with CFSE-stained *Borrelia* strains for various time points. a) The CFSE positive DC gate was determined based on the CFSE expression of a positive control of DCs incubated with mixed *Borrelia* compared to DCs alone (left histogram). The CFSE MFI of DCs increased over time for moDCs incubated with all three *Borrelia* strains at an MOI of 50 (right three histograms). b) The phagocytosis index ((% CFSE+ DCs x MFI at 37°C) minus (% CFSE+ DCs x MFI at 4°C)) increased over time for all three *Borrelia* strains, at an MOI of 5 and 50. Graphs

- 410 Wift at 4 C)) increased over time for all time *Borrelia* strains, at all wor of 5 and 50. Gr
- 411 show one representative experiment of two independent experiments.

## 412 Fig. 2: DCs become mature and produce cytokines in response to *B. miyamotoi*

MoDCs were stimulated with *Borrelia* strains for 48 hours. a) The MFI of maturation markers 413 414 CD83, CD86 and HLA-DR increased on DCs stimulated with each of the three Borrelia strains at an MOI of 5 and 50 compared to unstimulated DCs. Histograms are from one respresentative 415 416 experiment of eight. b) Fold difference in MFI of CD83, CD86 and HLA-DR compared to the 417 unstimulated control increased on DCs stimulated with each *Borrelia* strain. Graphs show the 418 mean (SEM) of eight independent experiments and donors. Asterisks denote significance of 419 difference compared to unstimulated control. c) DCs stimulated with B31, HT31 or LB-2001 at 420 an MOI of 50 produced higher amounts of IL-8, IL-6 and TNFa than unstimulated DCs. In addition, moDCs stimulated with the *B. miyamotoi* isolates HT31 and LB-2001 produced IL-12, 421 422 at significantly higher levels than B31-stimulated moDCs. HT31-stimulated moDCs additionally 423 secreted IL-1 $\beta$ . Graphs show the mean (SEM) of 8 independent experiments and donors. Asterisks denote significance of differences compared to unstimulated control. 424

## 425 Fig. 3: Skin DCs migrate out of the skin in response to inoculation with *B. miyamotoi*

Human skin was inoculated with *Borrelia* strains. a) Inoculation of HT31 or LB-2001 spirochetes induced migration of DCs out of the skin comparable to B31 and LPS injection and significantly greater than PBS injection. b) Skin injected with HT31 or LB-2001 produced significantly more IL-6 and IL-8 but not IL-1 $\beta$  or IL-10 than PBS-injected skin, as measured in the supernatant of biopsies. Graphs show one representative experiment of two independent experiments. Bars represent the SEM of triplicate measurements; each triplicate consists of three pooled biopsies. Asterisks denote significance of differences compared to PBS-injected skin.

## 433 Fig. 4: Blocking TLR2 partially supresses DC responses to B. miyamotoi

434 MoDCs were incubated with a blocking  $\alpha$ TLR2 antibody prior to stimulation with *B. miyamotoi*. 435 a) The increase in CD83 expression in response to HT31 or LB-2001 stimulation at an MOI of 5 or 436 50, and the increase in CD86 expression in response to HT31 stimulation, were impaired by 437 preincubation with  $\alpha$ TLR2. The response to LPS was unaffected, whereas CD83 upregulation by 438 Pam3CSK4 (Pam) was abrogated by blocking TLR2. b) A trend was observed toward lower 439 production of cytokines IL-12 and IL-6, as well as TNF $\alpha$  and IL-8 (not shown), in response to 440 stimulation after preincubation with  $\alpha$ TLR2. Graphs show one representative experiment of 441 three independent experiments. Error bars represent SEM of triplicate measurements. Asterisks 442 denote significant differences between stimulated moDCs with and without blocking TLR2, for 443 each condition.

# 444 Fig 5: *B. miyamotoi*-stimulated DCs induce greater proliferation of PBLs than *B. burgdorferi*-445 stimulated DCs

446 DCs stimulated with B31, HT31 and LB-2001 at an MOI of 50 for 48 hours were co-cultured with allogenic CFSE-stained PBLs for 4 days. a) The lower CFSE signal peak in proliferating PBLs 447 448 stimulated with  $\alpha$ CD28 and  $\alpha$ CD3 compared to PBLs alone was used to determine the gate for proliferating cells. b) Higher percentages of PBLs proliferated in response to HT31- and LB-2001-449 450 stimulated DCs than B31-stimulated DCs. c) CD4+ T cells had a lower CFSE signal after PBL coculture with HT31- and LB-2001-stimulated DCs. This was less pronounced after co-culture with 451 452 B31-stimulated DCs. d) Pooled results of four experiments using different DC:PBL donor 453 combinations revealed that a significantly higher percentage of PBLs were induced to proliferate by HT31- and LB-2001-stimulated DCs compared to unstimulated DCs. The percentage of 454 proliferating PBLs induced by B31-stimulated DCs was not significantly different to 455 unstimulated. This was also true for the CD4+ and CD8+ T cell populations of PBLs. Furthermore, 456 HT31- and LB-2001 stimulated DCs induced significantly higher percentages of proliferating total 457 PBLs and CD8+ T cells compared to B31-stimulated DCs and HT31-stimulated DCs induced a 458 459 significantly higher percentage of proliferating CD4+ T cells than B31-stimulated DCs. Graphs show the mean (SEM) of pooled results of the best DC:PBL ratio of each of four independent 460 461 DC:PBL donor combinations. Asterisks denote significance of difference compared to unstimulated control DCs. 462

























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