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1 **Title:** Induction of partial immunity in both males and females is sufficient to protect females  
2 against sexual transmission of *Chlamydia*

3  
4 **Running title:** Male and female immunity synergize for protection

5  
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15 **Conflict of Interest Statement**

16 No conflict

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39 **Abstract**

40 Sexually transmitted *Chlamydia trachomatis* causes infertility, and because almost 90% of  
41 infections are asymptomatic, a vaccine is required for its eradication. Mathematical modeling  
42 studies have indicated that a vaccine eliciting partial protection (non-sterilizing) may prevent  
43 *Chlamydia* infection transmission, if administered to both sexes prior to an infection.  
44 However, reducing chlamydial inoculum transmitted by males and increasing infection  
45 resistance in females through vaccination to elicit sterilizing immunity has yet to be  
46 investigated experimentally. Here we show that a partially protective vaccine (chlamydial  
47 major outer membrane protein – MOMP and ISCOMATRIX – IMX) provided sterilizing  
48 immunity against sexual transmission between immunized mice. Immunizing male or female  
49 mice prior to an infection reduced chlamydial burden and disease development, but did not  
50 prevent infection. However, infection and inflammatory disease responsible for infertility  
51 were absent in 100% of immunized female mice challenged intravaginally with ejaculate  
52 collected from infected immunized males. In contrast to the sterilizing immunity generated  
53 following recovery from a previous chlamydial infection, protective immunity conferred by  
54 MOMP/IMX occurred independent of resident-memory T cells. Our results demonstrate that  
55 vaccination of males or females can further protect the opposing sex, whereas vaccination of  
56 both sexes can synergize to elicit sterilizing immunity against *Chlamydia* sexual  
57 transmission.

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### 73 **Introduction**

74 In Australia, *C. trachomatis* accounts for 81% of all notifiable sexually transmitted infections  
75 and cases have increased from 14,045 in 1999 to 74,305 in 2010 (1). Women are 2 to 3 times  
76 more likely than men to contract an infection (2) and also have a higher incidence of  
77 asymptomatic infection, 70 – 90% compared to 40 – 60% in men (3). As the financial burden  
78 of *Chlamydia* infection is determined by the ease and cost of diagnosis, treatment and  
79 management of associated diseases (e.g. infertility), women account for 80% of the total  
80 economic cost of *Chlamydia* infections (4). Consequently, developing a vaccine that protects  
81 females from infection and disease has been the primary focus, despite studies showing the  
82 benefits of targeting both sexes (5).

83 We are reliant on animals to model infection transmission, and chlamydial research is most  
84 commonly conducted using the murine model, in which female mice require progesterone  
85 pretreatment to enhance their susceptibility to infection (6). Unfortunately, female mice  
86 become less receptive to mating during progesterone-induced diestrus (7), which prevents the  
87 study of sexual transmission between infected males and infection-susceptible female mice.  
88 Researchers instead utilize direct intravaginal inoculation of progesterone-primed female  
89 mice with *Chlamydia* to model sexual transmission. However, the infective dose can be  
90 highly variable between research groups, which can influence the kinetics of infection,  
91 immunity and disease development (8, 9). The inoculums of *Chlamydia* often used to infect  
92 mice are also considerably higher than the amount identified in guinea pigs and humans to be  
93 passed naturally through sexual transmission (10-12). Direct intravaginal inoculation of  
94 female mice with *Chlamydia* in sucrose buffer also cannot replicate the effects male ejaculate  
95 has on female genital tract immunity (13). Therefore, there is a need to improve the current  
96 model of *Chlamydia* sexual transmission in mice to test intervention strategies like vaccines.

97 The chlamydial major outer membrane protein (MOMP) is the most commonly used antigen  
98 in chlamydial vaccine research (14). However, the superior quality of MOMP-specific  
99 immunity that develops following a natural infection with *Chlamydia*, as opposed to simple  
100 immunization with MOMP, suggests that current adjuvant technology is failing to fully  
101 exploit the protective capabilities of MOMP. Similar in structure to ISCOMs (15),  
102 ISCOMATRIX® (IMX) is simply mixed with the antigen but does not need to encapsulate it  
103 to have an adjuvant effect, as the addition of phosphatidyl (choline or ethanol amine) into  
104 IMX during manufacturing provides exposed functional groups that allows for antigen  
105 conjugation through non-covalent interactions. IMX is safe and well-tolerated in humans and  
106 shown to elicit long-lived T and B cell responses in clinical trials (15). As protection against  
107 a chlamydial genital tract infection is dependent on effective antigen presentation by dendritic  
108 cells and the induction of both T and B cells (16), the combination of MOMP and IMX  
109 together with intranasal (IN) administration to elicit a mucosal response could formulate an  
110 effective vaccine against a genital tract infection.

111 By challenging infection-susceptible (progesterone-primed) female mice with ejaculates  
112 collected from infected males we could simulate *Chlamydia* sexual transmission in mice.  
113 Immunization with MOMP/IMX elicited partial protection against infection in both males

114 and females; however, infection and disease were completely absent from immunized female  
115 mice challenged with infectious prostatic fluid collected from immunized males, indicating a  
116 synergism between the partial immunity in both sexes.

## 117 **Results**

### 118 ***MOMP-specific T and B cell responses generated following immunization***

119 A MOMP-specific cell-mediated and humoral immune response contributes significantly  
120 towards protection against a *Chlamydia* genital tract infection (17). We therefore isolated  
121 splenocytes from immunized mice and quantified the expansion and cytokine production of  
122 their T helper cells (Th) (CD3+CD4+) and cytotoxic T cells (Tc) (CD3+CD8+)  
123 (Supplementary Figure 1) cells following *in vitro* re-stimulation with MOMP. Th, but not Tc  
124 cells isolated from MOMP/IMX immunized animals showed a significantly higher level of  
125 proliferation following re-stimulation with MOMP ( $p<0.001$ ) when compared to the  
126 unimmunized control (Fig. 1A). MOMP-specific Th cells isolated from MOMP/IMX  
127 immunized mice also showed significant staining for IFN $\gamma$  ( $p<0.001$ ), TNF $\alpha$  ( $p<0.001$ ), IL-  
128 17 ( $p<0.001$ ) when compared to unimmunized controls (Fig. 1B). Similarly, Tc cells  
129 displayed significant staining for IFN $\gamma$  ( $p<0.05$ ) and IL-17 ( $p<0.001$ ) when compared to  
130 unimmunized controls. Further analysis of the MOMP-specific Th cells isolated from the  
131 MOMP/IMX immunized animals indicated that approximately 60% were multifunctional  
132 (IFN $\gamma$ +TNF $\alpha$ +IL-17+, TNF $\alpha$ +IL-17+ IFN $\gamma$ +IL-17+,  $p<0.001$ ) compared to <10% in the  
133 unimmunized control groups (Fig. 1C). Tc cells in MOMP/IMX immunized mice were  
134 primarily of the single-positive phenotype (IFN $\gamma$ +, IL-17+ or TNF $\alpha$ +) and not significantly  
135 different from the unimmunized control. Immune responses generated following  
136 immunization were identical in male and female mice (data not shown).

137 Immunization with MOMP/IMX generated MOMP-specific sera and vaginal IgG and IgA  
138 (Fig 1D) and these were significantly greater than those detected following an intravaginal  
139 infection (LIC,  $p<0.01$ ). Sera from MOMP/IMX immunized mice significantly neutralized  
140 *Chlamydia* infectivity *in vitro* when compared to naïve sera ( $p<0.05$ ) (Fig. 1E). Stimulation  
141 of RAW264.7 with *Chlamydia* opsonized with sera from MOMP/IMX immunized also  
142 promoted expression of CXCL1 (7-fold), CXCL2 (1.5-fold) and IL-1 $\beta$  (2-fold) mRNA (Fig  
143 1F) after 24 hrs of stimulation.

### 144 ***Determining the infectious dose of Chlamydia transmitted by male mice***

145 The bacterial burden in the prostatic fluid, prostate and testes remained constant from week 1  
146 to week 10 post-infection (p.i) (Fig. 2A and B). The bacterial burden in the penis peaked at  
147 week 10 p.i, although this had little effect on the total amount of viable *Chlamydia* collected  
148 in the prostatic fluid over the course of the infection. Males produced  $4.7 \pm 1.7 \mu\text{L}$  (BALB/c)  
149 and  $4.4 \pm 2.4 \mu\text{L}$  (C57BL/6) of prostatic fluid (data not shown). The chlamydial load in the  
150 prostatic fluids contained  $49 \pm 20$  IFU (BALB/c) and  $64 \pm 35$  IFU (C57BL/6) of *Chlamydia*.  
151 Infected males were mated with receptive females (non-progesterone primed) at weeks 1 and  
152 2 p.i and the transmissible chlamydial load in their ejaculate was determined indirectly from  
153 the females (Fig. 2C and D). Samples collected from female mice that were mated with males  
154 at week 1 p.i contained the most *Chlamydia* IFU. Vaginal lavages collected from mated  
155 BALB/c females contained  $251 \pm 27$  IFU ( $\sigma$ BALB/c) and  $258 \pm 34$  IFU ( $\sigma$ C57BL/6), whilst

156 the vaginal plugs contained  $225 \pm 153$  IFU ( $\delta$ BALB/c) and  $165 \pm 50$  IFU ( $\delta$ C57BL/6) of  
157 *Chlamydia*. For both direct and indirect sample collection methods, the transmissible dose  
158 was determined to be 18 – 503 IFU/ejaculate (Supplementary Figure 2).

159 ***Protection against infection and disease in female mice following challenge with sexually***  
160 ***transmissible doses of Chlamydia***

161 Vaginal shedding was significantly reduced in MOMP/IMX immunized animals challenged  
162 with  $5 \times 10^1$  IFU on day 3 (15-fold) ( $p < 0.001$ ) and 6 p.i (5-fold) ( $p < 0.001$ ) when compared to  
163 the unimmunized control group (Fig. 3A). All unimmunized control animals challenged with  
164  $5 \times 10^1$  IFU of *C. muridarum* developed a cultivatable vaginal infection, whereas, 20% of  
165 immunized animals remained culture negative from the point of challenge up until days 12 –  
166 15 days post-challenge. This suppressive effect on infection in the vagina was abrogated  
167 when immunized animals were challenged with doses of  $1 \times 10^2$  IFU or greater. However, the  
168 duration of infection and chlamydial shedding was still significantly reduced in the vagina of  
169 MOMP/IMX immunized animals when compared to the unimmunized control groups on day  
170 6 p.i (3-fold) when mice were challenged with  $1 \times 10^2$  IFU ( $p < 0.001$ ) (Fig. 3B), as well as on  
171 day 3 (4-fold) ( $p < 0.001$ ) when mice were challenged with  $5 \times 10^2$  IFU (Fig. 3C). Immunized  
172 mice began clearing the infection in the lower genital tract from day 9 – 15 p.i, compared to  
173 day 21 – 35 p.i for the unimmunized group. Immunization elicited a significant level of  
174 protection against the severity of oviduct pathology following challenge with  $5 \times 10^1$  ( $p < 0.001$ )  
175 (Fig. 3A),  $1 \times 10^2$  ( $p < 0.001$ ) (Fig. 3B) and  $5 \times 10^2$  ( $p < 0.01$ ) (Fig. 3C), when compared to  
176 unimmunized controls. Mice recovered from a previous intravaginal chlamydial infection  
177 elicit sterilizing immunity against re-challenge. A prior infection (live infection control –  
178 LIC) elicited sterilizing immunity against all challenge doses, but no protection against the  
179 development of oviduct pathology.

180 ***Immune response in the genital tract of female mice following challenge with sexually***  
181 ***transmissible doses of Chlamydia***

182 By isolating cells from the entire genital tract (sans ovaries) (Fig. 4), we could identify  
183 differences in immune cell recruitment between immunized (MOMP/IMX), unimmunized  
184 (PBS) and immune mice (LIC) and link each cell infiltrate profile with its respective  
185 infection/disease protection profile. Leukocytes (CD45+) (Fig. 4A), neutrophils (CD45+Gr-  
186 1+) (Fig. 4B), macrophages (CD45+F4/80+) (Fig. 4C), B cells (CD45+CD19+) (Fig. 4D) Th  
187 cells (CD45+CD3+CD4+) (Fig. 4E) and Tc cells (CD45+CD3+CD8) (Fig. 4F) were analyzed  
188 (Supplementary Figure 3). Recruitment of B, Th and Tc cells into the genital tracts occurred  
189 on day 8 following challenge in both immunized and unimmunized animals. Although  
190 numbers of B, Th and Tc cells recruited to the genital tract were consistently greater in  
191 immunized versus unimmunized mice, levels were not significantly different between groups.  
192 Tc cells were however present in significant numbers in the genital tract of the LIC group  
193 prior to re-challenge (day 0) ( $p < 0.001$ ) and on days 2 and 4 p.i ( $p < 0.05$ ) when compared to  
194 the unimmunized controls. Th cell numbers were also significantly increased in the female  
195 genital tracts of mice in the LIC when compared to unimmunized mice on days 2 and 4  
196 ( $p < 0.05$ ) following challenge, indicating a rapid recruitment of Th cells in the LIC group.  
197 Neutrophil and macrophage numbers were not significantly different between immunized and

198 unimmunized mice. Uterine horn tissue sections were also stained for neutrophils (Gr-1+) 199 (Fig. 4G), macrophages (F4/80+) (Fig. 4H), B cells (B220+) (Fig. 4I) and T cells (CD3+) 200 (Fig. 4J), which matched those determined by flow cytometry. Interestingly, 8 – 15% of Tc 201 cells isolated from the genital tract of mice from the LIC on day 2 p.i were also CD103+, 202 indicating they were tissue resident memory T cells ( $T_{RM}$ ) (18) (Fig. 5). However,  $T_{RM}$  203 (CD3+CD8+CD103+) diminished to background levels within one week of the secondary 204 challenge.

### 205 ***Gene expression of immune cells sorted from the genital tract of female mice following*** 206 ***challenge with a sexually transmissible dose of Chlamydia***

207 As no significant differences in the anamnestic response were observed in the genital tract 208 between immunized and unimmunized female mice, we assessed gene expression in whole 209 oviduct tissues (Fig. 6A) and leukocytes recruited into the genital tract, sorted 210 (Supplementary Figure 3) into macrophages (Fig. 6B), neutrophils (Fig. 6C), Tc cells (Fig. 211 6D) and Th cells (Fig. 6E) to identify functional differences. Th and Tc cells isolated on day 212 8 p.i from MOMP/IMX immunized mice expressed significantly more IFN $\gamma$  (8- and 6-fold, 213 respectively), TNF $\alpha$  (2- and 6-fold, respectively) and IL-17A (10- and 12-fold, respectively) 214 than unimmunized mice. Similarly, Th and Tc cells isolated on day 2 p.i from LIC mice 215 expressed significantly more IFN $\gamma$  (6- and 2-fold, respectively), TNF $\alpha$  (2- and 3-fold, 216 respectively) and IL-17A (3- and 13-fold, respectively) than unimmunized mice on day 8 p.i. 217 Neutrophils and macrophages isolated on day 8 p.i from MOMP/IMX immunized mice 218 expressed significantly more CXCL1 (2-fold) and IL-1 $\beta$  (2- and 4-fold, respectively) than 219 unimmunized mice. Similar increases in gene expression were also detected in the whole 220 oviduct gene expression. This pattern of cytokine and chemokine expression in the oviduct of 221 MOMP/IMX immunized mice on day 2 p.i was similar to the gene expression RAW264.7 222 cell line stimulated with opsonized *C. muridarum* (Fig 1F).

### 223 ***In vivo depletion and passive immunization***

224 To confirm the roles of T and B cells in protection in female mice, we depleted Th (CD4+), 225 Tc (CD8 $\beta$ +) and B cells (CD20+) following immunization but prior to challenge. Th, Tc and 226 B cells (Supplementary Figure 4) were successfully depleted from the spleen and the genital 227 tract following administration of  $\alpha$ CD4,  $\alpha$ CD8 and  $\alpha$ CD20, respectively. MOMP-specific Ig 228 was not reduced in the serum or vaginal lavages of MOMP/IMX immunized mice following 229 B cell depletion (Supplementary Figure 4). Naïve mice were also passively immunized with 230 sera collected from MOMP/IMX immunized mice. Serum antibody levels were found to be 231 comparable between passively immunized and MOMP/IMX immunized mice 232 (Supplementary Figure 4). Depletion of CD4+ and CD8+ but not CD20+ cells from 233 MOMP/IMX immunized mice prior to challenge, inhibited protection against infection and 234 oviduct pathology (Fig. 7A). Depletion of CD4+ but not CD8+ or CD20+ cells from 235 unimmunized mice prior to challenge inhibited the natural resolution of the infection, but did 236 not exacerbate oviduct pathology (Fig. 7B). Depletion of CD8+ and CD4+ but not CD20+ 237 cells prior to re-challenge mitigated sterilizing immunity in LIC mice, but did not exacerbate 238 oviduct pathology (Fig. 7C). Passive immunization of naïve mice with serum from

239 MOMP/IMX reduced vaginal shedding between days 3 and 6 p.i when compared to infusion  
240 of sera collected from unimmunized mice, but did not affect oviduct pathology (Fig. 7D).

### 241 ***Protection against Chlamydia infection transmission between male and female mice***

242 Immunization conferred partial protection against infection in female mice, although it is  
243 unclear if vaccinating males prior to their infection could prevent or reduce the risk of  
244 infection transmission to immunized female mice. Unimmunized female mice challenged  
245 with the prostatic fluid collected from unimmunized males (week 1 p.i) developed a  
246 comparable vaginal infection and disease to females challenged with known doses of  $5 \times 10^1$ ,  
247  $1 \times 10^2$  and  $5 \times 10^2$  IFU of *C. muridarum* (Fig. 8A). Immunization of male mice prior to  
248 challenge significantly reduced the amount of *Chlamydia* detectable in the penis and testes,  
249 but not in prostatic fluid (Supplementary Figure 5). However, unimmunized female mice  
250 challenged with the infectious prostatic fluids from immunized males exhibited no reduction  
251 in infection or disease when compared to infection with unimmunized male ejaculates (Fig.  
252 8B). Half of immunized female mice showed sterilizing immunity against challenge with the  
253 infectious prostatic fluid from unimmunized males ( $p < 0.001$ ), but these mice showed no  
254 protection from disease when compared to challenge of unimmunized females (Fig. 8C).  
255 Interestingly, no immunized female mice challenged with the infectious ejaculate of  
256 immunized males acquired an infection or developed any oviduct pathology ( $p < 0.001$ ) (Fig.  
257 8D), indicating complete protection against *Chlamydia* sexual transmission between  
258 immunized males and females.

### 259 **Discussion**

260 The ability of an animal model to replicate sexual transmission of infection is crucial for the  
261 development of a human vaccine against *Chlamydia*. To date, only large and/or expensive  
262 animal models (non-human primates, pigs and guinea pigs) have been shown to transmit a  
263 chlamydial infection sexually (16). In this study, we determined the transmissible infectious  
264 dose in the ejaculate of male mice (18 to 503 IFU) in order to develop a mouse model of  
265 sexual transmission. This dose range determined for mice is comparable with Guinea pigs  
266 that transmit approximately 100 IFU of *C. caviae* (11, 12). Semen from *C. trachomatis*-  
267 infected human males has also been shown to contain between 675 –  $1.6 \times 10^4$  copies/mL of  
268 chlamydial DNA (10). With an average ejaculate volume of 4.66 mL (10), a transmissible  
269 infectious dose for humans would range between 3146 –  $7.5 \times 10^4$  copies/ejaculate, which  
270 closely matches the amount contained in mouse prostatic fluid (2080 –  $1.16 \times 10^5$   
271 copies/prostatic fluid). Therefore, we suggest that an inoculum of *C. muridarum* between 50  
272 and 500 IFU should be used for all future vaccine studies as this dose is not only capable of  
273 establishing an infection in all test animals, but it also reflects a physiologically relevant  
274 inoculum that could be passed naturally during sexual transmission.

275 Surprisingly, 20% of immunized female mice initially resisted the establishment of a genital  
276 tract infection for 12 – 15 days post-challenge with the sexually transmissible dose of 50 IFU.  
277 Despite the expansion of MOMP-specific and multifunctional Th and Tc cells following  
278 immunization, these could not be detected in the genital tract until day 8 p.i. Conversely,  
279 significant levels of MOMP-specific antibodies were generated following immunization,  
280 which were also shown to neutralize *Chlamydia* infectivity *in vitro* and reduce vaginal



281 shedding on days 3 and 6 p.i following infusion into naïve animals. In support of finding by  
282 others (19), this indicates that induction of humoral immunity through immunization is  
283 crucial for neutralizing the initial challenge inoculum, an important effect that was masked  
284 when using higher, non-physiological challenge doses of *C. muridarum* ( $>5 \times 10^3$  IFU) (data  
285 not shown).

286 Expression of chemokines (CXCL1, CXCL2 and CXCL5) and cytokines (IFN $\gamma$ , TNF $\alpha$  and  
287 IL-1 $\beta$ ) were up-regulated in the oviducts of immunized compared to unimmunized mice.  
288 Macrophages and neutrophils isolated from the genital tracts of immunized mice also showed  
289 a similar up-regulated pattern of cytokine and chemokine expression as the oviducts,  
290 indicating an increased level of activation, antimicrobial and chemotactic activity (16). This  
291 occurred during the early stages of the infection (day 2 p.i) where the presence of MOMP-  
292 specific antibodies in the genital tract of immunized animals was the only observed  
293 difference from unimmunized mice. Opsonization and complement are known to promote  
294 chemokines and cytokine production by APCs (20) and we too found that *Chlamydia*  
295 incubated with sera containing MOMP-specific antibodies enhanced expression of cytokines  
296 and chemokines by monocyte/macrophages *in vitro*. This indicated that in addition to their  
297 neutralizing effects, antibodies generated following immunization might also contribute  
298 towards protection by enhancing innate and adaptive responses through the induction of  
299 cytokines and chemokines.

300 The recruitment of T cells into the genital tract on day 8 p.i coincided with the resolution of  
301 the infection in immunized animals. However, unimmunized mice failed to resolve an  
302 infection from day 8 p.i despite mounting a T cell response comparable in both speed and  
303 magnitude to immunized animals. This indicated that functional differences existed between  
304 T cells recruited in immunized and unimmunized animals. Immunization induced expansion  
305 of MOMP-specific splenic Th and Tc cells, which produced IFN $\gamma$ , TNF $\alpha$  and IL-17A upon *in*  
306 *vitro* re-stimulation. Th and Tc cell isolated from the genital tracts of immunized mice on day  
307 8 p.i were also found to express more IFN $\gamma$ , TNF $\alpha$  and IL-17A than T cells isolated from  
308 unimmunized animals. These cytokines have been consistently shown to contribute to  
309 protection against infection and disease (21). Protection against vaginal shedding was  
310 alleviated following depletion of CD4 $^+$  and CD8 $^+$  cells, indicating an important role for Th  
311 and Tc cells in protection elicited by MOMP/IMX.

312 Animals recovered from a previous infection with *Chlamydia* (LIC group) demonstrated  
313 sterilizing immunity against re-challenge; therefore, it is important to compare the  
314 mechanisms of protection between MOMP/IMX immunized and LIC mice as this can  
315 highlight potential inadequacies in vaccine design. T cells were not detectable in the genital  
316 tract of MOMP/IMX immunized until day 8 p.i, whereas Tc cells were present in the genital  
317 tract of LIC mice prior to re-challenge and Th cells were rapidly recruited by day 2 p.i. Some  
318 Tc cells present in the genital tract of LIC mice prior to challenge were CD103 $^+$  and  
319 localized in the epidermis, which is consistent with the phenotype of T<sub>RM</sub> in female genital  
320 tract of mice during a HSV-2 infection (22). However, T<sub>RM</sub> were not detected in the genital  
321 tract of MOMP/IMX immunized mice even following challenge (day 2 – 8 p.i.). Tc cells  
322 isolated from the genital tracts of mice in the LIC group immediately prior to re-challenge  
323 (day 2 p.i) expressed increased levels of cytokines, which were associated with the rapid

324 recruitment of Th cells. This is also consistent with existing literature regarding the  
325 functionality of  $T_{RM}$ , that they provide a means of “sense and alarm” (23) by secreting  
326 cytokines and chemokines, which recruit effector Th cells necessary to eradicate the  
327 infection. Th and Tc cells recruited in the genital tract of MOMP/IMX immunized mice  
328 exhibited a similar pattern of increased cytokine expression as T cells from LIC, but were  
329 recruited later at day 8 p.i. Reduction of Tc cells numbers, including the  $T_{RM}$ , in the genital  
330 tract of LIC mice prior to re-challenge abrogated sterilizing immunity, which indicated that  
331 Tc cells are capable of preventing infection from sexually transmissible doses of *Chlamydia*.  
332 Th cells were also shown to contribute to sterilizing immunity in LIC mice, however; it is  
333 unclear if Th cells had a direct effect or if their depletion during the resolution of the primary  
334 infection inhibited the development of  $T_{RM}$  (24). Further phenotypic characterization,  
335 functionality and longevity studies will also be required to definitively characterize  $T_{RM}$ ,  
336 although this to our knowledge is the first indication that a small number of  $T_{RM}$  may be  
337 generated following a chlamydial infection. Comparison between immunized and LIC mice  
338 indicate that strategies to accelerate T cell recruitment into the genital tract, like the “prime-  
339 pull” approach (25), must be implemented if future vaccines are to achieve immunity  
340 equivalent to that which develops following a natural infection.

341 Mathematical modeling has been invaluable in determining the attributes a vaccine must  
342 possess in order to impact on infection transmission (5, 26). Assertions that non-sterilizing  
343 vaccines, if targeted towards both sexes, could interrupt infection transmission had not yet  
344 been investigated experimentally (5). Challenging female mice with prostatic fluids collected  
345 from infected males to model sexual transmission (Fig. 9A), we determined that the  
346 immunization status (immunized or unimmunized) of both sexes was critical for the induction  
347 of sterilizing immunity against infection and disease. Immunizing females was more effective  
348 against preventing infection transmission than immunizing males (Fig. 9B and C), as reported  
349 elsewhere (5). This could be attributed to the inherent difficulties in resolving an infection in  
350 the immune-privileged male genital tract and the vaccine eliciting a potent cell-mediated  
351 response as opposed to the more male protective humoral response (27). However, male  
352 vaccination status protected against the development of oviduct pathology in immunized  
353 females (Fig. 9D). As this reduction in severity and incidence of disease was not detected in  
354 unimmunized females, it is likely to be independent of bacterial burden. Instead, additional  
355 adaptive immune components like antibodies, chemokines or cytokines present in the  
356 ejaculate of immunized males may have provided cross talk with immunity in immunized  
357 females (13, 28). This indicates that anti-chlamydial immunity against infection will be  
358 dependent on a vaccine for females, whereas a vaccine targeting males may be required for  
359 protection against diseases associated with infertility.

360 In this study, analysis of the local immune response in the genital tract following a  
361 *Chlamydia* infection provides the first evidence that  $T_{RM}$  confer/are essential for sterilizing  
362 immunity against challenge and should be the target of future vaccines. Induction of  $T_{RM}$  in  
363 females may be required to elicit complete protection against an infection transmitted from an  
364 unimmunized male. A partially protective vaccine targeting females would have a greater  
365 impact on *Chlamydia* infection and disease than a male vaccine; however, sterilizing  
366 immunity, i.e. prevention of sexual transmission, would require vaccination of both sexes. It

367 has been consistently shown in the most at risk adolescent age group that sexual behavior and  
368 perception of risk to other sexually transmitted infections is unaltered by vaccination (29);  
369 therefore, the introduction of a non-sterilizing *Chlamydia* vaccine together with existing  
370 surveillance and antibiotic treatment programs could facilitate a decline in infection rates.

## 371 **Materials and Methods**

### 372 *Ethics statement*

373 Animal care and use protocols in this study adhered to the National Health and Medical  
374 Research Council guidelines to promote the wellbeing of animals used for scientific  
375 purposes. This study was approved by the Queensland University of Technology Animal  
376 Ethics Committee (QUT UAEC No. 1100000588 and 1400000010) and carried out in strict  
377 accordance with any recommendations. All animals were euthanized humanely by  
378 intraperitoneal injection with sodium pentobarbital (200 mg/kg).

### 379 *Animals and Chlamydia*

380 Mice (C57BL/6 and BALB/c) were sourced from the Animal Resource Centre (ARC,  
381 Canningvale, Australia) at 6 weeks of age. Animals were given food and water *ad libitum*. *C.*  
382 *muridarum* (Weiss strain), was a generous gift from Catherine O'Connell and cultured from  
383 McCoy cells and purified as previously described (30).

### 384 *Immunization*

385 Recombinant *C. muridarum* MOMP-MBP was purified and IN immunizations were  
386 performed as previously described (31). The vaccine contained recombinant MOMP (50 µg)  
387 and ISCOMATRIX® (10 µg) (Zoetis, Florham Park, USA) mixed in a 10 µL volume, 5 µL  
388 applied to each nare. Animals were immunized once on four separate occasions (days 0, 7, 14  
389 and 28).

### 390 *Antibodies and flow cytometry*

391 Cells were washed with 2%v/v fetal calf serum (FCS)/phosphate-buffered saline (PBS)  
392 before blocking with αCD16/CD32 – FcγR (Clone: 24G2) for 15 min at 4°C. Cells were  
393 labeled with Zombie Green Viability Dye (Cat No. 423112, Biolegend, San Diego, USA)  
394 prior to staining with αCD3 (Clone: 145-2C11, Cat No 100330 – Biolegend, San Diego,  
395 USA), αCD4 (Clone: RM4.5, Cat No. 562314 – BD Bioscience, San Jose, USA), αCD8α  
396 (Clone: 53.6-7, Cat No. 25-0081 – eBioscience, San Diego, USA), αCD19 (Clone: 1D3, Cat  
397 No. 551001 – BD Bioscience), αCD45 (Clone: 30-F11, Cat No. 562129 – BD Bioscience),  
398 αCD103 (Clone: 2E7, Cat No. 121408 – Biolegend), αGr-1 – Ly6C/Ly6G (Clone: RB6-8C5,  
399 Cat No. 108412 – Biolegend) and αF4/80 (Clone: BM8, Cat No. 123110 – Biolegend) in a  
400 volume of 50 µL for 15 min at 4°C. Cells were washed with 2% FCS/PBS before fixing using  
401 4%w/v paraformaldehyde for 10 min at 4°C. Fixed cells were permeabilized for intracellular  
402 cytokine staining using Perm/Wash (Cat No. 554723 – BD Bioscience) according to the  
403 manufacturer's instructions. Permeabilized cells were stained with αIFNγ (Clone: XMG1.2,  
404 Cat No. 554413 – BD Bioscience), αTNFα (Clone: MP6-XT22, Cat No. 554418 – BD  
405 Bioscience) and αIL-17 (Clone: TC11-18H10, Cat No. 559502 – BD Bioscience) diluted in  
406 Perm/Wash buffer for 40 min at 4°C. Stained cells were analyzed on the FACS Aria III (BD  
407 Bioscience) using Flowjo version X software (Tree Star Inc, Ashland, USA).

408 ***In vivo depletion and passive immunization***

409 Depletion of CD4<sup>+</sup> (Clone: GK1.5, IgG2b, 1 x 200 µg), CD8β<sup>+</sup> (Clone: 53-5.8 (32), IgG1, 1  
410 x 250 µg IP, 2 x 50 µg intravaginally) (BioXcell, West Lebanon, USA) and CD20<sup>+</sup> cells  
411 (Clone: 5D2, IgG2a, 1 x 250 µg) (Genentech Inc, San Francisco, USA) was performed  
412 following IP injection of each monoclonal antibody on day 13 prior to intravaginal challenge.  
413 Identical doses of αTNP (Clone: 2A3, IgG2a), αHRP (Clone: HRPN, IgG1) or αKLH (Clone:  
414 LTF-2, IgG2b) (BioXcell) were used as isotype controls. Passive immunization of naïve mice  
415 was performed one day prior to intravaginal challenge by IP infusion of 500 µL of serum  
416 collected from MOMP/IMX or PBS immunized mice.

417 ***Sample collection and tissue processing***

418 Vaginal lavages and serum were collected as previously described (21). Spleens processed as  
419 described previously (21) and cultured in complete Dulbecco's minimal essential medium  
420 (DMEM) (5%v/v fetal calf serum (FCS), 4 mM L-glutamine, 50 µg/mL gentamicin, 100  
421 µg/mL streptomycin sulfate) (Invitrogen, Carlsbad, USA) containing 50 µM β-  
422 mercaptoethanol. Genital tract tissues were digested in complete DMEM containing 500  
423 U/mL of collagenase I and 120 µg/mL DNase I (Invitrogen) for 1 hr shaking at 37°C.

424 ***MOMP-specific T cell proliferation and cytokine production***

425 Cells were stained with carboxyfluorescein succinimidyl ester (CFSE) (Sigma-Aldrich, St.  
426 Louis, USA) (5 µM) for 5 min at room temperature. Labeled splenocytes were stimulated  
427 with media containing recombinant MOMP (10 µg/well) for 96 hrs at 37°C with 5% CO<sub>2</sub>.  
428 Cells were incubated for an additional 12 hrs in complete DMEM containing brefeldin A  
429 (Sigma-Aldrich) (10 µg/mL) before intracellular cytokine staining.

430 ***MOMP-specific antibody quantification, in vitro neutralization of Chlamydia infectivity  
431 and opsonization***

432 Antigen-specific antibodies were measured by enzyme-linked immunosorbent assay (ELISA)  
433 and *in vitro* *C. muridarum* neutralization assay were performed as previously described (21).  
434 Monocytes/macrophage cell line RAW264.7 (ATCC TIB-71) were grown in RPMI 1640,  
435 HEPES, 10%v/ FCS, 4 mM L-glutamine, 50 µg/mL gentamicin, 100 µg/mL streptomycin. *C.*  
436 *muridarum* was incubated with a 1/20 dilution of serum for 1 hr at 37°C with 5% CO<sub>2</sub>, then  
437 added to RAW264.7 cells at an MOI of 1 and incubated for 2, 4, 6 and 24 hrs.

438 ***Intravaginal and intravaginal infections***

439 Intravaginal challenges were performed on mice by exposing the glans and inoculating the  
440 exterior of the urethra with 1x10<sup>6</sup> IFU of *C. muridarum* in 5 µL of sucrose-phosphate-  
441 glutamine (SPG) (219 mM sucrose, 3.8 mM KH<sub>2</sub>PO<sub>4</sub>, 8.6 mM Na<sub>2</sub>HPO<sub>4</sub>, 4.9 mM glutamic  
442 acid, pH 7.35).

443 Female mice received 2.5 mg of medroxyprogesterone (DepoProvera) subcutaneously, seven  
444 days prior to challenge with *C. muridarum*. Mice anaesthetized with ketamine (Parnell  
445 Laboratory, Alexandria, Australia) (100 mg/Kg) /xylazine (Bayer, Leverkusen, Germany) (10  
446 mg/Kg) were challenged intravaginally with *C. muridarum* in 20 µL of SPG two weeks  
447 following immunization. Mice recovering from a primary infection of 5x10<sup>1</sup>, 1x10<sup>2</sup> or 5x10<sup>2</sup>

448 IFU (6 weeks) were also re-challenged with the same dose as the primary infection to act as a  
449 live infection control (LIC) group.

#### 450 ***Prostatic fluid collection, mating and sexual transmission***

451 Male mice were maintained under isoflurane-induced anesthesia and placed on their backs for  
452 collecting prostatic fluid. Urine was drained from the bladder using a syringe prior to  
453 collection. Ejaculation was induced following electro-stimulation of the prostate using a  
454 custom made probe and the square wave generator (Universal Kymograph – Harvard  
455 Apparatus, Massachusetts, USA) (Supplementary Figure 6). The prostatic fluid was collected  
456 into a capillary tube and expelled into a tube before making the volume up to 20  $\mu$ L with cold  
457 SPG.

458 Infected males were mated with non-progesterone primed females during the peak of their  
459 infection (day 7 – 14 post-infection – p.i) at the beginning of the 12 hr dark cycle. Paired  
460 mice were observed for 1 hr and vaginal lavages were collected following the first instances  
461 in which a female mated with a male for  $\geq 10$  sec. Paired mice were then left for the  
462 remaining dark cycle and inspected for vaginal plugs at the beginning of the following 12 hr  
463 light cycle. Vaginal plugs were excised and homogenized in 800  $\mu$ L of SPG using the OMNI  
464 TH tissue homogenizer. The uterine horns were flushed with 200  $\mu$ L of the same SPG  
465 containing the vaginal plug.

#### 466 ***Quantification of C. muridarum from secretions, tissues and swabs***

467 Infected tissues ( $\leq 250$  mg) were excised and homogenized in 800  $\mu$ L of SPG. Vaginal swabs  
468 were collected using a sterile nasopharyngeal swab (Copan, Murrieta, USA) and stored in  
469 500  $\mu$ L of SPG containing two glass beads. The SPG supernatant was used to quantify  
470 chlamydial content by culture as previously described (21). Ejaculate was also screened by  
471 culture and *ompA*-specific qRT-PCR as described previously (31).

#### 472 ***RNA extraction and RT-PCR***

473 Genital tract tissues were equilibrated in RNAlater (QIAGEN, Venlo, Netherlands) then total  
474 RNA was extract using the RNeasy Tissue Mini Kit (Cat. No. 74704) and treated with  
475 RNase-free DNase (QIAGEN). RNA was extracted immediately from sorted cells using  
476 Trizol (Invitrogen) and 10  $\mu$ g of glycogen carrier protein (Cat No. AM9510, Applied  
477 Biosystems, Foster City, USA) according to manufacturer's instructions. cDNA was  
478 synthesized using High Capacity Reverse Transcriptase Kit (Applied Biosystems) (Cat. No.  
479 4368814) as per the manufacturer's instructions. Each reaction contained 10 ng of cDNA, 1  
480  $\mu$ M of forward/reverse primers (Supplementary Figure 7) (Sigma-Aldrich), 200  $\mu$ M dNTP's,  
481 1.5 mM MgCl<sub>2</sub>, 1X buffer, 0.15X SYBR green, and 5 U of Platinum Taq polymerase  
482 (Invitrogen) made up to a final 20 $\mu$ L volume using sterile endonuclease-free water. RT-PCR  
483 was performed using the Corbet Rotorgene Q (QIAGEN). Gene expression was analyzed  
484 using Hierarchical Clustering Explorer 3.5 Analysis Tool (University of Maryland, USA).

#### 485 ***Immunohistochemistry***

486 Immunohistochemistry was performed by the HistoTechnology Facility (Queensland Institute  
487 of Medical Research, Brisbane, Australia) as described previously (21).

488 **Gross oviduct pathology**

489 The oviduct diameter was measured on day 35 p.i to assess the severity of pathology (33, 34).  
490 The incidence of oviduct pathology (presence or absence) was the number of mice presenting  
491 with hydrosalpinx (unilateral and bilateral).

492 **Power calculations and statistical analysis**

493 Sample sizes were determined *a priori* using a one-tailed, Proportions: Inequality, two  
494 independent groups Fischer's exact test in G\*Power 3.1.7 software (Institute for  
495 Experimental Psychology, Dusseldorf, Germany). Statistical analysis of graphical data was  
496 performed using GraphPad Prism® version 5.00 (GraphPad, La Jolla, USA).

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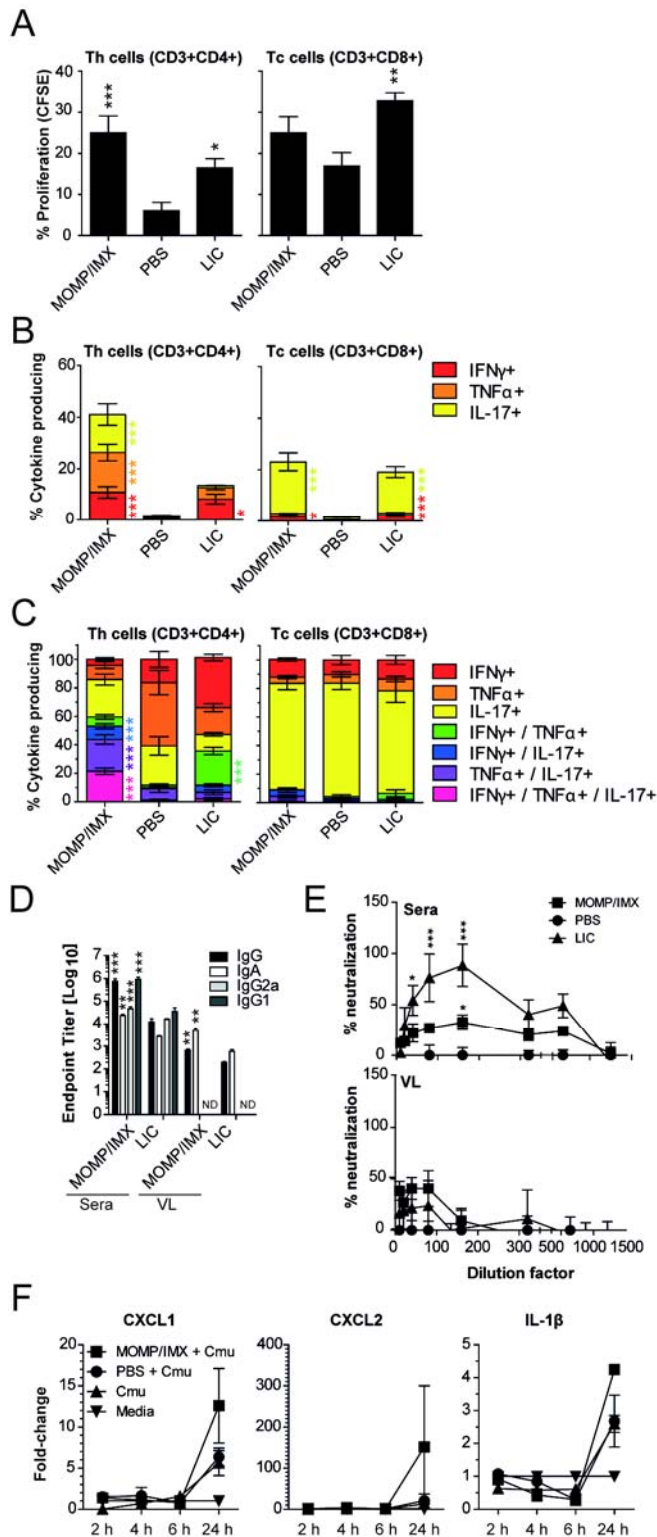
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**Figure 1: MOMP-specific T and B cell responses following immunization**

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Cells were isolated from spleens of immunized (MOMP/IMX) or nonimmunized (PBS) mice (7 day post-immunization) or infected mice (intravaginally challenged with  $5 \times 10^4$  IFU of *C.*

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*muridarum*, 21 day p.i), stimulated with MOMP for 5 days and analyzed by flow cytometry

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for antigen-specific proliferation and cytokine production. Live cells (viability stain) were

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gated initially on single cells (FSC-A/FSC-H) and lymphocytes (FSC/SSC) before gating

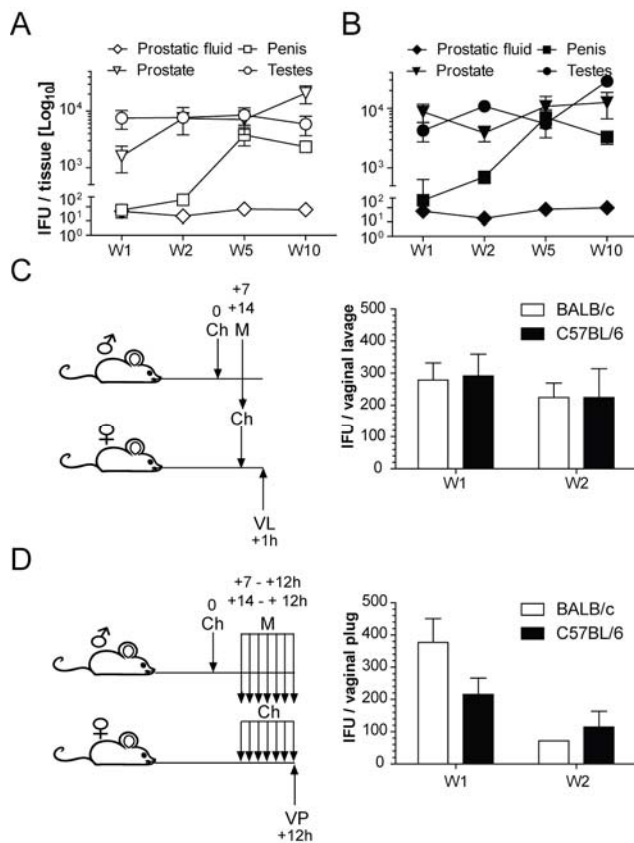
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CD3+ cells on Th (CD3+CD4+) and Tc cells (CD3+CD8+). (A) Cells were labeled with

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623 CFSE and stimulated with MOMP to measure the percentages of MOMP-specific Th and Tc  
624 cell proliferation (n = 10). (B) MOMP-specific Th cells staining positive for IFN $\gamma$ , TNF $\alpha$  and  
625 IL-17 are represented as a percentage of total Th cells (n = 10). (C) MOMP-specific Th cells  
626 staining positive for TNF $\alpha$  were assessed for multifunctional phenotype (IFN $\gamma$  and IL-17) are  
627 represented as a percentage of total cytokine producing Th cells (n = 10). (D) MOMP-  
628 specific IgG, IgA, IgG2a and IgG1 endpoint titers were quantified in serum and vaginal  
629 lavage (VL) samples by ELISA. (E) *In vitro* neutralization of *Chlamydia* infectivity was  
630 determined for serum and vaginal lavage by culture (n = 10). (F) The ability of serum  
631 collected from MOMP/IMX immunized mice to opsonize *Chlamydia* and promote CXCL1,  
632 CXCL2 and IL-1 $\beta$  production by monocytes/macrophages (RAW264.7 cell line) was  
633 determined using RT-PCR. Results are presented as the mean  $\pm$  SEM. ND refers to “not  
634 done”. Significant differences were determined using a one-way ANOVA with Tukey’s post-  
635 test. Significance was set at  $p < 0.05$  for all tests.  $p > 0.05$  (not shown), 0.01-0.05 (\*), 0.001-  
636 0.01 (\*\*) and  $< 0.001$  (\*\*\*).

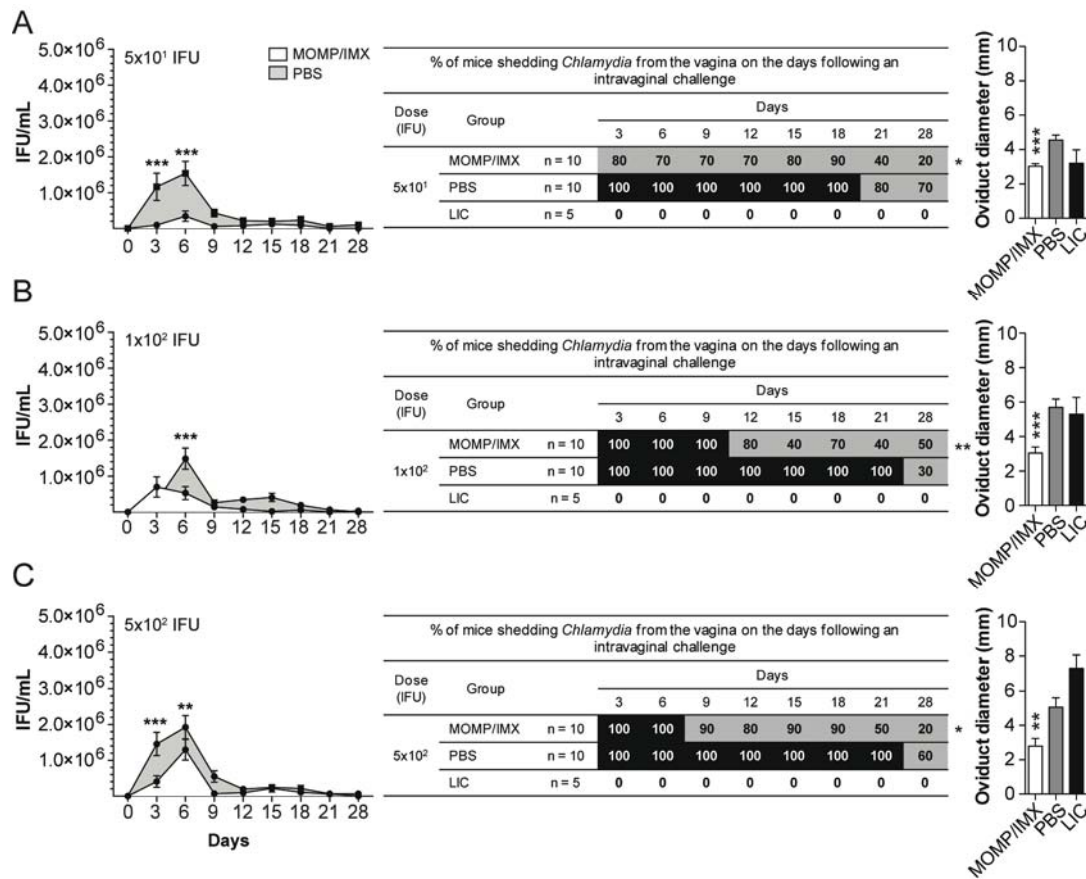
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**Figure 2: Quantification of the chlamydial dose sexually transmitted by male mice.**

(A) Male BALB/c and (B) C57BL/6 mice were challenged with  $1 \times 10^6$  IFU *C. muridarum* and the chlamydial burden was quantified from the prostatic fluid, penis, prostate and testes collected on week 1, 2, 5 and 10 p.i by culture (n = 5). (C) Male mice were challenged (Ch) with *C. muridarum* and mated (M) with non-progesterone primed females (receptive) and the chlamydial load transmitted was determined from vaginal lavages (VL) (n = 11) and (D) vaginal plugs (VP) (n = 15) during weeks 1 and 2 of their infection. Results are presented, as the mean  $\pm$  SEM. Significant differences were determined using a one-way ANOVA with Tukey's post-test. Significance was set at  $p < 0.05$  for all tests.  $p > 0.05$  (not shown), 0.01-0.05 (\*), 0.001- 0.01 (\*\*) and  $< 0.001$  (\*\*\*)



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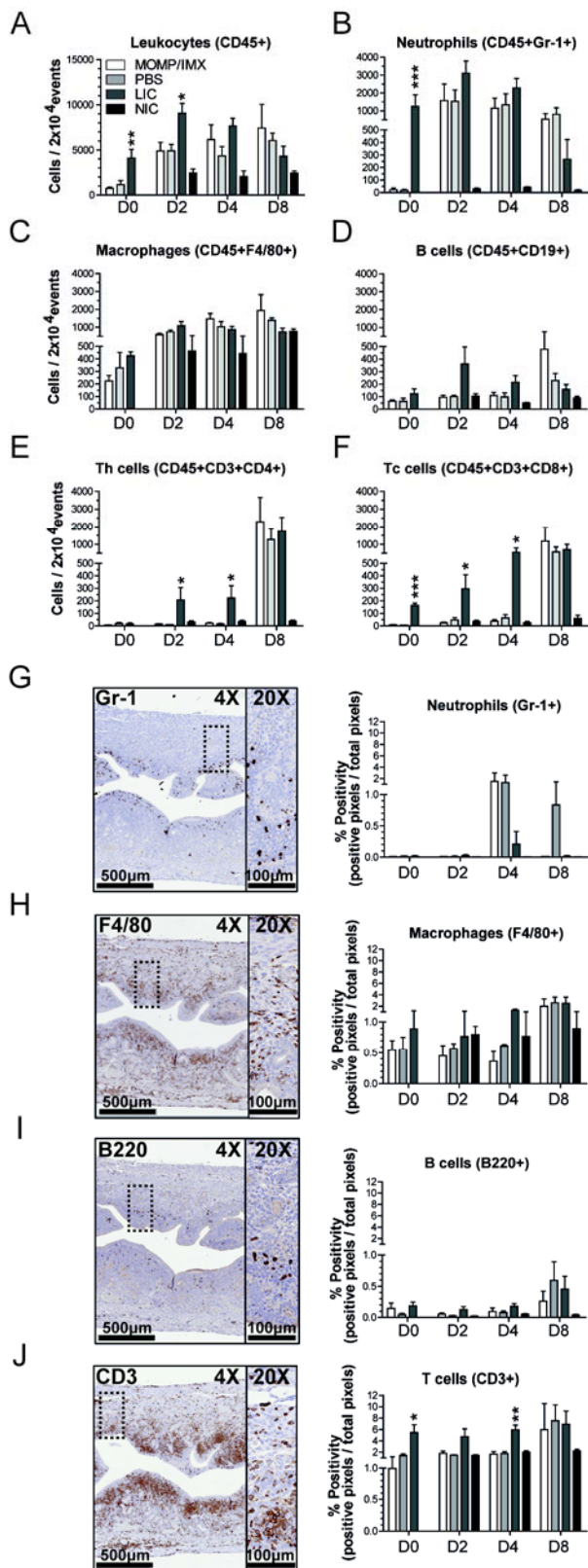
662 **Figure 3: Protection against genital tract infection and oviduct pathology in female mice**  
 663 **following immunization and challenge with sexually transmissible doses of *C.***  
 664 ***muridarum*.**

665 Female BALB/c mice were immunized (MOMP/IMX), nonimmunized (PBS) or infected (1°  
 666 infection, intravaginally challenged with 5x10<sup>1</sup>, 1x10<sup>2</sup> and 5x10<sup>2</sup> IFU of *C. muridarum*) then  
 667 challenged with (A) 5x10<sup>1</sup>, (B) 1x10<sup>2</sup>, (C) 5x10<sup>2</sup> IFU of *C. muridarum*. Vaginal swabs were  
 668 collected over 28 days p.i to quantify the amount of *Chlamydia* shed (IFU/mL) by culture.  
 669 Significant levels of bacterial burden were determined using a two -way ANOVA with  
 670 Tukey's post -test. Results are presented as the mean ± SEM. Percentages of mice shedding  
 671 *Chlamydia* from the vagina on the days following challenge are also presented. Significant  
 672 differences were determined using a Kaplan -Meier survival curve and the log rank post -test.  
 673 The severity of pathology was determined by measuring the diameter of the oviducts post-  
 674 mortem on day 35 p.i. The solid line across the graph represents the mean width of unaffected  
 675 oviducts (2 mm). Significance of disease severity was determined using a one -way ANOVA  
 676 with Tukey's post -test. Significance was set at  $p < 0.05$  for all tests.  $p > 0.05$  (not shown),  
 677 0.01-0.05 (\*), 0.001- 0.01 (\*\*), and  $< 0.001$  (\*\*\*)

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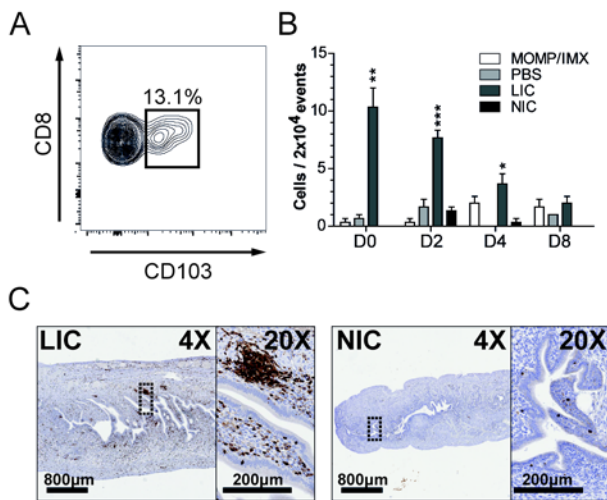


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 682 **Figure 4: Kinetics of leukocyte, neutrophil, macrophage, B, Th and Tc cell recruitment**  
 683 **following immunization and challenge with a sexually transmissible dose of *C.***  
 684 ***muridarum*.**

685 Female BALB/c mice were immunized (MOMP/IMX), nonimmunized (PBS) or infected (1°  
 686 infection, intravaginally challenged with  $5 \times 10^1$  IFU of *C. muridarum*) then challenged with

687  $5 \times 10^1$  IFU of *C. muridarum*. Tissues were collected from immunized mice (MOMP/IMX and  
688 PBS) prior to challenge (day 0) and on days 2, 4 and 8 following challenge (n = 6). Tissues  
689 were also collected from uninfected animals (no infection control – NIC) on days 2, 4 and 8  
690 and mice recovering from an intravaginal infection (day 0 = 1° infection, day 21 p.i) and on  
691 days 2, 4 and 8 p.i following re-challenge (live infection control – LIC) (n = 6). Cells were  
692 isolated from the uterine horns and cervix/vagina by digestion for analysis by flow cytometry.  
693 Live cells (viability stain) were gated initially on single cells (FSC-A/FSC-H) and (A)  
694 leukocytes (CD45+) before gating on (B) neutrophils (Gr-1+), (C) macrophages (F4/80+),  
695 (D) B cells (CD19+), (E) Th cells (CD3+CD4+) and (F) Tc cells (CD3+CD8+). Results are  
696 presented as number of cells per  $2 \times 10^4$  events. Uterine horns were also fixed for analysis by  
697 immunohistochemistry (n = 6). Serial tissue sections of the uterine horns were stained for the  
698 presence of (G) neutrophils (Gr-1+), (H) macrophages (F4/80+), (I) B cells (B220+), (J) and  
699 T cells (CD3+). Representative images (4x and 20x magnifications) are shown for each stain.  
700 ImageScope software was used to quantify the percentage positivity (positive pixels/total  
701 pixels) from a high-resolution image of each section. Results are presented as the mean  $\pm$   
702 SEM. Significant differences were determined using a one-way ANOVA with Tukey's post-  
703 test. Significance was set at  $p < 0.05$  for all tests.  $p > 0.05$  (not shown), 0.01-0.05 (\*), 0.001-  
704 0.01 (\*\*) and  $< 0.001$  (\*\*\*)).

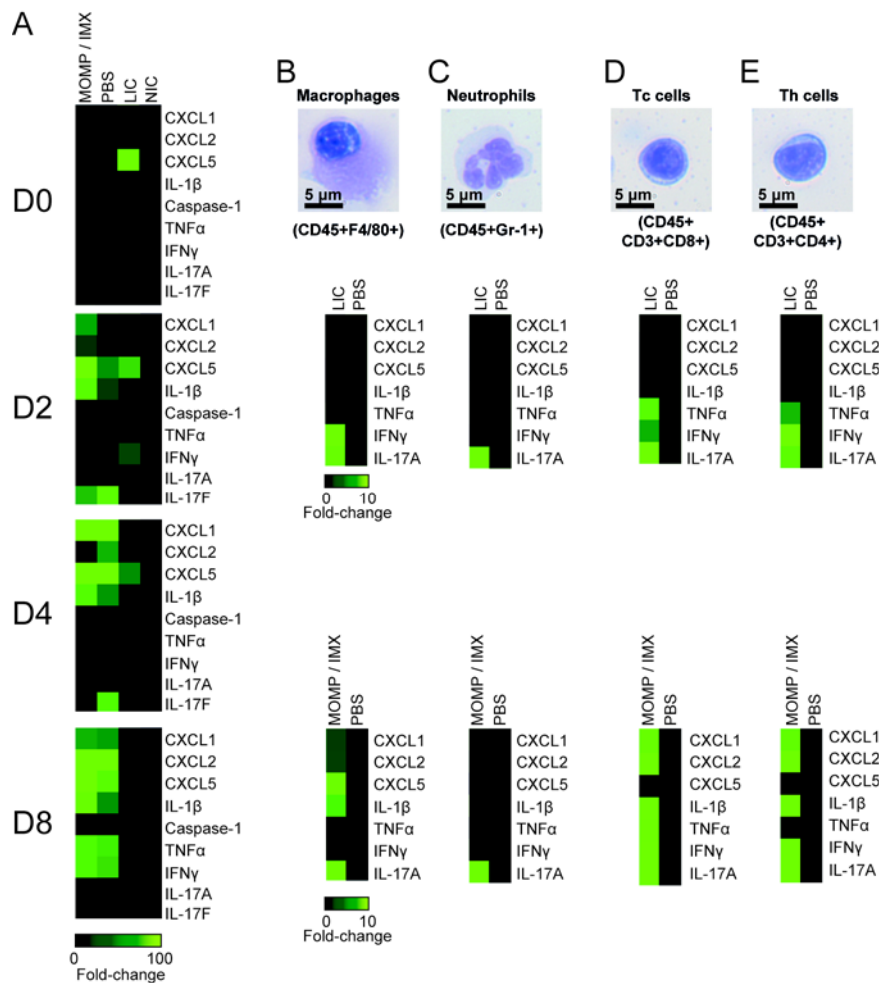
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**Figure 5: Kinetics of T<sub>RM</sub> cells (CD3+CD8+CD103+) following immunization and challenge with a sexually transmissible dose of *C. muridarum*.**

Female BALB/c mice were immunized (MOMP/IMX), nonimmunized (PBS) or infected (1° infection, intravaginally challenged with 5x10<sup>1</sup> IFU of *C. muridarum*) then challenged with 5x10<sup>1</sup> IFU of *C. muridarum*. Tissues were collected from immunized mice (MOMP/IMX and PBS) prior to challenge (day 0) and on days 2, 4 and 8 following challenge (n = 3). Tissues were also collected from uninfected animals (no infection control – NIC) and mice recovering from an intravaginal infection (day 0 = 1° infection, day 21 p.i) and on days 2, 4 and 8 p.i following re-challenge (live infection control – LIC) (n = 3). Cells were isolated from the uterine horns and cervix/vagina by digestion for analysis by flow cytometry. (A) Resident memory Tc were identified by gating on lymphocytes (FSC/SSC), single cells (FSC -A/FSC -H), leukocytes (CD45+), Tc cells (CD3+CD8+) and finally CD103+. (B) Cells per 2x10<sup>4</sup> events are presented for each group. Results are presented as the mean ± SEM. (C) CD3 staining of tissue sections from uterine horns collected from LIC and NIC groups prior to challenge. Significant differences were determined using a one -way ANOVA with Tukey’s post -test. Significance was set at *p* < 0.05 for all tests. *p*>0.05 (not shown), 0.01-0.05 (\*), 0.001- 0.01 (\*\*), and <0.001 (\*\*\*).



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740 **Figure 6: Gene expression in whole oviducts, macrophages, neutrophils, Tc and Th cells**  
 741 **isolated from the oviducts following immunization and challenge with a sexually**  
 742 **transmissible dose of *C. muridarum*.**

743 Female BALB/c mice were immunized (MOMP/IMX), nonimmunized (PBS) or infected (1°  
 744 infection, intravaginally challenged with  $5 \times 10^1$  IFU of *C. muridarum*) then challenged with  
 745  $5 \times 10^1$  IFU of *C. muridarum*. Tissues were collected from immunized mice (MOMP/IMX and  
 746 PBS) prior to challenge (day 0) and on days 2, 4 and 8 following challenge. Tissues were also  
 747 collected from uninfected animals (no infection control – NIC) and mice recovering from an  
 748 intravaginal infection (day 0 = 1° infection, day 21 p.i) and on days 2, 4 and 8 p.i following  
 749 re-challenge (live infection control – LIC) (n = 3). (A) Gene expression was performed on  
 750 pooled oviducts (n = 3) and fold change (scale 0 – 100-fold) was calculated relative to the  
 751 uninfected control (NIC). Live cells were isolated from the genital tract on days 2 and 8 p.i  
 752 and separated into (B) macrophages (CD45+F4/80+) (C) neutrophils (CD45+Gr-1+), (D) Tc  
 753 (CD45+CD3+CD8+) and (E) Th cells (CD45+CD3+CD4+) using flow cytometry (n = 10).  
 754 Cytospin and Geimsa staining was performed on sorted cell fractions to compare surface  
 755 marker phenotype with nuclear morphology (representative image shown). Gene expression  
 756 was performed on pooled cells isolated from the oviducts, uterine horns and cervix vagina (n  
 757 = 10), and fold changes (scale 0 – 10-fold) were calculated relative to an unimmunized  
 758 control. Significant differences were determined using a one-way ANOVA with Tukey's



759 post-test. Significance was set at  $p < 0.05$  for all tests.  $p > 0.05$  (not shown), 0.01-0.05 (\*),  
760 0.001- 0.01 (\*\*) and  $< 0.001$  (\*\*\*)

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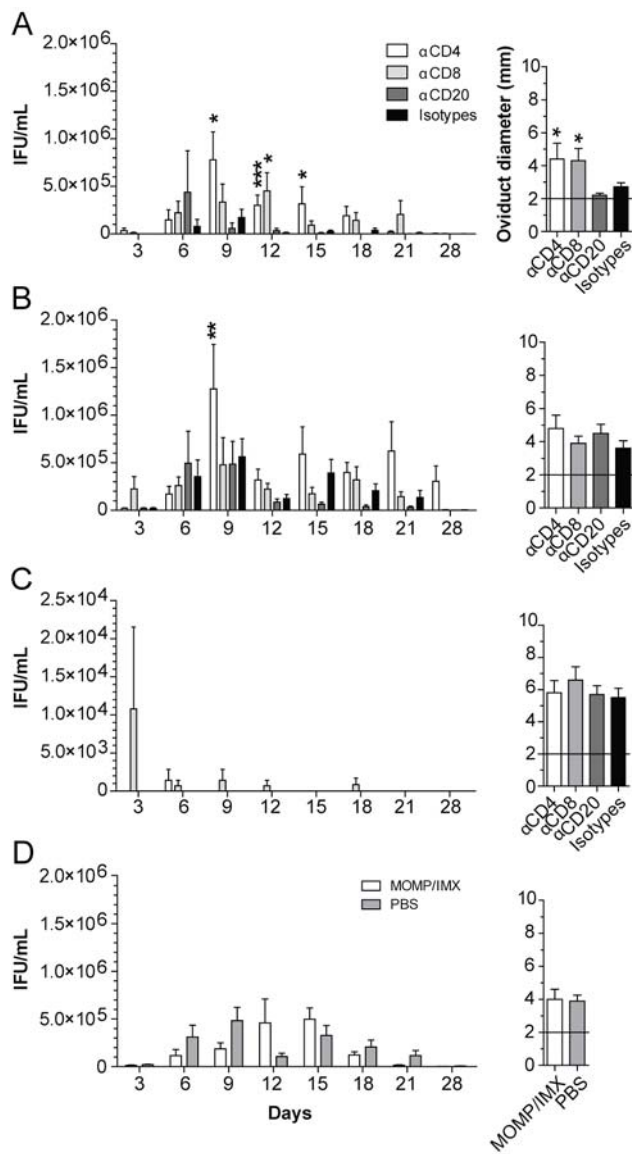
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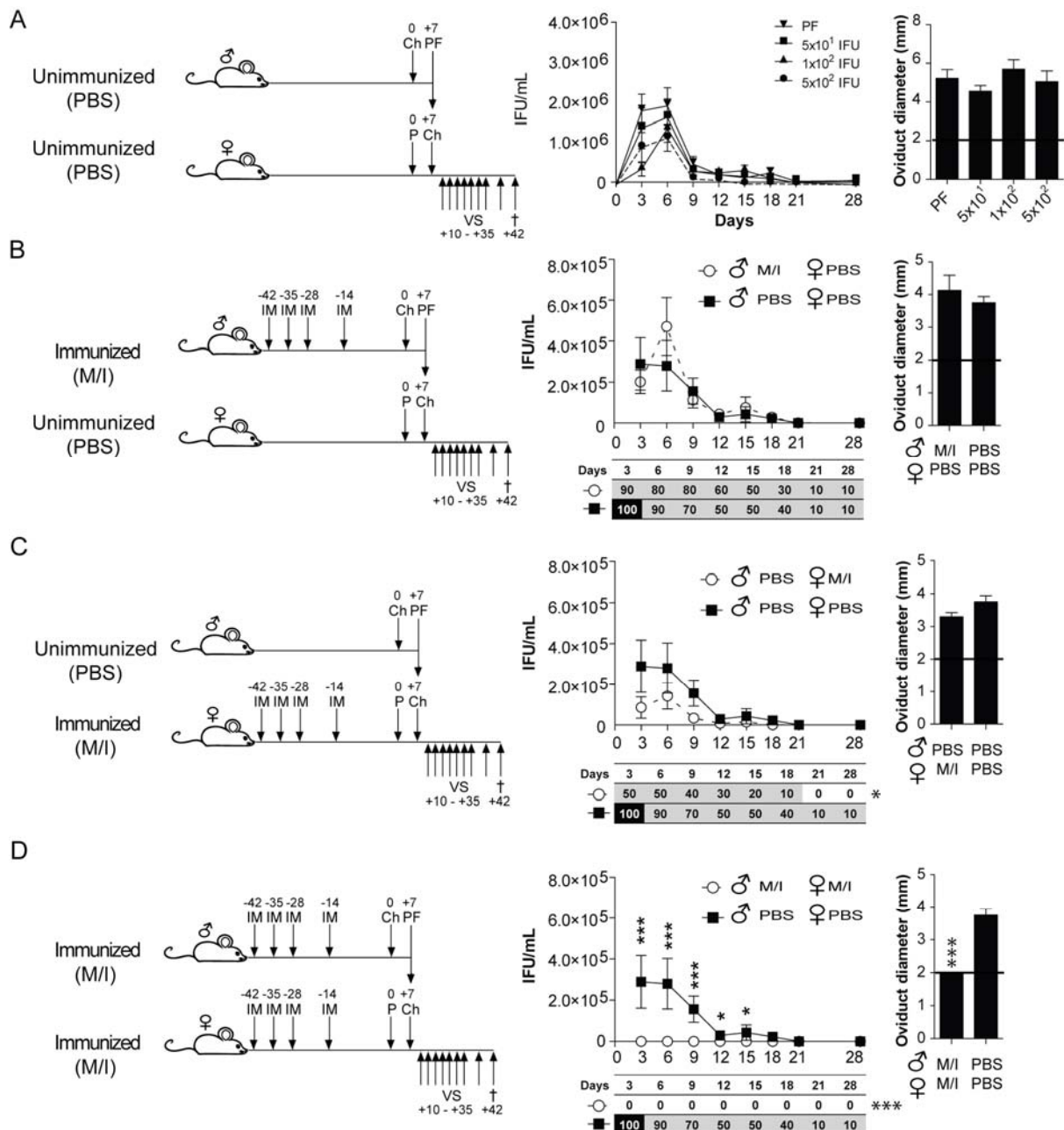


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777 **Figure 7: *In vivo* depletion (CD4+, CD8+ and CD20+) and passive immunization prior**  
 778 **to challenge with a sexually transmissible dose of *C. muridarum*.**

779 Female BALB/c mice were immunized (MOMP/IMX), nonimmunized (PBS) or infected (1°  
 780 infection, intravaginally challenged with 5x10<sup>4</sup> IFU of *C. muridarum*) and depleted of CD4,  
 781 CD8 and CD20 cells (n = 5). Unimmunized mice were also passively immunized with serum  
 782 collected from MOMP/IMX immunized mice (n = 5). Depleted and passively immunized  
 783 mice were then challenged with 5x10<sup>1</sup> IFU of *C. muridarum*. Vaginal infection and oviduct  
 784 pathology were quantified from (A) MOMP/IMX, (B) unimmunized, (C) LIC and (D)  
 785 passively immunized naïve mice. Vaginal swabs were collected for 28 days p.i to quantify the  
 786 amount of *Chlamydia* shed (IFU/mL) by culture. The severity of pathology was determined  
 787 on day 35 p.i by measuring the diameter of the oviducts. The solid line across the graph  
 788 represents the mean width of unaffected oviducts (2 mm). Results are presented as the mean  
 789 ± SEM. Significant levels of bacterial burden and disease severity were determined using a  
 790 one-way ANOVA with Tukey's post-test. Significance was set at  $p < 0.05$  for all tests.  
 791  $p > 0.05$  (not shown), 0.01-0.05 (\*), 0.001- 0.01 (\*\*), and <0.001 (\*\*\*)).

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**Figure 8: Transmission of *Chlamydia* between male and female mice with different vaccination statuses.**

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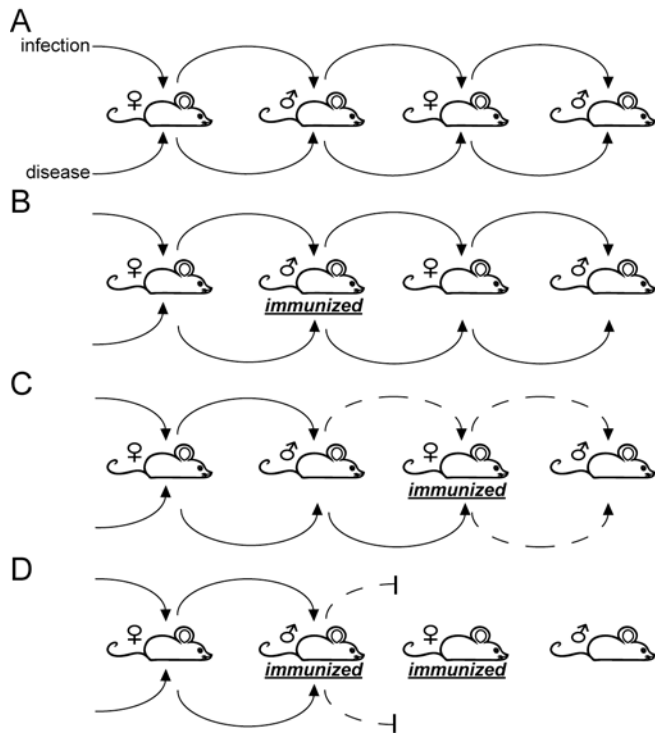
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(A) Male mice were challenged (Ch) with  $1 \times 10^6$  IFU *C. muridarum* and their prostatic fluids (PF) were collected 7 days p.i (n = 10). Female mice were progesterone-primed (P) 7 days prior to being challenged intravaginally with prostatic fluid collected from the infected males (n = 5). Vaginal shedding and development of hydrosalpinx following challenge with PF was compared to challenge of female mice with  $5 \times 10^1$ ,  $1 \times 10^2$  and  $5 \times 10^2$  of *C. muridarum*. This model of sexual transmission was repeated using (B) males, (C) females and (D) males and females that had been immunized (IM) with MOMP/IMX (M/I) prior to challenge with PF (n = 10). Vaginal swabs (VS) were collected over 28 days p.i to quantify the amount of *Chlamydia* shed (IFU/mL) by culture. Significant levels of bacterial burden were determined using a two-way ANOVA with Tukey's post-test. Results are presented as the mean  $\pm$  SEM. Percentages of mice shedding *Chlamydia* from the vagina on the days following challenge

807 are also presented. Significant differences were determined using a Kaplan -Meier survival  
808 curve and the log rank post-test. Protection against the severity of oviduct pathology  
809 (hydrosalpinx) in female mice challenge with prostatic fluid was determined post mortem (†)  
810 on day 35 p.i by measuring oviduct diameter. The solid line across the graph represents the  
811 mean width of unaffected oviducts (2 mm). Results are presented as the mean ± SEM.  
812 Significant levels of bacterial burden and disease severity were determined using a one -way  
813 ANOVA with Tukey's post -test. Significance was set at  $p < 0.05$  for all tests.  $p > 0.05$  (not  
814 shown), 0.01-0.05 (\*), 0.001- 0.01 (\*\*), and  $< 0.001$  (\*\*\*)).

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835 **Figure 9: Prevention of *Chlamydia* sexual transmission by vaccination.**

836 (A) *Chlamydia* infection transmission occurs unimpeded through unimmunized mice. (B) Risk of  
 837 infection transmission to unimmunized females is unaffected by the male vaccination status. (C)  
 838 Vaccinating female mice reduces their risk of acquiring an infection from unimmunized males by  
 839 50%, but not their risk of developing disease. As half of immunized females were protected from  
 840 an infection, risk of infection transmission and disease developing in unimmunized males is also  
 841 reduced. (D) No transmission of infection occurs between immunized males and females.  
 842 Immunized females show no signs of disease. Further transmission to unimmunized males is also  
 843 prevented.

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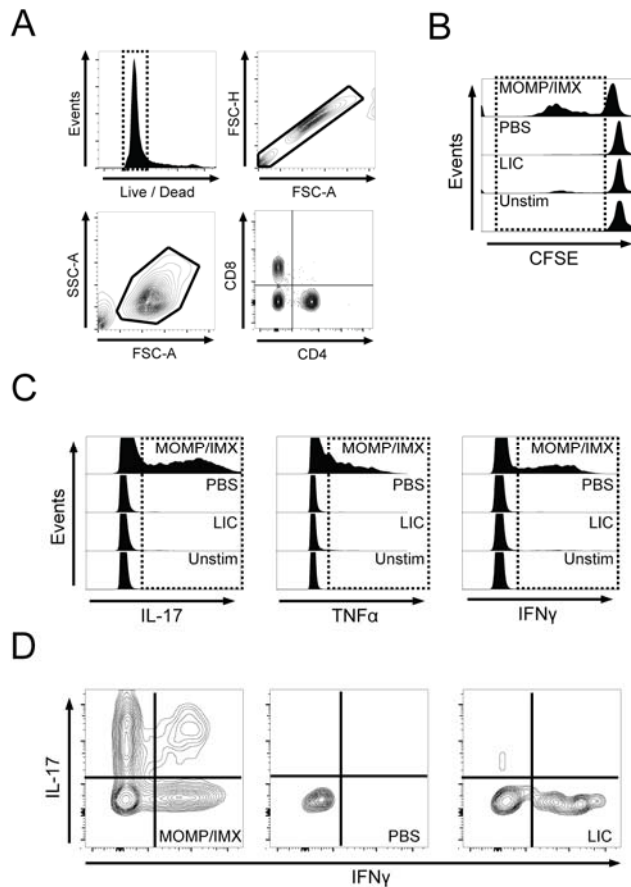
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851 **S1: *Chlamydia* quantified from infectious male mouse ejaculate**

Sample	Prostatic fluid	Vaginal lavage	Vaginal plug
Min – Max (IFU)	18 – 224	135 – 449	72 – 503
Mean ± SD (IFU)	68 ± 45	261 ± 98	220 ± 150
Min – Max (ompA copies)	2080 – 1.2x10 <sup>5</sup>	ND	1.7x10 <sup>5</sup> – 3x10 <sup>7</sup>
Mean ± SD (ompA copies)	3x10 <sup>4</sup> ± 665 <sup>7</sup>	ND	5x10 <sup>6</sup> ± 8.5x10 <sup>6</sup>

852 Minimum, maximum and mean amount of *Chlamydia* collected from prostatic fluid, vaginal  
 853 lavage and vaginal plugs determined by culture (IFU) and RT-PCR (*ompA* copies). Results  
 854 are a combination of both strains (BALB/c and C57BL/6) and time points (W1 and W2). ND  
 855 abbreviation refers to “not done”.

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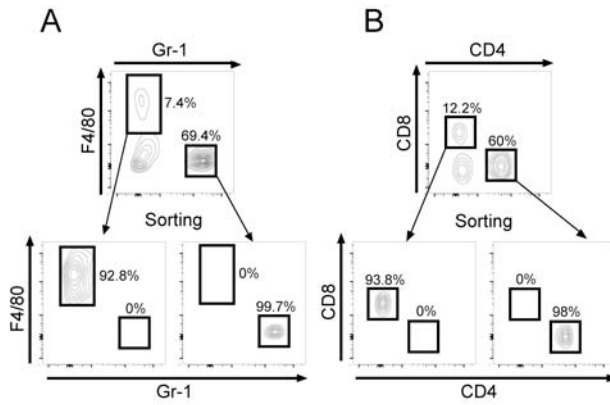
866 **S2: Gating strategies for MOMP-specific T cell responses following immunization**

867 (A) Live cells (viability stain) were gated initially on single cells (FSC-A/FSC-H) and  
 868 lymphocytes (FSC/SSC) before gating CD3+ cells on Th (CD3+CD4+) and Tc cells  
 869 (CD3+CD8+). (B) Cells were labeled with CFSE and stimulated with MOMP to measure the  
 870 percentages of MOMP-specific Th and Tc cell proliferation. (C) MOMP-specific Th cells  
 871 staining for IFN $\gamma$ , TNF $\alpha$  and IL-17. (D) MOMP-specific Th cells staining positive for TNF $\alpha$   
 872 were assessed for multifunctional phenotype (IFN $\gamma$  and IL-17) and represented as a  
 873 percentage of total cytokine producing Th cells.

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878 **S3: Gating strategies for macrophages, neutrophils, Tc and Th cells isolation and purity**

879 Live cells were isolated from the genital tract on days 2 and 8 p.i and separated into (A)  
 880 macrophages (CD45+F4/80+) and neutrophils (CD45+Gr-1+), (B) Tc (CD45+CD3+CD8+)  
 881 and Th cells (CD45+CD3+CD4+) using flow cytometry. Representative plots for  
 882 macrophages (92.8%) and neutrophils (99.7%), Tc (93.8%) and Th cells (98%) purity are  
 883 shown.

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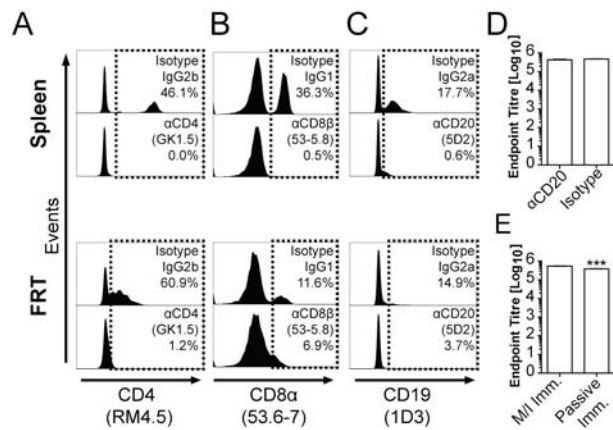
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896 **S4: Confirmation of *in vivo* depletion (CD4+, CD8+ and CD20+) and passive**  
 897 **immunization**

898 Splens and genital tracts were collected 2 days following intraperitoneal infusion of (A)  
 899 αCD4, (B) αCD8 and (C) αCD20 and depletion was confirmed using flow cytometry.  
 900 Depletion was compared against isotype control infused mice. (D) Serum was screened for  
 901 the presence of MOMP-specific Ig in MOMP/IMX immunized mice by ELISA following  
 902 αCD20 infusion. (E) MOMP-specific Ig in the serum collected from MOMP/IMX immunized  
 903 donor mice (M/I Imm.) and serum from recipient naïve mice passively immunized with  
 904 MOMP/IMX serum (Passive Imm.) were analyzed by ELISA.

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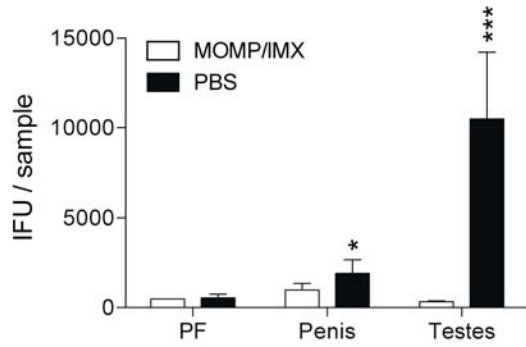
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915 **S5: Protection against genital tract infection in male mice following immunization and**  
 916 **challenged with *C. muridarum*.**

917 Male mice were challenged with  $1 \times 10^6$  IFU *C. muridarum* and their prostatic fluid (PF),  
 918 penis and testes were collected 7 days p.i (n = 5). The amount of *Chlamydia* presented in  
 919 each sample (IFU/sample) was quantified by culture. Results are presented as the mean  $\pm$   
 920 SEM. Significant levels of bacterial burden and disease severity were determined using a one  
 921 -way ANOVA with Tukey's post -test. Significance was set at  $p < 0.05$  for all tests.  $p > 0.05$   
 922 (not shown), 0.01-0.05 (\*), 0.001- 0.01 (\*\*), and  $< 0.001$  (\*\*\*).

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948 **S6: Electro-stimulation schedule for collection of prostatic fluid**

	<b>Frequency (Hz)</b>	<b>Pulse width (ms)</b>	<b>Intensity (V)</b>	<b>Duration (min:sec)</b>
*1.	1.0	1.0	0.5	0:30
2.	1.0	1.0	1.0	0:30
3.	1.0	1.0	1.5	0:30
4.	1.0	1.0	2.0	0:30
5.	2.0	1.0	2.5	0:30
6.	2.0	1.0	5.0	0:30
#7.	2.0	1.0	7.5	0:30
8.	2.0	1.0	10.0	0:30
†9.	2.0 – 10.0 – 2.0	1.0	10.0	0:04 x 5
10.	2.0 – 10.0 – 2.0	1.0	15.0	0:04 x 5
11.	2.0 – 10.0 – 2.0	1.0	20.0	0:04 x 5
			<b>Total</b>	<b>5:00</b>

949 \*Urine contamination can be produced early during the procedure and must be discarded.  
 950 #Prostatic fluid is typical produced from this point onwards. †Frequency is increased for 1  
 951 sec, held at maximum for 2 sec and decreased for 1 sec. This cycle is repeated five times for  
 952 each intensity setting

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977 **S7: Primers for RT-PCR**

<b>Gene</b>	<b>Accession No.</b>	<b>Forward Primer (5' to 3')</b>	<b>Reverse Primer (5' to 3')</b>
CXCL1	NM_008176.3	ACCCAAACCGAAGTCATAGCC	TTGTCAGAAGCCAGCGTTCA
CXCL2	NM_009140.2	CCCAGACAGAAGTCATAGCCAC	CGAGGCACATCAGGTACGAT
CXCL5	NM_009141.3	GCCCCTTCCTCAGTCATAGC	AGCTTTCTTTTTGTCAGTCCCC
IL-1 $\beta$	NM_008361.3	TGCCACCTTTTGACAGTGATG	AAGGTCCACGGGAAAGACAC
Caspase-1	NM_009807.2	TGAAAGAATTTGCTGCCTGCC	CCTTGTTTCTCTCCACGGCAT
IFN $\gamma$	NM_008337.3	AGGAACTGGCAAAGGATGGT	TCATTGAATGCTTGCGCTG
IL-17A	NM_010552.3	TGAGTCCAGGGAGAGCTTCA	CATTGCGGTGGAGAGTCCAG
IL-17F	NM_145856.2	CGTGAAACAGCCATGGTCAAG	GGGGTCTCGAGTGATGTTG
TNF $\alpha$	NM_013693.3	ACGTCGTAGCAAACCACCAA	ATAGCAAATCGGCTGACGGT
ActB	NM_007393.3	AGAGGGAAATCGTGCGTGAC	CAATCGTGATGACCTGGCCGT

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