1 Supplementary Figures



FIGURE S1 DCs differentiation and viability A. Flow cytometry plot showing the percentage of CD11c⁺ cells induced by GMCSF/IL-4 for 8 days in each group. **B** Bar graph showing the mean percentage of CD11c expression between the control and the infection groups. **C.** Flow cytometry plot shows 7-AAD⁺ cells in the X-axis and Annexin V⁺ cells in the Y-axis. The percentage of live cell population (cells that are negative for Annexin V and 7-AAD) are found in the lower left quadrant of the plot in each group. **D**. Bar graph showing the mean percentage of live cells between the control and the infection groups. The experiment was repeated three times in triplicate.





FIGURE S2 Maturation and cytokine induction of control DCs, cocultured with *E. faecalis* after differentiation A. Scattergrams from flow cytometry of (control) DCs treated with LPS or *E.*

after differentiation A. Scattergrams from flow cytometry of (control) DCs treated with LPS or *E. faecalis* after differentiation with GM-CSF/IL-4. CD11c⁺ on the X-axis and MHCII⁺ (Top), CD86⁺ (Middle) or CD80⁺ (Bottom) on the Y-axis. **B.** Bar graphs of mean fold change in MHCII⁺, CD86⁺ and CD80⁺CD11c⁺ DCs in the indicated groups. **C.** Bar graphs of mean fold change of mRNA expression of IL1 β , TNF α , IL12 & TGF β 1 by RT-PCR, normalized relative to GAPDH, of control DCs, LPS stimulated control DCs and differentiated DCs infected with *E. faecalis*. The experiment was repeated three times in triplicate.

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