



OPEN **V γ 9+V δ 2+ T cell control of *Listeria monocytogenes* growth in infected epithelial cells requires butyrophilin 3A genes**

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Intracellular bacteria produce antigens, which serve as potent activators of $\gamma\delta$ T cells. Phosphoantigens are presented via a complex of butyrophilins (BTN) to signal infection to human V γ 9+V δ 2+ T cells. Here, we established an in vitro system allowing for studies of V γ 9+V δ 2+ T cell activity in coculture with epithelial cells infected with the intracellular bacterial pathogen *Listeria monocytogenes*. We report that the V γ 9+V δ 2+ T cells efficiently control *L. monocytogenes* growth in such cultures. This effector function requires the expression of members of the BTN3A family on epithelial cells. Specifically, we observed a BTN3A1-independent BTN3A3 activity to present antigen to V γ 9+V δ 2+ T cells. Since BTN3A1 is the only BTN3A associated with phosphoantigen presentation, our study suggests that BTN3A3 may present different classes of antigens to mediate V γ 9+V δ 2+ T cell effector function against *L. monocytogenes*-infected epithelia.

The Gram-positive bacterium *Listeria (L.) monocytogenes* is causative agent of food-borne diseases. Infection via contaminated food may cause systemic spread of the bacteria and result in life-threatening meningoenitis or, in case of maternofetal infections, fetal abortion (reviewed in¹). Systemic bacterial spread results from *L. monocytogenes*' ability to cross the intestinal, blood–brain and placental barriers. Virulence factors encoded by the *L. monocytogenes* pathogenicity islands endow the bacterium with the ability to force uptake into the cytoplasm of epithelial cells as well as intracellular replication and cell-to-cell spread. In phagocytic hosts *L. monocytogenes* escapes from phagosomes to replicate in the cytoplasm and spread between host cells (reviewed in^{2,3}).

Defense mechanisms against intracellular bacterial infections include macrophage activation and the mobilization of cytotoxic T cells (reviewed in⁴). In humans, a special cytotoxic T cell subset of V γ 9+V δ 2+ T cells recognizes bacterial antigens (reviewed in⁵). V γ 9+V δ 2+ T cells expand in the peripheral blood in patients with *L. monocytogenes* infection, exceeding the levels of age-matched controls by five to sixfold on average⁶. The V γ 9+V δ 2+ T cell expansion among peripheral blood mononuclear cells (PBMCs) could be recapitulated in vitro by exposing the cells to either viable *L. monocytogenes*⁷ or extracts from heat-killed bacteria^{6,8,9}. Consistent with this, V γ 9+V δ 2+ T cell expansion was reproduced in non-human primate *L. monocytogenes* infection models¹⁰. Tanaka and colleagues identified (E)-4-Hydroxy-3-methyl-but-2-enyl Pyrophosphate (HMB-PP) as the immunostimulatory agent for the V γ 9+V δ 2+ T cell expansion. HMB-PP belongs to the group of phosphoantigens. It is a metabolite of the deoxyxylulose pathway for isoprenoid synthesis in *L. monocytogenes*¹¹ and various other bacterial or parasitic pathogens⁵. Further support for the importance of HMB-PP was provided by extracts from bacterial mutants with increased HMB-PP levels that strongly stimulated V γ 9+V δ 2+ T cell growth in a PBMC culture¹² and from HMB-PP-deficient *L. monocytogenes* strains that revealed weaker V γ 9+V δ 2+ T cell activating potential. The data underline a non-exclusive but important role of HMB-PP¹³.

BTN3A1 was identified and confirmed as important antigen-presenting molecule of HMB-PP and other phosphoantigens^{14–17}. BTN3A is a member of the butyrophilins, plasma membrane-resident proteins related to the B7 costimulator family. The BTN3A family consists of three genes, BTN3A1, BTN3A2 and BTN3A3, however only BTN3A1 contains a positively charged pocket in its 30.2 domain, that was shown to bind

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phosphoantigens^{18,19}. Very recently another butyrophilin family member, BTN2A1, was described to bind both BTN3A1 and the V γ 9 T cell receptor (TCR) and to play an essential role in V γ 9+V δ 2+ T cell recognition^{20–22}.

The majority of experimental approaches to study the impact of V γ 9+V δ 2+ T cells on *L. monocytogenes* infection is based on the use of bacterial extracts, synthetic phosphoantigens or drugs enhancing intracellular phosphoantigen levels. Noteworthy exceptions are studies in non-human primates⁵ and a report of V γ 9+V δ 2+ T cell cytotoxicity against infected monocyte-derived dendritic cells⁹. However, a system exploring the fate of *L. monocytogenes*-infected epithelial cells exposed to V γ 9+V δ 2+ T cells is missing. Epithelial cells represent the primary targets of gastrointestinal infection. The ability of host organisms to limit replication and spread of *L. monocytogenes* at epithelial barriers determines whether potentially life-threatening systemic infection will ensue.

The aim of this study was the establishment of an in vitro model to investigate consequences of subjecting infected epithelial cells to co-culture with V γ 9+V δ 2+ T cells. This system enabled us to monitor viability and the bacterial burden of target cells as well as measuring changes in the cytokine milieu. Gene deletion allowed us to scrutinize the roles of BTN3A genes. We observed a BTN3A1-independent BTN3A3 activity allowing V γ 9+V δ 2+ T cells to control bacterial growth in cultures of infected cells, thus rescuing them from complete killing by *L. monocytogenes* infection. Our model extends the experimental options to address the role of V γ 9+V δ 2+ T cells in human infectious disease and identifies both BTN3A1 and BTN3A3 as antigen presenting molecules.

Results

V γ 9+V δ 2+ T cells rescue cell populations from *L. monocytogenes*-induced killing

To fully understand the role of V γ 9+V δ 2+ T cells in *L. monocytogenes* infection, we established and optimized a co-culture system using the ‘real-time cell analysis’ (RTCA) method as a read-out. The RTCA system allows for continuous, label-free monitoring of cell adherence to study the time-resolved effects of V γ 9+V δ 2+ T cells on the growth and viability of infected target cells. Data are shown as % cell indices, where 100 represents the impedance (viability) of the culture at the onset of measurement. As a control for V γ 9+V δ 2+ T cell-induced cytotoxicity we used HDMAPP, a phosphoantigen and synthetic version of the bacterial V γ 9+V δ 2+ T cell activator HMB-PP. The detergent Triton X was used to mimic the effect of complete cell killing. As depicted in Fig. 1a, V γ 9+V δ 2+ T cells alone caused an initial drop of viable RKO colon cancer cells, however, the cells recovered with progressive culture. In contrast, combination of both V γ 9+V δ 2+ T cells and the activator HDMAPP caused a dramatic drop in the cell index within the first 24 h of co-culture and no subsequent recovery. This is in line with current literature showing a cytotoxic effect of V γ 9+V δ 2+ T cells on their target cells when treated with phosphoantigen^{23,24}. In the *L. monocytogenes* model, about 5% of cells were infected with *L. monocytogenes*, at start of co-culture under the conditions used (Supplementary Fig. S1). The cell index of *L. monocytogenes* infected target cells decreased steadily over 7 days as the infection was spreading from cell-to-cell. Strikingly, this drop was not observed in a co-culture system of V γ 9+V δ 2+ T cells and target cells infected with *L. monocytogenes*. Rather, a significant increase of the cell index was recorded, surpassing the cell index of *L. monocytogenes*-infected cells approximately 48 h post infection. This V γ 9+V δ 2+ T cell-induced effect on infected target cells will henceforth be designated as the ‘rescue effect’. To determine whether the rescue effect is specific for RKO cells we performed RTCA assays with the ovarian cancer cell line COV362 (Fig. 1b). V γ 9+V δ 2+ T cells exerted basal cytotoxicity on these cells, but a drop in cell viability in the presence of the phosphoantigen HDMAPP was similarly strong as in RKO cells. Upon *L. monocytogenes* infection, the cell index quickly dropped, and this effect was again rescued in co-culture with V γ 9+V δ 2+ T cells. To test a potential donor-specificity in the observed rescue effect, we performed the RTCA assay using a second V γ 9+V δ 2+ T cell donor and observed similar results (Fig. 1c). In conclusion, V γ 9+V δ 2+ T cells rescue cell cultures from *L. monocytogenes*-induced cell death, thus enabling proliferation of the target cells over time. This effect is consistent across two different human epithelial cell lines as well as two independent V γ 9+V δ 2+ T cell donors.

The V γ 9+V δ 2+ T cell-mediated rescue of *L. monocytogenes*-infected target cells is BTN3A-dependent

Next, we addressed the question whether the observed rescue of *L. monocytogenes* infected cultures by V γ 9+V δ 2+ T cells requires members of the BTN3A family. BTN3A pan knockout RKO cells (lacking BTN3A1, BTN3A2 and BTN3A3) were validated for the reduction of BTN3A1 and BTN3A3 protein levels using the Simple Western™ Automated Western Blot Systems size separation (Supplementary Fig. S2 and S3). Consistent with current literature^{18,22,25,26} BTN3A pan knockout cells failed to succumb to the cytotoxic effect of V γ 9+V δ 2+ T cells in the presence of HDMAPP (Fig. 2a). Likewise, *L. monocytogenes*-infected BTN3A pan knockout RKO cells co-cultured with V γ 9+V δ 2+ T cells did not exhibit the ‘rescue effect’ as their wildtype counterparts. The knockout cells experienced similar decreases of viability as *L. monocytogenes*-infected cells in absence of V γ 9+V δ 2+ T cells. Comparable results were obtained using BTN3A1/3 COV362 double-knockout cells (Fig. 2b, Supplementary Fig. S2 and S3). Thus, rescue of *L. monocytogenes*-induced cell death requires BTN3A1 or BTN3A3-dependent antigen presentation to V γ 9+V δ 2+ T cells in two human epithelial cell lines.

Cytokine production does not reveal clear correlations with BTN3A-dependent clearance of *L. monocytogenes* infection

Cytokines are key modulators of inflammation, shape the cellular environment during infections and reflect immune cell activation. In addition, they establish cell-intrinsic innate immunity against pathogens^{27–29}. Therefore, we sought to understand whether the rescue effect could be linked to steady-state cytokine production by expanded V γ 9+V δ 2+ T cell batches, or whether it requires further V γ 9+V δ 2+ T cell activation. We harvested cell supernatants 48 and 72 h post infection to measure cytokine levels produced by infected wildtype as well as BTN3A pan knockout RKO cells, either with or without co-cultured V γ 9+V δ 2+ T cells. V γ 9+V δ 2+ T cell batches

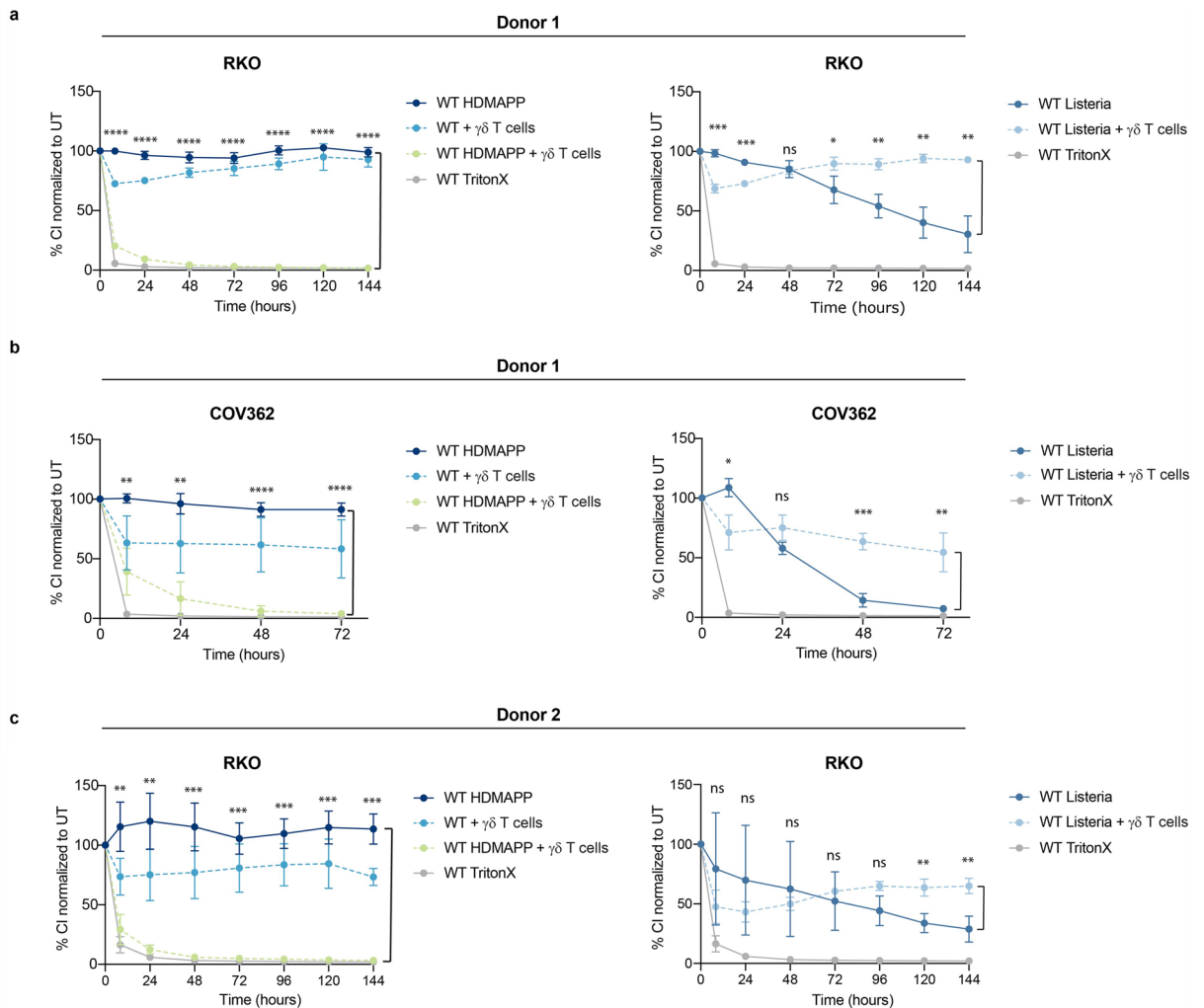


Figure 1. RTCA was performed with RKO (a,c) or COV362 (b) cells co-cultured with $V\gamma 9+V\delta 2+$ T cells donor 1 (a,b) or donor 2 (c). Target cells were either treated with HDMAPP (left panel) or infected with *L. monocytogenes* (right panel). Triton-X was used as positive control for cytotoxicity. Cell Index of the treatments was normalized to the untreated sample (100%). Two-tailed unpaired Student's t-test was performed using GraphPad Prism comparing HDMAPP with HDMAPP + $V\gamma 9+V\delta 2+$ T cells or *L. monocytogenes* with *L. monocytogenes* + $V\gamma 9+V\delta 2+$ T cells. P-values: ns $P > 0.05$; * $P \leq 0.05$; ** $P \leq 0.01$; *** $P \leq 0.001$; **** $P \leq 0.0001$. Experiments were performed in biological triplicates.

cultured in the presence of IL-2 during expansion as well as within the co-culture assays, released considerable amounts of cytokines in an uninfected co-culture system. (Supplementary Fig. S4). Differences in the amounts of secreted cytokines were not statistically significant when comparing cocultures of *L. monocytogenes*-infected WT and BTN3A pan knockout cells. This suggests that under our experimental conditions the presence of BTN3A is not required to stimulate or increase cytokine production by $V\gamma 9+V\delta 2+$ T cells. Infection with *L. monocytogenes* alone failed to increase cytokine production by epithelial cells. This is consistent with a very low fraction of initially infected cells (Supplementary Fig. S1) and, possibly, transient and moderate release of cytokines by the infected cells or single $V\gamma 9+V\delta 2+$ T cells. Taken together, these experiments do not support the hypothesis that the investigated cytokines contribute to the BTN3A-dependent effector mechanisms involved in the killing of infected epithelial cells in our co-culture system.

Limitation of *L. monocytogenes* growth in epithelial cell- $V\gamma 9+V\delta 2+$ T cell coculture requires BTN3A

In contrast to the treatment of epithelial cells with HDMAPP, their infection with *L. monocytogenes* is initially very inefficient and increases gradually via cell-to-cell spreading³⁰. We hypothesized that the small number of initially infected target cells were subject to BTN3A-dependent $V\gamma 9+V\delta 2+$ T cell cytotoxicity. In contrast, uninfected cells escape the cytotoxic activity of $V\gamma 9+V\delta 2+$ T cells, proliferate, and thus increase the cell index. To monitor the bacterial burden of cells under our coculture conditions, we performed colony-forming unit (CFU) assays in wildtype and BTN3A pan knockout cells infected with *L. monocytogenes* with or without co-cultured $V\gamma 9+V\delta 2+$ T cells (Fig. 3a). Intracellular bacterial numbers were measured every 24 h over a period of 6 days. While bacterial numbers in the wildtype situation increased immensely over time, they remained at the levels

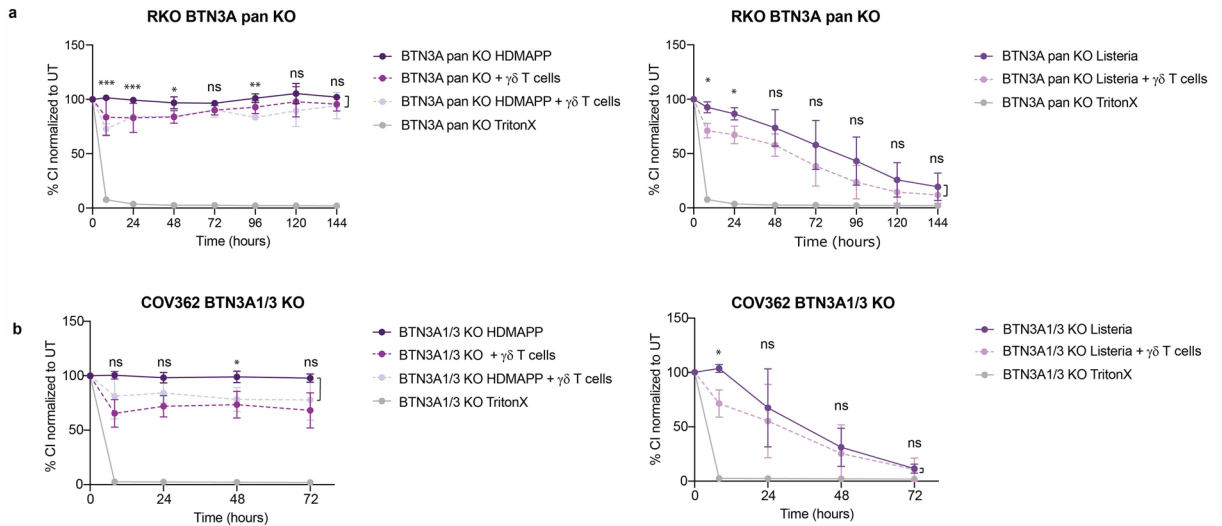


Figure 2. RTCA was performed with BTN3A pan knockout RKO (a) or BTN3A1/3 double knockout COV362 (b) cells in co-culture with $V\gamma 9+V\delta 2+$ T cells from donor 1. TritonX was used as positive control for cytotoxicity. Target cells were either treated with HDMAPP (left panel) or infected with *L. monocytogenes* (right panel). Cell Index of the treatments was normalized to the untreated sample (100%). Two-tailed unpaired Student's *t*-test was performed using GraphPad Prism comparing HDMAPP with HDMAPP + $V\gamma 9+V\delta 2+$ T cells or *L. monocytogenes* with *L. monocytogenes* + $V\gamma 9+V\delta 2+$ T cells. P-values: ns $P > 0.05$; * $P \leq 0.05$; ** $P \leq 0.01$; *** $P \leq 0.001$; **** $P \leq 0.0001$. Experiments were performed in biological triplicates.

of infection onset in the co-culture. Consistent with the results in Fig. 2, this effect was BTN3A-dependent, as the BTN3A pan knockout did not show any difference in bacterial growth when comparing cells infected with *L. monocytogenes* or co-cultured with $V\gamma 9+V\delta 2+$ T cells. Fluorescence microscopy using a GFP-labeled *L. monocytogenes* strain revealed a minor fraction of cells to be infected with *L. monocytogenes* after 24 h (Fig. 3b), consistent with the low initial MOI (Supplementary Fig. S1). The fraction of infected cells was even lower in co-culture with $V\gamma 9+V\delta 2+$ T cells. After 6 days of infection, *L. monocytogenes* spread throughout the culture, leading to enormous cell stress and cell death. In contrast, the co-cultured cells recovered from the infection, with only very low amounts of intracellular bacteria. Consistent with results in Fig. 2, co-culture of BTN3A pan knockout cells with $V\gamma 9+V\delta 2+$ T cell showed no discernible rescue effect. The epithelial cells exhibited high GFP levels, visible cellular stress, and cell death 6 days after culture alone or together with $V\gamma 9+V\delta 2+$ T cells.

The results are in line with the hypothesis that bacterial clearance occurs at rather early stages of infection by $V\gamma 9+V\delta 2+$ T cells targeting infected cells. Upon cell killing, uninfected cells start to outgrow and recover from the infection, which is not the case for the BTN3A knockout.

BTN3A3 controls *L. monocytogenes* infection during epithelial cell- $V\gamma 9+V\delta 2+$ T cell coculture in absence of BTN3A1

According to published results, $V\gamma 9+V\delta 2+$ T cell-induced killing upon HDMAPP stimulation requires the butyrophilin 3A gene BTN3A1^{20–22,25,26}. To investigate whether the same is true for the killing of cells infected with *L. monocytogenes*, we used single knockouts of either BTN3A1 or BTN3A3, validated using Simple Western™ (Supplementary Fig. S5 and S6 and Supplementary Tables S1 and S2). As the COV362 double knockout of BTN3A1 and BTN3A3 abolished the 'rescue effect', we excluded BTN3A2 as either not involved or functionally redundant for the rescue effect.

As expected, a single knockout of BTN3A1 prevented $V\gamma 9+V\delta 2+$ T cell-induced killing upon HDMAPP stimulation (Fig. 4a). In contrast, the knockout of BTN3A3 showed similar results as the wildtype cells (Fig. 4b). Surprisingly, the single knockouts of either BTN3A1 or BTN3A3 did not eliminate the ability of $V\gamma 9+V\delta 2+$ T cells to rescue the cultures infected with *L. monocytogenes*. The data support the conclusion that both BTN3A1 and BTN3A3 have the ability to mediate antigen recognition and killing of *L. monocytogenes*-infected target cells by $V\gamma 9+V\delta 2+$ T cells.

Discussion

In this study we established an in vitro system to investigate responses of $V\gamma 9+V\delta 2+$ T cells to the infection of epithelial cells with *L. monocytogenes*. With this approach, we advance previous studies mimicking *L. monocytogenes* infection using synthetic phosphoantigens or extracts from heat-killed bacteria. In line with recent literature^{6,12,14,16,31}, we observed that cells exposed to the synthetic HMB-PP analogue HDMAPP were completely eradicated by coculture with $V\gamma 9+V\delta 2+$ T cells. In contrast, *L. monocytogenes*-infected cultures were protected from complete eradication and showed a "rescue effect" of the culture resulting from the elimination of a sizeable fraction of infected cells (Fig. 5). Our infection system and the results we obtained differ from those of a recent study with monocyte-derived dendritic cells (Mo-DC)⁹. Contrasting epithelial cells, Mo-DC resisted the

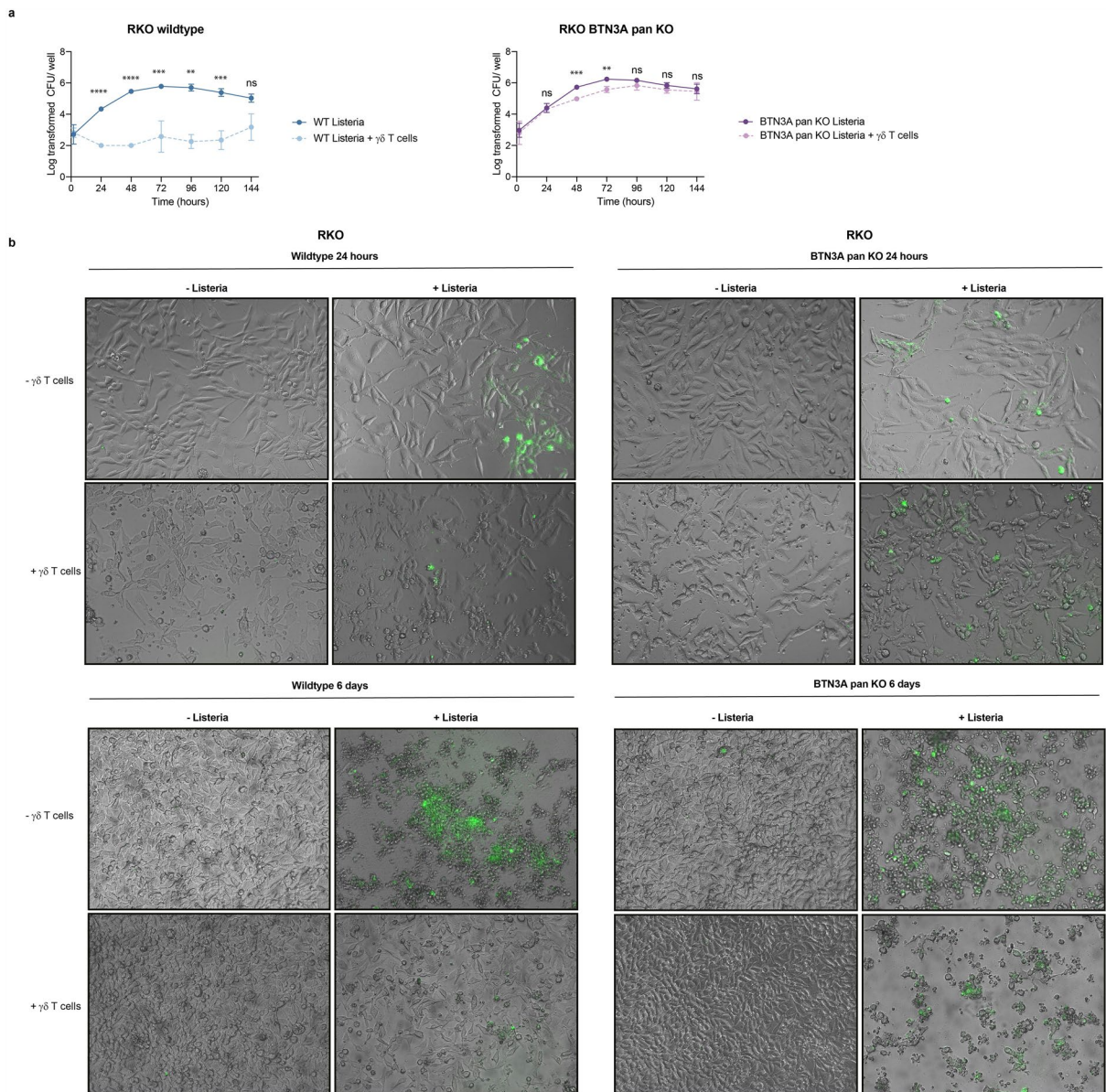


Figure 3. (a) Colony forming unit assay was performed using wildtype and BTN3A pan knockout RKO cells infected with *L. monocytogenes* alone or in co-culture with V γ 9+V δ 2+ T cells from donor 1. Cells were lysed every 24 h over a period of 144 h and intracellular bacteria were determined by CFU assay. Values represent the log transformed CFU per well of 3 biological replicates. Two-tailed unpaired Student's t-test was performed using GraphPad Prism comparing *L. monocytogenes* with *L. monocytogenes* + V γ 9+V δ 2+ T cells. P-values: ns P > 0.05; *P \leq 0.05; **P \leq 0.01; ***P \leq 0.001; ****P \leq 0.0001. (b) Wildtype and BTN3A pan knockout RKO cells were infected with GFP-labelled *L. monocytogenes* and/or co-cultured with V γ 9+V δ 2+ T cells from donor 1. Target cells and GFP-labelled bacteria were imaged via microscopy in a live system after either 24 h or 6 days.

lethal effects of *L. monocytogenes* infection. However, co-culture of infected Mo-DCs with V γ 9+V δ 2+ T cells resulted in their virtually complete elimination by $\gamma\delta$ T cell cytotoxicity. Since Mo-DC are capable of phagocytosing bacteria³² the different results most likely reflect more efficient infection of phagocytes and a limited potential of noninfected Mo-DCs to expand and rescue the culture. Besides *L. monocytogenes*, BTN3A/ $\gamma\delta$ T cell-mediated immunity was recently demonstrated in studies with the intracellular bacterium *Coxiella burnetii*, the causative agent of Q fever³³. The findings further emphasize the importance of $\gamma\delta$ T cell antigen recognition via butyrophilins and the importance of this mechanism in the combat of infections with intracellular bacteria.

Immune responses to *L. monocytogenes* are directed by a plethora of cytokines⁴. Broadly, these can be classified as proinflammatory (TNF- α), anti-inflammatory, promoting cell-autonomous immunity³⁴ (IFN- γ), or the recruitment (CCL3, CCL4, IL-16, IL-8) or growth (GM-CSF) of effector cells. Supplementary Fig. S4 shows that members of these cytokine categories are produced by $\gamma\delta$ T cells when co-cultured with infected cells and may contribute to the innate response of the infected cells. However, our findings do not support the notion that

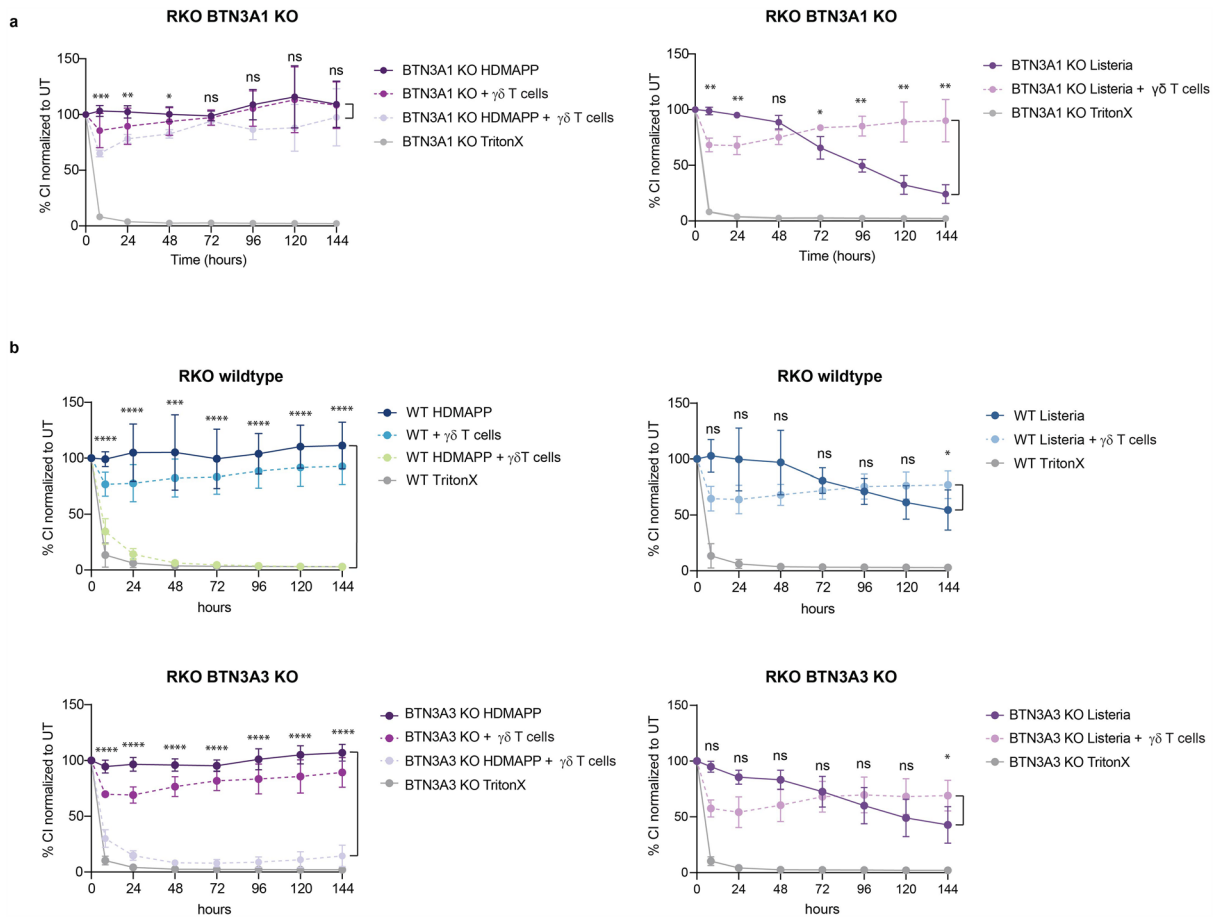


Figure 4. RTCA was performed with BTN3A1 corresponding to (a) and BTN3A3 (b) knockout RKO cells as well as wildtype RKO cells (B) either in co-culture with V γ 9+V δ 2+ T cells from donor 1 as well as controls with V γ 9+V δ 2+ T cells alone and cells treated with TritonX. Target cells were either treated with the HDMAPP (left panel) or infected with *L. monocytogenes* (right panel). Cell adherence was monitored over 144 h. Cell Index of the treatments was normalized to the untreated sample (100%). Two-tailed unpaired Student's t-test was performed using GraphPad Prism comparing either HDMAPP with HDMAPP + V γ 9+V δ 2+ T cells or *L. monocytogenes* with *L. monocytogenes* + V γ 9+V δ 2+ T cells. P-values: ns $P > 0.05$; * $P \leq 0.05$; ** $P \leq 0.01$; *** $P \leq 0.001$; **** $P \leq 0.0001$. Mean and STDEV from biological triplicates are shown.

antigen presentation by BTN3A increases the production of these cytokines. While this is an interesting negative result, it should be noted that the room for interpretation of our coculture system has clear limitations. On the one hand, the panel of investigated cytokines may lack an important contributor. More importantly, infection of the epithelial cells and their subsequent elimination from the culture is a gradual process occurring over several days. This asynchrony may obscure bursts of cytokine production by single cells in the process of infection and/or interaction with $\gamma\delta$ T cells. Time-resolved studies of single cells will be needed to examine this possibility.

Cytotoxicity against infected cells requires the activation of V γ 9+V δ 2+ T cells by antigen-activated BTN3A, as indicated by CRISPR/Cas9-mediated KO of individual, or groups of BTN3A genes in epithelial cells (Fig. 5). Cell cultures lacking all BTN3A family members, or the BTN3A1/3 genes failed to be rescued by V γ 9+V δ 2+ T cells. This rules out an independent role of BTN3A2. Both BTN3A1/3 double KO and BTN3A pan KO additionally lack the BTN2A3p, a pseudogene that is transcribed but not further processed and translated. Therefore, a role of BTN2A3p in the presentation of antigens to $\gamma\delta$ T cells is highly unlikely. Surprising in the light of recent reports, the rescue effect was observed in both single KO of BTN3A1 and BTN3A3. Contrasting the presentation of the model phosphoantigen HMB-PP, BTN3A1 and BTN3A3 appear to share the ability to present relevant *L. monocytogenes* antigens to V γ 9+V δ 2+ T cells. This finding is in line with data of Alice and colleagues⁹ showing phosphoantigen-independent V γ 9+V δ 2+ T cell activation by *L. monocytogenes*. The authors tested BTN3A dependency using the 103.2 monoclonal antibody. This antibody binds to all ectodomains of the BTN3A family members, and not exclusively to BTN3A1¹⁵. Its ability to block V γ 9+V δ 2+ T cell activation by *L. monocytogenes*-infected Mo-DC is therefore not contrary to the results we obtained with gene knockouts.

Reportedly, HMB-PP binds to the positively charged pocket of the cytoplasmic B30.2 subunit of BTN3A1, and a single amino acid change in the positively charged pocket of BTN3A3 prevents phosphoantigen reactivity¹⁹. Therefore, our finding supports the assumption that HMB-PP is not the only antigen inducing the rescue effect in our system. Similar conclusions were drawn by Alice et al.⁹ from results with mutant *L. monocytogenes* strains producing different levels of HMB-PP but eliciting similar levels of $\gamma\delta$ T cell-mediated cytotoxicity against

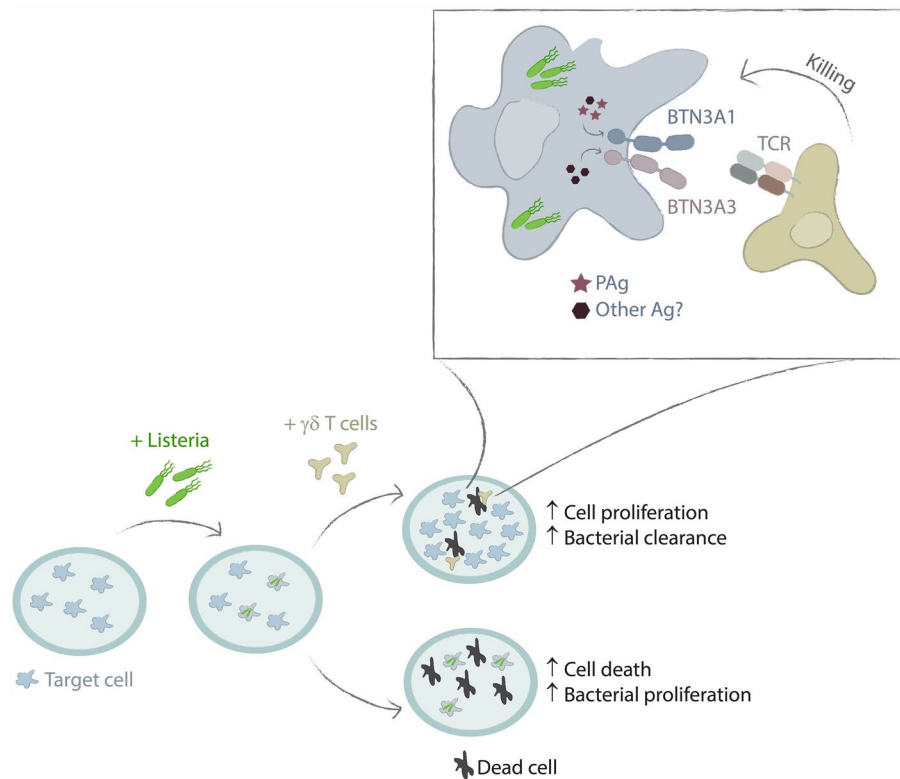


Figure 5. Illustration of *L. monocytogenes*-infected target cells with and without co-culture of V γ 9+V δ 2+ T cells. Upon infection, only a minor amount of target cells takes up *L. monocytogenes*. Over time, *L. monocytogenes* proliferates inside the target cells and spreads to the neighboring cells. This results in high amounts of bacteria as well as death of the target cells due to cell-to-cell spreading. In contrast, when co-culturing with V γ 9+V δ 2+ T cells, infected target cells will be immediately attacked by the V γ 9+V δ 2+ T cells reducing the spreading of *L. monocytogenes* to neighboring cells. Thus, infected cells are selectively killed, and uninfected cells will continue growing. For the selective killing of *L. monocytogenes*-infected target cells, expression of BTN3A1 or BTN3A3 on the target cells is essential. BTN3A1 might be activated by phosphoantigens (PAg) or an alternative antigen, that could also bind to and activate BTN3A3.

infected Mo-DCs. Overall, the results obtained in experiments with infected cells differ from those generated with heat-killed *L. monocytogenes* extracts and challenge the notion of an exclusive role for HMB-PP as the activating antigen for $\gamma\delta$ T cells^{6,11}. In this regard, *L. monocytogenes* extracts not subjected to heat treatment were reported to induce increased outgrowth of V γ 9+V δ 2+ T cells when compared to heat-treated lysates⁸. This may indicate that heat treatment reduces the potency of HMB-PP, or, alternatively, that it eliminates the activating potential of additional antigens. Moreover, Morita⁵ identified the presence of alkylamine antigens in *L. monocytogenes*^{11,35}. Similar to phosphoantigens, alkylamine antigens require cell–cell contact, but not professional antigen presentation via MHC class I, II or CD1 to stimulate V γ 9+V δ 2+ T cells. In light of these findings, our observations may indicate an unknown role of BTN3A3 in antigen presentation and the host defense against infections.

Materials and methods

Cell lines

Human colon cancer cell line RKO obtained from the American Type Culture Collection (Cat. No. CRL-2577) were maintained in RPMI 1640 GlutaMAX™ culture medium (Gibco, Cat. No. 61870-010) supplemented with 10% (v/v) heat-inactivated fetal bovine serum (FBS) (Gibco, Cat.No. 26150), 1X Minimum Essential Medium Non-Essential Amino Acids (MEM NEAA; Gibco, Cat. No. 11140-068) and 1× Sodium Pyruvate 100 mM (Gibco, Cat. No. 11360-088). Human ovarian cancer cell line COV362 obtained from European Collection of Authenticated Cell Cultures (Cat. No. 07071910) were maintained in DMEM (Sigma, Cat. No. D6429) culture medium supplemented with 10% (v/v) FBS. Lenti-X™ 293T cells line obtained from Clontech (Cat. No. 632180) were cultured in DMEM supplemented with 10% Tet System Approved FBS (Takara, Cat. No. 631106). All cell lines were cultivated at 37 °C in a humidified atmosphere with 5% CO₂.

Generation of primary V γ 9+V δ 2+ T cell batches

For the generation of human V γ 9+V δ 2+ T cell batches, buffy coats from healthy donors were purchased from the Austrian Red Cross, who always obtain samples under informed consent in accordance with relevant guidelines, regulations, and internal approvals to ensure ethics and informed consent of donors. PBMCs were isolated

from buffy coats by density-gradient centrifugation. In brief, buffy coats were diluted 1:1 in HBSS (Gibco Cat. No. 14170-088) and centrifuged in a Lymphoprep (Axis-Shield, Cat. No. 1114544) containing Leucosep tube (Greiner, Cat. No. 227290) to isolate PBMCs. After three washes with HBSS and erythrocyte lysis with ACK lysing buffer (Gibco Cat. No. A10492-01) PBMCs aliquots of 10^8 cells/ml were stored at -150°C until further use. For $\text{V}\gamma 9+\text{V}\delta 2+$ T cell expansion and isolation, PBMCs were thawed and seeded at a cell density of 5×10^6 cells/ml in RPMI 1640 GlutaMAX™ culture medium (Gibco, Cat. No. 61870-010) supplemented with 10% (v/v) heat-inactivated fetal bovine serum (FBS) (Thermo Scientific, #SH30071.03). Cells were treated over night with $0.2 \mu\text{M}$ HDMAPP*Li (Sigma-Aldrich, Cat.No. 95098) and 125 ng/ml recombinant human IL-2 (Peprotec, Cat. No. AF200-02) and were then expanded over 2 weeks with 125 ng/ml IL-2. At d14, $\text{V}\gamma 9+\text{V}\delta 2+$ T cells were isolated by negative selection using the $\gamma\delta$ T cell isolation kit from StemCell (Cat.No. 19255) according to manufacturer's protocol. For donor 1 a d1TCR antibody cocktail (Cat.No. 18309-A038) was added in addition. Quality and purity of isolated cells was estimated by flow cytometry after surface staining for CD3 (Biolegend, Cat.No. 300428), and g9 TCR (Biolegend, Cat.No. B235686) or d2 TCR staining (BD, Cat. No. 0038917) on a FACS CantoII. For both used donors we harvested 99% pure $\gamma\delta$ T cells batches, with 88% $\text{V}\gamma 9+\text{V}\delta 2+$ content in donor 1 and 91,5% $\text{V}\gamma 9+\text{V}\delta 2+$ content in donor 2.

Human primary $\text{V}\gamma 9+\text{V}\delta 2+$ T cells from different donors were thawed the day prior to the experiment in RPMI 1640 GlutaMAX™ culture medium supplemented with 10% (v/v) heat-inactivated fetal bovine serum (FBS). 10 million cells were resuspended in 10 ml medium and stimulated with 25 ng/ml recombinant human interleukin-2 (IL-2) (Peprotech #AF200-02). $\text{V}\gamma 9+\text{V}\delta 2+$ T cells were kept overnight at 37°C and 5% CO_2 .

Gene-edited cell lines

Human colon carcinoma RKO cell line was stably transduced with pRRL.SFFV-rtTA3-IRES-EcoR-PGK-Hygro and pRRL.TRE3-Cas9-P2A-BFP for tetracycline-induced Cas9 expression. To generate RKO cell lines with a gene knockout for components of the BTN3A family, the following guides were used:

```
gControl GGCAGTCGGTTCGGTTGATAT
gBTN3Apan AGAACTTCGATTCTGCGGGA
gBTN3A1 CCTGGACGTCTCCTTCTCTG
gBTN3A2 GATGCAGGATACACCCTCCC
gBTN3A3 ATAAAGTGGAGCGACACCAA
```

Guides were synthesized and cloned into pLenti-sgLib-U6-IT-EF1a-mKate2-P2A-Neo by Genscript. Plasmids were packed into lentiviral particles using the Lenti-X cell line and Lenti-X Packaging Single Shots (VSV-G) (Takara, Cat. No. 631276) according to manufacturer's protocol. After 48 h, we harvested viral particle containing supernatants, filtered through a $45 \mu\text{m}$ SFC filter and aliquoted and stored them at -80°C until further use. RKO with dox-inducible Cas9 were transduced with the BTN3A guide containing particles and were neomycin selected and tetracycline-induced for 2 weeks to generate polyclonal KO cell batches.

Double knockout clones for BTN3A1 and BTN3A3 were generated by GenScript. Briefly, Cov362 were electroporated with guide containing pSpCas9 (BB)-2A-GFP (PX458), cloned out by single cell dilution and screened for double KO by PCR. The guides used for double KO generation in Cov362 were:

```
gBTN3A1_doubleKO CCTGAGAACTACTAGATGAT
gBTN3A3_doubleKO CTGACGTCCCAGTTGTTCCC
```

Protein quantification by automated Western Blot

2 or 0.33×10^6 cells/well were seeded in a six well dish and harvest after 1 or 2 days, respectively. For harvesting, cells were washed with PBS and lysed in $200 \mu\text{l}$ RIPA lysis buffer (EMD Millipore, Cat. No. 20-188) supplemented with complete protease inhibitor cocktail (Roche, Cat. No. 11697498001) and 0.5% *n*-dodecyl- β -D-Maltoside (DDM, Avanti Polar Lipids, Cat. No. 850520P). Lysates were collected by scraping, sonicated with 35 mA for 40 s in a water bath and cleared from debris by centrifugation for 10 min at $14,000g$ at 4°C . The protein content of the cleared lysates was estimated using Pierce BCA Protein Assay Kit (Thermo Scientific, Cat. No. 216513) according to manufacturer's recommendations. BTN3A1 and BTN3A3 protein levels were analyzed using the Simple Western™ Automated Western Blot System from Bio-Techne. In brief, $1 \mu\text{g}/\mu\text{l}$ of cleared cell lysates was loaded onto a ProteinSimple 12–230 kDa 25-Capillary Cartridge (Bio-technique, Cat. No. SM W004). Size separation, immunoprobings and chemiluminescence-based detection was fully automated. Samples were run side-by-side on separate capillaries of the same cartridge. For adequate comparison of different samples, internal size standards and housekeeping gene GAPDH were detected in addition to BTN3A1 or BTN3A3 in each capillary. Capillaries were incubated for 60 min with antibodies specific to GAPDH (Cell Signaling, #2118, 1:1000), BTN3A1 (Novusbiologicals, Cat. No. NBP1-90750 1:200) or BTN3A3 (Novusbiologicals, Cat. No. 15896-1-AP, 1:200). After further 30-min incubation with the ProteinSimple Anti-Rabbit Detection Module (Chemiluminescence, Bio-Techne, Cat. No. DM-001), signals were developed using chemiluminescence. Chromatograms from immunoreactive and chemiluminescent signals are automatically detected and depicted in a Western Blot-like visualization. For quantitative evaluation of the KO efficiency, area under the curve (AUC) is calculated for the probed proteins from chromatographic peaks using a Gaussian bell curve fit.

Bacterial preparation and infection

L. monocytogenes strain LO28 was thawed and grown overnight a day prior to the experiment in Brain–Heart Infusion (BHI) medium at 37°C with continuously shaking. After reaching stationary phase, optical density at 600 nm (OD_{600}) was measured and desired multiplicity of infection (MOI) ratio could be determined. An MOI of 100 was used for experiments with RKO cells, while an MOI of 10 was used for experiments performed with COV362 cells. Bacteria were centrifuged 3 min at $10,000g$, washed twice with phosphate-buffered saline (PBS)

and resuspended in RPMI 1640 medium supplemented with 10% (v/v) FBS. Bacteria were added dropwise on top of the target cells. After 1 h of infection, the medium was exchanged to a medium containing a high concentration (50 µg/ml) Gentamycin (MP Biomedicals, Santa Ana, US). After a total of 2 h of *L. monocytogenes* infection, the medium was changed again, and cells kept in a low concentration of gentamycin (10 µg/ml).

Real time cell analysis (RTCA)

The RTCA is a method that allows continuous monitoring of adherent cells. The wells of the associated 96-well E-plates are coated with golden electrodes which monitor the impedance of adherent cells which is further translated into a unit called the Cell Index (CI). The wells of the E-plate were coated beforehand with 50 µl fibronectin (1 mg/ml) a day prior the experiment, washed twice with PBS and kept overnight at 4 °C. The day of the experiment, the background was measured with 50 µl of clean cell medium (3 swipes every 1 min). Target cells were counted and seeded into the wells of the E-plate. The plate was incubated for 30 min at room temperature before being placed back into the monitoring unit which was always kept at 37 °C and 5% CO₂. The interval of the swipes was set to every 5 min. Target cells were let to adhere to the wells for 7 h. Before the treatment, Cell Index of all wells was normalized to 1. At this point, cells were infected with *L. monocytogenes* at the desired MOI (see Section Bacterial Preparation and Infection). After one hour of infection, medium was exchanged to medium containing a high concentration of Gentamycin (50 µg/ml). After another hour, cells were either left untreated or treated with 1% (v/v) TritonX-100, HDMAPP (1 mg/ml) and/or human primary Vγ9+Vδ2+ T cells with an Effector to Target ratio of 2.5:1. In all conditions, medium was supplemented with a low concentration of Gentamycin as well as recombinant human IL-2 (125 ng/ml). The seeding density of the RKO cells line was 2 × 10⁴ cells/well, whereas we used 1.6 × 10⁴ cells/well for the COV362 cell line. All treatments were performed in technical triplicates.

MSD cytokine analysis

RKO cells were seeded with a density of 2 × 10⁴ cells/well in a 96-well plate. After 7 h, cells were infected with *L. monocytogenes* with an MOI of 100. After 1 h, medium was exchanged to medium containing 50 µg/ml Gentamycin. One hour later, Vγ9+Vδ2+ T cells were added to the target cells. Cells were kept in medium supplemented with 10 µg/ml Gentamycin and 125 ng/ml recombinant human IL-2 at 37 °C and 5% CO₂. After either 48 or 72 h, supernatant was harvested. Supernatant was centrifuged 5 min at 10,000g, transferred to a fresh tube and kept at -80 °C for 2 weeks to remove any viable bacteria. Afterwards, supernatant was analyzed using the Meso Scale Discovery (MSD) Multi Spot Assay system and the V-PLEX Human Cytokine 30-Plex Kit (MSD, Cat. No. K15054D2) according to manufacturer's instruction.

Colony forming unit assay (CFU)

RKO cells were seeded with a density of 2 × 10⁴ cells/well in a 96-well plate. After 7 h, cells were infected with *L. monocytogenes* with an MOI of 100. After 1 h, medium was exchanged to medium containing 50 µg/ml Gentamycin. One hour later, Vγ9+Vδ2+ T cells were added to the target cells. Cells were kept in medium supplemented with 10 µg/ml Gentamycin and 125 ng/ml recombinant human IL-2 at 37 °C and 5% CO₂. At the given timepoints, cells were washed twice with PBS and lysed with sterile water. The lysate was further serially diluted in PBS and dilutions were plated on BHI plates. Plates were kept overnight at 37 °C and 5% CO₂ and colonies were counted the next day.

Fluorescent microscopy

RKO cells were grown on a chambered eight-well microscopy slide (Ibidi, Cat. No. 80807) and kept in phenol red free medium. Target cells were infected for 1 h with the fluorescent GFP-labelled *L. monocytogenes* strain EGD-cGFP obtained from Olivier Disson, the Institut Pasteur, Unité des Interactions Bactéries-Cellules, Paris F-75015, France³⁶. Afterwards, medium was exchanged to medium containing 50 µg/ml Gentamycin. After another hour, Vγ9+Vδ2+ T cells were added with medium supplemented with low concentration of gentamycin (10 µg/ml) as well as 125 ng/ml recombinant IL-2. Samples were analyzed with a Zeiss Axio Observer Z1 microscope using a 20× objective. Images were further processed with ImageJ software.

Data availability

All data generated or analyzed during this study are included in this published article (and its Supplementary Information files).

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Author contributions

K.F., V.S. and T.D. designed the study and wrote the manuscript. K.F., M.B. and V.S. performed experiments and analysed the data.

Competing interests

The authors declare no competing interests.

Additional information

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