



Published in final edited form as:

Mucosal Immunol. 2015 November ; 8(6): 1349–1359. doi:10.1038/mi.2015.24.

Immunization with Ty21a live oral typhoid vaccine elicits cross-reactive multifunctional CD8⁺ T cell responses against *Salmonella enterica* serovar Typhi, *S. Paratyphi A* and *S. Paratyphi B* in humans

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Abstract

Previously we have extensively characterized *Salmonella enterica* serovar Typhi (*S. Typhi*)-specific cell-mediated immune responses (CMI) in volunteers orally immunized with the licensed Ty21a typhoid vaccine. In this study we measured *Salmonella*-specific multifunctional (MF) CD8⁺ T cell responses to further investigate whether Ty21a elicits cross reactive CMI against *S. Paratyphi A* and *S. Paratyphi B*, which also cause enteric fever. Ty21a elicited cross-reactive CMI against all three *Salmonella* serotypes were predominantly observed in CD8⁺ T effector/memory (T_{EM}) and, to a lesser extent, in CD8⁺CD45RA⁺ T_{EM} (T_{EMRA}) subsets. These CD8⁺ T cell responses were largely mediated by MF cells co producing IFN- γ , MIP-1 β and expressing CD107a with or without TNF- α . Significant proportions of *Salmonella*-specific MF cells expressed the gut homing molecule integrin $\alpha_4\beta_7$. In most subjects similar MF responses were observed to *S. Typhi* and *S. Paratyphi B*, but not to *S. Paratyphi A*. These results suggest that Ty21a elicits MF CMI against *Salmonella* which could be critical in clearing the infection. Moreover, because *S. Paratyphi A* is a major public concern and Ty21a was shown in field studies not to afford cross-protection to *S. Paratyphi A*, these results will be important in developing a *S. Typhi/S. Paratyphi A* bivalent vaccine against enteric fevers.

Keywords

Ty21a; Cross-reactive CD8 T cells; Multifunctional; *Salmonella*; Human

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Disclosures. The authors declare no conflict of interest. The content of this article is solely the responsibility of the authors and does not necessarily represent the official views of the National Institute of Allergy and Infectious Diseases or the National Institutes of Health.

Introduction

Typhoid fever caused by *Salmonella* enterica serovar Typhi (*S. Typhi*), is responsible for an estimated 21.7 million cases and 200,000 deaths per year worldwide.^{1,2} Other significant causative agents of enteric fevers are *S. Paratyphi A* and *S. Paratyphi B*, and rarely *S. Paratyphi C*.³ Recent reports indicate that the incidence of paratyphoid A fever is on the rise in areas of endemicity (e.g., South and Southeast Asia, China) and among travelers returning from those areas.^{1,4-7} The emergence of multiple antibiotic-resistant *Salmonella* strains has further increased the health risks posed by these infections.⁸

To prevent typhoid fever, three licensed vaccines are available, i.e., live attenuated oral vaccine Ty21a (Ty21a), parenteral polysaccharide Vi-vaccine (Vi vaccine) and Vi conjugated vaccine. In contrast, no vaccines are available against paratyphoid fevers. Although a high degree of homology at the DNA level exists among *S. Typhi*, *S. Paratyphi A*, and *S. Paratyphi B*, a critical virulence factor, the *S. Typhi* Vi polysaccharide, is not expressed either by *S. Paratyphi A* or *S. Paratyphi B*.⁹ Therefore, the parenteral Vi vaccines are not expected to provide cross-protection against paratyphoid A and B fevers. The possibility that the live attenuated Ty21a confers cross-protection against *S. Paratyphi A* and/or *S. Paratyphi B* has been studied in several field studies.¹⁰⁻¹² Those studies indicate that Ty21a does not protect against *S. Paratyphi A*, while it does confer a moderate degree of protection against *S. Paratyphi B* disease.¹³

Because of the recent increased incidence of enteric fever caused by *S. Paratyphi A*, the need for an effective vaccine against paratyphoid A fever has been emphasized.¹⁴ However, the need for an effective vaccine against *S. Paratyphi A*, as well as more effective vaccines to typhoid fever, requires a better understanding of the immunological basis of the cross-reactive and cross-protective responses induced by Ty21a. This is complicated by the fact that *S. Typhi* is a human host-restricted pathogen and animal models do not faithfully recapitulate human disease. Nevertheless, the *S. Typhimurium* “typhoid” mouse model has led to important insights into the role that various innate and adaptive effector mechanisms might play in protection from *Salmonella* infection. For example, resistance to virulent challenge with *S. typhimurium* by immunized mice, requires production of interferon (IFN)- γ and tumor necrosis factor (TNF)- α by both CD4+ and CD8+ T cells.¹⁵⁻¹⁸

In humans, humoral, and most importantly cell-mediated immune (CMI) responses that are induced following vaccination of healthy volunteers with Ty21a and other live oral candidate vaccine strains (i.e., CVD 908, CVD 908-*htrA*, CVD 909, MZH09) have been studied extensively by us and others.¹⁹⁻³⁰ Typically we observed that following immunization with live oral *S. Typhi* vaccines, both CD4+ and CD8+ T cell responses, including cytotoxic T cells (CTL), were observed depending on the nature of the stimulant used in *in-vitro* or *ex-vivo* experiments.^{23, 24, 27, 29-35} While typically CD4+T cells responses were more pronounced to soluble antigens (e.g., flagella), CD8+T cells were the predominant responders against *S. Typhi*-infected targets.^{24, 27, 30, 32}

Depending on the expression of defined markers, T memory (T_M) cell subsets have been broadly divided into T central memory (T_{CM} : CD45RA-CD62L+), T effector/memory

(T_{EM}; CD45RA-CD62L), and RA+T_{EM} (T_{EMRA}; CD45RA-62L-).^{36,37} Of note, T_{EMRA}, considered to be “terminally differentiated T_{EM} cells”,^{36, 37} were found to express high levels of perforin and granzyme and have been implicated in protection against viral (e.g., HIV, Cytomegalovirus and Epstein-Barr virus) and bacterial (*M. tuberculosis*) infections.³⁸⁻⁴² Regarding immunity to *S. Typhi*, we have shown that oral immunization with attenuated typhoid vaccines elicits *S. Typhi*-specific CD8+ T cell responses, mostly involving the T_{EM} and T_{EMRA} subsets, although lower magnitude responses were also observed in the CD8+ T_{CM} subset.^{22, 25, 26, 30} Of interest, we showed that a significant portion of *S. Typhi*-specific T cells co-expressed the gut homing molecule integrin $\alpha_4\beta_7$, suggesting their potential to migrate to the primary site of infection.^{25, 30, 43} Taken together, these observations strongly suggest that live oral typhoid vaccines elicit CD8+ CTL and other CMI responses likely to be the primary mediator(s) of protective immunity, both in clearing acute infection, as well as in providing long-term protection against *S. Typhi*.^{32, 44} Recent studies further showed that antigen specific multifunctional (MF) T cells (cells producing two or more cytokines and/or expressing CD107a (a marker of cytotoxic activity),⁴⁵ induced in response to various vaccines,⁴⁶ including Ty21a,^{22, 26} might play a key role in long-term protective immunity.

However, in spite of the considerable progress in uncovering *S. Typhi* specific responses, very limited information is available on the cross reactive responses induced against *S. Paratyphi A* or *S. Paratyphi B* by live oral typhoid vaccines. Recently, we and others have described cross-reactive humoral responses induced by Ty21a against *S. Typhi*, *S. Paratyphi A* and *B*.⁴⁷⁻⁴⁹ Humoral responses induced following immunization with Ty21a were directed predominantly against *S. Typhi*; however, cross-reactive responses were also recorded against *S. Paratyphi A* and *B*. We further observed the induction of cross-reactive functional vaccine induced antibodies which were, nevertheless, not sufficient to clear *Salmonella* infections once they become intracellular.^{21, 48} Taken together, these observations support the notion that in addition to humoral immunity, CMI might be critical for the efficient control of *S. Typhi*,^{22-27, 29, 30, 32, 44} as well as for *S. Paratyphi A* or *S. Paratyphi B* infections.^{47, 48}

To address the gaps in knowledge regarding the mechanisms of cross-protective immunity among enteric fevers, we compared the ability of Ty21a to induce cross-reactive CMI responses among *S. Typhi*, *S. Paratyphi A* and *B*. We observed, for the first time, that the predominant cross-reactive *Salmonella* specific responses were observed in the CD8+ T_{EM} subset, whilst lower magnitude responses were also observed in CD8+ T_{EMRA} cells. Moreover, we identified the dominant subsets of MF cells that mediate cross-reactive *Salmonella*-specific responses and show that *Salmonella*-specific CD8+ T_M populations are composed of cells that express, or not, the gut-homing molecule integrin $\alpha_4\beta_7$. Finally, of importance, we observed that Ty21a-elicited CMI responses against *S. Typhi* were found to be similar to those observed against *S. Paratyphi B*-, but not *S. Paratyphi A*-infected targets. These observations provide a plausible immunological explanation for the observations of cross-protection between typhoid and paratyphoid B fever in Ty21a vaccinated subjects in field trials.

Results

The PBMC samples used in this study were collected from volunteers before (day 0) and after (days 42/84) immunization with Ty21a as described in Methods. Routine Complete Blood Counts (CBC) performed in these blood specimens were used to estimate the absolute numbers of lymphocytes and CD3+CD8+ cells. We observed that the percentages and absolute lymphocyte counts were similar (i.e., not statistically different, $p>0.3$) when pre-vaccination (day 0) and post-vaccination (days 42/84) were compared (**Fig. S1**).

Furthermore, we measured the percentages of CD3+CD8+ T cells in PBMC by flow-cytometry and converted these percentages into approximate absolute counts of CD8+T cells using available absolute lymphocyte counts from CBC analyses. Again, no statistically significant differences ($p>0.15$) were observed in the calculated absolute counts for CD8+T cells among specimens collected at days 0, 42 or 84 (**Fig. S1**).

To measure *Salmonella*-specific responses, PBMC were stimulated *ex-vivo* with *S. Typhi*-, *S. Paratyphi A*- and *B*-infected autologous EBV-B cells (**Fig. S2**) as described in Methods. Activated CD8+ T cells (i.e., CD8+CD69+ cells) produced IFN- γ (CD69+INF γ +) and/or expressed CD107a (**Fig. S3**). Activated cells resided predominantly in the CD62L- T_M-sub populations, i.e., T_{EM} and T_{EMRA} (**Fig. S3**). A similar phenomenon was also observed in TNF- α -producing cells (data not shown). Based on these observations, subsequent analyses were focused in the CD8+ T_{EM} and T_{EMRA} T cell subsets.

Evaluation of *Salmonella*-specific multifunctional CD8+T cells

In response to *S. Typhi*-specific stimulation, activated effector CD8+ T cells from Ty21a vaccinees are capable of producing single cytokines or expressing CD107a only (single positives) or concomitantly producing two or more cytokines and/or expressing CD107a (Multifunctional, MF)^{22, 26} (**Fig.S4**). We observed that Ty21a immunization elicited increases in CD8+ T cells that produce IFN- γ and/or express CD107a following stimulation with *S. Typhi*-, as well as with *S. Paratyphi A*- or *B*-infected targets (**Fig. 1**). A significantly higher proportion of these *Salmonella*-specific IFN- γ producing cells were MF when compared to single positive IFN- γ + cells in both T_{EM} (**Fig. 1A**) and T_{EMRA} (**Fig. 1B**) subsets. Similarly, significantly higher percentages of *Salmonella*-specific MF cells expressing CD107a were also observed in CD8+ T_{EM} subsets (**Fig. 1C**). However, in CD8+ T_{EMRA} subsets, significant increases in *Salmonella*-specific MF CD107a responses were only observed after stimulation with *S. Typhi*-infected targets ($p<0.01$), while a trend was observed with *S. Paratyphi B* ($p=0.08$). No dominance of MF CD107a responses in the T_{EMRA} subset was observed for *S. Paratyphi A* ($p=0.23$) (**Fig. 1D**). The above described post-vaccination increases in *Salmonella*-specific MF CD8+ T_{EM} and CD8+T_{EMRA} T_M subsets for each individual volunteer are shown in **Fig. S5**.

Characterization of *Salmonella*-specific multifunctional CD8+ T_{EM} cells

As described above, results indicated that vaccination with Ty21a elicits cross-reactive, predominantly MF CD8+T_{EM} and T_{EMRA} CMI responses against *S. Typhi*-, *S. Paratyphi A*- and *S. Paratyphi B*-infected targets. To further characterize these MF responses, we first categorized these MF cells into double (2+), triple (3+) or quadruple (4+) positives subsets,

based on whether they produce IFN- γ , TNF- α and/or IL-2, and/or express CD107a. Results showed that among CD8+ T_{EM} cells the percentages of *Salmonella*-specific MF cells followed the hierarchy 2+=4+>3+ (Fig. S6 A,B,C). In contrast, among CD8+T_{EMRA} cells, the hierarchy of *Salmonella*-specific MF cells was 2+>3+>4+ (Fig. S6 D,E,F). We next evaluated whether unique MF profiles (e.g., production of particular cytokines and/or expression of CD107a combinations) were elicited by Ty21a against *S. Typhi*-, *S. Paratyphi A*- and *S. Paratyphi B*-infected targets. Of all possible combinations (16 for the 4 parameters evaluated), we focused our studies on the 5 dominant MF subsets in CD8+T_{EM} and T_{EMRA}, all showing net increases of >0.05% positive cells (Figs. 2 and 3). Of note, when combined, these 5 selected “high frequency” MF subsets typically represented >80% of all MF cells within both, the T_{EM} and T_{EMRA} T_M subsets. In CD8+T_{EM} subsets, post-vaccination increases showed a dominance of *S. Typhi*-specific IFN- γ +CD107a+TNF- α -IL-2 cells over the next 4 most frequent *S. Typhi*-specific MF subsets, i.e., IFN- γ +CD107a-TNF- α +IL-2 (p<0.01), IFN- γ +CD107a+TNF- α +IL-2 (p=0.07), IFN- γ +CD107a+TNF- α +IL-2 (p=0.09) and IFN- γ +CD107a+TNF- α +IL-2+ (p=0.02) (Fig. 2A). Moreover, following Ty21a immunization, the induction of *S. Typhi* specific CD8+ T_{EM} IFN- γ +CD107a+TNF- α -IL-2 cells (0.46 \pm 0.18), was significantly higher than those specific to *S. Paratyphi A* (0.06 \pm 0.03, p=0.01) or *S. Paratyphi B* (0.13 \pm 0.06, p=0.04) (Fig. 2A). Of importance, the percentages of subjects (n=16) who were considered responders for IFN- γ +CD107a+TNF- α -IL-2-CD8+ T_{EM} specific to *S. Typhi* (56.3%) were similar to those responding to *S. Paratyphi B* (43.8%, p=0.5) and both were significantly higher than the 12.5% of volunteers responding to *S. Paratyphi A* (p<0.01 and p<0.05 as compared to *S. Typhi* and *S. Paratyphi B*, respectively) (Fig. 2B).

Characterization of *Salmonella*-specific multifunctional CD8+ T_{EMRA} cells

A similar analysis to the one described above for T_{EM} cells was used to characterize the Ty21a-induced cross-reactive MF responses in CD8+ T_{EMRA} subsets (Fig. 3). The specific responses observed in CD8+ T_{EMRA} subsets were generally of lower magnitude; however, their MF profiles were similar to those observed in T_{EM} subsets. The post-vaccination increase of *S. Typhi*-specific CD107a+IFN- γ +TNF- α -IL-2-cells were moderately higher than the other subsets among CD8+T_{EMRA} MF cells, although unlike CD8+T_{EM} cells, these differences did not reach statistical significance (Fig. 3).

Comparison of MF cells between CD8+T_{EM} and CD8+T_{EMRA} subsets

We have described above that Ty21a immunization elicited increases in *Salmonella*-specific responses in CD8+T cells in both T_{EM} and T_{EMRA} subsets (Figs. 1-3). We next compared those responses induced in these two T_M subsets of CD8+ T cells (Fig. 4, S5). These comparative analyses showed that IFN- γ + and CD107a+ MF responses specific to *S. Typhi* (Fig. 4A) and *S. Paratyphi B* (Fig. 4C), but not those to *S. Paratyphi A* (Fig. 4B), were significantly higher in T_{EM} than the corresponding increases in CD8+ T_{EMRA} subsets. Interestingly, increased percentages of single CD107a expressing cells in T_{EM} over T_{EMRA} were observed in *S. Paratyphi B*-, but not *S. Typhi*- and *S. Paratyphi A*-infected targets (Fig. 4 A,B,C). As described above, *Salmonella*-specific MF cells can be categorized according to their “functional” characteristics into 2+, 3+ and 4+ subsets. A comparative analysis showed

that the 4+ MF cells specific to all three *Salmonella* strains were elicited at a significantly ($p < 0.01$) higher percentage in CD8+T_{EM} as compared to T_{EMRA} subsets (**Fig. S6**). In contrast, 2+ MF CD8+T_{EM} cells specific to *S. Typhi* ($p = 0.04$) and *S. Paratyphi B* ($p = 0.08$), but not *S. Paratyphi A* ($p = 0.2$) were induced in lower percentages than the corresponding 2+ MF CD8+T_{EMRA} cells (**Fig. S6**). Moreover, post vaccination increases observed in *S. Typhi*- ($p = 0.04$) and *S. Paratyphi B*- ($p = 0.02$) specific IFN- γ +CD107a+TNF- α -IL-2- MF cells were higher in T_{EM} compared to T_{EMRA} CD8+ subsets (**Fig. 4D**). Although similar effects were also observed with *S. Typhi* and *S. Paratyphi B*- specific IFN- γ +CD107a+TNF- α +IL-2- CD8+ T_{EM} MF cells, these increases did not reach statistical significance ($p < 0.1$) (**Fig. 4E**). Of importance, while induction of *S. Typhi*- or *S. Paratyphi B*-specific MF cells were higher in T_{EM} compared to the corresponding responses in CD8+T_{EMRA} subsets, no such differences were observed with *S. Paratyphi A* (**Figs. 4B,D, E and S6**). Taken together, these comparative analyses between CD8+ T_{EM} and T_{EMRA} subsets revealed that the response patterns elicited to *S. Typhi* were remarkably similar to those to *S. Paratyphi B*, but different than those to *S. Paratyphi A*.

Cross-reactive *Salmonella*-specific MIP-1 β and IL-17 responses

MIP-1 β and IL-17 are two critical chemokines/cytokines which have been recently implicated in protection against infections.^{50, 51} Therefore, we evaluated the induction of MIP-1 β and IL-17 production in response to *Salmonella*-infected targets by PBMC obtained from Ty21a vaccinees ($n = 8$). To this end, we used an optimized 14-color flow cytometry panel (described in Methods) that included additional mAbs against MIP-1 β and IL-17. Similar to the results described above regarding induction of IFN- γ or CD107a+ MF (**Fig. 1**), Ty21a immunization elicited *Salmonella*-specific, predominantly MF, MIP-1 β + cells in CD8+ T_{EM} (**Fig. 5A**) and T_{EMRA} (**Fig. 5B**) subsets. Results from individual volunteers are shown in **Fig. S7**.

To further our understanding of the MF capabilities of MIP-1 β + cells we evaluated their ability to concomitantly produce other cytokines/chemokines and/or express CD107a. Ty21a elicited MIP-1 β + specific MF cells consisted of 6 dominant MF subsets, which were identified as those exhibiting 2+ (MIP-1 β +IFN- γ +CD107a-TNF- α -IL-2-IL-17- and MIP-1 β +IFN- γ -CD107a+TNF- α -IL-2-IL-17-), 3+ (MIP-1 β +IFN- γ +CD107a+TNF- α -IL-2-IL-17-), 4+ (MIP-1 β +IFN- γ +CD107a+TNF- α +IL-2-IL-17- and MIP-1 β +IFN- γ +CD107a+TNF- α -IL-2+IL-17-) or 5+ (MIP-1 β +IFN- γ +CD107a+TNF- α +IL-2+-IL-17-) functional responses (data not shown). Interestingly, the common characteristics among these different CD8+T_{EM} MIP-1 β + MF subsets was that, besides MIP-1 β , all of them concomitantly produced/expressed one or both of the two key markers of CTL, i.e., IFN- γ and/or CD107a. Virtually identical profiles of MIP-1 β + MF cells were induced following stimulation with *S. Typhi*- or *S. Paratyphi B* infected targets, both in CD8+T_{EM} and in CD8+ T_{EMRA} subsets. However, as described above (**Fig. 2**), there was a trend (no statistically significant) towards lower magnitude responses to *S. Paratyphi A*-infected targets (data not shown). Of note, overall, the magnitude of all *Salmonella*-specific MF subsets in CD8+T_{EM} cells responses were higher than those observed in T_{EMRA} cells (data not shown).

We also included a mAb against IL-17 in our flow cytometry panel to evaluate its role in Ty21a elicited cross-reactive immunity. Post-vaccination increases in total IL-17 producing cells above 0.1% in CD8+T_{EM} or T_{EMRA} subsets following stimulation with *Salmonella*-infected targets were observed in 3 out of 8 volunteers (37.5%) (data not shown). However, because of the low magnitude of IL-17 responses, these data was not deemed adequate for detailed analyses for MF properties of IL-17 producing cells.

Characterization of the gut/homing potential of *Salmonella*-specific MF CD8+T_{EM} cells

Mucosal immunity elicited in the gut microenvironment following immunization with Ty21a is thought to be an important component of the protective immune response against enteric fevers.^{30, 43, 44, 52} Specific effector T cells with the potential to migrate to the gut mucosa can be measured by evaluating the expression of integrin $\alpha_4\beta_7$. Thus, we examined integrin $\alpha_4\beta_7$ expression by *Salmonella*-specific IFN- γ +, CD107a+, and MIP-1 β + single and MF cells in PBMC from 12 volunteers immunized with Ty21a. We focused our studies on the gut homing patterns of cells in the CD8+ T_{EM} subset since this was found to be the dominant subset of *Salmonella*-specific CD8+ T cells responding to Ty21a immunization. We observed that Ty21a immunization elicited increases in both single and MF *Salmonella*-specific CD8+ T_{EM} cells expressing, or not integrin $\alpha_4\beta_7$ (**Fig. 6**). Integrin $\alpha_4\beta_7$ co-expressing IFN- γ + (**Fig. 6A**) and CD107a+ (**Fig. 6B**) CD8+ T_{EM} MF cells elicited by Ty21a immunization were equally responsive to *S. Typhi*-, *S. Paratyphi A*, or *S. Paratyphi B* infected targets. In contrast, integrin $\alpha_4\beta_7$ negative IFN- γ + MF cells specific to *S. Typhi* (p<0.05) or *S. Paratyphi B* (p=0.12) showed higher post-vaccination increases compared to those specific to *S. Paratyphi A* (**Fig. 6A**). Similarly, although the magnitude of integrin $\alpha_4\beta_7$ CD107a MF cells specific to *S. Typhi* (p=0.26) or *S. Paratyphi B* (p=0.14) showed higher post-vaccination increases than those specific to *S. Paratyphi A*, these differences did not reach statistical significance (**Fig. 6B**). Of note, trends were also observed for integrin $\alpha_4\beta_7$ negative IFN- γ + MF cells specific to *S. Typhi* (p=0.12) and *S. Paratyphi B* (p=0.2), but not to *S. Paratyphi A* (p=0.8), to exhibit higher post-vaccination increases compared to the corresponding integrin $\alpha_4\beta_7$ + subsets (**Fig. 6A**). On the other hand, in response to all three *Salmonella*-infected targets, integrin $\alpha_4\beta_7$ - cells co-expressing CD107a, showed an higher post-vaccination increase compared to corresponding cells expressing integrin $\alpha_4\beta_7$ (**Fig. 6B**). Specific MIP 1 β responses were also observed in integrin $\alpha_4\beta_7$ negative and positive cells (data not shown).

We then investigated integrin $\alpha_4\beta_7$ expression by the dominant *Salmonella*-specific CD8+T_{EM}MF subsets described in **Fig. 2** (e.g., IFN- γ +CD107a+TNF- α -IL-2 and INF- γ +CD107a+TNF- α +IL-2-). We observed that although a significant proportion of those also expressed integrin $\alpha_4\beta_7$, most cells were $\alpha_4\beta_7$ negative (data not shown). This dominance of integrin $\alpha_4\beta_7$ negative MF cells was not observed in MIP-1 β + subsets.

Discussion

Ty21a and other attenuated *S. Typhi* oral vaccine strains elicit a wide array of CMI responses in immunized volunteers^{23,26, 32, 33, 44, 53} including the induction of *S. Typhi*-specific multifunctional CD8+ T cells.^{22, 26} In this study we investigated whether Ty21a

immunization elicits cross-reactive CMI responses against two closely related *Salmonella enterica* serovars, i.e., *S. Paratyphi A* and *S. Paratyphi B*. In addition, by comparing the CD8⁺ T cell responses to these three *Salmonella* serovars following Ty21a immunization we explored whether defined effector CMI responses might help explain field observations showing that Ty21a provides significant cross-protection against *S. Paratyphi B*, but not against *S. Paratyphi A*.¹³ We used PBMC samples collected from healthy volunteers before (day 0) and after (day 42 and/or day 84) immunization with the live oral typhoid vaccine Ty21a. Measurements of the absolute numbers of lymphocytes and CD3⁺CD8⁺ cells based on CBC counts and the proportions of these cells obtained by flow cytometry revealed that immunization with Ty21a did not significantly affect the levels of these cells in circulation. Thus, it is unlikely that the observed post-vaccination increases in the percentages of *Salmonella*-specific CD8⁺ T cell subsets have been influenced by fluctuations in absolute cell counts following vaccination.

We have recently reported that healthy subjects who have neither a previous history of exposure to *S. Typhi*, including vaccination, nor have travelled to endemic areas, have variable background immune responses to this organism.⁵⁴ These background responses are thought to be the result of the presence of cross-reactive T cells acquired during previous infections with other gram negative enteric pathogens or by natural exposure to other gram negative organisms which form part of the normal gut microbiota. Similar pre-vaccination responses were observed in the present studies. Thus, as in previous studies, we determined Ty21a elicited specific “recall” responses by subtracting the background responses before immunization in individual subjects from each post-vaccination time point (days 42/84).^{24, 27, 30, 53}

Post-vaccination increases in specific CD8⁺ T cell responses were observed against all three *Salmonella*-infected targets (i.e., *S. Typhi*, *S. Paratyphi A* or *S. Paratyphi B*), predominantly in the T_{EM} and T_{EMRA} subsets. In contrast, very low *Salmonella*-specific responses were observed in T_{CM} and, as expected, almost no responses in T naïve cells. These and previous studies^{22, 25, 26, 30} provided the rationale for focusing most our current studies on multifunctional T_{EM} and T_{EMRA} CD8⁺ T subsets.

CD8⁺ T cells mediate effector functions by producing various cytokines (e.g., IFN- γ , TNF- α , IL-2, IL-17), chemokines (e.g., MIP-1 β) or by releasing perforin and/or granzymes (indirectly measured by the expression of CD107a).^{45, 55} At the single cell level, T cells are capable of producing single cytokines/chemokines or simultaneously producing 2 or more cytokines/chemokines and/or expressing CD107a. The latter have been termed multifunctional (MF) cells. It has been shown that these MF T cells produce higher levels of individual cytokines, exhibit enhanced function and are more likely to correlate with protection from disease when compared to single cytokine producing cells.^{56, 57} In fact, induction of MF T cells at a higher magnitudes than single cytokine secreting cells have also been shown in other disease models, i.e., HIV, CMV, vaccinia and EBV infections,⁵⁸ including the evaluation of candidate vaccines against *M. tuberculosis*^{46, 59} and Ebola virus in humans.^{60, 61}

MF cells that produce IFN- γ together with other critical cytokines (e.g., TNF- α , IL-2) and/or express CD107a can enhance the killing of intracellular bacteria more efficiently than single cytokine producing T cells.^{58, 62} Moreover, specific MF responses by T_{EM}, as well as by T_{EMRA} CD8⁺ T cells are thought to be associated with protection against various viral and bacterial infections.^{63, 64} Therefore, the quality of T cell responses, as measured by their MF capabilities, have the potential to provide a more revealing assessment of vaccine induced immune responses than single parameter functional measurements (e.g., only IFN- γ production).⁵⁸ In the present study we found that the dominant subsets of specific MF cells were 2+ *S. Typhi*-specific cells, which were largely comprised of IFN- γ +CD107a+TNF- α -IL-2-; IFN- γ +CD107a-TNF- α +IL-2 and IFN- γ -CD107a+TNF- α +IL-2- subsets. However, a significant proportion of 3+ MF cells were also induced. Of note, most the CD8+ MF cells produced IFN- γ , co-producing/expressing CD107a+ or TNF- α , whilst a smaller subset also co produced IL-2 (4+). These results markedly extend those reported in other infectious diseases showing that antigen specific MF T_{EM} or T_{EMRA} CD8+T cells that produce IFN- γ also contained subsets co-producing TNF- α , but very few co producing IL-2.^{58, 65} Recently, it has been proposed that during antigen specific memory cell proliferation and differentiation TNF- α and IL-2 producing clones may fade earlier than those secreting IFN- γ . Thus, terminal effector CD8+ memory subsets are comprised of mostly IFN- γ secreting cells with less functional heterogeneity.⁵⁸ In this context, our observations of a dominance of *S. Typhi*-specific 2+ (IFN- γ +CD107a+TNF- α -IL2-) and 3+ (IFN- γ +CD107a+TNF- α +IL-2-) CD8+ T_{EM} cells may indicate that Ty21a immunization elicits a heterogeneous population of activated CD8⁺ T_{EM} that secrete IFN- γ and TNF- α with cytolytic activity (CD107a+), which subsequently become terminal effector cells, maintaining their ability to produce IFN- γ and express CD107a in the absence of TNF- α production.⁵⁸ Interestingly, we observed that post vaccination increases in *S. Paratyphi A*-specific IFN- γ +CD107a+TNF- α -IL-2- MF cells were less pronounced than those observed to *S. Typhi* or *S. Paratyphi B*. However, the significance of this observation is unclear since the exact role of the IFN- γ +CD107a+TNF- α -IL-2- CD8+T cells in protection remains undefined.

The observations that *S. Typhi* and *S. Paratyphi B*-specific IFN- γ + and CD107a+ MF cells, as well as double and triple positive MF subsets were induced at a higher magnitude in T_{EM} subsets than in T_{EMRA} cells, are similar to our previous observations in Ty21a vaccinees.^{22, 25, 26, 30, 32, 44} In contrast, MF cells specific to *S. Paratyphi A* were induced at lower magnitudes in most subjects and without such predominance of responses in T_{EM} subsets. In the absence of known correlates of protection or knowledge on the functional role of MF cells in protection from *S. Typhi* infection, the significance of these differences observed between *S. Paratyphi A* and *S. Paratyphi B* at present is unclear. However, it is reasonable to speculate, the similarities between *S. Typhi* and *S. Paratyphi B* specific CMI responses, as well as the differences with *S. Paratyphi A*, may help explain field trials with Ty21a reporting cross protection against *S. Paratyphi B* but not from *S. Paratyphi A*.¹³ Of note, although similar immune responses were observed in the majority of participants, these responses were, to a certain extent, heterogeneous, with a few volunteers exhibiting different dominant patterns. These results highlight the importance of considering cumulative responses, as well as those from individual volunteers, when interpreting data derived from human studies. Further studies are needed to fully understand the role of these T_M subsets in

protection from enteric fevers. Production of β -chemokines (i.e., RANTES, MIP-1 α , MIP-1 β) by CD8⁺ T cells has been shown, among others, to play an important role in CTL activity.⁶⁶ For example, HIV antigen specific CD8⁺ MIP-1 β ⁺ cells co producing IFN- γ were related with non-progressors suggesting that they might play a role against the infection.^{50, 67} A recent report also showed that in response to *S. Typhi*-antigens, PBMC obtained from *S. Typhi* infected convalescent patients produced MIP 1 β .⁶⁸ We have previously shown that MIP-1 β is co-produced with other cytokines, i.e., IFN- γ , TNF- α and IL-2 following vaccination with Ty21a.²² In this study, we further characterized Ty21a elicited CD8⁺ MF MIP-1 β T cells specific to *S. Typhi*, and provide the first evidence that these cells are cross-reactive to *S. Paratyphi A* and *S. Paratyphi B*. We observed that the majority of these *Salmonella*-specific MIP-1 β ⁺ MF cells co produced/expressed IFN- γ and CD107a, suggesting that Ty21a elicited MF cells co-producing MIP 1 β are likely an important component of a protective CTL response against enteric fevers.

Effector immune responses in the gut microenvironment are expected to be important in protecting the host against *S. Typhi* and other enteric infections, including those caused by *S. Paratyphi A* and *S. Paratyphi B*. In previous studies we have demonstrated that, as expected, a substantial component of the *S. Typhi*-specific IFN- γ ⁺ CD4⁺ and CD8⁺ T cells elicited by Ty21a and CVD 909 had the potential to home to the gut as measured by expression of the integrin $\alpha_4\beta_7$ gut homing molecule.^{25, 30} In this study we extended these observations by demonstrating that *S. Typhi*-specific CD8⁺ MF T cells producing/expressing IFN- γ ⁺, CD107a⁺ and/or MIP-1 β ⁺ elicited by Ty21a immunization consisted of cells expressing, or not, integrin $\alpha_4\beta_7$ and are cross reactive with *S. Paratyphi A* and *S. Paratyphi B*. Of note, *Salmonella*-specific integrin $\alpha_4\beta_7$ ⁺ CD8⁺ MF T_{EM} cells were present in circulation at a lower magnitude than integrin $\alpha_4\beta_7$ ⁻ cells. This observation is likely the consequence of the migration of *Salmonella*-specific integrin $\alpha_4\beta_7$ ⁺ cells to the gut mucosa, resulting in a decrease in circulation.

The present studies have a few limitations. These include the relatively limited number of volunteers studied and the availability of only two time points after vaccination (Days 42 and 84). The latter might have limited our ability to detect post vaccination increases *Salmonella*-specific IL17⁺ cells in the majority of individuals.

In summary, the present investigations provide insights into the immunological basis underlying the observed cross-protection against *S. Paratyphi B*, but not *S. Paratyphi A*, observed in Ty21a field studies.¹³ Overall, these observations support the notion that a bivalent *S. Typhi/S. Paratyphi A* vaccine might be required to protect against enteric fevers.

Methods

Subjects, immunizations and isolation of peripheral blood mononuclear cells (PBMC)

Sixteen healthy adult volunteers (median age 42 years, range 23 to 52 years) from the Baltimore, MD/Washington, DC, area and the University of Maryland Baltimore (UMB) community, who had no history of typhoid fever were recruited for the study with the approval of UMB Institutional Review Board. They received four recommended spaced doses of Ty21a vaccine (Vivotif enteric coated capsules [Crucell]).⁴⁷ Blood samples were

drawn pre-vaccination (day 0) and 42 (day 42) or 84 (day 84) days post-vaccination. PBMC were isolated immediately after blood draws by density gradient centrifugation and were cryopreserved in liquid nitrogen.^{33, 53}

Target/stimulator cell preparation

Epstein Barr virus (EBV)-transformed B-LCL (EBV-B cells) were generated from PBMC obtained from Ty21a vaccinees as previously described.^{27, 53} *Salmonella* strains, i.e., wild-type *S. Typhi* strain (ISP-1820, Vi+, a clinical isolate from Chile), *S. Paratyphi A* (CV 223, ATCC# 9150), and *S. Paratyphi B* (CV 23, a clinical isolate from Chile) were obtained from the Center for Vaccine Development, University of Maryland, USA (CVD) reference stocks. EBV-B cells were incubated with *Salmonella* strains, at an MOI of 10:1 (bacteria:cell) as previously described and rested overnight.^{27, 53} Infected cells were gamma-irradiated (6,000 rad) before being used as “targets” for ex vivo PBMC stimulation. To confirm the adequacy of the infection with *S. Typhi*, *S. Paratyphi A* or *S. Paratyphi B*, infected EBV-B cells were stained with anti-*Salmonella* common structural Ag (CSA-1)-FITC (Kierkegaard & Perry, Gaithersburg, MD) and analyzed by flow cytometry using a customized LSR-II instrument (BD, Franklin Lakes, NJ, USA). The percentage of cells infected with *S. Typhi* was recorded for each experiment. Infected targets were only used if the infection was detected (CSA-1 positive) in >40% of viable cells (**Fig. S2**).

Ex-vivo PBMC stimulation

Frozen PBMC were thawed, rested overnight and stimulated with autologous *S. Typhi*-, *S. Paratyphi A*- or *B*- infected targets at a ratio of 10:1 (PBMC:target). After 2 hours, the protein transport blockers Monensin (1 µg/ml, Sigma) and Brefeldin A (2 µg/ml; Sigma) were added to the PBMC and cultures were continued overnight at 37°C in 5% CO₂. Media alone and uninfected autologous EBV-B cells were used as negative controls. Staphylococcal enterotoxin B (SEB) (10 µg/mL; Sigma) was used as a positive control.

Surface and intracellular staining

Surface and intracellular staining was performed as described previously.²² Briefly, following ex-vivo stimulation, PBMC were first stained for live/dead discrimination using LIVE/DEAD fixable violet dead cell stain kit (Invitrogen, Carlsbad, CA) and then surface stained with a panel of fluorochrome conjugated monoclonal antibodies (mAbs) that included CD14-Pacific Blue (TuK4, Invitrogen), CD19-Pacific Blue (SJ25-C1, Invitrogen), CD3-Qdot 655 (UCHT1, BD), CD4- PerCP-Cy5.5 (SK3, BD), CD8-Qdot 705 (HIT8A, Invitrogen), CD45RA- biotin (HI100, BD), CD62L- APC-EF780 (Dreg 56, Invitrogen), integrin α4β7-Alexa 488 (clone ACT-1; conjugated in house) and CD107a-A647(eBioH4A3, eBiosciences, San Diego, CA). Of note, to maximize the detection of anti-CD107a mAb was added during the overnight ex-vivo stimulation. The cells were then fixed and permeabilized with Fix & Perm cell buffers (Invitrogen) according to the manufacturer's recommendations and was followed by intracellular staining with mAbs against IFN-γ-PE Cy7 (B27, BD), TNF-α-Alexa 700 (MAb11, BD), IL-2-PE (5344.111, BD) and CD69-ECD, (TP1.55.3, Beckman Coulter, CA, USA). For some experiments a modified panel of mAbs (14 colors) was used to concomitantly detect two additional cytokines, i.e., MIP-1β, and

IL-17. This modified panel of mAbs included surface staining with Live/DEAD fixable yellow dead-cell staining kit (Invitrogen), CD14-Brilliant violet (BV) 570 (TuK4, Invitrogen), CD19- BV570 (HIB19, Biolegend, San Diego, CA, USA), CD3- BV650 (OKT3, Biolegend), CD4- PE Cy5 (RPA T4, BD), CD8 PerCP Cy5.5 (SK1, BD), CD45RA-biotin (HI100, BD), CD62L-APC-EF780 (Dreg 56, eBioscience), CD107a-FITC (H4A3, BD) and integrin $\alpha 4\beta 7$ -A647(ACT 1; conjugated in house). Secondary staining was performed with streptavidin Qdot 800 (Invitrogen), followed by intracellular staining with IFN- γ -PE-Cy7 (B27, BD), TNF- α -Alexa 700 (MAB11, BD), IL-2 BV605 (MQ1 17H12, Biolegend), IL 17A BV421 (BL168, Biolegend), MIP-1 β -PE (24006, R&D, Minneapolis, MN, USA) and CD69 ECD or PE (TP1.55.3, eBioscience). After staining cells were fixed in 1% paraformaldehyde and stored at 4 °C until analyzed. Flow cytometry was performed using a customized LSRII flow cytometer (BD) and data were analyzed using WinList version 7 (Verity Software House, Topsham, ME, USA). Of note, in preliminary experiments we optimized the multichromatic panels used in these studies by performing titration of mAbs alone or in combination, as well as fluorescence minus one (FMO) staining, to minimize spectral overlap and compensation (data not shown).

Gating protocol

T cell responses in different live CD8⁺ (CD3⁺, CD8⁺ CD4⁻) T cell memory (T_M) subsets were evaluated by their expression of CD45RA and CD62L into T central memory (T_{CM} ; CD62L⁺ CD45RA⁻), T effector memory (T_{EM} ; CD62L⁻ CD45RA⁻) and T effector memory CD45RA⁺ (T_{EMRA} ; CD62L⁻ CD45RA⁺). Naïve T cells (T_N) were defined as CD62L⁺ CD45RA⁺ (**Fig. S2**). The FCOM analysis tool (WinList version 7) was used to classify events based on combinations of selected gates in multidimensional space (i.e., whether cells express single or multiple intracellular cytokines and/or CD107a alone or in all possible combinations) for the detection of single or MF cells. Flow cytometric analyses were performed in 300,000 500,000 events collected for each sample, of which 161,700 (128,023 - 208,752) (median and interquartile range in parenthesis) were within the live lymphocyte gate (**Fig. S3 panel A1**).

Statistical analyses

The statistical tests used to analyze each set of experiments are indicated in the Figure Legends. P values of <0.05 were considered significant.

Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

Acknowledgments

We are indebted to the volunteers who allowed us to perform this study. We also thank Robin Barnes and the staff from the Recruiting Section of Center for Vaccine Development for their help in collecting blood specimens; Regina Harley, Catherine Storrer, Haiyan Chen and Shah Zafar for excellent technical assistance. This paper includes work funded, in part, by NIAID, NIH, DHHS grants R01-AI036525 (to M.B.S.), U19 AI082655 (Cooperative Center for Translational Research in Human Immunology and Biodefense [CCHI], to M.B.S.), and U54-AI057168 (Regional Center for Excellence for Biodefense and Emerging Infectious Diseases Research Mid-Atlantic Region [MARCE] and U19-AI109776 (Center of Excellence for Translational Research [CETR], to M.M.L. and M.B.S).

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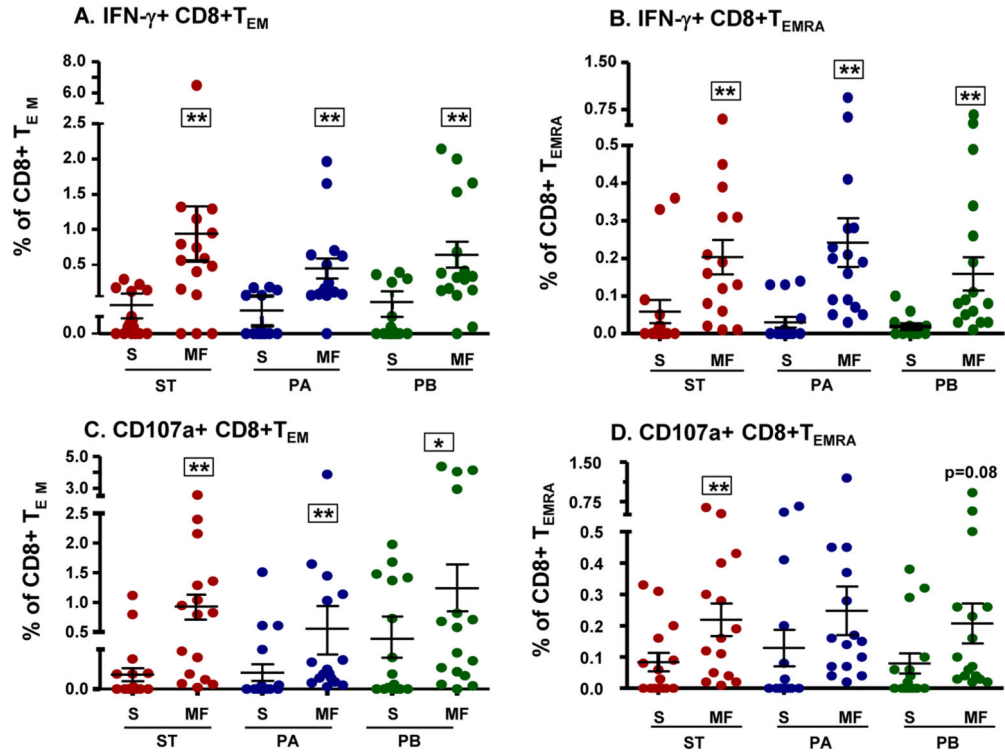


Figure 1. Induction of multifunctional cells in Ty21a vaccinees

PBMC collected from Ty21a vaccinees (n=16) were stimulated with *S. Typhi*-infected targets and the data analyzed using FCOM (described in the text). Shown are the peak post-vaccination increases in single positive (S) and IFN- γ + (panels A,B) and CD107+ (panels C,D) total multifunctional (MF, the sum of all multifunctional subsets) cells in CD8+T_{EM} (panel A,C) and CD8+T_{EMRA} (panels B,D) subpopulations, specific for *S. Typhi* (ST)-, *S. Paratyphi A* (PA) or *S. Paratyphi B* (PB)-infected targets.

Post-vaccination peaks: Peak of the responses at days 42 or 84 minus pre vaccination [day 0] levels

Horizontal bars represent mean \pm SEM.

**p<0.01. *p<0.05 compared to corresponding Single positive cells by Wilcoxon signed rank test, 2-tail

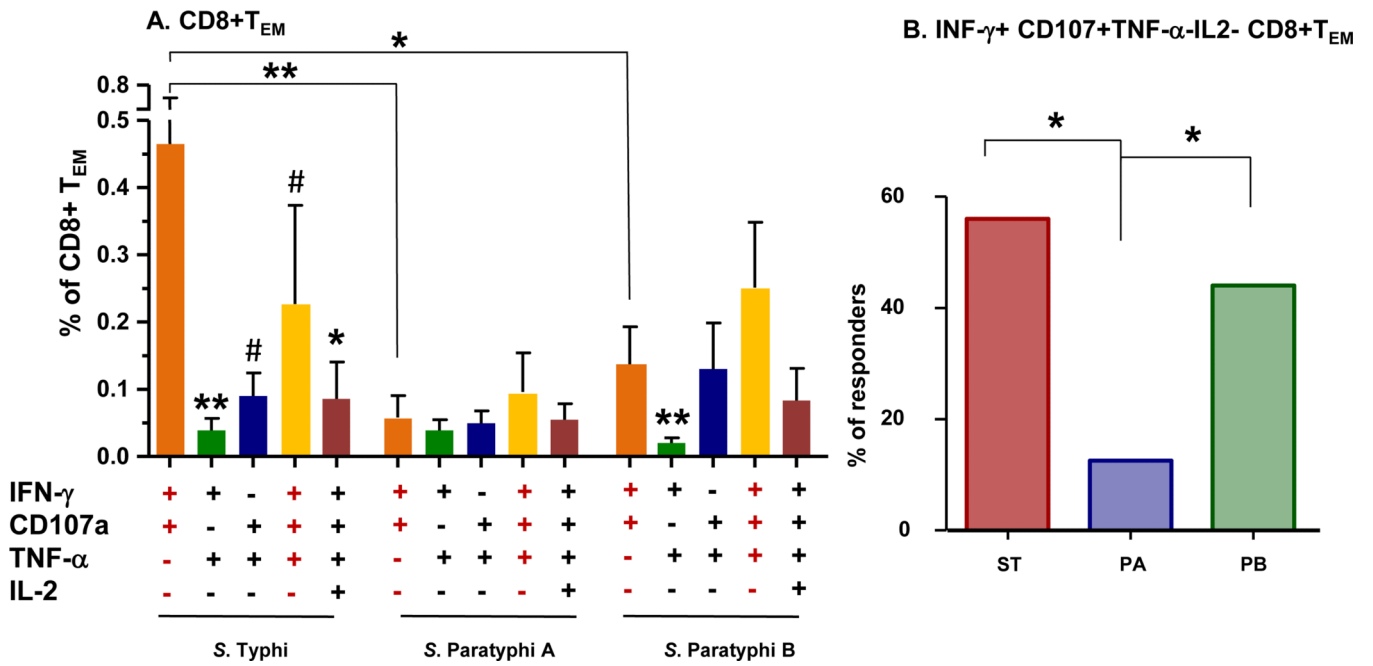


Figure 2. Post-vaccination increases in multifunctional (MF) CD8+TEM cells in PMBC from Ty21a vaccinees (n=16)

Induction of multiple cytokine (IFN- γ , TNF- α and/or IL-2)-producing and/or expressing CD107a CD8+TEM cells following stimulation with targets infected with *S. Typhi* (ST), *S. Paratyphi A* (PA) or *S. Paratyphi B* (PB). Data was analyzed using FCOM as described in the text. The post-vaccination peak increases (peak level at days 42 or 84 post-vaccination minus the corresponding pre-vaccination levels) in dominant subpopulations are shown as mean \pm SEM (**panel A**). The percentage of responders was calculated as: (number of volunteers with peak post vaccination increases \geq 0.1% in IFN- γ +CD107a+TNF- α -IL-2 subsets)/(total volunteers [n=16])X100 (**panel B**).

Horizontal bars represent mean \pm SEM

p<0.01; *p<0.05; #p<0.1 compared to IFN- γ +CD107a+TNF- α -IL-2- MF cells for each corresponding *Salmonella*-infected target. Other significance values relate to the indicated sets. Wilcoxon signed rank test, 2-tail (panel A**). *p<0.05, by Chi square test 2-tail (**panel B**)

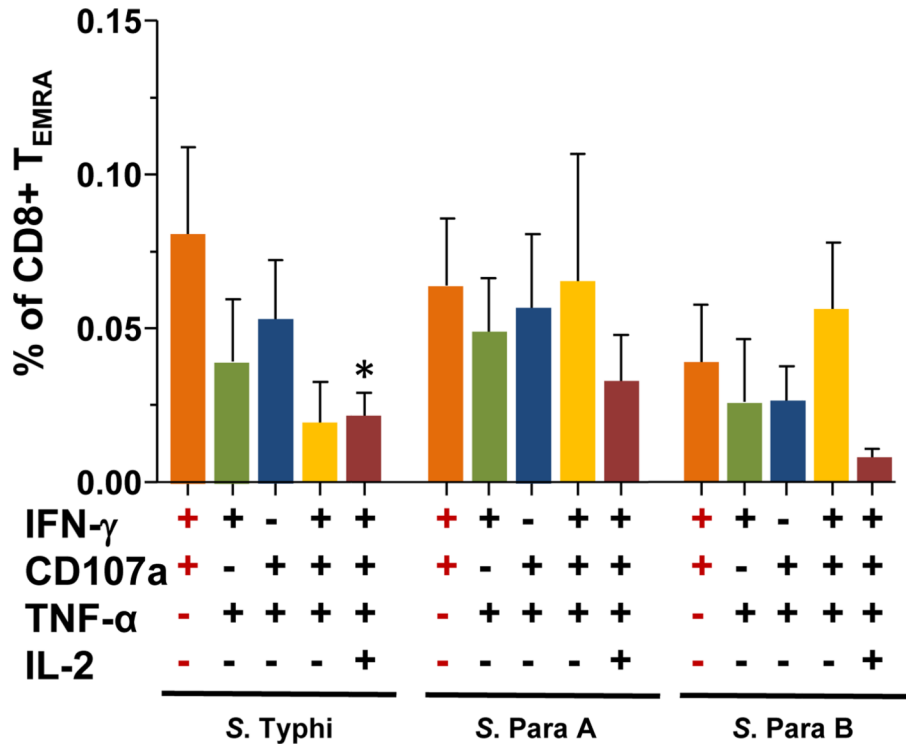


Figure 3. Post-vaccination increases in Multifunctional (MF) CD8+ T_{EMRA} subsets in PMBC from Ty21a vaccinees (n=15)

Shown are post-vaccination peak increases (peak at days 42 or 84 post-vaccination minus the corresponding pre-vaccination levels) in the 5 dominant MF subpopulations following stimulation with targets infected with *S. Typhi*, *S. Paratyphi A* (*S. Para A*) or *S. Paratyphi B* (*S. Para B*). Bars indicate mean \pm SEM.

*p<0.05, compared to IFN- γ +CD107a+TNF- α -IL2- MF cells for *S. Typhi*-infected targets. Wilcoxon signed rank test, 2-tail.

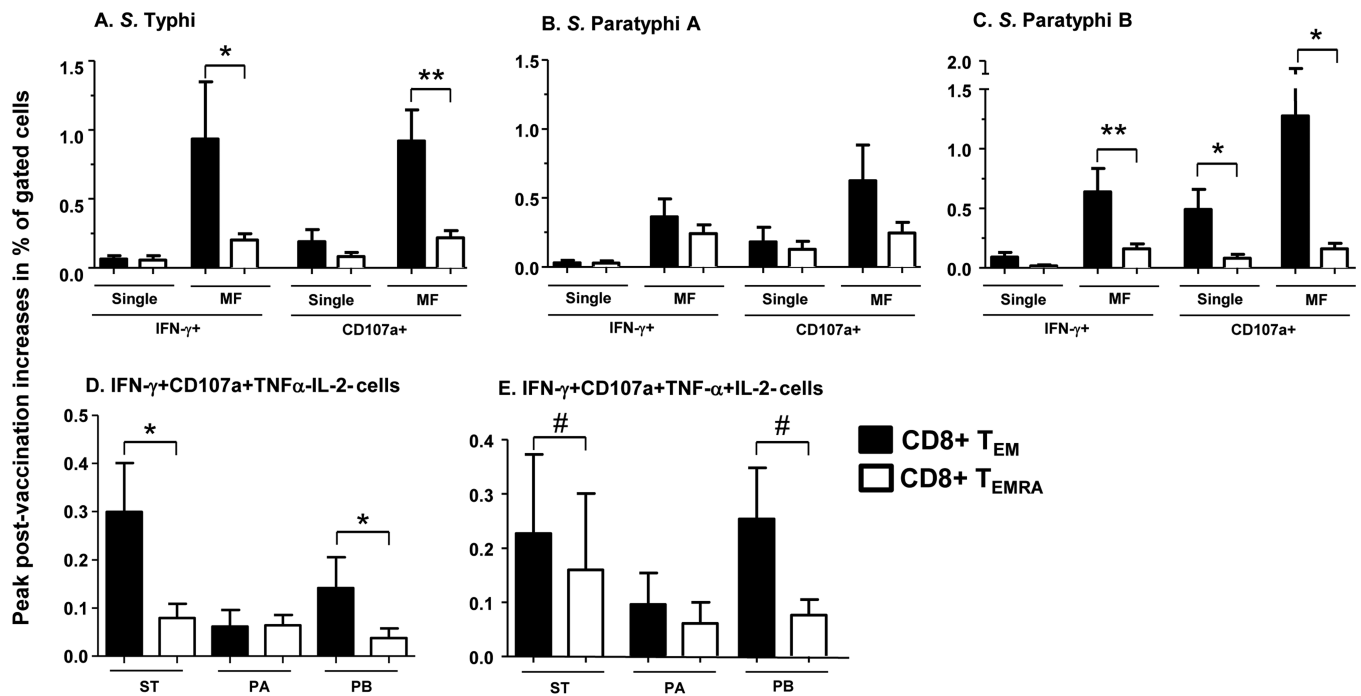


Figure 4. Comparison of the induction of cross/reactive multifunctional cells between CD8T_{EM} and CD8T_{EMRA} subpopulations

Post-vaccination peak increases (peak level at days 42 or 84 post-vaccination minus the corresponding pre-vaccination levels) in IFN- γ ⁺ and CD107a⁺ single and MF subsets in CD8T_{EM} and CD8T_{EMRA} subsets, in response to *S. Typhi* (ST), *S. Paratyphi A* (PA) and *S. Paratyphi B* (PB) were compared in **panels A, B and C**, respectively. Similar comparisons with IFN- γ ⁺CD107a⁺TNF- α -IL-2- and IFN- γ ⁺CD107a⁺TNF- α +IL-2- MF subsets are shown in **panels D and E**, respectively.

Bars indicate mean \pm SEM.

* $p < 0.05$, # $p < 0.10$ compared to corresponding CD8 T_{EM} subsets. Wilcoxon signed rank test, 2-tail.

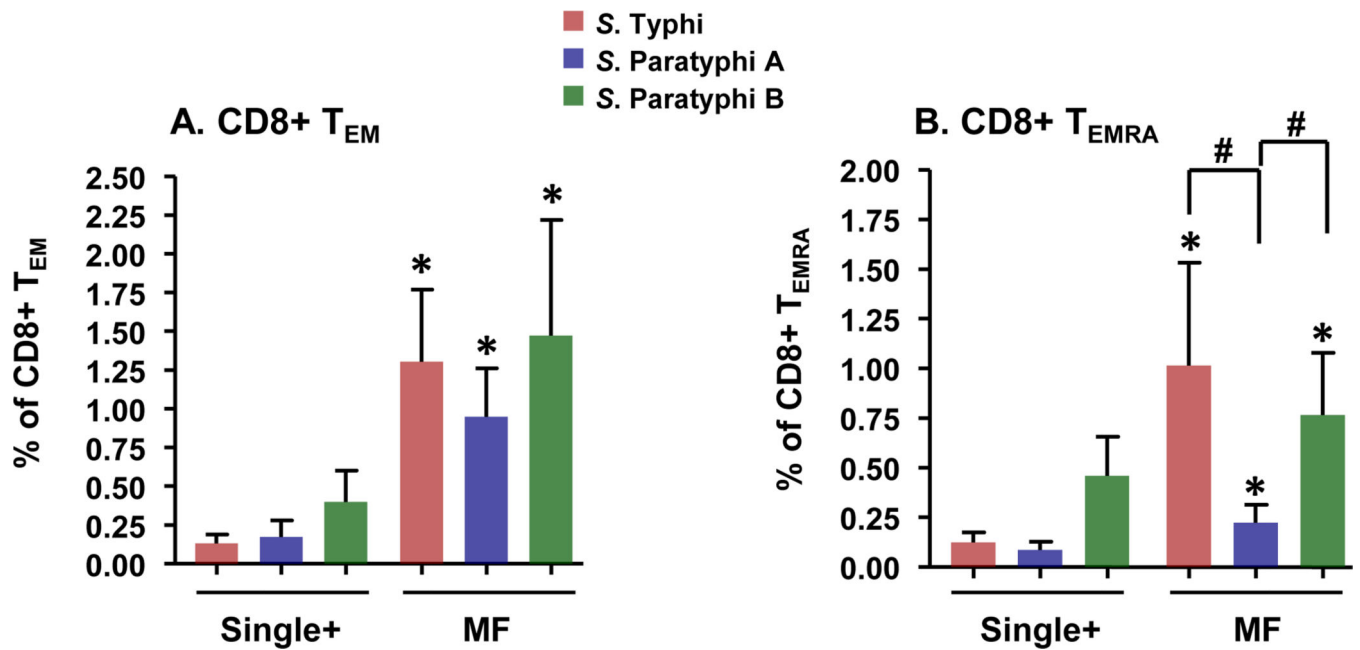


Figure 5. Post-vaccination increases in MIP-1 β producing CD8+ cells in response to *S* Typhi-, *S.* Paratyphi A- and *S.* Paratyphi B-infected targets

Post-vaccination peak increases in MIP-1 β production by CD8+T_{EM} (**panel A**) and CD8+ T_{EMRA} (**panel B**) in single positive (single+) for MIP-1 β (MIP-1 β +IFN- γ -TNF- α -IL-2-IL-17-) and all other MIP-1 β + that are multifunctional (MF) were analyzed by FCOM in PBMC obtained from Ty21a vaccinees (n=8) Bars indicate mean \pm SEM.

*p<0.05, MF compared to the corresponding single positive cells. #: P<0.12. Wilcoxon signed rank test, 2-tail.

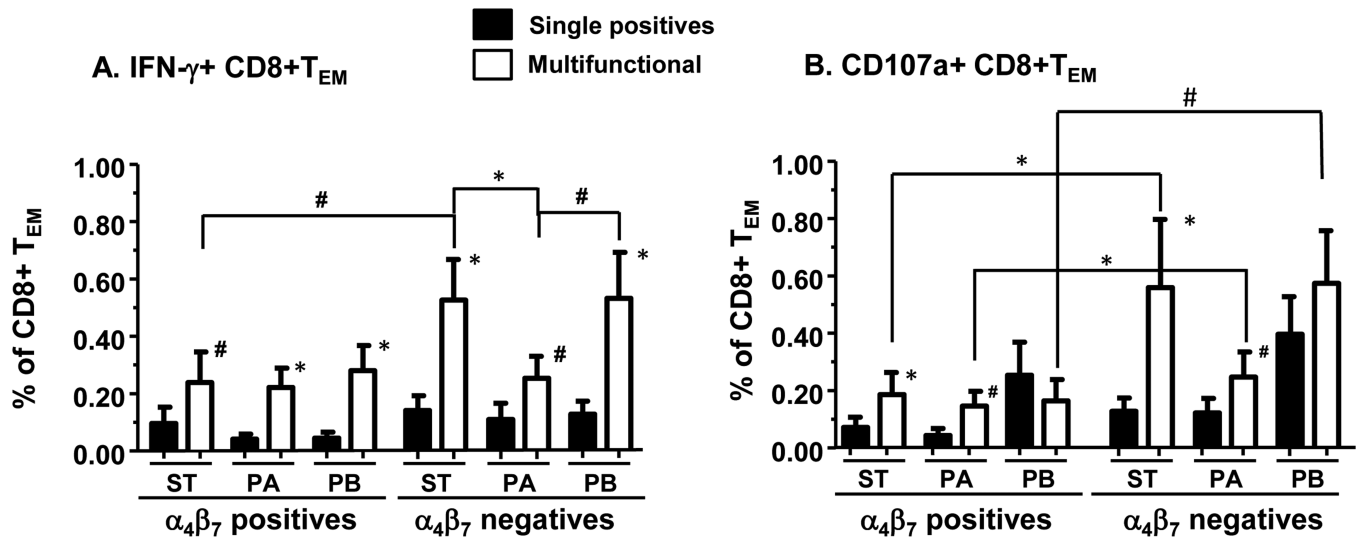


Fig. 6. Concomitant expression of the gut homing molecule integrin $\alpha_4\beta_7$ by Salmonella-specific single and multifunctional CD8+T_{EM} cells in Ty21a vaccinees
 PBMC collected from Ty21a vaccinees were stimulated with *S. Typhi* (ST), *S. Paratyphi A* (PA) and *S. Paratyphi B* (PB)-infected targets and the data were analyzed using FCOM (described in the text). Shown are the peak post-vaccination increases in antigen-specific IFN- γ + (**panel A**; n=12), CD107a+ (**panel B**; n=12), single positive (closed bars) and the sum of all multifunctional (open bars) cells in CD8+ T_{EM} subpopulations expressing integrin $\alpha_4\beta_7$ ($\alpha_4\beta_7$ positives) or not ($\alpha_4\beta_7$ negatives). Post-vaccination peaks: Peak responses at days 42 or 84 minus pre vaccination [day 0] levels Bars indicate mean \pm SEM. *p<0.05, #p 0.1 compared to corresponding Single positive cells by Wilcoxon signed rank test, 2-tail. Other significance values relate to the indicated sets.