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Bacille Calmette Guerin vaccination of human newborns induces T cells with complex cytokine and phenotypic profiles

Andreia P. Soares^{*}, Thomas J. Scriba^{*}, Sarah Joseph^{*}, Ryhor Harbacheuski[†], Rose Ann Murray[†], Sebastian J. Gelderbloem^{*}, Anthony Hawkridge^{*}, Gregory D. Hussey^{*}, Holden Maecker[‡], Gilla Kaplan[†], and Willem A. Hanekom^{*}

*South African Tuberculosis Vaccine Initiative, University of Cape Town, South Africa

[†]Laboratory of Mycobacterial Immunity and Pathogenesis, Public Health Research Institute, Newark, NJ

[‡]BD Biosciences, San Jose, California

Abstract

The immune response to vaccination with Bacillus Calmette-Guerin (BCG), the only tuberculosis vaccine available, has not been fully characterized. We used multiparameter flow cytometry to examine specific T cell cytokine production and phenotypic profiles in blood from 10-week old infants, routinely vaccinated with BCG at birth. Ex vivo stimulation of whole blood with BCG for 12 hours induced expression of predominantly IFN- γ , IL-2 and TNF- α in CD4+ T cells, in 7 distinct cytokine combinations. IL-4 and IL-10 expression were detected in CD4+ T cells at low frequencies, and only in cells that did not co-express Type 1 cytokines. Specific CD8+ T cells were less frequent than CD4+ T cells, and produced mainly IFN- γ and/or IL-2, and less TNF- α , IL-4 and IL-10. Importantly, many mycobacteria-specific CD4+ and CD8+ T cells did not produce IFN- γ . The predominant phenotype of BCG-specific Type 1 T cells was that of effector cells, i.e., CD45RA-CCR7-CD27+, which may reflect persistence of *M. bovis* BCG in infants until 10 weeks of age. Among 5 phenotypic patterns of CD4+ T cells, central memory cells were more likely to be IL-2+, and effector cells more likely to be IFN- γ +. We concluded that neonatal vaccination with BCG induces T cells with a complex pattern of cytokine expression and phenotypes. Measuring IFN- γ production alone underestimates the magnitude and complexity of the host cytokine response to BCG vaccination, and may not be an optimal readout in studies of BCG and novel tuberculosis vaccination.

Keywords

Human; Th1/Th2 Cells; T cells; Cytokines; Memory; Vaccination

Introduction

Nearly one-third of the global population is latently infected with *Mycobacterium tuberculosis*, and approximately 2 million people die of tuberculosis (TB) disease every year (Tuberculosis Fact Sheet, 2005, World Health Organization, Geneva, Switzerland). Bacille Calmette-Guerin (BCG), the only vaccine against tuberculosis currently available, has variable efficacy in preventing pulmonary disease (1), but 80% efficacy in preventing childhood miliary disease and meningitis (2). Our knowledge of immunity induced by BCG

Disclosures None. vaccination is incomplete, particularly after human newborn vaccination. However, infants will be targets of novel, safer and more efficacious tuberculosis vaccines in the future, and a better understanding of the immune response induced by newborn BCG vaccination is likely to facilitate development of improved vaccines.

Experimental evidence suggests that both CD4+ and CD8+ T cells are important for protection against mycobacteria (3-6). In humans, the role of CD4+ T cells has been highlighted by a increased risk of disease after infection with *M. tuberculosis* when CD4+ T cell numbers decline in HIV-infected persons (7). A primary function of both CD4+ and CD8+ T cells is to produce the Type 1 cytokine IFN- γ . The critical role of this cytokine has been demonstrated by characteristic severe mycobacterial disease in patients with mutations of the IFN- γ receptor (8, 9). Other Type 1 cytokines, such as TNF- α and IL-2, may also be important in protection against tuberculosis: the role of TNF- α has been underscored by high rates of reactivation of latent tuberculosis following treatment of rheumatoid arthritis patients with specific inhibitors of this cytokine (10, 11); T cell IL-2 expression has been associated with long-term memory (12, 13), which is the aim of protective immunity. BCG vaccination of human newborns does indeed induce specific CD4+ and CD8+ T cells, capable of producing IFN- γ (14-16) Previous studies have also shown that BCG vaccination of infants induces TNF- α which is detectable in plasma by ELISA (17). However, the pattern of production of all Type 1 cytokines, on a single cell basis, has not been delineated.

Our aim was to also describe expression of Type 2 cytokines such as IL-4, thought to reflect a suboptimal immune response to mycobacteria (18, 19). Although BCG vaccination of infants has been shown to induce low levels of Type 2 cytokines (15, 16, 20), the detection was in plasma, and cell-associated expression has not been reported. We also wished to assess T cell IL-10 expression, as this cytokine is likely to be an important regulator of effector T cell responses against tuberculosis (21) and is induced by newborn BCG vaccination (15, 16, 22).

The memory phenotype of T cells induced by BCG vaccination of newborns has not been described. Antigen-experienced cells may be categorized based on expression of surface markers (13, 23-26). Central memory cells express CCR7 but not CD45RA and are likely to represent a long-lived population, which expands rapidly in lymph nodes following subsequent antigen encounter (23, 27). In contrast, effector cells are both CCR7- and CD45RA- and act immediately following antigen exposure, but have limited proliferative capacity (13, 23). A third subset, terminally differentiated memory cells, are CD45RA+ and CCR7- and the most differentiated subpopulation, based on short telomere length and function (26, 28). Naïve, or non-antigen-experienced T cells, characteristically express both CD45RA and CCR7 (13, 26, 28). Combination of markers other than CCR7 and CD45RA may also differentiate subsets of antigen-experienced cells. Fritsch et al. recently proposed the phenotypic classification of CD4+ T cell populations based on expression of CD27 and CCR7 (26). Central memory T cells were defined as CD27+ and CCR7+, effectors as CD27+ and CCR7-, and terminally differentiated T cells as CD27- and CCR7- (26). Our aim was to evaluate expression of all these markers among antigen-experienced T cells induced by BCG. Although several investigators have characterized mycobacteria-specific immune responses by four color flow cytometry (29-31), these studies could only measure two cytokines or phenotypic markers at a time and thus most likely underestimated the complexity of the response.

Our hypothesis was that BCG vaccination of newborns would induce both CD4+ and CD8+ T cells capable of producing multiple cytokines, and that a central memory phenotype of specific cells would be dominant 10 weeks after vaccination. We used an intracellular cytokine assay with multiparameter flow cytometry to comprehensively characterize these

variables. To achieve our goals, we established clinical structures and optimized techniques (32) to overcome hurdles common to investigation of immunity in infants.

Materials and Methods

Study participants and blood collection

Healthy 10-week old infants, routinely vaccinated intradermally with BCG (Statens Serum Institut, Copenhagen) at birth, were enrolled in the Cape Town region of South Africa. This area has a very high TB disease incidence in under 5 year-olds, exceeding 2% per year in certain areas (32). Infants born to HIV-positive mothers, infants known to be HIV positive, infants with suspected or confirmed TB disease, infants with possible exposure to TB disease and infants with any other active or chronic illnesses at the time of enrollment, were excluded. Human participation was according to the US Department of Health and Human Services and good clinical practice guidelines. This included protocol approval by the University of Cape Town Research Ethics Committee and written informed consent. Sodium heparinized blood was collected for assessment with 2 different flow cytometric protocols (see below): 29 infants to assess cytokine expression of T cells, and 27 infants to assess the phenotype of IFN- γ and IL-2-expressing T cells.

Antigens and Antibodies

BCG was reconstituted from the vaccine vial (SSI) at 1.8×10^6 organisms/mL, as previously described (32). The positive control streptococcal enterotoxin B (SEB, Sigma) was used at 10μ g/ml. The co-stimulatory antibodies anti-CD28 and anti-CD49d (both from BD Biosciences) were each used at 1μ g/ml. Cytokine profiles of BCG-specific T cells were examined using the following conjugated antibodies: anti-CD3 Amcyan (SK7) and anti-IL-2 Alexa 610-PE (5344.111), both custom-conjugated at BD Biosciences; and anti-CD4 AlexaFluor 700 (RPAT4), anti-CD8 Cy5.5-PerCP (SK1), anti-IL-4 FITC (MP4-25D2), anti-IFN- γ PE (25723.11), anti-IL-10 APC (JES3-19F1), and anti-TNF- α Cy7-PE (MAb11), all obtained from BD Biosciences. A separate protocol was used for assessing T cell phenotypes, using the following conjugated antibodies: anti-CD3 PacBlu (UCHT1), anti-CD4 Cy5.5PerCP (SK3), anti-CD8 Cy5.5PerCP (SK1), anti-CD45RA Cy7PE (L48), anti-CD27 PE (MT271), anti-IFN- γ AlexaFluor 700 (B27) and anti-IL-2 FITC (5344.111), all obtained from BD Biosciences, and anti-CCR7 APC (150503), obtained from R&D Systems.

Whole Blood Intracellular Cytokine Detection Assay

To determine cell-associated cytokine production, 1mL heparinized whole blood was incubated with BCG and anti-CD28 and anti-CD49d, as described before (32). Blood incubated with SEB and co-stimulants, or with co-stimulatory antibodies alone (UNS), served as positive and negative controls, respectively. Brefeldin A (10μ g/mL, Sigma) was added during the last 5 hours of incubation to capture cytokines intracellularly. After a total incubation of 12 hours, red blood cells were lysed and white cells fixed with FACS Lysing Solution (BD Biosciences), followed by cryopreservation.

Cell staining and flow cytometric analysis

To detect intracellular cytokines, cryopreserved cells were thawed, washed in PBS, permeabilized with Perm/Wash solution (BD Biosciences) and incubated at 4°C with fluorescence-conjugated antibodies for an hour. To assess T cell memory phenotypes, a two-step staining method, resulting in optimal staining, was used: cells were thawed, washed in PBS, and permeabilized with Perm/Wash solution, and then incubated with surface marker

antibodies for an hour at 4°C, followed by an additional hour with antibodies specific for intracellular cytokines. Flow cytometric acquisition was completed on a LSRII flow cytometer (BD Biosciences) configured for 3 lasers and 12 detectors. All the cells in the tube were acquired. Analysis was performed using FACSDiva software (BD Biosciences). Although automated compensation with mouse IgG κ -beads was applied, compensation settings were assessed manually after acquisition, and adjusted if necessary. Multiparameter panel development (not shown) included evaluation of appropriate staining controls, of antibody and fluorochrome interactions and of spectral overlap, using control blood samples (33, 34).

Cut-offs to determine positive cytokine expression in CD4+ and CD8+ T cells from blood incubated with BCG were set using cells from blood incubated with co-stimulatory antibodies alone (negative control) (Fig. 1A). Angled cut-off lines were necessary for some fluorochromes, because of data spread at higher fluorescence intensities, following instrument compensation (e.g., Figure 1). SEB was an excellent positive control for induction of all cytokines, except IL-4. IL-4-expressing HICK-2 cells (BD Biosciences), processed as per manufacturer's protocol, were used as positive control for IL-4. Where distinction between positive and negative surface marker populations was not clear, isotype-matched control antibody staining was used to set cut-offs for phenotypic markers.

Plasma Cytokine Detection

Plasma was collected from the stimulated whole blood after 7 hours, and cryopreserved. Later, thawed plasma was used to measure levels of IL-2, IL-4, IL-10 and IFN- γ with multiplex beads following the manufacturer's instructions (Bio-Rad Laboratories), and read on a luminometer (Luminex). The range of detection for all cytokines was 1.95–32,000pg/mL. The optimal plasma dilution for our assay, determined in pilot experiments, was 1:4. Background cytokine levels measured in plasma harvested from unstimulated blood were subtracted from BCG-stimulated blood.

Statistical considerations

Negative control (background) values for Type 1 cytokine expression were not subtracted from BCG-induced responses, as the median backgrounds for all Type 1 CD4+ T cell subsets was 0.001% (range, 0.000%-0.01%) and for CD8+ T cell subsets 0.000% (range, 0.000%-0.01%). We used an empiric cut-off value of 0.01% as positive: given that a median of 508,509 CD4+ T cells and 188,498 CD8+ T cells were collected, this cut-off was predicted to be >90% different from background, at an alpha of 0.05 (35). IL-4 and IL-10 expression were reported with backgrounds (see below). Nonparametric tests were used to compare differences in cytokine expression and phenotypic profiles between CD4+ and CD8+ T cells. Associations between cellular expression of cytokines and plasma levels of these were assessed by the nonparametric Spearman test. Data were considered statistically significant when p<0.05. Statistical analysis was performed using GraphPad Prism 4 (GraphPad Software).

Results

BCG-specific CD4+ and CD8+ T cell Type 1 cytokine production

Intracellular expression of 3 Type 1 cytokines thought to be critical for protective immunity against mycobacteria, IFN- γ , IL-2 and TNF- α (36, 37), was evaluated by incubating blood from 29 BCG-vaccinated infants with BCG for 12 hours (Figure 1). The frequencies of CD4+ T cells expressing either IFN- γ or IL-2 or TNF- α were similar (Figure 2A). Lower frequencies of CD8+ T cells expressed IFN- γ or IL-2, compared with CD4+ T cells (p<0.05), while TNF- α was expression was very low in CD8+ T cells (Figure 2A). There

was a strong positive correlation between the frequencies of CD4+ and CD8+ T cells expressing IFN- γ , and between the frequencies of these cells expressing IL-2 (R=0.770 R=0.879, respectively, both p<0.0001, Spearman test).

Analysis of simultaneous expression of IFN- γ , IL-2 and TNF- α on a single cell level revealed 7 distinct Type 1 cytokine-expressing CD4+ T cell populations (Figure 2B). Most cytokine-producing CD4+ T cells expressed 2 of the 3 cytokines. Among CD8+ T cells, the dominant population expressed IFN- γ only; 3 other populations were discernable (Figure 2C). Importantly, a substantial proportion of CD4+ T cells expressing IL-2 and/or TNF- α did not co-express IFN- γ (Figure 2 D). Similarly, among CD8+ T cells, a proportion of IL-2 expressing T cells also did not co-express IFN- γ (Figure 2 D).

All 3 Type 1 cytokines could also be detected in plasma (data not shown). There was a significant correlation between plasma levels of IFN- γ , IL-2 and TNF- α and frequencies of CD4+ T cells producing these cytokines (Table I). Plasma IFN- γ and IL-2 also correlated with frequencies of CD8+ T cells producing IFN- γ and IL-2, respectively (Table I).

Taken together, we concluded that BCG vaccination of newborns induces multiple Type 1 T cell subsets defined by expression of distinct cytokine combinations.

BCG-specific IL-10 and Type 2 cytokine production

The frequency of T cells expressing IL-10 or IL-4 following incubation of whole blood with BCG was low (Figure 3). Few donors had responses above 0.01%, our cut-off for a positive Type 1 cytokine response, but IL-10 and IL-4 production were consistently above the background expression levels found in blood not incubated with BCG (Figure 3B and 3D). CD4+ T cell expression of both cytokines was slightly higher than that of CD8+ T cells (CD4+ T cell expression shown in Figure 3; CD8+ T cells: a median 0.004%, range 0.001%-0.024%, expressed IL-10 and 0.005%, 0.000%-0.011%, IL-4). IL-10 and IL-4 were never co-expressed by cells making Type 1 cytokines (Figure 3A and 3C). Intracellular IL-4 could be readily detected in HICK-2 cytokine-expressing cells, which served as positive control (Figure 3C). IL-4 and IL-10 were detected at low levels in plasma of whole blood incubated with BCG (Figure 4).

We concluded that BCG vaccination of newborns induces low levels of IL-4 and IL-10 expression.

Phenotypic profiles of specific Type 1 T cell subsets

Studies in other infectious disease models have shown that distinct populations of antigenexperienced T cells may be associated with long-lived protection (38). We therefore examined the phenotypic profiles of BCG-induced T cells. Specific CD4+ T cells were defined as either IFN- γ or IL-2-expressing (Figure 5A-D); frequencies of cells expressing other cytokines were too low to reliably delineate the phenotype. Five major and distinct antigen-experienced CD4+ T cells subsets could be discerned, based on expression of CD45RA, CCR7, and CD27 (Figure 6A-C). By far the most common phenotype of both IFN- γ and IL-2-expressing CD4+ T cells was CD45RA–CCR7–CD27+ (Figure 6A), a phenotype that has been reported to be characteristic of effector T cells (13). The second most common phenotype among IFN- γ -expressing CD4+ T cells was CD45RA–CCR7– CD27–, also characteristic of effector T cells (Figure 6A). Among IL-2-expressing CD4+ T cells, the latter population was significantly less frequent (Figure 6A), while central memory phenotypes were more common: IL-2+ cells were more likely to be CD45RA– CCR7+CD27+, CD45RA+CCR7–CD27+, or CD45RA+CCR7+CD27+, compared with IFN- γ -expressing CD4+ T cells (Figure 6A). CD4+ T cells that expressed both IFN- γ and IL-2 were predominantly effector memory cells (CD45RA–CCR7–CD27+ or CD45RA–CCR7+CD27+; Figure 6B).

Phenotypes of BCG-specific CD8+ T cells could reliably be detected only for IFN- γ producing cells, as the frequencies of IL-2-producing CD8+ T cells were too low. CD8+IFN- γ + T cells also displayed a pre-dominant CD45RA–CCR7–CD27+ effector phenotype (Figure 6D). Unlike CD4+IFN- γ + T cells, a central memory population (CD45RA+CCR7+CD27+) was the second most common.

We concluded that the majority of specific T cells induced by BCG vaccination of newborns has an effector phenotype, and that IL-2 expression is more likely to be associated with a central memory phenotype.

Discussion

We showed that BCG vaccination of human newborns induces a diverse set of T cells, delineated by distinct cytokine production and phenotypic profiles. BCG-specific T cells produced mainly Type 1 cytokines, as has been demonstrated before (14-16, 20, 22). However, we showed that not only IFN- γ is produced. A considerable number of IFN- γ -negative CD4+ T cells were present, expressing the other Type 1 cytokines IL-2 and TNF- α . Similarly, many CD8+ T cells produced IL-2 in the absence of IFN- γ . The most commonly used measure of mycobacteria-induced immunity today is IFN- γ production, be this to diagnose latent infection via IFN- γ release assays (39, 40) or to describe human immune responses to novel tuberculosis vaccines (41). Recent experimental data suggest that measuring IFN- γ may not correlate with vaccination-induced protection against tuberculosis (42-44). This strongly supports measurement of all 3 cytokines on a cellular level to delineate a mycobacteria-specific Type 1 response. Measuring a single component of the immune response may underestimate the magnitude and complexity of BCG-induced immunity.

A BCG-induced CD8+ T cell response was readily detectable, as Type 1 cytokine-producing cells. Murine studies suggest that CD8+ T cells play an important role in control of M. *tuberculosis* infection and contribute substantially to total IFN- γ production (4). Although a BCG-induced CD8+ T cell response has been described before (14, 45), we now show that the response is characterized by both IFN- γ and IL-2 producing subsets. An interesting observation was that CD8+ T cells produced very little TNF-a. Smith, et al., detected similar frequencies of CD8+ T cells expressing TNF-a and IFN-y following incubation of PBMC with BCG for 6 days (45). The contrasting low frequency of TNF- α expression observed in our study could be due to differences in assays, as we measured cytokine expression 12 hours after incubation of whole blood with BCG. In longer term assays such as those performed by Smith, et al., TNF-a production may be derived from newly differentiated effector T cells, whereas our short term assay measures cytokine producing potential directly ex vivo. Our results also contrast with data from HIV-infected adults, whose HIV-specific CD8+ T cells readily express TNF-a in short-term intracellular cytokine assays (46, 47), implying that BCG-specific CD8+ T cells in newborns express little TNF-a.

The memory phenotype of BCG-induced T cells has not been reported. We identified five phenotypically distinct subsets within BCG-specific Type 1 T cells, based on expression of CD45RA, CCR7 and CD27. The predominant phenotype of both IFN- γ and IL-2-expressing CD4+ and CD8+ T cells was CD45RA–CCR7–CD27+. CCR7– effector cells are characteristic of persistent activation of T cells seen in chronic viral infections where antigen is not cleared (48-50). Similarly, increased numbers of effector T cells are observed in

children with active TB (51). It is possible that *M. bovis* BCG had persisted up to 10 weeks of age, the time at which blood was collected from infants, resulting in this predominant phenotype.

Distinct differences in phenotypes were observed among CD4+ IFN- γ and IL-2-expressing cells. IL-2-expressing cells were significantly more likely to have a central memory phenotype, e.g., CD45RA–CCR7+CD27+, compared with IFN- γ -expressing cells. IL-2expressing T cells induced by BCG vaccination therefore follows patterns similar to those previously described for purified human central memory populations, which are likely to express IL-2 (13). However, our results, from 10 weeks after vaccination with BCG, contrast with results obtained 8 weeks after vaccination with tetanus toxoid; specific IL-2-expressing central memory cells were present but in a relatively small population, whereas this population was predominant after tetanus vaccination (52). Tetanus toxoid is not a persistent antigen; we therefore hypothesize that viable *M. bovis* BCG persisted following vaccination, driving differentiation predominantly into IFN- γ -expressing effector cells, and preventing differentiation into IL-2-expressing central memory cells. An alternate hypothesis is that continuous exposure to environmental mycobacteria may contribute to chronic immune activation, and therefore a predominance of effector T cells. The infants in this study were from a region with high rates of environmental mycobacteria, and unpublished data suggest that exposure may occur even within the first 10 weeks of life. It has been proposed that exposure to environmental mycobacterial antigens in tropical regions may undermine protective immunity induced by BCG (53). A third hypothesis is that the time point at which we measured the host response was too early for full differentiation into the central memory phenotype. A central memory/IL-2 phenotype has been associated with improved prognosis of chronic human viral infections such as HIV (49), and in experimental settings of chronic intracellular bacterial infection with long-lived protection (38). It remains to be determined whether a central memory phenotype is also associated with successful vaccination against tuberculosis.

We found that a significant amount of specific CD4+ and CD8+ T cells had a phenotype traditionally regarded as naïve, i.e., CD45RA+ and CCR7+. In a recent study of children with tuberculosis, Caccamo et al. also described this population, identified as specific by MHC Class I pentamers of antigen-85A (54). We propose that this CD45RA+CCR7+ population reflects early differentiation into antigen-specific cells, prior to loosing CD45RA expression.

We could detect only very low intracellular expression of the Type 2 cytokine IL-4. Expression of Type 2 cytokines has been associated with a suboptimal immune response to mycobacteria (55). For example, Ordway, *et al.*, showed that long-term control of latent *M. tuberculosis* infection in humans appeared to be associated with optimal Type 1 cytokine production and absence of detectable Type 2 cytokine production (56). They showed that high percentages of IL-4 expressing CD8+ and $\gamma \delta$ T cells soon after M.tb infection were associated with ultimate development of TB disease. The IL-4-expressing cells were detected after incubation of PBMC for 6 days, which contrasted with our 12-hour assay; longer-term assays may be required to detect these cells in the setting of BCG vaccination of the newborn. Intracellular expression of IL-10 could also be detected in our cohort, was low, and, like IL-4 expressing T cells were induced regulatory T cells, which are expected to be present at low frequencies. Regulatory T cells control conventional effector immune responses, are induced by infections (57), and likely also by vaccination with BCG.

Our results clearly demonstrate the advantages of complex multiparameter flow cytometric analysis for deciphering a vaccination-induced immune response. However, this technology

is not readily available. It is therefore important to note, from our findings, that when detection of IFN- γ or IL-2 is the aim and flow cytometry is not available, plasma levels may serve as surrogates of T cell cytokine production. We showed that plasma levels of Type 1 cytokines correlated strongly with intracellular expression despite the fact that non-T cells also have the ability to make these cytokines (58, 59). When only 4-color flow cytometry is available, inclusion of all 3 Type 1 cytokines in one color channel for detecting the total Type 1 response may be the most useful. This proposal is supported by observations in HIV infection, where presence of "polyfunctional" CD8+ T cell populations, i.e., HIV-specific CD8+ T cells that co-express multiple cytokines, was associated with better clinical outcome (46). The observation in a mouse model of Leishmania major infection that "polyfunctional" T cell induction is also associated with the best outcome (60) suggests that measurement of these cells may also be important following BCG vaccination, as the mechanisms of immune protection are very similar for Leishmania and mycobacteria. Regardless, measurement of all 3 Type 1 cytokines in one color would still not delineate the complexity of cytokine expression profiles. This may be important, as other studies HIV infection showed that IL-2 expression, rather than IFN- γ or dual expression, correlated with the best clinical outcome (49). We therefore propose that delineation of multiple cytokine-expressing subsets, individually, will be important to investigate protective immunity against tuberculosis, either following natural infection, or following vaccination.

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Figure 1.

Flow cytometric detection of CD4+ T cell cytokine expression in whole blood incubated with BCG for 12 hours, from a single 10-week old BCG-vaccinated infant. The cut-off gates for cytokine expression were determined using unstimulated T cells from whole blood incubated with co-stimulants only (A). Cytokine expression in CD4+ T cells from whole blood incubated with BCG (B), and with SEB (C) is shown. IL-2/TNF-a subset gating was based on patterns of cytokine expression in SEB-stimulated whole blood. Dotplots are gated on CD3+CD4+ T cells, and are representative of 29 infants.



Figure 2.

Cytokine profiles of BCG-specific T cells in 10-week old infants, vaccinated at birth. (A) Frequency of CD4+ and of CD8+ T cells expressing individual Type 1 cytokines following incubation of blood with BCG for 12 hours, in 29 infants. Responses above 0.01% were considered positive. The horizontal line indicates the median and the whiskers the interquartile range. (B) Frequency of BCG-specific CD4+ T cells expressing different combinations of Type 1 cytokines. (C) Frequency of BCG-specific CD8+ T cells expressing different combinations of Type 1 cytokines. (D) Representative staining of intracellular Type 1 cytokines in BCG-specific CD4+ T cells, from a single 10-week old infant. (E) Comparison of frequency of CD4+ and of CD8+ T cells expressing IFN- γ (dark bars) with frequency of T cells expressing IL-2 and/or TNF- α without IFN- γ (light bars), in 29 BCG-vaccinated infants.



Figure 3.

Expression of IL-10 (A) and of IL-4 (C) in CD4+ T cells from whole blood from a single vaccinated infant, incubated with co-stimulants (UNS), BCG or SEB. Frequency of CD4+ T cells expressing IL-10 (B) and of IL-4 (D) following incubation of blood with co-stimulants only (UNS) (light bars) and BCG (dark bars) for 12 hours, in 29 infants.



Figure 4.

Levels of IL-4 and IL-10, in plasma from whole blood incubated with BCG for 7 hours. The horizontal line represents the median. Background cytokine levels were subtracted.



Figure 5.

Phenotype of BCG-specific CD4+ T cells in 10-week old infants, vaccinated at birth. Antigen-specific CD4+ T cells were identified by expression of intracellular IFN- γ (A), IL-2 (B), or both cytokines (C), and the expression of CD45RA, CCR7 and CD27 determined in each case. The plots illustrate the distribution of cytokine-expressing cells (in color; foreground) in relation to the entire CD4+ T cell population (grey; background, also shown in D).



Figure 6.

Frequency of different subsets of BCG-specific CD4+ T cells, based on expression of CD45RA, CCR7 and CD27 among IFN- γ + and IL-2+ (A), and IFN- γ +IL-2+ (B), expressing CD4+ T cells. (C) Frequency of these cells among cytokine-negative, i.e., mostly mycobacteria-non-specific, CD4+ T cells. The horizontal line represents the median and the whiskers the interquartile range. *, p<0.05; ns, p 0.05.

Table I

Association between type 1 cytokine levels in plasma, and frequencies of CD4+ or CD8+ T cells expressing these cytokines, after incubation of whole blood with BCG. A Spearman test was used to assess correlation in 29 infants

	CD4+ T cells expressing cytokine		CD8+ T cells expressing cytokinea	
	r	р	r	р
Plasma IFN-γ	0.6793	< 0.0001	0.6305	0.0004
Plasma IL-2	0.6441	0.0003	0.6013	0.0009
Plasma TNF-a	0.4034	0.01	0.3187	0.1052