

Rapid flow cytometry-based assay for the evaluation of $\gamma\delta$ T cell-mediated cytotoxicity

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Abstract. The effector function of natural killer, lymphokine-activated killer cells and T lymphocytes is commonly evaluated by radioactive chromium-release cytotoxicity assays. In addition to this indirect method, fluorescence assays have been described for the assessment of *in vitro* cell-mediated cytotoxicity. In the present study, target cells were stained with 5-(and-6)-carboxyfluorescein diacetate succinimidyl ester (CFSE), which is a stable integrated fluorescent probe that allows target and effector cells to be distinguished from one another. Staining of target THP-1 cells with 8 μ M CFSE revealed high and stable loading of fluorescence and no effect of the viability of cells. After 4 h of *in vitro* co-culture between $\gamma\delta$ T cells and CFSE-labeled infected or uninfected THP-1 cells, staining with propidium iodide (PI) was performed to distinguish between vital and dead cells. During sample acquisition, target cells were gated on the CFSE positivity and examined for cell death based on the uptake of PI. CFSE and PI double positive cells were recognized as the dead target cells. The percentage of cytotoxicity in the CFSE-gated cell population was calculated by subtracting the value obtained for non-specific PI-positive target cells, which was measured in a control group that did not contain effector cells. The present study describes a simple and convenient assay that is based on the direct quantitative and qualitative analysis of cell damage at a single cell level utilizing a two-color flow cytometric assay. In conclusion, the flow cytometric-based assay described in the current study is a simple, sensitive and reliable tool to determine the cytolytic activity of $\gamma\delta$ T lymphocytes against mycobacteria. Therefore, the present study may provide

valuable information concerning the methods employed to investigate the function of $\gamma\delta$ T cells and potentially other lymphocyte subsets.

Introduction

According to a 2014 report by the World Health Organization (WHO), one-third of the global population is infected with *Mycobacterium tuberculosis* (*M.tb*), the agent responsible for human tuberculosis. Furthermore, WHO estimates that there are ~1.5 million deaths worldwide from this disease annually and an increasing number of drug-resistant cases are reported each year (1). Tuberculosis remains a concerning global health issue, which is partially due to a limited understanding of the details underlying the host protective immune response to the infection.

Human $\gamma\delta$ T cells are a subset of T cells that express the $\gamma\delta$ T cell receptor (TCR) and are a unique lymphocyte population with a specific tissue distribution and antigen (Ag) recognition pathway (2,3). Although these cells are thought to make up only a small proportion (3-10%) of the circulating CD3⁺ T cells in peripheral blood, $\gamma\delta$ T cells have essential roles in host defenses against invasive pathogens (4,5). The involvement of $\gamma\delta$ T cells in the host response to mycobacterial infections was described as early as 1989 (6). The expansion and activation of $\gamma\delta$ T cells was reported during *M.tb* invasion, which indicates the importance of $\gamma\delta$ T cells in the control of infection (7). Consistent with this report, the suppression of $\gamma\delta$ T cells by chronic tuberculosis infection leads to a poor outcome (8).

$\gamma\delta$ T cells participate in early immunity against infection through multiple pathways, including the production of potent, soluble proinflammatory molecules (such as interferon- γ and tumor necrosis factor- α) and chemokines, rapid cytotoxicity or crosstalk with other cell types that are involved (such as natural killer cells, macrophages and dendritic cells) (9). $\gamma\delta$ T cells exhibit different degrees of cytolytic activity in response to various types of pathogens, such as influenza viruses, which include human seasonal H1N1 and avian H5N1 and H9N2 viruses (10-12).. Wallace *et al* (13) reported that $\gamma\delta$ T cells from patients infected with human immunodeficiency virus (HIV) exhibited antiviral potential through their cytolytic

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functions. Therefore, the direct killing of infected cells or invasive pathogens is the most important mechanism of $\gamma\delta$ T cell-mediated immune responses to infection. The cytotoxicity of $\gamma\delta$ T cells involves numerous pathways, including Fas-Fas ligand interactions and the secretion of perforin, granzyme B and granzyme M (14,15). Although defects in these cytotoxic mechanisms lead to the suppression of $\gamma\delta$ T cell activity in various diseases, Poccia *et al* (16) revealed that stimulation with phosphoantigen enhances the function of $\gamma\delta$ T cells and may suppress HIV infection through cell-release antiviral factors, which may become a novel target for effective therapeutic strategies to control the infection.

Based on our understanding of $\gamma\delta$ T cell function and their importance in infectious disease, determining the number of $\gamma\delta$ T cells is not sufficient and a suitable clinical assay for $\gamma\delta$ T cell activity is required. The traditional method for cytotoxicity assays involves radioisotope labelling of target cells with ^{51}Cr , following which the $\gamma\delta$ T cell-specific activity is determined by measuring ^{51}Cr release following co-culture of target cells with the effector ($\gamma\delta$ T) cells (17). This assay is commonly used despite it being associated with certain disadvantages, including low sensitivity and the use of hazardous radiation. Additionally, the method is operationally complex and time consuming. In recent years, several novel assays using flow cytometry have been developed to measure natural killer cytotoxicity without the use of radioactivity (18-19). The objective of the present study was to optimize a rapid flow cytometry-based assay to assess the cytolytic activity of $\gamma\delta$ T cells following *M.tb* infection.

Materials and methods

***M.tb* culture and infection.** In the present study, *M.tb* (H37Ra) (cat no. 93020-5; General Microbiological Culture Collection Center, Beijing, China) was cultured in Sauton's medium supplemented with 0.05% Tween-80 (Hefei Real Biotechnology Co., Ltd., Hefei, China) for 4-6 weeks to late log phase, and the mycobacterial cells were harvested and washed three times with normal saline. Prior to infection, bacteria were incubated with RPMI-1640 (Gibco; Thermo Fisher Scientific, Inc., Waltham, MA, USA) medium and rocked for 30 min at 37°C, clumps were disrupted by multiple passages through a 25-gauge needle, as demonstrated in Fig. 1A. Subsequently, bacteria were resuspended in a physiologic solution at 3×10^8 colony-forming units (CFU)/ml.

To evaluate the phagocytosis of *M.tb* by THP-1 cells, bacteria were labeled with fluorescein isothiocyanate (FITC) as follows: 3×10^8 CFU/ml *M.tb* were incubated with 0, 20, 50 and 100 μg FITC per ml at 37°C for 2 h. Subsequently, labeled bacteria were washed three times with PBS (0.05% Tween-80) to remove unbound FITC. Thereafter, cells were resuspended in RPMI-1640 at 3×10^8 CFU/ml and kept at 4°C. *M.tb* incubated with 0 $\mu\text{g}/\text{ml}$ was used as blank control. During data analysis, values of each group (20, 50 and 100 $\mu\text{g}/\text{ml}$) were obtained by subtracting this background.

Differentiation of THP-1 cells. The THP-1 monocytic leukemia cell line (Shanghai Bogoo Biotechnology Co. Ltd., Shanghai, China) was cultured at 37°C with 5% CO_2 and 95% humidity in RPMI-1640 medium supplemented with 10%

fetal bovine serum (FBS; Gibco; Thermo Fisher Scientific, Inc.), 2 mM L-glutamine, 100 U/ml streptomycin and penicillin. Cells were seeded at a density of $10^5/\text{cm}^2$ and allowed to adhere and differentiate. Monocytic differentiation was induced with phorbol 12-myristate 13-acetate (PMA) at a final concentration of 40 ng/ml for 24 h at 37°C in a humidified atmosphere with 5% CO_2 . To evaluate the expression of CD14 on THP-1 cells, cells were harvested and incubated with FITC anti-human CD14 monoclonal antibody (eBioscience; Thermo Fisher Scientific, Inc.) on ice for 30 min and then washed twice with PBS. Cells were subsequently analyzed by flow cytometry and CellQuest™ software version 3.3 (BD Biosciences, Franklin Lakes, NJ, USA) and WinMDI 2.8 (J. Trotter, The Scripps Research Institute, La Jolla, CA, USA).

Phagocytosis of M.tb by THP-1 cells. Quantitative analysis of phagocytosis was determined by using the fluorescence quenching technique. Briefly, FITC-labeled bacterial suspensions were added to 3×10^5 PMA-differentiated THP-1 cells at different multiplicity of infections (MOIs; 1:1-100:1) and incubated for 1 and 2 h, or incubated for different durations (1-6 h) at an MOI of 10:1, at 37°C. Those in blank control group were treated with no bacteria (MOI 0) or no incubation duration (0 h). Subsequently, cultures were washed and resuspended in 0.125% trypan blue (pH 4.4) for 3 min to quench the fluorescence of non-ingested, but membrane-associated bacteria. The percentage of cells that exhibited positive FITC expression was analyzed as the phagocytosis rate by subtracting the value of control group.

Preparation of M.tb heat-treated Ag. (M.tb-HAg). *M.tb*-HAg were prepared according to the methods described in a previous report (20). Briefly, *M.tb* were cultured, collected, washed, resuspended in double volumes of ultra-pure water and heated at 120°C for 30 min. The supernatants from heat-treated *M.tb* cells were harvested and concentrated to 1 mg/ml prior to use for stimulating $\gamma\delta$ T cells.

Expansion of effective $\gamma\delta$ T cells. Peripheral blood mononuclear cells were isolated from the freshly heparinized venous blood of healthy volunteers by density gradient centrifugation using Ficoll-Hypaque (21). Ethical approval was obtained from the ethical committee of Bengbu Medical College, Bengbu, China. Each healthy donor signed informed consent according to institutional guidelines. Written informed consent was obtained from each healthy donor according to federal and institutional guidelines. Cells were maintained at 1.5×10^6 cells/ml, with 1 ml/well in 24-well cell culture plates in complete RPMI-1640 medium supplemented with 10% (v/v) FBS and 50 $\mu\text{g}/\text{ml}$ gentamycin (Beijing Biodee Biotechnology Co., Ltd., Beijing, China). These cells were cultured at 37°C in 5% CO_2 in the presence of 50 U/ml recombinant human interleukin (IL)-2 (rhIL-2; Biomics Biotechnologies Co., Ltd., Nantong, China) and 5 $\mu\text{g}/\text{ml}$ *M.tb*-HAg for 7-10 days and fed with fresh medium and cytokines every 3-4 days. The *M.tb*-HAg and rhIL-2-stimulated T cells were described as *M.tb*-HAg activated T cells, which were stained for surface markers with FITC anti-human CD3 and phycoerythrin (PE) anti-human TCR $\gamma\delta$. Both fluorochrome-conjugated monoclonal antibodies were purchased from eBioscience; Thermo

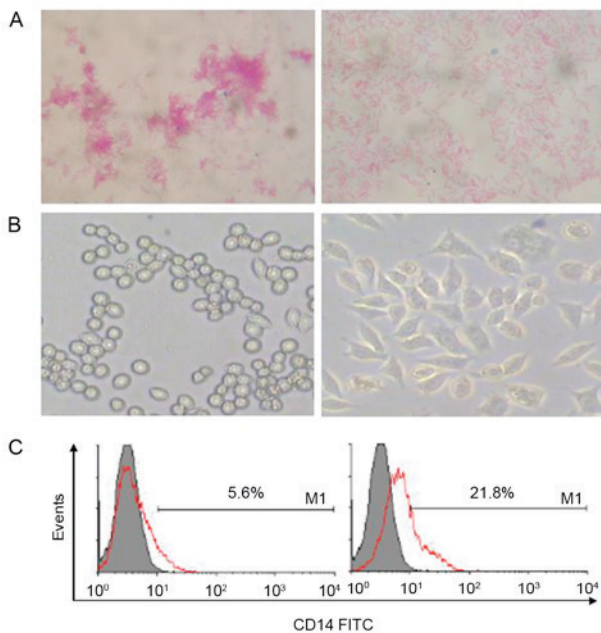


Figure 1. Alteration of morphology and phenotype in PMA-differentiated THP-1 cells. (A) Aggregate (left) and dispersed (right) *M.tb* stained with routine acid-fast staining (magnification, x1,000). (B) Morphology of THP-1 (left) and PMA-differentiated THP-1 (right) cells (magnification, x400). (C) Phenotype of THP-1 cells. THP-1 (left) and PMA-differentiated THP-1 (right) cells were cultured and stained with FITC anti-human CD14 antibody (red) or an isotype control antibody (gray) and examined by flow cytometry. The results are representative of three independent experiments that gave similar results. PMA, phorbol 12-myristate 13-acetate; *M.tb*, *Mycobacterium tuberculosis*; FITC, fluorescein isothiocyanate.

Fisher Scientific, Inc. $\gamma\delta$ T cells account for ~70-85% of the total *M.tb*-HA_g activated T cells, which were used as effective $\gamma\delta$ T cells. Effector cells were adjusted to a final concentration of $1-2 \times 10^6$ cells/ml.

Infection of THP-1 cells. For infection of cells, 1×10^6 THP-1 or PMA-differentiated THP-1 cells were washed three times and replenished with culture medium without antibiotics and infected with *M.tb* at an MOI of 50 bacteria per cell. Bacteria were centrifuged onto THP-1 or differentiated THP-1 cells at $400 \times g$ for 5 min at $4^\circ C$. After a period of 2 h at $37^\circ C$, non-ingested bacteria were removed by extensive washing with PBS and cells were collected and used as targets in cytotoxicity experiments. Meanwhile, Ziehl-Nielson acid-fast staining and laser scanning confocal microscopy (LSCM) was used to confirm the presence of intracellular bacteria. THP-1 or PMA-differentiated THP-1 cells (1×10^6) were seeded on coverslips and co-cultured with *M.tb* (for Ziehl-Nielson acid-fast staining) or FITC-labeled *M.tb* (for LSCM) at an MOI of 50:1 and incubated for 2 h at $37^\circ C$. Non-ingested bacteria were gently removed in warm PBS containing 0.5% bovine serum albumin (BSA) (Hefei Real Biotechnology Co., Ltd.). For analysis by confocal microscopy, cells on glass coverslips were fixed in 2% formaldehyde for 20 min on ice, then washed with PBS and blocked with 5% BSA for 20 min at room temperature. Subsequently, the samples were incubated with PE anti-human Toll-like receptor-2 (eBioscience; Thermo Fisher Scientific, Inc.) for 30 min at room temperature, then rinsed gently in distilled water for ~1 min and mounted on

microscopic slides. Images were captured using a laser confocal microscope (CI-Digital Eclipse, Nikon, Tokyo, Japan).

Cytotoxicity assay. To confirm the cell viability prior to cytotoxicity testing, Fluorescein diacetate (FDA) was used as a vital stain to assay membrane integrity (cell viability). 1×10^5 cells ($\gamma\delta$ T cells, THP-1 cells or THP-1 cells infected with *M.tb*) were incubated at room temperature for 5 min in the presence of 50 ng/ml FDA, followed by analysis by flow cytometry. The cytotoxicity assay was performed when cell viability was >85%. Target THP-1 cells were labeled with carboxyfluorescein diacetate succinimidyl ester (CFSE; Invitrogen; Thermo Fisher Scientific, Inc.) as follows: CFSE (2, 4 and $8 \mu\text{mol/l}$) was added to the target cell suspension ($1 \times 10^6/\text{ml}$) at $37^\circ C$ in 5% CO_2 for 10 min. Following labeling, the cells were washed and analyzed in a flow cytometer. CFSE-labeled target cells were resuspended in RPMI-1640 to a final concentration of $1 \times 10^5/\text{ml}$ for further investigation.

The cytotoxic function of $\gamma\delta$ T cells was evaluated by adding CFSE-labeled infected or uninfected THP-1 cells to yield effector:target (E:T) ratios of 20:1, 10:1, 5:1 and 1:1. Control tubes were also assayed to determine the spontaneous cell death. Tubes were gently mixed, centrifuged at $120 \times g$ for 2 min at $4^\circ C$ and incubated at $37^\circ C$ in 5% CO_2 . After 4 h of co-culture, cells were stained with $4 \mu\text{l}$ propidium iodide (PI; $100 \mu\text{g/ml}$) and were placed on ice for 5 min. The fluorescence intensity of cultured cells was measured by flow cytometry. During sample acquisition, a 'target cell gate' was set on the CFSE-positive cell population using a FL1-histogram, and ~2,000 target cells were collected. For data analysis, the CFSE-positive target cells were examined for cell death by uptake of PI. CFSE and PI double positive cells were considered to be dead target cells. The percentage of cytotoxic activity was subsequently calculated using the following equation:

$$\text{Cytotoxicity (\%)} = \frac{\text{Target cell deaths} - \text{spontaneous deaths}}{100 - \text{spontaneous deaths}} \times 100$$

The cytotoxicity of $\gamma\delta$ T cells against THP-1 cells was assessed also by measuring cytosolic lactate dehydrogenase (LDH) release into culture supernatants using a LDH cytotoxicity assay kit (Hefei Bomei Biotechnology Co., Ltd., Hefei, China).

Statistical analysis. Data obtained from independent experiments are presented as the mean \pm standard deviation. Statistical analysis was performed using one-way analysis of variance and the Student-Newman-Keuls test. Data from experiments in Fig. 5 were analyzed using Student's t-test, and the correlation between the cytotoxic activity obtained by flow cytometry-based and LDH cytotoxicity assays was estimated by the nonparametric Spearman's Rank correlation test. $P < 0.05$ was considered to indicate a statistically significant difference. The statistical software package SPSS 15.0 (SPSS, Inc., Chicago, IL, USA) was used for all the data analysis.

Results

Alterations in the morphology and phenotype of PMA-differentiated THP-1 cells. As demonstrated in Fig. 1B,

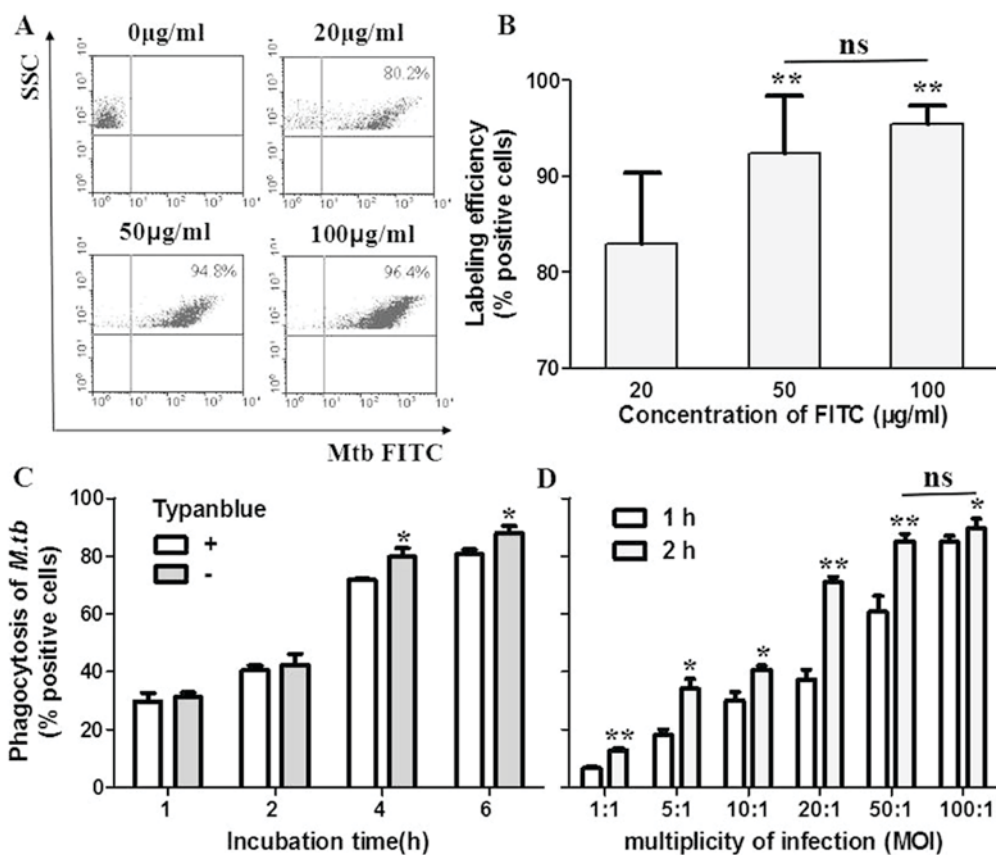


Figure 2. Phagocytosis of FITC-labeled *M.tb* by THP-1 cells. (A) Representative FACS dot plots demonstrate labeling efficiency at different concentrations of FITC. The numbers represent the percentage of FITC positive cells. (B) Mean labeling efficiency at different FITC concentrations in four independent experiments. ** $P < 0.01$ vs. 20 $\mu\text{g/ml}$ FITC. (C) Effect of trypan blue on phagocytic rate. PMA-differentiated THP-1 cells were exposed to FITC-labeled *M.tb* at the ratio of 10:1 for 1-6 h. Following incubation, surface fluorescence emitted by non-internalized bacteria was quenched by exposure to trypan blue. * $P < 0.05$ vs. trypan blue-quenching group at same time-point. (D) PMA-differentiated cells were incubated with increasing concentrations of *M.tb* for 1 or 2 h. The percentage of cells phagocytosing *M.tb* was measured by FACS analysis. The results are representative of two independent experiments. * $P < 0.05$ and ** $P < 0.01$ vs. 1 h at same MOI. FITC, fluorescein isothiocyanate; *M.tb*, *Mycobacterium tuberculosis*; FACS, fluorescence-activated cell sorting; PMA, phorbol 12-myristate 13-acetate; MOI, multiplicity of infection.

the shape of untreated THP-1 cells was round and cells did not adhere to the cell culture plate surface (left). In the presence of the common macrophage differentiation factor PMA for 24 h, the cells became flat and amoeboid in shape, and adhered to the bottom of the plate (Fig. 1B, right). Cell cultures treated with PMA (Fig. 1B, right) consisted of bigger cells at a lower density compared with the cultures that were not treated with PMA (Fig. 1B, left).

Subsequently, the present study evaluated whether PMA was able to induce the expression of CD14, which is considered to be a potential receptor for *M.tb*. Flow cytometry analysis revealed that CD14 was upregulated with macrophage differentiation (Fig. 1C). Upon treatment with PMA, a substantial increase in the proportion of CD14⁺ cells (21.8%; Fig. 1C, right) was observed compared with untreated THP-1 cells (5.6%; Fig. 1C, left; $P < 0.05$).

Phagocytosis of *M.tb* by THP-1 cells. Flow cytometry and FITC-labeled bacteria were used to measure the phagocytosis of *M.tb* by THP-1 cells. *M.tb* labeled with FITC at different concentrations were easily visualized by fluorescence microscopy, and no loss of viability was observed. The labeling efficiency of *M.tb* was determined by flow cytometry and was increased with increasing FITC concentrations. Of the various

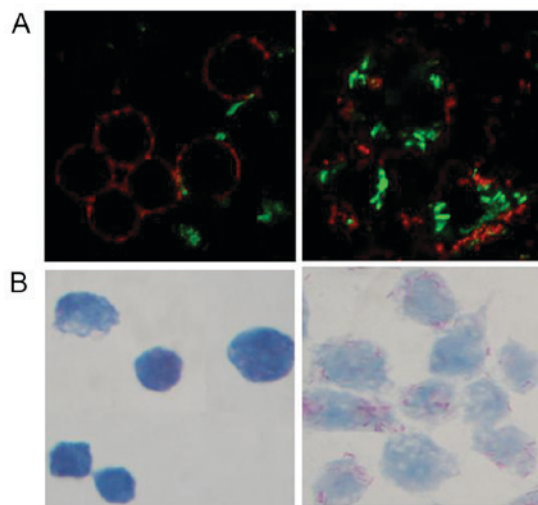


Figure 3. Use of microscopy to confirm the presence of intracellular bacteria following phagocytosis. (A) Fluorescence micrographs of THP-1 (left) and PMA-differentiated THP-1 (right) cells with phagocytosed fluorescein isothiocyanate-labeled *M.tb* and toll-like receptor 2-phycoerythrin antibody staining. Intracellular bacteria fluoresce green while the cell surface remains red (magnification, $\times 1,000$). (B) Light micrograph of THP-1 (left) and PMA-differentiated THP-1 (right) cells with phagocytosed *M.tb* stained with routine acid-fast staining (magnification, $\times 1,000$). PMA, phorbol 12-myristate 13-acetate; *M.tb*, *Mycobacterium tuberculosis*.

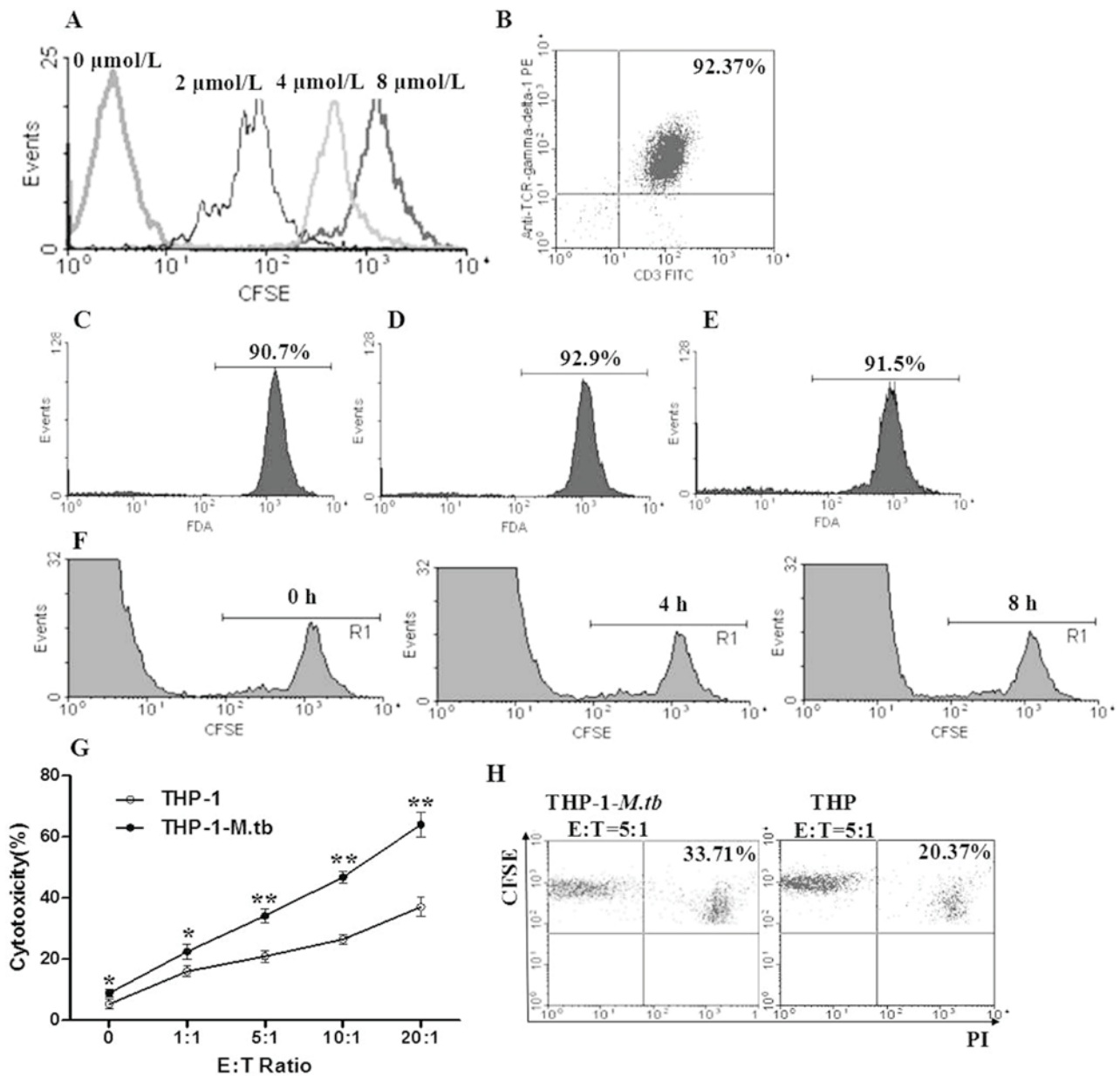


Figure 4. Flow cytometric analysis of $\gamma\delta$ T cell cytotoxicity assay. (A) Histogram overlay demonstrates flow cytometric analysis of unlabeled target cells and target cells immediately after CFSE labeling. (B) Representative flow cytometry panel demonstrates the phenotype of recombinant human interleukin-2- and *M.tb* heat-treated antigen-expanded T cells in which $\gamma\delta$ T cells account for ~92.35% of total cells. Histogram profiles of (C) $\gamma\delta$ T cells (effectors), (D) THP-1 cells and (E) *M.tb*-infected THP-1 cells (targets) treated with FDA. Samples with 50 ng/ml FDA were incubated at room temperature for 5 min prior to analysis by flow cytometry to confirm the cell viability. (F) Histogram indicates the level of target cell division, as assessed by examining differences in CFSE fluorescence between cells immediately after labeling and at 4 and 8 h after labeling with CFSE. (G) Evaluation of $\gamma\delta$ T cell cytotoxicity at different E:T ratios by the flow cytometry-based assay. Data are presented as the mean cytotoxicity value \pm standard deviation from one representative experiment of four independent experiments. * $P < 0.05$ and ** $P < 0.01$ vs. THP-1 cells at same E:T ratio. (H) Representative fluorescence-activated cell sorting dot plot represents $\gamma\delta$ T cell activity at an E:T ratio of 5:1 and dead target cells appear in the upper right quadrant. CFSE, 5-(and-6)-carboxyfluorescein diacetate succinimidyl ester; *M.tb*, *Mycobacterium tuberculosis*; E:T, effector:target; TCR, T cell receptor; PE, phycoerythrin; FITC, fluorescein isothiocyanate; PI, propidium iodide; FDA, fluorescein diacetate.

FITC concentrations employed, the maximum proportion of FITC positive cells was obtained using 50 $\mu\text{g/ml}$ (92%) and 100 $\mu\text{g/ml}$ (95%) FITC (Fig. 2A and B). No significant difference between the labeling efficiency at the concentrations of 50 and 100 $\mu\text{g/ml}$ was observed. Therefore, bacteria were labeled with 50 $\mu\text{g/ml}$ FITC for further experiments.

As demonstrated in Fig. 2C and D, THP-1 macrophages were exposed to bacteria at an MOI of 10:1 for different durations (1, 2, 4 and 6 h; Fig. 2C) or a range of bacterial concentrations

(MOI 1:1-100:1) for 1 and 2 h (Fig. 2D). The maximal loading of THP-1 cells with bacteria was obtained at the ratio of 100:1 at 2 h; however, a ratio of 50:1 at 2 h was chosen as this MOI was associated with less cytotoxicity ($85.20 \pm 2.59\%$; Fig. 2D) or 10:1 for 6 h ($80.90 \pm 1.47\%$). Following quenching of surface FITC fluorescence with trypan blue, the frequency of positivity decreased by 1% in the 1 h treatment group ($P > 0.05$) and 7% in the 6 h treatment group ($P < 0.05$; Fig. 2C). Routine acid-fast staining and LSCM confirmed the presence of intracellular

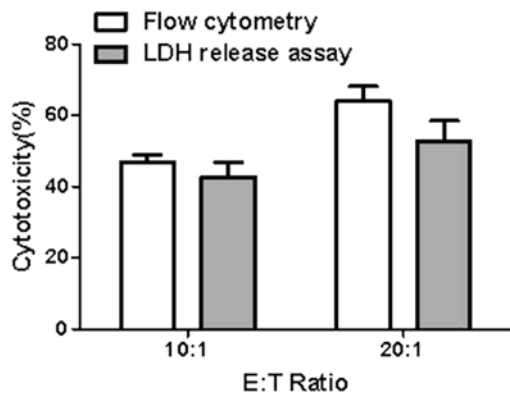


Figure 5. Comparison of $\gamma\delta$ T cell cytotoxicity as assessed by flow cytometry-based and LDH release assays. The values for $\gamma\delta$ T cell cytotoxicity obtained at various E:T ratios using flow cytometry and LDH release assays were compared. The mean of triplicate results is presented for E:T ratios of 10:1 and 20:1. LDH, lactate dehydrogenase; E:T, effector:target.

bacteria. Representative photomicrographs of infected cells with *M.tb* are presented in Fig. 3.

Optimization of target cell labeling. For THP-1 cell (target cells) staining, three different concentrations (2, 4 and 8 μM) of CFSE were employed and compared. Labeling with these concentrations of CFSE all allowed homogenous staining of >90% of THP-1 cells. As the fluorescence intensity of CFSE per cell is naturally reduced upon cell division, an optimal separation between stained and unstained cells was observed at the concentration of 8 μM CFSE (Fig. 4A).

Detection of $\gamma\delta$ T cell cytotoxicity. Effective $\gamma\delta$ T cells were collected from peripheral blood and the purity (CD3-TCR $\gamma\delta^+$) was >90% on day 10 after incubation (Fig. 4B). Prior to using in cytotoxicity tests, cell labeling with FDA confirmed that cell viability was >85% for $\gamma\delta$ T cells, THP-1 cells and THP-1 cells infected with *M.tb* (Fig. 4C-E, respectively).

The performance of $\gamma\delta$ T cells in the cytotoxic assay was determined by adding CFSE-labeled infected or uninfected THP-1 cells at different E:T ratios. The loss of intensity was investigated by comparing freshly stained target cells with target cells that were stained and incubated for different durations, and no decrease in fluorescence intensity was observed up to 8 h of incubation (Fig. 4F). When $\gamma\delta$ T cells were co-cultured with target cells for 4 h at ratios between 1:1 and 20:1, the cytotoxicities of $\gamma\delta$ T cells against targets increased from 16.05±1.74% to 64.06±4.07% (Fig. 4G). As demonstrated in Fig. 4G, the resulting cytotoxic activities increased with increasing E:T ratios, and, when infected THP-1 cells were used as targets, the maximum $\gamma\delta$ T cell cytotoxicity reached 64.06±4.07%. Comparing between the assays that employed infected and uninfected THP-1 cells, the cytotoxic activities of $\gamma\delta$ T cells against infected THP-1 cells were significantly higher compared with against uninfected THP-1 cells at each E:T ratio ($P < 0.05$). Fig. 4H illustrated that samples were analyzed by 2-color flow cytometry with CFSE gating to determine the cytotoxic activities at an E:T ratio of 5:1. CFSE and PI double positive cells (the upper right quadrant) were considered to be dead target cells.

The flow cytometry-based cytotoxicity assay was compared with the LDH cytotoxicity assay, which were run in parallel. All results obtained at E:T ratios 10:1 and 20:1 are presented in Fig. 5. The resulting cytotoxic activity was similar in both assays. In the flow cytometry-based cytotoxicity assay, the mean cell lysis at 10:1 and 20:1 ratios was 47.74±2.94% and 64.07±5.35%, respectively, while the LDH cytotoxicity assay detected cell lysis with means of 42.59±5.32% and 52.78±6.39%, respectively. The results of the two assays were significantly correlated ($r = 0.886$; $P < 0.05$). The responses obtained at ratios 10:1 were, in general, lower compared with those at 20:1, in the flow cytometry-based cytotoxicity assay (1.3-fold) and the LDH cytotoxicity assay (1.2-fold).

Discussion

$\gamma\delta$ T cells are a class of unconventional T cells that connect the innate and adaptive immune systems and constitute a crucial defense against microbial agents and tumors. The evaluation of $\gamma\delta$ T cells is essential whenever investigation into inflammation, autoimmunity or allergy occurs. Participation in early immunity against *M.tb* is also a feature of $\gamma\delta$ T-cells, and the aim of the present study was to describe the optimization of a flow cytometry-based $\gamma\delta$ T-cell assay developed for the assessment of *M.tb*-specific cytolytic activity.

Macrophages, a primary host cell niche for the intracellular growth and persistence of *M.tb*, are responsible for the activation of protective immune responses (22). As the *in vitro* growth and expansion of primary tissue macrophages is notoriously difficult, certain cell lines are commonly used to model macrophage function. Early studies indicated that THP-1, a leukemia cell line derived from a patient with acute monocytic leukemia, exhibit primary monocyte and macrophage morphological and functional properties. They are highly plastic and homogeneous, as a consequence of a minimal degree of variability in the cell phenotype (23). PMA is a common stimulus employed to induce THP-1 differentiation. The present study demonstrated that treating THP-1 cells with 40 ng/ml PMA for 24 h drove cells towards full macrophage differentiation, including the expansion of the cytoplasm, a higher rate of cell adherence, increased expression of the cell surface marker CD14 and high phagocytic capacity. These results indicate that, as reported in previous studies (24-26), that PMA-treated cells may represent a valuable model with which to investigate tissue macrophages.

The phagocytic capacity of macrophages is central to their function (27). In the current study, PMA-treated THP-1 cells exhibited efficient phagocytosis of *M.tb* (H37Ra) and the maximal percentage of phagocytosis reached 89%. By contrast, undifferentiated cells incubated with identical MOIs of bacteria exhibited only marginal internalization of bacteria. When analyzing the results based on the phagocytosis, the bacteria:cell ratio of 50:1 for 2 h was suitable for the subsequent preparation of target cells employed in cytotoxicity assays. THP-1 monolayers remained stable until the end of the experiments.

The traditional 'gold standard' for the determination of cytotoxicity is the ^{51}Cr release assay. However, this method is not favored clinically due to certain limitations, including

the use of radioactive reagents. Over recent years, flow cytometry-based assays have been developed in an attempt to avoid the problems associated with the ^{51}Cr -assay (28-29). Flow cytometric methods enable rapid analysis to be conducted at the single cell level and may be more convenient for routine application rather than indirectly by the release of a preloaded marker. At present, the flow cytometry-based assays employ forward and side scatter characteristics for the discrimination of different cell populations and distinction between dead and viable cells, or use differential labeling of target cells with various fluorescent dyes (19,30).

In the assay described in the present study, the discrimination of targets from effectors entirely depends on the cell labeling, therefore, it is crucial to stain the target cells with a fluorescent dye that is sensitive, does not leak into target cells and does not affect the cell viability. The low cost of CFSE with a simple and rapid labeling protocol makes it a convenient dye for clinical use in laboratories, and, it is suitable for use in combination with a second fluorescent label, such as PI, to determine the target cell viability. This was therefore selected for the labeling of target cells in the current study. CFSE and PI are readily distinguished by flow cytometry using the FL1- and FL3-channels, respectively, without spectral overlap. The specific lysis was measured via the uptake of PI in the CFSE-positive target cells following incubation with effector cells.

In conclusion, the present study describes a convenient assay that is based on the direct quantitative and qualitative analysis of cell damage at a single cell level utilizing a two-color flow cytometric assay. The final format of this simple flow cytometry-based assay for the determination of the cytolytic activity of $\gamma\delta$ T lymphocytes against mycobacterium will employ rhIL-2- and *M.tb*-HAg-expanded T cells as effective $\gamma\delta$ T cells, use an MOI of 50:1 for 2 h in the preparation of target cells, require 8×10^4 target cells per sample and include the collection of 2,000 target events per sample. Flow cytometry for the determination of $\gamma\delta$ T cell activity is a promising alternative to the conventional ^{51}Cr release assay. The present study may provide valuable information concerning the methods for the investigation of $\gamma\delta$ T cell function, and potentially the function of other lymphocyte subsets.

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