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Research paper

## Qualification of a whole blood intracellular cytokine staining assay to measure mycobacteria-specific CD4 and CD8 T cell immunity by flow cytometry<sup>☆</sup>

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## ABSTRACT

**Background:** Qualified or validated assays are essential in clinical trials. Short-term stimulation of whole blood and intracellular cytokine staining assay is commonly used to measure immunogenicity in tuberculosis vaccine clinical trials. Previously, the short-term stimulation process of whole blood with BCG was optimized. We aimed to qualify the intracellular cytokine staining process and assess the effects of long-term cryopreservation. Our hypotheses were that the assay is robust in the measurement of the mycobacteria-specific T cells, and long-term cryopreservation of fixed cells from stimulated whole blood would not compromise reliable measurement of mycobacteria induced CD4 T cell immunity.

**Methods:** Whole blood from healthy adults was collected in sodium heparinized tubes. The blood was left unstimulated or stimulated with mycobacterial antigens or mitogens for 12 h. Cells were harvested, fixed and multiple aliquots from each participant cryopreserved. Later, mycobacteria-specific CD4 and CD8 T cells expressing IFN- $\gamma$ , TNF- $\alpha$ , IL-2 and IL-17 were quantitated by flow cytometry. Assay performance characteristics evaluated included limit of quantification and detection, reproducibility, precision, robustness, specificity and sensitivity. To assess the effects of long-term cryopreservation, fixed cells from the stimulated bloods were analysed one week post-cryopreservation and at 3-month intervals over a 3-year period.

**Results:** The limit of quantification for the different cytokines was variable: 0.04% for frequencies of IFN- $\gamma$ - and IL-2-expressing T cells and less than 0.01% for TNF- $\alpha$ - and IL-17-expressing T cells. When measurement of the mycobacteria-specific T cells was assessed at levels above the detection limit, the whole blood intracellular cytokine assay showed high precision that was operator-independent.

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The assay was also robust: variation in staining conditions including temperature (4 °C or 20–23 °C) and time (45, 60 or 90 min) did not markedly affect quantification of specific T cells. Finally, prolonged periods of cryopreservation also did not significantly influence quantification of mycobacteria-specific CD4 T cells.

**Conclusions:** The whole blood intracellular cytokine assay is robust and reliable in quantification of the mycobacteria-specific T cells and is not significantly affected by cryopreservation of fixed cells.

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## 1. Introduction

Control of the global tuberculosis (TB) pandemic, which still has significant morbidity with more than 8 million people getting the disease every year (World Health Organization, 2013), is an urgent priority. Epidemiological modelling suggests that elimination of TB can only be achieved with an effective vaccination strategy, coupled with better diagnosis and more effective treatment of persons infected with *Mycobacterium tuberculosis* (*M.tb*) and with TB disease (Abu-Raddad et al., 2009; Dye et al., 2013). Immunization with Bacille Calmette-Guerin (BCG), the only licensed TB vaccine, confers protection against severe forms of TB in infants, such as miliary TB and TB meningitis, however efficacy against adult and childhood pulmonary disease is variable and mostly poor (Evans et al., 2013). A more efficacious vaccine is urgently needed and 16 new TB vaccines are currently in clinical testing (Scriba et al., 2012; Day et al., 2013; Abel et al., 2010a; Khader et al., 2007).

Measurements of vaccine immunogenicity, or vaccine take, in preclinical development and human trials routinely include quantitation of antigen-specific Th1 and Th17 cells (Ota et al., 2011; Abel et al., 2010b; van Dissel et al., 2010). A number of distinct immunological assays, among them IFN- $\gamma$  ELISPOT, intracellular cytokine staining (ICS) assays and ELISA-based quantification of IFN- $\gamma$ -release, have been used to measure immunogenicity of the different TB vaccine candidates (Hanekom et al., 2004). Results of outcomes generated by these different methodologies could be usefully compared only if qualified or validated assays are used. Until this is achieved it is difficult to include immunogenicity criteria when gauging the potential value of a given vaccine against those of other candidates. There is a clear and urgent need to optimize, standardize, qualify and validate assays that are suitable for measuring antigen-specific T cell responses.

We previously optimized a whole blood (WB) ICS assay for measurement of antigen-specific Th1 responses in clinical studies (Hanekom et al., 2008), and have applied this WB-ICS assay to evaluate immunogenicity in many clinical studies of TB vaccines in infants and adults ((FDA) FaDA, 1996). Advantages of the WB-ICS assay over PBMC-based assays include a requirement for smaller blood volumes, and immediate antigen stimulation of fresh cells leading to less cell death and lower sensitivity of fixed cells to freezing and thawing procedures ((FDA) FaDA, 1996). This assay is thus suitable for resource-limited settings where TB, HIV/AIDS and malaria are endemic and are the ideal sites for clinical trials.

Here we describe qualification of the WB-ICS assay readout by flow cytometry. Assay qualification precedes assay validation. Validation is defined as an evaluation of the method on its fitness for the intended applications (Standards USPatNfCl, 2007). The assay validation process involves evaluating

performance characteristics of several parameters to ensure that the assay limitations are known when applied to measure predefined outcomes. Assay validation parameters include, among others, accuracy, specificity, limit of detection, reproducibility or inter- and intra-assay co-efficient of variation (CV), precision, linearity and robustness (Hanekom et al., 2004). Establishing accuracy (how close the measured value is to the true value (Hanekom et al., 2004)) in ICS assays is impractical because the true value is unknown. We aimed to qualify the WB-ICS assay and established some of the performance characteristics of the WB-ICS assay.

## 2. Methods

### 2.1. Study participants and blood collection

Healthy adults were recruited at the Institute of Infectious Disease and Molecular Medicine (IDM) of the University of Cape Town in South Africa, or at the South African TB Vaccine Initiative (SATVI) field site in Worcester, 110 km from Cape Town. All participants reported BCG vaccination at birth. The study protocol was approved by the University of Cape Town Research Ethics Committee, and all procedures adhered to the guidelines of the National Health Research Ethics Council. Good clinical practice guidelines were adhered to including written informed consent. Venous blood was collected from study participants in sodium heparin tubes and processed within 60 min.

### 2.2. Antigens used in WB-ICS

Bacillus Calmette Guerin (BCG) Danish 1331 (Statens Serum Institut, Copenhagen) was reconstituted in the vaccine vial with RPMI and used at a final concentration of  $1.2 \times 10^6$  CFU/mL of blood, a dose previously optimized (Hanekom et al., 2004). Purified protein derivative (PPD, Statens Serum Institut, Copenhagen) was used at a final concentration of 10  $\mu$ g/mL. In some experiments, peptide pools spanning ESAT-6 and CFP-10 or Ag85A mycobacterial proteins (15-mers, overlapping by 10 amino acids, each at 2  $\mu$ g/mL; Peptide Protein Research Ltd.) were used. For positive controls, stimulation was done with either Staphylococcal enterotoxin B (SEB, Sigma) or Phytohemagglutinin (PHA, HA16, Sigma), at final concentrations of 10  $\mu$ g/mL or 5  $\mu$ g/mL, respectively, as previously described (Scriba et al., 2011).

### 2.3. Short-term whole blood stimulation and cryopreservation

One millilitre of whole blood, pipetted into Sarstedt tubes, was either left unstimulated (negative control) or stimulated with mycobacterial antigens (as described above), or positive

156 controls of SEB or PHA (either but not both positive controls), in  
 157 the presence of co-stimulatory antibodies (anti-CD28 and anti-  
 158 CD49d, each at 1 µg/mL; BD Biosciences). These co-stimulatory  
 159 antibodies have been shown to increase cytokine expression in  
 160 specific T cells (Scriba et al., 2011). Blood was incubated at 37 °C  
 161 for 12 h, and Brefeldin-A (Sigma-Aldrich, 10 µg/mL) was added  
 162 during the last 5 h of incubation. The blood was then harvested  
 163 with EDTA (Sigma, 2 µM) red blood cells were lysed and white  
 164 blood cells fixed with FACS lysing solution (BD Biosciences).  
 165 Fixed white cells were pelleted and multiple aliquots were  
 166 cryopreserved with 10% DMSO (Sigma) in 40% foetal calf serum  
 167 (BioWest) in RPMI at –190 °C in the vapor phase of liquid  
 168 nitrogen.

#### 169 2.4. Thawing and permeabilisation of cryopreserved fixed cells

170 Cryovials containing the stimulated, fixed and frozen white  
 171 cells from whole blood were retrieved from liquid nitrogen  
 172 tanks and thawed in a water bath at 37 °C for 2 min. Thawed  
 173 cells were transferred from cryovials to labelled tubes contain-  
 174 ing 2 mL of phosphate buffered saline (PBS, BioWhittaker).  
 175 Thereafter the cells were centrifuged at 215 g for 5 min. Next,  
 176 the cells were permeabilised by adding 2 mL Perm/Wash  
 177 solution (BD Biosciences) and incubated at room temperature  
 178 for 10 min (unless specific incubation temperatures were  
 179 investigated).

#### 180 2.5. Intracellular cytokine staining (ICS) and flow cytometry

181 Thawed cryopreserved fixed cells were washed in PBS and  
 182 immediately stained with cocktails of monoclonal antibodies  
 183 for 60 min at 4 °C, unless otherwise indicated. Two different  
 184 flow cytometry antibody panels were used: One monoclonal  
 185 antibody panel was for multiparameter flow cytometry,  
 186 utilizing a BD LSR II cytometer. For these experiments, the  
 187 cells were thawed, permeabilised in BD Perm/Wash buffer and  
 188 stained with previously optimized antibody-fluorochrome  
 189 combinations to the following markers: CD3-PacBlue (BD  
 190 Biosciences, clone MOPC-21), CD4-QDot605 (Invitrogen,  
 191 S3.5), CD8-PerCPy5.5 (BD Biosciences, SK1), IFN-γ-Alexa700  
 192 (BD Biosciences, B27), TNF-α-PeCy7 (eBioscience, Mb11), IL-2-  
 193 FITC (BD Biosciences, 5344.111), IL-17-Alexa647 (eBioscience,  
 194 SCPL1362) and the Ki67-PE (BD Biosciences, B1). Cytometer  
 195 Setting and Tracking (CST) beads (BD Biosciences) were  
 196 acquired before each experiment to ensure that cytometer  
 197 parameters remained consistent across all experiments.  
 198 Stained samples were acquired with a standard stopping gate  
 199 set at 200,000 CD3 lymphocytes. Single stained and negative  
 200 compensation beads (BD Biosciences) were acquired for each  
 201 experiment, before sample acquisition, and used to calculate  
 202 the compensation matrix.

203 To measure effects of long-term cryopreservation on ICS  
 204 outcomes in fixed white blood cells, a second monoclonal  
 205 antibody panel comprising of CD4-APC (SK3) and IFN-γ-PE  
 206 (25723.11; both from BD Biosciences) was acquired on a  
 207 FACSCalibur (BD Biosciences). For these experiments, cells  
 208 were thawed, permeabilized in BD Perm/Wash buffer and  
 209 stained as indicated above before acquisition. At least 40,000  
 210 CD4 T cells were acquired.

#### 211 2.6. IFN-γ ELISpot assay

212 We compared frequencies of IFN-γ expressing cells detect-  
 213 ed by WB-ICS and IFN-γ ELISpot assay from samples collected  
 214 in a previously completed clinical trial of the candidate TB  
 215 vaccine, MVA85A (Mazurek et al., 2010). We analysed data  
 216 from a subset of 36 healthy infants enrolled into the TB014 trial,  
 217 who received a single intradermal vaccination of  $5 \times 10^7$  pfu of  
 218 MVA85A (clinicaltrials.gov NCT00679159). The WB-ICS assay  
 219 was performed as described above. Whole blood and PBMC  
 220 were stimulated in parallel with a single pool of peptides  
 221 spanning the Ag85A protein (15-mers, overlapping by 10  
 222 amino acids, each at 2 µg/mL; Peptide Protein Research Ltd.).  
 223 For ELISpot assay, medium alone served as negative control and  
 224 PHA, (10 µg/mL) as a positive control. ELISpot plates, contain-  
 225 ing  $3 \times 10^5$  peripheral blood mononuclear cells (PBMC) per  
 226 well, were incubated with antigens for 18 h at 37 °C and  
 227 developed according to the manufacturer's protocol  
 228 (Mabtech), as previously described (Weinberg et al., 2000;  
 229 Nomura et al., 2000). Assays were performed in duplicate wells  
 230 and the average (with background subtracted) was used for  
 231 analysis.

#### 232 2.7. QuantiFERON-TB Gold In-Tube assay (QFT)

233 We also compared frequencies of IFN-γ expressing CD4 T  
 234 cells detected by ICS upon blood stimulation with ESAT-6 and  
 235 CFP-10 peptide pools with levels of antigen-specific IFN-γ  
 236 release measured by the validated QFT (Mazurek et al., 2010).  
 237 The QFT was performed according to the manufacturer's  
 238 protocol (Cellestis).

#### 239 2.8. WB-ICS assessments

240 To assess the lower limit of quantification (LOQ) of the WB-  
 241 ICS assay, blood from two healthy adult volunteers was  
 242 stimulated with BCG or left unstimulated. Stimulated cells  
 243 were serially diluted with autologous unstimulated cells,  
 244 ranging from 2 to 4096-fold. Unstimulated, BCG-stimulated  
 245 and serially diluted samples were then stained with an 8-colour  
 246 antibody panel, as described above, and cytokine expression by  
 247 CD4 and CD8 T cells was analyzed on a LSR II cytometer.  
 248 Expected frequencies were derived by dividing the frequencies  
 249 of BCG-specific cytokine-expressing CD4 and CD8 T cells in the  
 250 undiluted samples by the dilution factor.

251 To assess intra-assay variability, 3 to 5 aliquots of  
 252 unstimulated and BCG stimulated whole blood generated  
 253 in the same experiment (performed in 3 donors) were  
 254 thawed, stained and acquired on 4 separate days.

255 To assess inter-operator variability, the same procedure  
 256 outlined above to assess intra-assay variability was conducted  
 257 by two different operators.

258 To assess robustness, two operators independently re-  
 259 trieved multiple aliquots of stimulated and fixed cells derived  
 260 from 3 adult volunteers and stained the cells for 45, 60 and  
 261 90 min at either room temperature (20–23 °C) or at 4 °C.

262 To assess effects of long-term cryopreservation on the WB-  
 263 ICS assay, whole blood from 11 healthy adults was stimulated  
 264 with BCG, PPD, and SEB, or was left unstimulated, processed  
 265 and cryopreserved. Fixed cells from stimulated WB were

266 thawed, stained and acquired one week after cryopreservation  
267 and then every 3 months over a 4-year period.

### 268 2.9. Data analysis

269 All flow data analyses were performed using FlowJo  
270 software (version 9.0.2, TreeStar, Inc.). Statistical analysis was  
271 done with Microsoft Excel (2008) and GraphPad Prism (version  
272 5a). Frequencies of cytokine-expressing CD4 and CD8 T cells are  
273 reported after subtracting frequencies of cytokine-expressing  
274 cells detected in unstimulated controls. Ki67-expression by  
275 both CD4 and CD8 T cells was too low for reliable quantifica-  
276 tion; therefore, the molecule was excluded from the analysis.  
277 To assess precision, reproducibility and robustness, we calcu-  
278 lated the mean, standard error of the mean and the coefficient  
279 of variation of frequencies of specific cytokine-expressing CD4  
280 and CD8 T cells. Variance component analysis was performed to  
281 estimate the inter-run variance and the intra-run variance. The  
282 overall precision of the assay is the variability calculated from  
283 the results of two operators.

284 To assess influence of time of cryopreservation on WB-  
285 ICS readout, intra-individual coefficient of variation was  
286 calculated across multiple experiments performed over time  
287 of cryopreservation.

288 Associations between WB-ICS, ELISpot and QuantiFERON  
289 readouts were assessed using a Spearman's correlation test.

## 290 3. Results

### 291 3.1. Establishment of the lower limit of quantification 292 for the WB-ICS assay

293 Before conducting experiments to qualify the WB-ICS assay,  
294 we established the lower LOQ for the WB-ICS assay, by  
295 performing serial dilutions of stimulated WB with autologous  
296 unstimulated WB samples. Our aim was to determine the  
297 lowest frequencies of mycobacteria-specific IFN- $\gamma$ , TNF- $\alpha$ , IL-2  
298 and IL-17-expressing T cells that could be detected reliably  
299 using this assay. We used the gating strategy as shown in  
300 Supplementary Fig. 1. We calculated the percent deviation  
301 between expected and observed values of cytokine-expressing  
302 CD4 T cells. Observed frequencies of IFN- $\gamma$  and IL-2-expressing  
303 CD4 T cells appeared to vary marginally (5%–18% and 4%–21%  
304 respectively) from their respective expected values above  
305 0.04%, shown by the dashed horizontal lines (Fig. 1A and B).  
306 Marginal variation (3%–25% and 12%–29%) was observed for  
307 frequencies of TNF- $\alpha$  and IL-17 expressing CD4 T cells of 0.01% or  
308 above respectively (Fig. 1C and D). Similar analyses of antigen-  
309 specific CD8 T cells showed that reliable enumeration with  
310 similar marginal variability between observed and expected  
311 frequencies was only possible for IFN- $\gamma$ -expressing cells, above  
312 0.04% (data not shown). TNF- $\alpha$ , IL-17 and IL-2 were not detected  
313 or were expressed at very low frequencies in CD8 T cells,  
314 elevating variability in the measurement (data not shown).  
315 These results indicate that accuracy of enumeration of cytokine-  
316 expressing cells is dependent on the levels of expression, and  
317 possibly, the antibody-fluorochrome combination.

318 Next, we aimed to assess non-specific frequencies of  
319 cytokine positive cells (background levels) detected in this  
320 assay. We analysed frequencies of IFN- $\gamma$ -, TNF- $\alpha$ -, IL-2- and IL-  
321 17-expressing CD4 T cells detected in unstimulated, negative

control samples from the 54 samples generated in the 322  
experiment comparing the WB-ICS with QFT assays. Median 323  
frequencies of CD4 T cells expressing IFN- $\gamma$ , TNF- $\alpha$ , IL-2 and IL- 324  
17 in the unstimulated controls were 0.002 (range 0.0001– 325  
0.137), 0.002 (0.0001–0.071), 0.002 (0.0001–0.002) and 0.001 326  
(0.0001–0.038), respectively (Fig. 1I). Medians and upper 327  
interquartile range (IQR) values for the four cytokines fell 328  
below the established limit of reliable quantification of the flow 329  
cytometer (Fig. 1I). 330

### 329 3.2. Intra-assay variability 331

Next, we analyzed coefficients of variation (CVs), a measure 332  
of precision within or between repeated measurements. CVs 333  
below 30% have been previously reported and are considered 334  
acceptable in ICS assays (Weinberg et al., 2000; Nomura et al., 335  
2000). We used the gating strategy as shown in Supplementary 336  
Fig. 1. We calculated CVs from frequencies of BCG-specific CD4 337  
and CD8 T cells, measured in quadruplet stimulation conditions 338  
for each sample (intra-assay variation). Each experiment was 339  
also repeated four times, on different days. CVs for IFN- $\gamma$ , TNF- 340  
 $\alpha$ -, IL-2- or IL-17-expressing CD8 and CD4 T cells were 341  
calculated. CVs for the frequencies TNF- $\alpha$ -, IL-2- and IL-17- 342  
expressing CD8 T cells were mainly above 30% (black, blue and 343  
green circles) while CVs for IFN- $\gamma$ -expressing cells were all 344  
below 30% (red circles) (Fig. 1E and F). The four cytokine- 345  
expressing CD4 T cell subsets measured showed CVs below 30% 346  
(Fig. 1G and H). From these experiments, we concluded that 347  
measurement of BCG-specific IFN- $\gamma$ , TNF- $\alpha$ -, IL-2- or IL-17- 348  
expressing CD4 T cells and IFN- $\gamma$ -expressing CD8 T cells by WB- 349  
ICS assay was reliable. Therefore, our subsequent analyses 350  
focused on these outcomes. 351

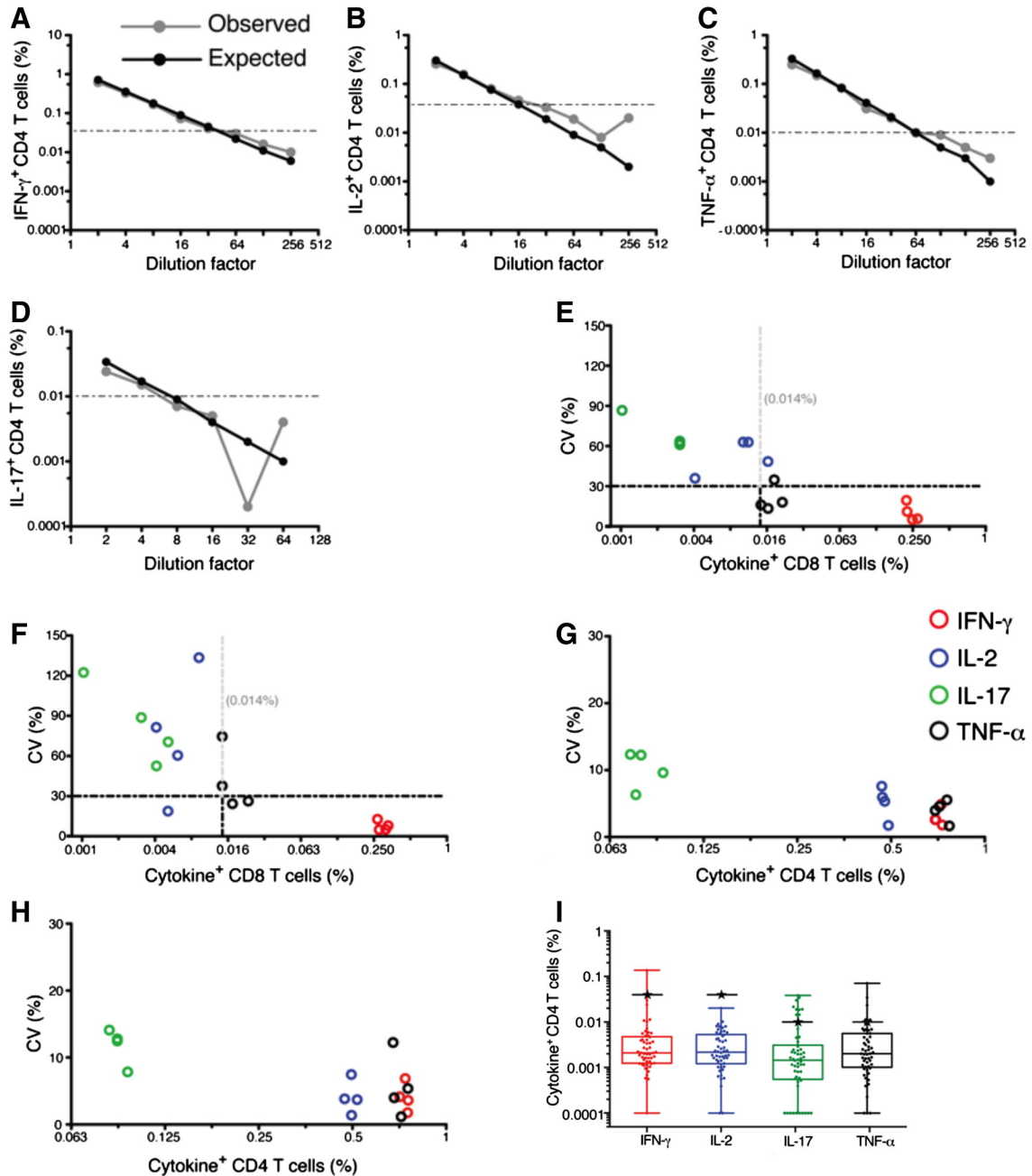
To measure intra-assay variability, CVs for each experiment 352  
and operator were computed separately. CVs below 15% were 353  
consistently observed for BCG-specific CD4 T cells expressing 354  
total IFN- $\gamma$ , TNF- $\alpha$  and IL-2, and for total IFN- $\gamma$ -expressing CD8 355  
T cells (Table 1). CVs for specific CD4 T cells expressing IL-17 356  
were greater than for the other cytokines due to the lower 357  
frequencies of these cells. Nevertheless, all the CVs were below 358  
30% except for a single day for one operator, when a CV of 45.9% 359  
was observed. This was an unexpected result and further 360  
investigations revealed high IL-17 expression in the experi- 361  
mental control, likely compromising the precision of the ICS. 362  
Nevertheless, in general, these results show that intra-assay 363  
variability of the WB-ICS assay is acceptable. 364

### 363 3.3. Inter- and intra- operator variability 365

Ideally, a single operator should perform sample processing, 366  
acquisition and analyses to minimise variability in immuno- 367  
logical assays. However, this is not always possible. We 368  
investigated inter-operator variability of the thawing, antibody 369  
staining and flow cytometry acquisition steps of the WB-ICS 370  
assay. Blood stimulation and processing until cryopreservation 371  
were performed by a single operator. Then, two operators 372  
thawed aliquots, applied identical reagents and antibody 373  
staining procedures to samples from 3 healthy donors and 374  
acquired the cells. Identical samples were stained and analysed 375  
on four different days. Data analysis, including gating was done 376  
by a single operator. Very little inter-operator variability was 377  
observed in frequencies of IFN- $\gamma$ , TNF- $\alpha$ , IL-2 and IL-17- 378

379 expressing CD4 T cells and IFN- $\gamma$ -expressing CD8 T cells across  
 380 the four experiments (Fig. 2). As expected, variability was  
 381 greatest when low frequencies of specific T cells were detected,

382 such as IL-17-expressing CD4 T cells (Fig. 2D). These findings  
 383 were supported by analyses of intra-operator CVs, which were  
 384 consistently below 30% (Table 1 and Fig. 3).



**Fig. 1.** Variability of the WB-ICS assay for detecting cytokine-expressing CD4 and CD8 T cells. Establishing the lower limit of quantification of the WB-ICS assay (A–D). Unstimulated cells were spiked with a series of 2-fold diluted BCG-stimulated cells, then stained and analysed. Expected frequencies of cytokine positive cells were derived by dividing the frequencies of BCG-specific cytokine-expressing CD4 and CD8 T cells in the undiluted samples by the dilution factor. The plots show the expected and observed frequencies of BCG-specific IFN- $\gamma$  (A), TNF- $\alpha$  (B), IL-2 (C) and IL-17 (D) CD4 T cells for each dilution factor. Establishing variability in detection of cytokine-expressing CD4 and CD8 T cells (E–H). Aliquots of fixed cells from a single BCG-stimulated (and unstimulated) whole blood sample were thawed and stained independently by two operators on 4 different days. For each operator on each day, 2–5 aliquots of cells were stained and acquired to calculate the coefficient of variation (CV) for frequencies of BCG-specific IFN- $\gamma$ , TNF- $\alpha$ , IL-2- or IL-17-expressing CD8 T cells (E, F) and CD4 T cells (G, H) for one operator are shown. The vertical dashed lines are a 30% CV mark. Frequencies of cytokine-expressing CD8 T cells with CVs less than 30% (IFN- $\gamma$ ) and more than 30% (TNF- $\alpha$ , IL-2- and IL-17) (panels E and F) are shown. For each repeat, frequencies detected in the unstimulated sample were subtracted from frequencies in the respective BCG-stimulated sample. (I) Frequencies of non-specific IFN- $\gamma$ , TNF- $\alpha$ , IL-2- or IL-17-expressing CD4 T cells, detected in unstimulated cells, in 54 adults. Horizontal lines within the boxes represent medians, boxes represent the IQR and whiskers represent the range. The black horizontal lines with stars represent the established limit of reliable quantification of the flow cytometer. (For interpretation of the references to colour in this figure legend, the reader is referred to the web version of this article.)

**Table 1**

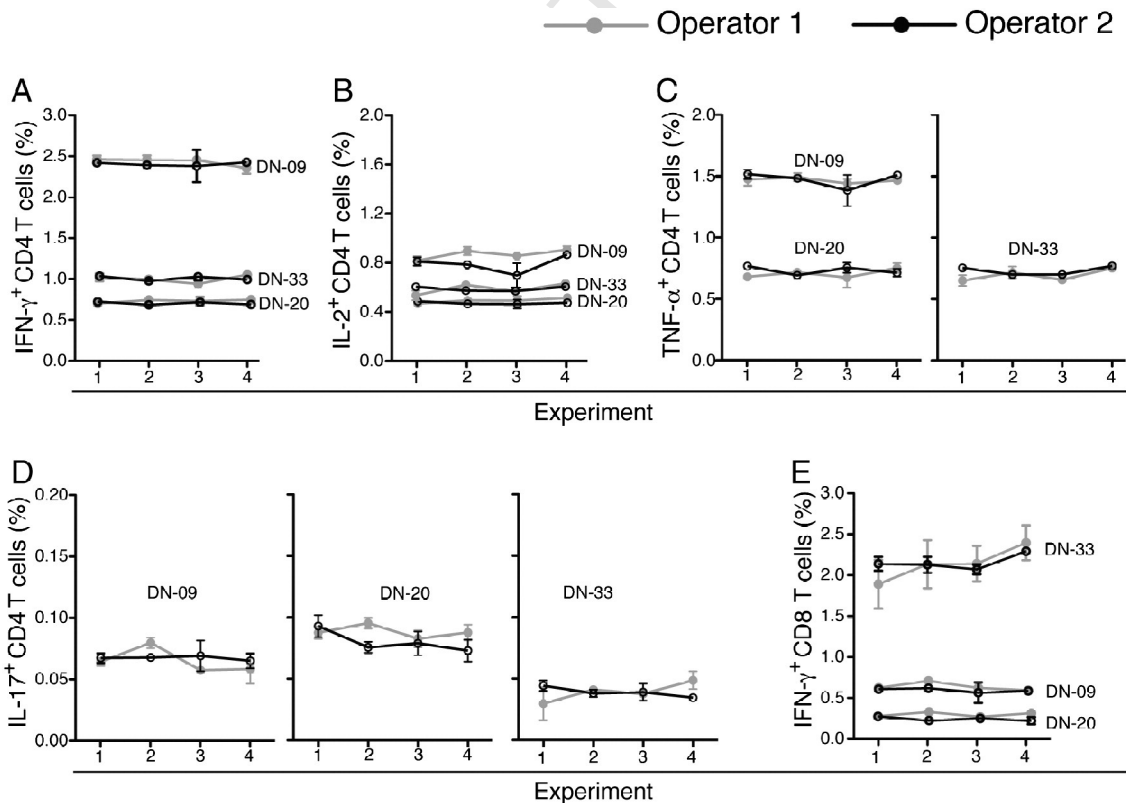
Intra-assay coefficient of variation using the WB-ICS assay: Two operators (A and B) independently processed, stained and acquired fixed unstimulated and BCG-stimulated cells derived from 3 adult healthy volunteers (DN-09, DN-20 and DN-33). Each operator performed each experiment (on 4 different days) using 2–5 separate aliquots of identical cells. Thereafter, flow cytometry data analyses were done by a single analyst and the CVs of frequencies of specific total CD4 T cells expressing IFN- $\gamma$ , IL-2, TNF- $\alpha$  or IL-17 and frequencies of specific total CD8 T cells expressing IFN- $\gamma$ , were calculated for each operator, donor and experiment (day). Unstimulated responses were subtracted from the respective BCG-stimulated responses for each sample and experiment.

		Intra-assay CV (%)										Number of repeats per run/day
		IFN- $\gamma$ <sup>+</sup> CD4 T cells		IL-2 <sup>+</sup> CD4 T cells		TNF- $\alpha$ <sup>+</sup> CD4 T cells		IL-17 <sup>+</sup> CD4 T cells		IFN- $\gamma$ <sup>+</sup> CD8 T cells		
Operator		A	B	A	B	A	B	A	B	A	B	A (B)
DN-09		1.9	1.2	4.6	4.2	3.5	2.5	5.0	5.9	6.2	6.5	5 (5)
		2.3	1.7	3.8	2.4	2.3	1.0	5.5	3.0	1.2	1.5	3 (3)
		1.2	8.3	1.3	14.2	2.6	9.1	2.7	18.2	2.7	22.1	3 (3)
		3.0	0.4	3.2	1.8	1.0	0.6	20.5	9.4	1.3	5.4	3 (3)
DN-20		4.1	1.8	3.8	1.8	4.0	1.6	12.9	9.5	5.0	5.8	5 (4)
		1.7	2.6	1.3	6.0	1.1	3.9	8.0	6.0	8.0	11.1	3 (3)
		6.8	4.9	7.4	7.7	12.2	5.5	14.3	12.1	12.7	4.8	3 (3)
		3.6	2.6	3.7	5.3	5.3	4.5	12.5	12.3	5.1	19.4	3 (3)
DN-33		4.7	1.8	7.3	2.6	6.8	0.7	<b>45.9</b>	9.7	15.7	4.0	5 (5)
		1.3	0.8	2.0	3.3	6.7	1.5	6.5	7.9	13.8	4.5	3 (3)
			4.2		5.1		3.6		17.8		2.8	2 (3)
		3.2		2.0		3.4		14.9		8.9		3 (2)

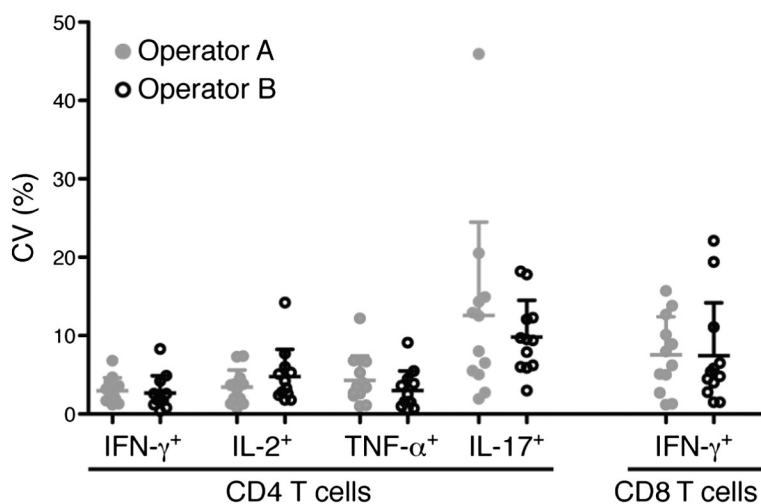
From these experiments, we concluded that the WB-ICS assay is not prone to major operator variability, provided that operators are suitably trained and adhere to standard operating procedures.

3.4. Robustness of the WB-ICS assay

Robustness is defined as the ability of an assay to withstand deliberate variation in assay conditions. We sought to determine



**Fig. 2.** Assay reproducibility. Four aliquots of fixed cells from 3 BCG-stimulated whole blood samples were thawed, stained and acquired by two operators. Analysis of flow cytometry data to determine frequencies of BCG-specific CD4 T cells expressing IFN- $\gamma$  (A), IL-2 (B), TNF- $\alpha$  (C) and IL-17 (D), and IFN- $\gamma$ -expressing CD8 T cells (E) were completed by a single analyst. Frequencies detected in unstimulated cells were subtracted from the respective BCG-induced frequencies. Mean and standard deviation of the 3 samples for each experiment are shown.



**Fig. 3.** Precision and intra-operator variability. CVs of frequencies of BCG-specific IFN- $\gamma$ , IL-2, TNF- $\alpha$  and IL-17-expressing CD4 T cells or IFN- $\gamma$ -expressing CD8 T cells, measured by 2 operators (grey and black dots). Each operator independently stained and acquired 3 sets of samples in 4 independent experiments. Frequencies detected in unstimulated cells were subtracted from respective BCG-induced frequencies for each sample and experiment. Then, the CVs for each operator and cytokine were computed. Lines on the plots represent means and SD.

392 effects of variation in duration and temperature of monoclonal  
 393 antibody staining on assay outcome. Two operators independ-  
 394 ently retrieved multiple aliquots of stimulated and fixed cells  
 395 derived from 3 adult volunteers and stained the cells for 45, 60  
 396 and 90 min at either room temperature (20–23 °C) or at 4 °C.  
 397 These staining conditions did not significantly affect quantitation  
 398 of frequencies of cytokine-expressing BCG-specific CD4 or CD8 T  
 399 cells; all CVs were below 9% (Table 2). These data were also used  
 400 to fit in a factorial design model. The data was fitted into the full  
 401 model and the analysis revealed that the interactions were not  
 402 significant (data not shown). From these analyses, we concluded  
 403 that WB-ICS was robust to withstand the evaluated changes in  
 404 the staining conditions.

### 3.5. Effects of long-term cryopreservation

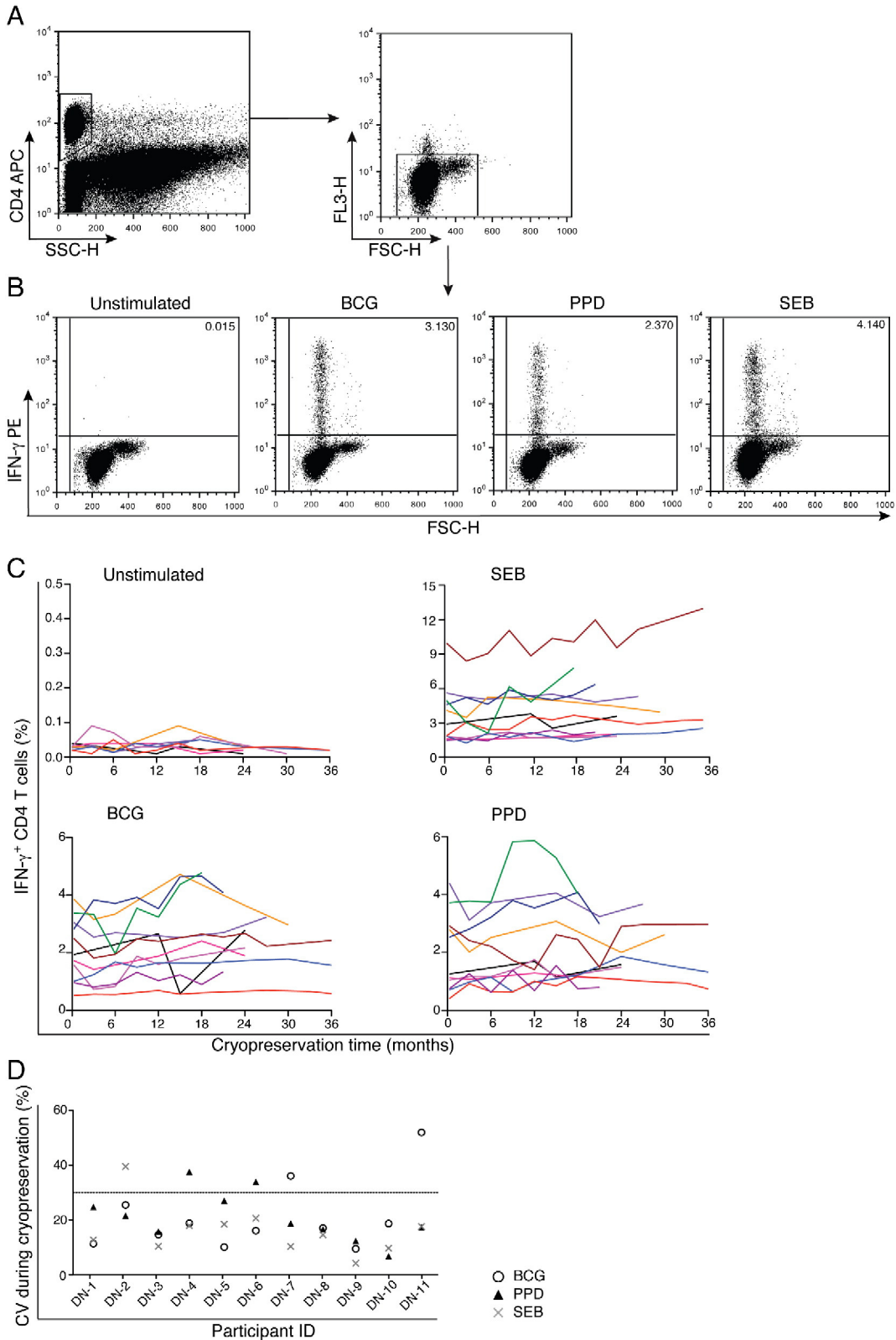
405

An advantage of the WB-ICS assay is that blood is stimulated 406  
 immediately and fixed lymphocytes are then cryopreserved, 407  
 allowing later staining and analysis in sample batches. Our next 408  
 experiments assessed effects of long-term cryopreservation on 409  
 measurement of mycobacteria-specific IFN- $\gamma$ -expressing CD4 T 410  
 cells. Whole blood from 11 healthy adults was stimulated with 411  
 BCG, PPD, and SEB, or was left unstimulated, then processed 412  
 and cryopreserved. Cells were thawed, stained and acquired on 413  
 a FACSCalibur one week after cryopreservation and then every 414  
 3 months over a 3-year period (Fig. 4A and B). Frequencies of 415  
 CD4 T cells expressing IFN- $\gamma$  either in the unstimulated or 416

t2.1 **Table 2**

t2.2 Robustness of the WB-ICS assay. Two operators (A and B) independently processed, stained and acquired fixed cells from either unstimulated or BCG-stimulated  
 t2.3 samples derived from 3 adult healthy volunteers. Variability in the temperature and duration of monoclonal antibody cocktail staining was assessed. Each operator  
 t2.4 performed the experiments independently. Thereafter, flow data analyses were done by one analyst and the mean frequencies of specific total CD4 T cells expressing  
 t2.5 IFN- $\gamma$ , IL-2, TNF- $\alpha$  or IL-17 and frequencies of specific total CD8 T cells expressing IFN- $\gamma$ , were calculated for each operator and each staining condition. Unstimulated  
 t2.6 responses were subtracted from the respective BCG-stimulated responses for each sample and experiment.

t2.7 ICS antibody staining t2.8 temperatures	Operator	ICS antibody staining duration (min)	Mean cytokine <sup>+</sup> CD4 T cells (%)				Mean cytokine <sup>+</sup> CD8 T cells (%)
			IFN- $\gamma$	IL-2	TNF- $\alpha$	IL-17	IFN- $\gamma$
t2.9 4 °C	A	45	1.070	0.487	0.823	0.057	0.791
t2.10 4 °C	B	45	1.104	0.530	0.819	0.056	0.728
t2.11 20–23 °C	A	45	1.110	0.556	0.909	0.060	0.737
t2.12 20–23 °C	B	45	1.135	0.565	0.899	0.067	0.718
t2.13 4 °C	A	60	1.181	0.550	0.888	0.056	0.739
t2.14 4 °C	B	60	1.150	0.587	0.886	0.057	0.780
t2.15 20–23 °C	A	60	1.073	0.526	0.847	0.064	0.678
t2.16 20–23 °C	B	60	1.151	0.566	0.892	0.064	0.716
t2.17 4 °C	A	90	1.277	0.535	0.896	0.066	0.825
t2.18 4 °C	B	90	1.174	0.591	0.927	0.069	0.834
t2.19 20–23 °C	A	90	1.146	0.576	0.975	0.062	0.816
t2.20 20–23 °C	B	90	1.186	0.611	0.925	0.071	0.730
t2.21 Overall mean			1.146	0.557	0.891	0.062	0.758
t2.22 Standard deviation			0.056	0.034	0.044	0.005	0.050
t2.23 CV			4.927	6.080	4.987	8.407	6.593





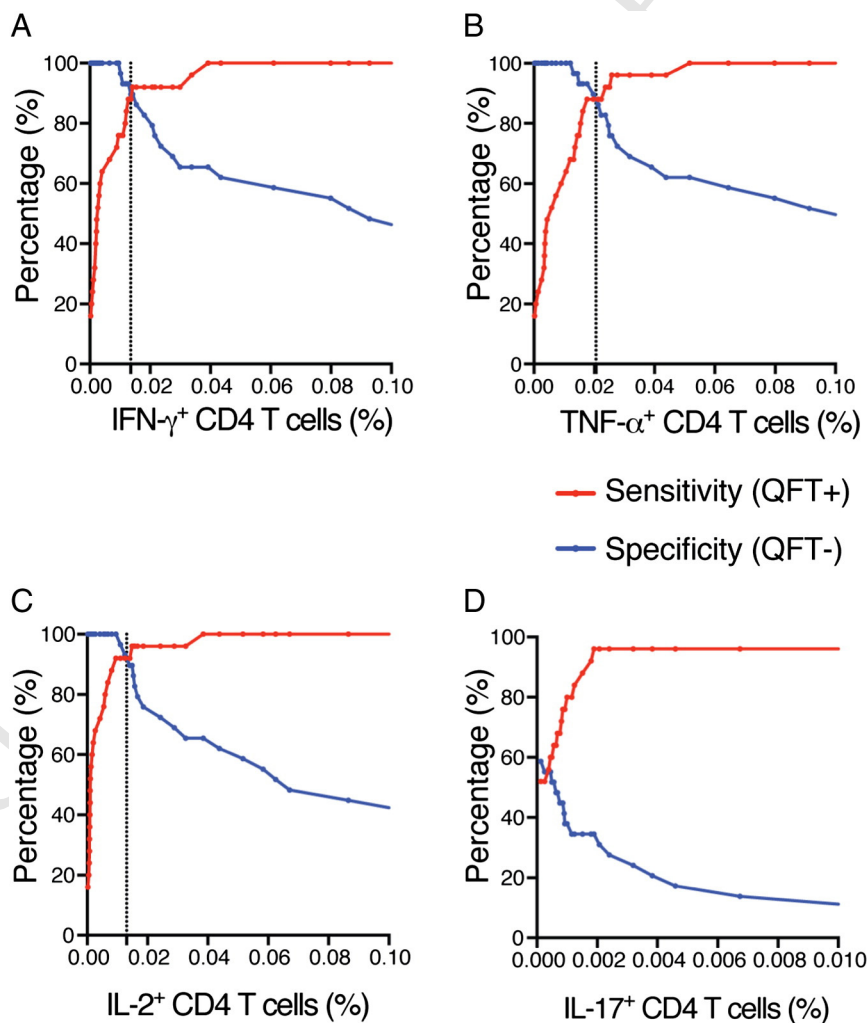
476 for diagnosis of latent *M.tb* infection (Scriba et al., 2012; Day  
477 et al., 2013; Abel et al., 2010b) (Fig. 5B). Of note, the magnitude  
478 of WB-ICS assay-detected specific T cells correlated well with  
479 the QFT-detected responses, even at very low frequencies.  
480 Interestingly, very strong correlations were also observed  
481 between frequencies of TNF- $\alpha$  or IL-2-expressing ESAT-6/CFP-  
482 10-specific CD4 T cells, detected by WB-ICS assay, and levels of  
483 soluble IFN- $\gamma$  detected by QFT (Fig. 5C and D).

484 These findings from our WB-ICS assay concur with those  
485 from the validated QFT, and further support the utility of the  
486 WB-ICS assay for accurate quantitation of antigen-specific  
487 cytokine-expressing T cells.

### 488 3.7. Sensitivity and specificity of the WB-ICS assay

489 Finally we were interested in assessing the sensitivity and  
490 specificity of the WB-ICS assay. In our study setting (Western  
491 Cape Province of South Africa), the population is highly  
492 exposed to *M.tb* and environmental non-tuberculous

mycobacteria, and BCG vaccination coverage is high (Corrigal, 493  
2005). As a result it is very rare to find persons without 494  
mycobacteria-specific T cell responses precluding evaluation of 495  
assay specificity. However, since the antigens ESAT-6 and CFP- 496  
10 measure T cell responses induced only by *M.tb*, and not BCG 497  
or most environmental mycobacteria, assay sensitivity and 498  
specificity could be assessed by comparing data from QFT 499  
negative and positive persons as shown in Fig. 5. The highest 500  
sensitivity and specificity pairing were observed for frequen- 501  
cies of IFN- $\gamma$ -, TNF- $\alpha$ -, and IL-2-expressing CD4 T cells at 502  
0.014%, 0.020% and 0.013%, respectively (Fig. 6A, B & C), 503  
suggesting that these frequencies could be used as assay 504  
positivity thresholds. We then applied these threshold fre- 505  
quencies to classify the QFT status from the 54 participants. 506  
Sensitivity and specificity values for IFN- $\gamma$ -, TNF- $\alpha$ -, and IL-2- 507  
expressing CD4 T cells were 90% and 92%; 90% and 88%; and 508  
93% and 92%, respectively. ESAT-6/CFP-10-specific IL-17- 509  
expressing CD4 T cells were more infrequent than Th1 cells, 510  
even in QFT-positive persons. This precluded reliable 511



**Fig. 6.** Sensitivity and specificity analysis of WB-ICS assay in QFT negative and positive adults. Curves showing paired sensitivity and specificity values at variable WB-ICS assay cut-offs for frequencies of ESAT-6/CFP-10-specific CD4 T cells expressing IFN- $\gamma$  (A), TNF- $\alpha$  (B), IL-2 (C), and IL-17 (D). Data are from 54 healthy adults with or without *M.tb* infection (QFT-positive or negative). The vertical lines represent paired sensitivity and specificity with the highest values, indicating the assay positivity threshold.

512 derivation of a positivity threshold for IL-17-expressing CD4 T  
513 cells and subsequent calculations of sensitivity and specificity.

#### 514 4. Discussion

515 Intact mycobacteria-specific T cell responses are necessary  
516 for human immunity against mycobacteria. However, exactly  
517 which functions such T cells should possess to protect against  
518 TB disease, is not known. As a consequence, assessment of  
519 vaccine-induced immune responses can only be used to infer  
520 vaccine immunogenicity or vaccine take. Regardless, regulatory  
521 agencies require data generated by qualified and/or validated  
522 assays for assessment of vaccine-induced immune responses to  
523 ultimately license candidate vaccines.

524 We report that the WB-ICS assay is a highly reproducible,  
525 reliable and robust method for quantifying antigen-specific  
526 cytokine-expressing CD4 and CD8 T cells. Variability in  
527 measured frequencies of specific T cells by WB-ICS, both within  
528 and across experiments was within accepted limits; CVs  
529 remained below 30%, even when performed by multiple  
530 operators and across different days.

531 The WB-ICS assay was originally optimized a decade ago,  
532 and has since been applied to many clinical studies (Roederer  
533 and Murphy, 1986). During the development and optimization  
534 process, many variables that may affect assay performance  
535 were tested, including blood volume, delay in initiation of  
536 antigen stimulation following blood collection and type of  
537 polypropylene tube used for stimulation (Scriba et al., 2011).  
538 Here, we build on these earlier optimization steps by  
539 investigating precision, robustness and effects of long-term  
540 cryopreservation to validate the WB-ICS assay readout for  
541 reliable measurement of mycobacteria-specific T cell frequen-  
542 cies. This assay is ideal for application in clinical field trials,  
543 where samples cannot be analysed in real time but rather in  
544 batches at a later time point, after cryopreservation. The  
545 sensitivity and accuracy of the WB-ICS assay are further  
546 supported by the strong correlation between *M.tb* antigen-  
547 specific frequencies of IFN- $\gamma$ , TNF- $\alpha$  and/or IL-2-expressing  
548 CD4 T cells and the quantitative value of IFN- $\gamma$  detected by the  
549 validated diagnostic QFT. Good agreement was also observed  
550 when compared with the highly quantitative IFN- $\gamma$  ELISpot  
551 assay. Furthermore, sensitivity and specificity curves generated  
552 with ESAT-6/CFP-10-specific T cell data derived from QFT  
553 positive and negative participants showed that the WB-ICS  
554 assay could reliably discriminate *M.tb* infection from non-  
555 infection, even at very low frequencies of mycobacteria-specific  
556 Th1 cells. Importantly, long-term cryopreservation of fixed cells  
557 from stimulated whole blood did not result in significant  
558 increases in assay variability. Our results suggest that the  
559 performance characteristics of the WB-ICS assay facilitate  
560 reliable measurement of antigen-specific T cells.

561 Whole blood-ICS assay experiments to establish sensitivity  
562 and specificity can be challenging in settings with high  
563 prevalence of sensitization to the antigen of interest. We applied  
564 frequencies of ESAT-6/CFP-10-specific T cells measured in QFT  
565 positive and negative participants to calculate the sensitivity and  
566 specificity of the WB-ICS, and derive assay positivity thresholds.  
567 Further, accuracy determination in WB-ICS assay is impractical  
568 because the true frequency of the outcome being quantified is  
569 often unknown. Comparing the WB-ICS assay with other  
570 validated assays such as ELISpot and QFT in quantifying specific

571 outcomes could be considered as an alternative evaluation of the  
572 accuracy. The WB-ICS assay allowed highly sensitive (sensitiv-  
573 ities of 90% to 93%) detection of cytokine-expressing ESAT-6/  
574 CFP-10-specific CD4 T cells, when compared with QFT assay. WB-  
575 ICS assay outcomes were highly correlated with QFT and ELISpot  
576 assay outcomes, even when including WB-ICS results below the  
577 lower LOQ formally suggested by the serial dilution experiments.  
578 An important factor that may have led to the underestimate of  
579 WB-ICS LOQ was the frequency of non-specific cytokine-positive  
580 T cells in the unstimulated sample. Because unstimulated  
581 samples were used to perform serial dilutions of stimulated  
582 cells, the lower LOQ for each cytokine was largely determined by  
583 the frequencies of cytokine expressing T cells in the unstimulated  
584 cells in this particular donor (0.04% for IFN- $\gamma$  and IL-2, 0.01% for  
585 TNF- $\alpha$  and IL-17 CD4 T cells, and 0.04% for IFN- $\gamma$  CD8 T cells).

586 To assess how inter-individual variability may have influ-  
587 enced the LOQ, we measured Th1 cell frequencies in  
588 unstimulated samples from a larger sample size ( $n = 54$ ).  
589 These mostly fell below the LOQ, suggesting that the sensitivity  
590 of quantification shown in Fig. 1 is likely underestimated. This  
591 possible underestimation of the LOQ is further supported by the  
592 positivity threshold frequencies of IFN- $\gamma$  (0.014%) and IL-2  
593 (0.02%), both below the cytometer's LOQ. However, a positivity  
594 threshold frequency of TNF- $\alpha$  (0.02%) was greater than the LOQ.

595 A major advantage of the WB-ICS assay is that cells can be  
596 fixed and cryopreserved after antigen stimulation is completed,  
597 thus eliminating any potentially detrimental effects of cryo-  
598 preservation on the function of antigen presenting and effector  
599 T cells. Further, because stimulated cells are fixed before  
600 cryopreservation, maintenance of cell integrity during cryo-  
601 preservation and thawing procedures is less important. A  
602 disadvantage of cryopreserving fixed cells from stimulated  
603 whole blood is that the choice of antigen for stimulation cannot  
604 be changed, limiting flexibility of the assay.

605 This highly reproducible, reliable and robust WB-ICS assay  
606 is ideal for detection and characterisation of antigen-specific  
607 cytokine expressing CD4 and CD8 T cells. This is essential for  
608 determining which attributes of antigen-specific T cell re-  
609 sponses may contribute to vaccine efficacy.

610 Supplementary data to this article can be found online at  
611 <http://dx.doi.org/10.1016/j.jim.2014.12.003>.

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