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1	Performance of an IAVI-African Network of Clinical Research Laboratories in
2	Standardized ELISpot and Peripheral Blood Mononuclear Cell Processing in
3	Support of HIV Vaccine Clinical Trials
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22 Running title: PBMC processing and ELISpot proficiency testing

24 Abstract

Immunological assays performed in different laboratories participating in multi-centre 25 clinical trials must be standardized in order to generate comparable and reliable data. This 26 entails standardized procedures for sample collection, processing, freezing and storage. 27 The International AIDS Vaccine Initiative (IAVI) partnered with local institutions to 28 29 establish Good Clinical Laboratory Practice (GCLP)-accredited laboratories to support clinical trials in Africa, Europe and Asia. Here we report on the performance of seven 30 laboratories based in Africa and Europe in the interferon-gamma enzyme-linked 31 immunospot (IFN- γ ELISpot) assay and peripheral blood mononuclear cell (PBMC) 32 processing over four years. Characterized frozen PBMC samples from 48 volunteer blood 33 packs processed at a central laboratory were sent to participating laboratories. For each 34 stimulus, there were 1751 assays performed over four years. 98% of these ELISpot data 35 36 were within acceptable ranges with low responses to mock stimuli. There were no significant differences in ELISpot responses at five laboratories actively conducting 37 immunological analyses in support of IAVI sponsored clinical trials or HIV research. In a 38 separate study, 1,297 PBMC samples isolated from healthy HIV-1 negative participants in 39 clinical trials of two prophylactic HIV vaccine candidates were analysed for PBMC yield 40 from fresh blood and cell recovery and viability following freezing and thawing. 94 % and 41 96 % of samples had fresh PBMC viabilities and cell yields within the pre-defined 42 acceptance criteria while for frozen PBMC, 99 % and 96 % of samples had acceptable 43 viabilities and cell recoveries respectively, along with acceptable ELISpot responses in 44 95%. These findings demonstrate the competency of laboratories across different 45 continents to generate comparable and reliable data in support of clinical trials. 46

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49 Importance.

There is a need for the establishment of an African network of laboratories to support large clinical trials across the continent to support and further the development of vaccine candidates against emerging infectious diseases such as Ebola, Zika and dengue viruses and the continued HIV-1 pandemic. This is particularly true in sub-Saharan Africa where the HIV-1 pandemic is most severe. In this report we have demonstrated by using standardized SOPs, training, equipment and reagents that GCLP-accredited clinical trial laboratories based in Africa and Europe can process clinical trial samples and maintain cell integrity and functionality demonstrated by IFN- γ ELISpot testing, producing comparable and reliable data.

72 Introduction

Clinical trials related to HIV, malaria and tuberculosis (TB) have been conducted in Africa 73 for decades (1-4). In order to harmonize the immunological data generated from these 74 clinical trials, laboratories responsible for clinical sample processing must establish 75 standardized procedures to meet International Conference on Harmonization (ICH) Good 76 Clinical Practice (GCP) and World Health Organisation (WHO) guidelines for collection, 77 processing and storage of samples and for immunological assays. The International AIDS 78 Vaccine Initiative (IAVI) has partnered with local institutions and established Good Clinical 79 Laboratory Practice (GCLP)-compliant laboratories across Africa, Europe and India to 80 conduct safety and immunogenicity assessments in support of clinical trials of HIV 81 vaccine candidates (5, 6). These laboratories are equipped to process and store samples 82 for later testing and are able to perform ELISpot and flow cytometry immunological 83 assays. IAVI has conducted over 20 phase 1 HIV vaccine trials (www.iavi.org/trials-84 database), with the majority in Africa (7-10). To ensure uniformity of data, IAVI sponsored 85 a central laboratory at Imperial College London (Human Immunology Laboratory) to 86 provide Standardised Operating Procedures (SOPs), training, critical assay reagents, 87 long-term centralised sample storage and to perform immunological assays where local 88 laboratories have no capability. IAVI and partners have developed the capacity of local 89 90 personnel professionally and academically, through technical training, mentoring and funding for investigator-initiated research projects. These GCLP-accredited-laboratories 91 have also been used as reference laboratories allowing local (and International) research 92 organizations to utilize existing facilities for their research and staff training in technical 93 assays and GCLP guidelines. 94

The performance of the IFN- γ ELISpot assay across multiple laboratories both within and across continents is critical to the generation of standardized data on vaccine immunogenicity (11). Previous studies (12) showed varied responses across laboratories
 while more recent ELISpot proficiency studies (13-15) have shown significantly improved
 results. These recent findings demonstrated that GCLP-accredited laboratories
 participating in proficiency testing over time can generate highly concordant results.

The majority of laboratories performing end-point ELISpot assays and enrolled in ELISpot proficiency schemes are based in Europe and USA (12, 15, 16). With increased focus on testing HIV vaccine products where the pandemic is most severe and with renewed interest in many "orphan" tropical infectious diseases (17), development of laboratory networks able to support large clinical trials across sub-Saharan Africa has gained importance. In order for IAVI-supported laboratories to meet international standards, they were enrolled into an IAVI-sponsored ELISpot proficiency scheme organised by Clinical Laboratory Services, South Africa.

119 Methods and Materials

120 Participating laboratories

All participants were IAVI-supported laboratories including: 1) Kenya AIDS Vaccine 121 Initiative-Institute of Clinical Research (KAVI-ICR) University of Nairobi, Nairobi, Kenya; 2) 122 IAVI Human Immunology Laboratory (HIL), Imperial College London, United Kingdom; 3) 123 Uganda Virus Research Institute (UVRI)-IAVI, Entebbe, Uganda; 4) Clinical Laboratory 124 Services (CLS), Witwatersrand University, Johannesburg, South Africa; 5) Kenya Medical 125 126 Research Institute Centre for Geographical Medicine Research Coast (KEMRI-CGMRC), Kilifi, Kenya; 6) Zambia EMORY HIV Research Project (ZEHRP), Lusaka, Zambia; and 7) 127 Projet San Francisco (PSF), Kigali, Rwanda. The laboratories were located in six countries 128 (Kenya, Uganda, Rwanda, Zambia, South Africa and UK). CLS is not supported by IAVI but 129 provide PBMC, pretesting and shipping services for the ELISpot proficiency testing. CLS 130 participate in quarterly testing of the ELISpot proficiency and are included in this analysis. 131

132 Laboratory Preparation

133 Process of establishing clinical trial laboratories under GCLP guidelines

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135 Comprehensive training programs, standardized GCLP-compliant serviced and calibrated 136 equipment and quality control (QC) measures were integral to establishing IAVI's 137 laboratories. The key elements for establishing the clinical trial laboratories are shown in 138 **Table 1** and described in detail in the Supplementary Information.

The quality systems and SOPs were designed to minimise failure, identify problems, initiate corrective actions and monitor resolutions. Two laboratories (HIL and CLS) have been designated as support and QC management. London (United Kingdom) and Johannesburg (South Africa) were ideal locations to support a global clinical trial program. Both are major international hubs in Europe and Africa, with direct flights to and from the above IAVIsupported laboratories, thereby reducing time, cost and risk to samples in transit.

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146 ELISpot proficiency panel design

IAVI-GCLP laboratories were enrolled in an ELISpot proficiency scheme coordinated by 147 CLS, South Africa. At CLS, PBMC samples from 48 volunteer blood packs were processed 148 and shipped to IAVI-supported laboratories for assessment of cell viability and IFN-y ELISpot 149 responses. Two peptide pools were used; a pool of 32 8-10mer peptides representing 150 immunodominant CD8+ T-cell epitopes from Cytomegalovirus. Epstein Barr virus and 151 Influenza virus (CEF) (18) and a pool of 138 15mer peptides overlapping by 11 amino acids 152 spanning the human Cytomegalovirus (CMV) pp65 protein at a final assay concentration of 153 1.5 µg/mL per peptide. A positive control of phytohaemagglutinin (PHA-L; Sigma L4144) at 154 10µg/mL and a mock stimulus (medium / dimethyl sulfoxide (DMSO) alone) were used. 155

156 **PBMCs for ELISpot Proficiency**

Each ELISpot proficiency panel consisted of 6 PBMC samples sufficient for monthly testing 157 of the same 6 samples for 6 months at all laboratories. Different PBMC batches were 158 provided from CLS each 6-month period over four years as shown in Figure 1. Four 159 laboratories; KAVI-ICR, UVRI-IAVI, PSF-Kigali and the HIL actively performing 160 immunological assays in support of IAVI clinical trials, conducted monthly ELISpot testing 161 while 3 laboratories; KEMRI-CGMRC, ZEHRP and CLS not actively performing 162 immunological assays conducted quarterly ELISpot testing. Raw ELISpot data were 163 submitted to the IAVI HIL central data repository for evaluation of responses and results 164 165 posted onto an access-restricted CLS website.

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168 Sample preparation and immunological assessment

169 Minimal Information About T-cell Assays (MIATA) guidelines were followed for transparent 170 reporting on immunological assays (19).

PBMCs and clinical trial samples: PBMCs used in ELISpot proficiency testing were 171 prepared by Ficoll density centrifugation of buffy coats obtained from 48 healthy, HIV-1 172 negative blood donors (South African National Blood Bank, Johannesburg, South Africa). 173 Clinical trial samples were obtained from heparinized blood from healthy HIV-1 negative 174 placebo and vaccine recipients participating in clinical trials of two prophylactic HIV vaccine 175 candidates at KAVI-ICR, UVRI-IAVI, PSF and ZEHRP (8, 10). Isolated PBMCs had to meet 176 177 the following pre-defined acceptance criteria: 1) viability above 90% and cell yield greater than 0.7 x10⁶ PBMC/mL blood for freshly isolated PBMCs, 2) viability above 80% for frozen 178 PBMCs following thaw and overnight rest and 3) processing of PBMCs from blood draw to 179 start of freezing to be completed within 6 hours. Cell counts and viabilities were determined 180 using a Vi-Cell XR (Beckman Coulter, UK) automated cell counter. PBMCs were frozen in 181 freezing media containing 10% DMSO diluted in fetal calf serum (FCS) using a rate-182 controlled freezer (Planer, Sunbury-On-Thames, UK) which lowered the temperature by 1°C 183 per minute from +4°C to -80°C then 10°C per minute to -120°C, before transfer to vapour 184 phase liquid nitrogen (LN) until assayed. PBMC for ELISpot proficiency were stored in 185 vapour-phase liquid nitrogen prior to shipment to participating laboratories using 186 temperature-monitored cryogenic dry shippers (MVE Jencons, United Kingdom). The 187 188 shippers were calibrated according to an SOP. Briefly, empty dry shippers were pre-weighed before filling with LN then left overnight for adsorption of LN and decanted the following day. 189 Shipper weight loss and temperatures were recorded over the next 5 days. The dry shippers 190

were also fitted with the temperature loggers where temperature data were downloaded upon receipt of the dry shippers at the participating laboratories. In order for the dry shippers to pass the calibration, the average 24-hour weight loss over the 5-day calibration was 0.6kg+/-10% and temperature <-190°C. Following receipt, PBMC were transferred to vapourphase liquid nitrogen until assayed.

IFN-y ELISpot Assay: Cell recovery and viability of samples thawed and rested overnight for 196 ELISpot testing were analysed. PBMCs were removed from LN storage and transported to 197 the laboratory in dry ice and immediately immersed in a 37°C water bath until a small amount 198 of ice remained. Cells were transferred to 10mL cell culture medium (RPMI 1640 199 supplemented with 10% heat-inactivated FCS, 1 mM L-glutamine, 100 units/mL penicillin, 200 100 µg/mL streptomycin, 1mM sodium pyruvate and 0.5 mM HEPES (R10)), centrifuged at 201 250g/10 mins, supernatants decanted, cell pellets disrupted and resuspended in 4mL RPMI 202 203 1640 supplemented with 20% heat-inactivated FCS, 1 mM L-glutamine, 100 units/mL penicillin, 100 µg/mL streptomycin, 1mM sodium pyruvate and 0.5 mM HEPES (R20). Cells 204 205 were transferred to wells of 24 well culture plates in a 37°C/5% CO₂ incubator overnight. The following day, cells were recovered and washed in 10mL R10, supernatants decanted, 1mL 206 of R10 added and cells counted. 207

RPMI 1640, L-glutamine, HEPES, penicillin/streptomycin, sodium pyruvate and heat-208 inactivated FCS were all purchased from Sigma-Aldrich (St Louis/Missouri, USA). All 209 laboratories used the same batch of FCS for PBMC isolation and ELISpot assay, which was 210 pretested for acceptable performance in procedures for PBMC isolation, freezing, recovery 211 from frozen and low background (mock) and acceptable PHA/CMV responses in ELISpot. 212 213 PVDF membrane plates were obtained from Millipore (MSIPS4510; United Kingdom). Antihuman IFN-γ antibody (clone 1-D1K, 1 mg/mL) and biotinylated anti-human IFN-γ antibody 214 (clone 7-B6-1, 1 mg/mL) were purchased from Mabtech, Sweden; peroxidase-avidin biotin 215

complex (ABC) from Vector Laboratories, Burlingame, CA, USA; dimethylformamide (DMF)
from VWR International, USA; PHA, 3-amino-9-ethylcarbazole (AEC) tablets (A6926), acetic
acid, sodium acetate, 30% hydrogen peroxide (H₂O₂) and sterile tissue culture water and
phosphate-buffered saline (PBS) were all purchased from Sigma-Aldrich. CEF and CMV
pp65 peptides were purchased from Anaspec Inc., CA, USA.

Prior to setting up each ELISpot assay, PVDF plates were treated with 50µL of 70% ethanol 221 for 2 minutes, washed three times with 200 µL/well sterile PBS, coated with 100µL/well of 222 anti-human IFN-y (1-D1K, 10 µg/mL in PBS) and stored overnight at 4°C. Plates were 223 washed three times with 200 µL/well sterile PBS, blocked for a minimum of 2 hours with 200 224 µL/well R10 at 37°C, 5% CO₂ incubator. The blocking medium was removed and 100 µL/well 225 of R10-diluted mock, CEF/CMV peptides (2.25 µg/mL) and PHA (15 µg/mL) were added to 226 their respective wells (following a designated plate plan). 100 µL CMV peptide was added to 227 a designated no cell well. Thawed and overnight-rested PBMC were added at 200,000 cells 228 in 50 µL R10 to each well, with each sample and condition plated in guadruplicate. The 229 plates were incubated for 16-24 hours (37°C, 5% CO₂). The following day, cells were 230 removed and plates washed six times with 200 µL/well PBS with 0.05% tween (PBS/T) using 231 an automated ELISA washer (Bio-Tek Instruments Inc., Winooski, VT, USA). 100 µL 232 biotinylated anti-human IFN-y antibody (7-B6-1, 1 µg/mL in PBS with 0.1% BSA) were added 233 234 for two hours at room temperature (RT). Plates were washed six times as above before addition of 100 µL/well peroxidase avidin-biotin complex (per manufacturer's instructions), for 235 one-hour at RT. Plates were washed three times with 200 µL/well PBS/T followed by three 236 washes with 200 µL/well PBS. 100 µL/well of 0.45µm filtered AEC substrate (AEC tablet 237 dissolved in 2.5mL DMF, added to 47mL sterile tissue culture water containing 280uL 2M 238 acetic acid and 180µL 2M sodium acetate and finally 25µL H₂O₂ added) was added for 4 min 239 at RT. Plates were emptied thoroughly and the reaction stopped under gently-running tap 240

water and the underdrain removed before leaving the plates to dry overnight protected from light. The acceptance criteria for the IFN- γ ELISpot was the mock wells should have less than 10 spots per well and the peptide/media alone (no cells) 5 or fewer spots per well.

244 **Data Acquisition and Analysis**: Spots were evaluated with an AID reader system 245 (AutoImmun Diagnostika, Germany) with software version 4.0. Each laboratory used the 246 same model of AID reader and defined spot parameters. Responses are expressed as spot-247 forming cells (SFC) per 10⁶ viable PBMCs as shown in **Figure 6**.

Our main outcomes included 1) the recovery and viability rates of frozen PBMCs, 2) ELISpot 248 results compared to mock, CMV and CEF stimuli and 3) comparison of ELISpot results 249 250 across laboratories. For each peptide repeated measures, Poisson regression model was fit on background-subtracted count (except mock), with counts from the same volunteer 251 assumed to be correlated. The resulting least squares parameter estimates are presented 252 together with their 95% confidence intervals adjusted for multiple comparisons using the 253 Bonferroni method. Each model included volunteer, laboratory and month. Pair-wise 254 255 comparisons between laboratories are shown as the ratio of the least squares estimates of mean count with corresponding adjusted (Bonferroni) 95% confidence interval. Statistical 256 significance is defined as a 95% CI for the ratio that excludes unity (i.e., entirely above or 257 below the value 1). Figures 1, 4 and 5 were generated by Graph Pad prism software version 258 7.01. Other figures and statistical analyses were performed using SAS Version 9.3. 259

260 **Results**

1. Performance in ELISpot assay across 7 laboratories

Each assay included 4 replicates for each peptide. Results based on less than 4 replicates were assumed to be less reliable and excluded from analysis. For each peptide and mock stimulus there were 1751 assays performed of which 50 were excluded (i.e., about 2.9%). The distribution of mock, CMV and CEF responses across laboratories over time is shown in **Figure 1A & B**. To compare ELISpot data across the 7 laboratories, the response was background-subtracted counts (except mock). The covariates in the model were sample, laboratory and month.

Across the seven laboratories, the geometric mean ELISpot counts (SFC/10⁶ PBMC) were 6
- 10 for Mock, 289 – 438 for CEF and 172 – 266 for CMV (S1A, S2A & S3A Tables
respectively).

272 Statistical differences were observed between laboratories in mock counts as shown in **S1B** 273 **Table** and **Figure 2B** (p=0.0007). For example, the mock count at CLS is estimated to be 274 1.73 times the mock count at ZEHRP. Also, the mock count at PSF is 0.78 times lower than 275 at UVRI-IAVI. ZEHRP tends to have lower mock counts than all other laboratories.

When comparing the responses against CEF peptides across laboratories, KEMRI-CGMRC had significantly higher counts than other laboratories (p=0.0045, **S2B Table & Figure 2B).** When data for KEMRI-CGMRC are excluded, the overall difference between laboratories is not statistically significant (p=0.11, **S2C Table**).

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When comparing the responses against CMV responses across laboratories, KEMRI-CGMRC again had significantly higher counts than other laboratories (p=0.012, **S3B Table & Figure 2B**). On excluding data for KEMRI-CGMRC the overall difference between laboratories is still statistically significant (p=0.033, **S3C Table**), due to the counts at ZEHRP being lower than CLS and KAVI-ICR.

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287 Inter-operator analysis

ELISpot data from 3 operators at KAVI-ICR were compared. Samples from 12 288 volunteers were analyzed by 3 operators on 2 occasions, each operator analyzing the 289 same set of samples at the same monthly time points. ELISpot counts were obtained 290 for mock and background-subtracted CMV and CEF peptide pools. The covariates in 291 the model were sample, operator and month across the three operators, the 292 geometric mean ELISpot counts (SFC/10⁶ PBMC) were 9 – 12 for Mock, 368 – 393 293 for CEF and 538 - 598 for CMV (S4A Table). The differences between operators 294 were not statistically significant (S4B Table and Figure 3). 295

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297 2. Viability and cell yield of freshly isolated PBMCs and recovery from frozen 298 PBMCs across 4 laboratories

A total of 1297 PBMCs isolated from clinical trial samples were analysed for cell 299 viability, recovery and cell yield in four laboratories supporting two IAVI-sponsored 300 clinical trials. 1220 of 1297 (94%) freshly-isolated PBMCs had viabilities above 90% 301 with a median of 95% (range 81-100%) while those below 90% had a median of 88% 302 (range 81-90%, Figure 4A). Over 96% of these samples had cell yields greater than 303 0.7x10⁶ PBMC/mL blood, within the pre-defined acceptability criteria with few 304 samples having low cell yield ranging from 0.13-0.56x10⁶ PBMC/mL blood (Figure 305 4B). A total of 1205 of these samples were tested in ELISpot assay and almost all 306 (99%) had viabilities above 80% following thaw and overnight rest (within 307 acceptability criteria) with only 9 samples having viabilities below 80% ranging from 308 66 to 78% as shown in Figure 4C. Cell recoveries for these samples were above 309 6.0x10⁶ PBMC/vial (PBMCs were frozen at 10-15x10⁶ PBMC/vial); data were 310 normalized to 10 million cells as shown in Figure 4D. For all samples tested, cells 311 were functional in ELISpot assay with over 95% of the samples having mock 312

responses <50 SFC/10⁶ PBMC, PHA>1000 SFC/10⁶ PBMC and a range of CMV
responses.

The length of time from blood draw to sample processing and freezing has been 315 shown to affect the integrity of PBMC (20-22). Nearly all of our samples were 316 processed within 6 hours with 81 (6 %) processed beyond 6 hours (range 6.1-9.5 317 hours, Figure 4E & 5E). To assess the impact of longer processing of these samples, 318 the cell yields and viabilities from fresh blood were analysed together with cell 319 recoveries and viabilities following freezing and thawing. All samples except one had 320 freshly-isolated cell viabilities and cell yields within the acceptable range, that is 321 >90% and >0.7x10⁶ PBMC per mL blood respectively (Figure 5A & B). Only one 322 sample had a slightly lower cell yield of 0.57x10⁶ per mL blood (98% viability). Post 323 PBMC freezing cell viabilities ranged from 93-100% and recoveries above 6x10⁶ 324 PBMC/vial in 71/81 (87%) samples (Figure 5C & D). We further tested these 325 samples in ELISpot assay to assess their cell functionality and all samples performed 326 well with the mock responses <50 SFC/10⁶ PBMC, PHA responses > 1000 SFC/10⁶ 327 PBMC and a typical range of CMV responses indistinguishable from samples 328 processed within 6 hours, as shown in Figure 5F. 329

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339 Discussion

This report compares data generated across a network of 7 IAVI-supported GCLPaccredited clinical trial laboratories based in Africa and Europe. These laboratories were assessed on ELISpot proficiency testing and PBMC processing.

ELISpot proficiency data were analysed and compared across and within sites. Acceptability criteria for mock and PHA controls and CMV and CEF responses had to be clearly positive or negative during pre-testing. We found that all laboratories correctly detected responses against CMV and CEF peptide stimuli with the exception of a few sporadic higher data points in mock stimulus which was seen across laboratories. As expected even for technically competent laboratories there were occasional discrepant data points.

In this report we have demonstrated that ELISpot data for CEF and CMV responses 350 from 5 laboratories were not significantly different and were overall comparable. Four 351 of these laboratories were actively performing ELISpot analysis in support of IAVI 352 clinical trials whilst a fifth laboratory (CLS) performed ongoing ELISpot analyses as 353 part of establishing the proficiency scheme and in training of staff at other 354 We did observe significantly different data for CEF and CMV laboratories. 355 responses from 2 laboratories that were not routinely conducting ELISpot assay in 356 support of IAVI clinical trials when compared to the other laboratories. Staff at these 357 laboratories only performed ELISpot analyses as part of the proficiency program 358 described in this report and therefore would have less ongoing technical experience 359

in ELISpot analysis compared to staff at laboratories with active participation in 360 clinical trials. To mitigate this, staff at the 2 laboratories were retrained and 361 competency assessed. This highlights the need for review of staff retraining and 362 continual monitoring of laboratories' performance with trouble-shooting and staff 363 training and re-training as required, especially for laboratories taking on new 364 activities or trials or where laboratories have not performed certain techniques in a 365 trial setting for some time. However, although the ELISpot responses observed at 366 these 2 laboratories were statistically significantly different, the range of estimated 367 368 least squares mean counts across the 7 laboratories was not high with 274 to 438 for CEF (**S2A Table**) and 172 to 266 SFC/10⁶ PBMC for CMV (**S3A Table**). Statistically 369 significant differences in mock values between laboratories were apparent which 370 may be expected as in effect the vast majority of mock responses were close to 371 zero. Across the seven laboratories, the geometric mean ELISpot mock counts were 372 6 - 10 SFC/10⁶ PBMC (**S1A Table**). High variability of low T cell responses has 373 been reported previously (23). 374

Operator-dependent variability in ELISpot is a known phenomenon (24) and we assessed this in this report. It was not possible to analyse inter-operator variability at all laboratories as some laboratories had either a lone operator throughout, or a change of operators during the study period. However, we report on one laboratory with 3 operators performing the ELISpot assay on a rotational basis. All operators detected correctly the expected responses for CMV and CEF stimuli. Their data were highly correlated and variability in data points was not significantly different.

Achieving accurate and reliable results when assessing the immunogenicity of vaccine candidates, especially for multi-site clinical trials, is essential. In order to achieve this, samples must be processed according to standardised SOPs following 385 GCLP-guidelines for data integrity. PBMC processing in four clinical trial laboratories 386 were analysed for processing time from blood draw to start of freezing, cell yield and 387 viability and post-freezing viability and recovery. We report that the vast majority of 388 freshly isolated PBMCs had viabilities and cell yields within the acceptable range 389 across all laboratories.

Proper freezing and storage of samples is critical in preserving cell integrity and functionality (25). In this report we assessed the integrity of PBMCs processed and frozen at the laboratories. Cells were thawed at the HIL (Central repository lab) for ELISpot testing and nearly all samples had cell viabilities and recoveries within the acceptance criteria, with cell functionality demonstrated by good performance in ELISpot assay, therefore demonstrating the competency of laboratories in isolation, freezing, storing and shipping of PBMC samples.

PBMCs processed beyond 8 hours have been shown to have reduced cell viability 397 and compromised cell functionality (26). Here, we report that the majority of samples 398 were processed within 6 hours with the exception of few samples processed beyond 399 6 hours, mainly due to delayed delivery to the laboratories from some clinics located 400 up to 50 miles away. For samples processed beyond 6 hours, corrective and 401 preventive action (CAPA) reports were written to minimise or prevent recurrence 402 where possible and monitored on a monthly basis. Cell yields, viabilities and 403 recoveries of these samples were assessed to determine the impact of longer 404 processing on their integrity. All samples performed well in ELISpot with responses 405 to Mock, CMV, CEF and PHA stimuli being in the expected ranges with data similar 406 to samples processed within 6 hours. Therefore, we show that PBMCs processed 407 longer than 6 hours (up to 9 hours) are still viable and functional in ELISpot assay 408 and similar to what other groups have shown (20-21). 409

Participating laboratories are audited regularly for GCLP compliance by internal and external independent auditors. The audit covers SOPs, ELISpot and flow cytometry proficiency, external quality assurance programs and data integrity. The GCLP audit by an external auditor from Qualogy Ltd, UK is conducted annually with an accreditation certificate issued to compliant laboratories.

In conclusion, we have demonstrated that using standardised SOPs, equipment and reagents and working in a GCLP compliant laboratory, clinical trial laboratories located in Africa and Europe can process clinical trial samples and maintain cell integrity and functionality through ELISpot testing, producing comparable and reliable data.

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586 Figures and Tables Legends

Figure 1. ELISpot spot forming cells (SFC) per million PBMC and variability 587 across laboratories represented in box plots. A) years 1-2 and B) years 3-4. 588 Each panel consists of the same 6 donor samples tested over a 6-month period by 7 589 laboratories. Box plots represent the quartiles, horizontal line the median and 590 whiskers represent the maximum and minimum values. Each point represents 591 average SFC/10⁶ PBMC from replicates per donor at each laboratory. Laboratories 592 are color-coded as follows: Red = KAVI-ICR, Blue = UVRI-IAVI, Green = PSF, 593 Purple = ZEHRP, Yellow = KEMRI-CGMRC, Cyan = CLS and Black = HIL. 594

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Figure 2. Comparisons of PBMC ELISpot responses: All pair-wise least squares 596 means and their significance, on a natural log scale, for mock, CEF and CMV 597 respectively. For each comparison a line segment, centred at the least squares-598 599 means in the pair, is drawn. The segment length corresponds to the projected width of a confidence interval for the least squares mean difference. Segments that fail to 600 cross the 45° reference line correspond to significant least squares mean 601 differences. The graph shows which site pairs are significantly different (blue lines) 602 and which are not (red lines). 603

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Figure 3. Inter-operator ELISpot comparison from 3 operators at KAVI-ICR: All pair-wise least squares means and their significance, on a natural log scale, for Mock, CEF and CMV respectively. For each comparison a line segment, centred at the lest squares-means in the pair, is drawn. The length of the segment corresponds to the projected width of a confidence interval for the least squares mean difference. Segments that fail to cross the 45° reference line correspond to significant least squares mean differences. None of the pairs of operators are significantly different(all lines cross the 45-degree reference line).

Figure 4. Cell recovery, viability and processing time of clinical trial samples: A) viability of freshly isolated PBMC; B) cell yield per mL of blood; C) viability after overnight rest; D) cell recovery of PBMC frozen at 10-15x10⁶ PBMC per vial; data were normalized to 10 million cells; E) PBMC processing time. Each point in the scatter plot represents a sample and the lines represent the median with interquartile range. Horizontal lines show the acceptance cut-off.

Figure 5. Cell recovery, viability and ELISpot responses of samples processed 619 beyond 6 hours: A) viability from freshly isolated PBMC; B) cell yield per mL blood; 620 C) viability from frozen samples; D) cell recovery of PBMC per 10 million cells frozen 621 622 following thaw and overnight rest; E) processing time from blood draw to freezing of PBMC; F) ELISpot responses of PBMC tested against mock, PHA and CMV stimuli 623 for PBMCs processed within 6 hours (red) and beyond 6 hours (black). Each point in 624 the scatter plot represents a sample and the median (horizontal line). Horizontal lines 625 represent the acceptance cut-offs. 626

Figure 6. PBMC ELISpot responses against CMV peptides. Representative well images (plate wells C9 to C12) of the CMV responses for sample 3522 performed in quadruplicates across the laboratories. The SFC counted per well are given in each well image.

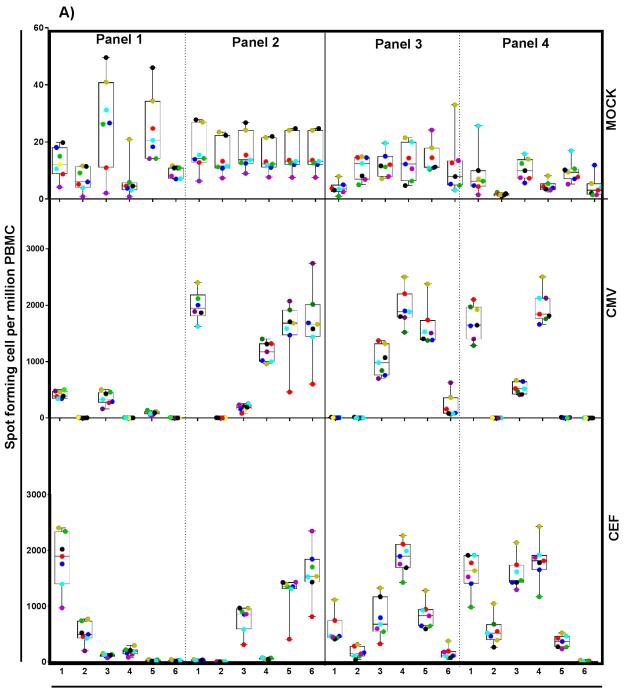
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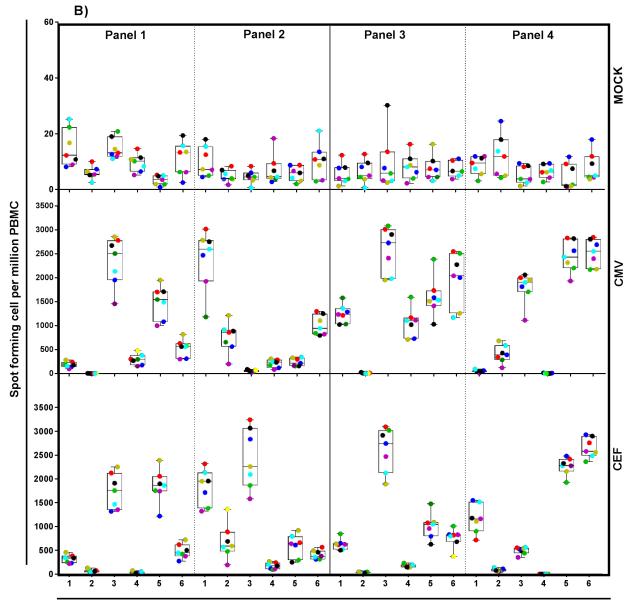
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Table 1: Summary of the process of establishing clinical trial laboratories underGCLP guidelines.



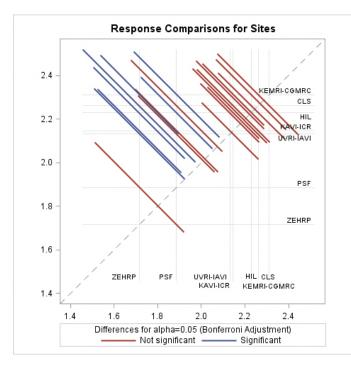
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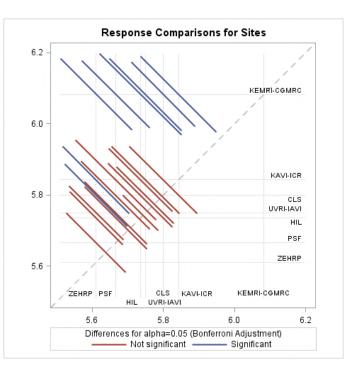


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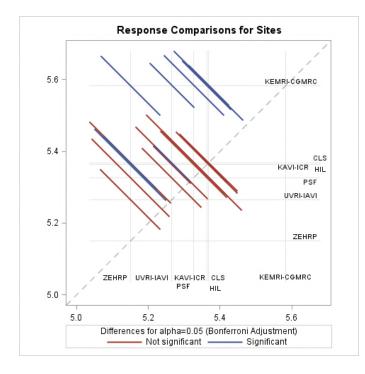
a. Mock





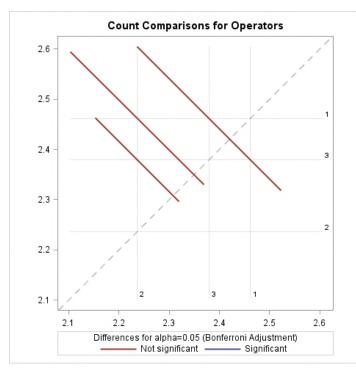


c. CMV

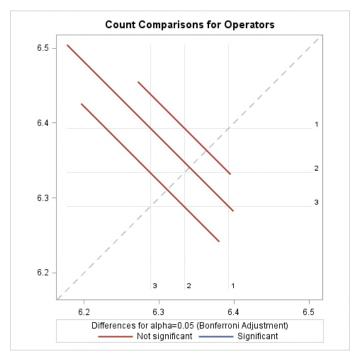


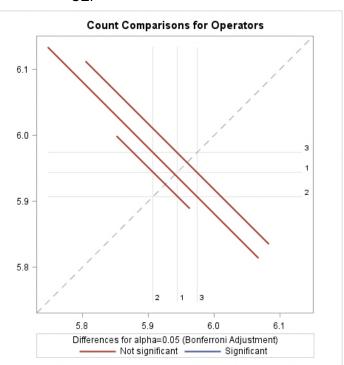
Mock

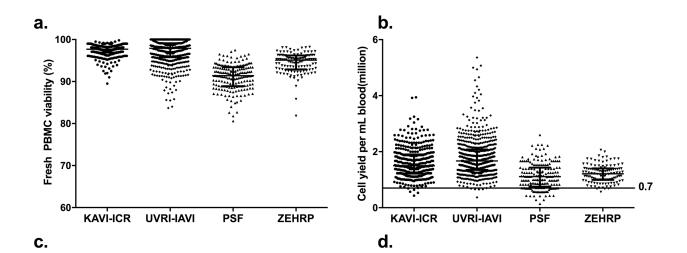
CEF

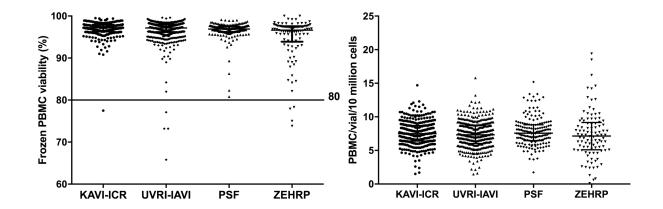


CMV

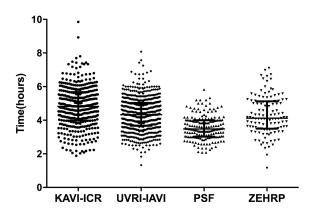


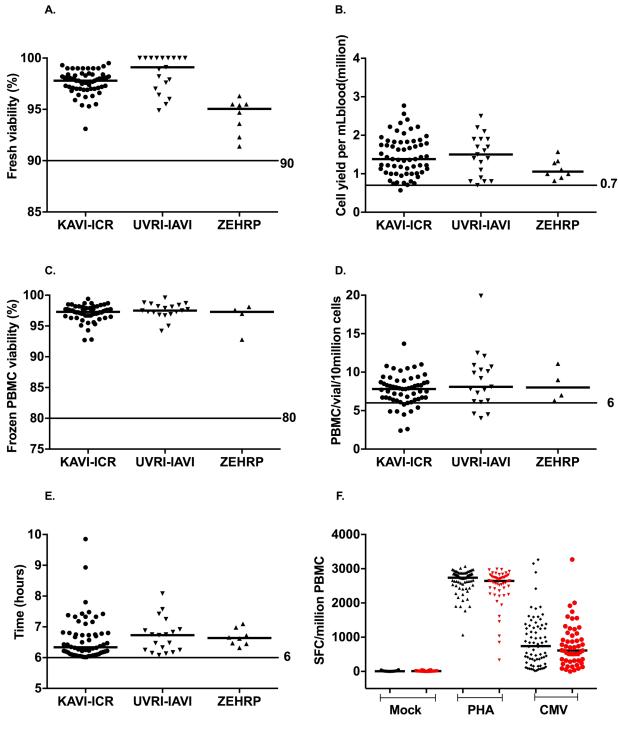






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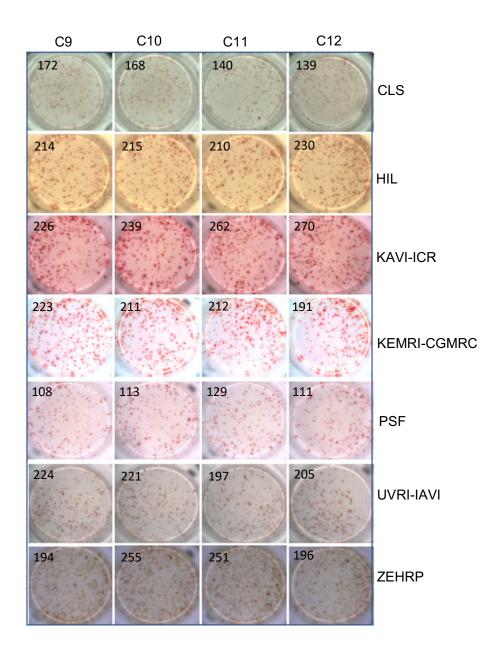




4 PHA

Mock





GCLP Guideline	Process
Development and	Assessment of laboratories' needs, develop required
qualification of collaborating	infrastructure, transfer and qualification of assays
laboratories	
Qualification and/or	Develop qualification and/or validation plans, generate
validation of equipment and	data and review compared against pre-defined criteria
assays	
Equipment service and	Develop calibration plans and document calibration
maintenance	and maintenance of all critical equipment
Development of essential	Development and review of SOPs and other
documents	supporting documents describing safety and
	immunogenicity assessments
Reagent and consumable	Critical reagents are purchased from approved
procurement	vendors according to an approved standardised
	specification
External Quality Assurance	Development of quality assessment program covering
Program	all safety testing parameters, processing, storage and
	shipment of PBMCs and the ELISpot assay
Training Program	GCLP and technical trainings to ensure compliance
	with international standards for conducting clinical
	trials
Evaluation and	GCLP compliance and acceptable technical
Accreditation	performance monitoring by a comprehensive audit
	programme
On-going technology	Transferring of new assays and establishment of
Transfer	separate research programs