1 What is the key question?

Are there serum host marker signatures, which are suitable for point-of-care tests that differentiate between active pulmonary TB and other conditions in individuals presenting with signs and symptoms suggestive of TB in primary health care settings in Africa?

5

6 What is the bottom line?

7 A seven-marker host serum protein biosignature consisting of CRP, transthyretin, IFN-γ, complement

8 factor H, apolipoprotein-A1, IP-10 and serum amyloid A, is promising as a diagnostic biosignature for

9 TB disease, regardless of HIV infection status or African country of sample origin.

10

11 Why read on?

12 The 7 serum marker biosignature identified in this large multi-centered study on 716 individuals with

13 signs and symptoms suggestive of TB could form the basis of a rapid, point-of-care screening test, and

14 with a sensitivity of 94% and negative predictive value of 96%, such a test would render about 75% of

15 the currently performed GeneXpert or TB cultures unnecessary.

16

- Diagnostic Performance of a Seven-marker Serum Protein Biosignature for the Diagnosis
 of Active TB Disease in African Primary Health Care Clinic Attendees with Signs and
 Symptoms Suggestive of TB
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- 21 Novel N. Chegou¹, Jayne S. Sutherland², Stephanus Malherbe¹, Amelia C. Crampin³, Paul
- 22 L.A.M. Corstjens⁴, Annemieke Geluk⁵, Harriet Mayanja-Kizza⁶, Andre G. Loxton¹, Gian van
- 23 der Spuy¹, Kim Stanley¹, Leigh A. Kotzé¹, Marieta van der Vyver⁷, Ida Rosenkrands⁸, Martin
- 24 Kidd⁹, Paul D. van Helden¹, Hazel M. Dockrell¹⁰, Tom H.M. Ottenhoff⁵, Stefan H.E.
- 25 Kaufmann¹¹, and Gerhard Walzl^{1#} on behalf of the AE-TBC consortium
- 26
- ¹DST/NRF Centre of Excellence for Biomedical Tuberculosis Research and SAMRC Centre
- 28 for Tuberculosis Research, Division of Molecular Biology and Human Genetics, Faculty of
- 29 Medicine and Health Sciences, Stellenbosch University, PO Box 241, Cape Town, 8000, South
- 30 Africa
- ³¹ ²Vaccines and Immunity, Medical Research Council Unit, Fajara, The Gambia
- 32 ³Karonga Prevention Study, Chilumba, Malawi
- ⁴Department of Molecular Cell Biology, Leiden University Medical Centre, PO Box 9600,
- 34 2300 RC Leiden, The Netherlands
- ⁵Department of Infectious Diseases, Leiden University Medical Centre, PO Box 9600, 2300
- 36 RC Leiden, The Netherlands
- ⁶Department of Medicine, Makerere University, Kampala, Uganda
- ⁷School of Medicine, Faculty of Health Sciences, University of Namibia, Namibia
- ⁸Department of Infectious Disease Immunology, Statens Serum Institut, Copenhagen 2300s,
- 40 Denmark
- 41 ⁹Centre for Statistical Consultation, Department of Statistics and Actuarial Sciences,
- 42 Stellenbosch University, Cape Town, South Africa
- ¹⁰Department of Immunology and Infection, London School of Hygiene and Tropical
 Medicine, Keppel Street, London WC1E 7HT, UK
- ¹¹Department of Immunology, Max Planck Institute for Infection Biology, Charitéplatz 1,
 10117 Berlin, Germany
- 47
- #Corresponding Author: Gerhard Walzl, DST/NRF Centre of Excellence for Biomedical
 Tuberculosis Research and SAMRC Centre for Tuberculosis Research, Division of Molecular
 Biology and Human Genetics, Faculty of Medicine and Health Sciences, Stellenbosch

- 51 University, PO Box 241, Cape Town, 8000, South Africa, Telephone: +27219389158, Fax:
- 52 +27219389863, E-mail: <u>gwalzl@sun.ac.za</u>
- 53

54 Alternate (Pre-publication) corresponding Author: Novel N. Chegou, DST/NRF Centre of

- 55 Excellence for Biomedical Tuberculosis Research and SAMRC Centre for Tuberculosis
- 56 Research, Division of Molecular Biology and Human Genetics, Faculty of Medicine and
- 57 Health Sciences, Stellenbosch University, PO Box 241, Cape Town, 8000, South Africa,
- 58 Telephone: +27219389069, Fax: +27219389863, E-mail:
- 59 <u>novel@sun.ac.za</u>/novel01@gmail.com
- 60
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69 ABSTRACT

70 Background

User-friendly, rapid, inexpensive yet accurate TB diagnostic tools are urgently needed at points-of-care in resource-limited settings. We investigated host biomarkers detected in serum samples obtained from adults with signs and symptoms suggestive of TB at primary health care clinics in five African countries (Malawi, Namibia, South Africa, The Gambia, and Uganda), for the diagnosis of TB disease.

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77 Methods

We prospectively enrolled individuals presenting with symptoms warranting investigation for pulmonary TB, prior to assessment for TB disease. We evaluated 22 host protein biomarkers in stored serum samples using a multiplex cytokine platform. Using a pre-established diagnostic algorithm comprising of laboratory, clinical and radiological findings, participants were classified as either definite TB, probable TB, questionable TB status or non-pulmonary TB.

84

85 **Results**

Of the 716 participants enrolled, 185 were definite and 29 were probable TB cases, six had questionable TB disease status, whereas 487 had no evidence of TB. A seven-marker biosignature of CRP, transthyretin, IFN- γ , CFH, apolipoprotein-A1, IP-10 and SAA identified on a training sample set (n=491), diagnosed TB disease in the test set (n=210) with sensitivity of 93.8% (95% CI, 84.0-98.0%), specificity of 73.3% (95% CI, 65.2-80.1%), and positive and negative predictive values of 60.6% (95% CI, 50.3-70.1) and 96.4% (95% CI, 90.5-98.8%) respectively, regardless of HIV infection status or study site.

93

94 **Conclusion**:

We have identified a seven-marker host serum protein biosignature for the diagnosis of TB
disease irrespective of HIV infection status or ethnicity in Africa. These results hold promise
for the development of a field-friendly point-of-care screening test for pulmonary TB.

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101 **INTRODUCTION:**

102 Tuberculosis (TB) remains a global health problem with an estimated 9.6 million people reported to have fallen ill with the disease and 1.5 million deaths in 2014¹. Sputum smear 103 microscopy, which has well described limitations, particularly sensitivity², remains the most 104 commonly used diagnostic test for TB in resource-constrained settings. Mycobacterium 105 106 tuberculosis (M.tb) culture, the reference standard test, has a long turn-around time², is 107 expensive, prone to contamination and is not widely available in resource-limited settings. The 108 GeneXpert MTB/RIF sputum test (Cepheid Inc, Sunnyvale, CA), arguably the most important 109 commercial recent advance in the TB diagnostic field yields results within 2hours, coupled 110 with the detection of rifampicin resistance. The Xpert test has been massively rolled out in 111 developed countries but limitations, including relatively high operating costs and infrastructural requirements³, hamper its use in resource-constrained settings. An important 112 limitation of diagnostic tests based on sputum, is that they are unsuitable in individuals, 113 114 particularly children, who have difficulty in providing good quality sputum⁴, and also in individuals with extrapulmonary TB. There is an urgent need for alternative diagnostic tests 115 116 that are suitable for use in all patient types, especially in resource-poor settings. Tests based on the detection of host inflammatory molecules^{5;6} may be beneficial, especially when applied to 117 easily available samples such as finger-prick blood or serum. 118

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In search of immunodiagnostic tools that could be useful for the diagnosis of active TB, attempts are being made to identify novel antigens⁷⁻⁹. Those currently used in the Interferongamma (IFN- γ) release assays (ESAT-6/CFP-10/TB7.7) cannot differentiate between latent and active TB. There is also a search for host markers other than IFN- γ , that are produced after overnight stimulation of blood cells with ESAT-6/CFP-10/TB7.7¹⁰⁻¹⁴, and antibodies against novel *M.tb* antigens^{15;16}.

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Although some T-cell-based approaches¹⁷ are promising for the diagnosis of active TB, overnight culture-based assays are not optimal as point-of-care tests. The importance of diagnosis of individuals with TB disease at the first patient contact and real-time notification to TB programs cannot be overemphasized, as delays in these steps lead to delays in the initiation of treatment and substantial loss to follow-up¹⁸. Therefore, diagnostic tests that can be easily performed at points-of-care by healthcare providers, without the need for sophisticated laboratory equipment will contribute significantly to the management of TBdisease.

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We conducted a study investigating the potential of protein serum host markers to identify pulmonary TB in primary health care clinic attendees from five African countries. Our aim was to further investigate the diagnostic potential of biosignatures identified in our own unpublished pilot studies in a relatively large cohort of study participants, from different regions of the African continent, as such biosignatures might be useful as point-of-care tests for TB disease.

142

143 **METHODS**

144 Study participants

145 We prospectively recruited adults who presented with symptoms requiring investigation for 146 pulmonary TB disease at primary health care clinics at five field sites in five African countries. 147 The clinics served as field study sites for researchers at Stellenbosch University (SUN), South 148 Africa; Makerere University (UCRC), Uganda; Medical Research Council Unit (MRC), The 149 Gambia; Karonga Prevention Study (KPS), Malawi; and the University of Namibia (UNAM), 150 Namibia, as part of the African European Tuberculosis Consortium (AE-TBC) for TB 151 Diagnostic Biomarkers (www.ae-tbc.eu). Study participants were recruited between November 152 2010 and November 2012. All study participants presented with persistent cough lasting ≥ 2 153 weeks and at least one of either fever, malaise, recent weight loss, night sweats, knowledge of 154 close contact with a TB patient, haemoptysis, chest pain or loss of appetite. Participants were 155 eligible for the study if they were 18 years or older and willing to give written informed consent 156 for participation in the study, including consent for HIV testing. Patients were excluded if they 157 were pregnant, had not been residing in the study community for more than 3 months, were 158 severely anaemic (haemoglobin <10g/l), were on anti-TB treatment, had received anti-TB 159 treatment in the previous 90 days or if they were on quinolone or aminoglycoside antibiotics 160 during the past 60 days. The study protocol was approved by the Health Research Ethics 161 Committees of the participating institutions.

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163 Sample collection and microbiological diagnostic tests

Harmonized protocols were used for collection and processing of samples across all study sites.
Briefly, blood samples were collected at first contact with the patient, in 4-ml plain BD
vacutainer serum tubes (BD Biosciences) and transported within 3 hours at ambient

167 temperature to the laboratory, where tubes were centrifuged at 2500 rpm for 10 minutes, after which serum was harvested, aliquoted and frozen (-80°C) until use. Sputum samples were 168 169 collected from all participants and cultured using either the MGIT method (BD Biosciences) 170 or Lowenstein-Jensen media, depending on facilities available at the study site. Specimens 171 demonstrating growth of microorganisms were examined for acid-fast bacilli using the Ziehl-172 Neelsen method followed by either Capilia TB testing (TAUNS, Numazu, Japan) or standard 173 molecular methods, to confirm the isolation of organisms of the *M.tb* complex, before being 174 designated as positive cultures.

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176 Classification of study participants and reference standard

177 Using a combination of clinical, radiological, and laboratory findings, participants were 178 classified as definite TB cases, probable TB cases, participants without pulmonary TB (no-179 PTB) or questionable disease status as described in table 1. Briefly, No-PTB cases had a range 180 of other diagnoses, including upper and lower respiratory tract infections (viral and bacterial 181 infections, although attempts to identify organisms by bacterial or viral cultures were not 182 made), and acute exacerbations of chronic obstructive pulmonary disease or asthma. In 183 assessing the accuracy of host biosignatures in the diagnosis of TB disease, all the definite and probable TB cases were classified as "TB", and then compared to the no-PTB cases, whereas 184 185 questionables were excluded from the main analysis (Figure 1).

186

Classification	Definition
	Sputum culture positive for MTB
Definite TB	OR
	2 positive smears and symptoms responding to TB treatment
	OR
	1 Positive smear plus CXR suggestive of PTB
	1 positive smear and symptoms responding to TB treatment
	OR
Probable TB	CXR evidence and symptoms responding to TB treatment
	Positive smear(s), but no other supporting evidence
	OR
	CXR suggestive of PTB, but no other supporting evidence.
Ouestionable	OR
X	Treatment initiated by healthcare providers on clinical suspicion only.
	No other supporting evidence
	Negative cultures, negative smears, negative CXR and treatment never
No-PTB	initiated by healthcare providers

187 Table 1: Harmonized definitions used in classifying study participants

- 188 Abbreviations: CXR, chest X ray; MTB, *Mycobacterium tuberculosis*; TB, pulmonary
- 189 tuberculosis, No-PTB, non-"pulmonary tuberculosis".
- 190

191 Multiplex immunoassays

Using the Luminex technology, we evaluated the levels of 22 host biomarkers including 192 193 interleukin-1 receptor antagonist (IL-1ra), transforming growth factor (TGF)- α , IFN- γ , IFN- γ -194 inducible protein (IP)-10, tumour necrosis factor (TNF)- α , IFN- α 2, vascular endothelial 195 growth factor (VEGF), matrix metallo-proteinase (MMP)-2, MMP-9, apolipoprotein A-1 196 (ApoA-1), Apo-CIII, transthyretin, complement factor H (CFH) (Merck Millipore, Billerica, 197 MA, USA), and C-reactive protein (CRP), serum amyloid A (SAA), serum amyloid P (SAP), 198 fibrinogen, ferritin, tissue plasminogen activator (TPA), procalcitonin (PCT), haptoglobulin 199 and alpha-2-macroglobulin (A2M) (Bio-Rad Laboratories, Hercules, CA, USA). Prior to 200 testing, samples for MMP-2 and MMP-9 were pre-diluted 1:100 following optimization 201 experiments. Samples for all other analytes were evaluated undiluted, or diluted as 202 recommended by the different manufacturers in the package inserts. The laboratory staff 203 performing the experiments were blinded to the clinical groups of study participants. All assays 204 were performed and read in a central laboratory (SUN) on the Bio-Plex platform (Bio-Rad), 205 with the Bio-Plex Manager[™] Software version 6.1 used for bead acquisition and analysis.

206

207 **Statistical analysis**

208 Differences in analyte concentrations between participants with TB disease and those without 209 TB were evaluated by the Mann-Whitney U-test for non-parametric data analysis. The 210 diagnostic accuracy of individual analytes was investigated by receiver operator characteristics 211 (ROC) curve analysis. Optimal cut-off values and associated sensitivity and specificity were selected based on the Youden's index¹⁹. The predictive abilities of combinations of analytes 212 were investigated by General discriminant analysis (GDA)²⁰ and random forests²¹, following 213 214 the training/test set approach. Briefly, patients were randomly assigned into the training set 215 (70% of study participants, n=491) or test set (30%, n=210), regardless of HIV infection status 216 or study site by the software used in data analysis (Statistica, Statsoft, Ohio, USA). These 217 training and test sets were selected using random sampling, stratified on the dependent (TB) 218 variable. The most accurate of the top 20 marker combinations identified in the training set 219 were then evaluated on the test sample set.

220

221 RESULTS

- A total of 716 individuals were prospectively evaluated in the current study. One study participant was found to be pregnant at the time of recruitment, and data for 8 other participants were not appropriately captured. These 9 individuals were excluded from further analysis (Figure 1). Table 2 shows participant characteristics.
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Table 2: Clinical and demographic characteristics of study participants. The number and
 characteristics of participants enrolled from the different study sites are shown

Study site	SUN	MRC	UCRC	KPS	UNAM	Total
Participants (n)	161	209	171	117	49	707
Age, mean±SD,	37.4±11.3	34.9 ± 12.1	33.1±10	39.9±13.6	36.5 ± 9.6	$36.0{\pm}11.8$
yr						
Males, n(%)	68(42)	123(59)	87(51)	59(50)	28(57)	365(52)
HIV pos, n (%)	28(17)	20(10)	28(16)	67(57)	27(55)	170(24)
QFT pos, n(%)	105(69)	83(41)	119(70)	44(38)	35(71)	386(56)
Definite TB,	22(14)	53(25)	59(35)	18(15)	33(67)	185(26)
n(%)						
Probable TB,	4(2)	13(6)	4(2)	3(3)	5(10)	29(4)
n(%)						
Total TB [#] , (n)	26	66	63	21	38	214
No-PTB, n (%)	133(83)	140(67)	108(63)	96(82)	10(20)	487(69)
Questionable,	2(1)	3(1)	0(0)	0(0)	1(2)	6(1)
n(%)						

Table notes: SUN, Stellenbosch University, South Africa; KPS, Karonga Prevention Study,
Malawi; MRC, Medical Research Council Unit, The Gambia; UCRC, Makerere University,
Uganda; UNAM, University of Namibia, Namibia; SD, standard deviation; QFT, Quantiferon
TB Gold In Tube; pos, positive; neg, negative; indet, indeterminate. *‡*Total TB cases = all the
Definite TB + Probable TB cases; TB, Pulmonary TB; No-PTB, non-"pulmonary
tuberculosis".

235

Using pre-established and harmonized case definitions (Table 1), 185 (26.2%) of the study participants were classified as definite pulmonary TB cases, 29 (4.1%) were probable TB cases, representing the active TB group (214 participants; 30.3%), whereas 487 (68.9%) were No-PTB cases and 6 (0.8%) had an uncertain diagnosis (Table 2). The characteristics of the different patient subgroups are shown in Table 3.

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Table 3: Characteristics of TB and no-PTB cases and individuals with "Questionable TB" disease status.

	Definite TB (n=185)	Probable TB (n=29)	ALL TB (n=214)	No-PTB (n=487)	Questionable TB (n=6)
Age, mean±SD, yr	33.8±9.6	36.3±9.6	34.1±9.6	36.8±12.6	36.5±12.0
Males, n(%)	118(64)	14(48)	132(62)	229(47)	4(67)
HIV pos, n(%)	47 (25)	8(28)	55(26)	114(23)	1(17)
QFT pos, n(%)	144 (78)	19(66)	164(78)	221(47)	2(33)
QFT neg, n(%)	28 (15)	10(34)	38(18)	235(49)	3(50)
QFT Indet, n(%)	8 (4)	0(0)	8(4)	19(4)	1(17)

SD, standard deviation; QFT= Quantiferon TB Gold In Tube; pos, positive; neg, negative;
 indet, indeterminate.

253

254 Utility of individual serum biomarkers in the diagnosis of TB disease

255 All serum markers investigated showed significant differences (p<0.05) between the TB cases 256 and No-PTB cases except A2M and MMP-2 (Supplementary Table 1), irrespective of HIV 257 infection status. Concentrations of CFH, CRP, ferritin, fibrinogen, haptoglobulin, IFN-a2, IFN-258 γ , IL-1ra, IP-10, MMP-9, PCT, SAA, SAP, TGF- α , TNF- α , TPA, and VEGF were significantly 259 higher in the TB cases while those of ApoA-1, Apo-CIII, and transthyretin were higher in the 260 no-PTB cases (Supplementary Table 1). When the accuracy for the diagnosis of TB disease 261 was investigated by ROC curve analysis, the areas under the ROC curve (AUC) were between 262 0.70 and 0.84 for 10 analytes: CRP, ferritin, fibrinogen, IFN- γ , IP-10, TGF- α , TPA, transthyretin, SAA and VEGF (Figure 2). Sensitivity and specificity were both >70% for six 263 264 of these analytes, namely; CRP, ferritin, IFN-γ, IP-10, transthyretin and SAA (Supplementary 265 Table 1).

266

Supplementary Table 1: Median levels of analytes detected in serum samples from individuals with pulmonary TB disease (n=214) or no-PTB disease (n=487), and accuracies in the diagnosis of TB disease

Host	No-PTB	ТВ	P-value	AUC	Cut-off	Sensitivity	Specificity
marker	(IQR)	(IQR)			value	(%)	(%)
IL-1ra	8 (0-40)	35 (0-77)	<0.0001	0.63 [0.58-	>33.9	52.2 [45.1-	71.9
				0.68]		59.2]	[67.6-
							75.9]
TGF-α	3 (1-6)	7 (3-13)	<0.0001	0.73 [69.1-	>5.6	62.8 [55.8-	76.0 [72.0-
				77.4]		69.4]	79.8]

248

IP-10	368 (209-	1712 (808-	<0.00001	0.82 [0.79-	>651.7	81.2 [75.2-	75.0 [71.0-
	652)	3558)		0.86]		86.3]	78.8]
TNF-α	7 (3-12)	14 (8-27)	<0.0001	0.69 [0.65-	>9.5	67.2 [60.3-	65.0 [60.6-
	~ /			0.74]		73.5]	69.3]
IFN-α2	0 (0-6)	7 (0-19)	<0.0001	0.67 [0.62-	>2.9	59.4 [52.4-	71.3 [67.0-
				0.71]		66.2]	75.3]
IFN-γ	1 (0-3)	9 (3-21)	<0.0001	0.80 [0.76-	>2.8	78.3 [72.0-	74.2 [70.0-
				0.84]		83.7]	78.0]
VEGF	158 (19-	341 (144-	<0.0001	0.70 [0.65-	>269.8	60.4 [53.4-	72.5 [68.3-
	286)	624)		74]		67.1]	76.5]
MMP-2	175792	92881	0.091	0.54 [0.49-	<254965	64.3 [57.3-	44.6 [40.1-
	(28693-	(22348-		0.59]		70.8]	49.2]
	474927)	312697)					
MMP-9	401540	651549	0.0004	0.59 [0.53-	> 525174	0.56 [0.49-	0.62 [0.58-
	(155072-	(43831-		0.64]		0.63]	0.66]
	756297)	1299700)					
ApoA-1	2593900	1999800	<0.0001	0.69 [0.65-	<	0.57 [0.50-	0.72 [0.68-
	(2101500-	(1493900-		0.73]	2.17e+006	0.64]	0.76]
. ~	3847700)	2604300)	0.0004	0 65 50 61	0 65 400	0 50 50 60	
Apo C-	261321	180967	<0.0001	0.65 [0.61-	< 265480	0.70 [0.63 -	0.50 [0.45-
111	(178708-	(115790-		0.70]		0.76]	0.55]
T	418395)	297779)	.0.0001	0 70 10 74	· 200505	0.72 [0.66	0 72 10 69
I ranstn	411528	184107	<0.0001	0.78 [0.74-	< 280585	0.73 [0.00-	0.75 [0.08-
yreun	(201039 - 501772)	(10/320 - 201498)		0.82]		0.79]	0.70]
CEU	591775)	291400) 760622	0.0012	0.58 [0.53	< 683022	0.61 [0.54	0 53 [0 40
СГП	(515681	(500474	0.0013	0.58 [0.55-	> 083022	0.01 [0.34 -	0.53 [0.49
	(313081-	1008200)		0.02]		0.08]	-0.58]
Δ2M	1770000	1380700	0 141	0 54 [0 49-	<	0 48 [0 41-	0 61 [0 57-
	(712956-	(501530-	0.141	0.54 [0.42	$1.26e \pm 0.06$	0 55]	0.61
	3273400)	3284700)		0.00]	1.2001000	0.55]	0.00]
Hantogl	955718	2774400	0.0001	0.62 [0.57-	>	0.47 [0.40-	0.71 [0.67
obulin	(287186-	(443581-	0.0001	0.66]	6.17e+006	0.541	-0.751
	26796000)	60000000)					1
CRP	1731 (321-	59195 [´]	<0.0001	0.84 [0.81-	> 7251	0.82 [0.76-	0.73 [0.68-
	9686)	(14047-		0.87]		0.87]	0.76]
		136520)					
SAP	46609	63664	0.0011	0.58 [0.53-	> 63321	0.50 [0.43-	0.67 [0.63-
	(23028-	(20776-		0.63]		0.57]	0.71]
	81115)	129181)					
РСТ	4259	6807	<0.0001	0.68 [0.63-	> 5245	0.70 [0.63-	0.61 [0.56-
	(2474-	(4399-		0.72]		0.76]	0.65]
	6776)	10000)					
Ferritin	33894	158610	<0.0001	0.78 [0.75-	> 69684	0.71 [0.64-	0.70 [0.66-
	(13921-	(61712-		0.82]		0.77]	0.74]
	83571)	365165)			_		
ТРА	1638 (931-	2977	<0.0001	0.72 [0.68-	> 2163	0.70 [0.63-	0.66 [0.61-
	2604)	(1949-		0.76]		0.76]	0.70]
		4317)					

Fibrinog	2466	3987	0<0.0001	0.73 [0.69-	> 2854	0.80 [0.74-	0.57 [0.53-
en	(1804-	(2991-		0.77]		0.85]	0.62]
	4182)	6555)					
SAA	771 (279-	6778	<0.0001	0.83 [0.80-	> 3113	0.86 [0.80-	0.71 [0.67-
	3985)	(4265-		0.86]		0.90]	0.75]
		9689)					

Abbreviations: CFH, complement factor H; A2M, alpha-2-macroglobulin; CRP, C-reactive protein; SAP, serum amyloid P; SAA, serum amyloid A; PCT, procalcitonin; TPA, tissue plasminogen activator; AUC, area under the ROC curve; ROC, receiver operator characteristics. Both HIV-infected and -uninfected individuals were included in the analysis. The values shown for IFN- α 2, IFN- γ , IL-1ra, IP-10, TGF- α , TNF- α , VEGF, ferritin, PCT and TPA are in pg/ml. All other analyte concentrations are in ng/ml. The values in brackets under AUC, sensitivity and specificity are the 95% Confidence Intervals.

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279 Accuracy of individual host markers in HIV-uninfected study participants

280 We stratified the study participants according to HIV infection status and repeated the ROC 281 curve analysis. No differences were observed in the AUCs for ApoA-1, PCT and MMP-9 in 282 HIV-positive versus HIV-negative participants. However, the AUCs for some of the acute-283 phase proteins including A2M, CRP, ferritin, haptoglobulin, SAP and TPA, were higher in 284 HIV-positive individuals. This was in contrast to the observations for the classical pro-285 inflammatory host markers (IFN- γ , IP-10, TNF- α); the growth factors (TGF- α and VEGF); the 286 blood clotting protein fibrinogen, the thyroxin and retinol transporting protein; transthyretin 287 and CFH, which performed best in HIV-uninfected individuals (Figure 3).

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289 Utility of serum multi-analyte models in the diagnosis of TB disease

290 General discriminant analysis (GDA) models showed optimal prediction of pulmonary TB 291 disease with seven-marker combinations. The most accurate seven-marker biosignature for the 292 diagnosis of TB disease, regardless of HIV infection status, was a combination of ApoA-1, 293 CFH, CRP, IFN-y, IP-10, SAA and transthyretin. Without any model "supervision", this 294 biosignature ascertained TB disease with a sensitivity of 86.7% (95% CI, 79.9-91.5%) and 295 specificity of 85.3% (95% CI, 81.0-88.8%) in the training dataset (n=491; 168 TB and 323 no-296 PTB), and a sensitivity of 81.3% (95% CI, 69.2-89.5%) and specificity of 79.5% (95% CI, 297 71.8-85.5%) in the test dataset (n=210, 77 TB and 133 No-PTB). To improve test performance, 298 we optimised the model for higher sensitivity at the expense of lower specificity, which would 299 allow the test to be used as a screening tool. The amended cut-off values ascertained TB disease 300 with a sensitivity of 90.7% (95% CI, 84.5-94.6%) and specificity of 74.8% (95% CI, 69.879.2%) in the training dataset (n=491), and sensitivity of 93.8% (95% CI: 84.0-98.0) and
specificity of 73.3% (95% CI, 65.2-80.1%) in the test dataset (n=210). The positive and
negative predictive values (NPV) of the biosignature were 60.6% (95% CI, 50.3-70.1 %) and
96.4% (95% CI, 90.5-98.8%), respectively (Table 4). The AUC for the seven-marker
biosignature (determined on the training sample set) was 0.91 (95% CI, 0.88-0.94) (Figure 4).

The random forest modelling approach gave similar prediction accuracies for TB and no-PTB as GDA (87% sensitivity and 83% specificity in the training sample set, and 83% sensitivity and 89% specificity in the test sample set), without selection of any preferred cut-off values. In addition to the seven analytes included in the optimal GDA biosignature, Apo-CIII, ferritin, fibrinogen, MMP-9 and TNF- α were also identified as important contributors to top models by the random forest analysis (Figure 4).

313

Table 4: Accuracy of the seven-marker serum protein biosignature (ApoA-1, CFH,
 CRP, IFN-γ, IP-10, SAA, transthyretin) in the diagnosis of TB disease regardless of HIV
 infection status.

317

Training set (n=491)

	Sensitivity	Specificity	PPV	NPV
%, (n/N)	86.7 (130/150)	85.3 (291/341)	72.2	93.6
95% CI	(79.9-91.5)	(81.0-88.8)	(65.0-78.5)	(90.1-95.9)
Test set (n=210)				
%, (n/N)	81.3(52/64)	79.5(116/146)	63.4	90.6
95% CI	(69.2-89.5)	(71.8-85.5)	(52.0-73.6)	(83.9-94.8)

Accuracy of biosignature after selection of cut-off values optimized for sensitivity

Training	set
(n=491)	

Sensitivity	
-------------	--

Specificity

NPV

PPV

%, (n/N)	90.7 (136/150)	74.8 (255/341)	61.3	94.8
95% CI	(84.5-94.6)	(69.8-79.2)	(54.5-67.6)	(91.2-97.0)

Test set (n=210)

%, (n/N)	93.8 (60/64)	73.3 (107/146)	60.6	96.4
95% CI	(84.0-98.0)	(65.2-80.1)	(50.3-70.1)	(90.5-98.8)

318

320

319 Accuracy of the seven-marker biosignature in smear and culture negative patients

321 We evaluated the accuracy of the biosignature in classifying all study participants as TB disease 322 or "no-TB" regardless of the results of the reference standard, and particularly focused on 323 patients who were missed by the microbiological tests (smear and culture) but diagnosed with 324 TB disease based on clinical features including chest X-rays and response to TB treatment 325 (Table 1). The biosignature correctly classified 74% (17/23) of patients who were smear negative but culture positive, and 67% (6/9) of patients who were both smear and culture 326 327 negative. However, the biosignature only correctly classified 88% (86/98) of all the smear 328 positive TB patients, but correctly diagnosed 91% (80/88) of these patients if the smear results 329 were culture confirmed.

330

Accuracy of serum biosignatures in individuals without HIV infection

333 In the absence of HIV infection the GDA procedure indicated optimal diagnosis of TB disease 334 when markers were used in combinations of four with ApoA-1, IFN- γ , IP-10 and SAA 335 constituting the top model with sensitivity of 76.5% (95% CI, 67.5-83.7%) and specificity of 336 91.1% (95% CI 86.7–94.1) in the training sample set (n=372, 115 TB and 257 no-PTB), and a 337 sensitivity of 77.3% (95% CI, 61.8-88.0) and specificity of 87.1% (95% CI, 79.3-92.3%) in 338 the test dataset (n=160, 44 TB and 116 no-PTB). The positive and NPV of the four-marker 339 model in the test set were 69.4% (95% CI, 54.4-81.3%) and 91.0% (95% CI, 83.7-95.4%), 340 respectively.

341

342

343 **DISCUSSION**

We investigated the potential value of 22 host serum protein biomarkers in the diagnosis of TB disease in individuals presenting with symptoms suggestive of pulmonary TB disease at peripheral-level healthcare clinics in five different African countries. Although most of the analytes showed promise individually, the most optimal discriminatory profile was a sevenmarker biosignature comprised of ApoA-1, CFH, CRP, IFN-γ, IP-10, SAA and transthyretin, which might be useful in the rapid diagnosis of TB disease regardless of HIV infection statusor ethnicity in Africa.

351

352 Diagnostic tests based on the detection of host protein biomarkers in ex vivo samples might be 353 more beneficial than antigen stimulation assays as results can potentially be obtained rapidly if 354 lateral flow technologies are employed. Besides the markers that were included in our final 355 seven-marker biosignature (ApoA-1, CFH, CRP, IFN- γ , IP-10, SAA and transthyretin), other 356 analytes including ferritin, fibrinogen, PCT, TGF-a, TNF-a, TPA, and VEGF showed 357 diagnostic potential for TB disease and could have equally been included in the final model in place of any of the seven selected markers. Most of these markers are well known, disease non-358 359 specific markers of inflammation and have been extensively investigated in diverse disease 360 conditions.

361

362 IFN- γ , IP-10 and TNF- α , together with other markers including IL-2 (reviewed in¹⁷), are 363 amongst the most investigated host immunological biomarkers for the diagnosis of *M.tb* 364 infection and disease. Both IFN- γ and IP-10 showed potential in this study. The inclusion of 365 these markers in the seven-marker model is not surprising, given their widely accepted roles in 366 the pathogenesis of *M.tb* infection.

367

368 CRP, ferritin, fibrinogen, SAA, and TPA are acute-phase proteins. The circulating levels of 369 these proteins, as well as those of complement and clotting factors, are known to change by at 370 least 25% in response to inflammatory stimuli, in keeping with their roles in host defense²². CRP (reviewed in²²) is predominantly produced by hepatocytes. The association between 371 372 serum levels of CRP, SAA and TB has long been established, including for treatment 373 response²³. Ferritin is widely known as a biomarker for iron deficiency 24 , and is essential in 374 iron homeostasis in $M.tb^{25}$. Although high levels of ferritin have been observed in many non-375 communicable diseases including cancers, disseminated *M.tb* disease is a common cause of 376 hyperferritinemia^{26;27}.

377

PCT, the precursor molecule of calcitonin is a general inflammatory response marker that is
secreted in healthy individuals by the C cells of the thyroid and by leukocytes via alternate
pathways, including induction by cytokines and bacterial products after microbial infection²⁸.
Although mainly known as a diagnostic marker for bacteremia²⁹, PCT levels have been shown

to be potentially useful in discriminating between pulmonary TB and community-acquired
 pneumonia in HIV-positive individuals³⁰.

384

385 ApoA-1, the major protein component of high-density lipoproteins, and CFH, a crucial 386 regulator of the alternative complement pathway, were also amongst the markers included in our final seven-marker biosignature. ApoA-1 is one of the most important biomarkers for 387 cardiovascular disease³¹, but is rarely investigated as a biomarker in TB. Like the other markers 388 389 investigated in this study, ApoA-1 may not play any specific role in the pathogenesis of TB. 390 The low levels obtained in TB patients may be a result of the many changes in lipid metabolism, 391 which are believed to occur after the generation of the acute-phase response following an 392 inflammatory condition³¹. One of the ways that CFH recognizes host cells is by binding to host markers expressed on the surfaces of cells undergoing apoptosis³². With the help of these 393 394 markers, including CRP and pentraxin 3, CFH ensures proper opsonization of these cells for 395 efficient removal without excessive complement activation during the process, thus limiting 396 immunopathology³². This process is however believed to be exploited by M.tb, to limit opsonization and therefore avoid killing³³. Like ApoA-1, lower levels of CFH were observed 397 398 in the TB cases in this study.

399

Transthyretin (reviewed in³⁴) is a protein that is secreted by the liver into the blood and by the choroid plexus into the cerebrospinal fluid and has been widely investigated as a biomarker for nutritional status³⁴. In previous TB studies, higher levels of transthyretin were observed in TB patients in comparison to uninfected controls³⁵, whereas lower levels were obtained in TB patients as compared to patients with lung cancer, with serum concentrations in TB patients increasing over the course of treatment³⁶. Our observation is in agreement with these reports.

Combinations between transthyretin, CRP, SAA and neopterin ascertained TB disease with 407 78% accuracy in a previous proteomic finger-printing study³⁷. In our study, a biosignature 408 409 containing transthyretin, CRP, SAA and markers involved in Th1-related immunity to TB (IFN- γ , IP-10), an apolipoprotein and CFH showed excellent promise as a diagnostic tool for 410 TB. Although most of these markers are promising individually^{17;23;26;27;30;35-37}, single host 411 412 markers have many shortcomings in predicting TB disease due to poor specificity. As observed 413 in this large multi-centered pan-African study, the accuracy of different host markers is affected differentially by HIV infection. A biosignature containing different classes of biomarkers, 414 415 produced by different cell types such as the classical Th1 immune-related markers plus acute416 phase proteins, complement and apolipoproteins appears to offset the non-specific response 417 patterns of individual or smaller groups of analytes. As a result, markers that perform relatively 418 well in HIV-infected individuals such as the acute phase proteins, help in identifying patients 419 who are missed by markers that may be more often affected by HIV infection such as IFN- γ 420 and IP-10. The resultant test performance with relatively high sensitivity (93.8 %) and high 421 NPV (96.4 %) appears promising as a screening test for active TB disease. Our data indicates 422 that a test based on this biosignature will be superior to smear microscopy and may identify 423 some patients who might be missed by culture.

424

425 The current study stands out in that the investigations were performed in a large number of 426 individuals recruited from peripheral level health care clinics in high-burden settings in 427 multiple countries from different ethnic regions of the African continent. Although there is a 428 need to evaluate the performance of the biosignature in other high TB burdened regions, the 429 inclusion of study participants from these different ethnic regions of the African continent 430 implies that the signature identified in this study may be highly relevant across Africa and 431 perhaps even globally. A limitation of this study was the lack of firmly established alternate 432 diagnoses in the no-PTB group, which is difficult in primary health care settings. This however 433 has no bearing on the importance of our findings as the goal of any TB diagnostic test is to 434 distinguish individuals with TB disease from those presenting with similar symptoms due to 435 conditions other than TB. The utility of this approach in difficult-to-diagnose TB groups such 436 as paediatric and extra-pulmonary TB has to be investigated in future studies. As the HIV 437 infected individuals in this study were not extensively staged with CD4 counts and viral loads, 438 it is not certain whether severe HIV infection might have any influence on the performance of 439 the biosignature. Therefore the influence of severe HIV infection on test performance as well 440 as the effect of anti-retroviral therapy should be investigated in future studies. Future studies 441 should also include samples from confirmed non-TB infectious or inflammatory diseases such 442 as non-TB pneumonia and patients with sarcoidosis and other systemic inflammatory disorders, 443 as such patient groups will be important in ascertaining the specificity of the biosignature for 444 TB.

445

The biosignature identified in the current study warrants further development into a fieldfriendly point-of-care screening test for active TB, potentially based on lateral flow technology^{38;39} and adapted for finger-prick blood. To allow appropriate point-of-care testing in remote settings, the final prototype would include a lightweight portable strip reader with 450 built-in software including an algorithm to interpret results obtained with LF strips comprising multiple cytokine test lines. Such a device is an improvement of the recently investigated UCP-451 LF platform in a multi-site evaluation study in Africa⁴⁰. A cheap point-of-care test, with a high 452 NPV of 96.4%, would identify patients who require confirmatory testing with gold standard 453 tests such as culture and the GeneXpert, which are technically more demanding and have to be 454 455 conducted in a centralized manner. A test with performance characteristics as demonstrated 456 here would render about 75% of the GeneXpert tests currently performed in presumed TB cases 457 for example in South Africa unnecessary, as most of the 70 to 75% of individuals that present 458 with symptoms, are tested, and in whom TB disease is ruled out, would be identified by the 459 point-of-care test, thereby leading to cost savings. The GeneXpert and culture tests could then 460 be used as confirmatory tests in individuals with positive point-of-care test results and for drug 461 susceptibility testing.

462

463 Conclusion

We have identified a promising seven-marker serum host protein biosignature for the diagnosis of active pulmonary TB disease in adults regardless of HIV infection status or ethnicity. These results hold promise for further development into a field-friendly point-of-care test for TB.

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501 **COMPETING INTERESTS**

502 Chegou NN, Walzl G and Mihret A are listed as inventors on an international patent application
503 on the work reported in this manuscript, application no: PCT/IB2015/051435, Filing date:
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634 FIGURE LEGENDS

Figure 1: STARD diagram showing the study design and classification of study
participants. CRF, case report form; TB, Pulmonary tuberculosis; No-PTB, Individuals
presenting with symptoms and investigated for pulmonary TB but in whom TB disease was
ruled out; ROC, Receiver operator characteristics.

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Figure 2: Levels of host markers detected in serum samples from pulmonary TB cases (n=214) and individuals without TB disease (n=487) and receiver operator characteristics (ROC) plots showing the accuracies of these markers in the diagnosis of pulmonary TB disease, regardless of HIV infection status. Representative plots for CRP, SAA, IP-10, ferritin, IFN- γ and transthyretin are shown. Error bars in the scatter-dot plots indicate the median and Inter-quartile ranges.

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Figure 3: Areas under the ROC curve for individual analytes. AUCs obtained after data from pulmonary TB and no-PTB patients were analysed after stratification according to HIV infection status is shown as histograms (A) or 'Before and after' graphs (B). Host markers that performed better in HIV infected individuals are indicated by an asterix.

652

653 Figure 4: Inclusion of different analytes into host biosignatures for the diagnosis of TB 654 disease. (A) Frequency of analytes in the top 20 most accurate GDA seven-marker biosignatures for diagnosis of TB disease regardless of HIV infection status. (B) Importance 655 656 of analytes in diagnostic biosignatures for pulmonary TB disease, irrespective of HIV infection as revealed by random forests analysis. (C) ROC curve showing the accuracy of the finally 657 658 selected seven-marker GDA biosignature in the diagnosis of pulmonary TB disease irrespective 659 of HIV status. (D) Frequency of analytes in the top 20 GDA biosignatures for diagnosis of TB 660 disease in HIV-uninfected individuals. The ROC curve for TB Vs. No-PTB, regardless of HIV 661 (C) was generated from the training dataset.

662