

1 **Diagnostic yield of tuberculosis investigations on bone**
2 **marrow biopsy samples in HIV positive participants at**
3 **Groote Schuur Hospital**

4
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24
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39 **Abstract:**

40 [246 /250 words]

41 **Background:**

42 Acid fast bacilli (AFB) staining on bone marrow samples has low sensitivity for diagnosing HIV-
43 associated tuberculosis and Tuberculosis (TB) culture results may be delayed. The GeneXpert®
44 MTB/RIF Ultra assay may provide a more sensitive diagnostic test on bone marrow biopsy samples.

45 **Methods:**

46 We conducted a two-stage study in a tertiary hospital in South Africa, initially assessing the
47 retrospective yield of TB diagnoses on bone marrow biopsies in adult HIV-positive participants
48 retrospectively from 01-01-2019 to 31-07-2020. Subsequently, determining the prospective yield and
49 diagnostic performance of the GeneXpert® MTB/RIF Ultra assay on bone marrow aspirate and
50 peripheral blood samples in adult HIV-positive participants undergoing bone marrow biopsy from 11-
51 08-2020 to 31-01-2021.

52 **Results:**

53 One hundred and twenty-two biopsies were analysed, of which 59/122 were performed for
54 haematological malignancy staging. Granulomata with AFB were detected in six samples, and nine
55 new lymphoma diagnoses were made. Bone marrow TB culture detected only one non-tuberculous
56 mycobacterial infection. All 17 participants who had TB diagnosed from another clinical site were
57 bone marrow TB culture negative.

58 TB treatment was confirmed in 11/33 participants recruited prospectively. One trace positive
59 GeneXpert® MTB/RIF Ultra result on peripheral blood was detected. All TB cultures on bone
60 marrow aspirates and peripheral blood were negative.

61 **Conclusion:**

62 In a tertiary care hospital in South Africa, the utility of TB culture and GeneXpert® MTB/RIF Ultra
63 on bone marrow aspirate specimens in HIV-positive participants was limited. We postulate that the
64 initiation of empiric anti-tuberculosis treatment could have resulted in false negative results.

65

66 **INTRODUCTION**

67 [453 words]

68 Tuberculosis (TB) is a leading cause of death from infectious diseases in South Africa.⁽¹⁾ Participants
69 with TB and HIV co-infection experience diagnostic delays and have high early mortality.⁽¹⁾
70 Participants with disseminated TB may present with cytopenias, and bone marrow biopsy may be

71 used as part of the diagnostic evaluation in the hospital setting. *Mycobacterium tuberculosis* (MTB)
72 can be identified on bone marrow trephine biopsies using the Ziehl-Neelsen (ZN) stain for acid fast
73 bacilli (AFB), however, this has inadequate sensitivity and specificity.⁽²⁾ TB culture of bone marrow
74 aspirates may take up to 6 weeks for a result. Bone marrow biopsy is an invasive investigation with
75 associated costs. Assessing the yield and diagnostic performance of TB investigations on bone
76 marrow samples is therefore important.

77 The GeneXpert® MTB/RIF assay (Cepheid) is a rapid, cartridge based, real-time polymerase chain
78 reaction (PCR) assay used for the diagnosis of TB on sputum as well as certain extra-pulmonary
79 samples. Its sensitivity on sputum is reduced in participants with HIV co-infection as their sputum
80 may be paucibacillary.⁽³⁾ Its performance on extra-pulmonary samples is variable, with some sample
81 types, such as serous effusions, showing poor sensitivity and yield. A previous study using the
82 GeneXpert® MTB/RIF on bone marrow aspirate samples reported sensitivity to be only 11% when
83 compared to histological findings.⁽⁴⁾ The World Health Organization (WHO) recently recommended
84 the GeneXpert® MTB/RIF Ultra, which has higher sensitivity, to replace the GeneXpert® MTB/RIF
85 in diagnostic algorithms⁽³⁾. South Africa has transitioned to the routine use of the GeneXpert®
86 MTB/RIF Ultra.⁽⁵⁾

87 The GeneXpert® MTB/RIF Ultra assay utilises the same platform as the GeneXpert® MTB/RIF
88 assay, but with a different cartridge and software. The Ultra cartridge makes use of molecular probes
89 to simultaneously detect MTB and rifampicin resistance by targeting two different multi-copy
90 amplification targets (the IS6110 and IS1081 insertion elements) and the Rifampicin Resistance
91 Determining Region of the MTB *rpoB* gene.⁽³⁾

92 Peripheral blood and bone marrow aspirate samples contain high numbers of red blood cells which
93 may inhibit PCR reactions. Recent studies have described techniques which address sample
94 preparation to optimise these types of samples for PCR.^(6,7) A protocol has been validated to remove
95 haem and lysis-concentrate blood without significant loss of bacilli, allowing blood samples to be
96 analysed using the GeneXpert® MTB/RIF Ultra.⁽⁷⁾

97 We conducted a two-stage study at Groote Schuur Hospital in South Africa. First, we evaluated the
98 yield of TB cultures on bone marrow aspirate samples taken from HIV-positive participants
99 retrospectively, over an 18-month period. We then prospectively assessed the yield and diagnostic
100 performance of the GeneXpert® MTB/RIF Ultra assay on bone marrow aspirate and peripheral blood
101 samples from HIV-positive participants undergoing bone marrow biopsy in the routine clinical
102 services at the hospital.

104 **METHODS**

105 [946 words]

106 **Study design and population**

107 **Retrospective review of bone marrow biopsies**

108 For the retrospective cohort study, all bone marrow biopsies performed between 1 January 2019 and
109 31 July 2020 at Groote Schuur Hospital, a tertiary hospital in Cape Town, South Africa were
110 analyzed. Data was collected from the National Health Laboratory Service (NHLS) laboratory
111 records. No clinical folder reviews were performed for the retrospective study. The requirement for
112 informed consent was waived by the ethics committee for this component of the study.

113 **Prospective evaluation of the GeneXpert® MTB/RIF Ultra assay**

114 The GeneXpert® MTB/RIF Ultra assay was evaluated on bone marrow aspirate and peripheral blood
115 samples in a cohort of consecutively enrolled adult HIV-positive participants undergoing bone
116 marrow biopsy over a 6-month period (11 August 2020 to 31 January 2021), at the same hospital. The
117 six-month period was chosen for feasibility as this study was conducted as part of a mini dissertation.
118 The decision to conduct the bone marrow biopsy was made by the attending clinicians and not the
119 study investigators.

120 Informed consent was obtained from all participants successfully recruited into the prospective study.
121 Adult (>18 years old) HIV-positive participants admitted in or referred to Groote Schuur Hospital for
122 a bone marrow biopsy, irrespective of the indication, were eligible for the study. Participants who
123 were already on TB treatment were included in the study.

124 Participants were excluded from the study if they were younger than 18 years of age, or where
125 aspirate samples could not be obtained or were of insufficient volume. HIV-positive participants
126 booked for bone marrow biopsy as part of the routine initial staging and standard of care for known
127 haematological condition were included, however follow-up bone marrow biopsies were excluded.

128 The University of Cape Town Human Research Ethics Committee approved both components of the
129 study (HREC REF: 032/2020). Research activities were performed in accordance with the Declaration
130 of Helsinki and Good Clinical Practice guidelines. Approval from Groote Schuur Hospital
131 management was obtained before data collection.

132 **Data collection**

133 Study data were collected and managed using the REDCap electronic data capture tool (REDCap
134 9.5.36 - © 2021 Vanderbilt University, Nashville, Tennessee, USA) hosted at the University of Cape
135 Town. ⁽⁸⁾

136 **Retrospective review of bone marrow biopsies**

137 Participant data including demographic information, clinical characteristics, HIV results (including
138 recent viral load and CD4 count), tissue histology, staging bone marrow biopsy, all TB culture and
139 GeneXpert® MTB/RIF Ultra results from any specimen within one month (before or after) of the
140 bone marrow biopsy, and information on previous TB (including treatment, start date, duration, and
141 treatment outcome) if available on the laboratory information system, were collected.

142 **Prospective evaluation of the GeneXpert® MTB/RIF Ultra assay**

143 Participants in the prospective cohort had full blood count values, results of radiological
144 investigations (e.g., abdominal ultrasound, chest radiographs, Computed Tomography (CT) and
145 Magnetic Resonance Imaging (MRI) scan reports) and information on TB treatment (including start
146 date, duration, and treatment outcomes if available) recorded, in addition to similar data collected as
147 described for participants in the retrospective study.

148 To minimise experimental bias, individuals performing the GeneXpert® MTB/RIF Ultra assay
149 (laboratory scientists at UCT Division of Medical Microbiology) were blinded to the results of the
150 culture and other TB test results.

151 **Sample collection and processing for the prospective study**

152 We collected bone marrow aspirate and peripheral blood samples from participants who were HIV-
153 positive and referred to our unit for bone marrow biopsy. We used residual bone marrow aspirate
154 samples and collected an additional peripheral blood sample in EDTA tubes. These EDTA tubes were
155 stored at -20 °C and batch-processed on the GeneXpert® MTB/RIF Ultra (Cepheid, Sunnyvale, CA,
156 USA) within 3 weeks of sample collection (Figure 1). We removed red blood cells (RBC) using
157 0.22µm filtered RBC lysis buffer, followed by 2 wash steps and vortexing at 3000-4200g for 25-30
158 minutes (see supplementary material).

159 GeneXpert Ultra testing was performed using the GeneXpert DX System Version 4.8 (Cepheid,
160 Sunnyvale, CA). Samples for GeneXpert were processed in accordance with a previously validated
161 protocol ⁽⁷⁾ (see supplementary material).

162 We attempted to collect 5 to 10 ml of bone marrow aspirate and 5 to 10 ml of peripheral blood in a
163 mycobacterial culture bottle (Myco/F Lytic bottles, Becton Dickinson BD, New Jersey USA)
164 provided by the NHLS. The Myco/F Lytic bottles were placed in the BACTEC incubator unit (Becton
165 Dickinson BD, New Jersey USA) within 6 hours of sample collection, where they were monitored for
166 42 days, and positive bottles screened for mycobacterial growth as per the laboratory standard
167 operating procedure (SOP) (see supplementary material).

168 **Definitions and statistical analysis**

169 Descriptive analysis was used to summarise the study population (e.g. means, medians and
170 interquartile ranges). A TB case was defined as: a positive TB culture or positive GeneXpert®

171 MTB/RIF Ultra on any specimen collected within one month (before or after) of bone marrow sample
172 collection, morphological features of TB on the bone marrow trephine biopsy (the presence of
173 granulomata and AFB on ZN staining which are also negative for PAS), or a strong clinical suspicion
174 of TB. Strong clinical suspicion of TB was defined as the clinical diagnosis of TB made by a medical
175 doctor and a decision made to start TB treatment.

176 Our original aim in the prospective study was to assess and compare the sensitivity and specificity of
177 the GeneXpert® MTB/RIF Ultra on bone marrow aspirate and peripheral blood samples, using TB
178 culture as a reference standard. Because all cultured samples were negative, this was not possible, and
179 therefore report the yield instead. We include the index test on blood and bone marrow when
180 calculating the diagnostic yield. Diagnostic yield was calculated as a proportion with a 95%
181 confidence interval.

182 In instances where there was missing data, group totals were adjusted for analyses (indicated in the
183 text where appropriate).

184

185 **RESULTS**

186 **[959 words]**

187 **Retrospective review of bone marrow biopsies**

188 A total of 760 bone marrow biopsies were performed between 01 January 2019 and 31 July 2020 at
189 Groote Schuur Hospital. Of these, 122 bone marrow biopsies were from confirmed HIV-positive
190 individuals with an accompanying bone marrow TB culture registered and were included in this study.
191 Participant characteristics are summarised in Table 1.

192 Almost half of the included bone marrow biopsies were performed for the staging of a haematological
193 malignancy (n=59), mostly lymphoma (Table 2 and Figure 2). Among these participants, classical
194 Hodgkin Lymphoma (n=23; 38%) and Diffuse Large B-cell Lymphoma (DLBCL) (n=19; 31%) were
195 the most common lymphoma diagnoses. The bone marrow was involved in six (23%) of Hodgkin
196 Lymphoma (HL) participants and nine (24%) of non-Hodgkin Lymphoma (NHL) participants.

197 Forty percent (n=25) of the remaining 63 bone marrow biopsies, which were performed for diagnostic
198 purposes, yielded a unique diagnosis. Of these, nine new cases of lymphoma were diagnosed on bone
199 marrow biopsy and confirmed by immunohistochemical stains; the majority of these were classical
200 HL (n=5) (Figure 3). No non-haematological malignancies were identified. There were four cases of
201 pure red cell aplasia (PRCA).

202 There were 17 participants who had microbiologically confirmed tuberculosis in the cohort. Bone
203 marrow granulomata were identified in seven of these participants, all of whom were on TB treatment
204 before the bone marrow biopsy was performed, and three of them were positive for AFB on ZN

205 staining of the bone marrow trephine. None of these TB diagnoses were made on the bone marrow
206 alone. Concurrent TB and lymphoma were found in five participants, four of whom had Hodgkin
207 Lymphoma, and one who had a diagnosis of Diffuse Large B-cell Lymphoma (DLBCL).

208 When looking at the entire cohort, granulomata were observed in 15% (n=18) (Table 2). These
209 participants had low CD4 counts (median 42 cells/uL), were mostly male (67%, n=12) and had bone
210 marrow biopsies performed for diagnostic purposes. Four of these participants had lymphoma (HL
211 (n=3) and DLBCL (n=1)) and one participant had Kaposi's Sarcoma. Eight of the 18 participants were
212 on TB treatment prior to the bone marrow biopsy (Figure 2 and 3), only four of whom had
213 microbiologically proven TB. No clear cause for the granulomata was identified in the other five
214 participants.

215 Five (4%) participants had bone marrow granulomata on the trephine together with AFB on ZN
216 staining which were negative on Periodic Acid-Schiff (PAS) staining. Only one new diagnosis of
217 probable TB (positive for AFB on ZN staining) was made in the absence of current TB treatment.
218 One participant was diagnosed with non-tuberculous mycobacterial (NTM) infection on bone marrow
219 biopsy (curved bacilli identified on the trephine which were positive for both ZN and PAS staining),
220 which was later confirmed on bone marrow aspirate TB culture.

221 Further testing of the single positive bone marrow NTM culture in our cohort led to the identification
222 of *M. genavense* / *M. triplex* with a PCR/Line probe assay. The assay used (GenoType
223 Mycobacterium AS, Hain Lifescience, Germany) is based on DNA STRIP technology, and due to the
224 same banding pattern of both species, is unable to differentiate between *M. genavense* and *M. triplex*.
225 ⁽¹⁷⁾ PCR and DNA sequencing and were unsuccessful in differentiating the two organisms.

226 No fungal infections were identified on bone marrow histology and only one participant had a
227 diagnosis of lymphoma (HL) and concurrent TB in our cohort.

228 In the cohort of participants who had a diagnostic bone marrow biopsy, a quarter (n=15, 24%) of
229 participants had a new unique diagnosis from the diagnostic bone marrow biopsy. The majority of
230 these participants were on antiretroviral therapy (ART) (n=13) with age (median =35 years, IQR 32-
231 39), a male predominance and median CD4 count of 226 cells/ul.

232 Bone marrow TB culture (n=122) only detected one active case of NTM, and none of the 17
233 microbiologically confirmed cases of TB in the retrospective cohort (Table 3).

234 In summary, there were 17 participants with microbiologically confirmed tuberculosis in our cohort,
235 and two additional participants with presumed TB based on the presence of granulomata and AFB on
236 ZN stain and negative PAS stain.

237 **Prospective testing using GeneXpert® MTB/RIF Ultra assay**

238 Fifty-two individuals were eligible for the study, but only 33 individuals were successfully recruited
239 prospectively, with the others refusing study participation. The majority were inpatients (n=26; 79%),
240 male (n=19; 58%), on ART and had had previous TB (n=18; 55%) (Table 4).

241 Ten participants were on TB treatment for a median duration of 36 days (IQR 6.75-49.5, Range 1-50
242 days), before the bone marrow biopsy was performed. Only six of these participants had
243 microbiologically proven TB: five of these participants had a positive GeneXpert® MTB/RIF Ultra
244 result on sputum, and an additional participant was only sputum TB culture positive (and GeneXpert®
245 MTB/RIF Ultra negative). One positive result (trace- semi-quantitation) was detected using the
246 GeneXpert® MTB/RIF Ultra assay on a peripheral blood sample. This participant had a positive
247 sputum GeneXpert® MTB/RIF Ultra and had been started on treatment 1 week prior to the bone
248 marrow biopsy. All other GeneXpert® MTB/RIF Ultra assays performed on bone marrow aspirate
249 samples were negative. There were four participants where only a bone marrow aspirate was collected
250 without a peripheral blood sample for the GeneXpert® MTB/RIF Ultra assay. One participant's
251 peripheral blood sample was clotted and could not be processed for the GeneXpert® MTB/RIF Ultra
252 assay. No error results were noted on the bone marrow GeneXpert® MTB/RIF Ultra. The diagnostic
253 yield on the GeneXpert® MTB/RIF Ultra assay using peripheral blood was nine percent (1/11), and
254 zero (n=11) using bone marrow aspirate. The diagnostic yield after excluding participants already on
255 TB treatment before samples were collected is still zero using bone marrow aspirates (n=2).

256 All of 66 TB cultures on bone marrow aspirate and peripheral blood samples were negative.

257 **DISCUSSION:**

258 **[855 words]**

259 We have shown there remains a role for bone marrow biopsy in HIV participants in the era of
260 widespread ART use. Diagnoses other than tuberculosis was identified frequently enough in the bone
261 marrow biopsies of participants with cytopenias to warrant routine use of the investigation in clinical
262 practice. In the prospective study, we found only one positive GeneXpert® MTB/RIF Ultra using
263 peripheral blood and none using bone marrow, which may have been the results of the fact that many
264 participants commenced TB treatment prior to sample collection.

265 A previous retrospective review reported on 147 bone marrow biopsies in HIV-positive participants at
266 our hospital who underwent a bone marrow biopsy for the investigation of fever and/or cytopenias
267 (bone marrow biopsies performed for staging of known haematological malignancies were excluded).
268 ⁽⁹⁾One-fifth of these participants were on TB treatment before the bone marrow biopsy, compared to
269 12% in our cohort. ⁽⁹⁾It is worth noting that this analysis was performed in a different era, where not
270 all participants with HIV were eligible for initiation of ART. New bone marrow diagnoses of Classic
271 Hodgkin Lymphoma in our study were comparable to the earlier cohort of participants at the same
272 centre (n=5 vs 4). ⁽⁹⁾

273 The cases of PRCA we diagnosed on bone marrow biopsy were thought to be drug-related, with
274 common viral causes including Parvovirus B19 excluded. Recent reports have implicated
275 emtricitabine as a cause of PRCA, which was thought to be the probable cause in our participants.

276 ^(13,14)

277 There were a surprisingly low number of biopsies with granulomata, which was half the proportion
278 reported in the previous study at our hospital (15%; n=18 vs 33%; n=49)⁽⁹⁾ but similar to a recent
279 study at another hospital in Cape Town, South Africa, that found similarly low levels of TB in their
280 bone marrow samples with granulomas.⁽¹⁵⁾ This was somewhat surprising in a setting that has a very
281 high incidence of TB in communities, in participants who are immunocompromised, though a
282 selection bias may contribute to these findings as only participants who have had several
283 investigations including sputum GeneXpert® MTB/RIF Ultra and imaging (e.g. chest x-rays,
284 abdominal ultrasounds) and still do not have a diagnosis would be referred for a bone marrow biopsy.
285 Bharuthram *et al.*, showed similarly low findings of TB in their Johannesburg cohort.⁽¹⁶⁾ In a recent
286 large retrospective series from Western China- a region with a high prevalence of TB, over 11 000
287 cases were reviewed, with only 0.97% of cases displaying bone marrow granulomata.⁽¹⁷⁾

288 Bone marrow granulomata were identified in seven participants who had microbiologically confirmed
289 TB and were on treatment before the bone marrow however, none of these tuberculosis diagnoses
290 were made on the bone marrow findings alone. Despite the higher percentage of participants on TB
291 treatment before the bone marrow biopsy, in the prior study at our hospital, they still reported 12% of
292 bone marrow TB cultures positive for MTB.⁽⁹⁾ We found that the yield of TB culture on bone marrow
293 aspirate samples was limited in our contemporary study population. This may have been due to the
294 number of participants who were on TB treatment before the bone marrow biopsy, and those whose
295 diagnosis was made purely on radiological and clinical findings.

296 In the prospective arm of the study, we intended to recruit approximately 100 participants, however,
297 we were not able to achieve these numbers due to insufficient referrals. Recruitment for the study
298 began during COVID, and we believe that this did contribute by increasing the threshold for referrals
299 to minimize risk to those performing the procedures. In addition, a few (19) participants did not
300 consent to participation in the study. Among the 33 participants included in this assessment, none had
301 a positive TB culture on bone marrow or blood. We were therefore unable to assess the diagnostic
302 performance of GeneXpert® MTB/RIF Ultra on bone marrow aspirate or peripheral blood, as
303 originally planned, given the lack of positive reference standard tests. Furthermore, these findings
304 suggest that TB culture on bone marrow aspirate samples has limited yield in highly selected
305 participants (those who are hospitalized and have had multiple previous investigations for TB which
306 were negative) or those already on TB treatment. We would have expected TB DNA to persist.

307 We considered that inhibition with haem and/or EDTA could have interfered with the GeneXpert®
308 MTB/RIF Ultra assay performance and thus the diagnostic yield. However, this is unlikely given the
309 negative TB cultures as well as the work done by Boloko *et al* validating the sample preparation
310 protocols used in the study.^(7,18) A pilot study performed at our institution (data not shown) showed
311 minimal inhibition from haem and EDTA in bone marrow samples without the RBC lysis treatment,
312 displaying similar PCR cycle threshold (Ct) values with varying volumes of bone marrow aspirate in
313 EDTA. Comparable results were found in another South African study.⁽⁴⁾ Digital droplet PCR

314 (ddPCR) has shown promise in detecting low levels of circulating *Mycobacterium tuberculosis* DNA
315 in whole blood, noting that ddPCR is highly resistant to the PCR reaction inhibitors such as EDTA,
316 and may be a direction in which future research will be focused.⁽¹⁹⁾

317 **Strengths and limitations of the study**

318 **[242 words]**

319 The retrospective review was conducted over a brief time (18 months) and did not include a clinical
320 folder review which may have limited this study. The clinical information for participants included in
321 the study was variable and based solely on the data provided by the requesting clinician at the time of
322 scheduling the procedure. No consistent criteria for the requesting of diagnostic bone marrow biopsies
323 could be found, and thus procedures were performed based on the clinical decision of the requesting
324 physician in consultation with the haematopathologist.

325 For the prospective study, we could not calculate the performance of the GeneXpert® MTB/RIF Ultra
326 assay in this study due to the lack of positive TB cultures. Sample volumes were not formally
327 measured for the TB cultures, and this may have influenced the diagnostic yield of the tests.
328 Increasing the sample volume of the TB blood cultures may have increased diagnostic yield.^(2, 18) The
329 average sample volume processed for the GeneXpert® MTB/RIF Ultra assay was 2.5 ml. The volume
330 recommended by the manufacturer is 1ml for sputum,⁽²⁰⁾ and the recommended minimum volume is
331 3-5ml according to the protocol by David Barr (see supplementary material). Boloko et al were able to
332 show that using 3–7 mL whole blood in EDTA was sufficient, and the GeneXpert® MTB/RIF Ultra
333 assay was positive in 37% (n= 165 of 447) participants with confirmed tuberculosis by the
334 microbiological reference standard in their cohort.⁽⁷⁾

335 Groote Schuur Hospital is a large referral hospital based in Cape Town, which is affiliated to the
336 University of Cape Town, and participants are mostly referred for advanced specialist care. Its
337 participant population is unlikely to represent the general HIV population as seen in the community,
338 and thus these results may only be generalizable to other tertiary care hospitals in the region.

339 **Conclusion**

340 **[259 words]**

341 The retrospective study highlighted the contemporary utility of the bone marrow biopsy in HIV
342 positive participants, with 17 new unique diagnoses made based on the biopsy, though there were
343 fewer new diagnoses of TB than expected. In the prospective study, TB culture on bone marrow
344 aspirate samples yielded no positive results in this highly selected cohort, one third of whom were
345 already established on TB treatment. In keeping with this, the GeneXpert® MTB/RIF Ultra on bone
346 marrow aspirate and peripheral blood was negative in almost all participants in the prospective study.
347 We postulate that the reason for these findings is that participants with cytopenias due to HIV-
348 associated TB in our setting are diagnosed on other clinical specimens or empirically started on TB

349 treatment, often at primary or secondary levels of care, and therefore not referred for biopsy or only
350 referred if they deteriorate on TB treatment. The diagnoses made on bone marrow biopsy are
351 therefore mainly non-TB diagnoses.

352 Future research on the diagnostic performance and yield of GeneXpert® MTB/RIF Ultra on bone
353 marrow aspirate could involve prospectively recruiting HIV-positive participants with cytopaenias
354 who are admitted to hospital and are being investigated for TB, but in whom initial sputum and urine-
355 based TB diagnostics are negative, and who have not yet been started on TB treatment. This may
356 provide a higher yield of diagnoses using this assay on bone marrow samples.

357 Our findings suggest that routine conduct of TB culture and GeneXpert® MTB/RIF Ultra on bone
358 marrow samples in HIV positive participants where TB is not suspected, may not be warranted.

359

360

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365 work and running the experiments.

366 **Competing interests**

367 The authors have no competing interests (financial, personal, or otherwise).

368 **Authors' contributions**

369 C.S and C.M devised the project, G.M., G.M., D.B and E.V assisted with developing the main
370 conceptual ideas and proof outline. C.S. supervised the project. C.M. assisted with measurements and
371 helped carry out the experiments. S.B performed the numerical calculations and data analysis. S.B.
372 took the lead in writing the manuscript. All authors provided critical feedback and helped shape the
373 research, analysis, and manuscript.

374

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387 applied a CC BY public copyright license to any Author Accepted Manuscript version arising from
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392 **Data availability statement**

393

394 Data will be governed by a Creative Commons license, and researchers will have privileged use of the
395 data for two years to enable the publication of research results, after which it will be made freely
396 available under this license. The other data will be retained for 10 years.

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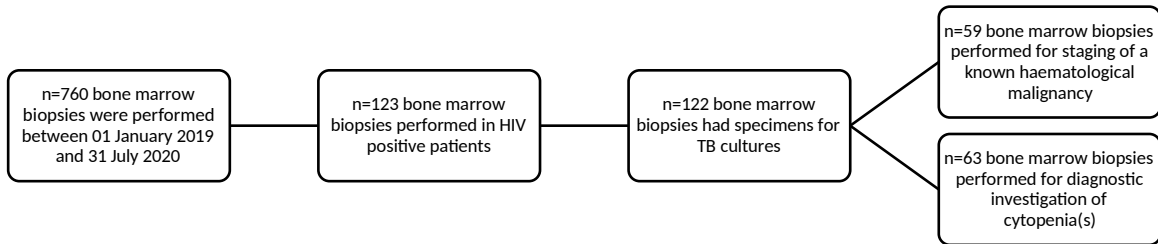
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461

462 **TABLES AND FIGURES**

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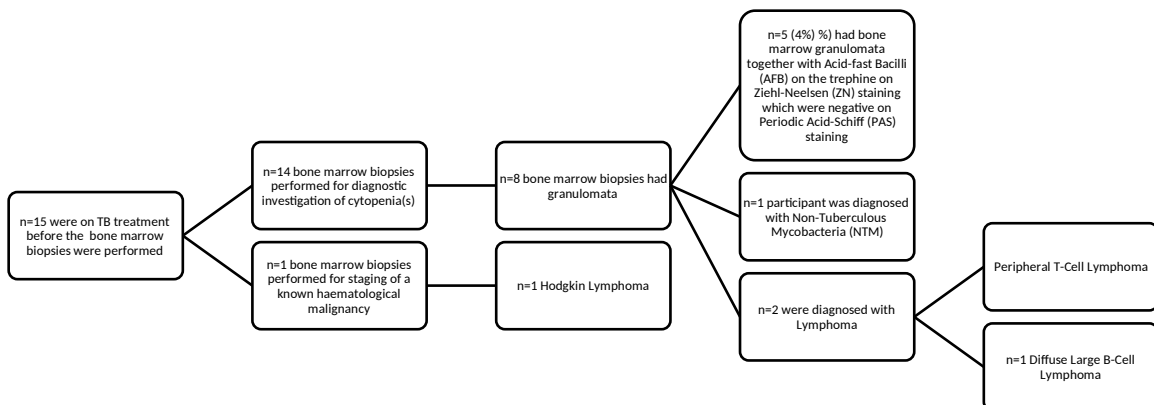


465

466 **Figure 1: Consort diagram of retrospective review**

467 We undertook a retrospective review of all bone marrow biopsies performed between 01 January
 468 2019 and 31 July 2020. One hundred and twenty-three bone marrow biopsies were performed in
 469 confirmed HIV positive individuals. Of these, only one participant did not have a bone marrow TB
 470 culture registered. Just over half (n=63; (52%)) of all the bone marrow biopsies were performed to
 471 investigate the cause of cytopenias in HIV positive individuals.
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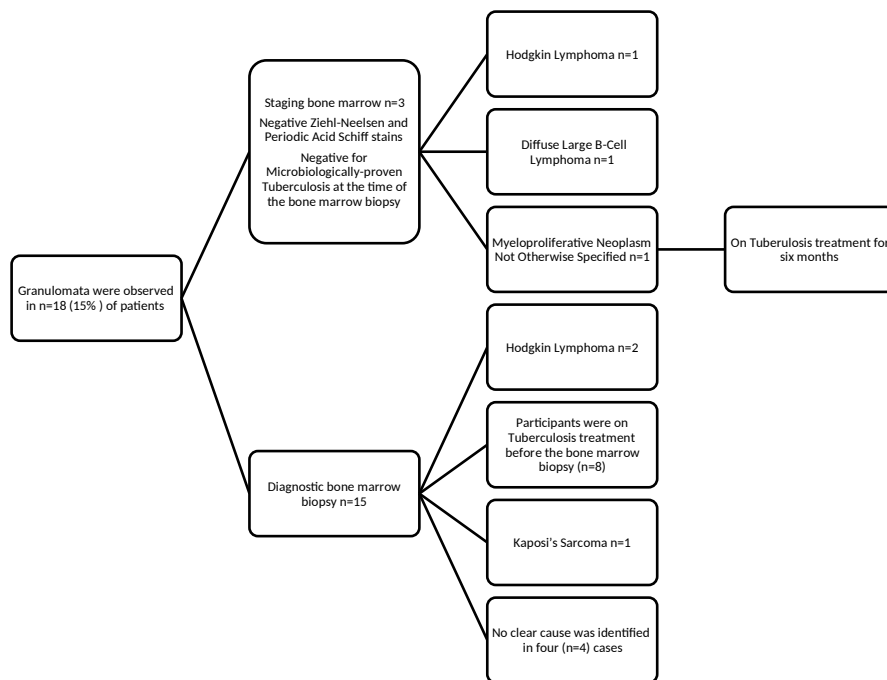
474 **Figure 2: Participants who had been started on TB treatment before the bone marrow biopsy in**
 475 **the retrospective arm of the study**

476 Most participants who were on TB treatment (n=15) before the bone marrow biopsy were referred for
 477 a diagnostic rather than staging bone marrow procedure. Of these participants, 5 (4%) had bone
 478 marrow granulomata together with Acid-fast Bacilli (AFB) on the trephine on Ziehl-Neelsen (ZN)
 479 staining which were negative on Periodic Acid-Schiff (PAS) staining. One participant was diagnosed
 480 with Non-Tuberculous Mycobacteria (NTM) on bone marrow biopsy (curved bacilli identified on the
 481 trephine which were positive for Ziehl-Neelsen (ZN) staining and Periodic Acid-Schiff (PAS)
 482 staining), which was later confirmed on TB bone marrow culture.

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487 **Figure 3: Participants who had granulomata on the bone marrow biopsy in the retrospective**
 488 **arm of the study.**

489 This figure illustrates those participants who had granulomata on their bone marrow. Four of these
 490 participants had Lymphoma (HL (n=3) and DLBCL), one participant had Kaposi's Sarcoma. No clear
 491 cause could be identified in four cases. Eight participants of the 18 were on TB treatment before the
 492 bone marrow biopsy. DLBCL= Diffuse Large B-Cell Lymphoma; HL=Hodgkin Lymphoma MPN
 493 NOS= Myeloproliferative Neoplasm Not Otherwise Specified; PAS= Periodic Acid Schiff stain;
 494 TB=Tuberculosis; ZN= Ziehl-Neelsen stain

495

497 **Table 1: Participant characteristics in the retrospective review of bone marrow TB cultures in**
 498 **HIV positive participants (n=122)**

499 This table describes the participant characteristics of participants in the retrospective arm of the study.
 500 They were mostly male (n=70), virally unsuppressed with a median CD4 count of approximately 200
 501 cells/uL. The majority bone marrow biopsies performed in these participants were for diagnostic
 502 reasons (n=63). Thirteen percent (n=16) of participants had started TB treatment before the bone
 503 marrow biopsy was performed.

Characteristic	n (%) or median (IQR)
Median age in years (IQR)	38 (33-44)
Sex n (%)	
Male	70 (57)
Female	53 (43)
In/Outpatient Status n (%)	
Inpatient	91 (74)
Outpatient	31 (24)
Most recent CD4 count (IQR) n=122	187 cells/uL (42-271)
Most recent HIV viral load (IQR) n=101	569 651 copies/ml (0- 49 367)
Antiretroviral treatment (ART) status (%)	
Antiretroviral naive	12 (10)
Currently on Antiretroviral treatment	81 (67)
Interrupted Antiretroviral treatment	12 (10)
Unknown	17 (14)
Indication for the bone marrow n (%)	
Staging for a known haematological malignancy	59 (48)
Diagnostic investigation of cytopenia (s)	63 (52)
On TB treatment before the bone marrow n (%)	
Yes	15 (12)
No	107 (88)

504 IQR: interquartile range

505 **Table 2: Bone marrow findings in the retrospective study**

Bone marrow (BM) finding:	n (%)
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Granulomata	
Present	18 (15)
Absent	102 (83)
Lymphohistiocytic aggregates	2 (2)
Ziehl-Neelsen stains	
Positive for AFB	6 (5)
Negative	25 (20)
Not done	91 (75)
Periodic Acid Schiff stains	
Positive	1 (1)
Negative	25 (20)
Not done	96 (79)

506

507 **Table 3: TB culture results in the retrospective study**

Specimen type	Positive n (%)	Negative n (%)
TB culture on		
Sputum n=18	2 (11)	16 (89)
Bone marrow n=122	1 (0.8)	121 (99.2)
Blood n=31	3 (10)	28 (90)
Other n=73	4 (6)	69 (94)
CSF n=15		
Pleural fluid n=6	1 (17)	5 (87)
Urine n=12	1 (8)	11 (92)
Joint aspirate n=2	0 (0)	2 (100)
Tissue biopsy n=38	0 (0)	38 (100)
GeneXpert® MTB/RIF Ultra		
Sputum n=54	8 (15)	46 (85)
Other n=29	1 (3)	28 (97)
CSF n=12		

Pleural fluid n=4	0 (0)	4 (100)
Urine n=3	1 (33)	2 (67)
Joint aspirate n=2	0 (0)	2 (100)
Tissue biopsy n=14	0 (0)	14 (100)

508

509

CSF= Cerebrospinal fluid, Tissue = Core needle biopsy or excision biopsy of any organ including liver and lymph nodes

510

511 **Table 4: Prospective study participant characteristics**

Characteristics n=33	n (%) or median (IQR)
Sex	
Male	19 (58)
Female	14 (42)
Most recent CD4 count n=33	201 cells/uL (47-195)
Most recent HIV viral load n=28	493 861 copies/mL (0- 17 845)
Previous TB	18 (55)
ART (Anti-retroviral treatment) status	
Antiretroviral naive	4 (12)
Currently on ART	18 (55)
Interrupted ART	11 (33)
In/Outpatient Status	
Inpatient	26 (79)
Outpatient	7 (21)
TB cases ^a at time the bone marrow was performed:	11 (33)
Median duration of TB treatment (days) before the bone marrow biopsy n=10 (30%); In one patient the treatment duration was unknown	36 (6.75-49.5)
Indication for the bone marrow	
Staging for a known haematological condition	14 (42)
Diagnostic investigation of cytopenias	19 (58)
GeneXpert® MTB/RIF Ultra positive: peripheral blood	1 (trace) ^b in a participant with TB on treatment

GeneXpert® MTB/RIF Ultra positive: bone marrow	0
TB culture positive (Bone Marrow/Peripheral Blood)	0

512 a: A TB case was defined as: a positive TB culture or positive GeneXpert® MTB/RIF Ultra on any specimen collected within one month (before or after) of
513 bone marrow sample collection, morphological features of TB on the bone marrow trephine biopsy (the presence of granulomata and AFB on ZN staining which
514 are also negative for PAS), or a strong clinical suspicion of TB. Strong clinical suspicion of TB was defined as the clinical diagnosis of TB made by a medical
515 doctor and a decision made to start TB treatment.

516 b: Trace-mini semi-quantitation available on the GeneXpert® MTB/RIF Ultra

517

518 **Appendix A-SOP for Xpert-ultra on blood**

519 Developed and validated by David Barr david.barr@liverpool.ac.uk

520 Used with permission.

521 **Reagents / consumables**

522

523 • Sterile Ethylenediaminetetraacetic acid (EDTA) tubes

524 • RBC lysis buffer: [155mM NH₄Cl; 12mM NaHCO₃; 0.1 mM EDTA].

525 Made with 0.22µm Filtered water and 0.22µm filtered again prior to use.

526 • Sterile water: 0.22µm filtered deionised water.

527 • 50 and 15ml centrifuge (falcon) tubes.

528 • Pipettes and centrifuge.

529 • Xpert-ultra lysis buffer and cartridges

530 **A note on volume of blood to collect**

531 The volume to collect is arbitrary – larger volumes will increase sensitivity. The below SOP was
532 optimised for 3-5ml (based on the maximum volume we could safely take from the patient in the
533 experiment design which involved multiple other samples) but could be used for up to 10ml. If you
534 are processing more than 10ml blood the volumes of lysis buffers etc might be too small, and it will
535 affect the pelleting etc, so the simplest thing would be to split the sample, e.g., if you collect 15ml
536 could run that as two 7.5ml samples but combine them at the end for one cartridge.

537 **Method**

538

539 1. Collect blood in a sterile EDTA tubes (purple top).

540 2. Transfer blood to a 50ml falcon tube. Wash out any residual blood in the EDTA tube using
541 RBD lysis buffer and add this to the falcon tube (you do not want to miss any bacilli adhering
542 to the sides of the tube). Make up the falcon tube volume to 45ml with RBC lysis buffer.

543 3. Incubate at room temperature for 30 minutes with regular inversions to mix, then pellet at
544 3000g for 30min. Remove the supernatant by incredibly careful pouring (into a NaOH waste
545 pot in the hood). It is more important to not accidentally pour any of the pellet out than it is to
546 remove all the supernatant. The pellet is soft and easily disperses so I would leave 4-6ml
547 supernatant at the bottom.

548 4. Resuspend the pellet in the residual supernatant volume by repeated reverse pipetting (works
549 better than vortexing). It is essential to resuspend in this smaller volume before moving to
550 next step.

551 5. Make to volume up to 45ml using sterile water. Incubate at room temperature for 30 minutes
552 with regular inversions to mix, then pellet at 3000g for 30min. The pellet should now not be
553 red – should be pale pink or buff colour. Remove supernatant and resuspend in residual
554 volume as above.

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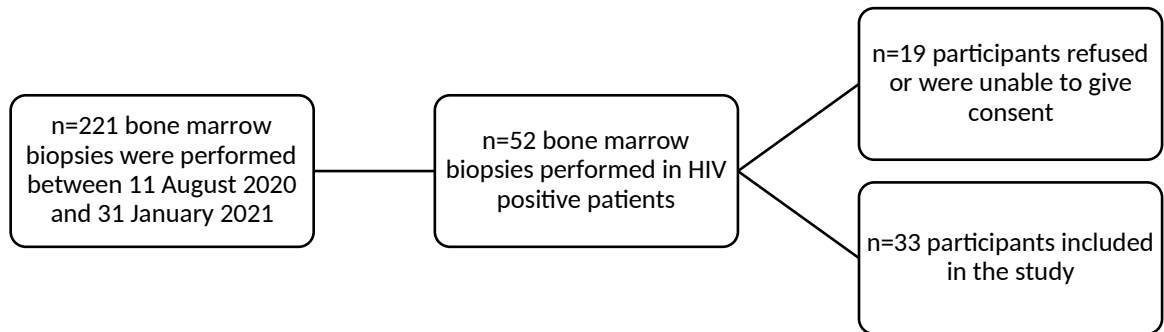
[Samples can be frozen for batching at this stage]

6. Add the sample to a 15ml falcon tube and make volume up to 10 ml using sterile water. Spin at 4200G for 25 minutes and remove 9.3 ml supernatant, leaving ~0.7ml sample. This should be resuspended in the 0.7ml by pipetting.
7. Add 1.5ml of the GeneXpert® MTB/RIF Ultra buffer to each (final volume ~2.2ml – *it should never be less than 2ml*). These are mixed by vigorous shaking 20 times (make sure lid is on) and then incubated at room temperature in hood for 10 minutes, followed by another 20 shakes and further 5-minute incubation.
8. Sample can now be added to the GeneXpert® MTB/RIF Ultra cartridge and processed as per manufacturer instructions.

566

Appendix B: Consort diagram of Prospective study

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Consort diagram of Prospective study

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The GeneXpert® MTB/RIF Ultra assay was evaluated on bone marrow aspirate and peripheral blood samples in a cohort of consecutively enrolled adult HIV-positive participants undergoing bone marrow biopsy over a 6-month period (11 August 2020 to 31 January 2021), at the same hospital. The six-month period was chosen for feasibility as this study was conducted as part of a mini dissertation. Two hundred and twenty-one bone marrow biopsies were performed during this period, 52 bone marrow biopsies performed in HIV positive patients and 33 participants were included in the final study. A total of 19 participants were unable to give informed consent or refused to participate in the study.