

Irreversible depletion of intestinal CD4⁺ T-cells is associated with T-cell activation during chronic HIV infection

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HIV infection in the human gastrointestinal (GI) tract is thought to be central to HIV progression, but knowledge of this interaction is primarily limited to cohorts within westernized countries. Here, we present a large cohort recruited from high HIV endemic areas in South Africa and found that people living with HIV (PLWH) presented at a younger age for investigation in the GI clinic. We identified severe CD4 T-cell depletion in the GI tract, which was greater in the small intestine than in the large intestine and not correlated with years on ART or plasma viremia. HIV-p24 staining showed persistent viral expression, particularly in the colon, despite full suppression of plasma viremia. Quantification of mucosal ARV drugs revealed no differences in drug penetration between the duodenum and colon. Plasma markers of gut barrier breakdown and immune activation were elevated irrespective of HIV, but peripheral T-cell activation was inversely correlated with loss of gut CD4 T-cells in PLWH alone. T-cell activation is a strong predictor of HIV progression and independent of plasma viral load, implying that the irreversible loss of GI CD4 T-cells is a key event in the HIV pathogenesis of PLWH in South Africa, yet the underlying mechanisms remain unknown.

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39

40 **Abstract**

41 HIV infection in the human gastrointestinal (GI) tract is thought to be central to HIV
42 progression, but knowledge of this interaction is primarily limited to cohorts within westernized
43 countries. Here, we present a large cohort recruited from high HIV endemic areas in South
44 Africa and found that people living with HIV (PLWH) presented at a younger age for
45 investigation in the GI clinic. We identified severe CD4 T-cell depletion in the GI tract, which
46 was greater in the small intestine than in the large intestine and not correlated with years on
47 ART or plasma viremia. HIV-p24 staining showed persistent viral expression, particularly in
48 the colon, despite full suppression of plasma viremia. Quantification of mucosal ARV drugs
49 revealed no differences in drug penetration between the duodenum and colon. Plasma
50 markers of gut barrier breakdown and immune activation were elevated irrespective of HIV,
51 but peripheral T-cell activation was inversely correlated with loss of gut CD4 T-cells in PLWH
52 alone. T-cell activation is a strong predictor of HIV progression and independent of plasma
53 viral load, implying that the irreversible loss of GI CD4 T-cells is a key event in the HIV
54 pathogenesis of PLWH in South Africa, yet the underlying mechanisms remain unknown.

55 **Introduction**

56 HIV-1 associated alterations within the gastrointestinal (GI) tract were first described in
57 1984 by Kotler and colleagues, who reported histologic abnormalities in the GI mucosa
58 of infected patients (1, 2). HIV-1 infection and replication is far more prevalent in the CD4
59 T-cells of the GI mucosa (3–6) than in blood CD4 T-cells (5, 7–9), leading to rapid and
60 severe loss of GI CD4⁺ T-cells following infection. The resulting intestinal damage and
61 microbial translocation (10) is thought to precipitate systemic inflammation and chronic
62 immune activation, which ultimately drives disease progression (10). Crucially, these
63 events are not fully reconstituted by antiretroviral treatment (ART) (11–15), and may have
64 long-term health consequences for the growing number of PLWH on ART. This includes
65 an elevated risk of non-infectious comorbidities (16, 17), such as cardiovascular disease,
66 renal failure, liver disease, neurocognitive disease, osteoporosis, diabetes and gut-
67 associated cancers (17–23). However, the direct link between HIV associated intestinal
68 epithelial barrier break down, CD4 loss in the GI tract and HIV pathology has yet to be
69 established, representing a major hindrance in the development of treatment strategies
70 to restore gut homeostasis.

71

72 The intestine is the largest outward facing organ in the body and it contains more activated
73 CD4 T-cells and tissue resident HIV reservoirs than any other human tissue site (24–26).
74 Persistent viral replication within the gut HIV reservoir and high levels of HIV DNA in the
75 gut CD4⁺ memory T-cells is a major obstacle for HIV eradication (6, 27). ART initiation
76 during early acute HIV-1 infection reduces immune activation and limits reservoir size,
77 but does not appear to prevent its formation or lead to its eradication after prolonged

78 treatment (26, 28–30). The reasons for these are not clear, but may include sub-optimal
79 ARV drug penetration into the gut mucosa (14, 27, 31, 32). Indeed, recent data suggests
80 that some regions of the gut may have little or no exposure to antiretroviral drugs during
81 ART (33).

82
83 The majority of studies investigating HIV related gut immunopathology have been
84 conducted on Caucasian populations and/or carried out in westernized countries.
85 Intestinal health and immunity, however, is influenced by genetics, diet, microbiome and
86 comorbidities, and thus varies greatly between different populations. Therefore, the
87 immunopathology of HIV in the GI tract of these study populations may be distinct from
88 Sub-Saharan African populations, where 2/3rd of the global HIV population live. In
89 addition, for practical reasons, previous work has focused primarily on the large intestine.
90 As a result, the impact of HIV infection on the small intestine, where the immune
91 composition, physiology and microbiome are distinct, remains less understood (34–37).

92
93 Here, we present data collected from a large cohort of more than 500 study subjects
94 undergoing clinically indicated endoscopy or colonoscopy that were recruited from
95 extremely high HIV-1 endemic areas within KwaZulu-Natal, South Africa. The HIV
96 infected participants presented to the GI clinic at a younger age and gut pinch biopsies
97 obtained from their colon, rectum and duodenum revealed severe and irreversible HIV
98 associated CD4 T-cell depletion, in particular within the small intestine. This was
99 associated with persistent HIV expression, as shown by histological staining of HIV-p24
100 in gut CD4 T-cells. This was greatest in the colon, in which CD4 depletion was less

101 severe, suggesting target cell availability may affect reservoir size in the GI tract, in
102 addition to the poor penetration of ART drugs observed. The degree of GI CD4 T-cell
103 depletion is directly correlated with immune activation of blood CD4 T-cells only in PLWH,
104 which strongly predicts HIV disease progression, independent of viral load (38). Plasma
105 markers of systemic immune activation, however, were elevated in all study participants
106 and did not correlate with GI CD4 T-cell levels, suggesting these biomarkers are not
107 directly linked to HIV pathology per se. This cohort represents a large and unique resource
108 to further study intestinal health and HIV reservoir dynamics in PLWH from areas of rural
109 Sub-Saharan Africa, which sits at the center of the global HIV epidemic.

110

111 **Results**

112 **Diverse GI diagnoses and increased risk of biliary pathology in PLWH**

113 To study the impact of HIV on the intestinal mucosa, we recruited 529 adult participants
114 from KwaZulu-Natal, South Africa, presenting in the GI clinic for either endoscopy,
115 including ERCP (small intestine) or colonoscopy (large intestine) procedures between the
116 years 2015 and 2019. Participants were from area of high HIV endemicity, and 170 (32%)
117 where infected with HIV (Table 1). Eighty four percent of HIV infected individuals had
118 undetectable plasma HIV viral load due to antiretroviral therapy (ART); with HIV
119 diagnosed for a median of 6.3 years of which 5.0 years were reported on ART and median
120 CD4 counts of 647 cells/mm³. The 29 HIV infected individuals with detectable viremia
121 (VL>20 RNA copies/ml plasma) were diagnosed for a median of 0.3 years with CD4 T-
122 cell counts of 297 cells/mm³ and median viral load of 29,350 RNA copies/ml plasma
123 (Table 1). Less than 3% presented with active Tuberculosis (TB) infection, but previous
124 TB infection were enriched in HIV infected individuals (4% vs 24%, P<0.0001)
125 (Supplemental Table 1) consistent with increased risk of TB infection in HIV infected
126 individuals.

127

128 In this cohort, clinical diagnoses were dominated by non-infectious morbidities such as
129 biliary pathology, gastrointestinal (GI) bleed, esophageal disorders and pain within the
130 chest and abdomen (Supplemental Figure 1). The most common indications for
131 colonoscopy were cancer, polyps, constipation/diarrhoea and ano-rectal disorders, while
132 for upper endoscopies these were dominated by biliary, esophageal and abdominal
133 complications. In 45 participants (9%) no obvious GI related diagnosis following

134 investigation were identified (Supplemental Figure 1). Stratifying participants based on
135 HIV status, showed that females and black ethnicity were enriched in the HIV positive
136 group, consistent with the demographic of the HIV epidemic in this region (Figure 1A).
137 Overall, the GI complication of biliary pathology were enriched in the HIV positive group
138 (OR 1.2, 25.3%, 39/154, P=0.02) by multivariate analysis (Figure 1A). Stratifying clinical
139 diagnoses by ART status, revealed that biliary pathology was enriched in ART treated
140 (OR 3.0, 95% CI 1.8–5.0, P<0.0001), but not in ART untreated individuals (OR 2.0, 95%
141 CI 0.7–5.7, P=0.27) (Figure 1B), while both HIV plasma viral suppressed and viremic
142 participants showed consistent enrichment for biliary pathology (both OR 2.5, 95% CI 1.4
143 to 4.3, P=0.0050) (Figure 1C). Taken together, these clinical diagnosis showed a diverse
144 spectra of GI complications of which frequencies of inflammatory disorders, including
145 inflammatory bowel disease, diverticulitis, prostatitis, ulcerative colitis and inflammation
146 not otherwise specified, were overall less frequent to that observed in Caucasian cohorts
147 (39) and highlights distinct GI related complications in this Sub-Saharan African cohort.
148 However, with the exception of biliary pathology, clinical presentation of HIV infected and
149 uninfected subjects was similar and, therefore, unlikely to significantly skew the
150 comparison of these groups.

151

152 **Early presentation of GI related complications in HIV infected individuals**

153 The HIV epidemic in South Africa is known to be discordant for ethnicity, age and gender
154 (41–43). We therefore stratified our cohort by ethnicity and found a skewed distribution of
155 non-black ethnicity compared to the background provincial data, which is likely to be at
156 least in part explained by differential access to tertiary healthcare units, in addition to

157 potential differences in genetic and environmental factors (Figure 2A, Supplemental Table
158 1). Overall, HIV prevalence was higher in women, reaching >60% infection in the 25-44
159 year age group (Figure 2B), which is consistent with recent population level studies from
160 the same region (40, 41). Strikingly, while females and males presented to the clinic for
161 GI complications at a similar age, HIV infected participants presented a median of 12
162 years earlier than HIV-1 uninfected participants (41 years, IQR, 34-52 vs. 53 years, IQR,
163 40-63 years; $P < 0.0001$) (Figure 2C) irrespective of ART treatment status, with females
164 and males presenting 13 and 6.5 years younger, respectively. We found no age
165 differences within the three indicated procedures (colonoscopy, upper endoscopy and
166 ERCP), but in each procedure, HIV infection was associated with an overall younger
167 population (Figure 2D).

168 In a subset of HIV infected participants full blood parameters where available, including
169 white and red blood cell analysis. Comparison with HIV uninfected controls, this revealed
170 reduced CD4 T-cell (median 858 vs 641 vs 297 cells/ul) and elevated CD8 T-cells (median
171 610 vs 670 vs 1040 cells/ul) counts in both viral suppressed and viremic groups compared
172 to HIV uninfected participants (Suppl Figure 2). In addition, HIV infection was associated
173 with elevated neutrophils and basophils, and with increased neutrophil to lymphocyte ratio
174 (NLR), suggesting overall increased inflammation irrespective of viral suppression
175 (Supplemental Figure 2A). Analysis of platelets and red cell parameters showed
176 significant changes including reduced red cell counts with skewed width distribution and
177 volume, and reduced mass and hemoglobin levels (Supplemental Figure 2B). These
178 data are consistent with HIV-1 infection leading to GI pathology in Sub-Saharan African

179 cohorts (40, 41) and suggests that GI complications may arrive earlier in HIV infected
180 individuals with reduced CD4 T-cell counts and overall increased systemic inflammation.

181

182 **Irreversible depletion of duodenal CD4⁺ T-cells despite long-term ART**

183 Next, to investigate alterations in CD4⁺ and CD8⁺ T-cell homeostasis within the small and
184 large intestine during HIV-1 infection, we quantified T-cells from fresh duodenum, colon
185 and rectum biopsies in 228 study participants (Figure 3A,B). The CD4:CD8 ratios in the
186 GI tract of HIV uninfected individuals were overall lower than those found in matched
187 uninfected blood samples, and showed compartment specific differences between the
188 small (duodenum) and large (colon/rectum) intestine ($P < 0.01$), but similar ratios between
189 colon and rectum (Figure 3C) (42). HIV-1 infection reduced CD4:CD8 T-cell ratios within
190 the duodenum, colon and blood that was not restored by full ART mediated viral
191 suppression in plasma (Figure 3D). Consistent with altered CD4:CD8 ratios, the
192 frequency of CD4⁺ T-cells, as % of CD45⁺ cells, was significantly reduced in the
193 duodenum, the colon/rectum and blood of viremic participants compared to HIV
194 uninfected participants ($P < 0.001$) (Figure 3E). The CD4 depletion remained highly
195 significant in the duodenum of virally suppressed participants ($P < 0.001$), but was less
196 pronounced in the colon and no longer reached statistical significance ($P = 0.1$). The blood
197 compartment overall showed consistent CD4 depletion (see Supplemental Figure 2A),
198 although the median change was lower than that observed in the colon for viral
199 suppressed individuals (Figure 3F). Duodenal CD4 T-cell depletion, irrespective of
200 plasma viral suppression, remained significant when we controlled for age, gender and
201 ethnicity with overall less impact in the large intestine (Figure 3G). Direct comparison of

202 CD4 T-cell levels in blood and duodenum/colon combined, showed significant positive
203 correlation in HIV infected (ART^{+/-}) participants ($r=0.39$; $P=0.0015$), but not among HIV
204 negative participants ($r=0.10$; $P=0.4$) (Figure 4A). In HIV viremic individuals, we found a
205 strong negative correlation between plasma VL and blood CD4 levels ($r=0.57$; $P=0.02$),
206 but not between plasma viral loads and gut CD4 levels ($r=0.002$; $P=0.99$) (Figure 4B). To
207 explore disconnect between plasma viremia and GI CD4 T-cells further, we used
208 available treatment initiation data from the GI clinic and found no overall correlation
209 between gut CD4 T-cell levels and ART duration ($r=0.05$; $P=0.7$) (Figure 4C). Breaking
210 this down by compartment, we observed a failure to reconstitute CD4 T-cells in the
211 duodenum even after 10 years of ART (2.5 fold reduction vs HIV uninfected, $P=0.04$; Figure
212 4C). The same trend was not observed in the colon, which might indicate improved
213 reconstitution in this compartment, although fewer samples were available for this
214 analysis. Blood CD4 T-cell levels showed a trend towards CD4 restoration ($r=0.24$;
215 $P=0.13$) with only a 1.3 fold reduction ($P=0.2$) in participants receiving ART more than 10
216 years (Figure 4D), which is consistent with a 1.1 reduction in absolute CD4 T-cell counts
217 (HIV⁻ vs HIV⁺ VL<20 ART>10years, median 858 vs 795 cells/mm³, $P=0.99$). Taken
218 together, CD4 T-cell frequencies within the small and large intestine from >200 GI patients
219 show that long-term ART fails to restore CD4 T-cell levels in the small intestine despite
220 long term and fully suppressive ART.

221

222 **Compartment specific HIV-1-p24 detection and CD4 depletion in the gut mucosa of**
223 **ART treated individuals**

224 Studies of non-SubSaharan African cohorts show that CD4 T-cells within the large
225 intestinal mucosa are a major site for viral infection and replication even during plasma
226 viral suppression (4, 43). We first quantified CD4 expressing cells in colon, after exclusion
227 of GALT-like structures (Figure 5A), and duodenum tissue sections (Figure 5B). We found
228 severe depletion of duodenal CD4 cells in both virally suppressed ($P < 0.05$) and viremic
229 ($P = 0.0001$) participants, corresponding to 4- and 10-fold reduction, compared to HIV
230 uninfected GI participant controls, respectively (Figure 5C), but no significant CD4
231 depletion in the colon. This is consistent with our findings from flow cytometric analysis of
232 these compartments with severe CD4 T-cell depletion in particular observed within the
233 duodenum (see Figure 3 above).

234

235 Next, we tested HIV-1 persistence in the gut by co-staining tissue sections for HIV-p24
236 and CD4 (Figure 5D,E) and (Supplemental Figure 3A). We found HIV-p24 protein in both
237 the duodenum and colon tissues of both viral suppressed and viremic individuals that co-
238 localised with CD4 expression and most often within the lamina propria (Figure 5D,E).
239 We detected more HIV-p24 protein in the colon of both plasma viral suppressed and
240 viremic individuals compared to that of the duodenum compartment, suggesting that
241 compartment specific differences exist in the gut for HIV detection (Figure 5D). We
242 confirmed HIV detection by digital droplet PCR and found a trend towards more HIV DNA
243 in the duodenum and colon compared to blood, despite low overall CD4 frequencies in
244 the gut compared to the blood compartment (Supplemental Figure 3B). To examine the
245 relative contribution of CD4 expressing macrophages to the overall HIV-p24 expressing
246 cells, we included CD68 staining and found relatively high frequencies of CD4 and CD68

247 co-expressing cells and with more HIV-p24⁺CD68⁺ cells co-expressing CD4 in the colon
248 compartment compared to the duodenum, suggesting that macrophages also may
249 contribute to overall increased level of HIV-p24 in the colon, at least in the plasma viral
250 suppressed participants (Supplemental Figure 3C-E). Thus, we found higher HIV-p24
251 expression in the colon and lower target cell availability in the duodenum in both
252 suppressed and viremic participants, suggesting that compartment specific effects may
253 contribute to the dynamics of the HIV reservoir in the gut.

254

255 **Poor ARV mucosal drug penetration levels in the GI tract does not explain**
256 **compartment specific differences in PLWH.**

257 Next we tested if the mucosal HIV-p24 protein detection and CD4 T-cell depletion in the
258 presence of plasma viral suppression could be linked to poor ARV penetration into the GI
259 tissue. For this, we established a liquid chromatography coupled to tandem mass
260 spectrometry (LC-MS/MS) approach to accurately quantify the concentration of the first
261 line ART regimen drugs used at the time of study (44); Tenofovir (TDF), Emtricitabine
262 (FTC), Efavirenz (EFV), in fresh mucosal tissue and matched plasma (Figure 6A). Using
263 this methodology, we observed significantly lower concentrations of FTC ($P < 0.0001$) and
264 EFV ($P < 0.0001$), but not TDF, in intestinal tissue compared to paired plasma, when
265 normalizing by tissue mass (see methods) (Figure 6B). In participants with detectable
266 plasma ARV, FTC was only detected in 45% (5/11) tissue samples, compared to 86%
267 (12/14) and 93% (14/15) for TDF and EFV, respectively (Figure 6B). Importantly,
268 however, levels of all three drugs were not different between the duodenum and colon
269 compartments (Figure 6C). Moreover, in a subset of participants for whom % CD4⁺ T-cell

270 frequencies and drug levels were available, we consistently found CD4 T-cell depletion
271 in both blood and tissue (Figure 6D) despite detection of all three ARVs in both
272 compartments (Figure 6E).

273 Although, it is not clear from these data whether the reduced levels of FTC and EFV
274 directly contribute to persistent viral replication or CD4 depletion in the GI tract, they do
275 indicate that the differences in CD4 T-cell depletion and HIV-p24 detection between the
276 duodenum and colon are not a consequence of differential ARV drug penetration into
277 these two distinct gut compartments. These data suggest that penetration of some ARV
278 drugs into the gut mucosa is reduced, but that this alone does not explain the lack of CD4
279 T-cell reconstitution observed within the small intestine.

280

281 **Persistent systemic T-cell activation correlates with gut CD4 T-cell depletion only** 282 **in PLWH**

283 Several studies report that both persistent T-cell activation and systemic inflammation,
284 despite ART viral suppression, is linked to increased risk of morbidity and mortality in
285 PLWH (43, 45–49). Therefore, to test the association between GI CD4 T-cell depletion
286 and activation and systemic T-cell activation in this GI cohort, we measured HLA-DR
287 expression as a surrogate marker for T-cell activation by flow cytometry (Figure 7A). As
288 all GI participants are undergoing medical examination (see Figure 1), we compared HLA-
289 DR expression in the blood of HIV⁻ participants from the GI cohort to that of healthy non-
290 GI HIV⁻ donors from an unrelated control cohort (42) and found no significant differences
291 (Figure 7A). By contrast, HLA-DR expression was upregulated on blood CD4 and CD8 T-
292 cell subsets in both HIV viral suppressed and viremic participants, confirming HIV driven

293 T-cell activation in these individuals (Figure 7B). Next we measured HLA-DR expression
294 on GI CD4 T-cells (Suppl Figure 4) and found overall higher steady state expression
295 compared to the blood compartment ($P < 0.001$) (Figure 7C), which is expected, given the
296 persistent exposure of GI T-cells to the gut microbiome. Surprisingly, we did not see
297 significantly increased HLA-DR expression on duodenum or colon CD4 T-cells in the viral
298 suppressed individuals, and activation was actually lower in the viremic group, reaching
299 significance in the colon ($P = 0.01$) (Figure 7D).

300 Next we tested the relationship between blood T-cell activation and CD4 T-cell frequency,
301 and found a direct inverse correlation in the blood of HIV infected individuals with CD4 T-
302 cell activation ($r = -0.61$; $P = 0.001$), that was not observed in the HIV uninfected group ($r =$
303 0.01 ; $P = 0.9$), or for CD8 T-cell activation in either group (Figure 7E). Next, we correlated
304 HLA-DR expression on blood CD4 and CD8 T-cells to mucosal CD4 T-cell frequency and
305 consistently found a strong inverse correlation in HIV infected individuals with activation
306 of both CD4 T-cells ($r = -0.51$; $P = 0.03$) and CD8 T-cells in the blood ($r = -0.61$;
307 $P = 0.06$) (Figure 6F). Again, this trend was not observed in HIV uninfected participants ($r =$
308 0.05 ; $P = 0.9$ and $r = 0.2$; $P = 0.5$). Therefore, these data suggest a potential mechanistic link
309 between HIV mediated CD4 T-cell depletion in the GI tract and systemic CD4 and CD8
310 T-cell activation, which is a strong predictor of HIV disease progression independent of
311 viral load (51).

312

313 **Elevated intestinal and inflammatory markers are not specific to HIV GI pathology**
314 **and do not correlate with HIV associated depletion of gut CD4 T-cells**

315 Several previous studies have investigated the association between HIV infection, loss of
316 gut barrier integrity and immune activation using plasma markers of innate immune
317 activation, soluble CD14 (sCD14), and of elevated gut epithelial turnover, intestinal fatty
318 acid-binding protein (I-FABP) (49–54). We measured these markers in our GI study
319 participants and additional samples from an unrelated HIV uninfected healthy participants
320 not undergoing GI investigation (Figure 8). Compared to these non-GI controls, we found
321 elevated sCD14 ($P<0.0001$) and I-FABP ($P<0.0001$) levels in all GI participants, and
322 detected no significant differences between HIV infected and uninfected individuals in the
323 GI cohort (Figure 8A). To confirm this, we measured additional plasma markers of
324 inflammation (IL-6, TNF- α), macrophage activation (sCD163), a further marker of
325 intestinal barrier break down Zonulin, and the anti-inflammatory soluble IL-33 receptor
326 sST2, (Supplemental Figure 5); and found no significant differences between HIV infected
327 and uninfected individuals from the GI cohort.

328

329 Taking all individuals together, we observed a positive correlation between sCD14 and I-
330 FABP, consistent with a mechanistic link between elevated gut epithelial turn over and
331 systemic innate immune activation in this cohort of subjects with GI complications ($r=0.4$;
332 $P=0.002$) (Figure 8B). However, there was no correlation between gut CD4 T-cell
333 frequency and matched plasma I-FABP or sCD14 levels, and no correlation with
334 activation of blood CD4 T-cells (Figure 8C). When individuals were grouped by HIV
335 status, we also found no correlation between these plasma markers and GI CD4 T-cell
336 frequency or blood CD4 T-cell activation in individuals with suppressed viremia (Table 2).
337 There was a significant negative association between I-FABP and GI CD4 T-cell

338 frequency, and a non-significant positive association with blood T-cell activation in the
339 small number of viremic subjects tested.

340 Taken together, these data suggest that gut barrier break down and systemic innate
341 immune activation are likely to be general markers of GI distress, are not unique to HIV
342 infected individuals and are unlikely to occur as a direct consequence of GI CD4 T-cell
343 depletion. Importantly, they also imply that systemic T-cell activation does not occur as a
344 direct result of GI barrier breakdown, at least in virally suppressed individuals.

345 **Discussion**

346 This study presents clinical and immunological changes associated with HIV infection in
347 a large cohort of individuals attending a tertiary GI clinic for investigation. We found an
348 overall HIV prevalence of 32%, that reached >60% in younger women; with 83% of all
349 PLWH being on fully suppressive ART. Flow cytometric analysis of gut biopsies from
350 more than 200 GI participants revealed severe depletion of CD4 T-cells associated with
351 HIV, in particular within the small intestine, compared to colon and blood compartments.
352 This was most severe in the small intestine and was not reversed by full plasma viral
353 suppression and long-term ART treatment. Greater CD4 T-cell depletion within the small
354 intestine compared to the colon of PLWH was confirmed *in situ*, by fluorescence
355 microscopy. Interestingly, higher CD4 T-cell levels within the large intestine in participants
356 under full plasma viral suppression, was accompanied by more abundant HIV infected
357 CD4 T-cell and CD68 macrophages, as shown by HIV-p24 protein detection. This implies
358 that target cell availability may play an important role in governing the HIV viral reservoir
359 in the GI tract. Importantly, although the level of both EFV and FTC appeared to lower in
360 the gut than the plasma, we found no evidence of discordant ARV drug penetrations
361 between the duodenum and colon, suggesting that ARV levels does not explain these
362 differential effects. CD4 T-cell frequency in the gut of PLWH was inversely correlated with
363 activation of circulating T-cells, a biomarker which correlates strongly with HIV disease
364 progression (55). This was not observed in HIV uninfected individuals, despite elevated
365 levels of plasma markers of innate immune activation and increased gut barrier
366 permeability, such as sCD14 or I-FABP, respectively. Previous studies examining HIV
367 associated gut pathology have observed elevated levels of these and other plasma

368 markers in relation to HIV negative healthy controls (49, 52, 56, 57). Uniquely, as these
369 data are generated from individuals all attending a clinic for intestinal complications, this
370 demonstrates that such markers are elevated in individuals with GI complications in
371 general, and not just those with HIV infection. It has been argued that the observation of
372 increased gut permeability with immune activation in PLWH suggests a mechanistic link
373 between HIV infection, increased gut permeability and the persistent immune activation
374 that drives HIV progression (54, 57–59). Here we show, however, that systemic immune
375 activation of both CD4 and CD8 T-cell, at least, is not linked to gut permeability per se,
376 but rather to depletion of GI CD4 T-cells. However, we cannot rule out that sustained GI
377 tract damage and microbial translocation may also contribute to persistent systemic
378 immune activation. To our knowledge, this is the largest study of GI participants recruited
379 within a high HIV prevalence area in Sub-Saharan Africa and unequivocally identifies the
380 lack of immune reconstitution throughout the intestinal mucosa as a persistent problem
381 in PLWH despite long-term ART.

382

383 Although this study was not powered to determine differences in clinical presentation in
384 PLWH, we observed a significant increase in biliary pathologies in this group. Although
385 further work would be needed to explore this further, it is in line with previous studies (60)
386 and suggest that HIV infection precipitates cholangiopathy (61, 62) despite plasma viral
387 suppression in most individuals. In addition, we detected a low frequency (<3%) of
388 inflammatory disorders, including bowel diseases, compared to non-African cohorts
389 undergoing clinically indicated GI investigation (39). This underline the potential
390 differences between these and Sub-Saharan African GI cohorts, and demonstrates the

391 importance to study HIV gut pathology in relevant populations for better representation of
392 the HIV epidemic.

393

394 Although ART has successfully decreased HIV associated morbidity and mortality (63,
395 64), we found that virally suppressed HIV infected participants in our cohort presented a
396 median of 12 years earlier to the GI clinic than HIV uninfected attendees, predominantly
397 with non-infectious comorbidities. This early GI pathology associated with HIV could imply
398 premature immune ageing in this group, irrespective of plasma viral suppression (15) that
399 precipitates GI complications. However, we cannot rule out that HIV infected participants
400 are better linked to care than uninfected participants, which can result in earlier clinical
401 investigation.

402

403 The severe CD4 T-cell depletion observed throughout the large and small intestine is
404 consistent with early observations in Caucasian cohorts (3, 4) and further underpins the
405 lack of immune reconstitution in the gut (65). We observed a more pronounced CD4 T-
406 cell depletion in the duodenum compared to the colon. This could be due to an increased
407 abundance of lymph node structures, such as peyers patches, cryptopatches or
408 isolated lymphoid follicles (ILF) in the lamina propria of the large intestine and requires
409 further validation, ideally in colon and duodenum samples from the same participants.
410 Strikingly, we observed absolutely no correlation between the duration of ART and GI
411 CD4 T-cells, suggesting that these cells are either continually being depleted, or that the
412 CD4 niche in this tissue has been irreparably damaged and cannot be repopulated.
413 Recent data show that intestinal lamina propria CD4 T-cells are essential for regulating

414 intestinal stem cells and epithelial cell differentiation in the crypt base (66, 67). The blood
415 compartment showed a trend towards CD4 reconstitution, which may indicate late ART
416 initiation in the majority of HIV infected GI participants (68) and further emphasize that
417 intestinal CD4 levels, in particular within the small intestine, do not recover despite long-
418 term ART (65) in contrast to treatment initiation during early acute HIV infection (69).

419

420 The observation of more pronounced CD4 T-cell depletion in the duodenum compared to
421 the colon was shown by both flow cytometry and histology. This is a potentially important
422 observation, as, for practical reasons, many studies on the impact of HIV on the GI tract
423 have limited their study to the colon. As CD4 loss was less pronounced in this
424 compartment and appears to be somewhat better restored by ART, limiting observations
425 to the colon may miss key aspects of persistent GI pathology in PLWH. It is important to
426 note, however, that we were unable to obtain paired duodenum and colon samples from
427 the sample individuals. It is therefore possible that compartment specific difference may
428 in part be explained by differences in the HIV patients presenting to the clinic for either
429 upper or lower GI tract endoscopy.

430

431 Although the life expectancy of people living with HIV has been prolonged by the
432 improvement in ART over the last few years (70), the gut has been implicated as a
433 potential reservoir for ongoing HIV-1 replication in the setting of ART suppressed viremia
434 (26). We tested if discordant ARV levels between duodenum and colon mucosal tissue,
435 with plasma matched controls, could explain the difference in CD4 T-cell depletion. We
436 found no evidence of compartment specific ARV levels, but that duodenal CD4 T-cell

437 depletion occurs in confirmed ARV detected mucosal tissue, although based on a few
438 individuals available. Thus, reduced ARV levels in mucosal tissue compared to plasma
439 levels (14), does not necessarily explain the lack of CD4 T-cell reconstitution in this
440 cohort. One caveat remains that comparing ARV levels in plasma to homogenized gut
441 tissue cells may be suboptimal instead of using tissue matched PBMCs and normalized
442 by cell numbers. However, previous studies have showed a correlation between plasma
443 and PBMC ARV levels (71–76). Whether ongoing viral replication could be linked to
444 reduction or absence of ARVs in gut mucosal tissue (33) remains to be tested in this
445 cohort. Finally, HIV-p24 detection in both the small and large intestine, despite full viral
446 suppression, may also be linked to ARV drug pharmacokinetics (77, 78). In plasma
447 samples, we found evidence of treatment failure in which 3 ARVs were detected in 40%
448 of viremic patients and highlights the vulnerability of this population where ART
449 adherence over long period of time is challenging (79) and also could accelerate GI
450 complications in this region (80).

451

452 Our GI cohort underpins the central role of the intestine in HIV-1 pathology and
453 demonstrates poor restoration of gut mucosal immunity, in particular within the small
454 intestine, by otherwise effective ART. This ongoing GI study cohort is a unique opportunity
455 to further reveal the unknown mechanisms of GI impairment that are central to both HIV
456 pathogenesis and its associated comorbidities in the growing population of individuals
457 receiving ART in Sub-Saharan Africa, a distinct and critical population that is under
458 studied. Future elected mucosal tissue sample collection from matched large and small
459 intestinal tissue sites, combined with better characterized participants with available long-

460 term clinical information from this region, will be a key resource for more in depth
461 understanding of the mechanisms underlying HIV pathology in the GI tract.

462 **Methods**

463 **Study participants and sample collection**

464 Patients presenting to the gastrointestinal (GI) surgical unit of Inkosi Albert Luthuli Central
465 Hospital (IALCH) were recruited into this study after obtaining written informed consent.
466 IALCH is an 846-bed central and tertiary hospital in the eThekweni district of KwaZulu-
467 Natal (KZN) Province in South Africa. It provides regional and tertiary services to KZN as
468 well as for referrals from the neighbouring Eastern Cape Province. Duodenum, colon and
469 rectum pinch biopsies with participants-matched blood samples were obtained during
470 endoscopy, endoscopic retrograde cholangiopancreatography (ERCP) and colonoscopy
471 procedures. Blood samples were also obtained from a healthy HIV uninfected volunteer
472 study cohort (42). Clinical information, including HIV status and demographic details of
473 these participants, was collected using a structured questionnaire. HIV status was
474 confirmed using the COBAS® TaqMan® HIV-1 Test (Roche). Full blood count and viral
475 load were done using the Sysmex XE-5000™ Automated Hematology Analyzer and
476 NUCLISENS EASYQ® HIV-1 (BioMerieux), respectively.

477

478 **Mononuclear cells isolated from blood and gut**

479 Blood was collected in BD vacutainers with sodium heparin (Becton Dickinson).
480 Peripheral blood mononuclear cells (PBMCs) were isolated using the Ficoll-Histopaque
481 1077 (Sigma-Aldrich) density gradient centrifugation. Gut biopsies (2 to 4 pinches) were
482 removed by the operating GI surgeon and transported to the laboratory in cold
483 Phosphate-Buffered Saline (PBS) (pH 7.2). The PBS was decanted from the tubes
484 containing the gut biopsies, which are about 5-8 mm in size, and they were incubated in

485 epithelial strip buffer (PBS; 0.5M EDTA; 1M DTT; FBS and Penicillin/Streptomycin) in a
486 37°C water bath for 10 minutes, with occasional agitation. Thereafter, the epithelial strip
487 buffer was removed, and the tissues were digested in a buffer containing Collagenase-D
488 (0.5 mg/ml; Roche) and DNase-I (20 µg/ml; Sigma-Aldrich) for 30 minutes in a 37°C water
489 bath with occasional agitation. Digested tissue was passed through a 70µM cell strainer
490 to isolate the cells, and these cells were washed with PBS.

491

492 **Flow cytometry**

493 PBMC and gut mononuclear cells were stained with antibody mixtures for a minimum of
494 20 minutes at room temperature. Cells were washed twice with PBS and acquired
495 immediately or fixed in 2% paraformaldehyde for later acquisition within 24 hours. The
496 BD FACSAria Fusion flow cytometer was used for acquisition of sample data. A minimum
497 of 100,000 total events was collected for each sample, and data analysis was done using
498 FlowJo software (version 9.7.2 or higher, TreeStar). The antibody mixture consists of
499 Live/Dead Fixable Near-IR Cell Marker (Invitrogen, Thermo Fisher Scientific) and an
500 optimized antibody cocktail for the different phenotypic staining that included the following
501 antibodies anti-CD45 (BD Biosciences 560777) anti-CD3 (BioLegend 317330) anti-CD4
502 (BD Bioscience 564651), anti-CD8 (BD Bioscience 563795), anti-HLA-DR (Biolegend
503 307606).

504

505 **Measurement of drug levels in gut and plasma**

506 The concentration of ARV drugs was simultaneously quantified in plasma and gut tissue
507 samples using liquid chromatography coupled to tandem mass spectrometry (LC-

508 MS/MS). The tissue samples were homogenized using a TH Omni international
509 homogenizer/soft tissue probe combination (Omni-International, Georgia, USA) with a
510 short pulse (15 sec) in 500 µl of cold acetonitrile: water (1:1, v/v) solution, then processed
511 for analysis. The drug concentration data were normalized to the average mass of the
512 tissue sample set and used a tissue density of 1.06 grams/mL to relate tissue weight to
513 tissue volume as previously described (75). The ARV drug analytes were extracted from
514 calibration standards (STDs), quality control (QCs) samples, and study samples using
515 protein precipitation followed by LC-MS/MS analysis. Human plasma was used as a
516 surrogate biological matrix for the preparation of quantitative STDs and QCs, due to the
517 limited availability of control (healthy donor/drug-free) gut tissue sample. Matrix effect
518 studies to compare drug quantitation between spiked plasma and homogenized gut tissue
519 samples were performed and showed equivalency with less than 15% variability in
520 analyte peak areas.

521 The extracted analytes were chromatographically separated on an Agilent Zorbax Eclipse
522 Plus C18 (2.1 x 50mm, 3.5 µm) HPLC column, at 40°C, using gradient elution with a
523 combination of mobile phase A, which consisted of water with 0.1% formic acid and B,
524 acetonitrile with 0.1% formic acid. A sample volume of 20 µl was injected, the flow rate
525 was set to 0.2 ml/minute and the total run time was 20 minutes. All ARV drug analytes
526 were analysed in the positive ionization mode, using 6-Aminonicotinic acid (6-ANA) as an
527 internal standard except for efavirenz which was analysed in the negative ionization mode
528 using deuterated efavirenz (EFV-d5) as an internal standard. Data were acquired using
529 selected reaction monitoring (SRM) and processed using AB Sciex Analyst Software,
530 version 1.6.2, on an AB Sciex Triple Quad™ 5500 mass spectrometer.

531

532 **Histology and fluorescent immunohistochemistry**

533 Multiplex fluorescent immunohistochemistry staining was performed using the Opal™ 4-
534 Color Manual IHC Kit 50 Slides (PerkinElmer, USA) as directed by the manufacturers.
535 Duodenum and colon tissue samples fixed in 4% formalin between 2 days and a month
536 were paraffin-embedded. 4µm sections were cut on glass slides and the slides were
537 baked at 60°C overnight. Then, the combined process of deparaffinization, rehydration
538 and antigen retrieval of the tissue sections was done using the PT-Link Pre-Treatment
539 instrument (Dako) and 1x Envision Target Retrieval Solution, High PH (Dako). Then,
540 slides were incubated for 1 minute in distilled water and equilibrated in 1x EnVision FLEX
541 Wash Buffer (Dako) for 5 minutes at room temperature. Then, the slides were incubated
542 in Peroxidase blocking solution (PerkinElmer) for 10 minutes and washed in 1x wash
543 buffer (Dako) immediately at room temperature. The slides were then incubated in Bloxall
544 blocking solution (PerkinElmer) for 10 minutes, and then in primary antibody-1 for 30
545 minutes at room temperature. Slides were then washed for 5 minutes in 1x wash buffer
546 and incubated in Secondary Opal Polymer Horseradish Peroxidase (HRP) Mouse and
547 Rabbit (PerkinElmer) for 30 minutes. The Opal Polymer HRP is recommended for human
548 tissues with a mouse or rabbit primary antibodies. Then, the slides were washed twice in
549 1x wash buffer, drained and the sections were incubated in Opal Fluorophore
550 (PerkinElmer) diluted 1:100 in amplification diluent (PerkinElmer) working solution for
551 signal amplification at room temperature for 10 minutes. The slides were then washed for
552 5 minutes in 1x wash buffer at room temperature. Afterwards, antibody stripping via
553 microwave treatment was done by placing the slides in a slide jar with pre-warmed buffer

554 1x AR6 (PerkinElmer). The jar was loosely covered and placed in a microwave for 2
555 minutes at 100% power, 10 minutes at 50% power and 5 minutes at 20% power. Slides
556 were allowed to cool in the dark by placing the slide jar on ice for 20 minutes and the
557 slides were rinsed in distilled water, followed by incubation in the 1x wash buffer for 5
558 minutes to equilibrate. The microwave step strips the primary-secondary-HRP complex
559 and allows the introduction of the next primary antibody. For the detection of the next
560 target (primary antibody-2), the protocol was restarted at the blocking step using Bloxall
561 blocking solution (PerkinElmer) for 10 minutes. After the third target was detected
562 (primary antibody-3), a working solution of DAPI (PerkinElmer) was applied to the
563 sections as the nuclear counterstain for 5 minutes. The slides were washed in 1x wash
564 buffer for 5 minutes, then in distilled water for 5 minutes and drained. Then, the sections
565 were coverslip with Fluorescence Mounting Medium (catalog number S302380-2; Agilent
566 Technologies) and the edges of the coverslip were sealed with nail polish. Slides were
567 stored in a humidity chamber at 2°C - 8°C until images are acquired.

568

569 The unlabelled primary antibodies used in this study were anti-HIV p24 (clone: Kal-1,
570 Dako) used as the first antibody in the staining cycle, diluted 1:10 in antibody diluent
571 (PerkinElmer), then anti-CD4 (clone: 4B12, Dako) which was the second antibody in the
572 staining cycle and it was pre-mixed, followed by anti-CD68 (clone: KP1, Dako) as the third
573 antibody in the cycle, diluted 1:200 in antibody diluent (PerkinElmer). The fluorophores
574 were FITC (product number FP1487001KT; PerkinElmer) for anti-HIV p24 (clone: Kal-1,
575 Dako), Texas-Red (product number FP1488001KT; PerkinElmer) for anti-CD4 (clone:
576 4B12, Dako) and Cy5 (product number FP1497001KT; PerkinElmer) for anti-CD68

577 (clone: KP1, Dako) signal generation. These fluorophores were diluted 1:100 in
578 amplification diluent. Images of the tissue sections were acquired using the TissueFAXS
579 software (TissueGnostics) connected with a Zeiss Axio Observer Z1 inverted microscope
580 (Olympus). The quantitative analysis of the cells of the different phenotypes within the
581 images was done using the TissueQuest quantitation software (TissueGnostics). Isotype
582 and non-antibody controls were included in the experiment to rule out non-specific
583 staining and autofluorescence.

584

585 **Measurement of blood and gut HIV-1 DNA using digital droplet PCR (ddPCR)**

586 Total PBMC, and duodenum and colon tissue mononuclear cells from study participant
587 were subjected to DNA extraction using DNeasy Blood & Tissue Kits (QIAGEN). Total
588 HIV-1 DNA and host cell concentrations in the DNA extracts were estimated using BIO-
589 RAD ddPCR, using primers and probes covering HIV-1 5' LTR-gag HXB2 coordinates
590 684–810 (81) (forward primer 5'-TCTCGACGCAGGACTCG-3', reverse primer 5'-
591 TACTGACGCTCTCGCACC-3' probe/56-
592 FAM/CTCTCTCCT/ZEN/TCTAGCCTC/31ABkFQ/, and human RPP30 gene (82)
593 forward primer 5'-GATTTGGACCTGCGAGCG-3', reverse primer 5'-
594 GCGGCTGTCTCCACAAGT-3', probe/56-
595 FAM/CTGACCTGA/ZEN/AGGCTCT/31ABkFQ/). ddPCR was performed using the
596 following thermocycler program: 95 °C for 10 minutes, 45 cycles of 94 °C for 30 seconds
597 and 60 °C for 1 minutes, 72 °C for 1 min. The droplets were subsequently read by the

598 BIO-RAD QX100 droplet reader and data were analyzed using QuantaSoft software
599 (BIO-RAD).

600

601 **Markers of microbial translocation and inflammation**

602 The concentration of some of the plasma markers was measured by ELISA using Human
603 soluble CD14 DuoSet ELISA (R&D Systems, catalogue number: DY383), Human
604 FABP2/I-FABP DuoSet ELISA (R&D Systems, catalogue number: DY3078) and Human
605 Zonulin/Haptoglobin ELISA kit (ARP American Research Products, catalogue number:
606 E01Z0004) per manufacturer's instructions. Also, Milliplex multiplex assays using
607 Luminex was used to determine the concentration of soluble ST2 and CD163 using the
608 Human Cardiovascular Disease Magnetic bead panel 5 (Milliplex, catalogue number:
609 HCVD5MAG-67K) per manufacturer's instructions. The concentration of IL-6 and TNF- α
610 was determined using the Human TH17 magnetic Bead Panel (Milliplex, catalogue
611 number: HTH17MAG-14K) per manufacturer's instructions.

612

613 **Statistics**

614 Descriptive and categorical data were represented as medians with interquartile ranges,
615 minimum and maximum values and differences were analyzed using the Mann-Whitney
616 U-test. The Kruskal-Wallis test was performed for multiple comparisons between HIV
617 uninfected and each of the HIV-1 infected groups. Bivariate correlations were determined
618 by the Spearman rank correlation test. A multivariate logistic regression model was used
619 to estimate the association between HIV-1 and GI comorbidities at a 95% confidence

620 interval (CI). Statistical analyses were performed using GraphPad Prism (version 8) and
621 STATA (version 15) software, and p-values <0.05 were considered statistically significant.

622 **Study Approval**

623 This study was approved by the Biomedical Research Ethics Committee (BREC) of the
624 University of KwaZulu-Natal (REF: BE 021/13) following its ethical standards and with the
625 1964 Helsinki declaration and its later amendments.

626

627 **Author contributions**

628 OA and AS performed experiments. FN and MN recruited study participants. FN and NM
629 consented participants in the GI ward. DR, FK coordinated human specimens. OA, AS,
630 AN, RF, YZ, KG, NH and KR contributed to experimental work. FM, VTM, FA contributed
631 surgical human tissues samples. NM, FN, FK and TN contributed sample collection. HK,
632 OA, AL prepared the manuscript. TN, ZP, JA, AL provided intellectual input. HK
633 conceptualized and provided funding.

634

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640 the African Academy of Sciences (AAS)'s Alliance for Accelerating Excellence in Science
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642 Planning and Coordinating Agency (NEPAD Agency) with funding from the Wellcome
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644 publication are those of the author(s) and not necessarily those of AAS, NEPAD Agency,
645 Wellcome Trust or the UK government. HK and AS was supported by SANTHE.

Table 1. HIV diagnosis and treatment duration in GI cohort (n=529)

	HIV ⁺ all, n=170/529 (32%)	HIV ⁺ ART ⁺ VL<20, n=101/120 (84%)	HIV ⁺ ART ⁻ , n=23/160 (14%)
Time HIV diagnosed, years (IQR)	6 (1.5-10.1)	6.3 (1.6-10.1)	0.3 (0.2-1.1)
Time on ART, years (IQR)	5.5 (1.7-8.0)	5.0 (1.6-7.7)	NA
Time ART naïve, years (IQR)	NA	2.1 (0.4-7.7)	NA
CD4 count cells/mm ² (IQR)	629 (380-810)	647 (465-813)	297 (155-363)
HIV RNA copies/ml plasma (IQR)	NA	NA	29,350 (205-147,500)

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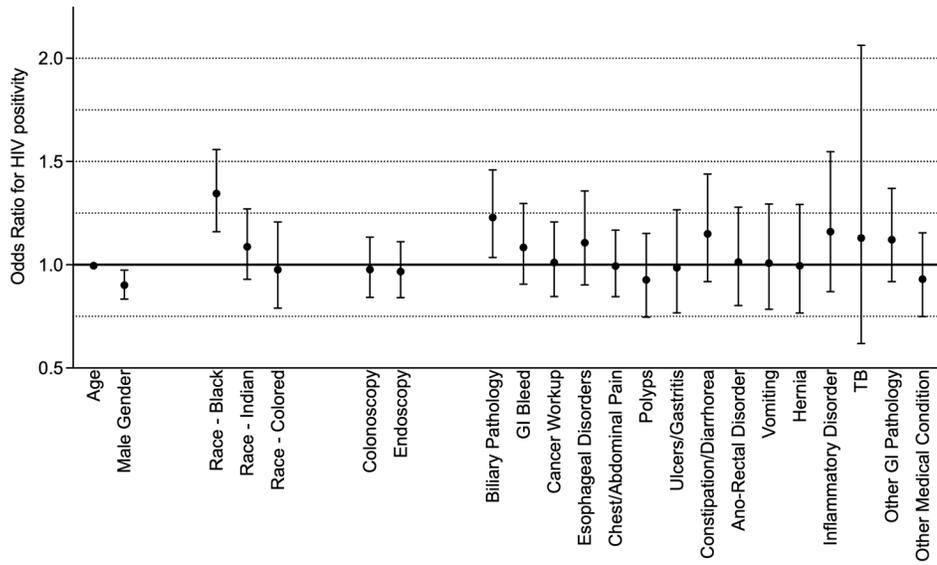
Table 2. Plasma marker correlations CD4 and HLA-DR

	*All	*HIV ⁻	*HIV ⁺ VL<20	*HIV ⁺ Viremic
sCD14 vs %CD4 in gut	0.9, -0.01 (40)	0.5, 0.16 (18)	0.2, 0.37 (15)	0.9, -0.04 (7)
IFABP vs %CD4 in gut	0.6, -0.08 (42)	0.14, 0.35 (19)	0.7, -0.09 (16)	0.006, -0.94 (6)
sCD14 vs %CD4 HLA-DR blood	0.6, 0.08 (51)	0.6, 0.11 (27)	0.6, -0.14 (19)	0.8, -0.14 (5)
I-FABP vs %CD4 HLA-DR blood	0.3, 0.13 (54)	0.9, -0.001 (30)	0.4, 0.19 (19)	0.1, 0.76 (5)

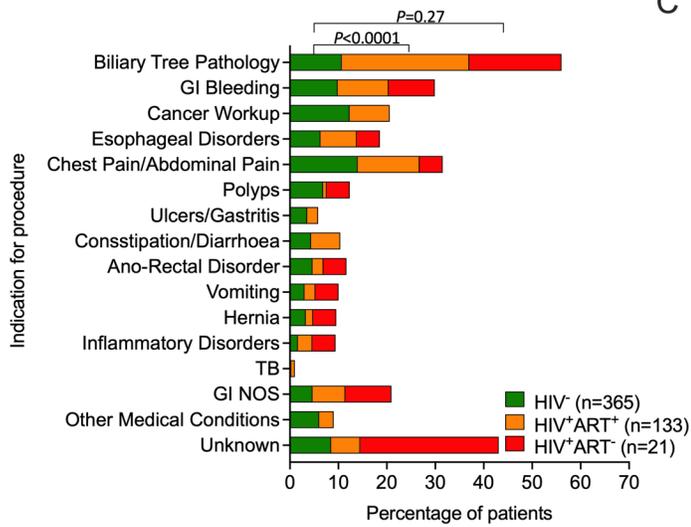
*Pearson correlation P-value, r –Pearson correlation coefficients, n – biological replicates

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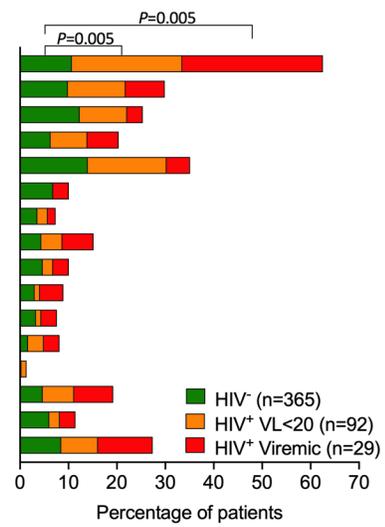
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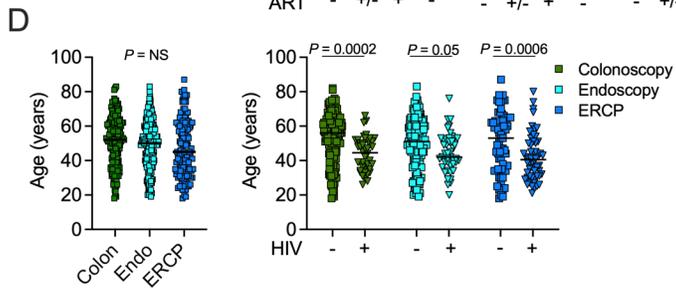
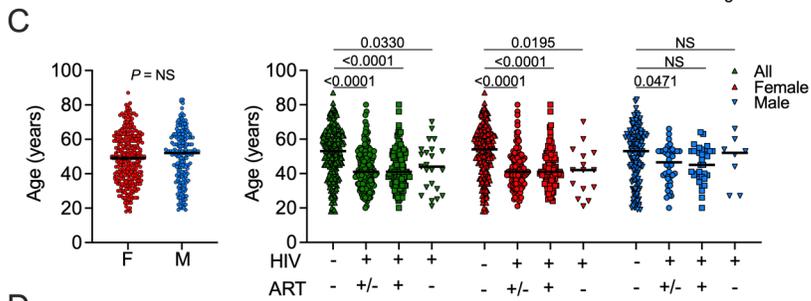
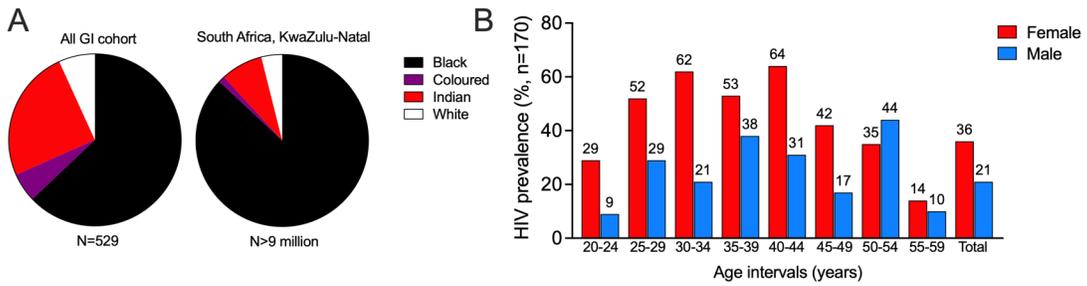


653 **Figure legends**

654 **Figure 1. Surgical indications of participants in GI study cohort. A.** Odds ratios for
655 HIV positivity associated with demographic variables and procedural indications.
656 **B.** Surgical indications stratified by HIV and ART status (n=519). **C.** Surgical indications
657 stratified by HIV and viral load status (n=486).

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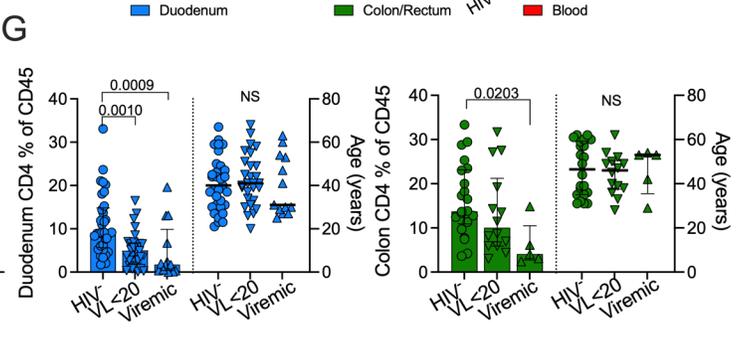
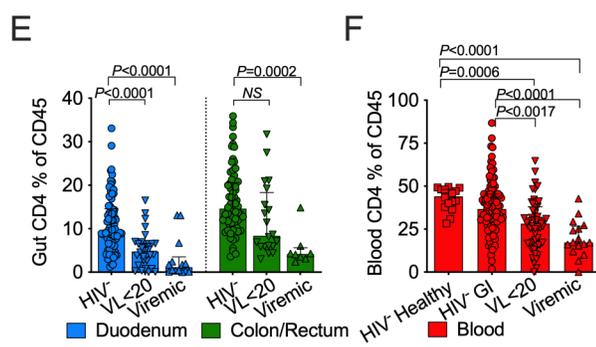
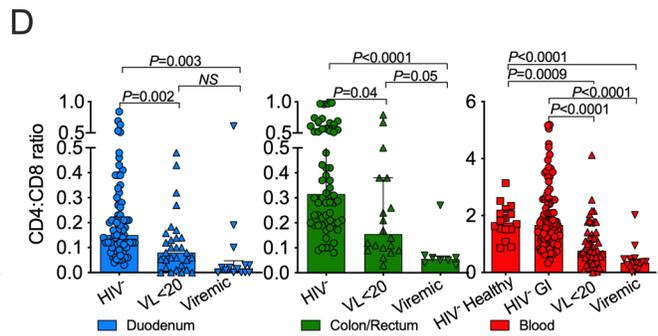
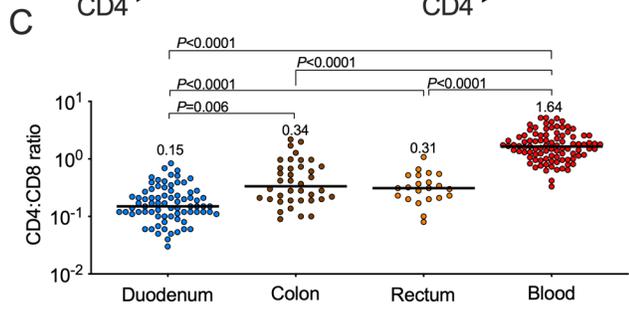
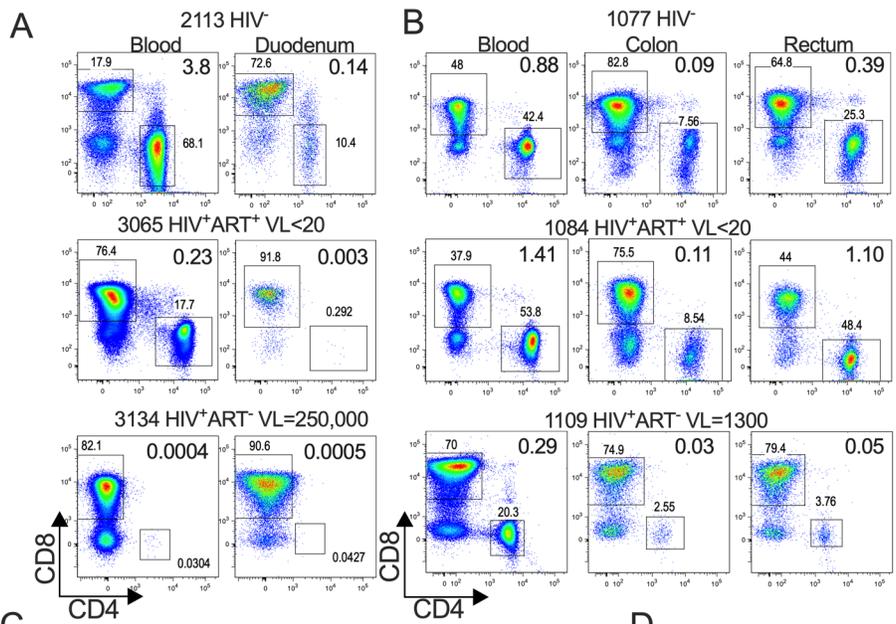
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660 **Figure 2. Characteristics of study participants. A.** Study participants (n=529) stratified
661 by ethnicity compared to the general population of the KwaZulu-Natal province. **B.** HIV
662 prevalence stratified by age group and gender, female (red) and male (blue) (n=170). **C.**
663 Age of presentation to the GI clinic stratified by gender, Female (red) and Male (blue);
664 HIV prevalence and ART status stratified by age and gender. All participants (green),
665 Female (red), Male (blue), p-values by Mann-Whitney U test and Kruskal-Wallis test. **D.**
666 Participants age of presentation for GI surgical procedures irrespective of HIV-1 status;
667 Participants' age of presentation for GI surgical procedure stratified by HIV-1 status. p-
668 values by Kruskal-Wallis test.

669

670



671 **Figure 3. Irreversible depletion of duodenal mucosal CD4⁺ T-cells despite long-term**
672 **ART. A.** Gating strategy to identify CD4⁺ and CD8⁺ T-cells in patient-matched blood and
673 duodenum from HIV uninfected and infected participants. **B.** Gating strategy to identify
674 CD4⁺ and CD8⁺ T-cells in participants-matched blood, colon and rectum from HIV
675 uninfected and infected patients. **C.** CD4: CD8 ratios of duodenum, colon, rectum and
676 blood from HIV uninfected participants. p-values by Kruskal-Wallis test. **D.** CD4: CD8
677 ratios of duodenum (blue), colon/rectum (green) and blood (red) from HIV uninfected,
678 HIV-1 infected suppressed and viremic participants. p-values by Kruskal-Wallis test. **E.**
679 CD4% of CD45 cells from the duodenum (blue) and colon/rectum (green) of HIV-1
680 uninfected, infected virally suppressed and viremic participants. **F.** CD4% of CD45 cells
681 from blood (red) of Healthy HIV uninfected non-GI study participants, HIV uninfected GI
682 study participants, HIV-1 infected suppressed and viremic from our GI cohort. p-values
683 by Kruskal-Wallis test. **G.** Duodenum (blue) and Colon (green) CD4% of CD45 cells
684 stratified by age from HIV uninfected GI study participants, HIV-1 infected suppressed
685 and viremic from our GI cohort. p-values by Kruskal-Wallis test.

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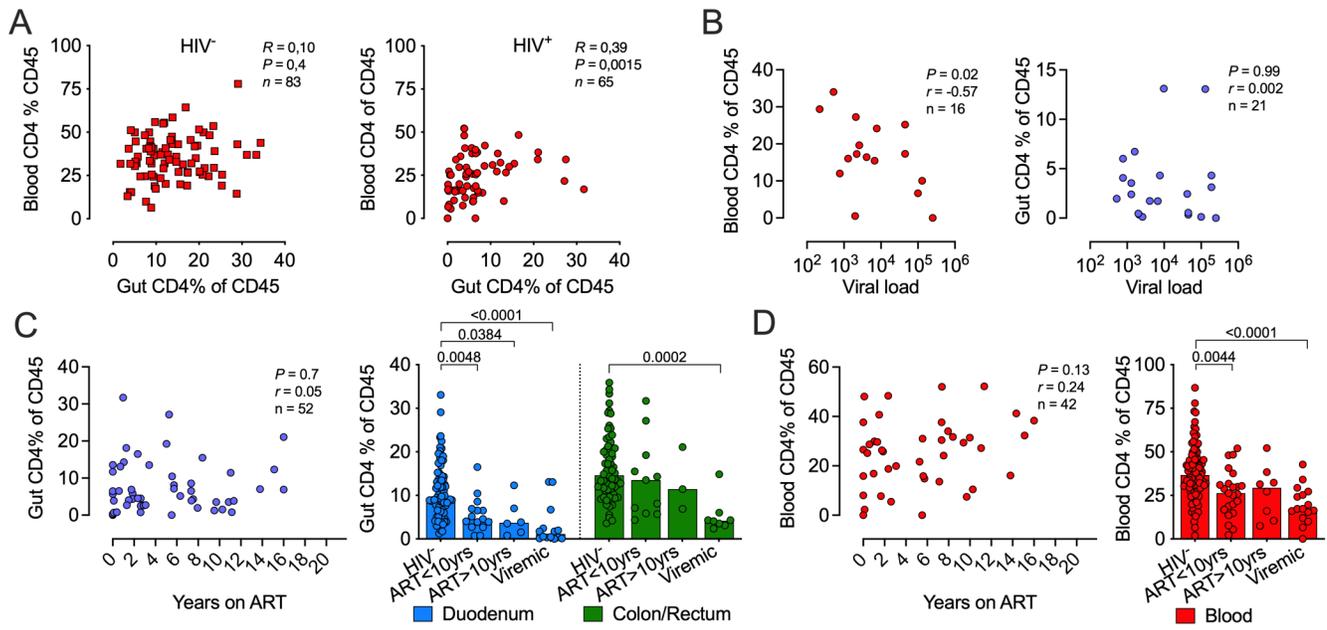
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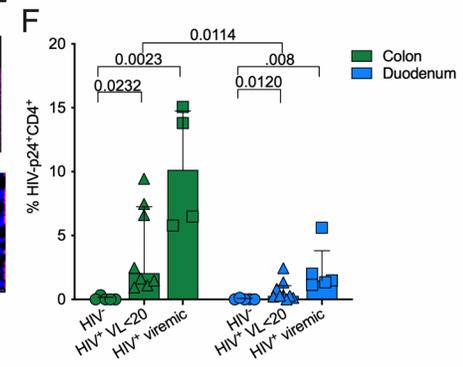
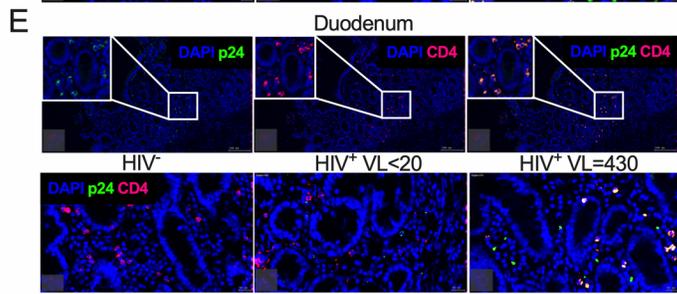
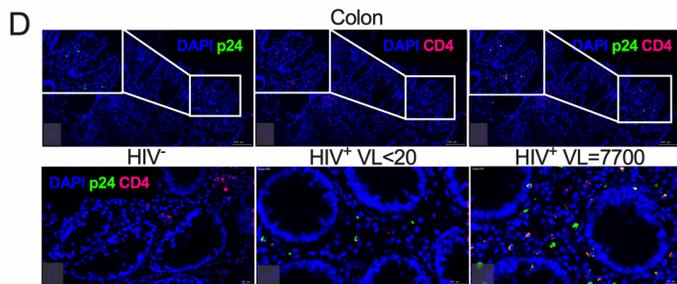
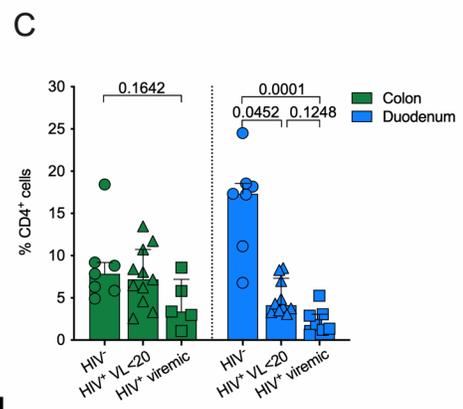
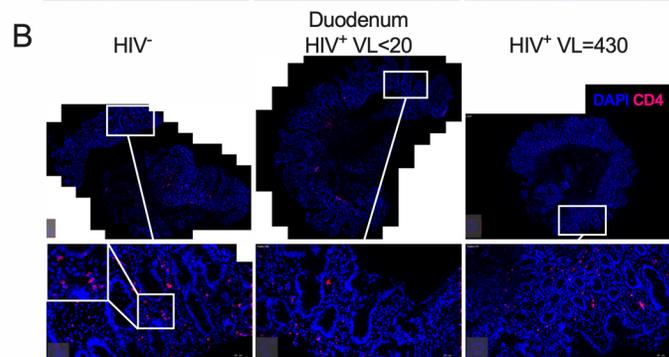
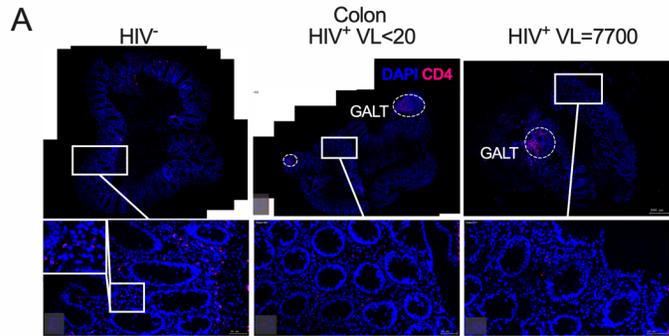
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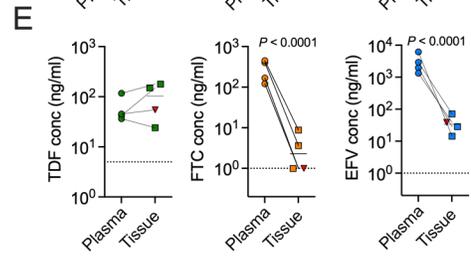
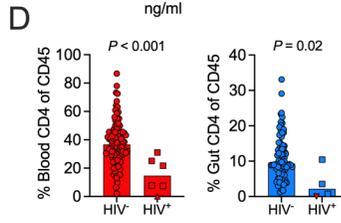
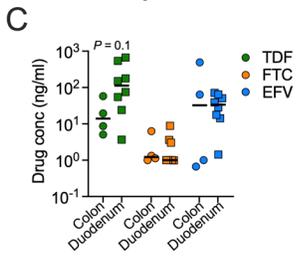
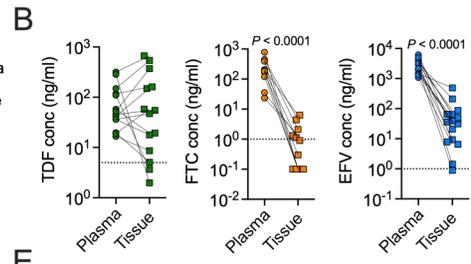
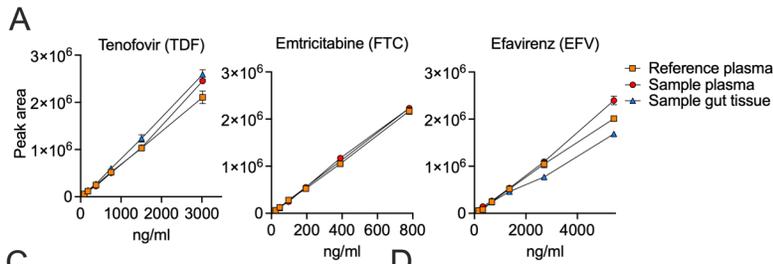
693 **Figure 4. Depleted duodenal CD4 T-cells fails to reconstitute even after 10 years of**
694 **ART. A.** Correlation of CD4% of CD45 cells of pooled gut tissues with that of blood among
695 HIV uninfected and infected participants. **B.** Correlation of viral load with CD4% of CD45
696 cells of blood (red) and gut tissues (blue). **C.** Correlation of participants' duration on ART
697 (years) with CD4% of CD45 cells of gut tissues (blue); CD4% of CD45 cells of duodenum
698 (blue) and colon/rectum (green) tissues stratified by HIV and ART status. **D.** Correlation
699 of participants' duration on ART (years) with CD4% of CD45 cells of blood (red),
700 correlations by spearman test; CD4% of CD45 cells of blood stratified by HIV and ART
701 status. Correlations by spearman test and P-values by Kruskal Wallis Test for multiple
702 comparisons.

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704



705 **Figure 5. Compartment specific HIV-1-p24 detection and CD4 depletion in the gut**
706 **mucosa of ART treated individuals. A.** Representative fluorescent
707 immunohistochemistry image of CD4 T-cells (red) 3proportion of total cells stained with
708 DAPI (blue) in colon tissue sections from HIV uninfected, HIV infected plasma virally
709 suppressed and viremic participants. Bars, 100 μ m for full image, 50 μ m and 20 μ m for
710 large and small inserted images respectively. **B.** Representative fluorescent
711 immunohistochemistry image of CD4 T-cells (red) proportion of total cells stained with
712 DAPI (blue) in duodenum tissue sections from HIV uninfected, HIV infected plasma virally
713 suppressed and viremic participants. Bars, 100 μ m for full image, 50 μ m and 20 μ m for
714 large and small inserted images respectively **C.** Quantification of CD4 T-cells in colon
715 (green) and duodenum (blue). Each dot is representative of a gut tissue section per
716 participants. p-values by Kruskal Wallis test corrected for multiple comparisons. **D.**
717 Representative fluorescent immunohistochemistry image of colon tissues showing HIV-
718 p24 (green), CD4 T-cells (red) and DAPI (blue) from HIV uninfected, virally suppressed
719 and viremic participants. Bars, 100 μ m for full image, 50 μ m and 20 μ m for large and
720 small inserted images respectively. **E.** Representative fluorescent immunohistochemistry
721 image of duodenum tissues showing HIV-p24 (green), CD4 T-cells (red) and DAPI (blue)
722 from HIV uninfected, virally suppressed and viremic participants. Bars, 100 μ m for full
723 image, 50 μ m and 20 μ m for large and small inserted images respectively. **F.**
724 Quantification of HIV-p24 and CD4 T-cells proportion of total cells in colon (green) and
725 duodenum (blue) stained with DAPI from HIV uninfected, virally suppressed and viremic
726 participants. P-values by Mann-Whitney U test.
727



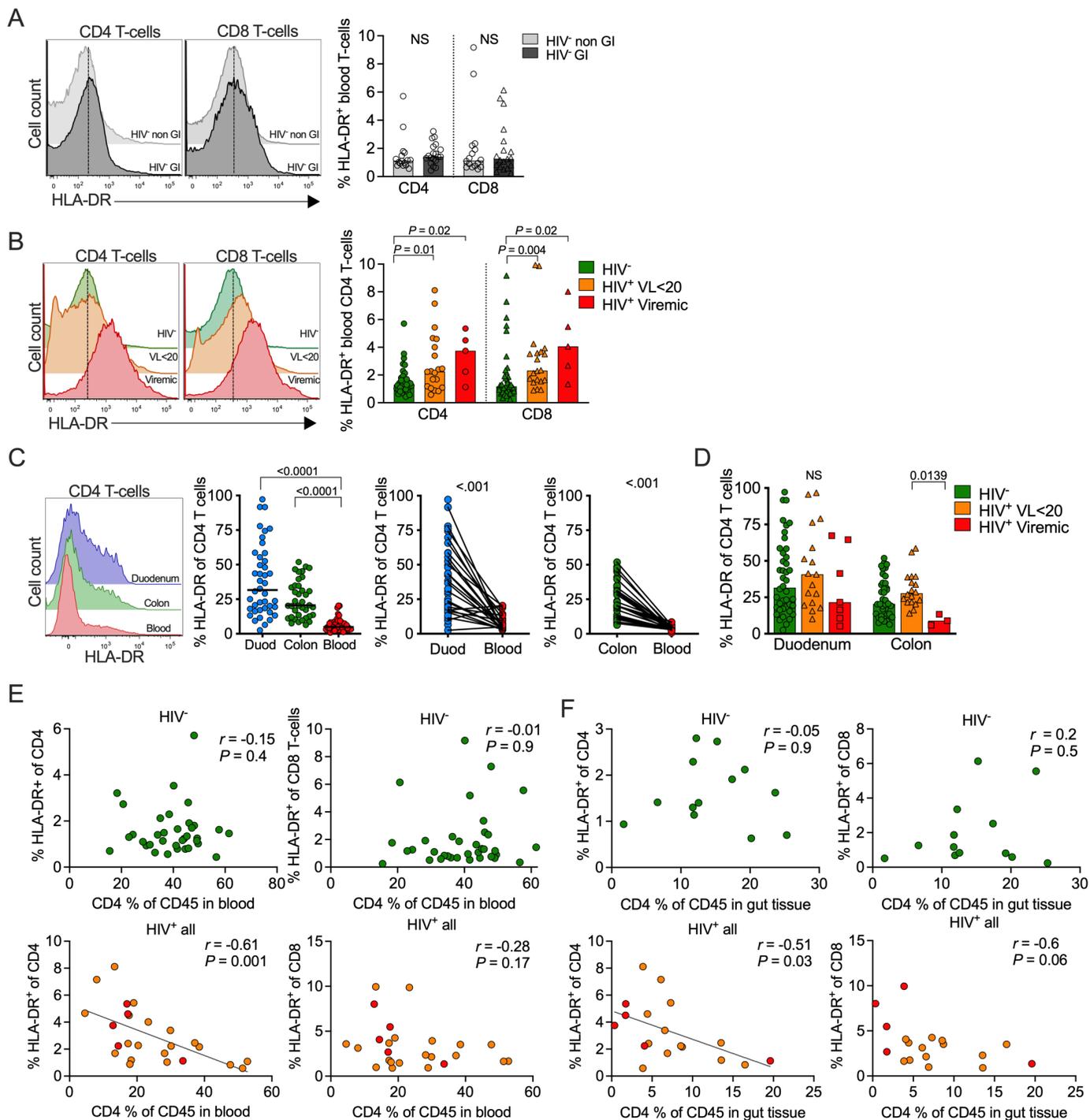
728

729 **Figure 6. ARV drug concentration within plasma and intestinal mucosal tissue sites**

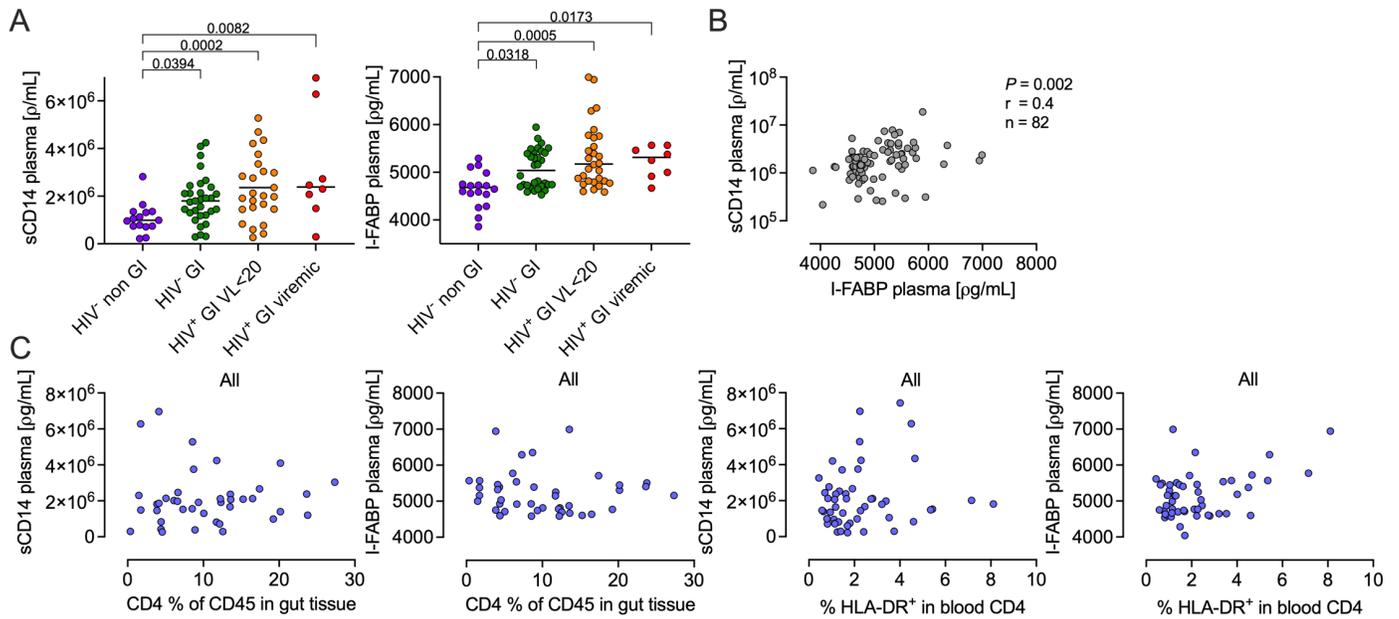
730 **A.** Quantitation of ARV drugs (TDF, FTC and EFV) in gut tissue and plasma using liquid
731 chromatography coupled to tandem mass spectrometry (LC-MS/MS), showing biological
732 matrix equivalency between spiked plasma and gut tissue samples. **B.** Comparison of
733 TDF, FTC and EFV concentrations between gut tissue (normalized by wet tissue mass)
734 and patient matched plasma with dotted line showing limit of detection. **C.** ARV
735 quantification stratified by colon and duodenum tissue. **D.** CD4 % of CD45 lymphocytes
736 in blood (red) and tissue (blue) of participants whose ARV concentration was tested,
737 stratified by HIV status. **E.** TDF, FTC and EFV concentrations between gut tissue
738 (normalized by wet tissue mass) and patient matched plasma in subset of participants
739 with available % CD4⁺ T-cell frequencies of CD45⁺ cells, with dotted line showing limit of
740 detection. TDF=Tenofovir, FTC=Emtricitabine, EFV=Efavirenz.

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743 **Figure 7. Persistent systemic T-cell activation correlates with gut CD4 T-cell**
744 **depletion. A.** Gating strategy to identify CD4⁺ and CD8⁺ T-cells activation marker, HLA-
745 DR in blood, comparing HIV uninfected non-GI and GI participants. **B.** T-cell activation
746 (CD4⁺HLA-DR⁺ and CD8⁺HLA-DR⁺) in blood from GI cohort participants stratified by viral
747 load. P-values by Kruskal-Wallis test. **C.** CD4 T-cell activation in duodenum (blue), colon
748 (green) and blood (red) from HIV uninfected GI participants. P-values by Kruskal-Wallis
749 test. **D.** CD4 T-cell activation in HIV uninfected GI participants (green) compared with
750 HIV-1 infected virally suppressed (orange) and viremic (red) participants from duodenum
751 and colon. P-values by Kruskal-Wallis test. **E.** Correlations of CD4% of CD45 cells in
752 blood with % of CD4⁺ and CD8⁺ T-cell activation among HIV uninfected and HIV infected
753 participants. **F.** Correlations of CD4% of CD45 cells in gut tissues with % of CD4⁺ and
754 CD8⁺ T-cell activation among HIV uninfected and HIV infected participants. P and r values
755 by pearson correlations.
756
757



758 **Figure 8. Elevated intestinal and inflammatory markers are associated with GI**
759 **pathology but do not correlate with HIV associated depletion of gut CD4 T-cells. A.**
760 Increased sCD14 and I-FABP among HIV-1 infected virally suppressed (orange) and
761 viremic participants (red) compared with healthy HIV uninfected participants (purple). P-
762 values by Kruskal-Wallis test. **B.** sCD14 correlates positively with I-FABP. P and r values
763 by pearson correlations. **C.** Lack of correlation between CD4% of CD45 cells in gut tissues
764 with plasma sCD14 and I-FABP; lack of correlation between % of activated CD4 in the
765 blood with plasma sCD14 and I-FABP.

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