

1 **TITLE:** T cell responses to nonstructural protein 3 distinguish infections by Dengue and
2 Zika viruses

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47 **ABSTRACT**

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49 The 2015-16 Zika virus (ZIKV) epidemic in the Americas and the Caribbean
50 demonstrates that clinical assays to detect, distinguish, and characterize immune
51 responses to flaviviral infections are needed. ZIKV and dengue virus (DENV) are
52 mosquito-transmitted flaviviruses sharing overlapping geographical distribution and
53 have significant sequence similarity that can increase the potential for antibody and T
54 cell cross-reaction. Using nonstructural protein 1-based enzyme-linked immunosorbent
55 assays (ELISAs), we determine the serostatus of individuals living in a DENV- and
56 ZIKV-endemic region in Brazil, identifying individuals with primary DENV (pDENV) and
57 ZIKV (pZIKV), ZIKV with primary DENV (ZIKVwpDENV), and secondary DENV
58 (sDENV) infections; pDENV and pZIKV were further confirmed by neutralization tests.
59 Development of an enzyme-linked immunospot (ELISPOT) assay for DENV and ZIKV
60 structural and nonstructural (NS) protein antigens enables us to distinguish infections by
61 these viruses based on T cells and to characterize those responses. We find that IFN- γ
62 and TNF- α T cell responses to NS3 differentiates DENV and ZIKV infections with 94%
63 sensitivity and 92% specificity. In general, we also show that pDENV and sDENV cases
64 and pZIKV and ZIKVwpDENV cases elicit similar T cell response patterns, and that HIV-
65 infected individuals have T cell responses that are lower in magnitude compared to HIV-
66 negative individuals. These results have important implications for DENV and ZIKV
67 diagnostic and vaccine development and provide critical insights into the T cell
68 response in individuals with multiple flaviviral infections.

69

70 **IMPORTANCE**

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72 The potential for antibody and T cell cross-reaction to DENV and ZIKV,
73 flaviviruses that co-circulate and can sequentially infect individuals, has complicated
74 diagnostic and vaccine development. Our serological data show that antibodies to
75 nonstructural protein 1 can distinguish sequential human infections by DENV and ZIKV.
76 The development of a simple and inexpensive assay also enables the differentiation of
77 DENV and ZIKV infections based on the characterization of T cell responses. Our T cell
78 data reveals strong response patterns that are similar in nature in individuals with one or
79 multiple DENV infections and in individuals with only primary ZIKV infection and ZIKV-
80 infected individuals with previous DENV exposure. The characterization of T cell
81 responses in a serologically-validated group of individuals is of relevance to the
82 development of vaccines and immunotherapeutics against these global threats.

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93 **INTRODUCTION**

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95 *Aedes* mosquitoes transmit globally relevant flaviviruses including dengue virus
96 (DENV) and Zika virus (ZIKV). DENV exists as four antigenic serotypes, DENV1 to
97 DENV4 (1). These viruses have a wide geographic distribution with approximately 390
98 million infections annually and more than a quarter of the world's population at risk (2).
99 Prior to 2015, ZIKV was considered obscure and known to circulate in Africa and
100 Southeast Asia as two separate viral lineages, African and Asian (3). While most
101 asymptomatic, the clinical presentation of ZIKV infection resembles that of dengue
102 including fever, rash, conjunctivitis, arthralgia, and myalgia (4). In early 2015, thousands
103 of Asian ZIKV cases appeared in northeast Brazil, with accompanying reports of severe
104 neuropathology including congenital microcephaly and Guillain-Barré syndrome (5, 6).
105 In February 2016, the World Health Organization declared ZIKV a public health
106 emergency of international concern (7). By June 2016, autochthonous transmission of
107 ZIKV had been reported in 40 countries and territories throughout South and Central
108 America and the Caribbean (8).

109 The emergence of ZIKV in DENV-endemic regions is of particular concern and
110 relevant for diagnostic and vaccine development. The cocirculation of these genetically
111 similar viruses can result in co-infection or sequential exposure, which has been shown
112 to potentiate cross-reactive immunity at both the antibody and T cell levels (9-12). The
113 envelope (E) protein is the major target of the antibody response in humans during
114 flaviviral infection (1). Antibody-based assays were found to detect extensive cross-
115 reactivity to ZIKV E protein with other flaviviruses, requiring confirmation by plaque

116 reduction neutralization tests (PRNTs) (11, 13-16). These tests, however, are
117 challenged in their ability to confirm infection in individuals with multiple flaviviral
118 infections especially during the acute and early convalescent phases. Several studies
119 have also shown that most dengue-immune sera or DENV E monoclonal antibodies
120 cross-react to ZIKV, but contain limited cross-neutralization activity and can instead
121 enhance ZIKV infection, known as antibody-dependent enhancement (ADE) (17-22). In
122 contrast, recent studies reported antibodies to ZIKV nonstructural protein 1 (NS1) were
123 able to discriminate infections by these viruses (23, 24). We previously showed that
124 combinations of DENV and ZIKV NS1-based enzyme-linked immunosorbent assays
125 (ELISAs) were capable of distinguishing confirmed cases with past and present flaviviral
126 infections including primary DENV (pDENV) and ZIKV (pZIKV), ZIKV with primary
127 DENV (ZIKVwpDENV), and secondary DENV (sDENV) infections (12). These ELISAs
128 are applicable for routine serological tests for DENV and ZIKV as well useful in
129 retrospective studies to identify individuals with primary and multiple flaviviral infections.

130 Pre-existing T cell responses to DENV have also been shown to react to
131 peptides encoded throughout the ZIKV proteome. DENV-naïve mice challenged with
132 ZIKV developed ZIKV-specific CD8⁺ T cells, whereas DENV-immune mice challenged
133 with ZIKV elicited cross-reactive CD8⁺ T cells that reduced infectious ZIKV (25). A study
134 in humans infected with Asian ZIKV demonstrated that DENV serostatus influences the
135 T cell response to ZIKV (10). DENV-immune individuals elicited CD4⁺ and CD8⁺ T cell
136 responses to ZIKV more rapidly and of greater magnitude compared to DENV-naïve
137 ZIKV-infected individuals. In addition, different patterns of immunodominant T cell
138 responses were observed in the case of DENV and ZIKV infections. While CD8⁺ T cell

139 responses against DENV target nonstructural (NS) proteins such as NS3, NS4B, and
140 NS5, ZIKV-specific CD8⁺ T cell responses target the structural proteins, capsid (C),
141 premembrane (prM), and E (10, 26). We previously developed a modified anthrax toxin
142 (LFn)-based enzyme-linked immunospot (ELISPOT) assay, which revealed long-term T
143 cell responses that were ZIKV- and DENV-specific to NS3 protease but cross-reactive
144 to NS3 helicase in individuals infected with DENV and African ZIKV (27). The impact of
145 cross-reactive immune responses in protection or development of ZIKV-mediated
146 neuropathology remains unclear.

147 In this study, we utilized our NS1-based ELISAs to determine the DENV and
148 ZIKV serostatus of individuals from Salvador, Brazil, a DENV-hyperendemic region with
149 one of the highest incidence rates of ZIKV during the 2015-16 epidemic (28). We then
150 tested the ability of our LFn ELISPOT assay to distinguish infections by DENV and
151 Asian ZIKV based on T cells and to characterize those responses.

152

153 **RESULTS**

154

155 **NS1-based ELISAs and neutralization test determine DENV and ZIKV serostatus**

156

157 During the ZIKV outbreak in Salvador, Brazil, acute-phase blood samples were
158 collected from hundreds of suspected ZIKV-infected patients attending HIV outpatient
159 clinics between November 2015 and May 2016. Serological testing for ZIKV-NS1 IgG
160 and DENV-E IgG was performed, revealing a high incidence of ZIKV infection in
161 presumed DENV-immune and -naïve individuals (28). Fifty of these patients were

162 included in the present study. Their median age was 43 (range: 23-72), 49% female,
163 and 76% were human immunodeficiency virus (HIV)-infected. All HIV-infected
164 individuals were on antiretroviral therapy; more than 92% had undetectable viral loads
165 and normal CD4 counts. Patient characteristics and acute serology data are
166 summarized in Table S1 and Table 1, respectively.

167 In order to determine the DENV and ZIKV serostatus among the study
168 participants who had potentially been dual exposed, we collected late convalescent-
169 phase blood samples and employed our previously developed ZIKV-NS1 and DENV-
170 NS1 IgG ELISAs (22). For samples positive for DENV-NS1, we calculated the ratio of
171 relative optical density (rOD) of ZIKV-NS1 to that of DENV-NS1 and used the rOD ratio
172 $< \text{or } \geq 0.24$ to determine sDENV or ZIKVwpDENV infection, respectively (22). Twelve
173 ZIKVwpDENV and 21 sDENV infections were identified (Table 1, Fig. 1A to C). Five
174 samples with ZIKV-NS1 positive and DENV-NS1 negative were pZIKV. Since these
175 samples were collected more than one year post-infection, some anti-NS1 antibodies
176 may have declined to levels below detection, we further tested with ZIKV and DENV E
177 protein-based ELISAs and identified four samples negative for both ZIKV and DENV in
178 all four ELISAs tested (Table 1). Based on the difference in rOD of ZIKV and DENV E
179 proteins ($\Delta\text{rOD} = \text{rOD of ZIKV} - \text{rOD of DENV}$), we identified four pZIKV ($\Delta\text{rOD} \geq 0.17$)
180 and four pDENV ($\Delta\text{rOD} < -0.17$) infections (Table 1, Fig. 1D to E). The negative, pZIKV
181 and, pDENV samples were further confirmed by micro-neutralization tests to DENV1-4
182 and ZIKV; all four negative samples had NT_{90} titers < 10 to DENV1-4 and ZIKV, and the
183 five pZIKV and four pDENV samples showed monotypic neutralization pattern to ZIKV
184 and to one of the four DENV serotypes, respectively (Table 1). For the remaining 33

185 samples, microneutralization tests to ZIKV, DENV1 and/or DENV2 or 3 were performed
186 to show that all 12 ZIKVwpDENV samples neutralize (NT_{90} titers ≥ 10) ZIKV plus at least
187 one DENV serotype, whereas all 21 sDENV samples neutralize at least two DENV
188 serotypes or DENV plus ZIKV; these patterns were compatible with unspecified
189 flavivirus infection according to CDC guidelines (16). An additional three samples
190 (ZK1004, 1005, 1008), which had ΔrOD between -0.17 and 0.17 , positive ZIKV-NS1
191 IgG at acute-phase sera but negative at late convalescent-phase, were classified as
192 undetermined (Table 1). Another sample (ZK0981), for which DENV acute-phase
193 serology was not performed, was negative for ELISAs using late convalescent-phase
194 serum and was also classified as undetermined.

195

196 **T cell responses to NS3 distinguish DENV and ZIKV infections**

197

198 We recently reported the development of a LFn ELISPOT assay based on NS3
199 protease and helicase to distinguish DENV and African ZIKV human infections (27). To
200 assess the ability of the assay to distinguish infection by DENV and Asian ZIKV, we
201 performed DENV and ZIKV homologous and heterologous LFn-NS3 protease and
202 helicase stimulation of date-matched late convalescent-phase PBMCs in an IFN- γ and
203 TNF- α ELISPOT among the serological-validated pDENV, pZIKV, sDENV, and
204 ZIKVwpDENV, and undetermined cases. Using a NS3 protease to helicase ratio cutoff
205 of 1.05 for the IFN- γ ELISPOT, pDENV and sDENV cases and pZIKV, ZIKVwpDENV
206 and the 3 out of the 4 serologically undetermined cases appeared to group together
207 (Fig. 2A). From the undetermined cases, 3 out of 4 (ZK1004, 1005, 1008) grouped with

208 the ZIKV-exposed individuals, while ZK0981 grouped with the DENV-infected
209 individuals. Using a ratio cutoff of 1.048 for the TNF- α ELISPOT, similar groupings were
210 observed (Fig. 2B). We were unable to distinguish sequential infections based on T cell
211 responses to NS3 protease and helicase.

212 Test data were further analyzed to define sensitivity (identifying true positives;
213 individuals who had been infected by DENV versus ZIKV) and specificity (true
214 negatives; DENV- or ZIKV-uninfected individuals). We evaluated sensitivity and
215 specificity as functions of the IFN- γ and TNF- α cutoff values, above which a sample was
216 considered positive and below which a sample was considered negative. We grouped
217 pDENV and sDENV and pZIKV and ZIKVwpDENV cases together based on the
218 clustering observed and excluded the 4 serologically undetermined cases from the
219 analysis. Receiver Operating Characteristic (ROC) curves and corresponding numerical
220 values illustrate the performance of the ELISPOTs as a function of the discrimination
221 threshold, plotted as sensitivity versus 1 – specificity. The areas of the ROC curves
222 represent test performance, where 1 represents a perfect test, and 0.5 represents a
223 random predictor. We measured areas of 0.96 and 0.97 for the IFN- γ and TNF- α
224 ELISPOTs, respectively (Table 2, Fig. 2 C-D). Using the cutoff values, the test
225 sensitivity and specificity for both the IFN- γ and TNF- α ELISPOTs were 94% and 92%,
226 respectively.

227

228 **LFn-DENV and -ZIKV structural and nonstructural proteins elicit robust T cell**
229 **responses and prior DENV exposure does not affect the response**

230

231 To assess the magnitude of T cell responses among the study participants, we
232 stimulated late convalescent-phase PBMCs in IFN- γ and TNF- α ELISPOTs using the
233 following six LFn fusion proteins: LFn-DENV-NS3-Protease (LFn-DV NS3-P), LFn-
234 DENV-NS3-Helicase (LFn-DV NS3-H), LFn-ZIKV-Capsid (LFn-ZV C), LFn-ZIKV-
235 premembrane (LFn-ZV prM), LFn-ZIKV-NS3-Protease (LFn-ZV NS3-P), LFn-ZIKV-
236 NS3-Helicase (LFn-ZV NS3-H). Individuals with pDENV and sDENV infections elicited
237 similar IFN- γ and TNF- α T cell response patterns (Fig. 3 A-B). These individuals had T
238 cell responses to LFn-DV NS3-H and LFn-ZV NS3-H that were stronger in magnitude
239 than to LFn-DV NS3-P and LFn-ZV NS3-P, respectively. Additionally, T cell responses
240 to LFn-DV NS3-P and NS3-H were stronger compared to LFn-ZV NS3-P and NS3-H.
241 The amount of T cell cross-reaction to the ZIKV structural proteins (LFn-ZV C and LFn-
242 ZV prM) was limited, compared to high cross-reactivity to LFn-ZV NS3-P and NS3-H.
243 Furthermore, individuals with pZIKV and ZIKVwpDENV infections elicited T cell
244 responses to LFn-ZV NS3-H and LFn-DV NS3-H that were stronger in magnitude
245 compared to LFn-ZV NS3-P and LFn-DV NS3-P, respectively (Fig. 3 C-D). While
246 individuals with pZIKV and ZIKVwpDENV infections had stronger IFN- γ T cell responses
247 to LFn-ZV NS3-H than to the ZIKV structural proteins, TNF- α responses to the ZIKV
248 structural proteins were stronger than to LFn-ZV NS3-P.

249 We further evaluated the impact of DENV immunity on the magnitude of T cell
250 responses. We compared the magnitude of the IFN- γ and TNF- α T cell responses
251 between individuals with pDENV and sDENV infections and pZIKV and ZIKVwpDENV
252 infections. In all cases, T cell responses in individuals with prior DENV exposure were
253 not significantly higher compared to individuals with a primary DENV or ZIKV infection

254 (Fig. 3 A-D). While IFN- γ and TNF- α responses appeared stronger to the ZIKV
255 structural proteins in individuals with sDENV infections than to individuals with pDENV
256 infections, these differences were not statistically significant. Similarly, individuals with
257 ZIKVwpDENV infection had comparable IFN- γ and TNF- α T responses to those with
258 pZIKV infections (Fig. 3 C-D).

259

260 **HIV influences the T cell response in DENV-exposed individuals**

261

262 We also compared the magnitude of the IFN- γ and TNF- α T cell responses in
263 DENV-exposed (grouping individuals with pDENV and sDENV infections together),
264 pZIKV, and ZIKVwpDENV individuals who were HIV-negative or HIV-infected. DENV-
265 exposed HIV-negative individuals had stronger IFN- γ responses to LFn-ZV C, -ZV NS3-
266 P, and -ZV NS3-H compared to HIV-infected individuals. IFN- γ responses to LFn-DV
267 NS3-P and NS3-H appeared to be stronger in the HIV-infected individuals, although
268 these differences were not statistically significant ($p=0.61$ and $p=0.13$, respectively)
269 (Fig. 4 A). A similar pattern of responses was observed for TNF- α (Fig. 4 D). In general,
270 ZIKVwpDENV HIV-negative individuals had stronger IFN- γ and TNF- α responses
271 compared to individuals that were HIV-infected (Fig. 4 B and E). In contrast, there was
272 largely no difference in the IFN- γ and TNF- α responses in pZIKV HIV-negative and HIV-
273 infected individuals (Fig. 4 C and F). There was an exception where the TNF- α
274 response to LFn-ZV NS3-H was stronger in individuals that were HIV-negative.

275

276

277 **DISCUSSION**

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279 We report on the characterization of late convalescent-phase antibody and T cell
280 responses in individuals from Salvador, Brazil, a DENV-hyperendemic region that was
281 burdened by the 2015-2016 ZIKV outbreak. Our study presents three major findings in a
282 serologically-validated group of DENV and/or ZIKV infected individuals. First, IFN- γ and
283 TNF- α T cell response ratios of ZIKV NS3 protease to DENV NS3 protease can
284 discriminate infections in individuals exposed to these viruses. Second, individuals with
285 pDENV and sDENV infections have similar T cell response patterns, with extensive
286 cross-reactivity to ZIKV NS3 helicase, whereas individuals with pZIKV and
287 ZIKVwpDENV infections have strong responses to both ZIKV structural and
288 nonstructural proteins, with high cross-reaction to DENV NS3 helicase. Third, HIV-
289 infection is associated with responses that are lower in magnitude in DENV exposed
290 individuals.

291 Our previous study of NS1-based ELISAs on convalescent-phase sera from RT-
292 PCR confirmed cases with pZIKV, pDENV, sDENV and ZIKVwpDENV infections
293 showed that sDENV infection panel cross-react to ZIKV-NS1 and the rOD ratio of ZIKV-
294 NS1 to DENV-NS1 in IgG ELISA can distinguish sDENV and ZIKVwpDENV infections
295 (22). Since anti-NS1 antibodies may decline over time and become undetectable
296 especially for those with primary infection, we further tested these samples with E
297 protein-based IgG ELISAs and identified four negative samples, five pZIKV and four
298 pDENV infections. All these 13 samples have been verified by neutralization test using
299 NT₉₀ ≥ 10 as cutoff based on the CDC guidelines (16), suggesting that Δ rOD based on

300 ZIKV-E and DENV-E IgG ELISAs can distinguish pZIKV and pDENV infections; this
301 could potentially be a useful tool for epidemiology and pathogenesis study in endemic
302 regions. However, the sample size is small and the Δ rOD of 0.17 was based on a single
303 serum dilution of 1:800, future studies involving larger sample size and different
304 dilutions or end-point titers are needed to further validate these observations.

305 The degree of amino acid sequence identity between DENV and ZIKV structural
306 and nonstructural proteins is 49% and 51%, respectively (10). Multiple sequence
307 alignment and homology determination of DENV and ZIKV NS3 demonstrates high
308 amino acid sequence identity of 67%, with protease and helicase homology of 58% and
309 72%, respectively, consistent with the higher degree of DENV/ZIKV cross-reaction in
310 NS3 helicase (Table 3, Fig. 5). Our recent characterization of acute- and convalescent-
311 phase T cells collected from individuals infected with DENV and African ZIKV in
312 Senegal, West Africa, revealed sustained DENV- and ZIKV-specific responses to NS3
313 protease and cross-reactive responses to NS3 helicase (27). Our findings in individuals
314 infected with DENV and Asian ZIKV are in agreement with our previous observations.
315 Although we were unable to distinguish sequential exposure, the LFn NS3 protease
316 ELISPOT differentiates infections between DENV- and ZIKV-infected individuals with
317 high sensitivity and specificity of 94% and 92%, respectively.

318 A relatively large body of epidemiological and laboratory-based evidence has
319 suggested that severe and often fatal forms of dengue disease occurs most commonly
320 during a secondary infection by a heterotypic DENV serotype (29, 30). Another
321 phenomenon, known as original antigenic sin (OAS), has been observed in antibody as
322 well as T-cell responses, in which less-effective T cells generated in response to a

323 primary DENV infection predominate during a subsequent infection with a different
324 DENV serotype, resulting in an inappropriate response and predisposing individuals to
325 severe disease (31, 32). The OAS hypothesis was challenged by a study in Sri Lankan
326 individuals infected with DENV, which showed that the phenomenon does not generate
327 less functional responses, but instead correlates with protective responses to conserved
328 viral sequences (26). Unexpectedly, we did not observe T cell responses that were
329 significantly higher in magnitude in individuals with prior DENV exposure. These results
330 are in contrast to our data on African ZIKV infections, which showed that previous
331 flavivirus exposure was associated with enhanced T cell responses (27). One possibility
332 is that the proportion of HIV infection among those with prior DENV exposure was
333 higher compared with DENV-naïve in this study (90.5% versus 50% comparing sDENV
334 and pDENV cases; 83.3% versus 60% comparing ZIKVwpDENV and pZIKV cases).
335 Nevertheless, as co-circulation of DENV, ZIKV, and other flaviviruses occurs throughout
336 many parts of the world, it is critical to continue to develop tools to better understand T
337 cell immunity in individuals exposed to multiple flaviviruses.

338 A recent study using human leukocyte antigen (HLA) transgenic mice infected
339 with DENV2 and Asian and African ZIKV strains revealed cross-reactive T cell
340 responses to HLA-restricted epitopes (25). Out of 8 ZIKV NS3 epitopes computationally
341 predicted to bind HLA class I molecules, only 3 epitopes elicited DENV2/Asian ZIKV
342 cross-reactive T cell responses. Of note, the cross-reactive epitopes were all positioned
343 within the helicase domain of NS3, further supporting our observations of high
344 DENV/ZIKV NS3 helicase T cell cross-reaction. Another study demonstrated ZIKV-
345 specific and ZIKV/DENV cross-reactive T cell responses in humans (10). T cell

346 responses generated in response to prior DENV exposure recognized peptides
347 sequences located throughout the ZIKV proteome. DENV serostatus also influenced T
348 cell immunity to ZIKV. DENV-naïve ZIKV-positive individuals had predominant CD8 T
349 cell responses directed against structural proteins. In contrast, a majority of CD8 T cells
350 responses were directed against nonstructural proteins in DENV-immune ZIKV-positive
351 individuals, suggesting that previous DENV exposure can alter the T cell response.

352 While the above studies used peptide stimulation to characterize the T cell
353 response, there are concerns around this approach (33). Some of the HLA-predicted
354 peptides may fail to stimulate strong T cell responses as expected. Longer and shorter
355 peptides have also been shown to elicit different types of responses (34-36). An
356 alternative to peptide stimulation is the anthrax LFn, which has the capability to deliver
357 full length antigen into the cytosol for native processing via the MHC pathways, and to
358 elicit better T cell responses compared to peptides in some cases (37-42). Our
359 adaptation of the LFn ELISPOT, not only allowed detection of human DENV and ZIKV
360 infections, but also characterization of the associated T cell responses to structural and
361 nonstructural proteins. We demonstrated that individuals with pDENV and sDENV
362 infections had similar IFN- γ and TNF- α T cell response patterns with high cross-
363 reactivity to ZIKV NS3 helicase, but low cross-reactivity to the ZIKV structural proteins.
364 A small number of individuals with sDENV infections had cross-reactive T cell
365 responses to ZIKV structural proteins. Interestingly, however, individuals with pZIKV
366 and ZIKVwpDENV infections had similar IFN- γ and TNF- α T cell responses patterns,
367 with strong responses to structural and nonstructural proteins. It is noteworthy that we
368 observed comparably strong T cell responses to the structural proteins in pZIKV and

369 ZIKVwpDENV cases, in contrast to Grifoni et al., suggesting that the most recent
370 infection may dictate the T cell response (10). Another possibility that cannot be
371 excluded is the differences in T cell stimulation strategies, which may be contributing to
372 the observed differences. Additionally, due to the limited collection of blood samples
373 from each patient, we were unable to characterize CD4- and CD8-specific responses.
374 Future characterization studies using the LFn delivery system on CD4 and CD8 T cells
375 will be important.

376 Our study of ZIKV seroprevalence in West Africa demonstrated continued human
377 transmission of the virus in HIV- and malaria-infected individuals (43). The co-infection
378 of flaviviruses with HIV or malaria could potentially impact pathophysiological
379 mechanisms, induce different clinical and laboratory findings, and interfere with
380 treatment. Previous studies have shown a suppression of HIV-1 replication during acute
381 DENV infection (44, 45). In this study, we demonstrated that DENV-exposed individuals
382 who are HIV-infected had T cell responses that were significantly lower in magnitude
383 compared to HIV-negative individuals except in individuals with pZIKV infections. We
384 also observed that DENV-exposed HIV-infected individuals have T cell responses that
385 were lower in magnitude to ZIKV proteins compared to DENV-exposed HIV-negative
386 individuals. Whether HIV infection in DENV-exposed individuals reduces the ability to
387 induce cross-reactive T cell responses has important implications. More studies with
388 larger sample sizes are needed to increase our limited understanding of the
389 epidemiological and immunopathogenesis interactions of flavivirus exposure in
390 individuals with HIV and other comorbidities.

391 In summary, despite high sequence homology between DENV and ZIKV,
392 diagnostic assays based on antibodies to NS1 and T cell responses to NS3 protease
393 are effective at distinguishing human infections by these viruses. The LFn ELISPOT
394 assay has enabled direct comparison of T cell characterization in DENV, Asian and
395 Africa ZIKV human infections. As vaccines against DENV and ZIKV are currently being
396 developed, the information generated from these characterization studies are of high
397 relevance. The results of these characterization studies may contribute to the design
398 and development of DENV and ZIKV vaccines and T cell based diagnostics.

399

400 **MATERIALS AND METHODS**

401

402 **Clinical samples and ethical statement**

403

404 Fifty late convalescent-phase blood samples were obtained from patients at
405 Professor Edgard Santos University Hospital, Federal University of Bahia, Salvador,
406 Brazil. These individuals were suspected ZIKV-infected during the 2015-2016 ZIKV
407 epidemic and their acute-phase sera were screened for ZIKV and DENV antibodies.
408 Late convalescent-phase peripheral blood mononuclear cells (PBMCs) were separated
409 from whole blood in EDTA tubes by Ficoll-Hypaque gradient density (Sigma-Aldrich, St.
410 Louis, MO, USA) and cryopreserved in freezing media (10% dimethyl sulfoxide [DMSO],
411 Sigma-Aldrich, St. Louis, MO, USA) at -80°C overnight prior to transfer to liquid
412 nitrogen. Convalescent-phase serum was aliquoted and immediately transferred to -
413 80°C.

414 The Federal University of Bahia Institutional Review Board (IRB), the Harvard
415 T.H. Chan School of Public Health IRB, and the University of Hawaii IRB approved the
416 primary studies under which the samples and data were collected. All patients provided
417 informed consent for the collection of samples. Excess samples and corresponding data
418 were banked, coded prior to analyses, and stored at the Federal University of Bahia.

419

420 **ELISAs**

421

422 For acute-phase sera, commercial ZIKV-NS1 and DENV-E based IgG ELISAs
423 (Euroimmun, Luebeck, Germany) were performed (27). For late convalescent-phase
424 sera, ZIKV- and DENV1-NS1 IgG ELISAs were performed as described previously (22). Briefly,
425 purified NS1 proteins (16 ng per well) were coated onto 96-well plates overnight, followed by
426 blocking and incubation with primary (serum at 1:400 dilution) and secondary (anti-
427 human IgG conjugated with HRP, Jackson) antibodies (22). The OD at 450 nm was
428 read with a reference wavelength of 650 nm. Each ELISA plate included two positives
429 (two confirmed-Zika or confirmed-dengue samples for ZIKV- and DENV-NS1 ELISAs,
430 respectively), four negatives (4 flavivirus-naïve sera), and tested samples (all in
431 duplicates). The OD values were divided by the mean OD value of positive controls to
432 calculate the rOD values. The cut-off was defined by the mean rOD value of negatives
433 plus 12 standard deviations as described previously (22). For samples positive for both
434 ZIKV- and DENV-NS1 ELISAs, the ratio of rOD ($=\text{rOD of ZIKV-NS1}/\text{rOD of DENV-NS1}$)
435 was calculated; rOD ratio $<$ or \geq 0.24 indicated sDENV or ZIKVwpDENV infection,
436 respectively (22).

437 E protein-based IgG ELISAs using DENV1 virion or ZIKV (MR766 strain) virus
438 like particles (VLP) were also tested for late convalescent-phase sera (46). Briefly,
439 DENV1 virions or ZIKV-VLP derived from ultracentrifugation of culture supernatants of
440 virus-infected Vero cells or pENTR-ZIKV prME plasmid-transfected 293T cells,
441 respectively, were UV inactivated (for virions) and coated on 96-well plates at 4°C
442 overnight, followed by blocking and incubation with primary (serum at 1:800 dilution)
443 and secondary antibodies as above. The rOD and cut-off rOD values were similarly
444 calculated. The difference in rOD of ZIKV and DENV E proteins ($\Delta rOD = rOD \text{ of ZIKV} -$
445 $rOD \text{ of DENV}$) was determined; $\Delta rOD \geq 0.17$ or < -0.17 was classified as pZIKV or
446 pDENV infection, respectively.

447

448 **Neutralization test**

449

450 PRNT was performed on acute-phase sera to detect neutralization antibody to
451 ZIKV as reported previously (47). For late convalescent-phase sera, a previously
452 described micro-neutralization test was performed (48). Briefly, flat-bottom 96-well
453 plates were seeded with Vero cells (3×10^4 cells per well) 24 h prior to infection. Four-
454 fold serial dilutions of serum (starting from 1:10) were mixed with 50 focus-forming units
455 of DENV1 (Hawaii strain), DENV2 (NGC strain), DENV3 (CH53489), DENV4 (H241
456 strain) or ZIKV (PRVABC59 strain) at 37°C for 1 h. The mixtures were added to each
457 well followed by incubation for 48 h (except 70 h for DENV1), removal of medium, and
458 fixation as described previously (46). After adding murine mAb 4G2 and secondary
459 antibody mixture (IRDye® 800CW-conjugated goat anti-mouse IgG at 1:10000 and

460 DRAQ5™ Fluorescent Probe at 1:10000), the signal (800 nm/700 nm fluorescence) was
461 detected by Li Cor Odyssey classic (LiCor Biosciences) and analyzed by Image Studio
462 software to determine percent neutralization at different concentrations and NT₉₀ as
463 described previously (46, 48).

464

465

466 **LFn fusion protein design**

467

468 Commercially synthesized gene fragments encoding the NS3 protease and
469 helicase of DENV2 and C, prM, and NS3 protease and helicase of Asian ZIKV and were
470 cloned into the LFn expression vector (pET15bLFn). The pET15bLFn vector contains a
471 T7 promoter, histidine tag (His₆), and the terminal domain of the anthrax lethal factor
472 (LFn; 255 amino acids). The pET15bLFn containing the coding sequences of the DENV
473 and Asian ZIKV proteins were transformed into *E. coli* BLR (DE3) (Millipore, Medford,
474 MA, USA). Selected clones were sequenced to verify the reading frame, and clones
475 containing the correct sequence were used for protein expression.

476 The LFn-DENV and -ZIKV fusion proteins and the LFn control were expressed
477 upon isopropylthiogalactoside ([IPTG], Sigma-Aldrich, St. Louis, MO, USA) induction in
478 5L Luria broth containing carbenicillin and chloramphenicol for 2-4 hours. Cells were
479 pelleted by centrifugation and resuspended in imidazole (1mM) binding buffer
480 (Novagen, Madison, WI, USA) in the presence of a protease inhibitor cocktail (Thermo
481 Fisher Scientific, Rockford, IL, USA). Cell pellets were sonicated, centrifuged at 4°C,
482 and the supernatants were loaded in an equilibrated nickel-charged column for affinity

483 purification. The bound proteins were eluted in 100-200 mM imidazole, desalted with a
484 Sephadex G-25M column (Sigma-Aldrich, St. Louis, MO, USA), and eluted in PBS
485 (Sigma-Aldrich, St. Louis, MO, USA). The PBS-eluted proteins were passed through
486 Detoxi-Gel (Thermo Fisher Scientific, Rockford, IL, USA). Protein concentrations were
487 determined samples were stored at -80°C.

488

489 **ELISPOT assay**

490

491 *Ex vivo* ELISPOTs were performed as previously described. Briefly, 96-well
492 polyvinylidene difluoride (PVDF)-backed MultiScreen_{HTS} (MSIP) microtiter plates
493 (Millipore, Medford, MA, USA) were treated with 100ul of 90% ethanol for 30 seconds
494 and washed 5 times with sterile PBS. Plates were coated with 100ul of capture
495 antibodies (Abs) in PBS. Plates containing capture Abs were incubated overnight at
496 4°C. Plates were then blocked with 1% bovine serum albumin ([BSA], Sigma-Aldrich, St.
497 Louis, MO, USA) in PBS and washed 6 times with PBS. Cryopreserved PBMCs were
498 thawed in R10 medium and incubated overnight at 37°C. PBMCs were washed 2 times
499 with PBS and seeded at 2×10^5 cells/well in a final volume of 100ul/well. LFn-DENV
500 and -ZIKV proteins were added to each well. As a positive control, PBMCs were
501 stimulated with phytohemagglutinin ([PHA], Sigma-Aldrich, St. Louis, MO, USA). As a
502 negative control, wells received LFn. After incubation for 24-28 hours at 37°C in 5%
503 CO₂, the cells were discarded and plates were washed 3 times with PBS and 3 times
504 with PBS with 0.05% Tween-20 ([PBST], Bio Rad Technologies, Hercules, CA, USA) to
505 remove cells. The detection antibodies were added and plates were incubated overnight

506 at 4°C. Plates were then washed 6 times with PBST, then incubated for 2 hours at room
507 temperature with mixtures containing the enzymatic conjugates. To develop spots,
508 plates were washed 4 times with PBST, three times with PBS, and 1 time with water.
509 Vector Blue substrate solution (Vector Laboratories, Burlingame, CA, USA) was added
510 for 5-15 mins before rinsing with water and air-drying. Digitized images were analyzed
511 for spots using CTL ImmunoSpot reader (Cellular Technology Limited, Cleveland, OH,
512 USA). DENV and ZIKV spots were calculated by subtracting the mean of the negative
513 control value from the mean value of the specific stimulation. Positive responses had to
514 be greater than 4 times the mean background, 3 standard deviations above the
515 background, and ≥ 55 spot-forming cells per (SFC)/ 10^6 PBMCs.

516

517 **ROC analysis**

518

519 The ELISPOTs were validated using PBMCs from individuals that were
520 confirmed DENV- and/or ZIKV-infected by ELISA and/or neutralization tests. DENV and
521 ZIKV NS3 protease to helicase values were calculated, resulting in normalized test
522 ratios (ZIKV NS3 protease divided by DENV NS3 protease) ranging from 0.15-2.95. On
523 the basis of these data, we determined the optimal cutoffs between 0.15 and 2.95 by
524 calculating the sensitivity (number of true positives divided by total confirmed positive
525 values) and specificity (number of true negatives divided by the total confirmed
526 negatives) at increasing 0.05 to the theoretical cutoffs. After calculating the sensitivity
527 and specificity values, the optimal cutoffs were defined as the highest sum of sensitivity
528 and specificity, such that the optimal cutoff values reflected the optimal sensitivity and

529 specificity. The optimal cutoffs obtained for the IFN- γ and TNF- α was 1.05 and 1.048,
530 respectively (Prism 7, GraphPad Software, San Diego, CA, USA).

531

532 **Multiple sequence alignment and percent homology analysis**

533

534 Multiple sequence alignment of DENV1-4 and ZIKV NS3 was performed using
535 the Clustal Omega program (EMBL-EB, Cambridgeshire, UK). Averages of DENV and
536 ZIKV NS3 protease and helicase proteins were calculated using the ExPASy
537 Bioinformatics Resource Portal (Swiss Institute of Bioinformatics, Lausanne,
538 Switzerland) and based on averages of the different homology values in the four DENV
539 serotypes and ZIKV. Average conservation was determined on a per-residue basis for
540 NS3 protease, helicase, and full-length protein.

541

542 **Statistical analysis**

543

544 Statistical analysis was performed using Prism 7 (GraphPad Software, San
545 Diego, CA, USA). Where appropriate, data were expressed as geometric positive
546 means on box whisker and bar graphs \pm standard deviation. Data comparisons were
547 conducted using the Wilcoxon rank sum test. A threshold of $p < 0.05$ was considered
548 statistically significant.

549

550

551

552 **Data Availability**

553

554 All relevant data has been included in the manuscript. We will provide any
555 additional data upon request.

556

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558

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566

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757 TABLES

758 Table 1 Results of Serological tests

ID	Acute-phase sera			Late convalescent-phase serum						Interpretation
	ZIKV-NS1 IgG ^a	DENV-E IgG ^a	PRNT test ^b	ZIKV-NS1 IgG ^c	DENV-NS1 IgG ^c	rOD ratio ^c	ZIKV-E IgG ^d	DENV-E IgG ^d	NT test ^f D1/D2/D3 /D4/ZIKV	
ZK0978	-	-	-	-	-	NA	-	-	<10/<10/<10/<10/<10	negative
ZK0982	-	-	-	-	-	NA	-	-	<10/<10/<10/<10/<10	negative
ZK0987	-	-	-	-	-	NA	-	-	<10/<10/<10/<10/<10	negative
ZK0999	-	-	-	-	-	NA	-	-	<10/<10/<10/<10/<10	negative
ZK0979	+	+		+	-	NA	+ ^e	+	<10/<10/<10/<10/>160	pZIKV
ZK0993	+	+		+	-	NA	+ ^e	+	<40/<40/<10/<10/>160	pZIKV
ZK0998	+	+		+	-	NA	+ ^e	+	<10/<10/<40/<10/>160	pZIKV
ZK1006	+	+		+	-	NA	+ ^e	+	<80/<40/<10/<40/640	pZIKV
ZK0996	+	-	+	+	-	NA	+ ^e	+	<10/<10/<10/<10/>160	pZIKV
ZK0966	-	+	-	-	-	NA	+	+ ^e	<40/160/<10/<40/<10	pDENV
ZK0980	-	+		-	-	NA	+	+ ^e	<40/<40/>160/<40/<10	pDENV
ZK0995	-	+		-	-	NA	+	+ ^e	>640/160/<10/<10/<10	pDENV
ZK0997	ND	ND		-	-	NA	+	+ ^e	<40/>160/<40/<40/<10	pDENV
ZK0972	+	+		+	+	≥0.24	+	+	>40/ND/ND/ND/>80	ZIKVwpDENV
ZK0975	+	+		+	+	≥0.24	+	+	>40/ND/ND/ND/>80	ZIKVwpDENV
ZK0989	+	+	+	+	+	≥0.24	+	+	>40/ND/ND/ND/>80	ZIKVwpDENV
ZK0991	+	+	+	+	+	≥0.24	+	+	>40/ND/ND/ND/>80	ZIKVwpDENV
ZK1000	+	+		+	+	≥0.24	+	+	>40/ND/ND/ND/>40	ZIKVwpDENV
ZK1009	+	+	+	+	+	≥0.24	+	+	>10/ND/ND/ND/>80	ZIKVwpDENV
ZK1011	+	+		+	+	≥0.24	+	+	>40/ND/ND/ND/>80	ZIKVwpDENV
ZK1012	+	+	+	+	+	≥0.24	+	+	>40/ND/ND/ND/>80	ZIKVwpDENV
ZK1014	+	+	+	+	+	≥0.24	+	+	>40/ND/ND/ND/>80	ZIKVwpDENV
ZK1015	+	+	+	+	+	≥0.24	+	+	>40/ND/ND/ND/>80	ZIKVwpDENV
ZK0968	+	ND		+	+	≥0.24	+	+	>40/ND/ND/ND/>80	ZIKVwpDENV
ZK0984	+	+		+	+	≥0.24	+	+	>320/>1280/<80/>1280/>320	ZIKVwpDENV
ZK0976	-	+		+	+	<0.24	+	+	>40/ND/ND/ND/<10	sDENV
ZK0986	-	+		+	+	<0.24	+	+	>40/ND/ND/ND/>80	sDENV

ZK0967	+	+		+	+	<0.24	+	+	>40/ND/ND/ND/>80	sDENV
ZK0969	+	+		+	+	<0.24	+	+	>40/ND/ND/ND/>40	sDENV
ZK0971	+	+		+	+	<0.24	+	+	>40/>40/ND/ND/<10	sDENV
ZK0977	+	+		+	+	<0.24	+	+	>40/ND/ND/ND/>80	sDENV
ZK0983	+	+	+	+	+	<0.24	+	+	>40/ND/ND/ND/>10	sDENV
ZL0985	+	+	+	+	+	<0.24	+	+	>40/ND/ND/ND/>40	sDENV
ZK0988	+	+		+	+	<0.24	+	+	>40/ND/ND/ND/>40	sDENV
ZK0992	+	+	+	+	+	<0.24	+	+	>40/ND/ND/ND/>80	sDENV
ZK0994	+	+	-	+	+	<0.24	+	+	>40/ND/ND/ND/>40	sDENV
ZK1001	+	+		+	+	<0.24	+	+	>40/>40/ND/ND/<10	sDENV
ZK1010	+	+		+	+	<0.24	+	+	>40/ND/ND/ND/>80	sDENV
ZK1013	+	ND		+	+	<0.24	+	+	>40/ND/ND/ND/>10	sDENV
ZK0973	-	ND		-	+	<0.24	+	+	>40/<80/>80/ND/>320	sDENV
ZK0974	-	+		-	+	<0.24	+	+	>40/>320/>80/ND/>10	sDENV
ZK1002	-	+	-	-	+	<0.24	+	+	>40/>80/<80/ND/<10	sDENV
ZK1003	-	+	-	-	+	<0.24	+	+	>40/>80/<80/ND/<10	sDENV
ZK1007	-	+		-	+	<0.24	+	+	>40/>80/>80/ND/<10	sDENV
ZK1016	-	+		-	+	<0.24	+	+	>40/>1280/>1280/ND/>10	sDENV
ZK0990	-	+	-	-	+	<0.24	+	+	>40/>1280/>80/ND/<10	sDENV
ZK1004	+	+		-	+	NA	+	+	ND/ND/ND/ND/ND	unknown
ZK1005	+	ND		-	+	NA	+	+	ND/ND/ND/ND/ND	unknown
ZK1008	+	+		-	-	NA	+	+	ND/ND/ND/ND/ND	unknown
ZK0981	-	ND		-	-	NA	-	-	ND/ND/ND/ND/ND	unknown

759

760 pZIKV: primary ZIKV infection; pDENV: primary DENV infection; sDENV: secondary DENV infection; ZIKVwpDENV: ZIKV infection

761 with previous DENV infection. ND: not determined; NA not applicable.

762 ^aEuroimmun ZIKV-NS1 and DENV-E IgG ELISAs were performed on acute-phase sera (28).

763 ^bPRNT was performed on acute-phase sera to detect neutralization antibody to ZIKV (47).

764 ^cZIKV-NS1 and DENV-NS IgG ELISAs were described previously (12). The rOD ratio (ZIKV-NS1/DENV-NS1) < or ≥0.24 was
765 classified as sDENV or ZIKVwpDENV (12).

766 ^deZIKV-E and DENV-E IgG ELISAs utilized ZIKV VLP and DENV virions, respectively (46). ΔrOD (rOD of ZIKV–rOD of DENV) ≥
767 0.17 or < –0.17 was classified as pZIKV or pDENV infection, respectively.

768 ^fMico-neutralization test (NT) were performed (with NT₉₀ titers shown) to confirm none, pZIKV or pDENV infection (46, 48).

769

770 **Table 2** Numerical values of ROC analysis and sensitivity and specificity results

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	IFN- γ ⁺ DENV vs. ZIKV	TNF- α ⁺ DENV vs. ZIKV
AUC ^a	0.96	0.97
95% CI ^b	0.91-1.02	0.92-1.01
Cutoff	1.055	1.048
Sensitivity	94%	94%
Specificity	92%	92%
^a Area under the curve, ^b Confidence interval		

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Table 3 Sequence homology of DENV and Asian ZIKV NS3^a

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Serotype	% homology to ZIKV		
	NS3 protease	NS3 helicase	Full-length
DENV1	55%	71%	66%
DENV2	58%	72%	67%
DENV3	58%	72%	67%
DENV4	59%	71%	67%
^a Homology analysis between Asian ZIKV (GenBank accession number: NC_035889.1) and DENV1 (ACO06157.1), DENV2 (JN819419.1), DENV3 (ACY70771.1), and DENV4 (AEW50183.1).			

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794 **Figure 1. ZIKV and DENV NS1-based and E-based IgG ELISAs.** (A) ZIKV-NS1, (B)
795 DENV-NS1 IgG ELISAs and (C) rOD ratio. (D) ZIKV-E, (E) DENV-E and (F) Δ rOD=rOD
796 of ZIKV–rOD of DENV. Dots lines indicate cut-off values, 0.24 for rOD ratio and 0.17 for
797 Δ rOD. pZIKV: primary ZIKV infection; pDENV: primary DENV infection; sDENV:
798 secondary DENV infection; ZIKVwpDENV: ZIKV infection with previous DENV infection.
799 NA not applicable.

800

801 **Figure 2. T cell responses to NS3 protease and helicase and ROC analysis of the**
802 **ELISPOT test.** Late convalescent-phase PBMCs from DENV- and/or ZIKV-infected
803 individuals were treated with homologous and heterologous LFn-DENV and -ZIKV NS3
804 protease and the specific IFN- γ and TNF- α T cell responses were detected by *ex vivo*
805 ELISPOTs. (A) Scatter plot of the ratios of ZIKV NS3 protease to DENV NS3 protease
806 IFN- γ responses versus ratios of helicase. (B) ROC analysis for the IFN- γ ELISPOT. (C)
807 Scatter plot of the ratios of ZIKV NS3 protease to DENV NS3 protease TNF- α T cell
808 responses versus ratios of helicase. (D) ROC analysis for the TNF- α ELISPOT. The
809 dashed line on (A) represents the optimal cutoff value of 1.05 and the dashed line on
810 (C) represents the optimal cutoff value of 1.048. Individual colored dots represent
811 serologically-validated DENV- and/or ZIKV-infected individuals and the undetermined
812 cases.

813

814 **Figure 3. T cell responses to ZIKV and/or DENV structural or nonstructural**
815 **proteins among subgroups with different DENV and ZIKV serostatus.** Late
816 convalescent-phase PBMCs from DENV- and/or ZIKV-infected individuals were treated

817 with homologous and/or heterologous LFn-DENV and -ZIKV capsid (ZV C),
818 premembrane (ZV prM), NS3 protease (DV or ZV NS3-P) and NS3 helicase (DV or ZV
819 NS3-H) and the specific IFN- γ and TNF- α T cell responses were detected by *ex vivo*
820 ELISPOTs. IFN- γ and TNF- α spot forming cells (SFC) were detected, counted, and
821 expressed as box plots with mean and standard deviations. Comparison of late
822 convalescent-phase (A) IFN- γ and (B) TNF- α T cell responses between individuals with
823 pDENV and sDENV infections. Comparison of late convalescent-phase (C) IFN- γ and
824 (D) TNF- α T cell responses between individuals with pZIKV and ZIKVwpDENV
825 infections. Individual colored plots represent serologically-validated DENV- and/or ZIKV-
826 infected individual. *, $p < 0.05$.

827

828 **Figure 4. Impact of HIV status on the T cell response.** Comparison of mean
829 convalescent-phase IFN- γ T cell responses expressed as bars and standard deviation
830 between (A) HIV-negative (open black bars) and HIV-infected (shaded black bars)
831 individuals with pDENV and sDENV infections grouped together, (B) HIV-negative
832 (open dark grey bars) and HIV-infected (shaded dark grey bars) individuals with
833 ZIKVwpDENV infections, and (C) HIV-negative (open light grey bars) and HIV-infected
834 (shaded light grey bars) individuals with pZIKV infections. Comparison of mean
835 convalescent-phase TNF- α T cell responses expressed as bars and standard deviation
836 between (D) HIV-negative (open black bars) and HIV-infected (shaded black bars)
837 individuals with pDENV and sDENV infections grouped together, (E) HIV-negative
838 (open dark grey bars) and HIV-infected (shaded dark grey bars) individuals with

839 ZIKVwpDENV infections, and (F) HIV-negative (open light grey bars) and HIV-infected
840 (shaded light grey bars) individuals with pZIKV infections. *, $p < 0.05$.

841

842 **Figure 5. Clustal Omega generated amino acid sequence alignment of DENV**
843 **serotypes 1 to 4 and Asian ZIKV.** The residues in yellow represent the NS3 protease
844 domain (amino acids 1-169) and the residues in orange represent the helicase domain
845 (amino acids 179-619). *, single, fully conserved residue. :, conservation between
846 groups of strongly similar properties – scoring > 0.5 in the Gonnet PAM 250 matrix. .,
847 conservation between groups of weakly similar properties – scoring ≤ 0.5 in the
848 Gonnet PAM 250 matrix.









