

1 HIV envelope gp120 alters T-cell receptor mobilization in the immunological
2 synapse of uninfected CD4 T cells and augments the T cell activation

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20 **Abstract**

21 HIV is transmitted most efficiently from cell to cell and productive infection occurs mainly
22 in activated CD4 T cells. It is postulated that HIV exploits immunological synapses
23 formed between CD4 T cells and antigen-presenting cells to facilitate the targeting and
24 infection of activated CD4 T cells. This study sought to evaluate how the presence of
25 the HIV envelope (Env) in the CD4 T cell immunological synapse affects synapse
26 formation and intracellular signaling to impact the downstream T cell activation events.
27 CD4 T cells were applied onto supported lipid bilayers (SLBs) that were reconstituted
28 with HIV Env gp120, anti-TCR antibody OKT3 and ICAM-1 to represent the surface of
29 HIV Env-bearing antigen-presenting cells. The results showed that the HIV Env did not
30 disrupt immunological synapse formation. Instead, the HIV Env accumulated with TCR
31 at the center of the synapse, altered the kinetics of TCR recruitment to the synapse, and
32 affected the synapse morphology over time. The HIV Env also prolonged Lck
33 phosphorylation at the synapse and enhanced TCR-induced CD69 upregulation, IL-2
34 secretion, and proliferation to promote virus infection. These results suggest that HIV
35 uses the immunological synapse as a conduit not only for selective virus transmission to
36 activated CD4 T cells, but also for boosting the T cell activation state, thereby
37 increasing its likelihood for undergoing productive replication in the targeted CD4 T cells.

38 **Importance**

39 There are about two million new HIV infections every year. A better understanding on
40 how HIV is transmitted to the susceptible cells is critical to devise effective strategies to
41 prevent HIV infection. Activated CD4 T cells are preferentially infected by HIV, although
42 how this is accomplished is not fully understood. This study examined whether HIV co-

43 opts the normal T-cell activation process through the so-called immunological synapse.
44 We found that the HIV envelope is recruited to the center of the immunological synapse
45 together with the T-cell receptor and enhances the T cell receptor-induced activation of
46 the CD4 T cells. The heightened cellular activation promotes the capacity of the CD4 T
47 cells to support productive HIV replication. This study provides evidence for the
48 exploitation of the normal immunological synapse and T-cell activation process by HIV
49 to boost the activation state of the targeted CD4 T cells and promote infection of these
50 cells.

51 Introduction

52 HIV infection leads to severe destruction of immune cells and functions. The
53 helper CD4 T cells are one of the main cell types profoundly affected by HIV (1, 2).
54 However, not all CD4 T cells are equally affected by HIV. Although HIV can infect
55 resting naïve CD4 T cells, these cells predominantly express the co-receptor CXCR4
56 and are less likely to express the co-receptor CCR5 required for the entry of the majority
57 of transmitted and circulating HIV-1 isolates. In contrast, many memory CD4 T cells
58 express the co-receptors CXCR4 and CCR5 (3, 4). The post-entry steps in the HIV life
59 cycle are also tightly linked to the activation status of the CD4 T cells. Reverse
60 transcription (5, 6), nuclear import (7), and integration (8) are inefficient unless the CD4
61 T cells are activated and enter the cell cycle. Virus transcription is triggered via NFκB
62 (9), which is activated as a result of the specific signaling cascade triggered upon TCR
63 engagement. Therefore, TCR-activated CD4 T cells are the optimal targets for HIV.
64 Indeed, the recruitment of activated CD4 T cells to the genital or rectal mucosa
65 associated with HSV-2, gonorrhea, and other STDs is considered to be one of the
66 factors that increase the risk for HIV acquisition (10-12). Studies of SIV and SHIV
67 infections in rhesus macaques also showed that the increased numbers of activated
68 CD4 T cells at the site of virus entry constitutes one of the correlates for the increased
69 infection (13, 14). However, the mechanisms by which HIV preferentially targets the
70 activated subsets of CD4 T cells are not fully understood.

71 CD4 T cell activation commences in an immunological synapse, a tight junction
72 at the contact site between a CD4 T cell and an antigen-presenting cell (APC) formed

73 when the CD4 T cell recognizes the cognate peptide-MHC class II (pMHC) complexes
74 on the APC (reviewed in (15, 16)). A CD4 T cell will stop migrating once it has formed
75 an immunological synapse (17). At the periphery of the synaptic area, pMHC/TCR
76 interactions form microclusters which quickly translocate to the center and converge to
77 become the central supramolecular activation cluster (cSMAC) (18, 19). At the same
78 time, ICAM-1/LFA-1 interactions begin clustering to form the peripheral SMAC (pSMAC).
79 A mature stable synapse is thus created with a fully segregated cSMAC and pSMAC
80 ring which arrests the cell migration for over one hour. Recent correlative
81 optical/electron microscopy analyses of the immunological synapse formed on
82 surrogate antigen-presenting cells based on lipid bilayers have provided higher
83 resolution pictures showing the cSMAC region as a cleft containing TCR-rich vesicles
84 (20). These images are highly reminiscent of the electron tomographic data showing
85 the accumulation of HIV virions in the contact area formed between a primary CD4 T
86 cell target and an infected T cell (21).

87 The cell-cell contact area implicated in facilitating the highly efficient HIV
88 transmission from one cell to another is called the virological synapse (22-26). This
89 synapse has similar morphology and utilize many components of the immunological
90 synapse, although the two types of synapses also display distinctive features (27). The
91 assembly of virological synapses is triggered by the binding of the HIV Env on infected
92 cells to CD4 on the target cells and can occur independently of TCR-pMHC
93 engagement (28). Similar to the immunological synapse, the formation of the HIV Env-
94 induced synapse also results in the arrest of CD4 T cell migration (28). During this time,
95 HIV Env-CD4 microclusters form and quickly converge to assemble a cSMAC-like

96 cluster surrounded by the pSMAC-like ICAM-1/LFA-1 ring (29). The TCR alpha beta
97 chains are also recruited to the cSMAC-like cluster albeit in the absence of cognate
98 pMHCs (29). However, when CD4 T cells encounter HIV-infected APCs, simultaneous
99 HIV Env-CD4 and TCR-pMHC engagements ensue and the consequences of such
100 interactions remain unclear.

101 As a result of TCR-pMHC engagement at the immunological synapse, T cells
102 receive a series of activation signals (reviewed in (15)). The TCR activation signals
103 begin with the rapid recruitment and activation of Src kinases, Lck and Fyn (18, 19, 30).
104 The signals propagate downstream leading to the phosphorylation of phospholipase C-
105 $\gamma 1$ (PLC- $\gamma 1$) and the hydrolysis of phosphatidylinositol 4,5-bisphosphate (PIP2). This
106 results in the production of the second messengers diacylglycerol (DAG) and inositol
107 trisphosphate (IP3), the activation of transcription factor NF κ B via PKC θ and the
108 MAPK/Erk pathways, the flux of intracellular Ca²⁺, and the transcription of IL-2 via Ca²⁺-
109 associated calmodulin, calcineurin and NFAT. Of note, the TCR-induced signaling
110 cascade is transient; it is initiated at the newly formed TCR-pMHC microclusters and
111 terminated at the cSMAC (18, 30). CD4 distribution is consistent with these data in that
112 CD4 initially associates with the TCR-pMHC microclusters yet does not accumulate in
113 the cSMAC (31). The lack of active signaling at the cSMAC is also explained by the
114 finding that TCRs associated with micro-vesicles are detached and released out of the
115 cell into the synaptic cavity from the cSMAC (20).

116 A similar but atypical signaling cascade is triggered in the HIV Env-induced
117 virological synapse (29). The HIV Env-CD4 interactions at the synapse lead to the
118 phosphorylation of Lck and the downstream TCR signalosome components CD3 ζ , ZAP-

119 70, LAT, SLP-76, Itk, and PLC- γ 1 (29). These signaling molecules accumulate in the
120 cSMAC-like central Env/CD4 cluster and retain their phosphorylation for more than one
121 hour, even after the synapse breaks and the cell resumes migration. Notably, the
122 protracted signaling does not lead to full T-cell activation as indicated by lack of PKC θ
123 activation and no elevation of Ca²⁺ or CD69. Nonetheless, the Env-induced signals
124 trigger actin polymerization to create an F-actin ring with a depleted central zone
125 implicated to serve as a channel for efficient transport of the HIV virions within the target
126 CD4 T cell (29). These signals also activate LFA-1 on naïve CD4 T cells (32).

127 In this study, we postulate that the aberrant HIV Env-triggered signaling exerts a
128 dominant effect on the TCR-induced activation signals at the immunological synapse to
129 impact the activation state of CD4 T cells targeted by the virus. We utilized a supported
130 lipid bilayer (SLB) system that has been used to monitor the immunological synapses
131 formed by CD4 T cells, CD8 T cells, and NK cells and the signaling molecules recruited
132 to the various synapses (18, 19, 28, 29, 33, 34). To evaluate the specific contribution of
133 the HIV Env in the CD4 T cell immunological synapse, we applied primary activated
134 CD4 T cells onto SLB constituted with OKT3 (an anti-CD3 monoclonal antibody that
135 engages TCR), ICAM-1 (a key adhesion molecule integral to the immunological
136 synapse), and plus or minus HIV Env gp120. Although the HIV Env did not disrupt the
137 immunological synapse formation, its presence at the synapse affected the kinetics of
138 TCR mobilization and the synapse morphology. Further, the HIV Env induced temporal
139 and spatial alterations to the most proximal TCR-associated signaling molecule Lck and
140 augmented CD4 T cell activation and proliferation. These data support the notion that
141 HIV co-opts the immunological synapse to disseminate to highly vulnerable CD4 T cells

142 undergoing TCR-induced activation and promote its replication by heightening the T cell
143 activation state.

144 **Materials and Methods**

145 **Cells.** Peripheral blood mononuclear cells were isolated from leukopacks of
146 anonymous healthy donors (purchased from New York Blood Center) using Ficoll-
147 Hypaque. Approval for the use of human specimens was obtained from the New York
148 University School of Medicine Institutional Research Board. CD4 T cells or CD8 T cells
149 were enriched by using CD4 or CD8 negative-selection magnetic bead kits (STEMCELL
150 Technologies). The T cells were then activated on plates coated with anti-CD3 and anti-
151 CD28 antibodies at 5 $\mu\text{g/ml}$ each. After 48 h, activated cells were transferred onto new
152 plates at a density of 1×10^6 cells/ml and supplemented with 20U/ml interleukin-2. Cells
153 were used in the SLB experiments 5 to 12 days later.

154
155 **Supported lipid bilayers (SLBs).** SLBs were prepared as described (35) with
156 some modifications. Liposomes containing DOPC (Avanti Polar Lipids) supplemented
157 with 12.5 % 1,2-dioleoylsn-glycero-3-[(N-(5-amino-1-carboxypentyl)iminodiacetic acid)-
158 succinyl] nickel salt (Ni-NTA-DOGS) and 0.005% 1,2-dipalmitoyl-sn-glycero-3-
159 phosphoethanolamine-N-(Cap biotinyl) were applied onto clean coverslips placed onto
160 sticky-Slide VI 0.4 chambers (Ibidi). After blocking with 5% casein containing 100 μM
161 NiCl_2 , His₁₂-tagged HIV gp120 SF162 (250/ μm^2) and ICAM-1 (200/ μm^2) were added to
162 bind Ni-NTA-DOGS, while mono-biotinylated OKT3 (30/ μm^2) was attached to Cap-
163 biotinyl. All proteins were also fluorescently labelled prior to addition to the SLBs. The
164 density of HIV gp120 tested on the SLBs was established in our previous studies (28).
165 For comparison, the estimated Env density on the surface of HIV virions is 8 to 14
166 envelope spikes per virion (36, 37), which correspond to 250 to 450/ μm^2 for virions with

167 a diameter of ~100 nm. On the cell surface, the HIV envelope glycoproteins are
168 expressed in patches (38, 39), and the confocal microscopy measurements of the local
169 densities of HIV gp120 at such patches on the HIV envelope-transfected cell surfaces
170 yielded an average density of 369 molecules/ μm^2 (median, 287 molecules/ μm^2 ; range,
171 0-2490 molecules/ μm^2) ((28) and data not shown). The ICAM-1 and OKT3 densities of
172 on the SLB were based on their expression levels on the surfaces of antigen-presenting
173 cells (28). The density of each protein was determined prior to use in the SLBs by
174 coating the same SLB preparations onto 5- μm -diameter silica beads and analyzing the
175 beads by flow cytometry using calibration beads (Bangs Laboratories Inc., IN). The
176 SLBs were then washed, and incubated with 0.5×10^6 cells in 200 μl HEPES-buffered
177 saline containing 1% human serum albumin, 2 mM MgCl_2 , and 1 mM CaCl_2 .

178 Immunofluorescence staining was used to detect total and phosphorylated Lck at
179 the immunological synapse. At the designated time points, cells on SLBs were fixed
180 with 2% paraformaldehyde, permeabilized with 0.1% Triton X-100, and stained with
181 antibodies specific for Lck phosphorylated on Y394 (R&D Systems) or Y505 (BD
182 Transduction Laboratories) or total Lck (BD™ Phosflow), followed with an appropriate
183 Alexa Fluor 546-labeled secondary antibody (Invitrogen,CA).

184 For antibody-blocking experiments, anti-gp120 mAbs obtained from Drs. Susan
185 Zolla-Pazner (Icahn School of Medicine at Mount Sinai) and Mirosław K. Gorny (NYU
186 School of Medicine) were used (40-42). SLBs containing OKT3, ICAM-1 and gp120
187 were treated with 10 $\mu\text{g}/\text{ml}$ of mAb for 30 minutes prior to introduction of the cells and
188 washed three times with the assay buffer to remove excess mAb. The mAbs 654, 2158,
189 and 2558 have potent neutralizing activities against HIV-1 SF162 from which the gp120

190 protein used in this study was derived (IC_{50} titers of <1, 6, <1 $\mu\text{g/ml}$, respectively (43)).

191 The control mAb 1418 is specific for parvovirus and does not neutralize HIV.

192

193 **Total Internal Reflection Fluorescence Microscopy.** T cells were loaded onto
194 SLBs and allowed to interact for approximately 1 hour or for the indicated times. Images
195 were acquired either from live cells or after fixation and staining. Fluorescence
196 microscopy and IRM were performed on a Nikon Ti microscope with a TIRF module and
197 an Andor DU897 back illuminated electron multiplier EMCCD camera. Solid-state
198 lasers provided illumination at 405, 488, 561, and 641 nm and narrow pass filters were
199 used for detection.

200

201 **Image analysis.** Acquisition settings were maintained constant throughout each
202 imaging procedure and between samples. Image analysis was performed with ImageJ.
203 To measure intensities, individual-cell contacts were traced with region of interest (ROI)
204 on the IRM channel. Images were subtracted for background, and then cells in the
205 subtracted images were marked with ROIs.

206 The densities of gp120 were calculated by measuring the fluorescence intensity
207 values from gp120-AF488 channel within each cell and subtracting them with the
208 background values from SLBs with no gp120. The gp120 densities on the
209 OKT3+ICAM-1+gp120 SLBs in the areas with no cell contacts were normalized to 250
210 molecules/ μm^2 . The densities of OKT3 were measured based on the fluorescence
211 intensity values from the OKT3-AF647 channel minus the background values. The
212 OKT3 background values were calculated from the zero time-point and normalized to 30

213 molecules/ μm^2 .

214 Measurement of signaling molecules in cells on the SLBs was done on the entire
215 cell-SLB contact area as detected in the IRM channel (29). Average fluorescence
216 intensity was measured in each of the areas designated above. The background
217 fluorescence was subtracted from the average intensity. The integrated fluorescence
218 intensity was then calculated by multiplying the average intensity by the total pixel area
219 measured in each cell. The pixel area was converted to μm^2 based on the pixel/ μm^2
220 ratio for the specific cameras and objectives that were used.

221

222 **CD69 upregulation and IL-2 secretion.** CD4 T cells were activated with anti-
223 CD3 and anti-28 mAbs as described above and used at least 7 days post-activation.
224 CD4 T cells (0.5×10^6 per condition) were incubated with silica beads of 5 μm in
225 diameter (Bangs Laboratories), the standard bead size for T cell stimulation and
226 expansion (44). The beads were coated with SLBs containing OKT3, ICAM-1, plus or
227 minus CD80 or HIV gp120. After 6 hours, cells were stained with anti-human CD69 PE
228 Abs (eBioscience) and analyzed by FACSCalibur (BD). At 24 hours, supernatants were
229 collected and IL-2 concentrations were measured by using human IL-2 ELISA Kit
230 (Thermo scientific).

231

232 **T cell proliferation and HIV replication.** To measure proliferation, CD4 T cells
233 freshly isolated from peripheral blood were labeled with CFSE, cultured on wells coated
234 with OKT3 plus or minus HIV gp120. Anti-CD4 mAb Leu3A was also tested to block
235 CD4 interaction with HIV gp120. After 5 days, cell proliferation was measured based on

236 CFSE dilution by flow cytometry, and the data were analyzed with FlowJo. These
237 stimulated CD4 T cells were also assessed for the ability to support virus replication.
238 After culture with OKT3, OKT3+HIV gp120, or OKT3+anti-CD28 mAb for 5 days, the
239 cells were infected with HIV-1 SF162 (moi=0.1 as measured in the MT4-TMZ-R5
240 reporter cells), washed and incubated in the presence of the same stimuli for an
241 additional 10 days, without the addition of exogenous cytokines. Virus production was
242 monitored in the supernatants at day 3 and day 10 post-infection by p24 ELISA
243 (XpressBio).

244

245 **Statistics.** The graphs and statistical analyses were performed with Microsoft
246 Excel and Graphpad Prism. Experiments were performed at least three times.

247

248 **Results**

249 **CD4 T cell immunological synapse formation and TCR recruitment to the synapse** 250 **in the presence of HIV Env gp120**

251 In order to study the effect of the HIV Env on the formation of TCR-induced
252 immunological synapse in primary human CD4 T cells, we used the SLB as an
253 experimental model. This system enables the real-time visualization of microclusters
254 and supramolecular activation clusters (SMAC) as they form in the TCR-induced
255 immunological synapse and in the TCR-independent HIV Env gp120-induced virological
256 synapse (18, 19, 28, 29). The SLBs were constituted with the anti-CD3 mAb OKT3,
257 ICAM-1 and HIV Env gp120 to represent the surface of APCs expressing HIV Env
258 following infection. A recombinant monomeric HIV-1 (SF162) gp120 with a His₁₂ tag at
259 its C terminus was labeled with the fluorescent dye Alexa Fluor 488 (AF488-gp120) and
260 attached to the SLB via Ni²⁺-chelating NTA lipids. ICAM-1 was also tagged with His₁₂
261 but was labeled with Alexa Fluor 405. Anti-CD3 mAb OKT3 was mono-biotinylated,
262 labeled with Alexa Fluor 647, and reacted sequentially with streptavidin and the biotin-
263 CAP present in the SLBs. Activated CD4 T cells were then added onto the SLBs and
264 live images were acquired by total internal reflection fluorescence microscopy (TIRFM)
265 for 1 hour (**Fig 1**). The previously activated CD4 T cells were used to allow the
266 evaluation of a relatively homogenous population of primary target cells that are
267 permissive to HIV infection and form well-defined immunological synapses on SLBs (18,
268 19, 28, 29).

269 Within 10 minutes after the CD4 T cells had interacted with OKT3 and ICAM-1 on
270 the SLBs, typical immunological synapses displaying the TCR-rich cSMAC and the

271 ICAM-1 pSMAC rings were assembled in the absence of the HIV Env (**Fig 1A, top**
272 **panel**). In the presence of HIV gp120 on the SLBs, the cells also formed synapses with
273 segregated cSMAC and pSMAC (**Fig 1A, bottom panel**). These synapses were similar
274 to those formed on SLBs without gp120. Notably, HIV gp120 accumulated in the center
275 of the synapse and coalesced together with TCR at the cSMAC (**Fig 1A**). The relative
276 distribution of HIV gp120 and TCR in the synapse was quantified by determining the
277 densities of gp120 and OKT3 along the cell radius. Data were acquired at the 10-
278 minute time point from a minimum of 30 cells per condition. These data confirmed the
279 visual observation in **Fig 1A** and demonstrated an increase of gp120 density from an
280 initial density of 250 molecules/ μm^2 on the SLBs to 400 molecules/ μm^2 at the center of
281 the cell (**Fig 1B left**). Measurements also showed an increase in OKT3 density towards
282 the center of the synapse and an accumulation of OKT3 at the cSMAC that was ~3x fold
283 higher ($p < 0.0001$) in the presence of HIV gp120 (**Fig 1B right**), indicating that HIV
284 gp120 may help augment TCR recruitment and clustering at the cSMAC.

285 For comparison, we also evaluated the effect of HIV gp120 on the immunological
286 synapse formed by CD8 T cells. Activated CD8 T cells were applied to the SLBs
287 containing OKT3, ICAM-1 and plus or minus gp120. The CD8 T cells formed
288 immunological synapses with the typical bull's eye appearance comprising of the TCR-
289 rich cSMAC and the ICAM-1 pSMAC ring on the SLBs with or without gp120 (**Fig 1C**
290 **upper panel and lower panel**). However, little gp120 was recruited to the centers of
291 the immunological synapses formed by the CD8 T cells (**Fig 1C lower panel and Fig**
292 **1D left**). The OKT3 accumulation at the c-SMAC was also not significantly increased
293 (**Fig 1D right**). These results indicate that HIV gp120 binds primarily to CD4 T cells and

294 augments TCR accumulation in the CD4 T cell immunological synapse.

295 To further examine the alteration of TCR recruitment by HIV gp120, we
296 compared the kinetics of OKT3 signal accumulation in the synapse over time in the
297 presence versus the absence of gp120. CD4 T cells were introduced to the SLBs
298 bearing OKT3, ICAM-1, and gp120 or with the SLBs with only OKT3 and ICAM-1, and
299 the OKT3 signals were measured for 50 minutes. In the absence of gp120, OKT3
300 signals increased to a plateau of approximately 60-70 molecules/ μm^2 at 7 minutes (**Fig**
301 **2A-B** (also see **Movie S1** for video images). This plateau was then maintained for up to
302 50 minutes. In contrast, when HIV gp120 was present, OKT3 signals increased more
303 dramatically, peaking at 150 molecules/ μm^2 at 7 minutes, but subsequently declined
304 over time (**Fig 2A-B and Movie S2**).

305 On the SLBs with OKT3, ICAM-1 and gp120, the kinetics of gp120 accumulation
306 was also evaluated. The HIV gp120 density increased to 400 molecules/ μm^2 in
307 approximately 10 minutes and continued to rise at a lower rate (**Fig 2C and Movie S2**).
308 The gp120 signals fluctuated during the 50-minute observation time, but they did not
309 show a steady decline as seen with OKT3. In fact, from 20 to 50 minutes the gp120
310 signals became more prominent than the OKT3 signals. Hence, the pattern of TCR
311 accumulation did not follow that of gp120. Taken together with the data in **Fig 1**, these
312 data demonstrate that the presence of HIV gp120 in the CD4 T cell immunological
313 synapse HIV gp120 does not prevent the synapse formation. However, the HIV gp120
314 is recruited to the cSMAC along with TCR, albeit with distinct patterns, and causes a
315 dramatic change in the kinetics of TCR recruitment and retention in the synapse.

316

317 **Effects of anti-gp120 antibodies on the recruitment of HIV gp120 and TCR to the**
318 **synapse**

319 To determine the interactions involved in HIV gp120 recruitment to the CD4 T-
320 cell immunological synapse, we pre-treated SLBs containing gp120, OKT3 and ICAM-1
321 with human mAbs directed to different regions of gp120. The CD4 T cells were then
322 added and allowed to interact with the SLBs. After 20 minutes, cells were fixed and HIV
323 gp120 densities in the synapses were measured. The results showed that mAbs
324 against the CD4-binding site (654), the V1V2 domain (2158) and the V3 loops (2558)
325 significantly reduced the densities of HIV gp120 at the synapses, while the irrelevant
326 mAb control (1418) did not have this effect (**Fig 3A**). Representative images of the
327 individual cells further showed that the anti-gp120 mAb treatment decreased the
328 amounts of HIV gp120 at the synapses but did not block clustering (**Fig 3C**). The
329 location of the HIV gp120 clusters near the TCR-rich cSMAC also remained unchanged.
330 Notably, low levels of gp120 interactions remained after treatment with the mAbs,
331 including mAb 654 that blocks CD4-gp120 binding. These findings are similar to our
332 past study in which the mAb against the CD4-binding site also did not completely
333 abrogate the CD4 T cell interactions with HIV gp120 on the SLBs presenting only gp120
334 and ICAM-1 (28). The data indicate that, in addition to CD4, other receptors are also
335 engaged by HIV gp120 at these synapses and these interactions cannot be completely
336 blocked by a single anti-gp120 mAb. Nonetheless, the non-CD4 receptors and their
337 relative contributions are yet to be delineated.

338 Measurements of OKT3 density showed higher levels of OKT3 at the synapses
339 in the presence of HIV gp120 (**Fig 3B**), which is consistent with data in **Fig 1 and 2**.

340 Pre-treatment with the anti-gp120 mAbs did not significantly affect the enhanced OKT3
341 accumulation. The OKT3 densities were slightly reduced by mAbs against the CD4-
342 binding site (654) and the V3 loop (2558), but the decrease did not reach statistical
343 significance. Hence, mAbs directed to the different sites of gp120 reduced HIV gp120
344 recruitment to the synapse without significantly curtailing the TCR accumulation. These
345 data suggest that the presence of very low amounts of HIV gp120 in the synapse is
346 sufficient for the increased OKT3 recruitment.

347

348 **Effects of HIV gp120 on the synapse morphology over time**

349 The ability of HIV gp120 to alter the kinetics of TCR accumulation and retention
350 in the synapse suggests that HIV gp120 may affect the synapse morphology over time.
351 To evaluate this idea, we monitored the synapses 10 and 30 minutes after CD4 T cells
352 made contact with SLBs presenting HIV gp120, OKT3, and ICAM-1 as compared to the
353 synapses on control SLBs containing only OKT3 and ICAM-1. On the control SLBs,
354 70% of the CD4 T cells formed the so-called mature immunological synapses within 10
355 minutes. These cells form stable contacts with the SLBs and the synapses displayed
356 the hallmark central OKT3 cluster (cSMAC) in the center surrounded by a symmetrical
357 peripheral ICAM-1 ring (pSMAC) (**Fig 4A, OKT3+ICAM-1: symmetrical**). The ICAM-1
358 ring keeps the migratory T cells from crawling across the SLBs (17, 28). These
359 synapses were maintained throughout the observation period; after 30 minutes the
360 percentage of the cells with mature synapses remained at 75% (**Fig 4B**). The
361 percentages of cells with off-center TCR clusters and asymmetrical pSMAC structures
362 also stayed at 25-30% throughout the observation period (**Fig 4A, OKT3+ICAM-1:**

363 **asymmetrical and Fig 4B).**

364 In contrast, on the SLBs with HIV gp120, OKT3 and ICAM-1, only 17% of the
365 cells displayed mature synapses with central TCR clusters at 30 minutes, although at 10
366 minutes the proportion was comparable to that in the absence of gp120 (**Fig 4A,**
367 **gp120+OKT3+ICAM-1: symmetrical and Fig. 4B).** Hence, after 30 minutes, the
368 number of cells with mature synapses on the SLBs with gp120 declined. Most of the
369 cells became elongated with off-center TCR clusters and no longer had well-defined
370 ICAM-1 rings (**Fig 4A, gp120+OKT3+ICAM-1: asymmetrical**), indicative of cells that
371 had resumed migration. These data demonstrate that the presence of HIV gp120 not
372 only alters TCR accumulation but also affects the propensity of the cells to migrate after
373 forming the synapses.

375 **Sustained Lck phosphorylation in the synapse in the presence of HIV gp120**

376 Given the ability of HIV gp120 to enhance TCR accumulation in the synapse, we
377 next evaluated whether HIV gp120 affected the activation of TCR-proximal signaling.
378 Since Lck activation is the first signal triggered upon TCR engagement and by gp120-
379 CD4 binding (18, 29, 45, 46), we performed immunostaining of CD4 T cells interacting
380 with OKT3 and ICAM-1 plus or minus gp120 on the SLBs with anti-Lck antibodies. The
381 cells were added to the SLBs for 10 or 45 minutes, fixed, permeabilized, and stained
382 with antibodies specific for total Lck, pLck (Y505), and pLck (Y394) (**Fig 5**). TIRFM was
383 then used to detect Lck phosphorylation and its recruitment to the synapses. TIRFM
384 allows the detection of fluorescence within a 100-200 nm slice and thus provides high
385 lateral and axial resolution of Lck or pLck signals recruited specifically to the membrane-

386 proximal areas at the T cell-bilayer interface (<200 nm of the T cell plasma membranes
387 at the cell-bilayer contact sites). The fluorescence signals beyond this thin area are
388 excluded from the measurements. The data from individual cells from one of the donors
389 are shown in **Fig 5A**, while the cumulative data from three different donors tested in
390 separate experiments are presented in **Fig 5B**.

391 In the cells on the control OKT3+ICAM-1 SLBs with no gp120, total Lck levels at
392 the synapses were comparable at 10 minutes and 45 minutes (**Fig 5A-B**). In the
393 presence of HIV gp120 a slightly higher level of total Lck staining was observed at 10
394 minutes, but overall the total Lck staining remained relatively constant over time in the
395 presence or absence of HIV gp120. In contrast, phosphorylated Lck staining (both
396 pY505 and pY394) showed a declining trend from 10 minutes to 45 minutes in the cells
397 on the control SLBs. This is consistent with the fact that the physiologic Lck activation
398 triggered via TCR is transient (18, 19, 46). In the presence of HIV gp120, however,
399 pLck(Y505) staining showed an altered pattern: it increased from 10 minutes to 45
400 minutes such that the pLck(Y505) staining at 45 minutes was higher in the presence vs.
401 absence of HIV gp120. pLck (Y394) staining did not rise from 10 minutes to 45 minutes
402 but was sustained at higher levels at both time points in the presence of HIV gp120.
403 **Fig 5C (middle and right panels)** further shows that, in the presence of HIV gp120,
404 most pLck staining accumulated at the cSMAC after 45 minutes and localized near
405 gp120 and OKT3. This pattern was not seen with the total Lck, which remained
406 dispersed as small puncta throughout the cell-bilayer contact areas (**Fig 5C left panel**),
407 indicating the vast majority of Lck molecules were not phosphorylated and were not
408 recruited to the membrane-proximal area of the gp120-containing cSMAC. In the

409 absence of gp120, pLck and total Lck also did not converge to the cSMAC or co-localize
410 with OKT3, consistent with past data (29).

411 These data indicate that, unlike the transient Lck activation induced by TCR in
412 the peripheral areas of the immunological synapse, the presence of HIV Env gp120 in
413 the synapse triggers a sustained level of phosphorylated Lck which accumulates in the
414 synapse along with gp120 and OKT3 for up to 45 minutes. This pattern is reminiscent
415 of the long-lasting Lck activation induced by HIV gp120 alone (29) and suggests the
416 dominant effects of HIV gp120 over TCR on Lck activation in the CD4 T cell
417 immunological synapse.

418

419 **HIV gp120-mediated enhancement of CD4 T cell activation and virus replication**

420 The ability of HIV gp120 to augment TCR-induced Lck activation led us to
421 investigate the downstream events associated with T cell activation. By itself HIV
422 gp120 only partially activates the early TCR signaling machinery on CD4 T cells, and
423 the signals do not propagate to result in intracellular Ca^{2+} elevation or increased CD69
424 expression (29). However, the presence of HIV gp120 during TCR engagement may
425 alter CD4 T cell activation. To test this idea, we compared the extent of CD69
426 upregulation, IL-2 secretion, and proliferation when CD4 T cells were triggered by OKT3
427 or by a combination of OKT3 and HIV gp120. To accommodate for the prolonged assay
428 periods required to detect CD69 upregulation and IL-2 secretion beyond the one hour
429 period for the TIRFM experiments above, SLBs were formed on the standard 5- μm
430 sterile silica beads and functionalized with OKT3, ICAM-1, and gp120 or no gp120,
431 which were then incubated with CD4 T cells for up to 24 hours. SLB-beads coated with

432 OKT3, ICAM-1, and CD80 (a co-stimulatory molecule binding to CD28) or with only
433 ICAM-1 were also tested for comparison. The beads have been shown to provide
434 equivalent stimulation to T cells as the planar surfaces (47). The cells were stained with
435 anti-CD69 mAb at 6 hours and 24 hours and analyzed by flow cytometry. The
436 supernatants were collected at 24 hours and tested for IL-2 in ELISA. Untreated cells
437 served as background control. T cell proliferation was also assessed based on CFSE
438 dilution 5 days after the cells were cultured on wells coated with OKT3 with or without
439 gp120.

440 On the CD4 T cells treated with OKT3 + ICAM1 beads, CD69 expression
441 increased but declined rapidly (**Fig 6A**). Stimulation with OKT3 + ICAM-1 beads also
442 induced no IL-2 production (**Fig 6B**). With the addition of HIV gp120, CD69
443 upregulation was augmented and sustained over the 24-hour observation period (**Fig**
444 **6A**). These results were comparable to those achieved with an OKT3 + CD80 co-
445 stimulation. HIV gp120 also significantly increased IL-2 production to the levels
446 observed with an OKT3 + CD80 co-stimulation (**Fig 6B**). Cells treated with ICAM-1-
447 beads showed a similar response to untreated cells and did not upregulate CD69
448 expression or secrete IL-2. For CD4 T cell proliferation assay (**Fig 6C**), CFSE-labeled
449 CD4 T cells were cultured for 5 days in wells coated with OKT3 plus or minus HIV
450 gp120 (strains SF162 or CH040). Enhanced proliferation was observed with CD4 T
451 cells stimulated with OKT3 plus HIV gp120 as compared with OKT3 alone, and the
452 enhanced response was inhibited by the Leu3A mAb that blocked HIV gp120-CD4
453 binding. These results indicate that HIV Env gp120 enhanced CD4 T cell activation to a
454 similar extent as the bona fide co-stimulatory ligand.

455 To determine whether the CD4 T cells stimulated by OKT3 in the presence of
456 HIV gp120 can support virus infection better than those activated by OKT3 alone, we
457 treated the activated cells with HIV-1 SF162 and measured virus replication based on
458 virus p24 in the culture supernatants. For comparison, p24 levels were also measured
459 in the supernatants of unstimulated cells that supported minimal virus replication.
460 These experiments were done without addition of exogenous cytokines. The data show
461 that the amounts of p24 produced by the stimulated cells increased from day 3 to day
462 10 post-infection, indicative of active virus replication, although the concentrations
463 varied among the different donors (**Fig 6D**). Notably, higher levels of p24 were
464 produced by infected CD4 T cells that had been stimulated with OKT3 in the presence
465 of HIV gp120 than in those activated only with OKT3. The greater p24 production
466 induced by OKT3 + HIV gp120 was detected as early as day 3 and became more
467 evident at day 10. About 2-5 fold enhancement was attained in the absence of any
468 exogenous cytokines and consistently observed with cells from the different donors.
469 Importantly, virus replication after stimulation with OKT3 + HIV gp120 was comparable
470 to or even higher than that seen in CD4 T cells stimulated with OKT3 and anti-CD28
471 mAb, indicating the HIV gp120's potent co-stimulatory activity. All together these results
472 show that in the CD4 T cell immunological synapse HIV Env may act as a co-stimulator
473 that heightens TCR activation of the CD4 T cells and imparts the cells with the capacity
474 to support higher levels of virus replication.
475

476 **Discussion**

477 This study demonstrated the ability of HIV Env gp120 to alter the kinetics of TCR
478 recruitment and the activation of TCR-proximal signaling at the CD4 T cell
479 immunological synapse. While most earlier studies examine the changes in the
480 immunological synapse formed by the CD4 T cells already infected with the virus (48-
481 50), this study evaluated the HIV Env interaction with target CD4 T cells in the
482 immunological synapse prior to infection. Notably, the presence of the HIV Env did not
483 interfere with the immunological synapse assembly. TCR accumulation at the cSMAC
484 of the synapse and its segregation from the adhesive pSMAC ring of ICAM-1/LFA-1
485 interactions proceeded similarly in the presence or absence of the HIV Env. The
486 proportion of cells forming the synapse was also comparable. However, the binding of
487 HIV Env to CD4 T cells resulted in the accumulation of HIV Env and phosphorylated Lck
488 at the cSMAC, although the cSMAC is normally devoid of Lck in the absence of the HIV
489 Env (18, 19, 31). Importantly, the HIV Env accumulation at the synapse enhanced
490 TCR-induced activation that predisposed the target CD4 T cells to become more
491 receptive to HIV infection and productive replication. It should be noted that this activity
492 was observed with the HIV Env gp120 monomers that cannot mediate virus infection.
493 However, these monomeric gp120 proteins have receptor-binding activities. Moreover,
494 when bound on membranes like the SLBs, the HIV Env gp120 monomers were able to
495 cross-link and cluster the HIV Env-receptor interactions to the cSMAC to result in Lck
496 activation at the cSMAC. Past studies also have reported the activation of Lck and the
497 downstream signals upon CD4 engagement by the HIV virions or the HIV Env
498 expressed on cells (51-53). On the surfaces of virions and infected cells,

499 heterogeneous species of the HIV Env glycoproteins are expressed (54-56), including
500 non-functional Env proteins that cannot partake in the virus entry process but may have
501 immunomodulatory activities demonstrated in this study.

502 The exact mechanism by which the HIV Env at the immunological synapse
503 augments TCR-induced activation of CD4 T cells is yet to be delineated. However, the
504 temporal pattern of TCR recruitment was significantly different in the presence of the
505 HIV Env. Thereby, in the presence of the HIV Env the TCR density in the synapse
506 reached a higher peak (~3 fold higher than control) and declined over time. In contrast,
507 in the absence of the HIV Env, the TCR level increased to a plateau that was
508 maintained throughout the one hour observation time. The alteration of TCR
509 mobilization was induced by the HIV Env binding to CD4 T cells. However, treatment
510 with the anti-Env mAbs directed to the CD4-binding site, the V1V2 domain, or the V3
511 loop reduced the HIV Env accumulation in the CD4 T cell synapses, without blocking
512 the enhanced TCR recruitment to the synapses. These data suggest the potential
513 involvement of multiple receptors in affecting the TCR recruitment and that a single anti-
514 Env mAb is not sufficient to block this activity. We also observed that none of the
515 individual anti-Env mAbs, including the 654 mAb efficiently blocking CD4-gp120
516 interaction, was able to totally prevent the HIV Env recruitment to the CD4 T cell
517 synapses. Moreover, low levels of HIV Env were found to interact with the CD8 T cells
518 on the SLBs, albeit without affecting the TCR recruitment. Hence, in addition to CD4,
519 the other HIV Env receptors such as the integrin $\alpha 4 \beta 7$ and the mannose receptors are
520 likely to be engaged at these synapses. Further, given that a partial inhibition of the
521 Env recruitment was also seen with a mAb against the V3 loop known to be important

522 for the chemokine receptor binding (57), these co-receptors may also play a role in the
523 synapses. Nonetheless, the relative contributions of the different non-CD4 receptors
524 remain to be determined.

525 The activity of the HIV Env to enhance TCR-induced T cell activation is rather
526 unique and is in contrast to the inhibitory effects seen more commonly with other virus
527 proteins. For example, the presence of HERV Env on dendritic cells reduces the ability
528 of the dendritic cells to form immunological synapses with T cells resulting in blockage
529 of T cell activation (58). Similarly, the Nef proteins of most SIV and HIV-2 have been
530 shown to downregulate TCR and CD3 expression in virus-infected T cells resulting in
531 the suppression of immunological synapse formation with antigen presenting cells (48).
532 In contrast, the HIV-1 Nef expression in infected CD4 T cells does not disrupt the
533 formation of immunological synapses between the infected CD4 T cells and APCs and
534 instead affects the synapse functions by inhibiting TCR-induced actin polymerization
535 and TCR-proximal signaling (48, 59). The HCMV pUL125 protein also reduces the
536 efficiency of the immunological synapse formation by modifying the actin cytoskeleton
537 required for the formation of immunological synapses with NK cells (60). Alternatively,
538 the hRSV nucleoprotein interferes with the assembly of the dendritic cell-CD4 T cell
539 immunological synapses by decreasing TCR-pMHC binding and TCR-proximal
540 signaling (61). The ability of these viruses to directly tamper with the immunological
541 synapse and suppress immune cell activation serves as an escape mechanism from
542 immune surveillance. By contrast, HIV via its Env manipulates the immunological
543 synapse to optimize the activation state of the uninfected CD4 T cells to increase the
544 capacity of these cells to support virus replication.

545 Although the significance of the HIV Env-induced alterations affecting the kinetics
546 of TCR recruitment and retention in the immunological synapse is not fully understood,
547 the data from this study show that the TCR accumulation at the cSMAC was
548 accompanied by HIV Env recruitment to the same central area of the synapse. In our
549 earlier studies, HIV Env-CD4 interactions were also found to be mobilized to the
550 synapse center to form a cSMAC-like structure independent of TCR engagement (28,
551 29). Importantly, the central Env-CD4 clusters were associated with long-lasting but
552 partial activation of membrane-proximal signals that began with Lck phosphorylation
553 (29). The present study also showed a similar pattern of durable Lck activation in the
554 Env-containing TCR-induced immunological synapse. These observations indicate the
555 dominant effect of the Env over the transient TCR-induced signaling.

556 The presence of HIV Env also enhanced TCR-induced CD4 T cell activation, as
557 measured by CD69 upregulation and IL-2 secretion, and induced a greater level of CD4
558 T cell proliferation. These results are in concordance with a previous report by
559 Zimmermann et al. (62) showing that a simultaneous engagement of TCR by anti-CD3
560 antibody and CD4 by the HIV Env boosted CD4 T cell activation. In contrast, the
561 interaction of CD4 T cells with soluble HIV gp120 retarded TCR-induced proliferation.
562 HIV Env-CD4 interactions that occurred separately prior to TCR engagement were also
563 found to undermine TCR-proximal signaling and downstream activation events (62).
564 Similarly, the pre-engagement of CD4 by HIV followed by treatment with anti-
565 CD3/CD28-coated beads resulted in the inhibition of Lck and F-actin recruitment to the
566 T cell immunological synapses (49). Taken together, these studies present evidence for
567 the multifaceted role of the HIV Env in shaping the activation state and functionality of

568 the host CD4 T cells. It is also important to note that such HIV Env-mediated immune-
569 modulation may play a part not only in the immune dysregulation seen during HIV
570 infection, but also in the inadequate elicitation and maturation of effector and memory
571 immune responses to many HIV Env-based vaccine candidates evaluated thus far (63-
572 65).

573 This study demonstrated the contribution of the HIV Env as would be expressed
574 on virus-bearing APCs in augmenting the TCR-induced activation of cognate CD4 T
575 cells that are targeted by the virus. We also observed that the HIV Env reduced the
576 duration of the otherwise long-lasting immunological synapse, consistent with the
577 transient nature of T cell migration arrest induced by the HIV Env reported previously
578 (28). A transient synapse may offer advantages for the virus. After a brief migration
579 arrest to allow cell-cell contact and virus transfer, a subsequent rapid detachment would
580 prevent cell-cell fusion that terminates the virus life cycle while promoting the spread of
581 the newly infected cells to other sites in the lymphoid tissues. Overall, the data
582 presented herein offers a model in which HIV, via its Env glycoproteins, alters the
583 immunologic synapse and the activation state of CD4 T cells undergoing TCR activation
584 in a way that is likely to facilitate viral replication and spread.

585

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597

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800

801 **Figure legends**

802 **Fig 1: Recruitment of HIV Env gp120 and TCR to the CD4 and CD8 T cell**

803 **immunological synapses. (A-B)** Activated CD4 T cells were introduced to SLBs
804 containing fluorescence-tagged anti-CD3 antibody OKT3, ICAM-1, and gp120 or no
805 gp120. Images of the cell contact area (as measured by interference reflection
806 microscopy or IRM) and fluorescence signals were collected for one hour. **(A)**
807 Representative images from the 10-minute time point are shown. Merge: combined
808 fluorescence signals from OKT3 and gp120. White bar: 5 μ m. **(B)** The distribution of
809 gp120 (left) and TCR (right) in the CD4 T cell synapse at the 10-minute time point. The
810 densities of gp120 and OKT3 were measured radially from the center of each of 30 cells
811 per experiment (white circle in panel A merge, top). Data from representative cells are
812 shown in the top right and left panels. The averages and standard errors from
813 experiments performed independently with cells from three different donors (>30 cells
814 per experiment) are shown in the bottom right and left panels. p values by the Mann
815 Whitney test. **(C-D)** Activated CD8 T cells were added onto SLBs presenting anti-CD3
816 mAb OKT3, ICAM-1, and gp120 or no gp120. Live images were acquired for one hour.
817 Data from the 10-minute time point were analyzed. **(C)** Images from representative
818 CD8 T cells on SLBs with or without gp120. IRM: interference reflection microscopy.
819 Merge: combined fluorescence signals from gp120 and OKT3. **(D)** The densities of
820 gp120 (left) and OKT3 (right) were measured along the cell radius as above. Data from
821 representative cells are shown in the top right and left panels. The averages and
822 standard errors from three different donors were similarly calculated and shown in the
823 bottom right and left panels. ns: not significant by the Mann Whitney test.

824

825 **Fig 2: Kinetics of TCR accumulation in the CD4 T cell immunological synapse in**
826 **the presence or absence of HIV Env gp120.** Activated CD4 T cells were introduced
827 to SLBs containing OKT3, ICAM-1, and plus or minus gp120. Live images were
828 collected for one hour. **(A)** Representative images from different time points during the
829 one-hour observation. Time points (green numbers) are in minutes. IRM: interference
830 reflection microscopy (contact area). Merge: combined fluorescence signals from
831 gp120 and OKT3. White bar: 5 μ m. **(B)** The kinetics of OKT3 recruitment into the
832 synapse in the cells interacting with gp120 or no gp120 on the SLBs. The OKT3
833 densities were analyzed for ~30 individual cells per condition for up to 50 minutes. The
834 averages and standard deviations from all cells analyzed are shown in the left panel,
835 while data from randomly selected individual cells are shown in the right panel. The
836 data from two donors are presented. Statistical analysis was done with two-tailed non-
837 parametric Wilcoxon test comparing OKT3 recruitment in the presence versus absence
838 of HIV gp120. **(C)** The recruitment of gp120 to the synapse over time in cells interacting
839 with gp120, OKT3, and ICAM-1 on the SLBs. The gp120 densities were analyzed as
840 performed for OKT3. The left panel shows the averages and standard deviations from
841 ~30 cells and the right panel shows the patterns displayed by representative individual
842 cells. The data from two donors tested in independent experiments are shown.

843

844 **Fig 3: Reduction of HIV Env gp120 versus TCR accumulation by human anti-**
845 **gp120 mAbs.** Activated human CD4 T cells were introduced into SLBs with gp120, anti-
846 CD3 mAb OKT3 and ICAM-1 pre-treated with anti-gp120 mAbs that target the CD4-

847 binding site (654), V2 (2158), and V3 (2558), or an irrelevant mAb (1418). The mAbs
848 were used at 10 μ g/ml, a saturating concentration for binding gp120 on the SLBs (28).
849 Cells were fixed at 20 minutes and images were collected. At least 30 cells were
850 analyzed for each condition. Statistical analysis was performed with one-way ANOVA.
851 Data from one of three independent experiments are shown. **(A)** The densities of
852 gp120 in the cell contact areas. ****, $p < 0.0001$; ***, $p < 0.001$; ns: not significant. Red
853 bars: mean values. **(B)** The densities of OKT3 in the cell contact areas. ***, $p < 0.001$
854 as compared to gp120 + OKT3 + ICAM plus or minus mAb; ns: not significant. Red
855 bars: mean values. **(C)** Representative images to show the distribution of gp120 and
856 OKT3 with or without anti-gp120 mAb. IRM: interference reflection microscopy to
857 determine the cell contact area. Merge: combined fluorescence signals from gp120 and
858 OKT3.

859
860 **Fig 4: Alteration of CD4 T cell synapse morphology over time in the presence of**
861 **HIV Env gp120.** Activated CD4 T cells were introduced to SLBs containing OKT3,
862 ICAM-1, and gp120 or no gp120 and monitored over time for changes in the synapse
863 morphology. **(A).** Representative images showing cells forming symmetrical versus
864 asymmetrical synapses as determined by the OKT3 cluster position relative to the
865 center of the cell. IRM: interference reflection microscopy. BF: bright field. Merge:
866 merged images from gp120, OKT3 and BF. White bar: 5 μ m. **(B)** The percentages of
867 cells forming symmetrical or asymmetrical synapses at 10-minute and 30-minute time
868 points on SLBs with or without gp120. The averages (shown by bar graphs) from
869 independent experiments performed with cells from five different donors (depicted by

the scatter plots) are shown. *, $p < 0.05$ by one-way ANOVA with Dunn's multiple comparison.

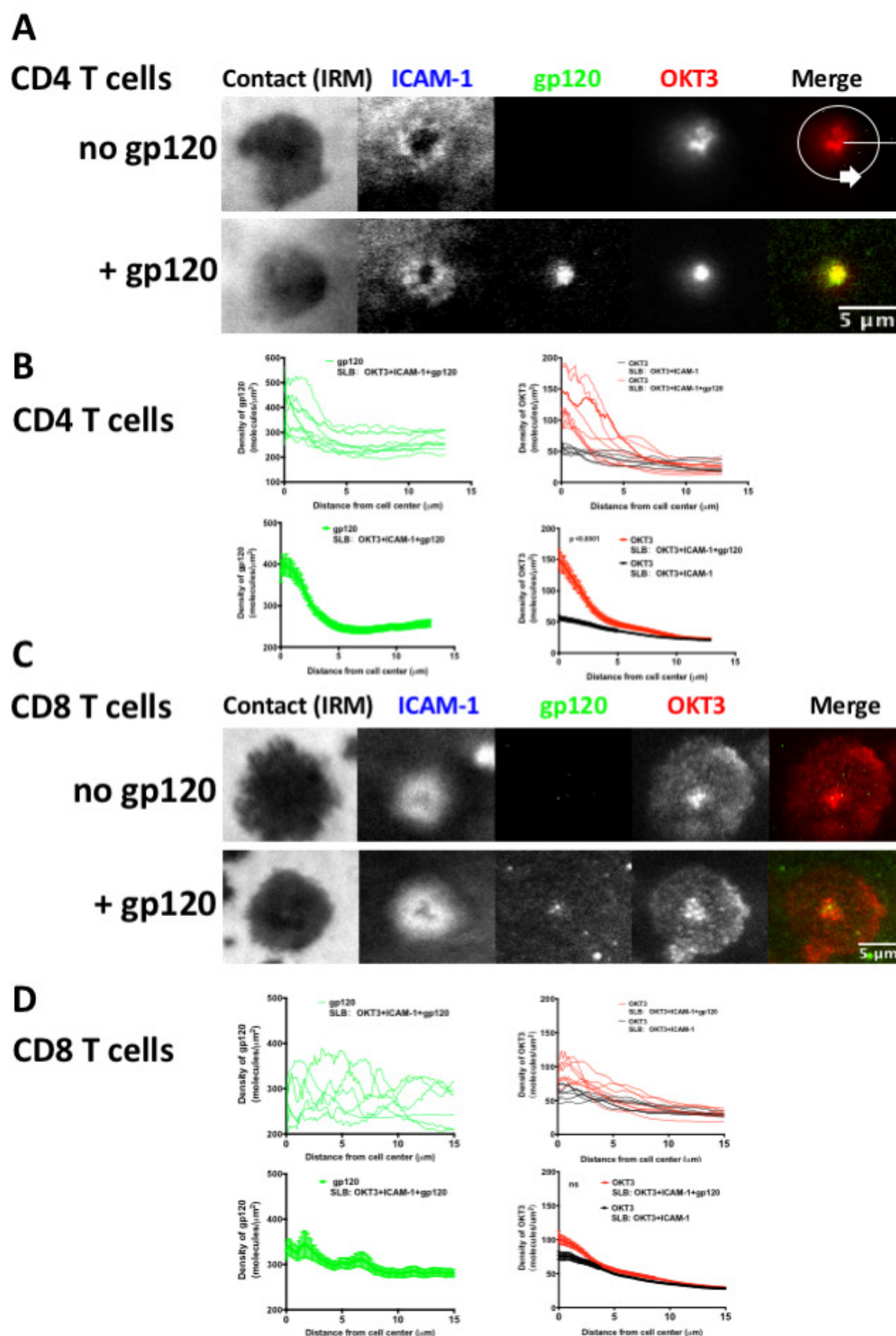
Fig 5: Altered Lck activation in CD4 T cell immunological synapse in the presence of HIV Env gp120. CD4 T cells were introduced onto SLBs containing anti-CD3 mAb OKT3, ICAM-1, and gp120 or no gp120. After 10 minutes or 45 minutes, the cells were fixed and stained for total Lck, pLck (Y505), and pLck (Y394). The integrated fluorescence intensities detected by TIRFM was measured. **(A)** A total of 30 to 50 cells were analyzed for each condition and the data from one representative donor are shown. **(B)** The average integrated intensities from independent experiments performed with cells from three different donors are shown. ****, $p < 0.0001$, ***, $p < 0.001$, **, $p < 0.01$, * < 0.05 , ns: not significant. Red bars: mean values. Iso Ctrl: staining with isotype control. Total and phosphorylated Lck staining at 10 minutes and 45 minutes in the presence or absence of gp120 was significantly higher than isotype control ($p > 0.05$). Statistical analysis was performed with two-tailed *t*-tests. **(C)** Images of representative cells on SLBs with gp120 or without gp120 to show the distribution of total Lck and phosphorylated Lck at the two time points. IRM: interference reflection microscopy. Merge: merged gp120, OKT3, and Lck signals.

Fig. 6: Enhanced TCR-induced activation in the presence of HIV Env gp120. CD4 T cells were incubated with silica beads coated with the SLBs containing OKT3, ICAM-1, and gp120 or no gp120. Cells were also treated with beads coated with SLBs containing OKT3, ICAM-1 and CD80 or ICAM-1 alone for comparison. Untreated cells

893 were used to establish the background levels. CD4 T cells were initially activated on
894 plates coated with anti-CD3 and anti-28 antibodies, and tested >7 days after activation.
895 **(A)** After 6 and 24 hours of incubation with the SLB-coated beads, the cells were fixed,
896 stained with anti-CD69 mAb, and analyzed by flow cytometry. **(B)** The culture
897 supernatants were also collected after 24 hours of incubation and tested in ELISA for IL-
898 2 secretion. The averages and standard errors shown in the graphs are from three
899 experiments performed independently with cells from different donors. **(C)** CFSE dilution
900 assay on CD4 T cells cultured for 5 days in wells coated with OKT3 plus or minus HIV
901 gp120 (strains SF162 or CH040). CD4 T cells freshly isolated from PBMCs were used
902 in the experiments. Anti-CD4 mAb Leu3a blocking CD4-gp120 binding was used to
903 affirm the gp120-mediated enhancement. Histograms from one representative
904 experiment are shown. Division indexes were calculated from the histograms with
905 FlowJo; the means and standard errors from 2-5 independent experiments with cells
906 from different donors are shown. Statistical analysis was done with the Mann Whitney
907 test. **(D)** Virus production in cultures of CD4 T cells grown in wells coated with OKT3,
908 OKT3+HIV gp120, or OKT3+anti-CD28 mAb for 5 days and infected with HIV-1 SF162.
909 CD4 T cells freshly isolated from PBMCs were used in the experiments. Virus p24
910 concentrations in the supernatants were measured by ELISA on day 3 and day 10 post-
911 infection. Data from the different donors are shown. Bkgrd: p24 concentrations in the
912 supernatants of unstimulated cells. ****, $p < 0.0001$, **, $p < 0.01$, * < 0.05 , ns: not
913 significant.

914

Fig. 1



A **no gp120** **OKT3** **IRM** **Merge**



Fig. 3

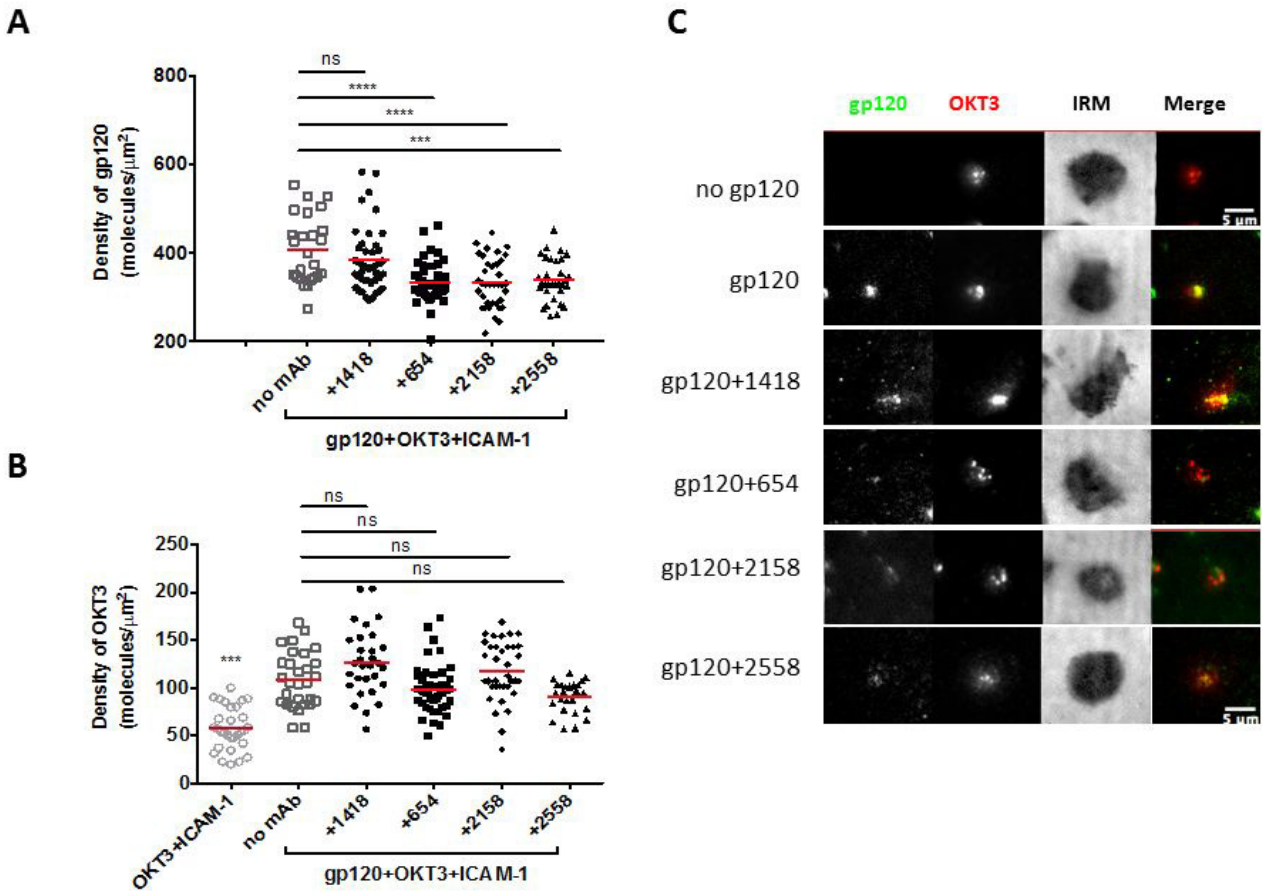


Fig. 4

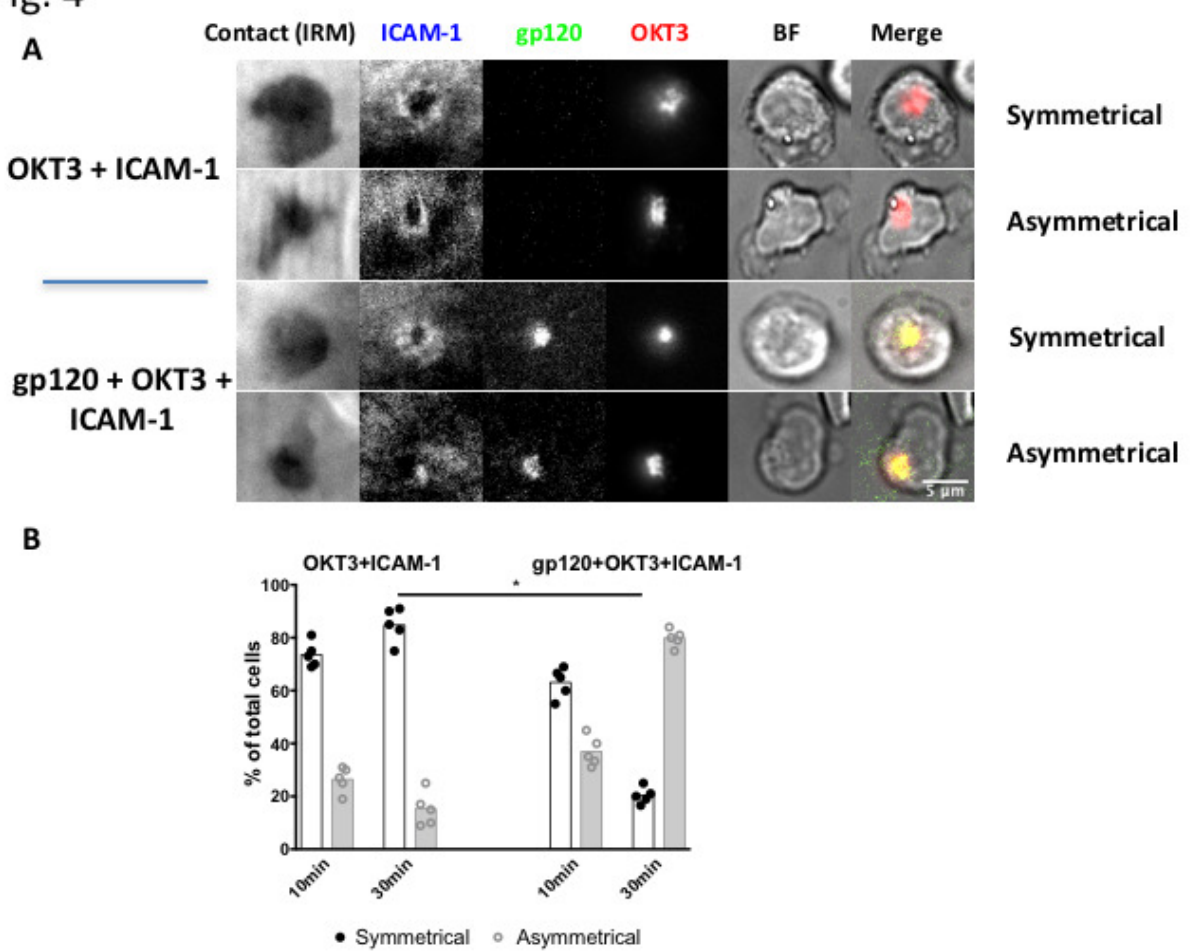


Fig. 6

