

1 **Supplementary Materials**

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3 **Supplemental Acknowledgements**

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24 **Supplemental Figures**

25 Supplemental Figure 1. ATI study design.

26 Supplemental Figure 2. Comparison of pDC frequency with pDC absolute count and between
27 treatment arms.

28 Supplemental Figure 3. Changes in mDC and monocyte frequencies after ATI.

29 Supplemental Figure 4. Comparison of surface marker expression between arms of RV397.

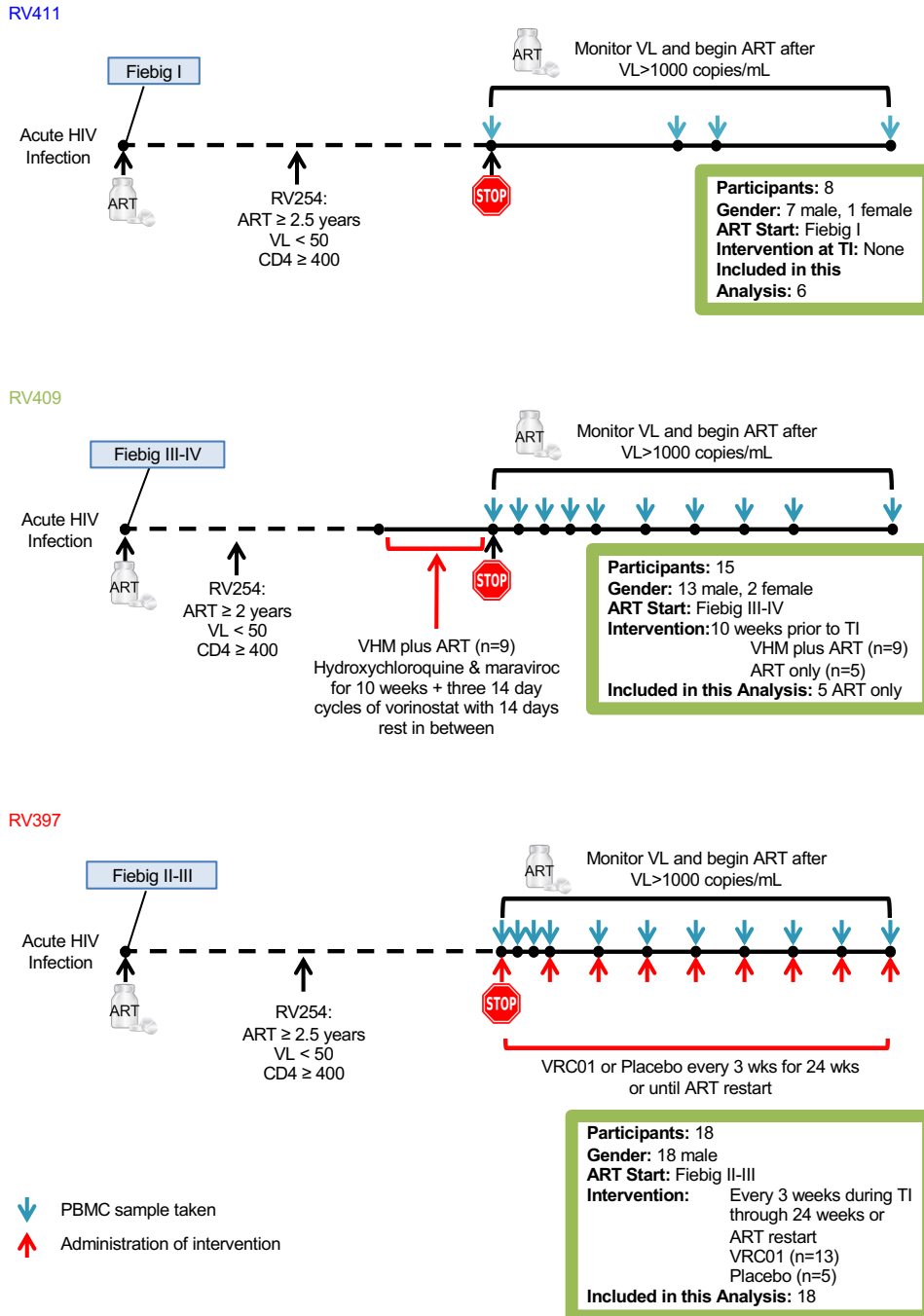
30 Supplemental Figure 5. Cytokine production by pDCs after in vitro activation.

31 Supplemental Figure 6. Changes in non-functional pDC phenotypes after ATI.

32 Supplemental Figure 7. Changes in signaling and gene expression in mDCs after ATI.

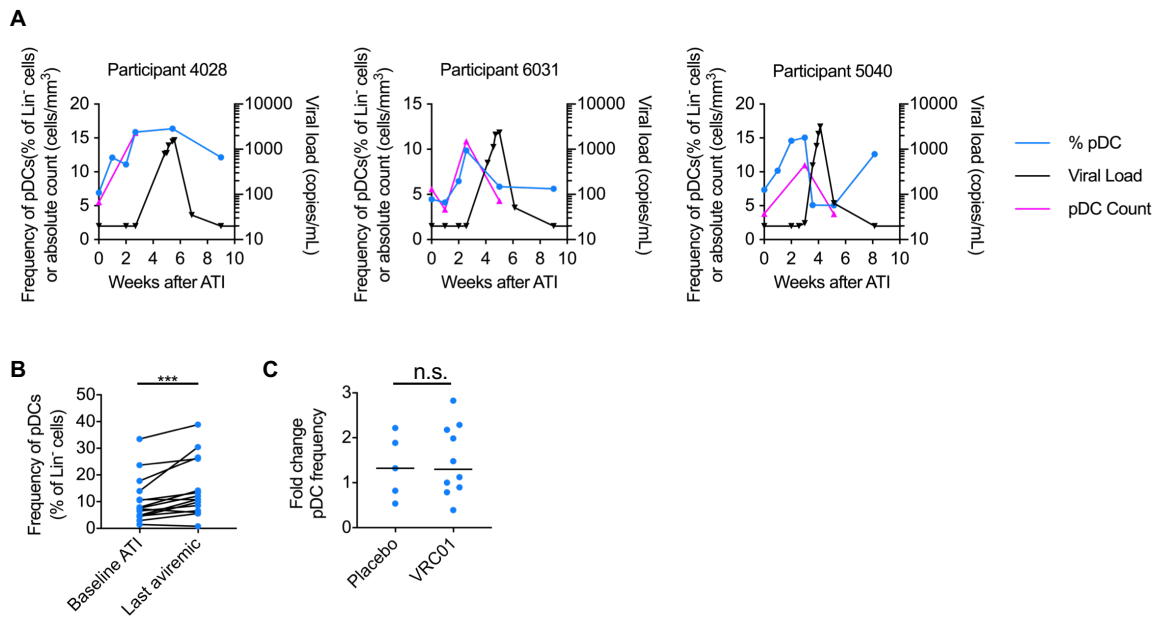
33 Supplemental Table 1. TaqMan Assay IDs

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 36 **Supplemental Figure 1. ATI study design.** Participants were enrolled in one of three ATI
 37 studies which occurred in Thailand. RV411 included 8 participants who were treated in Fiebig
 38 stage I of AHI and received no intervention at the time of ATI. RV409 included 15 participants
 39 who started ART in Fiebig stages III-IV, only 14 of whom interrupted ART. Nine participants in
 40 RV409 received vorinostat, hydroxychloroquine, and maraviroc in addition to ART for ten
 41 weeks prior to ATI. RV397 included 18 participants who started ART during Fiebig stages II-III
 42 of acute infection. Thirteen participants in this study received infusions of the VRC01 antibody
 43 at three-week intervals after ATI.

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46 **Supplemental Figure 2. Comparison of pDC frequency with pDC absolute count and**

47 **between treatment arms. (A)** pDC absolute count was determined in RV397 participants at time

48 points when the absolute CD4⁺ T cells count was measured at the time of sampling. Absolute

49 pDC count was calculated using the number of pDCs measured by flow cytometry and the ratio

50 of the absolute CD4⁺ T cell count to the number of CD4⁺ T cells measured by flow cytometry.

51 Shown are the pDC frequency (blue) and absolute pDC count (pink) for three participants for

52 whom pDC counts could be calculated prior to detection of plasma viremia. (B) The frequency of

53 pDCs was compared between the baseline time point at treatment interruption and the last

54 aviremic point after ATI in participants who received no therapeutic intervention. n = 15 (***)

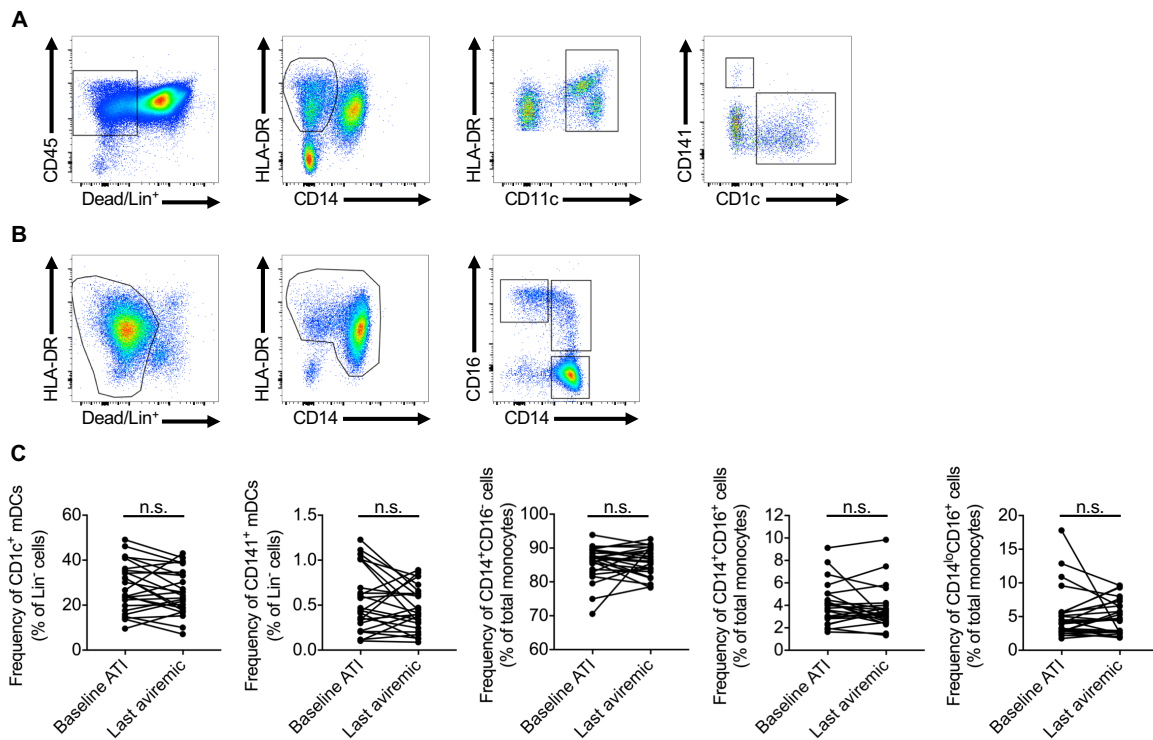
55 $p < 0.001$, Wilcoxon test). (C) The median fold change in pDC frequency between baseline ATI

56 and the last aviremic time point was compared between participants in the placebo and treatment

57 arms of RV397. Data were analyzed with the Mann-Whitney test. n = 5 placebo, 10 treatment.

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61 **Supplemental Figure 3. Changes in mDC and monocyte frequencies after ATI.**

62 (A) CD1c⁺ and CD141⁺ mDCs in the blood were identified within the CD45⁺HLA-DR⁺CD11c⁺

63 lineage negative cells (CD3⁻CD14⁻CD19⁻CD56⁻) by flow cytometry. (B) Classical

64 (CD14⁺CD16⁻), intermediate (CD14⁺CD16⁺), and nonclassical (CD14⁻CD16⁺) monocytes were

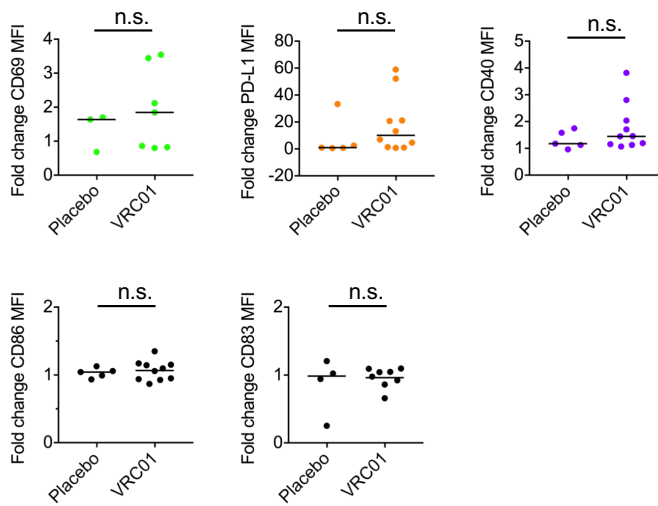
65 identified within the lineage negative (CD3⁻CD19⁻CD56⁻) HLA-DR⁺ population. (C) The

66 frequencies of dendritic cell and monocyte populations at baseline ATI and the last aviremic time

67 point are shown. Dendritic cell frequencies were measured as a percentage of the HLA-DR⁺

68 lineage negative cells. n = 25, 10 of whom received VRC01.

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71 **Supplemental Figure 4. Comparison of surface marker expression between arms of RV397.**

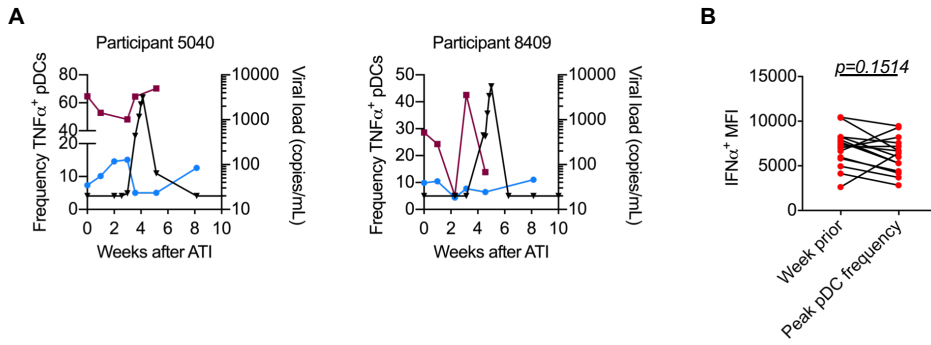
72 The median fold change in MFI of CD69, PD-L1, CD40, CD86, and CD83 between baseline

73 ATI and the last aviremic time point was compared between participants in the placebo and

74 treatment arms of RV397. Data were analyzed with the Mann-Whitney test. n = 5 placebo, 10

75 treatment.

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78 **Supplemental Figure 5. Cytokine production by pDCs after in vitro activation.**

79 (A) The percentages of pDCs that produced TNF α in response to imiquimod stimulation

80 (maroon) are shown. pDC frequency (blue) and HIV-1 viral load (black) are included for

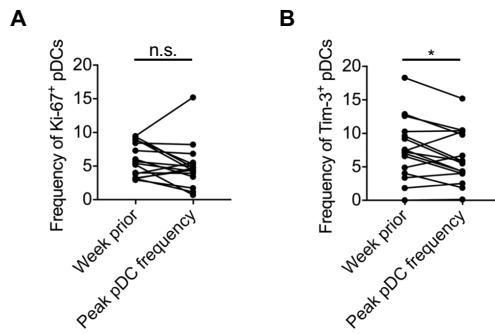
81 reference. (B) The mean fluorescence intensity of IFN α in pDCs after imiquimod stimulation is

82 compared between the time point before viremia at which the highest pDC frequency occurred

83 and the time point immediately prior. p-value was calculated with the Wilcoxon test. n = 14

84 participants from RV397, 11 of whom received VRC01.

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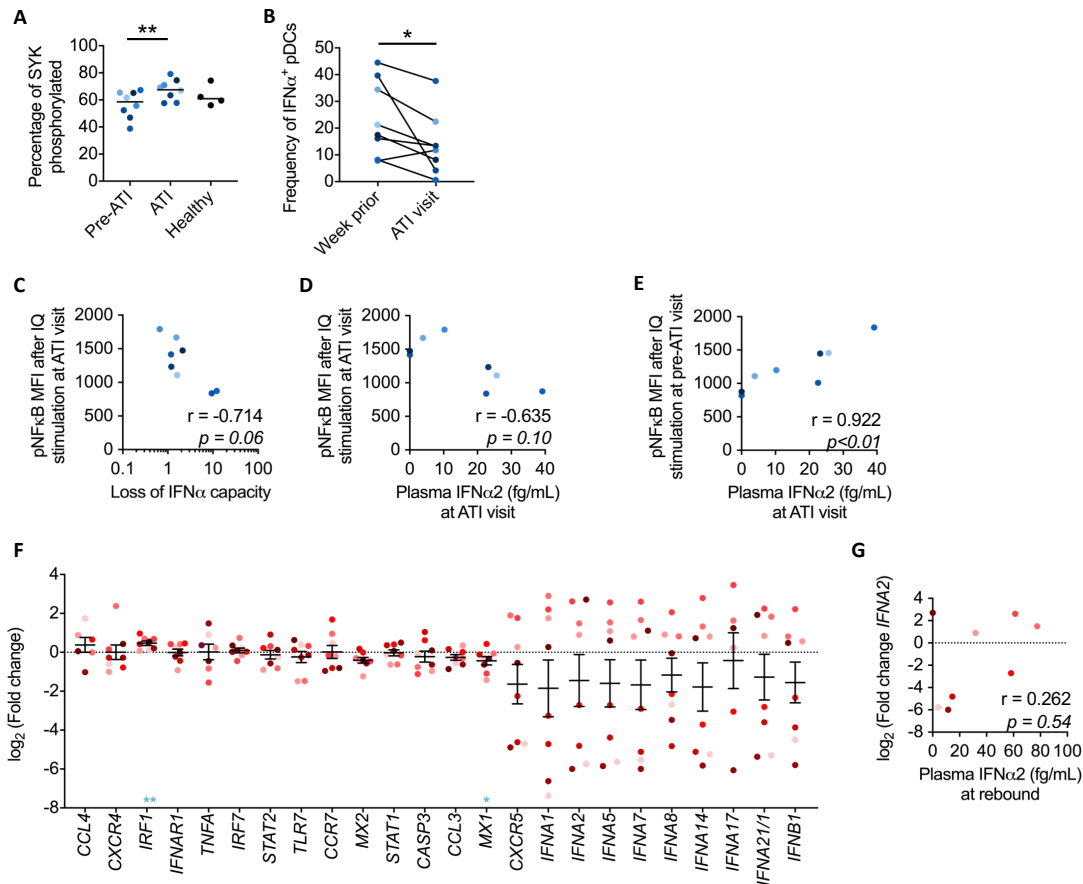


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87 **Supplemental Figure 6. Changes in non-functional pDC phenotypes after ATI.**

88 (A) The percentage of pDCs positive for Ki-67 was measured by flow cytometry and compared
 89 between the visit with peak pDC frequency and the visit immediately prior. (B) The frequency of
 90 pDCs expressing Tim-3 was measured by flow cytometry and compared between the week of
 91 peak pDC frequency prior to detectable viremia and the week prior (* $p < 0.05$, Wilcoxon test).

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 94 **Supplemental Figure 7. Changes in signaling gene expression in mDCs after ATI.**
 95 (A) The basal percentage of SYK that was phosphorylated was measured by flow cytometry
 96 immediately after thawing PBMCs from a visit prior to ATI (pre-ATI), after ATI (ATI), and
 97 from HIV⁻ healthy Thai individuals (bar represents median percentage, ** $p < 0.01$, Wilcoxon
 98 test). (B) The frequencies of pDCs producing IFN α after imiquimod stimulation are shown for
 99 the ATI visit when phosphorylation of IRF7 and NF- κ B was measured and the week prior (*
 100 $p < 0.05$, Wilcoxon test). Loss of IFN α capacity was calculated as the frequency of pDCs
 101 producing IFN α at the “Week Prior” visit divided by the ATI visit. (C) The phosphorylation
 102 levels of NF- κ B were measured in pDCs by flow cytometry after in vitro imiquimod stimulation
 103 of PBMCs. Spearman correlations were performed between the MFI of NF- κ B and the loss in
 104 IFN α producing capacity. (B-C) Plasma IFN α 2 levels were measured by SIMOA. Spearman
 105 correlations were performed between the plasma IFN α 2 levels at the ATI visit and the MFI of
 106 pNF- κ B in pDCs from the ATI visit (B) or the pre-ATI visit (C). (F) Gene expression was
 107 measured by BioMark in mDCs sorted from the pre-ATI and ATI visits. Shown is the difference
 108 in CT values between the ATI time point and pre-ATI time point after normalization to GAPDH
 109 (mean \pm SEM, * $p < 0.05$, ** $p < 0.01$, One sample Wilcoxon test, blue indicates values were no
 110 longer significant when corrected for a false discovery rate of 10% by the Benjamin-Hochberg
 111 Procedure). (G) Spearman correlation between the plasma IFN α 2 levels after viral rebound and
 112 the change in *IFNA2* levels in mDCs at the ATI time point. (A-E) $n = 8$ participants from RV397,
 113 all received VRC01 (F-G) $n = 8$ participants from RV397, 4 of whom received VRC01.
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115 **Supplemental Table 1. TaqMan Assay IDs**
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Gene	TaqMan Assay ID	Gene	TaqMan Assay ID
CASP3	Hs00234387_m1	IFNA7	Hs01652729_s1
CCL3	Hs00234142_m1	IFNA8	Hs00932530_s1
CCL4	Hs99999148_m1	IFNAR1	Hs01066116_m1
CCR7	Hs01013469_m1	IFNB1	Hs01077958_s1
CXCR4	Hs00237052_m1	IRF1	Hs00971965_m1
CXCR5	Hs00540548_s1	IRF7	Hs01014809_g1
GAPDH	Hs99999905_m1	MX1	Hs00895608_m1
IFNA1	Hs00855471_g1	MX2	Hs01550814_m1
IFNA14	Hs00533748_s1	STAT1	Hs01013996_m1
IFNA17	Hs00819693_sH	STAT2	Hs01013123_m1
IFNA2	Hs02621172_s1	TLR7	Hs00152971_m1
IFNA21/1	Hs00353738_s1	TNFA	Hs00174128_m1
IFNA5	Hs00818220_s1		

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