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Supplemental Information

**Neutralizing Antibodies Protect against
Oral Transmission of Lymphocryptovirus**

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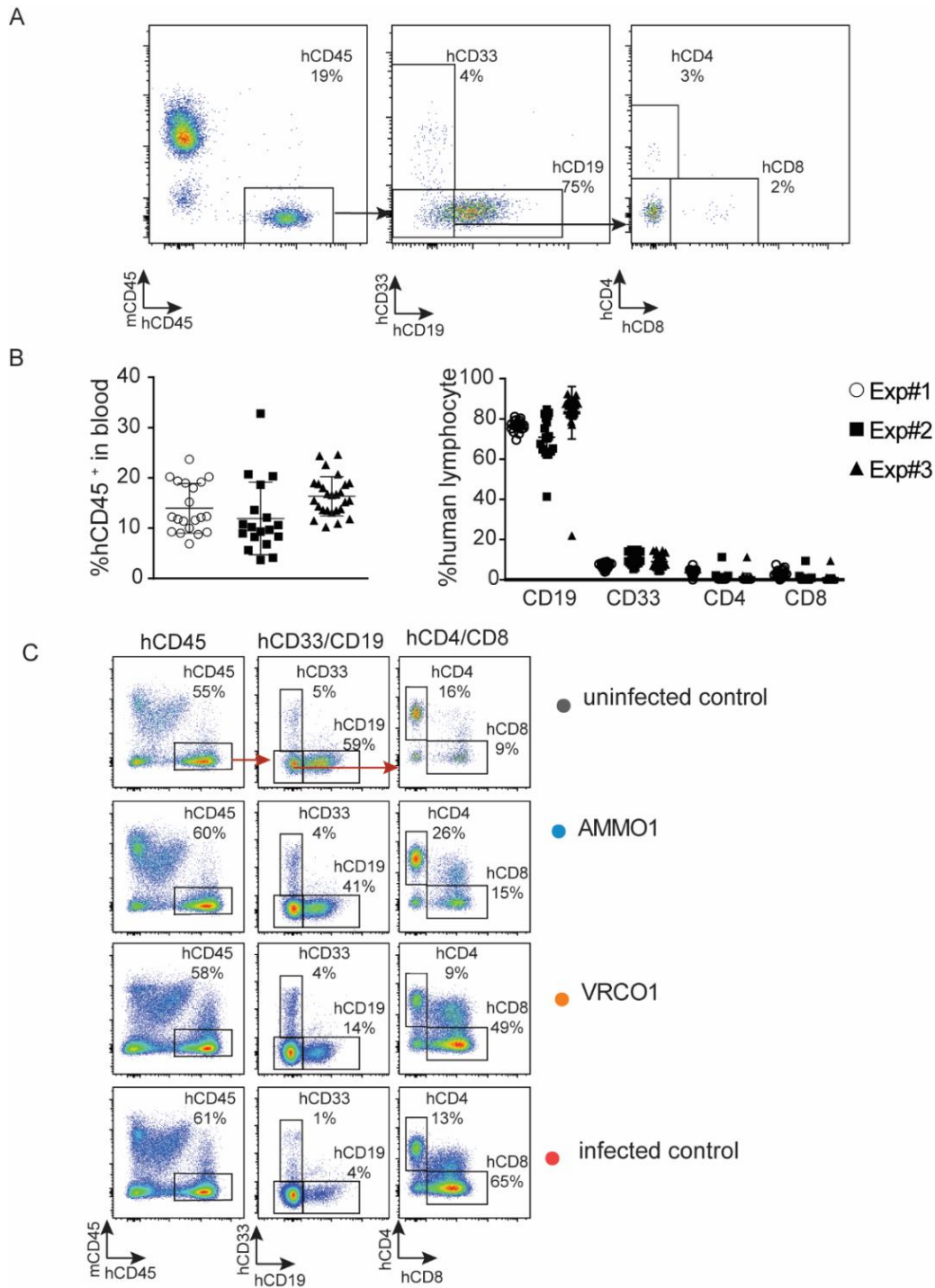


Figure S1. Human cell engraftment in NSG recipient mice prior to EBV infection, related to Figure 1. The frequency of human-derived cells in peripheral blood mononuclear cells from NSG mice were measured 12-14 weeks following HSC engraftment but prior to viral challenge. (A) Representative flow plot showing the gating strategy used to analyze peripheral blood and splenocytes. (B) Frequency of human (h) hCD45⁺ (left) and, hCD45⁺hCD19⁺hCD45⁺hCD33⁺, hCD45⁺hCD4⁺ and hCD45⁺hCD8⁺ cells (right) from 3 independent experiments. (C) Representative flow plot showing the gating and frequency of hCD45⁺ hCD19⁺, hCD45⁺hCD33⁺, hCD45⁺hCD4⁺ and hCD45⁺hCD8⁺ cells in the peripheral blood of mice from the indicated groups following viral challenge.

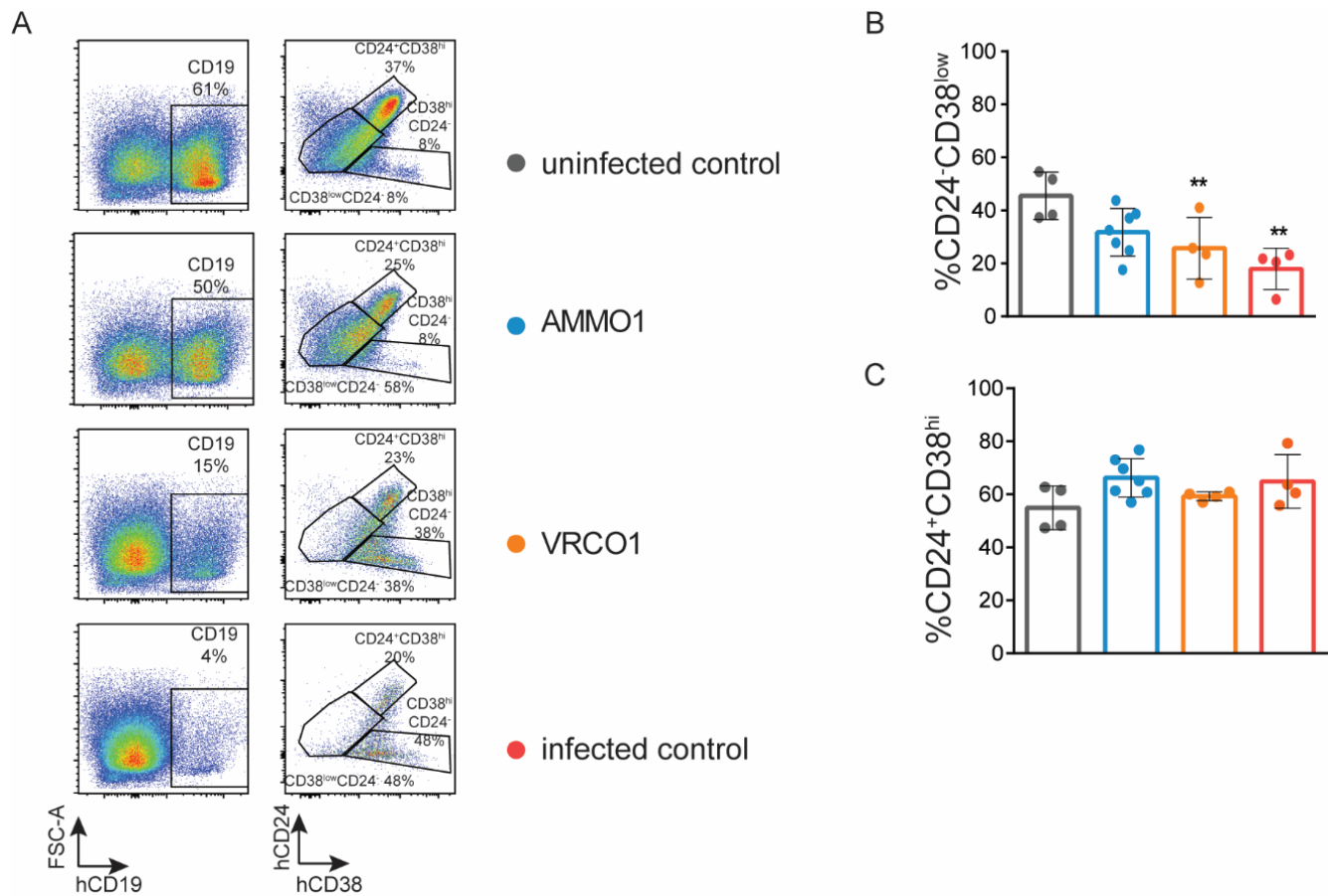


Figure S2. Human B cell phenotype in humanized mice following viral challenge, related to Figure 1. (A) Representative flow plot showing the staining and gating strategy to identify B cell subsets in splenocytes based upon staining with hCD19, hCD38 and hCD24 (pre-gated on the mCD45⁺hCD45⁺ human cell population). Frequency of **(B)** hCD45⁺hCD19⁺hCD24⁺hCD38^{low} and **(C)** hCD45⁺hCD19⁺hCD24⁺hCD38^{hi} cells from the animals from Figure 1. Viral infection is associated with a decrease in the overall proportion of hCD19⁺ cells and a greater proportion of hCD45⁺hCD19⁺hCD24⁺hCD38^{hi} cells. The frequency of hCD45⁺hCD19⁺hCD24⁺hCD38^{hi} cells is shown in Figure 1K. Asterisks represent groups that are statistically different from the uninfected controls determined by One Way ANOVA, Sidak multiple comparisons test **, $p \leq 0.002$.

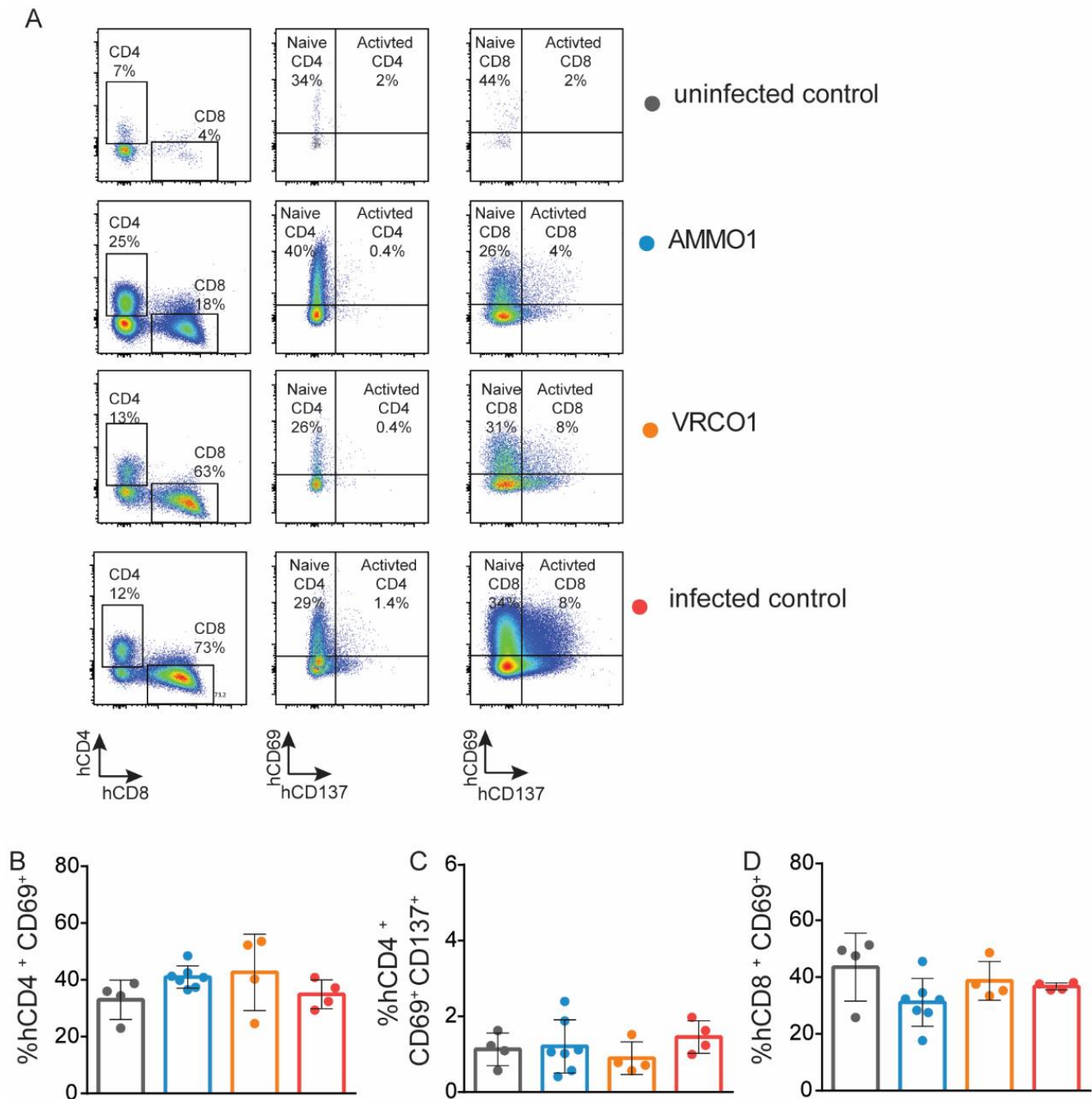


Figure S3. Splenic T cell phenotype in humanized mice following viral challenge, related to Figure 1. (A) Representative flow plot showing the staining and gating strategy of hCD4⁺, hCD8⁺, hCD69⁺ and hCD137⁺ T cells at necropsy (pre-gated on engrafted human cells based upon mCD45⁺hCD45⁺ staining). Frequency of **(B)** naïve hCD45⁺hCD69⁺hCD137⁻ and **(C)** activated hCD45⁺hCD69⁺hCD137⁺hCD4⁺ T cells, and the frequency of **(D)** naïve hCD45⁺hCD69⁺hCD137⁻hCD8⁺ T cells in the splenocytes of the animals from Figure 1. The frequency of activated hCD45⁺hCD69⁺hCD137⁺hCD8⁺ T cells is shown in Figure 1M.

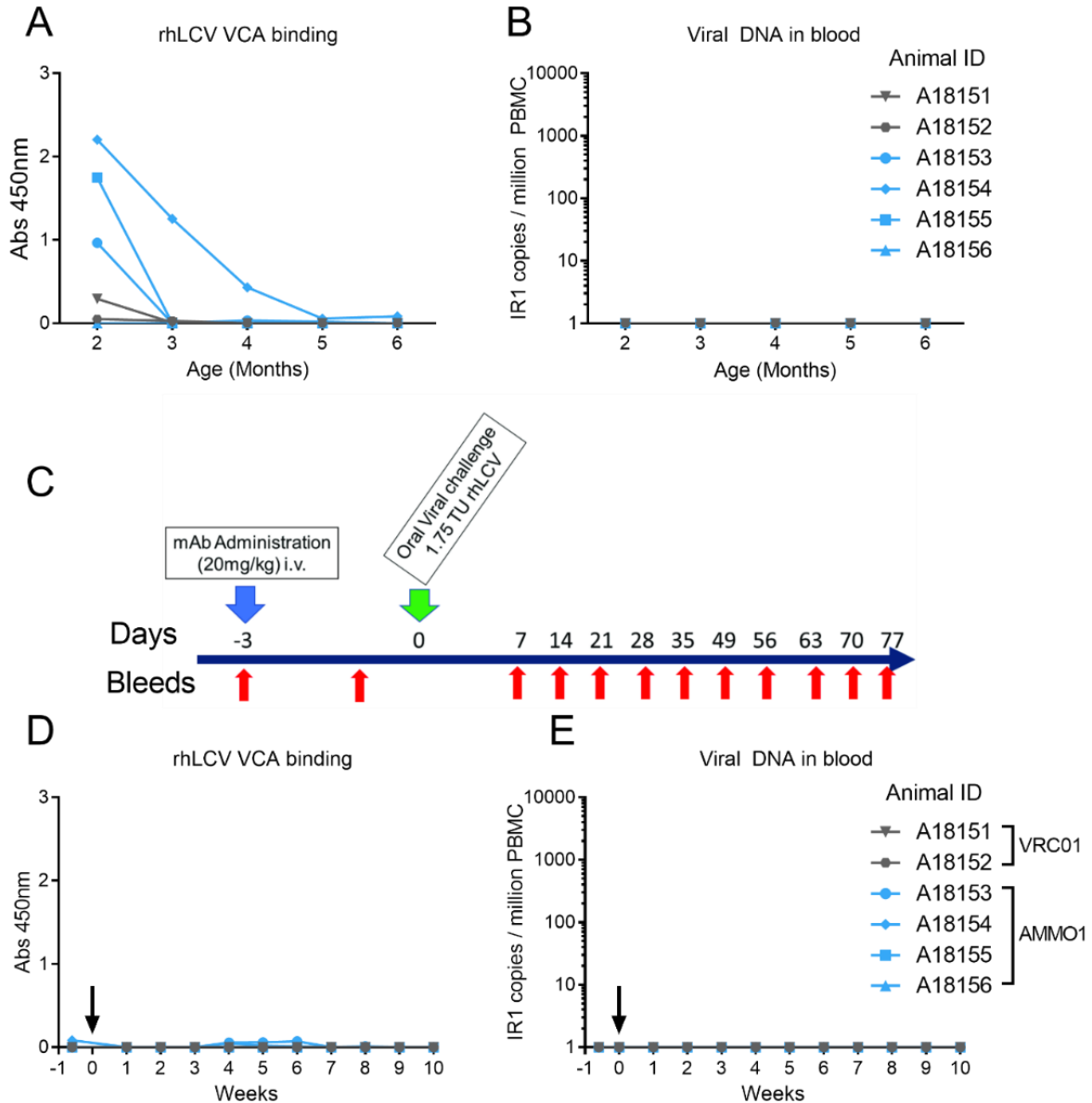


Figure S4. Monitoring for RhLCV infection in study macaques prior to and following an initial low-dose challenge, related to Figure 4. Rhesus macaques used in this study were obtained within 24-48 hours of birth. Beginning at 2 months of age, blood was collected monthly. **(A)** Plasma reactivity to pooled peptides derived from the rhesus lymphocryptovirus small viral capsid antigen (rhLCV VCA) was measured by ELISA. **(B)** Viral DNA was quantified in PMBC with a digital droplet PCR assay using primers against the internal repeat region 1 (IR1). Data from macaques that went on to receive infusions of AMMO1 or VRC01 are shown in blue and gray, respectively. **(C)** Timeline of initial challenge study. At ~6 months of age rhesus macaques were infused iv with 20 mg/kg of VRC01 (n=2, gray symbols) or AMMO1 (n=4, blue symbols). Three days later the macaques were challenged orally with 1.75 transforming units of rhLCV. Following challenge, blood was collected weekly for 10 weeks. **(D)** Plasma reactivity to pooled peptides derived from rhLCV VCA was measured by ELISA. **(E)** Viral DNA was quantified in PMBC using a digital droplet PCR assay with primers and probes against the internal repeat region 1 (IR1) of rhLCV. Black arrows indicate the time of viral challenge in **D** and **E**.