



# RELEVANCE OF CMV AND EBV CO-INFECTION TO CHRONIC INFLAMMATION IN HIV ELITE CONTROLLERS

Rayoun Ramendra<sup>1,2,3</sup>, Stéphane Isnard<sup>1,2</sup>, John Lin<sup>1,2</sup>, Brandon Fombuena<sup>1,2,3</sup>, Jing Ouyang<sup>1,2</sup>, Frank P. Dupuy<sup>1,2</sup>, Yonglong Zhang<sup>4</sup>, Malcolm Finkelman<sup>4</sup>, Ido Kema<sup>5</sup>, Cécile Tremblay<sup>6,7</sup>, Nicole F. Bernard<sup>2</sup>, Jean-Pierre Routy<sup>1,2,9</sup>

1. Chronic Viral Illness Service, McGill University Health Centre, Montreal, QC, Canada; 2. Infectious Diseases and Immunity in Global Health Program, Research Institute, McGill University Health Centre, Montreal, QC, Canada; 3. Department of Microbiology and Immunology, McGill University, Montreal, QC, Canada; 4. Associates of CapeCod Inc., Falmouth, MA, United States; 5. Department of Laboratory Medicine, University Medical Center, University of Groningen, the Netherlands; 6. Département de microbiologie, infectiologie et immunologie, Université de Montréal, Montréal, QC, Canada; 7. Centre de Recherche du Centre Hospitalier de l'Université de Montréal, Montréal, QC, Canada; 8. Division of Hematology, McGill University Health Centre, Montreal, QC, Canada

## Introduction

Elite controllers (EC) present with chronic inflammation and remain at increased risk of developing non-AIDS comorbidities. Microbial translocation is a contributor to chronic inflammation and CMV co-infection has been recently linked to increased microbial translocation. We previously reported that CMV seropositivity was associated with elevated epithelial gut damage and microbial translocation in ART-treated PLWH and HIV-uninfected controls. As Canada has one of the lowest CMV co-infection prevalence in the world, we evaluated the link between CMV seropositivity, microbial translocation, and inflammation among EC.

## Methods

Study samples were collected from 37 EC (25 CMV+, 12 CMV-). By HLA typing, we categorized participants with/without protective HLA alleles (B\*27, B\*57, n=16). We measured CD4 and CD8 T-cell counts, anti-CMV IgG and anti-EBV IgG titers, markers of epithelial gut damage REG3 $\alpha$  and I-FABP, markers of microbial translocation LPS, sCD14 and B-D-Glucan (BDG), as well as total IgG, IgM, IgA, IL-1 $\beta$ , IL-6 and kynurenine/tryptophan.

## Population Characteristics

Table 1, Participant Characteristics

	Elite controllers (n=35)
Age in years	
Median (IQR)	45 (39-50)
Range	26-72
Sex	
Male, n (%)	21 (60.0)
Female, n (%)	14 (40.0)
CD4 T-cells/ $\mu$ L	
Median (IQR)	650 (552-823)
Range	290-1090
CD8 T-cells/ $\mu$ L	
Median (IQR)	690 (409-968)
Range	180-1460
CD4/CD8	
Median (IQR)	1.1 (0.8-1.4)
Range	0.3-2.5
HIV VL (log <sub>10</sub> copies/mL)	
Median (IQR)	<1.7
Range	NA
Years on ART	
Median (IQR)	NA
Range	NA
CMV status	
Positive (%)	25 (71.4)
Anti-CMV IgG (IU/mL)	
Median (IQR)	23.9 (14.6-28.6)
Anti-EBV IgG (IU/mL)	
Median (IQR)	17.1 (8.3-22.1)

## Results

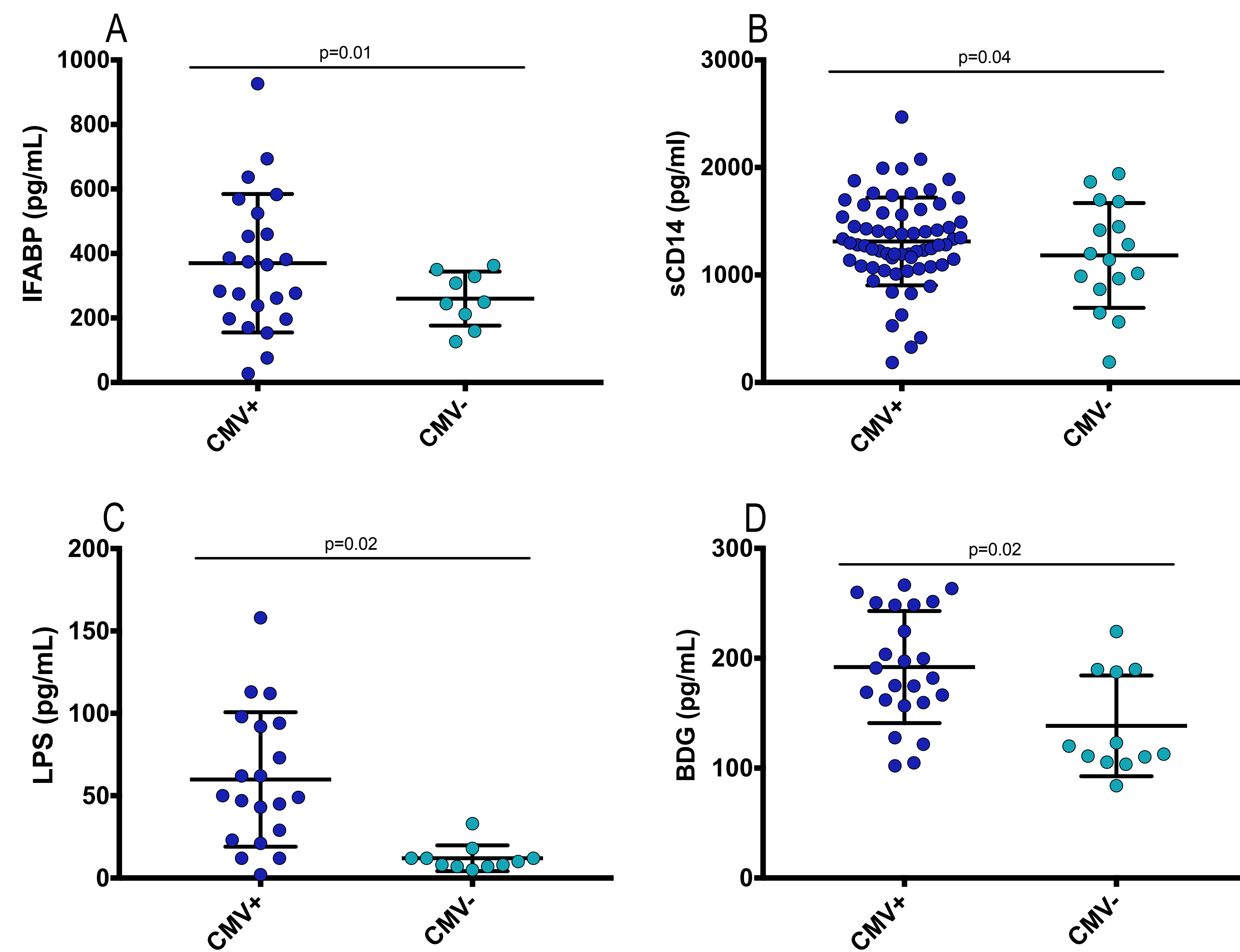


Figure 1. Plasma levels of markers of epithelial gut damage and microbial translocation in HIV elite controllers. CMV-seropositive elite controllers have elevated plasma levels of (A) I-FABP, (B) sCD14, (C) LPS, and (D) BDG compared to their CMV-seronegative counterparts.

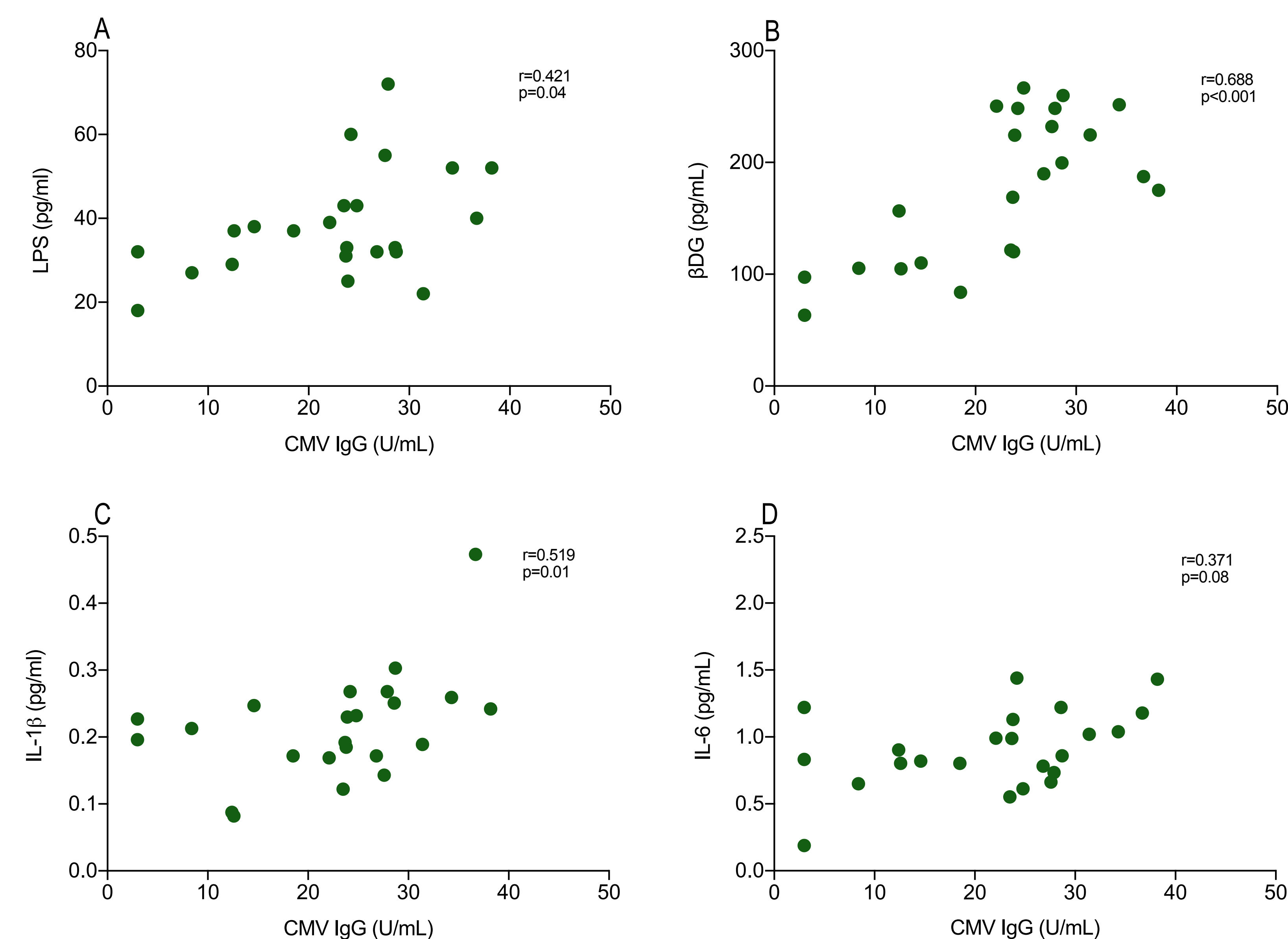


Figure 2. Correlations between anti-CMV IgG titers and plasma levels of (A) LPS, (B) BDG, (C) IL-1 $\beta$ , and (D) IL-6 in CMV seropositive EC.

## Results Cont.

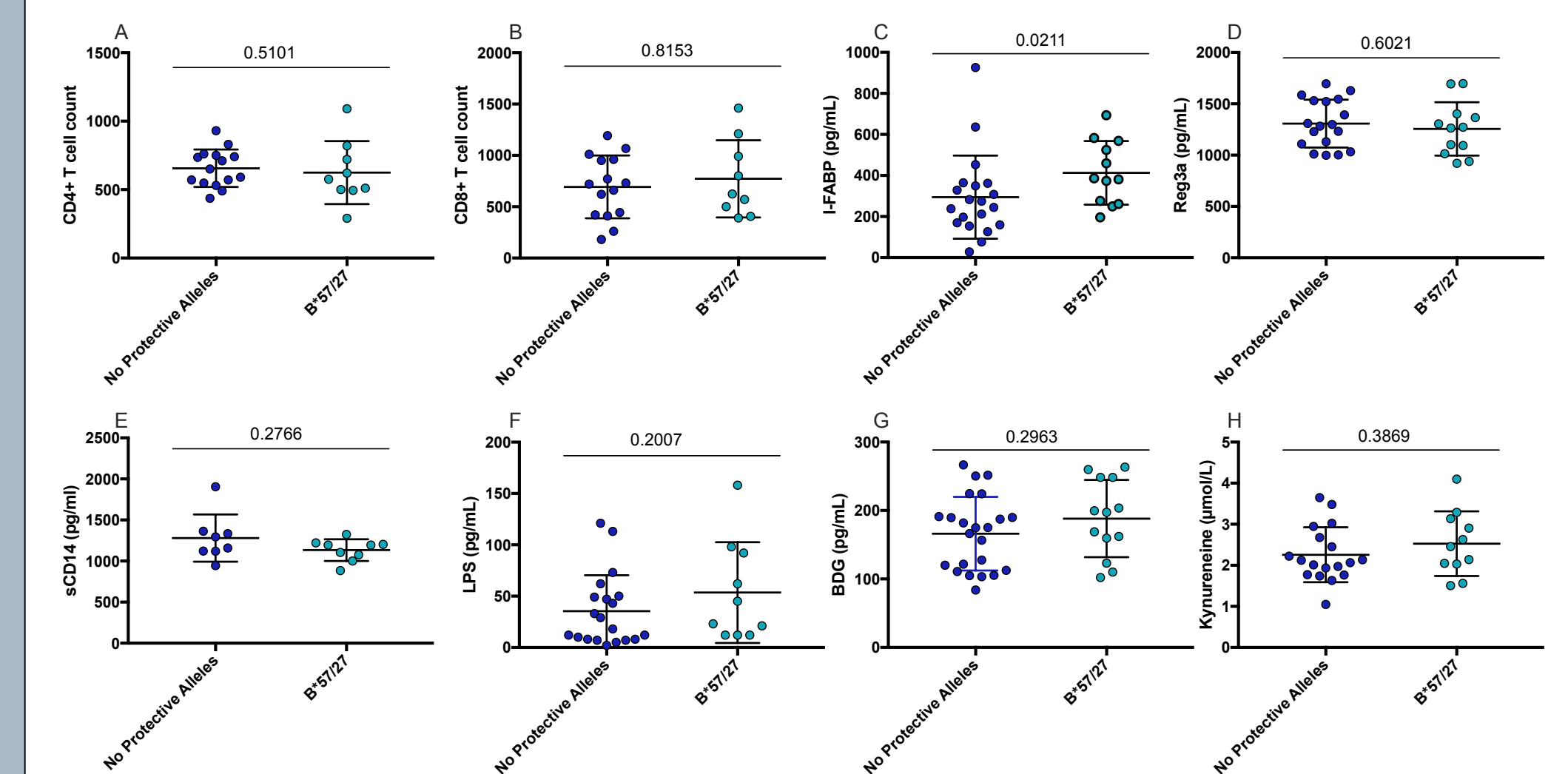


Figure 3. Comparison of (A) CD4+ T cell count, (B) CD8+ T cell count, (C) Intestinal Fatty Acid Binding Protein (I-FABP), (D) Reg3a, (E) sCD14, (F) LPS, (G) (1 $\rightarrow$ 3)-B-D-Glucan, and (H) Kynurenine amongst elite controllers with and without HLA-protective alleles.

## Conclusions

Markers of epithelial gut damage, microbial translocation, and inflammation were higher in CMV seropositive EC, irrespective of protective HLA alleles. Therefore, CMV co-infection emerges as an important contributor to gut damage and microbial translocation and may contribute to non-AIDS comorbidities in EC.

## Citations

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Ramendra R., Isnard S., et al. CMV seropositivity is associated with increased microbial translocation in people living with HIV and uninfected controls. *Clinical Infectious Diseases* 2020.

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