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Microbial Translocation of Fungal β -D-Glucan is Associated with Cardiovascular Risk in People Living with HIV

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Background

- **Microbial translocation** from the gut to circulation contributes to **immune activation** during HIV infection and is usually assessed by measuring plasma levels of bacterial lipopolysaccharide (LPS).
- Gut fungal colonization increases during HIV infection and **elevated systemic levels** of the **fungal polysaccharide β -D-Glucan (β DG)** have been reported in people living with HIV (PLWH). β DG represents one of the most abundant components of fungal cell walls and serves as a potent pathogen-associated molecular pattern in triggering antifungal immunity
- Despite antiretroviral therapy (ART), PLWH have an **increased risk of inflammatory comorbidities**. Gut damage and translocation of bacterial LPS and fungal β DG drive inflammation in ART-treated PLWH.

Herein, we investigated whether markers of gut damage and bacterial and fungal translocation are associated with cardiovascular risk in ART-treated PLWH.

Methods

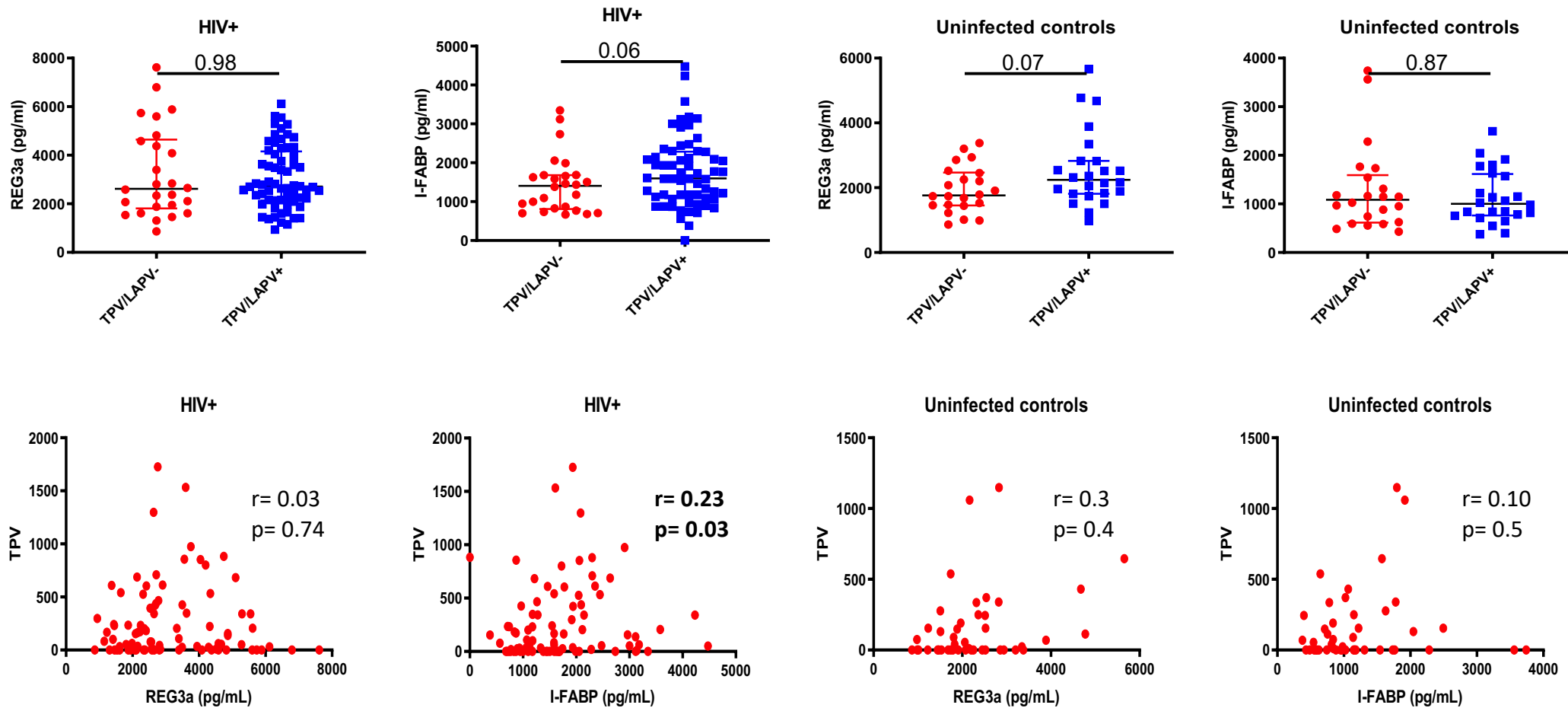
As part of the Canadian HIV and Aging Cohort (CHAC), we analyzed plasma from 147 participants > 40 y.o., 95 ART-treated PLWH with viral load <50 copies/ml, and 52 uninfected controls. Participants were free of clinical cardiovascular disease at baseline and underwent a cardiac computed tomography. Total coronary plaque volume (TPV) and low-attenuation coronary plaque volume (LAPV) were measured. Plasma levels of REG3 α , I-FABP and LPS were measured by ELISA. Plasma BDG levels were analyzed using the Fungitell[®] assay.

TPV/LAPV- \rightarrow coronary plaque = 0

TPV/LAPV+ \rightarrow coronary plaque volume >0



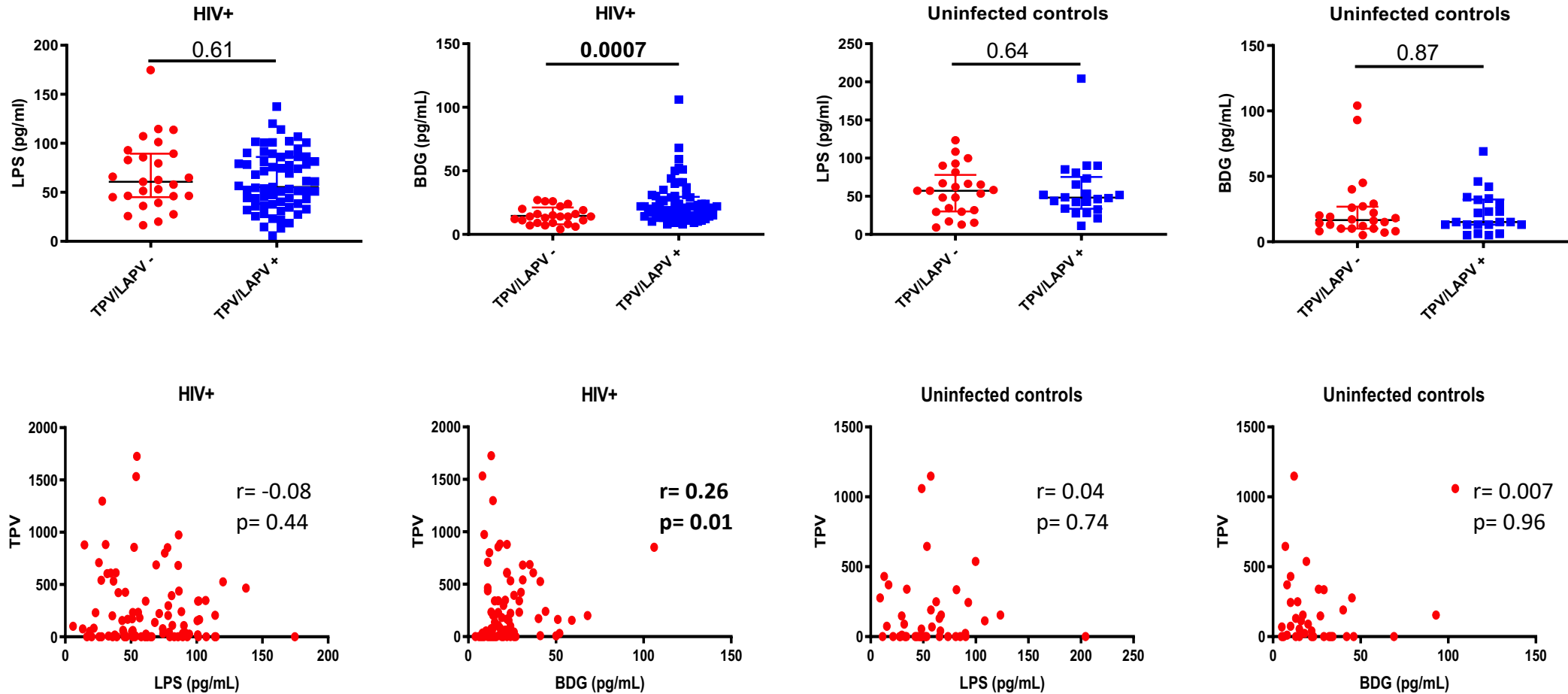
Gut Damage and Cardiovascular Risk



Plasma levels of I-FABP and Reg3α were slightly elevated although not significant in TPV/LAPV+ groups in comparison to TPV/LAPV- groups. Moreover, plasma levels of I-FABP and Reg3α correlated with TPV size in PLWH. In all participants (not depicted), TPV and LAPV correlated with gut damage markers REG3α ($r=0.2$, $p=0.03$ for both) and I-FABP ($r=0.2$, $p=0.001$ for both). In ART-treated PLWH, I-FABP but not REG3α levels correlated with TPV ($r=0.23$, $p=0.03$) and LAPV ($r=0.27$, $p=0.01$).



Microbial Translocation and Cardiovascular Risk



Plasma levels of BDG, but not LPS, were elevated in TPV/LAPV+ groups in comparison to TPV/LAPV- groups in PLWH. Moreover, Plasma levels of BDG, but not LPS, correlated with TPV size. In all participants (not depicted), BDG but not LPS levels correlated with TPV and LAPV ($r=0.2$, $p=0.04$ for both). In ART-treated PLWH, BDG but not LPS levels correlated with TPV ($r=0.26$, $p=0.01$) and LAPV ($r=0.25$, $p=0.010$).



Conclusion

Gut damage markers and translocation of fungal product BDG, but not bacterial LPS, were associated with the **burden of coronary atherosclerosis** in ART-treated PLWH.

Furthermore, following multivariate analysis, statin use and smoking habit (current or former) were also found to be associated with TPV in ART-treated PLWH ($p=0.024$ and $p=0.053$, respectively). However, **BDG proved to be a better marker** at predicting TPV risk ($p=0.004$). Per 10 units of BDG increase, there was shown to be a 3.17-fold increase in TPV risk.

Variable	N	Odds ratio	p
Age	84	1.06 (0.96, 1.18)	0.261
Sex			
Female	7	Reference	
Male	77	0.43 (0.02, 3.49)	0.489
BMI	84	0.93 (0.81, 1.05)	0.248
Statin use			
No	54	Reference	
Yes	30	5.37 (1.39, 27.92)	0.024
Smoking habit			
Never	21	Reference	
Current or former	63	3.90 (1.02, 16.81)	0.053
BDG	84	3.17 (1.62, 7.78)	0.004
(Intercept)		0.04 (0.00, 58.92)	0.383

More research is needed to appraise causality of the association; however, translocation of fungal products represents a potential therapeutic target to prevent cardiovascular disease in ART-treated PLWH.