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### Permalink

<https://escholarship.org/uc/item/2jj7k5j3>

### Journal

JAIDS Journal of Acquired Immune Deficiency Syndromes, 75(5)

### ISSN

1525-4135

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### Publication Date

2017-08-15

### DOI

10.1097/qai.0000000000001376

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Peer reviewed



# HHS Public Access

Author manuscript

*J Acquir Immune Defic Syndr.* Author manuscript; available in PMC 2018 August 15.

Published in final edited form as:

*J Acquir Immune Defic Syndr.* 2017 August 15; 75(5): e132–e141. doi:10.1097/QAI.0000000000001376.

## Higher Body Mass Index is Associated with Greater Proportions of Effector CD8+ T cells Expressing CD57 in Women Living with HIV

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### Abstract

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**Disclosures:** No relevant disclosures or conflicts of interest.

**AUTHOR CONTRIBUTIONS:** Research idea and study design: MJAR, SMB, ALL, EAF, PH, PCT and SDW; data acquisition: MJAR, SMB, LAS, ALL, MHC, PCT, SDW; data analysis/interpretation: MJAR, SMB, ALL, EAF, PH, PCT and SDW; statistical analysis: LAS; supervision or mentorship: EAF, PH, PCT, SDW. Each author contributed important intellectual content during manuscript drafting or revision and accepts accountability for the overall work by ensuring that questions pertaining to the accuracy or integrity of any portion of the work are appropriately investigated and resolved. SDW takes full responsibility that this study has been reported honestly, accurately, and transparently; that no important aspects of the study have been omitted; and that any discrepancies from the study as planned have been explained.

**Background**—A low proportion of CD28<sup>-</sup>CD8<sup>+</sup> T cells that express CD57 is associated with increased mortality in HIV infection. The effect of increasing BMI changes in the proportion of CD57<sup>+</sup>CD28<sup>-</sup>CD8<sup>+</sup> T cells among HIV-infected individuals on ART is unknown.

**Setting**—In a U.S. cohort of HIV-infected women, we evaluated associations of BMI and waist circumference with 3 distinct CD8<sup>+</sup> T cell phenotypes: % CD28<sup>-</sup>CD57<sup>+</sup>CD8<sup>+</sup> T cells, % CD57<sup>+</sup> of CD28<sup>-</sup>CD8<sup>+</sup> T cells and % CD28<sup>-</sup> of all CD8<sup>+</sup> T cells.

**Methods**—Multivariable linear regression analysis was used to estimate beta-coefficients for each of three T cell phenotypes. Covariates included HIV parameters (current and nadir CD4, current viral load), demographics (age, race, income, study site), and lifestyle (tobacco, alcohol use) factors.

**Results**—Of 225 participants, the median age was 46 years and 50% were obese (BMI>30 m<sup>2</sup>/kg). Greater BMI and waist circumference were both associated with higher %CD28<sup>-</sup>CD57<sup>+</sup>CD8<sup>+</sup> T cells and %CD57<sup>+</sup> of all CD28<sup>-</sup>CD8<sup>+</sup> T cells in multivariable analysis, including adjustment for HIV viral load (all p<0.05). The association between greater BMI and the overall proportion of CD28<sup>-</sup> CD8<sup>+</sup> cells in fully adjusted models (0.078, 95% CI: (-0.053 – 0.209) were not significant.

**Conclusions**—In this analysis, greater BMI and waist circumference are associated with greater expression of CD57 on CD28<sup>-</sup>CD8<sup>+</sup> T cells and a greater proportion of CD57<sup>+</sup>CD28<sup>-</sup> CD8<sup>+</sup> T cells. These findings may indicate that increasing BMI is immunologically protective in HIV-infected women. Future research is needed to understand the prognostic importance of these associations on clinical outcomes.

## Keywords

HIV; obesity; WIHS; CD57; immune senescence

## INTRODUCTION

The advent of highly active antiretroviral therapy (ART) has had a profound impact on HIV-associated morbidity and mortality.<sup>1</sup> While the prevalence of HIV-associated wasting has declined, however, the proportion of overweight and obese HIV-infected individuals is increasing.<sup>2</sup> While recent studies have failed to show a consistent effect for obesity in the pathogenesis of cardiovascular disease in HIV-infected individuals,<sup>3,4</sup> there is evidence demonstrating that obese HIV-infected individuals have a greater prevalence of metabolic diseases, including type 2 diabetes mellitus, compared to non-obese persons.<sup>5-7</sup> In addition, increasing BMI is associated with greater innate and adaptive immune activation, even in the setting of treated HIV infection.<sup>8,9</sup> As observed in the general population<sup>10</sup>, serum levels of systemic markers of inflammation such as C-reactive protein (CRP) and tumor necrosis factor (TNF)- $\alpha$ <sup>11-13</sup> are higher among HIV-infected adults with greater adiposity. There is also evidence that adiposity influences adaptive immune responses, with several recent studies suggesting that obesity independently influences immunologic recovery in individuals initiating ART.<sup>14-17</sup>

Whether increasing BMI also influences the development of other immune defects in HIV-infected individuals is less clear. Studies have demonstrated that HIV infection leads to numerous CD8+ T cell abnormalities, some of which are also seen in elderly populations (e.g., an expansion of CD28- CD8+ T cells) and some of which are distinct from those observed in the elderly (e.g., decreased terminal differentiation and a reduced proportion of CD28-CD8+ T cells that express CD57)<sup>18,19</sup>. It is possible that increasing adiposity may further affect T cell differentiation due to elevations in systemic low grade inflammation, heightened oxidative stress,<sup>20</sup> alterations in nutrition and micronutrients<sup>21,22</sup>, psychosocial factors such as depression and poor quality of life<sup>23</sup> or perturbations in serum concentrations of leptin, which has been shown to influence T cell activation and proliferation<sup>24</sup>.

The aim of our study was to investigate the relationship between BMI and CD8+ T cell phenotypes that have been previously linked to mortality in a diverse cohort of HIV-infected women, while adjusting for potential confounders such as socioeconomic factors, lifestyle and behavioral variables, and HIV viral load. Given the numerous deleterious inflammatory and metabolic sequelae of obesity<sup>9,11</sup> we hypothesized that greater BMI would be associated with a pattern of poor CD8+ T cell differentiation in HIV-infected adults that has been previously associated with mortality in this setting.

## METHODS

### Study design and population

The Women's Interagency HIV Study (WIHS) is a large, multicenter, ongoing prospective cohort of HIV-infected and at-risk women in the United States<sup>25,26</sup>, established in 1993. At semi-annual visits, participants are interviewed and examined, and serum specimens are collected and stored in a -80°C freezer. The enrolled women are representative of U.S. women living with HIV in terms of demographic and clinical parameters<sup>25</sup>. Specifically, this investigation was a cross-sectional study of WIHS women enrolled in the WIHS Food Insecurity Sub-study from six U.S. WIHS sites (Bronx, Brooklyn, Chicago, Los Angeles, San Francisco, and Washington, DC) who were seen from April 2013 through September 2013. Our analysis was limited to HIV-infected participants who were on ART, and did not have diagnoses of cancer, autoimmune diseases, or hepatitis B or C virus infection. All participants were fasting for laboratory studies.

### Laboratory methods

T cell immune senescence was characterized by polychromatic flow cytometry on frozen/thawed peripheral blood mononuclear cells (PBMCs) at Rush University Medical Center (Landay lab). Thawed cells were stained for cell viability with the Aqua Live/Dead cell stain kit (Invitrogen, Carlsbad, CA) prior to cell surface staining. Cell surface markers were stained with fluorochrome-conjugated monoclonal antibodies to CD3, CD4, CD8, CD28 and CD57 (BD Biosciences, San Jose, CA). Cells were acquired within 24 hours on a LSR2 flow cytometer using BD FACSDiva software (BD Biosciences, San Jose, CA). Analysis of flow cytometry data was performed using FlowJo software (Ashland, OR). Immune senescence

(CD57+ \ CD28-) analysis was performed and reported on singlet live (Aqua-) CD3+ \ CD8+ T cells.

### Predictor

The primary predictor in this study was body mass index (BMI, continuous in kg/m<sup>2</sup>) measured at the visit closest to the visit where serum was obtained for testing of cellular immune markers. We also examined BMI as a categorical measure with obesity defined as BMI > 30kg/m<sup>2</sup>. Anthropometric characteristics included waist circumference [in cm], hip circumference [in cm], and hip to waist ratio were also included as predictors.

### Outcome

The primary outcomes of interest were the CD8+ T cell phenotypes that characterize immunosenescence, specifically three separate CD8+ T cell phenotypes: the proportion of CD8+ T lymphocytes that were CD28-CD57+, the proportion of CD8+CD28- expressing CD57, and the proportion of CD8+ T lymphocytes that were CD28- of the total CD8+ T lymphocyte population. Each of these three phenotypes was considered as individual outcomes and measured continuously in cells/ $\mu$ L. The first T cell phenotype in our analysis, % CD28- CD57+ of CD8+ T cells, was calculated as the summation of two distinct CD8+ T cell populations identified on flow cytometry, namely the proportion of CD8+ T lymphocytes that were CD57- CD28- and the proportion that were CD57+ CD28-. The second T cell phenotype, % CD57+ of CD28- of CD8+ T cells, was reported output from the flow cytometry. The third output was derived from dividing proportion of CD28- CD57+ of CD8+ T cells by the proportion CD28- of CD8+T cells that were expressing CD57.

### Covariates

Candidate covariates included sociodemographic factors: age [in years at the visit], ethnicity [non-Hispanic White, African-American, Hispanic or other], educational attainment [less than high school, completed high school, some college or greater], employment status [employed or not], annual income [less than or equal versus greater than \$30,001] and, study site [Bronx and Brooklyn, NY, Washington, DC, Chicago, IL and San Francisco, CA]. Lifestyle factors considered were: smoking [none, current, or past], smoking duration [in years smoked], alcohol use [none; light drinking, defined as consumption of 1-15g of alcohol/day, moderate drinking, defined as consumption of 15-30g alcohol/day; or heavy drinking, defined as >30 g/day]. HIV-related factors included: current CD4+ T cell count (in cells/mm<sup>3</sup>), nadir CD4+ T cell count (in cells/mm<sup>3</sup>), HIV viral load (log transformed in copies/mL), history of clinical AIDS, and current use of antiretroviral therapy (ART). Variables were selected by *a priori* consideration of confounders on the relationship between BMI and the outcomes of interest, informed by previous literature<sup>15,27,28</sup>. Height was included as a covariate because use of height-squared as the denominator of BMI does not completely adjust for height.<sup>29</sup> CMV serostatus was not available for inclusion in the analysis, although based on other analyses performed in the WIHS cohort, we anticipate that prevalence of CMV seropositivity was very high.<sup>30</sup>

## Statistical analysis

Summary statistics were obtained for covariates. We compared sociodemographic and clinical characteristics within each of the three CD8+ T cell phenotypes using the Kruskal-Wallis test for continuous variables and the Fisher exact test for categorical variables. All continuous covariates were standardized by dividing by the interquartile range to improve comparison and interpretation. Given the exploratory nature of this study, stepwise model building was used to select among the available covariates, retaining only those variables with  $P < 0.2$ , and was performed individually for each of the three outcomes. Linear regression models were used to estimate beta-coefficients for the primary predictor and covariates selected by stepwise modeling, on each of the outcomes. An alpha of 0.05 was selected as the significance threshold. All analyses were conducted using STATA (version 13, StataCorp, College Station, TX). In an additional sensitivity analysis, separate models restricted to those with an undetectable viral load (defined as  $< 20$  or  $48$  copies/ml depending on the timing of the specimen) were also constructed to control for the immunologic effect of persistent viremia.

Finally, in separate models we evaluated other anthropometric measures as primary predictors of CD8+ T cell phenotypes. In models controlling for demographic, lifestyle factors and HIV-specific variables, we determined the association of waist circumference and waist to hip ratio with each of the CD8+ T cell phenotypes. In each case, fully adjusted models that did and did not adjust for HIV viral load were compared in order to determine the extent to which HIV viremia attenuated associations between body composition parameters and CD8+ T cell phenotypes.

## Ethics statement

All participants provided written and informed consent for participation in the Women's Interagency HIV Study (WIHS). The research conducted as part of the WIHS was approved by institutional review boards at all study sites. All studies were conducted according to the principles outlined in the Declaration of Helsinki.

## RESULTS

We present data on 225 women included in this analysis. Most participants identified as Black (Table 1). The median age was 46.5 years. Nearly two thirds of the participants had an annual income of less than \$30,000 and had completed 12 or fewer years of education. The median BMI was  $30 \text{ kg/m}^2$ . More than three quarters of study participants were classified as overweight and 50% were either obese or morbidly obese. The median duration of ART therapy was 13 years. The majority (74%) of participants had an undetectable viral load. The median current CD4 T cell count was  $626 \text{ cells/mm}^3$ , and the median CD4 T cell count nadir was  $310 \text{ cells/mm}^3$ .

### Association of body composition parameters with CD8+ phenotypes

There were significant positive correlations between increasing BMI and the proportion of CD8+ cells that were CD28-CD57+ ( $r=0.187$ ,  $p<0.05$ ) (Figure 1a) and the proportion of CD28- CD8+ T cells expressing CD57+ ( $r=0.185$ ,  $p<0.05$ ) (Figure 1b), but not the overall

proportion of CD28<sup>-</sup> CD8<sup>+</sup> T cells ( $r=0.053$ ,  $p=0.418$ ) (Figure 1c). Neither current CD4 count (Figure 1d) nor HIV viral load (Figure 1e) were correlated with BMI ( $R=0.021$ ,  $p=0.746$  and  $R=-0.08$ ,  $p=0.221$  respectively).

In regression analysis, each IQR increase in BMI was associated with a significantly greater proportion of CD28<sup>-</sup>CD57<sup>+</sup> of CD8<sup>+</sup> T cells (0.184, 95% confidence intervals [CI] 0.06, 0.307) and a greater proportion of CD28<sup>-</sup>CD8<sup>+</sup> T cells expressing CD57<sup>+</sup> in unadjusted analysis (0.175, 95% CI: 0.056, 0.293). Furthermore, in fully adjusted models, controlling for demographic variables (including annual income, study site location and highest level of education), height and HIV-specific variables (including current CD4 T cell count and viral load), the association between greater BMI and both the proportion of CD28<sup>-</sup>CD57<sup>+</sup> CD8<sup>+</sup> T cells and the proportion of CD28<sup>-</sup>CD8<sup>+</sup> T cells expressing CD57<sup>+</sup> remained significant (0.166, [95% CI: 0.04, 0.29] and 0.145, [0.026, 0.263]). The association between greater BMI and the overall proportion of CD28<sup>-</sup> CD8<sup>+</sup> cells in unadjusted, (0.051, 95% CI: -0.073, 0.175) and fully adjusted models (0.078, 95% CI: (-0.053 - 0.209) were not significant.

Comparing different measures of body composition in fully adjusted models (Table 3), greater BMI and waist circumference were associated with a significantly greater proportion of CD28<sup>-</sup>CD57<sup>+</sup> of CD8<sup>+</sup> T cells and the proportion of CD57<sup>+</sup> of CD28<sup>-</sup> CD8<sup>+</sup> T cells. These associations were only minimally attenuated by addition of viral load to the models, remaining significant in both cases. When examining BMI as a dichotomous variable comparing obese (BMI>30) to non-obese (BMI≤30), there was a positive association between obesity and the proportion of CD57<sup>+</sup> of CD28<sup>-</sup>CD8<sup>+</sup> T cells, however, the association was not significant ( $p=0.065$ ). Waist to hip ratio was not associated with significant differences in the proportion of CD28<sup>-</sup>CD57<sup>+</sup> of CD8<sup>+</sup> T cells or the proportion of CD28<sup>-</sup>CD8<sup>+</sup> T cells expressing CD57 regardless of adjustment for viral load (all  $p>0.01$ ).

In a sensitivity analyses restricted to individuals that were virologically suppressed ( $n=168$ ), the estimates for the associations of BMI and waist circumference with the CD8<sup>+</sup> phenotypes were similar to analyses with both virologically suppressed and unsuppressed participants included in the models, although the associations were no longer significant (all  $p>0.1$ ) (data not shown).

### Association of other variables with CD8<sup>+</sup> phenotypes

In fully adjusted multivariable analysis, older age was associated with a significantly greater percentage of CD8<sup>+</sup> T cells that were CD28<sup>-</sup>CD57<sup>+</sup> (0.134, 95% CI: 0.009, 0.259) and a significantly greater percentage of CD57<sup>+</sup> of CD28<sup>-</sup>CD8<sup>+</sup> T cells (0.154, 95% CI: 0.037, 0.272). Being Hispanic was also associated with a significantly greater proportion of CD57<sup>+</sup>CD28<sup>-</sup>CD8<sup>+</sup> T cells or CD57<sup>+</sup> of all the CD28<sup>-</sup> CD8<sup>+</sup> T cells (0.450, 95% CI: 0.069, 0.68) in multivariable analysis.

Among the HIV-related factors, higher HIV viral load was associated with a small, albeit significant, higher percentage of CD57<sup>+</sup> CD28<sup>-</sup> of CD8<sup>+</sup> T cells (0.014, 95% CI: 0.005, 0.023) and percentage of CD57<sup>+</sup> of CD28<sup>-</sup>CD8<sup>+</sup> T cells (0.012, 95% CI: 0.004, 0.021) in unadjusted analyses. These associations were attenuated and ceased to be significant after

adjusting for other factors. HIV viral load was not associated with a significant difference in the overall proportion of CD28<sup>-</sup> CD8<sup>+</sup> T cells in either unadjusted or fully adjusted models. Each IQR higher CD4 count was associated with a lower proportion of CD57<sup>+</sup>CD28<sup>-</sup>CD8<sup>+</sup> T cells (-0.241, 95% CI: -0.354, -0.128) and CD28<sup>-</sup> CD8<sup>+</sup> T cells that were CD57<sup>+</sup> (-0.215, 95% CI: -0.324, -0.107) as well as the overall proportion of CD28<sup>-</sup> CD8<sup>+</sup> T cells (-0.126, 95 CI: -0.273, -0.047). The associations between greater CD4 count and lower percentage of CD57<sup>+</sup>CD28<sup>-</sup>CD8<sup>+</sup> T cells count and the proportion of CD28<sup>-</sup>CD8<sup>+</sup> T cells that were CD57<sup>+</sup> remained significant even in fully adjusted models. By contrast each IQR higher CD4 nadir was not associated with significant differences in CD8<sup>+</sup> cell phenotypes, but longer duration on ART was associated with a significantly lower proportion of all CD8<sup>+</sup> cell phenotypes in unadjusted analysis (all  $p < 0.001$ ). Additional exploratory analyses, adjusting for total CD8<sup>+</sup> T cell population and CD8<sup>+</sup>/CD4<sup>+</sup> ratio, did not alter the associations between three CD8<sup>+</sup> phenotypes and BMI. Furthermore, including percent body fat data, available for a subset of 195 women, in multivariable regression models did not alter these associations either (data not shown).

## DISCUSSION

In this population of women living with HIV infection, most of who had been on ART for more than ten years and were overweight or obese, we found that greater BMI was associated with a greater proportion of CD57<sup>+</sup>CD28<sup>-</sup>CD8<sup>+</sup> T cells. We also demonstrated that, while BMI was not associated with a significant difference in the overall proportion of CD28<sup>-</sup>CD8<sup>+</sup> T cells, greater BMI was associated with a greater proportion of CD28<sup>-</sup>CD8<sup>+</sup> that expressed CD57 even after further adjustment for HIV viral load.

Contrary to our a priori hypothesis that adiposity promotes maturational CD8<sup>+</sup> T cell defects that predict increased mortality in treated HIV infection (e.g., a low proportion of CD28<sup>-</sup>CD8<sup>+</sup> T cells that express CD57), we found that greater BMI was associated with greater expression of CD57 by CD28<sup>-</sup>CD8<sup>+</sup> T cells. Furthermore, the association was only minimally attenuated by additional adjustment for HIV viral load. Given recent studies by Lee et al showing that *increased* expression of CD28<sup>-</sup>CD8<sup>+</sup> T cells that express CD57 was predictive of decreased mortality in HIV-infected individuals on ART,<sup>27,28</sup> we speculate that the association between BMI and increased proportions of effector cells expressing CD57 may indicate that increasing BMI is immunologically protective.

In Lee's analysis, several markers of innate immune activation, including sCD14 and IL-6 were correlated with decreased CD28<sup>-</sup>CD8<sup>+</sup> cells expressing CD57. A potential explanation for our findings of an association between greater BMI and *increased* CD28<sup>-</sup>CD8<sup>+</sup> T cells expressing CD57 could be that a greater BMI reflects less of an inflammation-associated catabolic state. If this were the case, it would be the converse of what is seen in HIV-uninfected elderly adults where higher frequencies of CD28<sup>-</sup>CD57<sup>+</sup> CD8<sup>+</sup> T cells are predictive of increased mortality not improving health.<sup>31</sup> These contradictory results may reflect fundamentally distinct immunologic pathways mediating the functional T cell defects that persist during treated HIV and those that characterize the aging process. Since increasing BMI in HIV-infected adults initiating ART reflects restorative changes in both lean muscle mass and adipose tissue,<sup>32</sup> the association between increasing expression of



CD57 and BMI may be evidence of how suppressive ART leads to healthy changes in both body composition and adaptive immunity. Further research is necessary to better understand the interplay of inflammation, changing body composition and T cell differentiation in this population.

Another possible explanation is that increased adiposity promotes expansion of CD57+ cells. This seems plausible, given that persons with congenital lipodystrophy also have lower proportions of CD57+CD8+ T cells than in the normal population.<sup>33</sup> Furthermore, in a study of HIV-uninfected adolescents by Spielmann et al, being at risk for obesity was associated with a higher proportion of CD28–CD8+ T cells, including the CD57+CD28–CD8 subset.<sup>34</sup> Given that untreated HIV-infection is associated with a lower proportion of CD57+CD28–CD8+ T cells, our findings suggest that some degree of adiposity may counteract the adverse changes in T cell differentiation seen in untreated HIV-infection. We found that BMI was only associated with an increase in the proportion of cells expressing CD57, rather than the larger proportion of CD28–CD8+ T cells, which were associated with increased obesity in Spielmann's analysis. We speculate that this difference may be more consistent with a decreased immune activation (associated with a reduced catabolic state and greater BMI) in HIV allowing for greater terminal differentiation of effector CD8+ T cells, in contrast to a scenario where increased immune activation in obese adolescents drives CD28–CD8+ T cells to express CD57. Nevertheless, we cannot exclude the possibility that the association between BMI and terminally differentiated CD8+ T cells are associated with long-term deleterious health effects. Whether uncontrolled factors, such as nutrition, food insecurity or physical activity mediate the association between increasing BMI and T cell differentiation in HIV-infection also remains unclear. Research in HIV-uninfected adults has shown that obesity is associated with a reduction in circulating regulatory T cells<sup>35</sup>. Since these cells serve to suppress immune responses and exert a suppressive function on effector T cells,<sup>36</sup> further research is necessary to determine if these regulatory cells mediate the association noted in our analysis. Furthermore, whether weight loss, among obese HIV-infected women, alters T cell differentiation warrants further exploration.

We did not find a significant association between CD57 expression and obesity when using a dichotomous variable comparing obese versus non-obese. However, the median BMI in our patient population was 30 m<sup>2</sup>/kg, and 75% were overweight or obese, and therefore lack of a significant association with obesity may reflect homogeneity across the study population. Nevertheless, whether increased adipokine-mediated inflammation independently alters CD8+ effector T cell populations in morbidly obese individuals is unclear. Further prospective studies may be helpful to better understand the interplay of weight gain, innate immune activation pathways and CD8+ T cell phenotypes in HIV-infected adults.

Contrary to expectation, there was an association between higher current CD4 count and a lower proportion of CD28–CD8+ cells expressing CD57. However, higher current CD4 count was also associated with a lower proportion of CD28–CD8+ T cells, an unexpected finding that may have been driven by unmeasured confounders. Whether other factors, such as co-morbid diseases, have a confounding effect on the association is unclear and warrants further evaluation.

Our study had several limitations. We did not include an HIV-uninfected control group for comparison and cannot comment on how associations between BMI and CD8+ T cell subtypes would be different in age-matched HIV-uninfected women. Furthermore, the cross-sectional design did not allow us to make conclusions on causality. Longitudinal studies relating changes in adaptive immunity to changes in body habitus are required, and such studies are currently underway. We acknowledge that BMI is an imprecise predictor that does not measure total fat mass or fat distribution, although monitoring BMI in a large sample size does provide insight into population characteristics<sup>37</sup>. Aerobic fitness is associated with lower age-related accumulation of terminally differentiated T cells in HIV-uninfected adults,<sup>38</sup> but we did not have available data on physical activity concurrent with our outcomes, and hence were unable to determine whether physical activity plays an independent role in adaptive immunity. Furthermore, we were unable to control for CMV infection, an important independent determinant of T cell differentiation, and increased CD57 expression<sup>39</sup>. While, we expect that most study participants were CMV co-infected<sup>30</sup>, we have no reason to suspect that CMV serostatus would have confounded the association of BMI with CD57 expression found in our analysis.

There were also many strengths to our study. We were able to conduct this work with ethnically diverse HIV-infected women in the United States. Research is fundamentally lacking in this group, and research in a wide-variety of settings will only improve our understanding of how ART alters adaptive immunity. The population was restricted to women, on antiretroviral therapy who mostly had suppressed viral loads, and excluded women with concomitant viral hepatitis, autoimmune disease and/or cancer – such restrictions reduced concerns about confounding from gender, co-morbid illness, or differential treatment –that might independently influence inflammation and T cell differentiation. The cohort size for this particular study, with immunological parameters measured on each participant, was relatively large, much more so than work that has been previously done. Furthermore, it included woman most of whom had been on ART for many years. Finally, our work is largely consistent with an evolving body of literature regarding the role of CD28–CD8+ T cells in HIV immunology.

In conclusion, in a relatively large cohort of HIV-infected women on ART, we found that greater BMI is associated with greater expression of CD57 on the proportion of CD28–CD8+ T cells, and increase in the proportion of CD57+CD28–CD8+ T cells, but not an accumulation of the overall CD28–CD8+ T cell subset. The impact of body composition on CD8+ T cell phenotype is complex in the setting of HIV infection given known effects of HIV on lean mass and subcutaneous tissue. Future research is needed using careful assessment of the various body composition compartments to understand the prognostic importance of these associations on clinical outcomes in HIV-infected individuals.

## Acknowledgments

**Support:** Part of this work was supported by R01-MH095683-01A1 to S.D.W. Data in this analysis were collected by the Women's Interagency HIV Study (WIHS). The contents of this analysis are solely the responsibility of the authors and do not represent the official views of the National Institutes of Health (NIH). WIHS (Principal Investigators): UAB-MS WIHS (Michael Saag, Mirjam-Colette Kempf, and Deborah Konkle-Parker), U01-AI-103401; Atlanta WIHS (Ighovwerha Ofotokun and Gina Wingood), U01-AI-103408; Bronx WIHS (Kathryn Anastos), U01-AI-035004; Brooklyn WIHS (Howard Minkoff and Deborah Gustafson), U01-AI-031834; Chicago

WIHS (Mardge Cohen and Audrey French), U01-AI-034993; Metropolitan Washington WIHS (Mary Young and Seble Kassaye), U01-AI-034994; Miami WIHS (Margaret Fischl and Lisa Metsch), U01-AI-103397; UNC WIHS (Adaora Adimora), U01-AI-103390; Connie Wofsy Women's HIV Study, Northern California (Ruth Greenblatt, Bradley Aouizerat, and Phyllis Tien), U01-AI-034989; WIHS Data Management and Analysis Center (Stephen Gange and Elizabeth Golub), U01-AI-042590; Southern California WIHS (Joel Milam), U01-HD-032632 (WIHS I – WIHS IV). The WIHS is funded primarily by the National Institute of Allergy and Infectious Diseases (NIAID), with additional co-funding from the Eunice Kennedy Shriver National Institute of Child Health and Human Development (NICHD), the National Cancer Institute (NCI), the National Institute on Drug Abuse (NIDA), and the National Institute on Mental Health (NIMH). Targeted supplemental funding for specific projects is also provided by the National Institute of Dental and Craniofacial Research (NIDCR), the National Institute on Alcohol Abuse and Alcoholism (NIAAA), the National Institute on Deafness and other Communication Disorders (NIDCD), and the NIH Office of Research on Women's Health. WIHS data collection is also supported by UL1-TR000004 (UCSF CTSA) and UL1-TR000454 (Atlanta CTSA). S.M.B. and M.J.A.R. were both supported by the UCSF Traineeship in AIDS Prevention Studies (US National Institutes of Health (NIH) T32 MH-19105). The funders did not play a role in study design, collection, analysis, interpretation of data, writing the report or the decision to submit the report for publication. PCT was also supported by K24 AI108516

We thank the Women's Interagency HIV Study (WIHS) participants who contributed data to this study. Data were collected by the WIHS Collaborative Study Group with centers (Principal Investigators at the time of data collection) at New York City/Bronx Consortium (Kathryn Anastos, MD); Brooklyn, NY (Howard Minkoff, MD); Washington DC, Metropolitan Consortium (Mary Young, MD); The Connie Wofsy Study Consortium of Northern California (Ruth Greenblatt, MD, Phyllis Tien, MD, and Bradley Aouizerat, PhD, MAS); Los Angeles County/Southern California Consortium (Alexandra Levine, MD); Chicago Consortium (Mardge Cohen, MD); and Data Coordinating Center (Stephen J. Gange, PhD). The contents of this publication are solely the responsibility of the authors and do not necessarily represent the official views of the US National Institutes of Health. We also thank Dr. Sulggi Lee for participating in helpful scientific discussion at the inception of this study.

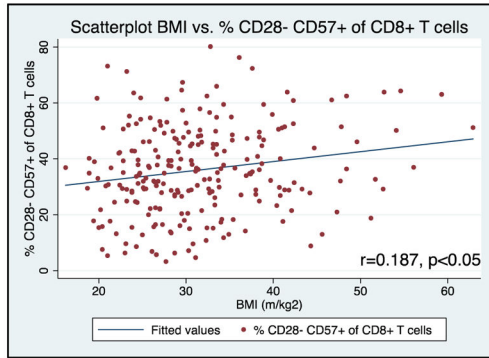
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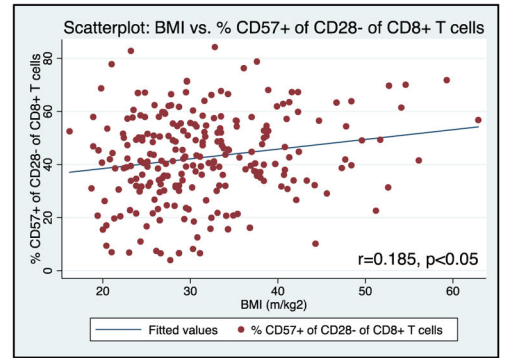
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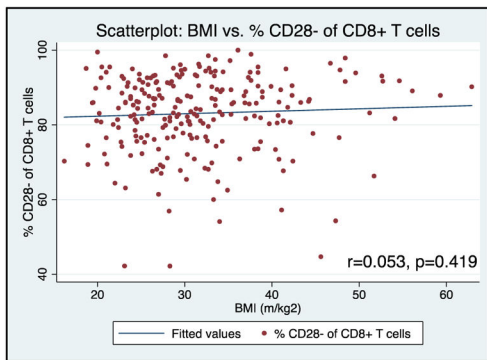
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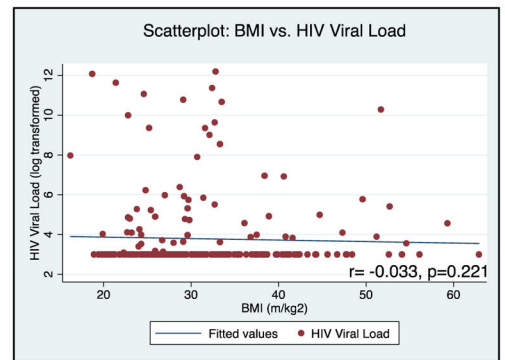
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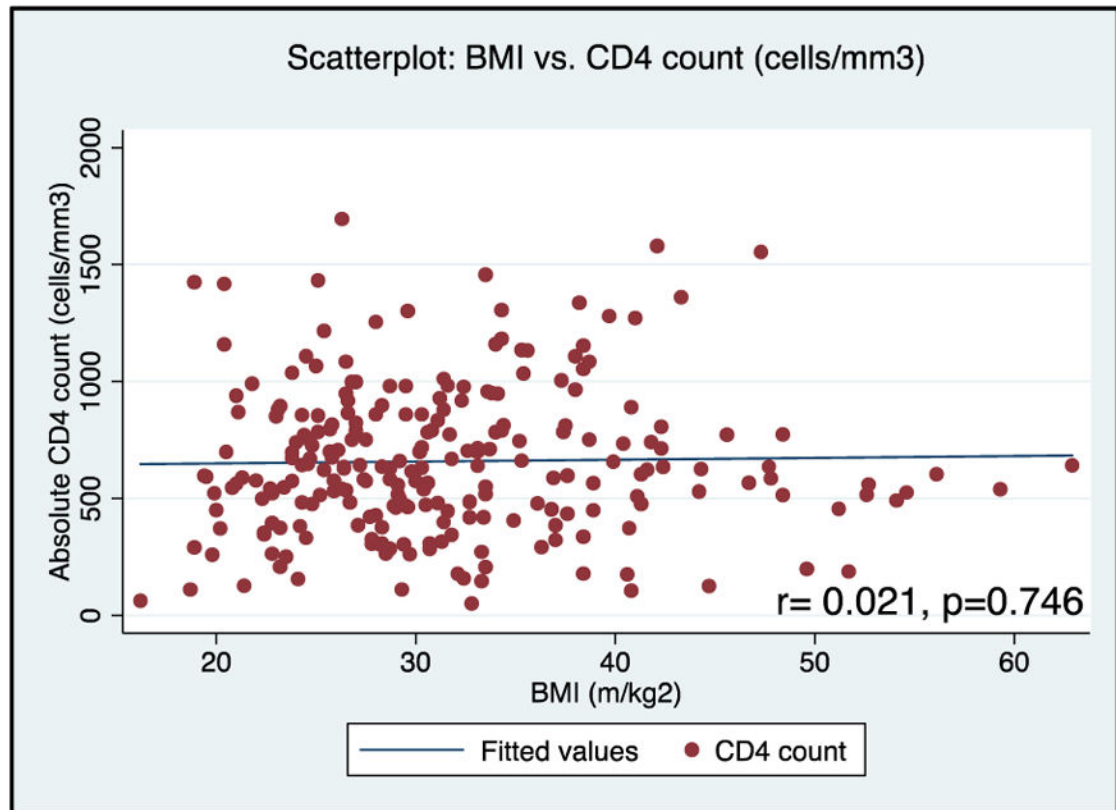
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**Figure 1. Scatterplots of BMI vs. CD8+ T cell phenotypes in HIV-infected women**

Figure 1a. BMI vs. % CD28<sup>-</sup> CD57<sup>+</sup> of CD8<sup>+</sup> T cells

Figure 1b. BMI vs. % CD57<sup>+</sup> of CD28<sup>-</sup> of CD8<sup>+</sup> T cells

Figure 1c. BMI vs. % CD28<sup>-</sup> of CD8<sup>+</sup> T cells

Figure 1e. BMI vs. HIV viral load

Figure 1d. BMI vs. current CD4<sup>+</sup> T cells

Demographic and clinical characteristics among 225 women with Human Immunodeficiency Virus (HIV) infection

**Table 1**

Variables	N (%) <sup>*</sup>
<b>Demographic Variables</b>	
Age in years (IQR)	46 (41, 51)
Site	
Bronx	28 (12%)
Brooklyn	82 (34%)
Georgetown	59 (24%)
San Francisco	11 (4%)
Chicago	62 (26%)
Race	
Non-Hispanic White	25 (10%)
Black/African American	173 (71%)
Hispanic	32 (13%)
Other	11 (6%)
Income	
\$30,000	144 (63%)
>\$30,001	84 (37%)
Education	
Less than high school	79 (33%)
Completed high school	71 (29%)
Some college or greater	91 (38%)
Employment status	
Employed	132 (55%)
Unemployed	109 (45%)
Lifestyle Variables	
Current Smoker	71 (29%)
Smoking history, (years)Median (IQR)	6.5 (0-24)



Variables	N (%) <sup>*</sup>
<b>Alcohol consumption</b>	
None	126 (53%)
>0-7 drinks/wk	100 (41%)
>7-12 drinks/wk	5 (2%)
>12 drinks/wk	9 (4%)
<b>Body Composition Variables</b>	
<b>Hip circumference, cm (IQR)</b>	108 (14.65)
<b>Waist circumference, cm (IQR)</b>	98 (88, 112)
<b>Height, cm (IQR)</b>	162 (158, 167)
<b>BMI (m/kg<sup>2</sup>) (IQR)</b>	30 (25, 36)
<b>HIV parameters</b>	
<b>Current CD4 count, median (IQR)</b>	626 (456-834)
<b>Nadir CD4 count, median (IQR)</b>	310 (173, 461)
<b>Log Viral Load, median (IQR)</b>	3.00 (3.0, 3.18)
<b>Viral suppression</b>	178 (74%)
<b>Duration on HAART, yrs (IQR)</b>	13 (6, 15)

\* Data are median (IQR) values, unless otherwise indicated.

Abbreviations: BMI, body mass index.

**Table 2**

Unadjusted and multivariable analysis factors associated with CD8+ T cell phenotypes in HIV-infected women

	% CD28- CD57+ of CD8+ T cells <sup>††</sup>			% CD57+ of CD28- of CD8+ T cells <sup>††</sup>			% CD28- of CD8+ T cells <sup>††</sup>		
	Unadjusted		Adjusted	Unadjusted		Adjusted	Unadjusted		Adjusted
	(95% CI)	p-value	(95% CI)	(95% CI)	p-value	(95% CI)	(95% CI)	p-value	(95% CI)
Age at visit	0.104 (-0.02, 0.228)	0.10	0.134 (0.009, 0.259)	0.135 (0.017, 0.253)	0.025	0.154 (0.037, 0.272)	-0.104 (-0.226, 0.018)	0.094	-0.068 (-0.198, 0.062)
Annual Income									
<=\$30,000	Ref.		Ref.	Ref.		Ref.	Ref.		Ref.
\$30,001+	-0.016 (-0.222, 0.189)	0.89	0.112 (-0.121, 0.344)	-0.016 (-0.211, 0.179)	0.87	0.141 (-0.078, 0.360)	-0.05 (-0.256, 0.155)	0.63	-0.068 (-0.31, 0.174)
Site									
Bronx	Ref.		Ref.	Ref.		Ref.	Ref.		Ref.
Brooklyn	-0.177 (-0.507, 0.152)	0.29	-0.111 (-0.456, 0.234)	-0.279 (-0.592, 0.035)	0.081	-0.169 (-0.493, 0.156)	0.307 (-0.013, 0.628)	0.60	0.193 (-0.166, 0.552)
Georgetown	0.02 (-0.325, 0.365)	0.91	0.092 (-0.28, 0.464)	-0.126 (-0.455, 0.203)	0.45	-0.015 (-0.365, 0.335)	0.437 (0.101, 0.773)	0.011	0.342 (-0.045, 0.729)
San Francisco	-0.18 (-0.716, 0.355)	0.51	-0.085 (-0.629, 0.459)	-0.409 (-0.919, 0.101)	0.12	-0.268 (-0.78, 0.243)	0.714 (0.193 - 1.235)	0.007	0.624 (0.058 - 1.189)
Chicago	-0.218 (-0.560 - 0.125)	0.21	-0.111 (-0.466 - 0.244)	-0.347 (-0.673 - -0.021)	0.037	-0.198 (-0.532 - 0.137)	0.411 (0.077, 0.744)	0.016	0.312 (-0.058, 0.681)
Education									
<High School	Ref.		Ref.	Ref.		Ref.	Ref.		Ref.
High School Degree	-0.05 (-0.295, 0.194)	0.69	-0.039 (-0.291, 0.212)	-0.039 (-0.271, 0.194)	0.74	-0.034 (-0.27, 0.203)	-0.049 (-0.292, 0.193)	0.69	-0.056 (-0.297, 0.225)
College or greater	-0.221 (-0.451, 0.009)	0.06	-0.082 (-0.341, 0.177)	-0.268 (-0.486, -0.049)	0.017	-0.155 (-0.399, 0.089)	0.053 (-0.175, 0.281)	0.65	0.153 (-0.117, 0.423)
Race/Ethnicity									
Non-Hispanic White	Ref.		Ref.	Ref.		Ref.	Ref.		Ref.
Black	0.373 (0.053 - 0.693)	0.022	0.27 (-0.088 - 0.627)	0.375 (0.069 - 0.680)	0.016	0.265 (-0.071 - 0.602)	0.129 (-0.190 - 0.447)	0.43	0.128 (-0.244 - 0.499)
Hispanic	0.408 (0.009, 0.807)	0.045	0.453 (0.01, 0.896)	0.45 (0.069, 0.831)	0.021	0.444 (0.027, 0.861)	0.048 (-0.349, 0.445)	0.81	0.19 (-0.271, 0.650)
Other	0.555 (0.014, 1.095)	0.045	0.51 (-0.030, 1.051)	0.507 (-0.010, 1.023)	0.055	0.458 (-0.051, 0.966)	0.324 (-0.215, 0.862)	0.24	0.322 (-0.241, 0.884)
Body Composition Parameters									
BMI <sup>††</sup>	0.184 (0.06 - 0.307)	0.004	0.166 (0.040 - 0.292)	0.175 (0.056 - 0.293)	0.004	0.145 (0.026 - 0.263)	0.051 (-0.073 - 0.175)	0.42	0.078 (-0.053 - 0.209)
Height (cm) <sup>††</sup>	0.031 (-0.103, 0.165)	0.650	0.02 (-0.122, 0.162)	0.019 (-0.109, 0.147)	0.77	0.028 (-0.105, 0.161)	0.037 (-0.095, 0.169)	0.58	-0.014 (-0.161, 0.134)
Waist Circumference <sup>††</sup> (cm)	0.139 (-0.009, 0.287)	0.065		0.15 (0.009, 0.290)	0.037		-0.031 (-0.175, 0.113)	0.67	
Hip Circumference <sup>††</sup> (cm)	0.177 (0.058, 0.296)	0.004		0.165 (0.052, 0.278)	0.005		0.062 (-0.055, 0.179)	0.30	
HIV Specific Parameters									
CD4 count	-0.241 (-0.354, -0.128)	<0.001	-0.188 (-0.312, -0.064)	-0.215 (-0.324, -0.107)	<0.001	-0.17 (-0.287, -0.054)	-0.16 (-0.273, -0.047)	0.006	-0.126 (-0.254, 0.003)

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	% CD28- CD57+ of CD8+ T cells <sup>§</sup>				% CD57+ of CD28- of CD8+ T cells <sup>§</sup>				% CD28- of CD8+ T cells <sup>§</sup>			
	Unadjusted		Adjusted		Unadjusted		Adjusted		Unadjusted		Adjusted	
	(95% CI)	p-value	(95% CI)	p-value	(95% CI)	p-value	(95% CI)	p-value	(95% CI)	p-value	(95% CI)	p-value
Viral Load (log <sub>10</sub> ) <sup>¶</sup>	0.014 (0.005, 0.023)	0.002	0.008 (-0.002, 0.018)	0.10	0.012 (0.004, 0.021)	0.006	0.007 (-0.003, 0.016)	0.16	0.011 (0.002, 0.020)	0.021	0.006 (-0.004, 0.017)	0.24
CD4 Nadir <sup>¶</sup>	0.083 (-0.037, 0.204)	0.18			0.073 (-0.042, 0.189)	0.21			0.069 (-0.050, 0.188)	0.25		
Duration of HAART <sup>¶</sup>	-0.243 (-0.404, -0.082)	0.003			-0.158 (-0.314, -0.002)	0.047			-0.373 (-0.528, -0.218)	<0.001		

Data are from generalized linear regression analyses.

<sup>§</sup>Values standardized by dividing individual value by Interquartile Range

Abbreviations: BMI, body mass index; HAART, highly active antiretroviral therapy

Factors associated with associated with CD8+ T cell phenotypes in HIV-infected women, in (1) multivariable analysis including and (2) multivariate analysis with additional adjustment for viral load.

**Table 3**

	% CD28- CD57+ of CD8+ T cells				% CD57+ of CD28- of CD8+ T cells				% CD28- of CD8+ T cells			
	Fully adjusted $\beta$ (95% CI)	p-value	Fully adjusted + viral load $\beta$ (95% CI)	p-value	Fully adjusted $\beta$ (95% CI)	p-value	Fully adjusted + viral load $\beta$ (95% CI)	p-value	Fully adjusted $\beta$ (95% CI)	p-value	Fully adjusted + viral load $\beta$ (95% CI)	p-value
BMI <sup>ab</sup>	0.161 (0.035, 0.287)	0.012	0.166 (0.041, 0.292)	0.01	0.141 (0.022, 0.259)	0.02	0.145 (0.026, 0.263)	0.017	0.074 (-0.057, 0.204)	0.27	0.0778 (-0.053, 0.209)	0.24
Waist Circumference, cm <sup>ab</sup>	0.168 (0.0160 - 0.321)	0.031	0.167 (0.0148 - 0.320)	0.032	0.164 (0.0222 - 0.306)	0.024	0.163 (0.0210 - 0.305)	0.025	0.0162 (-0.139 - 0.171)	0.84	0.0147 (-0.141 - 0.170)	0.85
Obesity <sup>a</sup>	0.185 (-0.012, 0.382)	0.065	0.185 (-0.011, 0.382)	0.065	0.151 (-0.0341, 0.336)	0.11	0.151 (-0.034, 0.335)	0.11	0.0886 (-0.114, 0.291)	0.39	0.0886 (-0.114, 0.292)	0.391
Waist Hip Ratio, cm <sup>ab</sup>	0.0289 (-0.142, 0.199)	0.74	0.0236 (-0.147, 0.194)	0.79	0.0482 (-0.110, 0.207)	0.55	0.0437 (-0.115, 0.202)	0.59	-0.0864 (-0.257, -0.0845)	0.32	-0.0907 (-0.262, 0.081)	0.3

<sup>a</sup>Model adjusting for age, race/ethnicity, income, education, WIHS site, height, CD4 count, and viral load

<sup>b</sup>Values standardized by dividing individual value by Interquartile Range