

Metabolomic Profiling of Left Ventricular Diastolic Dysfunction in Women With or at Risk for HIV Infection: The Women's Interagency HIV Study

Claudio A. Bravo, MD; Simin Hua, MHS; Amy Deik, BA; Jason Lazar, MD, MPH; David B. Hanna, PhD; Justin Scott, BS; Jin Choul Chai, PhD; Robert C. Kaplan, PhD; Kathryn Anastos, MD; Octavio A. Robles, BS; Clary B. Clish, PhD; Jorge R. Kizer, MD, MSc; Qibin Qi, PhD

Background—People living with HIV have an increased risk of left ventricular diastolic dysfunction (LVDD) and heart failure. HIV-associated LVDD may reflect both cardiomyocyte and systemic metabolic derangements, but the underlying pathways remain unclear.

Methods and Results—To explore such pathways, we conducted a pilot study in the Bronx and Brooklyn sites of the WIHS (Women's Interagency HIV Study) who participated in concurrent, but separate, metabolomics and echocardiographic ancillary studies. Liquid chromatography tandem mass spectrometry–based metabolomic profiling was performed on plasma samples from 125 HIV-infected (43 with LVDD) and 35 HIV-uninfected women (9 with LVDD). Partial least squares discriminant analysis identified polar metabolites and lipids in the glycerophospholipid-metabolism and fatty-acid-oxidation pathways associated with LVDD. After multivariable adjustment, LVDD was significantly associated with higher concentrations of diacylglycerol 30:0 (odds ratio [OR], 1.60, 95% CI [1.01–2.55]); triacylglycerols 46:0 (OR 1.60 [1.04–2.48]), 48:0 (OR 1.63 [1.04–2.54]), 48:1 (OR 1.62 [1.01–2.60]), and 50:0 (OR 1.61 [1.02–2.53]); acylcarnitine C7 (OR 1.88 [1.21–2.92]), C9 (OR 1.99 [1.27–3.13]), and C16 (OR 1.80 [1.13–2.87]); as well as lower concentrations of phosphocholine (OR 0.59 [0.38–0.91]). There was no evidence of effect modification of these relationships by HIV status.

Conclusions—In this pilot study, women with or at risk of HIV with LVDD showed alterations in plasma metabolites in the glycerophospholipid-metabolism and fatty-acid-oxidation pathways. Although these findings require replication, they suggest that improved understanding of metabolic perturbations and their potential modification could offer new approaches to prevent cardiac dysfunction in this high-risk group. (*J Am Heart Assoc.* 2020;9:e013522. DOI: 10.1161/JAHA.119.013522.)

Key Words: heart failure • HIV • left ventricular diastolic dysfunction • metabolomics

Progress in delineating risk factors for HIV infection, along with advances in treatment, have redefined the contemporary epidemiology of this condition.^{1–4} Incidence of HIV infection has been declining over time, yet the number of people living with HIV continues to rise, driven by the markedly improved survival conferred by antiretroviral therapy (ART).^{1–4} With ART's transformation of HIV infection into a chronic condition has come the recognition that people living

with HIV appear to be susceptible to a range of common aging-related disorders.^{4–6}

People living with HIV have been reported to have a higher risk of cardiovascular disease than uninfected individuals.⁷ Apart from a well-documented increased risk of myocardial infarction in persons with HIV,^{8,9} available evidence also points to a higher risk of heart failure in the absence of previous coronary artery disease,¹⁰ including heart failure with preserved

From the Division of Cardiology, Department of Medicine, Columbia University Medical Center, New York, NY (C.A.B.); Department of Epidemiology & Population Health, Albert Einstein College of Medicine, Bronx, NY (S.H., D.B.H., J.C.C., R.C.K., K.A., Q.Q.); Metabolomics Platform, Broad Institute of MIT and Harvard, Cambridge, MA (A.D., J.S., C.B.C.); Division of Cardiovascular Medicine, State University of New York Downstate Medical Center, Brooklyn, NY (J.L.); Public Health Sciences Division, Fred Hutchinson Cancer Research Center, Seattle, WA (R.C.K.); Department of Medicine, Albert Einstein College of Medicine, Bronx, NY (K.A.); Department of Biologic Sciences, Universidad de Chile, Santiago, Chile (O.A.R.); Division of Cardiology, San Francisco Veterans Affairs Health Care System, University of California San Francisco, San Francisco, CA (J.R.K.).

Accompanying Data S1, Tables S1 through S3, and Figures S1 and S2 are available at <https://www.ahajournals.org/doi/suppl/10.1161/JAHA.119.013522>

Correspondence to: Qibin Qi, PhD, Department of Epidemiology and Population Health, Albert Einstein College of Medicine, 1300 Morris Park Ave, Bronx, NY 10461. E-mail: qibin.qi@einstein.yu.edu

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Clinical Perspective

What Is New?

- This is the first investigation of metabolomic perturbations associated with left ventricular diastolic dysfunction in the context of HIV infection.

What Are the Clinical Implications?

- These findings highlight the potential role of impaired fatty acid metabolism, mitochondrial dysfunction, and increased oxidative stress in early-stage cardiac dysfunction.
- Pending replication, these findings suggest that further work to understand the pathways involved and testing pharmacological interventions could lead to new strategies to prevent heart failure, both among women with or at risk for HIV and more broadly.

ejection fraction.¹¹ Consistent with this finding, a meta-analysis of echocardiographic studies of HIV-positive individuals receiving ART (mean age, 41 years) showed a prevalence of left ventricular (LV) diastolic dysfunction (LVDD) of 43%,¹² far exceeding the prevalence of 8% reported in an older (mean age, 63 years) HIV-negative population.¹³ More-recent classification schemes used more-stringent criteria for LVDD, leading to lower prevalences,¹⁴ but among mostly middle-aged participants with or at risk for HIV in the WIHS (Women's Interagency HIV Study), frequencies of LVDD based on the latest guidelines^{15,16} were still substantial at 19% to 29%.¹⁷

The high prevalence of HIV-related LVDD reflects an increased burden of subclinical myocardial disease, with increased myocardial fibrosis and cardiomyocyte steatosis documented by magnetic resonance imaging in persons with HIV.¹⁸ The precise basis for such myocardial disease/dysfunction has not been determined, but various factors are likely implicated, including chronic inflammation and immune activation, adverse effects of ART, or increased behavioral and clinical risk factors for cardiovascular diseases.⁷

Women with or at risk for HIV have a high burden of obesity and metabolic dysregulation,¹⁹ which along with inflammation and oxidative stress can alter systemic and myocardial energetics.²⁰ In this context, the advent of metabolomics allowing untargeted assessment of a wide array of metabolic intermediates offers the opportunity to probe biochemical pathways involved in cardiac dysfunction.²¹ Given that LVDD reflects early myocardial functional derangements that can eventuate in clinical heart failure,²² and is of particular relevance to women and race/ethnic minorities,^{23,24} evaluating the metabolomic signatures of LVDD could lead to improved risk stratification, novel mechanistic insights, and potential new prevention

approaches for this important condition.²¹ In this pilot study, we leveraged availability of plasma metabolomics and lipidomics measured among women who completed echocardiograms in a substudy of the WIHS^{25,26} to investigate the plasma metabolite profiles of LVDD in this high-risk cohort.

Methods

The data that support the findings of this study are available from the corresponding author upon reasonable request.

Participants

The study was conducted in a sample of participants enrolled in the WIHS, an ongoing multicenter study of the natural history of HIV disease in women.^{25,26} Women in the present analysis were drawn from those participating in 2 WIHS ancillary studies, an echocardiographic study performed in the Bronx and Brooklyn sites in 2004–2005,²⁷ and a metabolomic study performed in nondiabetic participants with available carotid sonograms.²⁸ Relevant demographic, behavioral, anthropometric, and clinical information was obtained from the included participants, as previously reported.²⁸

The study was approved by the institutional review boards of both study sites, and all subjects provided written informed consent.

Echocardiography Procedures

Echocardiogram acquisition followed a standardized protocol involving 2-dimensional, M-mode, spectral, tissue-Doppler, and color-Doppler imaging, as previously reported.²⁷ Trans-mitral diastolic velocities were obtained by placing the sample volume at the leaflet tips. Tissue Doppler imaging was performed at the medial mitral annulus. Data were recorded on VHS tapes and analyzed by a single experienced echocardiographer (J.L.). LV wall thicknesses and internal diameters were determined in accordance with the American Society of Echocardiography (ASE) guidelines.²⁹ LV ejection fraction was calculated using the biplane method of discs from the apical-4-chamber and apical-2-chamber views and confirmed by visual estimation. Left atrial volumes were determined by the biplane method of discs.

For our primary evaluation of LV diastolic function, we used the 2009 ASE/European Association of Cardiovascular Imaging (EACVI) guidelines,¹⁶ applying the 2016 ASE/EACVI guidelines in a sensitivity analysis.¹⁵ This approach was chosen to enhance identification of metabolomic profiles associated with earlier and milder stages of LVDD occurring in our younger population, detectable with the less-restrictive criteria formulated by the 2009 ASE/EACVI classification.

The 2009 ASE/EACVI guidelines define LVDD based on septal and lateral mitral annular early diastolic (e') velocities and left atrial volume. We applied the 2009 ASE/EACVI criteria incorporating the presence of left atrial enlargement (≥ 34 mL/m²) and either septal e' velocity < 8 cm/s or lateral e' velocity < 10 cm/s to define LVDD, but modified these to include only septal e' because lateral e' was not obtained.¹⁷ Similarly, we adapted the 2016 ASE/EACVI guidelines¹⁵ to define LVDD as the presence of > 2 of the following echocardiographic findings: (1) early mitral inflow (E)/septal e' ratio > 15 ; (2) septal e' velocity < 7 cm/s; (3) tricuspid regurgitation velocity > 2.8 m/s; and (4) left atrial volume > 34 mL/m². For the 2016 ASE/EACVI guidelines, cases with indeterminate diastolic function were reclassified as LVDD if LV end-diastolic volume index was > 61 mL/m² or LV mass index was > 95 g/m². Grading of LVDD was done in accordance with each set of guidelines.^{15,17} For both LVDD definitions, we included all grades of LVDD.

Metabolomics

Fasting plasma samples were sent to the Broad Institute Metabolomics Platform (Cambridge, MA), where metabolomic technology was used to quantitatively profile metabolites. Metabolomic analysis was performed utilizing tandem liquid chromatography/mass spectrometry. Liquid chromatography affords reproducible separation of metabolites on the basis of their physical properties, and mass spectrometry enables further resolution of metabolites on the basis of mass-to-charge ratio (m/z). Two separate liquid chromatography/mass spectrometry methods were performed to measure polar metabolites and lipids in each sample, as previously described (Data S1).^{28,30,31} Raw data from Orbitrap mass spectrometers were processed using Progenesis Q1 software (NonLinear Dynamics, Durham, NC) for feature alignment, untargeted signal detection, and signal integration. Metabolites were quantified using area under the curve of the peaks. The targeted processing of a subset of known metabolites was conducted using TraceFinder software (version 3.1; Thermo Fisher Scientific; Waltham, MA). In the current analysis, a total of 325 metabolites (114 polar metabolites and 211 lipids) were included after excluding metabolites with coefficient of variation of $\geq 20\%$ or percentage of missing values $\geq 20\%$.

Statistical Analysis

Continuous variables were summarized with the mean and SD, if the data were normally distributed, and the median and interquartile range otherwise. Means were compared utilizing the parametric 2-tailed Student t test, and comparisons between medians were performed using the nonparametric

McNemar test. The chi-squared test was performed to compare proportions, and Fisher's exact test was utilized when the number in a cell was < 5 in the 2×2 table. Given that the data obtained from metabolomics are semiquantitative, that is, the values are not the actual levels with a unit, but are instead relative levels, we applied rank-based inverse normal transformation to all plasma polar metabolites and lipids to transform the measurements to a normal distribution. Given that the number of polar metabolites and lipids was greater than the number of participants in this study, we used the dimension-reduction technique, partial least squares discriminant analysis (PLS-DA), to identify polar metabolites and lipids with high contributions to group separation. Because the polar metabolites and lipids were analyzed in different platforms, we performed PLS-DA separately for each group of metabolites. Thereafter, we selected the top 10 polar metabolites and top 10 lipids with the highest loading coefficients and variable importance projection scores of the first principal component, using the mixOmics package (mixomics.org) for further analyses. We assessed the associations between these polar metabolites and lipids with the presence of LVDD using logistic regression, adjusting for age, race/ethnicity, level of education, smoking or drinking history, hepatitis C virus infection, and history of intravenous drug use (model 1); HIV status, baseline viral load, and ART status (included in model 2, in addition to variables in model 1); and total cholesterol, high-density lipoprotein cholesterol, systolic blood pressure, homeostasis model assessment of insulin resistance, and current use of lipid-lowering or antihypertensive medication (included in model 3, in addition to variables in model 2). The interpretation was the odds ratio of LVDD per SD of plasma polar metabolite and lipid levels. We tested the difference by HIV status in associations between polar metabolites/lipids and LVDD by adding an interaction term of polar metabolites/lipids and HIV status in the logistic models and reported P for interaction.

In order to evaluate the joint associations of the top 10 polar metabolites in addition to the top 10 lipids, we examined the score of each metabolite in association with LVDD using 3 models. We generated a weighted score of the 20 selected metabolites by summing the weighted rank-based inverse normal transformed levels of these metabolites. The weight for each metabolite is the coefficient of the metabolite generated from logistic regression, with LVDD as the dependent variable and each metabolite and covariates in model 1 as independent variables.

All analyses were conducted using SPSS (version 15; SPSS, Inc, Chicago, IL) and R software (version 3.4.2; R Foundation for Statistical Computing, Vienna, Austria). Visualization of pathway mapping was performed in Cytoscape (version 3.7.1; MetScape application version 3.1.3).³² Because of the pilot nature of the study, statistical significance was defined as

$P < 0.05$ without correction for multiple testing. The Benjamini–Yekutieli procedure was applied to explore whether any associations met significance by controlling the study-wide false discovery rate at $q \leq 0.05$.

Results

Baseline Characteristics

Among 160 participants included in this study, 52 (32.5%) women had echocardiographic evidence of LVDD. Of these, 42 women had grade I and 10 had grade II LVDD. A total of

125 women were HIV positive, of whom 43 (34.4%) had evidence of LVDD; and among the 35 HIV-negative women, 9 (25.7%) had evidence of LVDD. A small minority of participants overall had LV systolic dysfunction, 4 (2.5%), without significant difference by HIV status. This and other pertinent information are listed in Table 1.

Metabolomic and Lipidomic Profiles With LVDD

PLS-DA showed that using the first and second principal components of metabolomic and lipidomic profiles, respectively, there were some distinctions between participants with

Table 1. Characteristics of Participants

	All	LVDD		P Values
		No	Yes	
No. of participants	160	108	52	
Demographic and behavioral				
Age, y	41.5 [39.0–46.0]	41.0 [39.0–45.0]	44.5 [39.0–51.0]	0.02
Race/ethnicity				0.53
Non-Hispanic black	108 (68%)	75 (69%)	33 (63%)	
Hispanic	47 (29%)	29 (27%)	18 (35%)	
Non-Hispanic white/other	5 (3%)	4 (4%)	1 (2%)	
Body mass index, kg/m ²	27.9 [24.6–31.5]	27.8 [24.6–31.3]	28.4 [24.6–31.8]	0.79
Waist circumference, cm	88.7 [81.3–97.2]	88.0 [80.7–95.5]	91.6 [82.6–100.1]	0.31
Hip circumference, cm	99.1 [93.4–108.0]	101.2 [93.4–108.0]	96.4 [92.9–109.3]	0.13
Current crack/cocaine use	9 (6%)	7 (6%)	2 (4%)	0.72
History of intravenous drug use	38 (24%)	20 (19%)	18 (35%)	0.03
Current smoker	87 (54%)	61 (56%)	26 (50%)	0.44
Alcohol abuse (>7 drinks per wk)	18 (11%)	10 (9%)	8 (15%)	0.25
Laboratory parameters				
Total cholesterol, mg/dL	180.6 (37.7)	182.5 (39.0)	176.6 (34.7)	0.35
LDL cholesterol, mg/dL	105.0 (31.0)	106.6 (33.5)	102.0 (25.1)	0.37
HDL cholesterol, mg/dL	48.7 (17.2)	50.1 (17.8)	45.6 (15.5)	0.12
Triglycerides, mg/dL	108.0 [74.0–144.0]	110.5 [74.0–140.0]	105.0 [75.0–152.0]	0.63
HOMA-IR	2.4 [1.6–3.6]	2.4 [1.7–3.6]	2.8 [1.5–3.8]	0.44
eGFR, mL/min/1.73 m ²	93.4 (21.6)	95.7 (20.6)	88.5 (23.0)	0.05
hs-CRP, μ g/mL	2.0 [0.8–4.9]	2.2 [0.8–4.9]	1.8 [0.8–5.8]	0.96
Comorbidities				
Systolic blood pressure, mm Hg	119.3 (19.0)	117.3 (16.3)	123.2 (23.2)	0.1
Diastolic blood pressure, mm Hg	74.5 (11.4)	73.6 (11.1)	76.3 (11.9)	0.17
Current antihypertensive medication use	31 (19%)	19 (18%)	12 (23%)	0.41
Current lipid-lowering medication use	4 (3%)	2 (2%)	2 (4%)	0.6
Hepatitis C virus infection	48 (30%)	25 (23%)	23 (44%)	0.006

Continued

Table 1. Continued

	All	LVDD		P Values
		No	Yes	
HIV and related factors				
HIV infection	125 (78%)	82 (76%)	43 (83%)	0.33
CD4 ⁺ count, cell/mm ³	429 [282–602]	469 [299–636]	367 [267–564]	0.23
HIV ⁺ viral load, copies/mL	260 [80–5000]	195 [80–4000]	540 [80–7700]	0.57
Current antiretroviral therapy use	100 (80%)	64 (78%)	36 (84%)	0.45
Echocardiographic parameters				
LV ejection fraction (%)	59.8 (6.2)	60.1 (5.8)	59.2 (7.1)	0.4
LV ejection fraction <50%	4 (2.5)	2 (1.9)	2 (3.9)	0.60
LV mass index, g/m ²	82.1 (17.7)	80.2 (17.4)	86.1 (17.7)	0.05
Left atrial volume index, mL/m ²	27.5 (6.5)	26.4 (4.7)	29.8 (8.8)	0.01
LV end-diastolic volume index, mL/m ²	45.8 (10.6)	45.6 (10.6)	46.0 (10.6)	0.85
LV early filling velocity, E (m/s)	0.8 [0.7–0.9]	0.8 [0.7–0.9]	0.7 [0.6–0.9]	0.04
LV late filling velocity, A (m/s)	0.7 [0.6–0.8]	0.6 [0.5–0.8]	0.7 [0.6–0.8]	0.01
E/A ratio	1.2 [0.9–1.4]	1.3 [1.1–1.5]	1.0 [0.8–1.2]	0.0007
Septal e' at mitral annulus, ms	0.10 [0.08–0.13]	0.11 [0.10–0.14]	0.07 [0.06–0.10]	<0.0001
E/e' ratio	8.0 [6.2–10.0]	7.1 [5.5–8.8]	10.0 [7.5–12.0]	<0.0001

eGFR indicates estimated glomerular filtration rate; HDL, high-density lipoprotein; HOMA-IR, homeostasis model assessment of insulin resistance; hs-CRP, high-sensitivity C-reactive protein; LDL, low-density lipoprotein; LV, left ventricular; LVDD, left ventricular diastolic dysfunction.

and without LVDD (Figure 1) and also in the presence or absence of HIV (Figure S1), but these distinctions were at a relatively low level. We then focused on the top 10 polar

metabolites and top 10 lipids in the first principal components selected by the PLS-DA (Table 2 and Table S1). None of these top polar metabolites or lipids showed significant associations

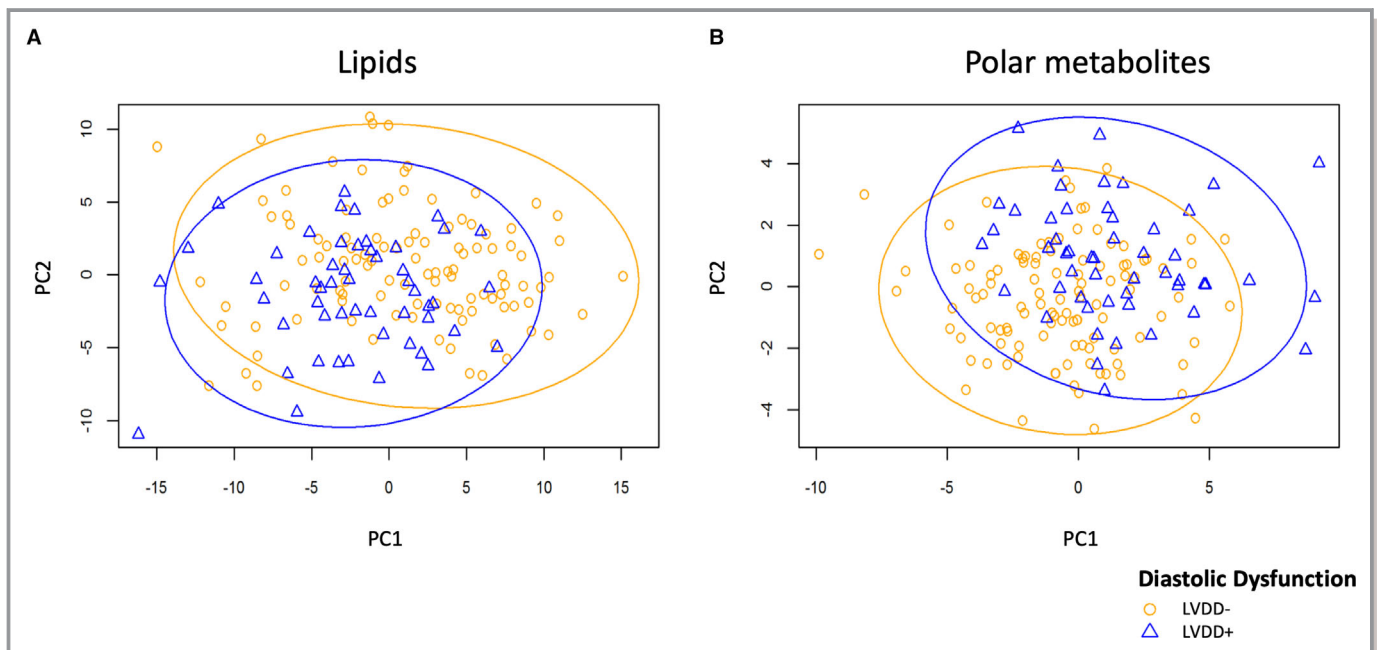


Figure 1. Scores plot of metabolites and lipids according to presence of LVDD. Scores plot showing metabolites from principal component 1 and 2 obtained with PLS-DA analysis of participants with LVDD (blue triangles and line) vs without LVDD (orange circles and line) for (A) lipids and (B) polar metabolites. LVDD indicates left ventricular diastolic dysfunction; PC, principal component; PLS-DA, partial least squares discriminant analysis.

Table 2. Relationship Between Top Polar Metabolites/Lipids Concentration and LVDD

	All		HIV ⁺	HIV ⁻	P-int*
	OR (95% CI)	P Value	OR (95% CI)	OR (95% CI)	
Polar metabolites					
Phosphocholine					
Model 1	0.59 (0.39–0.87)	0.008	0.56 (0.37–0.87)	0.65 (0.26–1.62)	0.26
Model 2	0.56 (0.37–0.85)	0.006	0.55 (0.35–0.86)	0.63 (0.25–1.59)	0.64
Model 3	0.60 (0.39–0.93)	0.022	0.52 (0.32–0.86)	1.06 (0.41–2.71)	0.92
C7-Carnitine					
Model 1	1.60 (1.09–2.36)	0.017	1.61 (1.04–2.50)	1.55 (0.68–3.54)	0.25
Model 2	1.62 (1.09–2.41)	0.018	1.63 (1.04–2.56)	1.58 (0.69–3.63)	0.38
Model 3	1.88 (1.18–2.98)	0.008	1.94 (1.16–3.25)	1.67 (0.68–4.06)	0.27
C9-Carnitine					
Model 1	1.73 (1.18–2.55)	0.006	1.96 (1.24–3.12)	1.23 (0.56–2.71)	0.35
Model 2	1.82 (1.21–2.72)	0.004	2.07 (1.28–3.35)	1.23 (0.56–2.72)	0.35
Model 3	1.96 (1.25–3.08)	0.004	2.22 (1.32–3.75)	1.27 (0.51–3.17)	0.62
C16-Carnitine					
Model 1	1.67 (1.13–2.48)	0.011	1.72 (1.10–2.70)	1.44 (0.58–3.60)	0.30
Model 2	1.73 (1.14–2.63)	0.010	1.81 (1.12–2.91)	1.47 (0.59–3.67)	0.37
Model 3	1.94 (1.18–3.21)	0.010	2.14 (1.22–3.75)	1.30 (0.47–3.63)	0.28
Lipids					
DAG 30:0					
Model 1	1.65 (1.10–2.49)	0.016	1.46 (0.93–2.30)	2.65 (1.05–6.73)	0.32
Model 2	1.74 (1.14–2.66)	0.011	1.56 (0.98–2.49)	2.64 (1.04–6.68)	0.28
Model 3	1.70 (1.05–2.76)	0.032	1.48 (0.87–2.53)	2.76 (1.03–7.37)	0.60
TAG 46:0					
Model 1	1.62 (1.08–2.41)	0.018	1.45 (0.94–2.23)	2.89 (1.06–7.90)	0.36
Model 2	1.69 (1.12–2.54)	0.012	1.51 (0.97–2.35)	2.89 (1.06–7.92)	0.34
Model 3	1.78 (1.10–2.87)	0.018	1.56 (0.93–2.62)	3.17 (1.09–9.23)	0.52
TAG 48:0					
Model 1	1.61 (1.08–2.40)	0.019	1.40 (0.91–2.14)	3.75 (1.28–11.00)	0.32
Model 2	1.69 (1.12–2.54)	0.012	1.46 (0.94–2.26)	3.76 (1.28–11.08)	0.37
Model 3	1.77 (1.08–2.88)	0.023	1.49 (0.89–2.51)	4.39 (1.32–14.63)	0.50
TAG 48:1					
Model 1	1.65 (1.10–2.49)	0.016	1.48 (0.96–2.27)	3.40 (1.09–10.59)	0.35
Model 2	1.70 (1.12–2.57)	0.012	1.53 (0.99–2.38)	3.39 (1.09–10.56)	0.32
Model 3	1.79 (1.06–3.03)	0.029	1.60 (0.93–2.77)	3.61 (1.06–12.32)	0.41
TAG 50:0					
Model 1	1.59 (1.08–2.35)	0.019	1.39 (0.91–2.12)	3.22 (1.16–8.96)	0.30
Model 2	1.65 (1.10–2.47)	0.015	1.44 (0.93–2.23)	3.25 (1.16–9.06)	0.33
Model 3	1.73 (1.05–2.83)	0.031	1.47 (0.86–2.50)	3.75 (1.19–11.74)	0.41

Regression was performed adjusted for age, race, education level, smoking, drinking habits, HCV infection, and intravenous drug use (model 1). Further adjustment was performed for HIV status, baseline viral load, and ART treatment status (model 2); and, additionally, for total cholesterol, HDL cholesterol, systolic blood pressure, HOMA-IR, lipid-lowering medication, and antihypertensive medication use (model 3). Metabolites underwent rank-based inverse normal transformation. The OR for LVDD for each 1-unit increase in the standardized plasma metabolite level (ie, per SD increase in the rank-based inverse-normal transformed level). ART indicates antiretroviral therapy; DAG indicates diacylglycerol; HCV, hepatitis C virus; HDL, high-density lipoprotein; HOMA-IR, homeostasis model assessment of insulin resistance; LVDD, left ventricular diastolic dysfunction; OR, odds ratio; TAG, triacylglycerol.

*P for interaction test.

with LVDD by the Benjamini–Yekutieli false discovery rate threshold.

Polar Metabolites

Of the top 10 polar metabolites that PLS-DA identified (Table S1), after adjustment for age, race/ethnicity, level of education, smoking or drinking history, hepatitis C virus infection, and history of intravenous drug use, 4 polar metabolites that belong to the glycerophospholipid-metabolism pathway (ie, phosphocholine) and the fatty-acid-oxidation

pathway (ie, acylcarnitines [C7, C9, and C16-carnitine]) were significantly associated with LVDD. The same polar metabolites were significantly different in participants with and without LVDD after additional adjustment for HIV-specific variables: HIV status, baseline viral load, and ART treatment status; and after further adjustment was performed using conventional cardiovascular risk factors: total cholesterol and high-density lipoprotein-cholesterol concentration, systolic blood pressure, and lipid-lowering and antihypertensive medication use (Table 2; Figure 2). According to a test for HIV-polar metabolite interaction, there was no significant

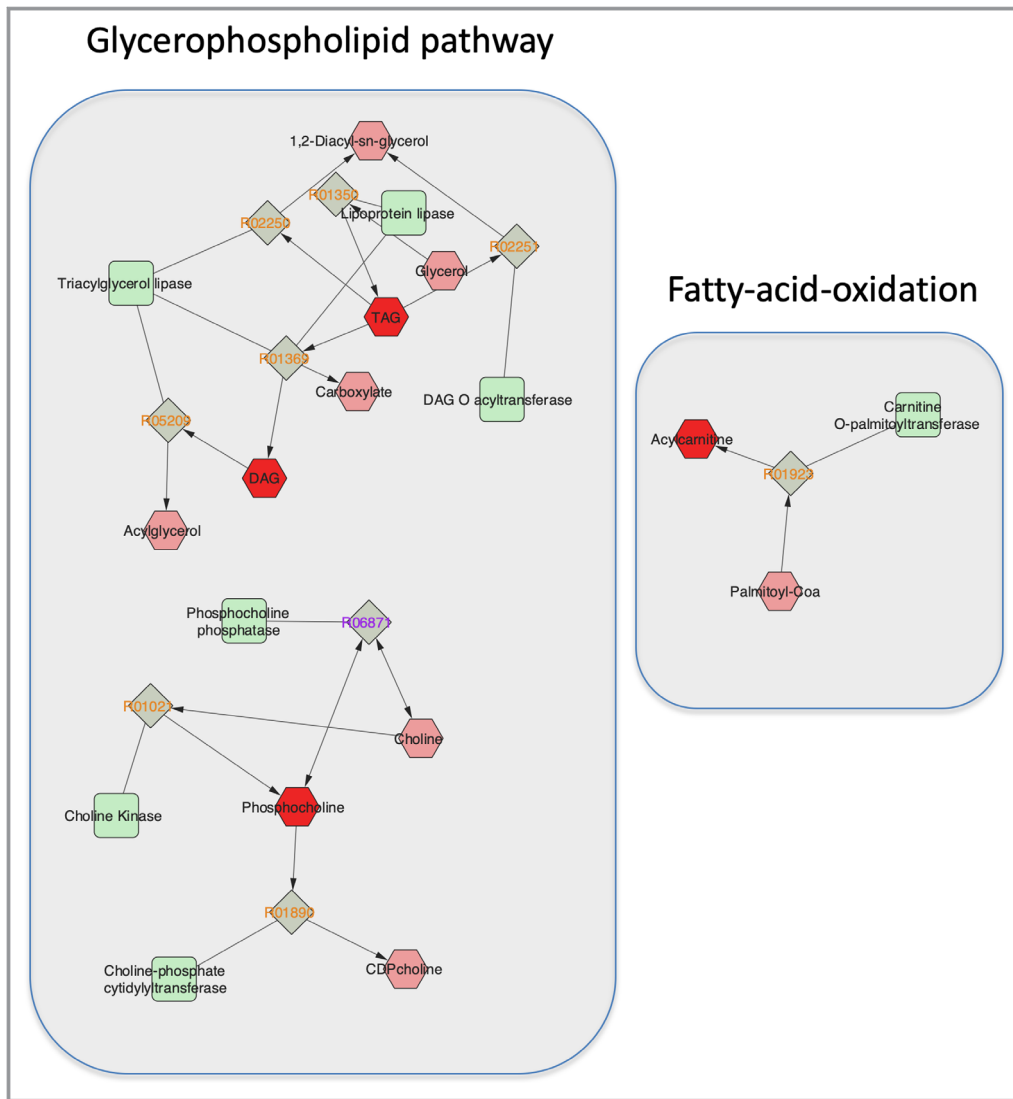


Figure 2. Glycerophospholipid and fatty-acid-oxidation pathways. Metabolic pathways constructed utilizing Cytoscape software (version 3.7.1; MetScape application version 3.1.3). The red-colored hexagons represent the individual metabolites and lipids that were associated with the presence of LVDD within the first principal component, and the light red hexagons are other relevant compounds of those pathways. Light green squares represent the enzymes, and gray rotated squares are the reactions (KEGG ID number) of key metabolic steps of those pathways. DAG indicates diacylglycerol; KEGG, Kyoto Encyclopedia of Genes and Genomes; LVDD, left ventricular diastolic dysfunction.

difference in the associations of these 4 polar metabolites with LVDD between HIV-positive and -negative participants (Table 2).

Lipids

Of the top 10 lipids that PLS-DA identified (Table S1), after adjustment for age, race/ethnicity, level of education, smoking or drinking history, hepatitis C virus infection, and history of intravenous drug use, 5 lipids that belong to the glycerophospholipid-metabolism pathway, that is, diacylglycerol (DAG; 30:0) and triacylglycerol (TAG; 46:0, 48:0, 48:1, and 50:0), were associated with LVDD. The same lipids were significantly different between participants with and without LVDD after additional adjustment for HIV-specific variables; and after further adjustment was performed using conventional cardiovascular risk factors (Table 2; Figure 2). No significant effect modification by HIV status was observed (Table 2).

Joint Associations of Polar Metabolites and Lipids

In an additional exploratory analysis of the joint associations involving the top 10 polar metabolites along with the top 10 lipids, we found that the score of the 20 metabolites was significantly related to an increased risk of LVDD (odds ratio, 1.25; 95% CI, 1.11–1.40; $P < 0.0001$). We did not find evidence of interaction for this association by HIV status (Table S2).

Sensitivity Analyses

In sensitivity analyses, we explored the associations of PLS-DA-identified polar metabolites and lipids above with LVDD defined using the 2016 ASE/EACVI guidelines,¹⁵ which identified 26 women with LVDD. After adjustment for age, race/ethnicity, level of education, smoking or drinking history, hepatitis C virus infection, and history of intravenous drug use, none of the polar metabolites were significantly associated with newly defined LVDD. After multivariable adjustment, 2 lipids that belong to the glycerophospholipid-metabolism pathway (ie, DAG 30:0 and TAG 50:0) were associated with the newly defined LVDD. However, the associations between lipids and LVDD did not remain statistically significant after additional adjustment for HIV-specific variables and conventional cardiovascular risk factors (Table S3).

Discussion

In this sample of early-middle-aged women with or at risk for HIV, mostly of black and Hispanic race/ethnicity and free of diabetes mellitus, there were no significant differences in

metabolite principal components between those with and without echocardiographic LVDD. Evaluation of the top 10 polar metabolites and top 10 lipids within the first principal component, however, showed that LVDD was associated with a lower concentration of phosphocholine and higher concentration of acylcarnitine, as well as higher concentrations of TAGs and DAG. Further exploration of the joint relationship of the top 10 polar metabolites and top 10 lipids with LVDD also showed that the corresponding metabolite score was significantly associated with this outcome. There was no evidence in our sample of effect modification of these associations by HIV status. Additional analysis using the least absolute shrinkage and selection operator method confirmed the findings obtained using PLS-DA.

To our knowledge, the present exploratory study is the first to evaluate the metabolomic signatures of LVDD in a sample of women with or at risk of HIV. Our investigation relied on the WIHS, a long-running study originally designed to characterize the natural history of HIV infection in women.^{25,26} The participants enrolled are at increased risk of cardiovascular diseases, exhibiting high frequencies of smoking, substance abuse, and other risk factors. These characteristics are commonly noted in people living with HIV, but as our sample shows, the WIHS counts as a strength that the population at risk for HIV included as a comparator has a similar frequency of risk behaviors and risk factors.³³

Previously, a metabolomic analysis of people with symptomatic heart failure with preserved ejection fraction of the HEART (Alberta Heart Failure Etiology and Analysis Research Team) project showed some similar metabolic features to those found in our study. Among the metabolomic differences uncovered, the study documented that patients with heart failure with preserved ejection fraction and heart failure with reduced ejection fraction had higher concentrations of acylcarnitines, but lower concentrations of phosphatidylcholines, as compared with controls.³⁴ In turn, metabolomic analysis of blood samples from the CATHGEN (Catheterization Genetics) study showed that patients with heart failure with preserved ejection fraction and, even more so, heart failure with reduced ejection fraction had higher concentrations of acylcarnitines compared with a control group without evidence of heart failure.³⁵

Like previous studies of patients with heart failure, the present study found that higher acylcarnitine levels were associated with LVDD. Through the carnitine shuttle, acylcarnitines are carriers of fatty acids from the cytoplasm into the mitochondria, where the fatty acids undergo beta-oxidation and ATP production (Figure S2). High plasma concentrations of acylcarnitines might reflect mitochondrial dysfunction, either in the myocardium, skeletal muscle, or both. Insulin resistance can also affect the fatty acid oxidation and plasma concentrations of acylcarnitine.³⁶ In

our study, however, there was no difference in homeostasis model assessment of insulin resistance between participants with LVDD versus those with normal LV diastolic function, as noted above.

Under physiological conditions, fatty-acid-beta oxidation is the source of 50% to 70% of cardiomyocyte energy production, and glucose is only a minor contributor.³⁷ Evidence from animal and human studies shows that heart failure, depending on the etiology, is associated with reprogramming of energy metabolism and change in fuel preference.^{38,39} In our cohort, we found that participants with LVDD had a higher plasma concentration of DAG and TAGs, which, in turn, are sources of fatty acids (Figure S2). Furthermore, TAG, acting through PPAR α (peroxisome proliferator-activated receptor α), can activate the transcription of genes that encode important regulatory fatty-acid-oxidation proteins, resulting in enhanced lipid metabolism.⁴⁰ Diabetes mellitus and obesity can be associated with higher concentrations of DAGs and TAGs, but in our study diabetic women were excluded. Furthermore, insulin resistance (homeostasis model assessment of insulin resistance), body mass index, and lipid profiles were similar between participants with LVDD versus those with normal LV diastolic function. It is possible, however, that the associations observed could reflect residual confounding, wherein greater lipolysis or hepatic DAG or TAG production may signal incompletely measured insulin resistance or related metabolic perturbations.

Another notable association identified by our study is that of lower phosphocholine with presence of LVDD. Despite the documented association of lower phosphocholine with clinical heart failure,³⁴ available findings on the relationship of this metabolite with different measures of LVDD are contradictory.⁴¹ Our finding lends support to deficiency of this metabolite, and not excess, as implicated in subclinical cardiac dysfunction or associated disorders. Phosphocholine is a metabolic intermediate in the synthesis of phosphatidylcholine from choline, the latter primarily derived from dietary sources. Choline levels are reduced by oxidative stress, which could account for the observed association of low phosphocholine with LVDD. Moreover, phosphatidylcholine is a key component of biological membranes, suggesting that reduced levels could alter membrane function in ways that are unfavorable to cardiac function or its determinants.

The overall higher availability of fatty acids and acylcarnitines could potentially lead to increased cardiomyocyte reliance on fatty acid oxidation, heightening myocardial susceptibility to ischemia, because ATP production by beta-oxidation requires greater oxygen consumption than that required by glycolysis.³⁷ Our participants were free of self-reported coronary heart disease, however, diminishing this as a contributing factor to LVDD in the present study. Higher fatty acid delivery to cardiomyocytes may lead to mismatch

between supply and mitochondrial oxidative capacity, resulting in cardiac steatosis and lipotoxicity as observed in diabetics and obese patients.⁴² In addition, higher concentration of acylcarnitines, apart from reflecting impaired fatty acid oxidation, could itself contribute to cardiac dysfunction by fostering inflammation and oxidative stress.⁴³

Echocardiographic LVDD is a measure of subclinical cardiac dysfunction that portends an increased risk of progression to overt heart failure. The echocardiographic diagnosis of LVDD has been a matter of debate for the past several years, as reflected by the different diagnostic criteria used in the 1998 European Society of Cardiology guidelines,⁴⁴ and subsequently in newer versions published on 2009 and 2016 by the ASE/EACVI.^{15,16} Observational studies have shown that compared with the 1998 and 2009 guidelines, prevalence of LVDD by the 2016 classification is lower, but appears to correlate better with clinical outcomes and N-terminal pro b-type natriuretic peptide plasma level.¹⁷ We applied the 2009 classification¹⁶ for this study, leveraging its higher sensitivity to improve our ability to detect early abnormalities in this modest sample that likely also portend progression to full-fledged LVDD and, ultimately, LVDD, albeit with lower specificity than the 2016 classification. In support of this approach, sensitivity analysis using the 2016 ASE/EACVI guidelines,¹⁵ which identified 28 women with LVDD (as compared with 52 women using the 2009 guidelines),¹⁶ also showed an association between LVDD and high concentrations of TAGs and DAG, though not acylcarnitines. The extent to which these differences might reflect differences in echocardiographic features captured by 2009 and 2016 guideline definitions will require further study.

The strengths of this exploratory study are its inclusion of participants from a well-characterized cohort of women with or at risk for HIV and predominantly of black or Hispanic race/ethnicity and leveraging of state-of-the-art metabolomic profiling and standardized echocardiographic assessments. Owing to the modest sample size, no individual metabolites remained significant after correction for multiple testing in this exploratory study. Accordingly, the findings of this study can only be considered as hypothesis generating, although, interestingly, the association of the metabolite score with LVDD proved to be highly statistically significant. As such, the results of this pilot study do suggest that perturbations in distinct pathways are related to early-stage myocardial abnormalities in high-risk middle-aged women. The study's modest size also prevented well-powered assessment of differences in metabolomic associations between HIV-positive and -negative groups, a question that requires investigations of substantially larger scale. Another important limitation is the study's cross-sectional observational design, which prevents determinations of direction of associations or their casual basis. Although we undertook extensive adjustment for

covariates, residual confounding by partly measured or unmeasured factors cannot be excluded. In addition, blood metabolites provide an integrative picture of metabolism, but cannot distinguish the tissue sources of the metabolites in question. Such determinations can only be accomplished through experimental studies focused on specific organs. Last, these exploratory results are not necessarily generalizable to men or other race/ethnic groups.

In conclusion, the present exploratory study shows that metabolite profiles reflecting glycerophospholipid metabolism and fatty acid oxidation are associated with subclinical LVDD in women with or at risk for HIV. These results suggest that better understanding of these metabolic abnormalities and their determinants could offer new therapeutic strategies for prevention of cardiac dysfunction and heart failure in high-risk individuals with or at risk for HIV.

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Disclosures

None.

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SUPPLEMENTAL MATERIAL

Data S1.

Supplemental Methods

Plasma levels of metabolites were profiled at Broad Institute using liquid chromatography-tandem mass spectrometry (LC-MS). LC-MS data were acquired using a Nexera X2 U-HPLC (Shimadzu Corp.; Marlborough, MA) coupled to an Exactive Plus mass spectrometer (Thermo Fisher Scientific; Waltham, MA).

Polar metabolites were extracted from plasma (10 μ L) using 90 μ L of 74.9:24.9:0.2 v/v/v acetonitrile/methanol/formic acid containing stable isotope-labeled internal standards (valine-d8, Sigma-Aldrich; St. Louis, MO; and phenylalanine-d8, Cambridge Isotope Laboratories; Andover, MA). The samples were centrifuged (10 min, 9,000 x g, 4°C), and the supernatants were injected directly onto a 150 x 2 mm, 3 μ m Atlantis HILIC column (Waters; Milford, MA). The column was eluted isocratically at a flow rate of 250 μ L/min with 5% mobile phase A (10 mM ammonium formate and 0.1% formic acid in water) for 0.5 minute followed by a linear gradient to 40% mobile phase B (acetonitrile with 0.1% formic acid) over 10 minutes. MS analyses were carried out using electrospray ionization in the positive ion mode using full scan analysis over 70-800 m/z at 70,000 resolution and 3 Hz data acquisition rate. Other MS settings were: sheath gas 40, sweep gas 2, spray voltage 3.5 kV, capillary temperature 350°C, S-lens RF 40, heater temperature 300°C, microscans 1, automatic gain control target 1e6, and maximum ion time 250 ms.

Lipid metabolites were extracted from plasma (10 μ L) using 190 μ L of isopropanol containing 1,2-didodecanoyl-sn-glycero-3-phosphocholine (Avanti Polar Lipids; Alabaster, AL). After centrifugation, supernatants were injected directly onto a 100 x 2.1

mm, 1.7 μ m ACQUITY BEH C8 column (Waters; Milford, MA). The column was eluted isocratically with 80% mobile phase A (95:5:0.1 vol/vol/vol 10mM ammonium acetate/methanol/formic acid) for 1 minute followed by a linear gradient to 80% mobile-phase B (99.9:0.1 vol/vol methanol/formic acid) over 2 minutes, a linear gradient to 100% mobile phase B over 7 minutes, then 3 minutes at 100% mobile-phase B. MS analyses were carried out using electrospray ionization in the positive ion mode using full scan analysis over 200–1100 m/z at 70,000 resolution and 3 Hz data acquisition rate. Other MS settings were: sheath gas 50, in source CID 5 eV, sweep gas 5, spray voltage 3 kV, capillary temperature 300°C, S-lens RF 60, heater temperature 300°C, microscans 1, automatic gain control target 1e6, and maximum ion time 100 ms. Lipid identities were denoted by total acyl carbon number and total number of double bond number.

Metabolite peaks were identified and confirmed using authentic reference standards. Metabolite peak were quantified using area-under-the-curve. Raw data were processed using TraceFinder software (Thermo Fisher Scientific; Waltham, MA) and Progenesis Q1 (Nonlinear Dynamics; Newcastle upon Tyne, UK).

Table S1. Relationship between top 10 polar metabolites/lipids concentration and LVDD.

		All		HIV +	HIV -	
		OR (95% CI)	P	OR (95% CI)	OR (95% CI)	P-int*
Polar metabolites						
N-carbamoyl-beta-alanine	Model 1	1.32 (0.90 - 1.92)	0.150	1.29 (0.84 - 1.96)	1.37 (0.54 - 3.46)	0.40
	Model 2	1.30 (0.87 - 1.95)	0.204	1.28 (0.82 - 2.00)	1.37 (0.54 - 3.45)	0.56
	Model 3	1.21 (0.77 - 1.90)	0.412	1.24 (0.76 - 2.04)	1.06 (0.39 - 2.88)	0.67
Phosphocholine	Model 1	0.59 (0.39 - 0.87)	0.008	0.56 (0.37 - 0.87)	0.65 (0.26 - 1.62)	0.26
	Model 2	0.56 (0.37 - 0.85)	0.006	0.55 (0.35 - 0.86)	0.63 (0.25 - 1.59)	0.64
	Model 3	0.60 (0.39 - 0.93)	0.022	0.52 (0.32 - 0.86)	1.06 (0.41 - 2.71)	0.92
DMGV [†]	Model 1	1.20 (0.82 - 1.74)	0.353	1.07 (0.69 - 1.64)	1.77 (0.74 - 4.23)	0.55
	Model 2	1.24 (0.84 - 1.83)	0.284	1.12 (0.71 - 1.76)	1.73 (0.72 - 4.15)	0.42
	Model 3	1.23 (0.77 - 1.97)	0.386	1.11 (0.65 - 1.91)	1.65 (0.65 - 4.21)	0.78
N-Acetyl putrescine	Model 1	1.30 (0.90 - 1.88)	0.164	1.45 (0.95 - 2.21)	0.79 (0.33 - 1.92)	0.38
	Model 2	1.32 (0.90 - 1.92)	0.152	1.50 (0.97 - 2.30)	0.78 (0.32 - 1.90)	0.31
	Model 3	1.47 (0.97 - 2.23)	0.068	1.73 (1.07 - 2.79)	0.81 (0.32 - 2.05)	0.82
Guanine	Model 1	1.42 (0.97 - 2.09)	0.070	1.58 (1.01 - 2.48)	0.81 (0.28 - 2.33)	0.41
	Model 2	1.51 (0.97 - 2.33)	0.066	1.74 (1.05 - 2.89)	0.80 (0.28 - 2.34)	0.25
	Model 3	1.37 (0.86 - 2.19)	0.182	1.59 (0.93 - 2.70)	0.73 (0.24 - 2.22)	0.65
C7-carnitine	Model 1	1.60 (1.09 - 2.36)	0.017	1.61 (1.04 - 2.50)	1.55 (0.68 - 3.54)	0.25
	Model 2	1.62 (1.09 - 2.41)	0.018	1.63 (1.04 - 2.56)	1.58 (0.69 - 3.63)	0.38
	Model 3	1.88 (1.18 - 2.98)	0.008	1.94 (1.16 - 3.25)	1.67 (0.68 - 4.06)	0.27
C9-carnitine	Model 1	1.73 (1.18 - 2.55)	0.006	1.96 (1.24 - 3.12)	1.23 (0.56 - 2.71)	0.35
	Model 2	1.82 (1.21 - 2.72)	0.004	2.07 (1.28 - 3.35)	1.23 (0.56 - 2.72)	0.35
	Model 3	1.96 (1.25 - 3.08)	0.004	2.22 (1.32 - 3.75)	1.27 (0.51 - 3.17)	0.62
C14-carnitine	Model 1	1.41 (0.97 - 2.05)	0.075	1.50 (0.98 - 2.31)	1.08 (0.48 - 2.43)	0.34
	Model 2	1.42 (0.96 - 2.08)	0.076	1.53 (0.98 - 2.37)	1.09 (0.48 - 2.44)	0.43
	Model 3	1.64 (1.02 - 2.64)	0.042	1.83 (1.07 - 3.13)	1.11 (0.44 - 2.78)	0.27
C16-carnitine	Model 1	1.67 (1.13 - 2.48)	0.011	1.72 (1.10 - 2.70)	1.44 (0.58 - 3.60)	0.30
	Model 2	1.73 (1.14 - 2.63)	0.010	1.81 (1.12 - 2.91)	1.47 (0.59 - 3.67)	0.37

	Model 3	1.94 (1.18 - 3.21)	0.010	2.14 (1.22 - 3.75)	1.30 (0.47 - 3.63)	0.28
C18-carnitine	Model 1	1.48 (1.01 - 2.17)	0.047	1.46 (0.94 - 2.25)	1.52 (0.67 - 3.46)	0.24
	Model 2	1.45 (0.98 - 2.13)	0.063	1.42 (0.91 - 2.21)	1.54 (0.68 - 3.52)	0.53
	Model 3	1.50 (0.96 - 2.33)	0.072	1.53 (0.93 - 2.49)	1.41 (0.58 - 3.45)	0.67
Lipids						
DAG [‡] 30:0	Model 1	1.65 (1.10 - 2.49)	0.016	1.46 (0.93 - 2.30)	2.65 (1.05 - 6.73)	0.32
	Model 2	1.74 (1.14 - 2.66)	0.011	1.56 (0.98 - 2.49)	2.64 (1.04 - 6.68)	0.28
	Model 3	1.70 (1.05 - 2.76)	0.032	1.48 (0.87 - 2.53)	2.76 (1.03 - 7.37)	0.60
DAG 34:0	Model 1	1.56 (1.04 - 2.34)	0.030	1.38 (0.88 - 2.15)	2.77 (1.01 - 7.60)	0.33
	Model 2	1.65 (1.08 - 2.54)	0.022	1.47 (0.92 - 2.35)	2.78 (1.01 - 7.61)	0.31
	Model 3	1.58 (0.97 - 2.59)	0.066	1.35 (0.80 - 2.30)	3.19 (1.08 - 9.41)	0.58
PC [§] 38:6 P	Model 1	0.64 (0.43 - 0.95)	0.029	0.69 (0.44 - 1.07)	0.47 (0.18 - 1.21)	0.31
	Model 2	0.64 (0.42 - 0.95)	0.029	0.68 (0.44 - 1.06)	0.47 (0.18 - 1.21)	0.54
	Model 3	0.69 (0.40 - 1.19)	0.179	0.76 (0.43 - 1.37)	0.45 (0.15 - 1.35)	0.70
TAG ^{**} 44:0	Model 1	1.57 (1.06 - 2.31)	0.023	1.45 (0.95 - 2.20)	2.39 (0.94 - 6.07)	0.33
	Model 2	1.63 (1.10 - 2.43)	0.015	1.50 (0.98 - 2.32)	2.38 (0.94 - 6.05)	0.32
	Model 3	1.66 (1.06 - 2.61)	0.028	1.49 (0.91 - 2.43)	2.61 (0.97 - 6.99)	0.51
TAG 46:0	Model 1	1.62 (1.08 - 2.41)	0.018	1.45 (0.94 - 2.23)	2.89 (1.06 - 7.90)	0.36
	Model 2	1.69 (1.12 - 2.54)	0.012	1.51 (0.97 - 2.35)	2.89 (1.06 - 7.92)	0.34
	Model 3	1.78 (1.10 - 2.87)	0.018	1.56 (0.93 - 2.62)	3.17 (1.09 - 9.23)	0.52
TAG 46:1	Model 1	1.59 (1.07 - 2.36)	0.022	1.43 (0.93 - 2.18)	2.73 (1.01 - 7.33)	0.35
	Model 2	1.63 (1.09 - 2.44)	0.017	1.48 (0.96 - 2.27)	2.71 (1.01 - 7.29)	0.32
	Model 3	1.69 (1.04 - 2.73)	0.034	1.50 (0.89 - 2.54)	2.69 (0.96 - 7.52)	0.44
TAG 48:0	Model 1	1.61 (1.08 - 2.40)	0.019	1.40 (0.91 - 2.14)	3.75 (1.28 - 11.00)	0.32
	Model 2	1.69 (1.12 - 2.54)	0.012	1.46 (0.94 - 2.26)	3.76 (1.28 - 11.08)	0.37
	Model 3	1.77 (1.08 - 2.88)	0.023	1.49 (0.89 - 2.51)	4.39 (1.32 - 14.63)	0.50
TAG 48:1	Model 1	1.65 (1.10 - 2.49)	0.016	1.48 (0.96 - 2.27)	3.40 (1.09 - 10.59)	0.35
	Model 2	1.70 (1.12 - 2.57)	0.012	1.53 (0.99 - 2.38)	3.39 (1.09 - 10.56)	0.32
	Model 3	1.79 (1.06 - 3.03)	0.029	1.60 (0.93 - 2.77)	3.61 (1.06 - 12.32)	0.41

TAG 50:0	Model 1	1.59 (1.08 - 2.35)	0.019	1.39 (0.91 - 2.12)	3.22 (1.16 - 8.96)	0.30
	Model 2	1.65 (1.10 - 2.47)	0.015	1.44 (0.93 - 2.23)	3.25 (1.16 - 9.06)	0.33
	Model 3	1.73 (1.05 - 2.83)	0.031	1.47 (0.86 - 2.50)	3.75 (1.19 - 11.74)	0.41
TAG 50:1	Model 1	1.59 (1.06 - 2.37)	0.025	1.38 (0.90 - 2.11)	4.87 (1.26 - 18.84)	0.39
	Model 2	1.63 (1.08 - 2.48)	0.021	1.43 (0.92 - 2.21)	4.89 (1.26 - 18.98)	0.35
	Model 3	1.61 (0.94 - 2.77)	0.086	1.43 (0.82 - 2.49)	5.17 (1.13 - 23.57)	0.46

Regression was performed adjusted for age, race, education level, smoking, drinking habits, HCV infection and intravenous drug use (model 1). Further adjustment was performed for HIV status, baseline viral load and ART treatment status (model 2); and additionally, for total cholesterol, HDL cholesterol, systolic blood pressure, HOMA-IR, lipid-lowering medication and anti-hypertensive medication use (model 3). Metabolites underwent rank-based inverse normal transformation. The OR for LVDD for each 1-unit change in the standardized plasma metabolite level.

* P for interaction

† Dimethylguanidino valeric acid

‡ Diacylglycerol

§ Phosphatidylcholine

** Triacylglycerol

Table S2. Joint effect of the first 20 metabolites of the first principal component.

	Model 1		Model 2		Model 3	
	OR (95% CI)	P*	OR (95% CI)	P*	OR (95% CI)	P*
All	1.20 (1.09 - 1.31)	<0.0001	1.21 (1.10 - 1.33)	<0.0001	1.25 (1.11 - 1.40)	<0.0001
Among HIV+	1.18 (1.07 - 1.30)		1.19 (1.08 - 1.31)		1.23 (1.09 - 1.39)	
Among HIV-	1.30 (1.05 - 1.59)	0.39	1.30 (1.06 - 1.60)	0.43	1.33 (1.06 - 1.66)	0.51

Weighted score of the first 10 polar metabolites and 10 lipids of the first principal component. Regression was performed adjusted for age, race, education level, smoking, drinking habits, HCV infection and intravenous drug use (model 1). Further adjustment was performed for HIV status, baseline viral load and ART treatment status (model 2); and additionally, for total cholesterol, HDL cholesterol, systolic blood pressure, HOMA-IR, lipid-lowering medication and anti-hypertensive medication use (model 3). The OR for LVDD for each 1-unit change in the standardized plasma metabolite level.* P for interaction

Table S3. Sensitivity analysis of relationship between top polar metabolites/lipids concentration and LVDD.

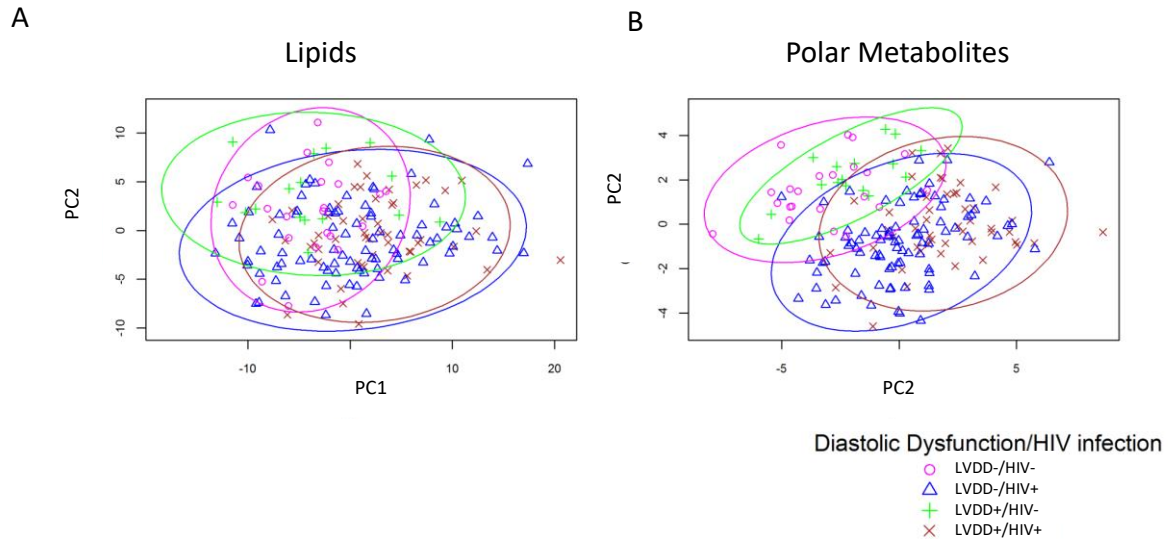
		OR (95% CI)	P
Polar Metabolites			
Phosphocholine	Model 1	0.94 (0.60 - 1.48)	0.78
	Model 2	0.95 (0.59 - 1.52)	0.82
	Model 3	0.94 (0.55 - 1.59)	0.81
C7-Carnitine	Model 1	1.16 (0.74 - 1.83)	0.52
	Model 2	1.11 (0.69 - 1.80)	0.67
	Model 3	0.99 (0.55 - 1.80)	0.98
C9-Carnitine	Model 1	0.97 (0.62 - 1.52)	0.90
	Model 2	0.86 (0.53 - 1.41)	0.56
	Model 3	0.81 (0.45 - 1.45)	0.48
C16-Carnitine	Model 1	1.04 (0.67 - 1.63)	0.85
	Model 2	1.10 (0.66 - 1.81)	0.72
	Model 3	1.16 (0.61 - 2.22)	0.65
Lipids			
DAG [†] 30:0	Model 1	1.75 (1.04 - 2.93)	0.04
	Model 2	1.79 (1.03 - 3.14)	0.04
	Model 3	1.50 (0.80 - 2.82)	0.21
TAG [†] 46:0	Model 1	1.61 (0.97 - 2.66)	0.06
	Model 2	1.55 (0.92 - 2.62)	0.10
	Model 3	1.34 (0.72 - 2.47)	0.35
TAG 48:0	Model 1	1.57 (0.95 - 2.59)	0.08
	Model 2	1.49 (0.89 - 2.52)	0.13
	Model 3	1.20 (0.64 - 2.23)	0.57
TAG 48:1	Model 1	1.55 (0.94 - 2.58)	0.09
	Model 2	1.58 (0.92 - 2.71)	0.10
	Model 3	1.29 (0.66 - 2.52)	0.46
TAG 50:0	Model 1	1.78 (1.08 - 2.92)	0.02
	Model 2	1.72 (1.02 - 2.89)	0.04

Sensitivity analysis of the top polar metabolites and lipids from the first principal component using the 2016 ASE/EACVI guidelines. Regression was performed adjusted for age, race, education level, smoking, drinking habits, HCV infection and intravenous drug use (model 1). Further adjustment was performed for HIV status, baseline viral load and antiretroviral therapy (ART) treatment status (model 2); and additionally, for total cholesterol, HDL cholesterol, systolic blood pressure, homeostatic model assessment of insulin resistance (HOMA-IR), lipid-lowering medication and anti-hypertensive medication use (model 3). Metabolites underwent rank-based inverse normal transformed. The odds ratio (OR) for LVDD for each 1-unit change in the standardized plasma metabolite level.

* Diacylglycerol

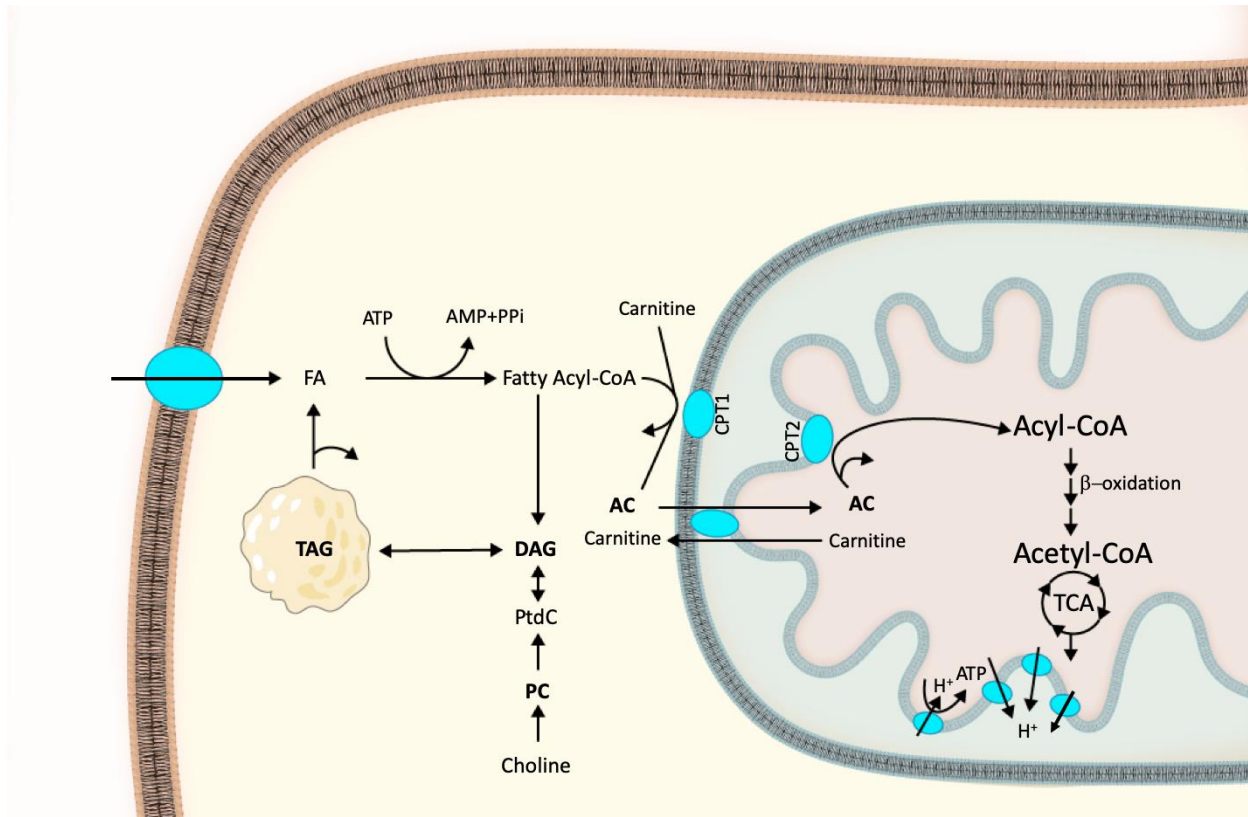
† Triacylglycerol

Figure S1. Scores plot of polar metabolites and lipids according to presence of LVDD and HIV status.



Scatterplot showing metabolites from principal component 1 and 2 obtained with PLS-DA analysis for (A) lipids and (B) polar metabolites of four subgroups of participants: 1. without LVDD and HIV-negative (pink circles); 2. without LVDD and HIV-positive (blue triangles); 3. with LVDD and HIV-negative (green crosses); and 4. with LVDD and HIV-positive (brown rotated crosses). LVDD: left ventricular diastolic dysfunction. PLS-DA: Partial least squares discriminant analysis.

Figure S2. Diagram of glycerophospholipid and fatty-acid-oxidation in the cardiomyocyte.



In women at risk or with HIV infection, LVDD was associated with higher levels of acylcarnitines, diacylglycerols and triacylglycerols, but lower levels of phosphocholine. Fatty acids can be obtained from the extracellular space by simple diffusion or via CD36/FATP transporters or from storages located in the cytoplasm in the form of triacylglycerol (TAG). On the other side, choline through the CDP-choline pathways choline, is metabolized into phosphocholine (PC), phosphatidylcholine (PtdC), and subsequently diacylglycerol (DAG), which finally can be stored as TAG. Free fatty acids can be esterified into fatty acyl-CoA which can be further esterified to form DAG or the acyl group transferred to carnitine forming acylcarnitine (AC) via carnitine palmitoyltransferase 1 (CPT1) which is located in the outer mitochondrial membrane. AC is transported into the mitochondria matrix where via CPT2 is converted back to fatty acyl-CoA, which is metabolized via beta-oxidation, producing NADH and FADH₂, as well as acetyl CoA which undergoes final metabolism via the tricarboxylic acid cycle (TCA).