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Plasminogen Activator Inhibitor-1 Predicts Negative Alterations in Whole-Body Insulin Sensitivity in Chronic HIV Infection

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Abstract

Plasminogen activator inhibitor type 1 (PAI-1), a key negative regulator of fibrinolysis, has been investigated to be one of the potential mechanisms of the development of impaired insulin sensitivity, insulin resistance, and diabetes mellitus. Because chronically stable HIV-infected individuals frequently develop abnormal glucose metabolism, including insulin resistance and diabetes mellitus, we postulated that PAI-1 could be one of the multifactorial pathogenic roles in the development of impaired insulin sensitivity and insulin resistance among chronic HIV-infected individuals. From our longitudinal cohort study, we selectively recruited chronically stable HIV-infected individuals without diagnosis of diabetes mellitus at baseline (N=62) to analyze the correlation of baseline inflammatory cytokines, including PAI-1 and whole-body insulin sensitivity, with 2-year follow-up, as measured by Matsuda Index. We found a negative correlation between baseline PAI-1 and Matsuda Index (r=-0.435, p=.001) and a negative correlation between baseline PAI-1 and Matsuda Index at 2 years (r=-0.377, p=.005). In a linear regression model that included age, total body fat mass percentage, serum amyloid A, and family history of diabetes mellitus, PAI-1 still remained significantly associated with Matsuda Index at 2-year follow-up ($\beta=-.397$, p=.002). Our longitudinal study suggests that PAI-1 is an independent predictor of impaired insulin sensitivity among chronic HIV-infected individuals.

Keyword: plasminogen activator inhibitor-1, insulin sensitivity, chronic HIV infection

Introduction

S INCE THE INTRODUCTION of antiretroviral therapy (ART), HIV-associated mortality and morbidity has remarkably declined and the life expectancy of HIV-infected individuals is close to the general population.¹ However, adverse metabolic effects such as insulin resistance, diabetes mellitus, and hyperlipidemia have been more frequently found in the chronically stable HIV-infected population than the general population.^{1–3} To date, the underlying pathophysiologic mechanism of impaired insulin sensitivity, insulin resistance, and diabetes mellitus in treated HIV-infected individuals is still not well understood. Plasminogen activator inhibitor type 1 (PAI-1), a crucial inhibitor of fibrinolysis in the body, may play an important role in the development of impaired insulin sensitivity, insulin resistance, and diabetes mellitus.^{4–7} Although the clear explanation of the relationship between increased PAI-1 levels and impaired insulin sensitivity and insulin resistance remains complicated, *in vitro* studies suggested that increased PAI-1 level may interfere with the insulin signaling through complex signaling molecules, including $\alpha v\beta 3$ integrin and the insulin receptor substrate. Hence, increased PAI-1 can potentially inhibit the full mechanism of action of insulin. PAI-1 concentration among HIV-infected patients who have HIV-associated lipodystrophy syndrome and

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obesity has been found to be elevated compared with HIVnegative controls.^{8,9} There are currently no reported studies of using inflammatory cytokines, including PAI-1, to predict impaired insulin sensitivity in patients with chronic HIV infection. The primary rationale of our study was to investigate the potential predictor of the development of impaired insulin sensitivity in patients with chronic HIV infection.

We included 62 chronically stable HIV-infected individuals and over time, we investigated the correlation between soluble biomarkers of inflammation, including PAI-1, and whole-body insulin sensitivity measured by the Matsuda Index. We hypothesized that PAI-1 would be closely associated with impaired insulin sensitivity in chronically HIV-infected patients on stable therapy.

Materials and Methods

Study design

This was a longitudinal study using baseline and 2-year follow-up data from participants enrolled into the Hawaii Aging with HIV-Cardiovascular cohort (HAHC-CVD). Complete details of the cohort study design were previously published.¹⁰ In brief, criteria for entry into the initial cohort were documented HIV infection, 40 years of age or older, and on stable ART for >3 months. The study was approved by the Committee on Human Subjects of the University of Hawaii, and written informed consent was obtained from all participants.

Subjects who had baseline and 2-year follow-up measures of the Matsuda Index and baseline characteristics, including monocyte subset and inflammatory biomarkers, were included in the analyses. Subjects diagnosed with diabetes mellitus, at baseline as determined by the American Diabetic Association guidelines, as well as those on diabetic medications, including Biguanide, insulin secretagogues, and insulin, were excluded.¹¹

Clinical assessment

Blood pressure, height, weight, and waist and hip measurement were obtained at study entry. Total fat mass percentage was used as an indicator of body fat mass instead of body mass index (BMI) because it provides a better measurement of body fat composition, which was measured by a dual-energy X-ray absorptiometry scan. BMI number generally provides a simple measurement of weight status, which was derived from the individual's mass (weight of fat, muscles, and tissues) and height. Fasting laboratory measurements, including complete blood count (CBC), CD4 cell count, plasma HIV RNA, glucose, insulin, total cholesterol, low-density lipoprotein cholesterol, high-density lipoprotein cholesterol, and triglycerides, were also obtained. Clinical history included age, gender, current medications, past medical and family histories, CD4 cell count nadir, and tobacco use. Metabolic syndrome status was determined by using the National Cholesterol Education Program's Adult Treatment Panel III report.12

Matsuda Index was used to measure whole-body insulin sensitivity and was calculated as $\frac{10,000}{\sqrt{.(G_0 \times I_0 \times G_{120} \times I_{120})}}$, using mean plasma glucose and insulin concentration from the oral glucose tolerance test (OGTT). Because this method reflects how well both hepatic and peripheral tissues respond to

glucose, the Matsuda Index provides more precise wholebody insulin sensitivity than HOMA2, which uses only fasting glucose and insulin concentrations.¹³

Plasma-soluble biomarkers

Plasminogen activator inhibitor (PAI-1) along with other soluble biomarkers, including soluble E-selectin (sE-selectin), soluble vascular cell adhesion molecule-1 (sVCAM-1), soluble intercellular adhesion molecule-1 (sICAM-1), matrix metalloproteinase (MMP)-9, myeloperoxidase (MPO), C-reactive protein (CRP), serum amyloid A (SAA), serum amyloid P (SAP), IL-1b, IL-6, IL-8, IL-10, TNF- α , MCP-1, vascular endothelial growth factor, MCP-1, and interferon (IFN)- γ , were assessed at baseline. The Milliplex Human Cardiovascular Disease panels (EMD Millipore) was used to assay plasmasoluble biomarkers as previously described.¹⁰ We did not obtain blood samples for the soluble biomarkers at the 2 year follow-up.

Statistical analysis

Demographic and clinical information was summarized by median (quartile 1, quartile 3) for continuous variables, and frequency (percentage) for categorical variables. Inflammatory cytokines were log-10 transformed to adjust for normality, where necessary. Pearson correlations were used to assess the correlation between variables. The associations of baseline and 2-year Matsuda Index with inflammatory cytokines were assessed by separate multivariable linear regression models, adjusting for insulin resistance-associated risk factors such as age, body fat percentage, and family history of diabetes mellitus. The outcome variable was the 2-year follow-up Matsuda Index. The two-year follow-up Matsuda Index was subsequently adjusted for the insulin resistance-associated risk factors. This approach, rather than using change in the Matsuda Index over 2 years, would provide better precision and power in small sample size.^{14,15} Stepwise backward elimination regression was performed to determine the predictors associated with whole-body insulin sensitivity, by using a selection criterion of p < .05 to remain in the model. All statistical analyses were conducted in SPSS (IBM, Version 23). A p-value of < .05 was regarded as statistically significant.

Results

Patient characteristics

Of the 160 subjects enrolled in the HAHC-CVD cohort, 62 had an available Matsuda Index at baseline and 2-year follow-up. Of the 62 subjects, soluble inflammatory cytokines and monocyte subsets at baseline were available in 54 subjects. The demographic, clinical, and immunologic characteristics of the patients are summarized in Table 1. The patients were predominately Caucasian (61.3%) and of male gender (90.3%). The median age was 51 years. Study enrollment inclusion criteria required patients to be on stable ART for at least 3 months, and 85% had suppressed viral load that was defined as HIV RNA <50 copies/ml. The median CD4 cell count was 513 cells/ μ l, and the median CD4 cell count nadir was 160 cells/ μ l. ART consisted of nucleoside reverse transcriptase inhibitors, non-nucleoside reverse transcriptase inhibitors, and protease inhibitors (PIs) at entry

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TABLE 1. E	Baseline	CHARACTERISTICS	OF THE SUBJECTS
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Baseline	characteristics

Ν	62
Age, year	51 [47–56]
Male, <i>n</i> (%)	56 (90.3%)
Caucasian race	38 (61.3%)
BMI, kg/m ²	26.07 [24.48–27.79]
Framingham risk score	0.06 [0.03–0.11]
History of clinical CVD events, n (%)
Hypertension	20 (32.26)
Stroke	2(3.2)
History of myocardial infarction	3 (4.8)
Family history of diabetes, n (%)	29 (46.77)
Smoking history, n (%)	37 (59.7)
Undetectable HIV RNA, n (%)	53 (85)
Current antiretroviral medications, r	ı (%)
Nucleoside reverse	61 (98.4)
transcriptase inhibitor	21(50)
transcriptase inhibitor	51 (50)
Protease inhibitor	30(484)
Matabalia sundroma u (07)	16 (25.9)
Metabolic syndrome, n (%)	10 (23.8)
Matsuda Index	07.3 [01.0-93.0] A 01 [3 A2_0 80]
Mean glucose during OGTT, mg/dl	117.13 [101.5 - 139.62]
Mean insulin during OGTT, mg/dl	50.86 [28.27-74.85]
Soluble markers	· · · · · [· · · · · · ·]
sE-Selectin ng/ml	36 [19-46]
sICAM 1. ng/ml	136 [109–158]
sVCAM 1, ng/ml	1174 [936–1343]
MMP9, ng/ml	55 [45-92]
MPO, ng/ml	16.1 [12.0-21.2]
SAA, ng/ml	10270 [3188-43826]
CRP, ng/ml	8186 [3081-26244]
IL-10, pg/ml	2.18 [0.93–4.65]
IFIN-y, pg/ml	0.00 [0.3/-1.30]
I 6 pg/ml	1 44 [1 02 2 42]
IL-0, pg/ml	3 38 [2 66 4 23]
MCP-1 ng/ml	146 [111–186]
VEGF. pg/ml	33 [14–55]
TNF-a, pg/ml	3.21 [1.54-4.34]
Year 2 data	
Fasting glucose, mg/dl	87,50 [83.00-96.00]
Matsuda Index	4.38 [2.82–7.62]
Mean glucose during OGTT, mg/dl	117.81 [99.75–140.00]
Mean insulin during OGTT, mg/dl	53.77 [39.15-71.12]

CVD, cardiovascular cohort; PAI-1, plasminogen activator inhibitor type 1; BMI, body mass index; VEGF, vascular endothelial growth factor; OGTT, oral glucose tolerance test.

by 98%, 50%, 48% of patients, respectively. None of the subjects were on integrase inhibitors.

Associations between Matsuda Index and soluble biomarkers

Baseline levels of measured soluble biomarkers are shown in Table 1.

At baseline, the median Matsuda Index was 4.91 and there were significant negative associations between baseline Matsuda Index and biomarkers, including PAI-1, CRP, IL-1b,

MPO, SAP, SAA, and svCAM. At 2-year follow-up, the median Matsuda Index was 4.38 and only PAI-1 and SAA were associated with the 2-year follow-up Matsuda Index. Two of the 62 patients had developed type 2 diabetes mellitus by the second year of this study.

We found a negative correlation between baseline Matsuda Index and PAI-1 (r=-0.435 and p=.001) and a negative correlation between 2-year Matsuda Index and baseline PAI-1 (r=-0.376, p=.004). In addition, a negative correlation between baseline Matsuda Index and SAA (r=-0.323, p=.017) and a negative correlation between year 2 Matsuda Index and baseline SAA were found (r=-0.278, p=.042). Neither baseline nor year 2 Matsuda Index was significantly different between genders, Table 2.

The relationship between 2-year Matsuda Index and the inflammatory cytokines, including PAI-1 and SAA, was examined further in a multivariable regression model with the Matsuda Index at 2-year follow-up as the dependent variable, Table 3. After adjusting for traditional risk factors of insulin resistance at baseline (age, total body fat percentage, family history of diabetes mellitus), baseline PAI remained significantly associated with the 2-year Matsuda Index after backward selection was performed ($\beta = -.397$, p = .002). Backward selection also determined that age was the most important risk factor for the 2-year Matsuda Index (p = .014). SAA, family history of diabetes mellitus, and total body fat percentage were no longer significantly associated with the Matsuda Index at 2-year follow-up after the backward elimination regression approach. Nine of the 62 subjects

TABLE 2. PEARSON'S CORRELATION OF BASELINE
Cytokines and Matsuda Index at Baseline
and at 2-Year Follow-up

	Matsuda Index at baseline		Matsuda Index at 2-year follow-up	
	Correlation coefficient, r	р	Correlation coefficient, r	р
Total monocyte counts at baseline	-0.262*	.040	069	.594
CRP ^a	-0.310*	.022	-0.261	.057
IFNy ^a	-0.072	.606	0.080	.567
IL10 ^a	-0.177	.201	-0.261	.056
IL1b ^a	-0.309*	.025	-0.070	.619
IL6 ^a	-0.158	.253	-0.151	.277
IL8 ^a	0.044	.754	-0.151	.276
MCP1 ^a	0.114	.412	0.033	.815
MPO ^a	-0.334*	.014	-0.206	.136
MMP9 ^a	-0 .206	.136	-0.255	.062
SAA ^a	-0.323*	.017	-0.278*	.042
Selectin ^a	-0.181	.191	-0.235	.086
SAP ^a	-0.333*	.014	-0.196	.155
SICAM ^a	-0.109	.431	-0.252	.066
SVCAM ^a	-0.341*	.012	-0.237	.084
TNFa ^a	0.076	.585	0.008	.953
PAI-1 ^a	-0.435**	.001	-0.377 **	.005
VEGF ^a	-0.050	.718	0.068	.626

^aVariable was log-10 transformed.

*Significant at p < .05.

**Significant at p < .01.

Bold, statistically significant correlation with Matsuda Index.

Model	Independent variable	Unstandardized coefficients beta	Standardized coefficients beta	Sig.
1	Age	-0.115	-0.274	0.033
	Total body fat percentage	-0.019	-0.057	0.672
	SAA ^a	-0.447	-0.130	0.324
	PAI-1 ^a	-4.87	-0.365	0.008
	Family history of DM	-0.444	-0.193	0.119
2	Age	-0.114	-0.271	0.033
	SĂA ^a	-0.492	-0.143	0.260
	PAI-1 ^a	-5.11	-0.383	0.003
	Family history of DM	-0.456	-0.199	0.105
3	Age	-0.128	-0.303	0.015
	PĂI-1 ^a	-5.55	-0.416	0.001
	Family history of DM	-0.453	-0.197	0.107
4	Age	-0.130	-0.310	0.014
	PĂI-1 ^a	-5.30	-0.397	0.002

TABLE 3. MULTIVARIABLE LINEAR REGRESSION OF MATSUDA INDEX AT TWO-YEAR FOLLOW-UP

Dependent variable: Matsuda Index at 2-year follow-up.

Models represent stages of backward selection.

^aVariable was log-10 transformed.

(14.5%) had changes in their ART regimen during the 2 year period. Changes to their ART regimens are as follows: Three had changed from non-nucleoside reverse transcriptase inhibitors (NNRTIs) to Integrase Strand Transfer Inhibitors (INSTIs); three had changed from PIs to INSTIs; one had addition of PI to an NNRTI-based regimen; and two had intensification of their ART regimens with Maraviroc. No obvious pattern in the Matsuda Index was seen with subjects who had changes in their ART regimen compared with those who had no change in ART. Removing the individuals who had changes in ART from the analysis did not change the results.

Discussion

In this prospective cohort study, we chronically analyzed stable HIV-infected patients to investigate the correlation between inflammatory cytokines and changes in whole-body physiological insulin sensitivity. Our results found that elevated PAI-1 levels were associated with a decreased Matsuda Index, suggesting that PAI-1 could independently predict future negative alterations in wholebody insulin sensitivity in patients with chronic HIV infection. This finding was consistent with our previous cross-sectional study, which found a correlation between PAI-1 and HOMA-IR.¹⁰

PAI-1 is a key regulator of fibrinolysis, which has been found to be associated with coronary artery disease, thrombosis, insulin resistance, and diabetes mellitus.^{4,5,8} PAI-1 concentration in chronically stable HIV individuals appears to be higher compared with HIV-negative individuals because of the increased incidence of abnormalities of body fat redistribution and the effect of ART.^{8,16,17} One article from the Official *Journal of The International AIDS Society* reported that ART could influence the endothelial activation by direct or indirect effects, which contribute to increased secretion of PAI-1 from the endothelial cells.¹⁶ Lipodystrophy and abnormal fat redistribution are often found in chronic HIV infection, especially in those who are on ART. An increase in PAI-1 gene expression was found in HIV-positive patients with lipodystrophy, thereby increasing PAI-1 concentration.^{8,18} Lipodystrophy is associated with increased liver fat content and the liver usually clears PAI-1, so PAI-1 concentration can be increased by clearance defect from fat accumulation in the liver.^{8,19}

Because of the enigmatic mechanism of the development of impaired insulin sensitivity and insulin resistance by enhanced PAI-1 levels, there have been numerous studies that attempted to elucidate the precise underlying mechanism.^{4,5,20} One hypothesis was the interference of insulin signaling by PAI-1. The complete insulin signaling required full activation of several signaling molecules such as $\alpha\nu\beta3$ integrin and insulin receptor substrate.²⁰ PAI-1 is capable of competing with $\alpha\nu\beta3$ integrin, which could potentially reduce tyrosine phosphorylation of the insulin receptor substrate. Several longitudinal studies in the general population found that subjects who developed type 2 diabetes mellitus had higher levels of PAI-1 at baseline compared with subjects who did not develop type 2 diabetes mellitus.²¹

Our study has some limitations that should be acknowledged. First and most notably, the inflammatory cytokines were not available for the 2-year follow-up. It would be helpful to have the inflammatory cytokines at 2-year followup to examine the changes of the inflammatory cytokines over time in terms of responsiveness of PAI-1 level among chronically HIV-infected individuals on ART, not to examine the causality of diabetes mellitus in chronic HIV infection. In the absence of inflammatory cytokine data at 2-year follow-up, we believe that using baseline inflammatory cytokines to predict the 2-year follow-up Matsuda Index provides a more precise and intuitive analysis compared with using the 2-year change in the Matsuda Index. Second, our study had no control group or HIV-negative individuals and this analysis was retrospective in nature. Hence, we could not account for unmeasured confounders and interpret the result as a causal factor of diabetes mellitus. Despite these limitations, the strength of this study was the robust association between the Matsuda Index and PAI-1 in this wellcharacterized cohort independently of potential confounding factors such as age, total body fat percentage, and family history of diabetes mellitus.

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This is the first study to investigate the potential predictor of the development of impaired insulin sensitivity among patients with chronic HIV infection. Our results indicated that an elevation of PAI-1 concentrations at baseline in chronically stable HIV infection was associated with negative alterations in whole-body insulin sensitivity over time. This could suggest that PAI-1 is a potentially independent predictor of impaired insulin sensitivity among chronically stable HIV-infected individuals. Exploring the underlying mechanism of PAI-1 production may offer insights into insulin resistance and insulin sensitivity in HIV-infected individuals, and it may be a novel therapeutic target for insulin resistance and diabetes mellitus in the future.

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Author Disclosure Statement

No competing financial interests exist.

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