

A High HIV DNA Level in PBMCs at Antiretroviral Treatment Interruption Predicts a Shorter Time to Treatment Resumption Independently of CD4 Nadir

Christophe Piketty, Laurence Weiss, Lambert Assoumou, Marianne Burgard, Aurelie Melard, Jean-Michel Ragnaud, Michele Bentata, Pierre Marie Girard,

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A High HIV DNA Level in PBMCs at Antiretroviral Treatment Interruption Predicts a Shorter Time to Treatment Resumption Independently of CD4 Nadir

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Abstract

3 This study aimed to evaluate the safety of antiretroviral treatment interruption 4 (TI) in HIV-infected patients who started treatment based on earlier guidelines, 5 and to identify baseline factors predictive of the time to reach fixed criteria for 6 treatment resumption.

Prospective, open-label, multicentre trial. Patients were eligible if they had a CD4 cell count >350/mm³ and plasma HIV RNA <50000 copies/ml when they first started antiretroviral therapy (ART); and if they had a CD4 count >450/mm³ and stable plasma HIV RNA <5000 copies/mL for at least six months prior to enrolment. The criteria for ART resumption were a CD4 cell count <300/mm³ and/or a CDC stage B or C event.

116 patients had received ART for a median of 5.3 years. The median CD4 cell count and plasma HIV RNA values at inclusion were 809/mm³ and 2.6 loa copies/mL, respectively. Median HIV DNA load at inclusion was 2.3 log copies/10⁶ PBMCs. Thirty-six months after TI, 63.9% of the patients had not yet reached the criteria for ART resumption, and 55.9% of patients had not resumed ART. In Cox multivariable analysis, a high HIV DNA level at TI, a low CD4 nadir, and pre-existing AIDS status were the only significant risk factors for reaching the criteria for ART resumption (hazards ratio: 2.15(1.02-4.53), 4.59(1.22-17.24) and 5.74(1.60-20.56), respectively).

Patients who started ART with a CD4 cell count above 350/mm³ were able to interrupt treatment for long periods without a high absolute risk of either AIDS or severe non-AIDS morbidity/mortality. A high PBMC HIV DNA level at treatment interruption was a strong predictor for more rapid treatment resumption.

- 1 (ClinicalTrials.gov Identifier, NCT00118677)
- 2 Key words: HIV, treatment interruption, HIV-DNA, CD4 nadir, ART

1 Introduction

Combined antiretroviral therapy (ART) increases CD4 cell numbers, reduces plasma HIV RNA levels and improves patient survival [Hogg et al., 1998; Hogg et al., 2001; Lee et al., 2001; Li et al., 1998; Mocroft et al., 2000; Murphy et al., 2001; Palella et al., 1998; Palella et al., 2003]. However, efficacy can be undermined by suboptimal adherence to treatment and by adverse effects. Treatment interruption appears to be feasible and safe in patients with high CD4 cell counts, a high CD4 nadir, and long-term undetectable plasma HIV RNA, as suggested by several small uncontrolled studies [Boschi et al., 2004; Fagard et al., 2003; Fernandez Guerrero et al., 2005; Giuntini et al., 2005; Krolewiecki et al., 2006; Maggiolo et al., 2004; Parish et al., 2002; Pellegrin et al., 2005; Pogany et al., 2007; Ruiz et al., 2000; Skiest et al., 2004; Skiest et al., 2007; Tarwater et al., 2003] and by two controlled trials [Ananworanich et al., 2006; Maggiolo et al., 2008]. However, recent large randomized trials indicate that, compared with patients who continue ART, unselected patients undergoing treatment interruption experience significantly worse clinical outcomes, including HIV disease progression and cardiovascular events [Danel et al., 2006; El-Sadr et al., 2006].

In view of the results of these latter studies, treatment interruptions are no
longer recommended. However, many patients are concerned with the toxicity
and potential long-term adverse effects of ART, and spontaneously stop taking
their treatment. Factors predictive of treatment resumption would therefore be
useful for clinicians managing such patients, but have not been studied
extensively.

1 The aim of this study was to evaluate the safety of treatment interruption in 2 chronically HIV-infected patients who started ART based on earlier guidelines 3 and who had a high CD4 nadir. Factors potentially predictive of the length of 4 treatment interruption, and changes in metabolic parameters were also 5 examined.

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1 Patients and Methods

3 Study design

The ANRS 116 SALTO trial was a prospective, open-label, multicenter trial of treatment interruption in patients who had started treatment early in the course of HIV infection, based on contemporary guidelines, and who wished to discontinue ART. The study protocol was approved by the ethics committee of Pitié Salpétrière Hospital, Paris, France, and all the patients gave their written informed consent before entering the study. The study conformed to the International Conference on Harmonization guidelines for Good Clinical Practices, and with the Helsinki Declaration.

13 Study population

The following patients were eligible: HIV-infected men and non pregnant women at least 18 years of age; who had a CD4 cell count above 350/mm³ and plasma HIV RNA below 50 000 copies/ml (4.7 log₁₀ copies/ml) at ART initiation; who had been on a stable ART regimen (at least two drugs) for 6 months prior to inclusion; who had had no more than one treatment failure; and who, at baseline, had a CD4 cell count above 450/mm³ and stable plasma HIV RNA below 5000 copies/ml.

The exclusion criteria were chronic hepatitis B treated with lamivudine or
tenofovir disoproxil fumarate; acute opportunistic infections; and anticancer
chemotherapy or immunotherapy.

Assessment and monitoring

3 Study design

ART was interrupted at enrolment in the study and the patients were monitored
for up to 36 months. Patients taking a non nucleoside reverse transcriptase
inhibitor were advised to discontinue it 7 days before their other antiretroviral
drugs.

The patients were evaluated at the screening visit (week -4), then at day 0 (TI), week 2, months 1, 2, 4 and 6, and every 3 months thereafter until month 36. Absolute numbers of CD4 T lymphocytes were determined by flow cytometry in whole blood. The plasma HIV RNA level was determined with the techniques routinely used in each participating center, all of which had a detection limit of 400 genome copies/mL. Each patient's samples were all tested with the same technique. The plasma HIV RNA level and the CD4 cell count were determined at each study visit.

The HIV DNA level in peripheral blood mononuclear cells (PBMCs) was measured in blood samples taken at day 0 and at months 6, 12 and 18. PBMC HIV-1 DNA was guantified by real-time PCR with amplification of the LTR gene, using the 8E5 cell line as an external standard [Avettand-Fenoel et al., 2007; Goujard et al., 2006; Rouzioux et al., 2005]. Briefly, PBMCs were separated by Ficoll-Hypague gradient centrifugation and aliquots were frozen until use. Total DNA extracted from each aliquot of PBMCs was first quantified with the Hoechst dye method (Pharmacia), 1 µg of DNA being considered equivalent to ~150 000 cells. This HIV-DNA PCR method guantifies total HIV-DNA (both integrated and non integrated). Final results were expressed as the number or

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log₁₀ number of HIV-1 DNA copies per 10⁶ PBMCs (quantification limit: 5
 copies/PCR, i.e. 30 copies/10⁶ PBMCs).

Frozen plasma samples were collected at months 1, 6, 12, 18 and 24. HIV
genotypic resistance mutations were studied before treatment resumption, as
described elsewhere [Mouroux et al., 2001].

6 Safety was assessed at each visit by means of an interview, a physical
7 examination, and collection of adverse events. Adverse events were coded with
8 the Medical Dictionary for Regulatory Activities (MedDRA) and their severity
9 was graded from 1 to 4 with the ANRS Adverse Event Grading Scale.

10 Lipodystrophy was scored clinically, and anthropometric measurements were 11 made (waist and hip circumference, waist/hip ratio, thigh circumference and 12 body mass index) on day 0 and at months 6, 24 and 36.

Total cholesterol, HDL cholesterol, LDL cholesterol and triglyceride serum
levels were measured after an overnight fast on day 0 and every 6 months
thereafter.

16 The criteria for ART resumption were a fall in the CD4 cell count to below 17 300/mm³ in two consecutive samples collected 15 days apart, and/or the 18 occurrence of a stage B or C AIDS-defining event. Immunologic and virologic 19 responses were assessed 1, 3 and 6 months after ART resumption.

21 Statistical analysis

22 Sample size

To estimate the percentage of patients who did not reach the ART resumption
criteria 12 months after antiretroviral treatment interruption, with a precision of
10%, 100 patients needed to be enrolled.

2 Statistical Analysis

The results are reported as proportions (and 95% confidence intervals when appropriate) for categorical variables, and as medians and interguartile ranges for continuous variables. Changes in the CD4 cell count during treatment interruption were analyzed by using mixed linear models, testing for changes in the slopes during the interruption, with cut points chosen after visual inspection of the median CD4 cell count at each time point. In these analyses, patients resuming ART were censored at the time of treatment resumption. The primary endpoint was the time to the first of two consecutive CD4 cell counts below 300/mm³, 15 days apart, or a CDC stage B or C AIDS-defining event. All events were reviewed by the Data and Safety Monitoring Board. Patients who reached the primary endpoint were analyzed as such even if they did not resume treatment, while patients who resumed treatment without reaching the primary endpoint were censored at time of treatment resumption in the intent-to-treat analysis. In the analysis of observational data, all patients resuming therapy were considered to have reached the primary endpoint. The percentage of patients reaching the primary endpoint was determined using Kaplan-Meier estimates. Factors associated with the occurrence of the primary endpoint were analyzed with univariate and multivariable Cox proportional hazards models. The following variables were tested: baseline age, gender, transmission group, AIDS status, the CD4 cell nadir, the pre-ART CD4 cell count, the baseline CD4 cell count, pre-ART plasma HIV-1 RNA, baseline PMBC HIV DNA, the duration of ART at treatment interruption, and the use of dual- or triple-drug therapy. Akaike's criteria were used to determine whether continuous variables were

best modelled as continuous variables, or after log transformation, or as three-class categorical variables based on the terciles. Based on the results of these analyses, the baseline CD4 cell count and the baseline HIV DNA level, expressed in log₁₀ values, were modelled as three-class variables, and the CD4 cell nadir and the pre-ART CD4 cell count expressed in log₂ were modelled as continuous variables in the final model. Variables with p values below 0.05 in univariate analysis were entered in the Cox multivariable model by using a backward procedure.

9 Changes in lipodystrophy and metabolic parameters were tested for 10 significance with Wilcoxon's paired test. The SPSS software package version 11 15.0 for Windows (SPSS Inc., Illinois, Chicago) and the SAS software package 12 version 9.1 (SAS institute, Cary, NC, USA) were used for all analyses.

1 Results

3 Study population

One hundred and sixteen patients (37% women) were enrolled at 21 sites between January 2003 and March 2004. Baseline characteristics of the study population are shown in Table 1. Follow-up during treatment interruption lasted a total of 270.2 person-years. Three patients had had AIDS-defining events prior to enrolment (pulmonary tuberculosis, esophageal candidiasis, cutaneous Kaposi's sarcoma). The interrupted treatments comprised a protease inhibitor-based regimen in 16% of cases, a non-nucleoside analogue regimen in 35% of cases, a triple nucleoside analogue regimen in 34% of cases (Trizivir in 17% of cases), a dual nucleoside analogue regimen in 14% of cases, and a non nucleoside analogue plus a protease inhibitor in 1% of cases. The median (IQR) CD4 cell count at inclusion was 806/mm³ (676 – 940) and the median CD4 nadir was 381/mm³ (341 – 490). Only 11% of the patients had a CD4 nadir below 300/mm³. The median pre-ART plasma HIV RNA level was 4.2 log₁₀ copies/mL (3.8 – 4.5). Ninety-one percent of the patients had a plasma HIV RNA level below 400 copies at inclusion. The median baseline HIV DNA value was 2.3 log copies per million PBMCs (1.9 - 2.8).

21 Patient disposition

Four patients were lost to follow-up, at months 17, 24 (2 patients) and 27. Four women left the study because they became pregnant, at months 2, 21, 24 and 32. One patient died, of a ruptured cerebral aneurysm at month 21.

Dynamics of CD4+ T cell count, plasma HIV-RNA and PBMC HIV-DNA values after treatment interruption

Changes in the median CD4 cell count after treatment interruption are shown in Figure 1a. The CD4 cell curve comprised three distinct slopes: a first steep decline (-127.2 \pm 21.4 cells) during the first month, then a slower decline (-15.4 \pm 2.2 cells/month) between month 1 and month 12, and a very slow decline (-3.1 \pm 0.8 cells/month) between month 12 and month 36. The median CD4 cell counts at months 12, 24 and 36 were respectively 497/mm³ (376 – 626), 479/mm³ (394 - 628) and 455/mm³ (356 - 587).

Changes in plasma HIV RNA load after treatment interruption are shown in Figure 1b. After a rapid increase from 2.6 to 4.0 log₁₀ copies per ml during the first month, the median plasma HIV RNA level stabilized. Median plasma HIV RNA values at months 12, 24 and 36 were respectively 4.1 log₁₀ copies/mL (3.6 -4.4), 4.1 log₁₀ copies/mL (3.7 - 4.5) and 4.3 log₁₀ copies/mL (3.7 - 4.6). Respectively 5%, 6% and 2% of patients still had plasma HIV RNA values below 400 copies/mL at months 12, 24 and 36 of treatment interruption. Median PBMC HIV DNA load in these patients at baseline and month 6 were respectively 1.0 (1.0 - 1.4) and 2.2 $(1.8 - 2.4) \log_{10}$ per million PBMCs.

Changes in HIV DNA load during the first 18 months of the study are shown in
Figure 1c. The HIV DNA level increased from 2.3 (1.9 -2.8) at baseline to 3.1
(2.7 - 3.4) at 18 months.

24 Incidence of the primary endpoint

Within 36 months after treatment interruption, 23 patients (27.0%) reached a
CD4 cell count below 300/mm³ and 20 of them (22.8%) resumed therapy.
Another nine patients (12.4%) resumed therapy because of CDC stage B or C
events, and 20 patients (20.0%) resumed therapy for other reasons (mainly at
the patient's request).

The nine CDC stage B/C events consisted of severe thrombocytopenia (2 cases), oral hairy leukoplakia (3 patients), oral candidiasis (2 patients) and cutaneous Kaposi's sarcoma (2 patients). One of the two patients who developed cutaneous Kaposi's sarcoma during treatment interruption had already had a first episode prior to enrolment in the study. The CD4 cell counts of these two patients at the onset of Kaposi's sarcoma during treatment interruption were 648/mm³ and 364/mm³. No AIDS-defining events occurred during the first 12 months of treatment interruption.

In the intent-to-treat analysis the proportions of patients who did not meet the criteria for ART resumption at months 12, 24 and 36 were respectively 97.3% (95% CI: 92.5 – 99.1), 86.5% (78.5 – 91.8), and 63.9% (52.9 – 73.6). The proportions of patient who had not resumed ART at months 12, 24 and 36 (observational data) were respectively 92.2% (85.8 – 95.8), 75.3% (66.5 – 82.4), and 55.9% (46.1 – 65.3).

21 Factors predictive of reaching the primary endpoint

The results of univariate and multivariable Cox proportional hazards models are shown in Table 2. In univariate analyses, HIV DNA load above 2.56 log₁₀ per million PBMCs, a low CD4 nadir, the pre-ART CD4 cell count, the baseline CD4

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cell count, and AIDS status at baseline were significantly associated with the time taken to reach the primary endpoint.

In multivariable analyses, HIV DNA load above the upper tercile of 2.56 log₁₀/10⁶ PBMCs (hazards ratio, HR=2.15), a low CD4 nadir (HR=4.59 per 50%) reduction in the CD4 nadir) and pre-existing AIDS status (HR=5.74) were all predictive of a shorter time to the first of two CD4 cell counts below 300/mm³. 15 days apart, or to a CDC stage B or C event. The Kaplan-Meier curves representing the time to the primary endpoint are shown in Figure 2 according to HIV DNA load (above or below 2.56 log₁₀ per million PBMCs) adjusted for the CD4 nadir.

Severe non AIDS-defining morbidity

Four cardiovascular events occurred during follow-up, including two cases of coronary artery disease (CD4 cell counts at diagnosis: 337 and 448), one stroke (CD4=346), and one ruptured cerebral aneurysm (CD4=559); the incidence of cardiovascular events was 1.5/100 person-years (95% CI: 0.6 - 3.2).

No cases of renal or hepatic failure and no non AIDS-related malignancies occurred during follow-up.

Changes in lipodystrophy and metabolic parameters

Changes in anthropometric measurements, lipodystrophy scores and metabolic parameters between day 0 and the last follow-up visit during treatment interruption are shown in Table 3. No significant change in anthropometric values occurred, but the proportion of patients with either lipoatrophy or

lipohypertrophy fell significantly (42% versus 23%, p< 0.001; and 43% versus
 16%, p< 0.001, respectively).

The median changes in the total serum cholesterol level, the serum HDL cholesterol level, and the serum LDL cholesterol level from baseline to the last follow-up visit off treatment were respectively -0.35 mmol/l (IQR -0.99, 0.08) (p < 0.001), -0.11 mmol/l (-0.33, 0.05) (p < 0.001) and -0.25 mmol/l (-0.78, 0.26) (p = 0.001). Serum glucose and triglycerides levels did not change significantly during treatment interruption.

10 Treatment resumption

Of the 47 patients who resumed therapy during the 36 months of follow-up, 31 patients (66%) were prescribed a regimen different from that received prior to enrolment. The median CD4 cell count was 345/mm³ (256 – 446) at resumption, and 475/mm³ (388 – 587) six months later. The median plasma HIV RNA level was 26470 copies/mL (4790 - 81430) at resumption, and 39 patients (81%) had plasma HIV RNA levels below 400 copies/mL six months later. Genotypic mutations conferring resistance to NRTIs and NNRTIs were identified in all 8 patients (19%) who had plasma HIV RNA levels above 400 copies/mL six months after resumption. The baseline profile was not available in three of these patients. Only one of the remaining five patients acquired resistance to a new drug (3TC). At treatment interruption, 6 of these 8 patients had plasma HIV RNA values below 400 copies/mL. The ART regimens were subsequently optimized, and plasma HIV RNA became undetectable (<400 copies/mL) after a further 6 months in 7 of the 8 patients.

25 No B or C events were diagnosed after treatment resumption.

1 Discussion

This study indicates that treatment interruption in selected patients with a high CD4 cell nadir is associated with a low rate of HIV disease progression after 36 months. Only 36% of the patients met the criteria for treatment resumption during the 36 months of follow-up, namely a CD4 cell count below 300/mm³ or a CDC stage B or C event. Only two CDC grade C clinical events (two cases of cutaneous Kaposi's sarcoma) occurred during the 36 months of follow-up. It should be noted that patients with AIDS were eligible for this study. One of the two patients who developed Kaposi's sarcoma during follow-up, when their CD4 cell counts were above 350/mm3, had already had an episode of Kaposi's sarcoma prior to inclusion. Both these events occurred after one year off treatment, and the incidence was 0.7/100 P-Y. This is within the range of incidence rates observed elsewhere in antiretroviral-naive patients with CD4 cell counts >350/mm³ [Guiguet et al., 2008; Phillips et al., 2007]. Guiguet et al. reported an incidence rate of 0.8/100 P-Y in such patients [Guiguet et al., 2008]. Our findings are in keeping with those of other studies of treatment interruption in similar populations [Ananworanich et al., 2006; Boschi et al., 2004; Maggiolo et al., 2008; Mata et al., 2005; Mussini et al., 2005; Pogany et al., 2007; Skiest et al., 2004; Skiest et al., 2007; Wit et al., 2005]. Among the 284 patients included in the treatment interruption arm of the Staccato trial [Ananworanich et al., 2006], only minor HIV-related signs and symptoms were observed, with no AIDS-defining events or AIDS-related deaths. In a recent trial [Maggiolo et al., 2008], 329 patients with a CD4 cell nadir above 200/mm³ and a CD4 cell count above 700/mm³ at inclusion were randomized to continue their ongoing antiretroviral treatment or to discontinue it until their CD4 cell count fell to

350/mm³. During 689 P-Y of follow-up, only one AIDS-defining event occurred in the STI arm and none in the control arm, while eight cases of pneumonia occurred in the STI arm, compared to one in the control arm. However, recent large studies have shown a higher rate of clinical events in patients undergoing CD4-guided treatment interruption than in patients who continue on treatment [Danel et al., 2006; El-Sadr et al., 2006]. These studies did not exclude patients with AIDS or HIV-related disorders, and included patients with different degrees of immune impairment and low CD4 cell nadirs. One study in which there was a high rate of events (mainly bacterial infections) during treatment interruption involved patients in Africa, where the background rate of opportunistic infections may be higher [Danel et al., 2006].

Among the patients enrolled in the SMART trial, opportunistic infections and deaths were significantly more frequent in the "drug conservation" arm, in which ART initiation or resumption was deferred until the CD4 cell count fell to 250/mm³), than in the continuous ART arm. Even in the subgroup of patients whose last CD4 cell count was above 350/mm³, the incidence of AIDS-related events in the "drug conservation" arm was 1.28 per 100 person-years [Lundgren] et al., 2008], a rate similar to that found in the present study (0.7/100 P-Y;IQR 0.2 - 2.0). However, clinical events were infrequent in our study, in which the patients had started first-line antiretroviral therapy with CD4 cell counts above 350/mm³. This further suggests that lengthy treatment interruption is less risky in AIDS-free patients with a high CD4 cell nadir.

The SMART study highlighted an increased risk of cardiovascular events during
treatment interruption [EI-Sadr et al., 2006]. A slight increase in cardiovascular
events was observed in the "drug conservation" arm of the SMART study, with

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an incidence of 1.3/100 P-Y, compared to 1.5/100 P-Y in this study. As HIV
replication induces immune activation, plasma HIV RNA rebound during
treatment interruption might lead to inflammation and endothelial dysfunction,
thereby increasing the risk of cardiovascular events [Blum et al., 2005; Solages
et al., 2006; Tebas et al., 2008].

As also observed elsewhere [Hatano et al., 2000], a significant improvement of
lipodystrophy and lipid abnormalities was observed during treatment
interruption. However, changes in lipodystrophy were assessed with an
unvalidated clinical score, and no significant change in anthropometric values
was observed during treatment interruption.

As reported previously by others, this study showed a rapid initial CD4 cell decline after ART interruption, followed by a more gradual loss during the first months (accompanied by a rapid increase in plasma HIV RNA), and then by CD4 cell count and plasma HIV RNA load stabilization [Boschi et al., 2004; Fagard et al., 2005; Maggiolo et al., 2004; Skiest et al., 2007; Tarwater et al., 2003; Tebas et al., 2002; Thiebaut et al., 2005; Wit et al., 2005]. The present study showed an increase in HIV DNA levels after treatment interruption, indicative of gradual replenishment of cellular reservoirs.

In Cox proportional hazards multivariable analyses, baseline factors predictive of outcome 36 months after treatment interruption were PBMC HIV DNA load above the upper tercile (2.56 log₁₀ copies/ml), the CD4 nadir, and AIDS status. It is commonly reported that the higher the CD4 cell nadir, the smaller the fall in the CD4 cell count and the longer the duration of treatment interruption [Boschi et al., 2004; Fernandez Guerrero et al., 2005; Giuntini et al., 2005; Maggiolo et al., 2004; Mata et al., 2005; Mussini et al., 2005; Poulton et al., 2003; Skiest et

al., 2004; Skiest et al., 2007; Tarwater et al., 2003; Thiebaut et al., 2005]. As also found in some other studies, the pre-ART plasma HIV RNA level was not predictive of the duration of treatment interruption in this study [Maggiolo et al., 2004; Tebas et al., 2002]. Interestingly, the PBMC HIV DNA levels above 2.56 log₁₀ copies/10⁶ PBMCs were an independent predictor of a shorter time to treatment resumption. PBMC HIV DNA load was also found to be a significant predictor of treatment resumption in a pilot study involving 62 patients who stopped their ongoing antiretroviral treatment [Sarmati et al., 2009]. The HIV-DNA level provides an estimate of the proportion of circulating HIV-infected cells and reflects the size of the viral reservoir. Central and transitional memory CD4+ T cells contribute to the HIV reservoir, the size of which correlates with the nadir CD4 cell count [Chomont et al., 2009]. PBMC HIV-1 DNA load shortly after infection is predictive of disease progression, independently of the plasma HIV-RNA level and the CD4 cell count [Goujard et al., 2006; Rouzioux et al., 2005]. The HIV reservoir is established soon after infection, and the degree to which it is diminished by ART, whether initiated during the acute or chronic phase of infection, is variable [Ngo-Giang-Huong et al., 2001; Strain et al., 2003]. While all patients in this study had low or undetectable plasma HIV-RNA at inclusion, their HIV-DNA levels varied over a broad range and were therefore informative. These results suggest that the larger the viral reservoir at the time of treatment interruption, the less time the patient can remain off treatment. In other words, a high level of HIV-DNA was indicative of a risk of progression in case of treatment interruption, whatever the prior duration of ART. Following the results of the SMART trial, treatment interruptions are not

25 currently recommended in HIV-infected patient management guidelines

[Hammer et al., 2008; Yeni, 2008]. In current practice, the number of patients who stop taking ART, for whatever reason, has fallen since the results of the SMART trial were first reported in January 2006, but still represented 5.7% of patients monitored in 2007 [Yeni, 2008]. Thus, it remains important to estimate the risk of AIDS and severe non-AIDS morbidity/mortality during treatment interruption, and to identify factors associated with the duration of interruption. The results of this study indicate that patients with a history of AIDS-defining events should not interrupt their treatment. In addition to a low CD4 nadir, a high PBMC HIV DNA level at time of treatment interruption is a strong predictor of more rapid treatment resumption, contrary to the pre-therapy plasma HIV-1

11 RNA level.

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Table 1: Characteristics of the study population

Ν		116		
Men, n (%)		73 (63%)		
Age (years), median (IQR)		39 (25 – 80)		
Transmission group, n (%)	Homo/bisexual men	57 (49%)		
	Heterosexuals	46 (40%)		
	Intravenous drug users	9 (8%)		
AIDS status, n (%)		3 (3%)		
CD4 nadir (cells/mm ³)				
- median (IQR)		381 (341 – 490)		
- CD4 nadir < $300/\text{mm}^3$, ,		
- GD4 nadir < 300/mm		13 (11%)		
CD4 cell count/mm ³ at inclu	sion, median (IQR)	806 (676 – 940)		
Pre-ART plasma HIV-RNA ((log ₁₀ copies/mL), median (IQR)	4.2 (3.8 – 4.5)		
Plasma HIV-RNA < 400 cop	ies/mL at inclusion, n (%)	106 (91%)		
HIV DNA (log ₁₀ copies/10 ⁶ F	PBMCs) at inclusion			
- median (IQR)		2.3 (1.9 – 2.8)		
- HIV-DNA > 2.56 log		38 (33%)		
Duration of ART (years), me	edian (IQR)	5.3 (4.2 – 6.1)		
Triple nucleoside-based reg	imen, n (%)	39 (34%)		
Double nucleoside-based re	egimen, n (%)	17 (14%)		
Non nucleoside-based regin	nen, n (%)	41 (35%)		
Protease inhibitor-based rec	18 (16%)			
Non nucleoside/protease inl	1 (1%)			

Table 2: Risk factors of reaching the criteria for treatment resumption in

univariate analysis and multivariable Cox proportional hazards models

7 8			Univariate	analysis		Multivariable anal	ysis
9 Variables 10 11		Number of (patients, events)	Hazard Ratio (CI 95)	р	AIC	Hazard Ratio (CI 95)	р
12 Total		(N=116, n=32)					
13 14 ^{Age (year)}			1.00 (0.96 – 1.03)	0.779			
15 Gender 16	Women Men	(N=43, n=12) (N=73, n=20)	1 (reference) 0.99 (0.48 – 2.08)	0.966			
17 18 19	No Yes	(N=59, n=16) (N=57, n=16)	1 1.04 (0.52 – 2.09)	0.905			
20 Duration of ART			0.96 (0.76 – 1.21)	0.709			
21 Type of ART 22	cART dual therapy	(N=99, n=31) (N=17, n=1)	1 0.17 (0.02 – 1.27)	0.085			
23 24 Prior AIDS status			6.32 (1.85 – 21.64)	0.003		5.74 (1.60 – 20.56)	0.007
25 Pre-ART CD4 (log2)§			7.14 (1.43 – 33.33)	0.017	270.4		
26 27 Pre-ART CD4 cells/mm3				0.116	272.8		
28 29	>519 405-519 <405	(N=39, n=8) (N=40, n=10) (N=37, n=14)	1 1.35 (0.53 – 3.43) 2.40 (1.00 – 5.72)	0.524 0.049			
30 31 Pre-ART VL (log10) [#]			2.04 (0.85 - 4.90)	0.109	271.7		
32 Pre-ART VL copies/ml							
33	<10000	(N=41, n=9)	1	0.324	272.4		
34 35	10000-27000 >27000	(N=36, n=9) (N=36, n=14)	1.35 (0.53 – 3.40) 1.89 (0.82 – 4.36)	0.529 0.138			
36 CD4 at inclusion (log2) § 37		(· · · /	3.03 (0.93 – 10.10)	0.065	273.6		
38 CD4 at inclusion cells/mm3				0.035	270.3		
39	>885 700-885	(N=35, n=7) (N=39, n=8)	1 1.22 (0.44 – 3.37)	0.700			
40 41	<700	(N=42, n=17)	2.77 (1.15 – 6.71)	0.024			
41 42 CD4 nadir (log2) [§]			6.25 (1.78 – 25.00)	0.004	268.8	4.59 (1.22 – 17.24)	0.024
43							
44 CD4 nadir cells/mm3	>448	(N=38, n=7)	1	0.169	273.0		
45 46	361-448	(N=40, n=12)	2.10 (0.82 - 5.35)	0.120			
47	<361	(N=38, n=13)	2.35 (0.94 – 5.89)	0.069			
48 HIV DNA (log10) copies/million 49 PBMCs			1.94 (1.13 – 3.35)	0.017	251.9		
50 HIV DNA (log10)	0.17	(NL 00 x 7)	4	0.031	250.8		
51 52	<2.17 2.17-2.56	(N=39, n=7) (N=36, n=6)	1 0.96 (0.32 – 2.86)	0.944			
53	>2.56	(N=38, n=17)	2.60 (1.08 – 6.27)	0.034			
54 HIV DNA (log10) > 2.56		(N=38, n=17)	2.65 (1.28 – 5.45)	0.008		2.15 (1.02 – 4.53)	0.044
55§ Hazard Rat	io per two-fold c	lecrease in the CD4	l cell count; [#] Hazard R	atio per t	en-fold inc	rease in plasma	
57			events during follow-up	•		·	
59	ct models (the be	est model has the lo	owest AIC).				

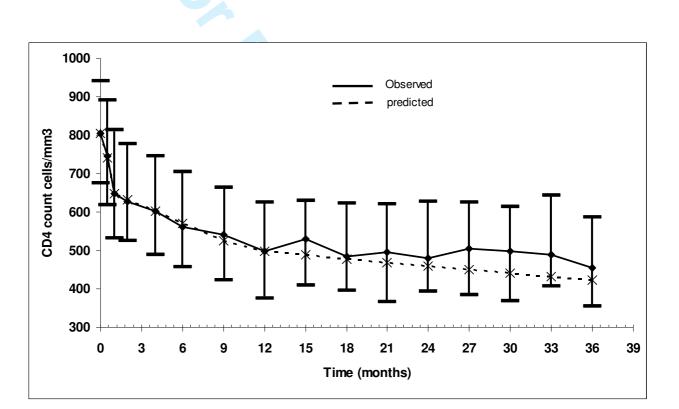
Table 3: Change in anthropometric measures, the lipodystrophy score andmetabolic parameters during treatment interruption

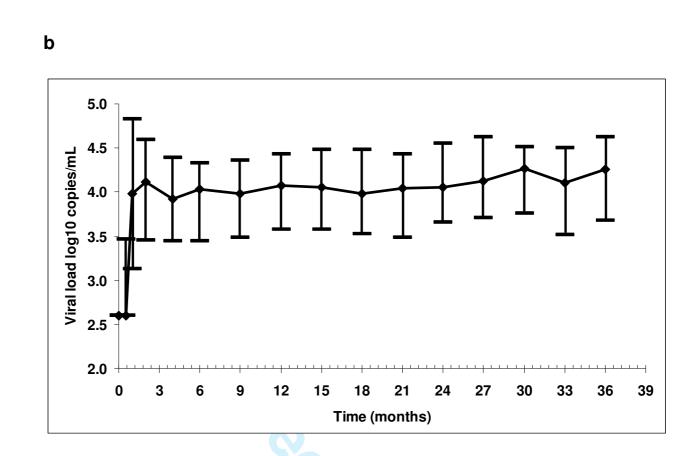
	Baseline	Last values	
		during	р
		interruption	
Hip circumference (cm), median (IQR)	90.8 (86.0 – 90.8)	91.0 (86.8 – 97.0)	0.010
Waist (cm)	83.0 (76.0 - 90.0)	83.0 (76.0 – 91.3)	0.251
Waist/hip ratio	0.92 (0.87 – 0.99)	0.91 (0.87 – 0.97)	0.203
Thigh circumference (cm)	47.0 (44.0 – 50.0)	47.0 (44.0 – 51.0)	0.258
Body mass index (kg/m ²)	22.9 (20.7 – 24.5)	22.8 (20.7 – 25.1)	0.468
Lipodystrophy score, median (IQR)	1.0 (0.0 – 4.0)	0.0 (0.0 - 1.0)	<0.00
- atrophy ≥1, n (%)	49 (42%)	25 (23%)	<0.00
- hypertrophy ≥1, n (%)	50 (43%)	18 (16%)	<0.00
Total fasting cholesterol serum level (mmol/l), median (IQR)	5.10 (4.38 – 5.74)	4.58 (4.11 – 5.22)	<0.00
Fasting HDL cholesterol serum level (mmol/l), median (IQR)	1.22 (1.02 – 1.49)	1.13 (0.95 – 1.36)	<0.00
Fasting LDL cholesterol serum level (mmol/l), median (IQR)	3.08 (2.48 - 3.72)	2.82 (2.36 – 3.33)	0.001
Fasting triglyceride serum level (mmol/l), median (IQR)	1.18 (0.80 – 1.86)	1.08 (0.78 – 1.60)	0.076
Fasting glucose serum level (mmol/l), median (IQR)	5.00 (4.54 - 5.40)	4.94 (4.43 - 5.50)	0.928

Figure 1: Changes in the CD4 cell count (a) and plasma HIV-RNA (b) during treatment interruption and in PBMC HIV DNA during the first 18 months of treatment interruption (c)

Data points represent median values and whiskers represent interquartile ranges in (a) and (b). The dotted curve in (a) represents predicted changes in the CD4 cell count predicted by mixed linear models. The box plots in (c) represent median and interquartile ranges and whiskers indicate minimum and maximum values.



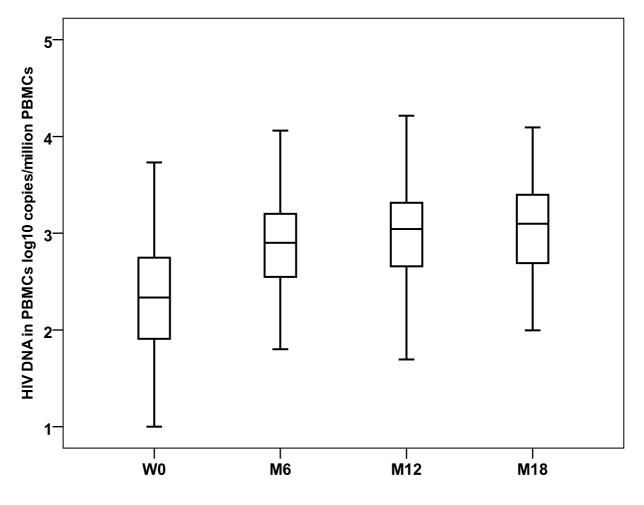




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John Wiley & Sons Figure 2: Kaplan-Meier curve of time to the endpoint according to the HIV DNA level adjusted for the CD4 cell nadir The curves represent the time taken to reach the criteria for treatment

resumption in patients with baseline HIV DNA load below and above 2.56 log copies/10⁶ PBMCs (upper tercile).

