MAJOR ARTICLE

Impact of Obefazimod on Viral Persistence, Inflammation, and Immune Activation in People With Human Immunodeficiency Virus on Suppressive Antiretroviral Therapy

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Background. Persistence of viral reservoirs has been observed in people with human immunodeficiency virus (HIV), despite long-term antiretroviral therapy (ART), and likely contributes to chronic immune activation and inflammation. Obefazimod is a novel drug that inhibits human immunodeficiency virus type 1 (HIV-1) replication and reduces inflammation. Here we assess whether obefazimod is safe and might impact HIV-1 persistence, chronic immune activation, and inflammation in ARTsuppressed people with HIV.

Methods. We evaluated obefazimod-related adverse events, changes in cell-associated HIV-1 DNA and RNA, residual viremia, immunophenotype, and inflammation biomarkers in blood and rectal tissue. We compared 24 ART-suppressed people with HIV who received daily doses of 50 mg obefazimod for 12 weeks $(n = 13)$ or 150 mg for 4 weeks $(n = 11)$ and 12 HIV-negative individuals who received 50 mg for 4 weeks.

Results. The 50- and 150-mg doses of obefazimod were safe, although the 150-mg dose showed inferior tolerability. The 150-mg dose reduced HIV-1 DNA (*P* = .008, median fold change = 0.6) and residual viremia in all individuals with detectable viremia at baseline. Furthermore, obefazimod upregulated miR-124 in all participants and reduced the activation markers CD38, HLA-DR, and PD-1 and several inflammation biomarkers.

Conclusions. The effect of obefazimod by reducing chronic immune activation and inflammation suggests a potential role for the drug in virus remission strategies involving other compounds that can activate immune cells, such as latency-reversing agents.

Keywords. HIV-1 persistence; immune activation; inflammation; miR-124; obefazimod (ABX464).

Antiretroviral therapy (ART) has been successful in maintaining long-term viral suppression in people with human immunodeficiency virus (PWH). However, the persistence of latently infected cells carrying intact, inducible proviruses has been observed in different cellular and anatomical reservoirs [\[1–3](#page-10-0)]. Activation of latently infected cells can lead proviral

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DNA to be transcriptionally active and capable of producing new viral particles [[4,](#page-10-0) [5](#page-10-0)], which may contribute to residual viremia and thus persistence of immune activation and inflammation, despite long-term ART suppression [[6](#page-10-0)]. Because viral persistence is associated with a variety of biological mechanisms, the success of human immunodeficiency virus (HIV) remission strategies will most likely depend upon combinatorial strategies [\[7](#page-10-0)]. Thus, new antiretroviral drugs aimed at preventing viral reactivation from latently infected cells need to be explored, including those that target viral transcription, processing, and/or maturation of viral transcripts [\[8\]](#page-10-0), which could simultaneously help reduce chronic immune activation and inflammation exhibited by ART-suppressed PWH.

Obefazimod (formerly ABX464) is an investigational small quinoline with a novel mechanism of HIV type 1 (HIV-1) inhibition based on interference with RNA biogenesis. In brief, by binding to the cap-binding complex (CBC), obefazimod targets

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viral Rev protein and prevents Rev-mediated nuclear export and maturation of unspliced viral RNA [\[9\]](#page-10-0). Obefazimod has been shown to be highly effective at inhibiting HIV-1 replication in human peripheral blood mononuclear cells (PBMCs) and macrophages in vitro. Moreover, obefazimod reduced viral replication in HIV-1–infected humanized mice and delayed viral rebound after drug discontinuation [\[9\]](#page-10-0). Furthermore, obefazimod demonstrated safety, tolerability, and potential reduction of viral replication in previous clinical trials [\[10–13\]](#page-10-0).

Of note, the CBC targeted by obefazimod is also involved in the biogenesis of endogenous small noncoding RNAs (miRNAs) such as miR-124. During miRNA processing, obefazimod specifically enables a quick release of miR-124 through splicing of its long noncoding RNA precursor (lncRNA 0599-205) [\[14](#page-10-0)]. Through the targeting of STAT3 mRNA, miR-124 functions as a negative regulator of inflammation, attenuating the production of proinflammatory cytokines to help maintain homeostasis, and is reported to be a critical modulator of immunological and inflammatory disorders [[15,](#page-11-0) [16\]](#page-11-0). Obefazimod has demonstrated strong anti-inflammatory effects by the selective induction of miR-124 in in vitro and in vivo studies, including in PWH [\[14](#page-10-0), [16](#page-11-0), [17](#page-11-0)]. Hence, due to these properties, this novel drug might be a promising therapeutic strategy to impact the HIV-1 reservoir and reduce chronic immune activation and inflammation in the context of sustained inflammation during ART, as well as in virus remission strategies involving other compounds that can activate immune cells. The present study aimed to further evaluate safety and impact of obefazimod on HIV-1 persistence and chronic immune activation and inflammation in blood and gut-associated lymphoid tissue (GALT) from ART-suppressed PWH.

METHODS

Study Design and Participants

This was a nonrandomized, open-label, phase 2b/2a clinical trial designed to evaluate safety, virological, and immunological parameters in 36 adult participants (24 PWH and 12 HIV-negative) at either 50 mg $(n = 13)$ once-daily oral doses of obefazimod for 12 weeks (HIV_{50mg/12w}) or 150 mg (n = 11) for 4 weeks ($HIV_{150mg/4w}$) or, in the case of HIV-negative individuals (n = 12), 50 mg for 4 weeks (HN_{50mg/4w}).

Seropositive participants were required to be on an integrase inhibitor–based regimen for at least 12 months prior to enrollment with undetectable plasma viremia (≤50 HIV-1 RNA copies/mL) during the 6 months prior to screening. In addition, they were required to have $CD4^+$ T-cell counts $>$ 250 cells/ μ L since diagnosis with at least 600 cells/μL at screening. All study participants were recruited from the University Hospital Germans Trias i Pujol (Badalona, Spain) from April 2017 to December 2018, and all provided signed informed consent. The study protocol (ABX464-005) was approved by the hospital ethics committee (AC-16-047-CEIM) and the Spanish Agency of Medicines and Medical Products (EudraCT: 2016-002797-12) and registered at ClinicalTrials.gov (identifier NCT02990325).

Safety Assessments

Safety and tolerability were evaluated based on individual values for clinical data (including signs and symptoms, laboratory toxicities, and clinical events), physical examinations, vital signs, and number and grade of treatment-emergent adverse events (AEs) at each visit as described in [Figure 1](#page-2-0)*A*. The data and safety monitoring board reviewed the safety data to recommend, if appropriate, enrollment of additional individuals and the continuation of the study.

Sample Collection and Analysis

Blood and GALT samples [\[18](#page-11-0)] were collected at baseline, end of intervention (EoI), and 4 weeks after drug discontinuation (EoI + 4w) from each participant in accordance with the schedule of study events [\(Figure 1](#page-2-0)*A*). Total HIV-1 DNA [\[18\]](#page-11-0), cell-associated (CA) HIV-1 RNA [[19](#page-11-0)], and ultrasensitive residual viremia [\[20](#page-11-0)] were measured to estimate changes in viral persistence. To evaluate the state of immune activation, T-cell subset distribution and expression of cell surface activation markers were measured by flow cytometry. Moreover, miR-124 was quantified in rectal tissue to evaluate intestinal inflammation, and 26 inflammation biomarkers were measured in plasma to assess systemic inflammation. Specific details on these assays and statistical analysis can be found in the [Supplementary Material.](http://academic.oup.com/jid/article-lookup/doi/10.1093/infdis/jiad251#supplementary-data)

RESULTS

Participant Characteristics and Disposition

Thirty-six participants were prospectively and sequentially enrolled into 3 groups as follows: 13 ART-suppressed PWH received 50 mg of obefazimod orally once daily for 12 weeks (HIV50mg/12w group), 11 ART-suppressed PWH received 150 mg for 4 weeks (HIV_{150mg/4w} group), and 12 HIV-negative individuals received 50 mg for 4 weeks ($HN_{50mg/4w}$ group) as shown in [Figure 1](#page-2-0)*B*. All study participants were White cisgender men with a median age of 44 years. Baseline demographic and clinical characteristics are summarized in [Table 1](#page-3-0). No differences between groups were observed, except for the time on viral suppression, which was longer in the $HIV_{150mg/4w}$ group than in the $HIV_{50mg/12w}$ group ($P = .02$). All participants were on suppressive integrase inhibitor–based ART at study initiation.

Obefazimod Is Safe Either Alone or in Combination With ART

Just over half of participants (53%) experienced any AE during obefazimod administration [\(Table 2](#page-4-0)). However, no AEs were severe (grade 3 or above), life-threatening, or disabling, and all symptoms and biochemical abnormalities normalized upon drug withdrawal. Only 3 participants, 1 in the HIV_{50mg}

A

Figure 1. Study design and flow diagram. A, Schedule of study events. B, Consort flow diagram. Abbreviations: ART, antiretroviral therapy; AE, adverse event; ART, antiretroviral therapy; CA, cell-associated; HIV-1, human immunodeficiency virus type 1; HIV_{50mg/12w}, HIV infected taking once-daily oral obefazimod 50 mg for 12 weeks; HIV_{150mg/4w}, HIV infected taking once-daily oral obefazimod 150 mg for 4 weeks; HN_{50mg/4w}, HIV negative taking once-daily oral obefazimod 50 mg for 4 weeks; VL, viral load.

Data are presented as median (interquartile range) unless otherwise indicated. *P* values ≤ .05 are shown in bold.

Abbreviations: ART, antiretroviral therapy; HIV⁻, human immunodeficiency virus negative; HIV⁺, human immunodeficiency virus positive; VL, viral load.

 $_{12w}$ group and 2 in the HIV_{150mg/4w} group, discontinued the study due to AEs and were excluded from subsequent analyses. The most frequently reported AEs were nervous system disorders (headache) (42%), gastrointestinal disorders (abdominal pain, diarrhea, and nausea) (39%), and musculoskeletal pain (back pain and myalgia) (19%). The total number of AEs was slightly higher at the 150-mg dose compared to the 50-mg dose (*P* = .3) [\(Table 2](#page-4-0)). Nevertheless, standard plasma viral load, and $CD4^+$ and $CD8^+$ T-cell counts showed no significant changes during obefazimod treatment, which suggests that obefazimod does not disrupt clinical control of HIV-1 infection and therefore does not negatively interact with ART [\(Supplementary Figure 1\)](http://academic.oup.com/jid/article-lookup/doi/10.1093/infdis/jiad251#supplementary-data).

Obefazimod Reduces HIV-1 Persistence

To evaluate the impact of obefazimod on proviral reservoir size, we measured total HIV-1 DNA in peripheral CD4⁺ T cells by droplet digital PCR (ddPCR). Total HIV-1 DNA was detected in all individuals. Data confirmed a significant reduction of total HIV-1 DNA at the 150-mg dose $(P = .008, \text{ median fold}$ change = 0.6) [\(Figure 2](#page-5-0)*A* and 2*B*). Indeed, the number of individuals with decreased total HIV-1 DNA was higher at the 150-mg dose compared to the 50-mg dose (88.9% vs 50%, respectively; *P* = .159, Fisher exact test) ([Figure 2](#page-5-0)*C* and 2*[D](#page-5-0)*). However, this was a transient effect and total HIV-1 DNA returned to basal levels after drug discontinuation [\(Supplementary Figure 2](http://academic.oup.com/jid/article-lookup/doi/10.1093/infdis/jiad251#supplementary-data)*A–D*). These results suggest that

obefazimod reduces total HIV-1 DNA in a dose-dependent manner, potentially impacting viral persistence.

To assess the effect of obefazimod on viral transcription, we measured CA HIV-1 RNA in peripheral $CD4^+$ T cells by reverse-transcription (RT) ddPCR. CA HIV-1 RNA was detected in all individuals. Data showed no consistent effect of obefazimod on viral transcription during the administration period (Figure 2*E–*2*H*). These data suggest that obefazimod did not significantly affect viral transcription initiation (as measured by our targeting of the 5′ end of the HIV-1 mRNA transcripts). However, CA HIV-1 RNA significantly increased after drug discontinuation at the 150-mg dose ($P = .02$, median fold chan $ge = 2.1$) and the 50 mg ($P = .02$, median fold change = 1.2) [\(Supplementary Figure 2](http://academic.oup.com/jid/article-lookup/doi/10.1093/infdis/jiad251#supplementary-data)*E* and 2*[F](#page-5-0)*). Indeed, the number of individuals who increased CA HIV-1 RNA after discontinuation was higher at the 150-mg dose compared to the 50-mg (100% vs 75%, respectively) [\(Supplementary Figure](http://academic.oup.com/jid/article-lookup/doi/10.1093/infdis/jiad251#supplementary-data) *2G* and 2*H*).

To evaluate the impact of obefazimod on residual viremia, we measured plasma viral load (VL) using an ultrasensitive VL (usVL) test (limit of detection <1 HIV-1 RNA copy/mL). Residual viremia was detectable in 10 (56%) individuals at baseline (6 in the HIV_{50mg/12w} group and 4 in the HIV_{150mg/4w} group) ([Figure 2I and 2](#page-5-0)*J*). Longitudinal analysis confirmed a substantial reduction of plasma HIV-1 RNA in all (100%) individuals who showed detectable residual viremia at baseline at the 150-mg dose compared to the 50-mg (56%) (Figure 2*I*–2*L*). Nonetheless, this effect was lost after drug discontinuation [\(Supplementary Figure 2](http://academic.oup.com/jid/article-lookup/doi/10.1093/infdis/jiad251#supplementary-data)*I–*2*L*). The very low levels of plasma

Data are presented as No. (%).

Abbreviation: AE, adverse event; HIV, human immunodeficiency virus; HIV⁻, human immunodeficiency virus negative; HIV⁺, human immunodeficiency virus positive.

HIV-1 RNA at baseline and high proportion of undetectable samples might have interfered to detect significant changes. Nevertheless, these results suggest that obefazimod reduces residual viremia in a dose-dependent manner.

Obefazimod Reduces Chronic Immune Activation in PWH

To assess the effect of obefazimod on immune activation, first we measured CD4⁺ T-cell subset distribution in PBMCs by flow cytometry. The dynamics of the CD4⁺ T-cell subsets did not significantly change throughout obefazimod administration, which suggests that there is no redistribution of cellular subsets [\(Figure 3](#page-6-0)*A*).

Then, the dynamics of cell surface activation markers (CD38⁺HLA-DR⁺ and PD-1⁺ cells) were analyzed in peripheral and rectal CD4⁺ and CD8⁺ T cells. Data showed no baseline differences between PWH and HIV-negative individuals except for lower proportion of rectal CD4⁺PD-1⁺ cells in HIV-negative individuals compared to the $HIV_{150mg/4w}$ group ($P = .04$; [Supplementary Table 1\)](http://academic.oup.com/jid/article-lookup/doi/10.1093/infdis/jiad251#supplementary-data). Data confirmed a significant reduction of peripheral CD8⁺PD-1⁺ and CD4⁺CD38⁺HLA-DR⁺ cells at week 12 (EoI) and $CD4+PD-1+$ cells at week 16 (EoI + 4w) in the HIV_{50mg/12w} group ($P = .007$, $P = .04$, and $P = .02$, respectively) ([Figure 3](#page-6-0)*B*, 3*F*, and 3*[G](#page-6-0)*; [Supplementary Table 1\)](http://academic.oup.com/jid/article-lookup/doi/10.1093/infdis/jiad251#supplementary-data). The fact that the reduction of immune activation in peripheral $CD4^+$ and $CD8^+$ T cells is only significant when obefazimod is given for at least 12 weeks suggests a lasting effect of obefazimod at reducing immune activation in ART-suppressed PWH. Of note, these results were reinforced by the findings

Figure 2. Effect of obefazimod on viral reservoir throughout the intervention period. $A-D$, Changes in total human immunodeficiency virus type 1 (HIV-1) DNA from peripheral CD4+ T cells measured by droplet digital PCR (ddPCR). *E–H*, Changes in cell-associated HIV-1 RNA from peripheral CD4+ T cells measured by reverse-transcription ddPCR. *I-L*, Changes in residual plasma viremia measured using an ultrasensitive viral load test. Undetectable plasma viral load was defined as <0.55 HIV-1 RNA copies/mL (broken line). Data from 3 individuals in the HIV-infected group taking 50 mg for 12 weeks were lost due to technical problems. *A–B*, *E–F*, *I–J* panels represent absolute values of each timepoint in each group. *C–D*, *G–H*, *K–L* panels represent waterfall charts to show the increase/decrease from baseline to end of obefazimod administration. Each symbol/shape represents a different individual. Horizontal lines in panels *A–B*, *E–F*, and *I–J* represent medians. Asterisks denote significant differences between timepoints detected using the Wilcoxon signed-rank test (**P ≤ .01). Abbreviations: BL, baseline; EoI, end of intervention; HIV-1, human immunodeficiency virus type 1; HIV_{50mg/12w}, HIV infected taking once-daily oral obefazimod 50 mg for 12 weeks; HIV_{150mg/4w}, HIV infected taking once-daily oral obefazimod 150 mg for 4 weeks; TBP, TATA-binding protein.

in GALT, as we observed a significant reduction in rectal CD4⁺CD38⁺HLA-DR⁺ and CD8⁺CD38⁺HLA-DR⁺ cells at week 8 (EoI + 4w) in the HIV_{150mg/4w} group ($P = .008$, median fold change = 1.9; and $P = .02$, median fold change = 2.4, respectively) ([Figure 3](#page-6-0)*D* and 3*E*; [Supplementary Table 1\)](http://academic.oup.com/jid/article-lookup/doi/10.1093/infdis/jiad251#supplementary-data). However, obefazimod intervention transiently increased rectal $CD8⁺CD38⁺HLA-DR⁺$ cells in the $HIV_{50mg/12w}$ group (*P* = .03) and $CD4+PD-1^+$ cells in HIV-negative individuals $(P=.01)$ [\(Figure 3](#page-6-0)*E* and 3*H*; [Supplementary Table 1](http://academic.oup.com/jid/article-lookup/doi/10.1093/infdis/jiad251#supplementary-data)). Altogether, these data suggest that obefazimod may reduce chronic immune activation in PWH.

Obefazimod Reduces Chronic Inflammation in PWH

To further investigate the effect of obefazimod on intestinal and systemic inflammation, we measured expression of miR-124 in rectal tissue by qRT-PCR and plasma levels of 26 inflammation

biomarkers using Luminex-based multiplex immunoassay and enzyme-linked immunosorbent assay. miR-124 was detected in all individuals. Basal expression of miR-124 was higher in the $HIV_{150mg/4w}$ group compared to the $HIV_{50mg/12w}$ ($P = .02$; [Figure 4](#page-7-0)*A*), perhaps because they have been on suppressive ART longer. Nevertheless, miR-124 expression significantly increased in all (100%) participants from all groups ($HN_{50mg/4w}$, $P = 0.0005$, median fold change = 2.2; HIV_{50mg/12w}, $P = 0.0005$, median fold change = 4.0; $HIV_{150mg/4w}, P = .004$, median fold change $= 6.7$) [\(Figure 4](#page-7-0)*B*). However, this effect was reversed after drug discontinuation. These data suggest that obefazimod may reduce intestinal inflammation by the selective upregulation of miR-124 in ART-suppressed PWH.

Regarding soluble inflammation biomarkers, interleukin (IL) 4, IL-6, IL-8, IL-12 (p40), IL-22, and liver-type fatty acid-binding protein were excluded from the analysis due to

Figure 3. Effect of obefazimod on T-cell activation. A, Upper graph shows the relative frequency of different CD4⁺ T-cell subsets in the study groups identified in peripheral blood mononuclear cells by flow cytometry. Bars represent medians of each subset. *B–I*, Immune activation levels determined as percentages of CD38+ HLA-DR+ (*B–E*) and PD-1⁺ (*F–I*) cells analyzed both in peripheral and rectal CD4⁺ and CD8⁺ T-cell populations by flow cytometry. *B* and *C*, Frequency of CD38⁺HLA-DR⁺ cells from peripheral CD4⁺ and CD8⁺ T cells. *D* and *E*, Frequency of CD38⁺HLA-DR⁺ cells from rectal CD4⁺ and CD8⁺ T cells. *F* and *G*, Frequency of PD-1⁺ cells from peripheral CD4⁺ and CD8⁺ T cells. *H* and CD8⁺ *I*, Frequency of PD-1+ cells from rectal CD4+ and CD8+ T cells. Median and interquartile range are represented. Asterisks denote significant differences between timepoints detected using the Wilcoxon signed-rank test (**P* ≤ .05; ***P* ≤ .01). Abbreviations: BL, baseline; EoI, end of intervention; EOI + 4w, 4 weeks after end of intervention; HIV_{50mg/} $_{12}$ w, HIV infected taking once-daily oral obefazimod 50 mg for 12 weeks; HIV_{150mg/4w}, HIV infected taking once-daily oral obefazimod 150 mg for 4 weeks; HN_{50mg/4w}, HIV negative taking once-daily oral obefazimod 50 mg for 4 weeks; T_{CM}, central memory T cells; T_{EM}, effector memory T cells; T_{EMRA}, effector memory RA T cells; T_N, naive T cells; T_{TM} , transitional memory T cells.

the high proportion of undetectable samples. Among the remaining 20 biomarkers, data showed a significant downregulation of granulocyte colony-stimulating factor (G-CSF), granulocyte-macrophage colony-stimulating factor (GM-CSF), CD14, and IL-7 in HIV-negative individuals $(P = .003, P = .01,$ $P = .02$, and $P = .05$, respectively); IL-10 in the HIV_{50mg/12w} group $(P = .05)$; and macrophage inflammatory protein-1beta (MIP-1β; CCL4) and monocyte chemoattractant protein-1 (MCP-1; CCL2) in the $HIV_{150me/4w}$ group ($P = .01$ and $P = .05$, respectively) throughout obefazimod administration [\(Figure 5](#page-8-0)*A*). These results reinforce the anti-inflammatory effect of obefazimod in both PWH and HIV-negative individuals.

Unexpectedly, C-reactive protein (CRP) significantly increased in the HIV_{150mg/4w} group ($P = .008$), although mean values were always <10 mg/L, the upper limit of the normality range. In addition, CRP significantly decreased in all groups after drug withdrawal $(HN_{50mg/4w}, P = .03; HIV_{50mg/12w}, P = .03;$ and HIV_{150mg/4w}, *P* = .004; [Supplementary Figure 3](http://academic.oup.com/jid/article-lookup/doi/10.1093/infdis/jiad251#supplementary-data)*B*). Nevertheless, significant differences remained present between PWH and HIV-negative individuals either at baseline or after obefazimod administration, such as the expression of G-CSF, IL-7, tumor necrosis factor–α, IL-18 bp, MIP-1β (CCL4), MCP-1 (CCL2), and CRP ([Supplementary Figure 4\)](http://academic.oup.com/jid/article-lookup/doi/10.1093/infdis/jiad251#supplementary-data). This fact supports the evidence that PWH exhibit chronic inflammation,

Figure 4. Effect of obefazimod on miR-124 expression. A, Baseline mir-124 expression of the study population measured in rectal tissue by reverse-transcription quantitative polymerase chain reaction. *B*, Fold changes in miR-124 expression throughout intervention and subsequent discontinuation of the study groups. Horizontal lines in panel A represent medians. Asterisks denote significant differences between timepoints detected using the Wilcoxon signed-rank test (**P* ≤ .05; ***P* ≤ .01; ****P* ≤ .001). Abbreviations: ΔCt, change in threshold cycle; BL, baseline; EoI, end of intervention; EoI + 4w, 4 weeks after end of intervention; HIV_{50mg/12w}, HIV infected taking once-daily oral obefazimod 50 mg for 12 weeks; HIV_{150mg/4w}, HIV infected taking once-daily oral obefazimod 150 mg for 4 weeks; HN_{50mg/4w}, HIV negative taking once-daily oral obefazimod 50 mg for 4 weeks.

which seems to be higher in the $HIV_{150mg/4w}$ group, perhaps due to the longer time living with HIV-1 and undergoing ART.

Of note, the tendency to reduce inflammation was even greater 4 weeks after drug discontinuation in PWH, especially when obefazimod was administered at 50 mg for 12 weeks, showing a significant downregulation of GM-CSF, IL-7, IL-1β, IL-12 (p70), IL-2, and IL-17α (CTLA-8) ($P = .003$, $P = .03$, $P = .02$, $P = .02$, $P = .02$, and $P = .05$, respectively) [\(Figure 5](#page-8-0)*B*). Overall, these results suggest that obefazimod may reduce chronic inflammation in ART-suppressed PWH, and that longer duration of the intervention may enhance lasting anti-inflammatory effects.

DISCUSSION

In this study we evaluated the safety and tolerability in addition to the impact of obefazimod on viral persistence, and chronic immune activation and inflammation in ART-suppressed PWH. We confirm previous findings showing that obefazimod is safe and well tolerated when used in combination with ART in PWH [\[13](#page-10-0)]. Furthermore, we observed a reduction in total HIV-1 DNA, residual viremia, and cell-activation and inflammation markers in blood and GALT. Overall, this study suggests that obefazimod may have a triple in vivo effect by reducing HIV-1 persistence, chronic immune activation, and chronic inflammation in PWH.

Our study showed a reduction in total HIV-1 DNA in peripheral CD4+ T cells and a trend toward decreased residual plasma viremia when obefazimod was administered at 150 mg for 4 weeks in ART-suppressed PWH. This effect might indirectly happened by cellular toxicity or changes in cellular distribution, since central and transitional memory CD4⁺ T cells are the major cellular reservoirs for HIV [[5](#page-10-0)]. However,

no toxicity related to obefazimod affecting CD4⁺ T-cell counts has been previously described [\[12](#page-10-0)] nor observed in our study [\(Supplementary Figure 1\)](http://academic.oup.com/jid/article-lookup/doi/10.1093/infdis/jiad251#supplementary-data), and no redistribution of memory CD4+ T-cell subsets was observed either [\(Supplementary](http://academic.oup.com/jid/article-lookup/doi/10.1093/infdis/jiad251#supplementary-data) [Table 1\)](http://academic.oup.com/jid/article-lookup/doi/10.1093/infdis/jiad251#supplementary-data). Therefore, our results suggest that obefazimod directly impacts the proviral reservoir size in a dose-dependent manner. Previous published studies with obefazimod have already confirmed a dose-dependent efficacy of the drug [\[12\]](#page-10-0). Indeed, similar trend in total HIV-1 DNA had previously been reported in PWH receiving 150 mg dose of obefazimod for 4 weeks [[13\]](#page-10-0) but, comparatively, our study provides statistically significant changes in total HIV-1 DNA and a greater proportion of responders. Furthermore, in a recently reported substudy of the 150-mg dose of obefazimod [[21\]](#page-11-0), intact proviral DNA was measured and showed a similar decreasing trend without reaching statistical significance. Altogether these data suggest that obefazimod decreases the total proviral reservoir and may also decrease the intact proviral reservoir when administered in combination with ART [\[21](#page-11-0)]. This finding has important implications given the failure of multiple interventions to achieve a measurable reduction of the viral reservoir size in other studies. Nevertheless, in both studies the effect on proviral DNA reversed after drug discontinuation, so it is tempting to speculate that obefazimod's ability to interfere with cell activation and inflammation might contribute to reducing T-cell proliferation as well.

In our study, we did not observe changes in the CA HIV-1 RNA during obefazimod treatment. However, as our assay is designed to measure all mRNAs resulting from HIV-1 transcription initiation and not specifically further mature RNA forms such as elongated or spliced molecules, this might have impeded the detection of the impact of obefazimod in late steps of HIV-1 RNA biogenesis. To further investigate the viral

Figure 5. Effect of obefazimod on inflammation. *A*, Heat map representing median fold changes in plasma levels of inflammation biomarkers throughout obefazimod intervention (baseline to end of intervention). *B*, Heat map representing median fold changes in plasma levels of inflammation biomarkers from baseline to 4 weeks after obefazimod discontinuation. Decrease and increase in median fold changes are shown in blue and red, respectively. C-reactive protein is outside the defined range. Median increase/decrease comparisons between timepoints were performed using the Wilcoxon signed-rank test. *P* values are shown. Abbreviations: CRP, C-reactive protein; G-CSF, granulocyte colony-stimulating factor; GM-CSF, granulocyte-macrophage colony-stimulating factor; HIV_{50/12}, HIV infected taking once-daily oral obefazimod 50 mg for 12 weeks; HIV_{150/4}, HIV infected taking once-daily oral obefazimod 150 mg for 4 weeks; HN_{50/4}, HIV negative taking once-daily oral obefazimod 50 mg for 4 weeks; IFN-γ, interferon gamma; IL, interleukin; IP-10, interferon-γ inducible protein-10; MCP-1, monocyte chemoattractant protein-1; M-CSF, macrophage colony-stimulating factor; MIG, monokine induced by gamma interferon; MIP-1β, macrophage inflammatory protein-1beta; TNF-α, tumor necrosis factor alpha.

transcription profile throughout obefazimod administration, we made an in-depth substudy of different viral transcripts in which we only analyzed samples from the HIV_{150mg/4w} group given the dose-dependent efficacy of the drug [\[21](#page-11-0)]. This substudy showed that obefazimod decreases viral transcription initiation and may decrease viral transcription elongation [[21\]](#page-11-0). Nevertheless, both studies showed an increase in viral transcription after drug discontinuation, like the increase in total HIV-1 DNA after drug withdrawal, newly suggesting that obefazimod withdrawal might also increase viral transcription through associated changes in immune cell activation and inflammation. These findings suggest that obefazimod may impact specific forms of HIV transcription but lacks lasting effect. Overall, our study showed the potential effect of obefazimod at reducing the persistence of the HIV-1 reservoir in PWH despite suppressive ART.

So far, no study has investigated the effect of obefazimod on immune activation in blood nor rectal tissue in ART-suppressed PWH. In agreement with previous published data, we reported

higher frequency of activated T cells in rectal tissue compared to blood $[22, 23]$ $[22, 23]$ $[22, 23]$. Indeed, the highest proportion of $CD4+PD-1+T$ cells was found within the rectal tissue, mainly in the $HIV_{150me/4w}$ group who had been living with HIV and taking long-term ART. Importantly, our study shows for the first time the lasting effect of obefazimod at reducing immune activation in ART-suppressed PWH by reducing the expression of cell surface activation markers (ie, CD38, HLA-DR, and PD-1), that occurs in peripheral and rectal $CD4^+$ and $CD8^+$ T cells during HIV-1 infection, which is associated with a progressive loss of effector function [\[24](#page-11-0), [25](#page-11-0)]. Therefore, it may have applicability for the treatment of chronic immune activation exhibited by PWH despite long-term ART suppression.

Consistent with previous studies [[14](#page-10-0), [16](#page-11-0), [26](#page-11-0)], we confirmed the strong anti-inflammatory effect of obefazimod in ART-suppressed PWH. Obefazimod upregulated the antiinflammatory miR-124 in all HIV-1–infected participants undergoing ART. miR-124 is a critical modulator of inflammation and immunity that is reported to provide therapeutic restitution of physiological pathways lost in multiple inflammatory diseases [[15\]](#page-11-0). By the selective induction of miR-124, obefazimod demonstrates anti-inflammatory effects in both preclinical models and patients with ulcerative colitis (UC) [[17,](#page-11-0) [27,](#page-11-0) [28](#page-11-0)], and is currently in phase 3 trials in UC, and phase 2 trials in Crohn disease and rheumatoid arthritis [[29\]](#page-11-0). The role of miR-124 in HIV-1 infection has only been preliminarily explored. Thus, it has been suggested that HIV-1 Tat could induce microglial activation via the promoter DNA hypermethylation of miR-124, leading to the exacerbated release of proinflammatory cytokines [[30\]](#page-11-0), a molecular mechanism that could be involved in neuroinflammation associated with HIV-1-associated neurocognitive disorders [[31\]](#page-11-0).

Consistent with miR-124 upregulation, obefazimod represses overexpression of some HIV-inductive soluble inflammation biomarkers, such as G-CSF, GM-CSF, CD14, IL-7, IL-10, MIP-1β (CCL4), and MCP-1 (CCL2). These effects on the immune system are reminiscent of some, but not all, of those observed in serum samples from UC patients treated with obefazimod (Tazi, personal communication), which emphasizes that inflammatory bowel diseases and HIV infection share some similarities, but that the inflammation seen in HIV-infected patients is very specific and that there are significant differences in terms of immunopathological changes between these 2 conditions [[32\]](#page-11-0). CRP was transiently upregulated only after obefazimod administration at the 150-mg dose $(P = .008)$, but the magnitude of change was not clinically meaningful. Actually, no transient elevation of CRP has been observed to date in >800 UC patients given 50 mg of obefazimod [\[29](#page-11-0)], and when a Spearman correlation of CRP with all variables included in this study was analyzed, no significant correlation was found that could explain this transient effect (data not shown).

In contrast with miR-124 upregulation, which was reversed after drug discontinuation, the greatest attenuation of inflammation biomarkers was observed 4 weeks after drug discontinuation, such as downregulation of GM-CSF, IL-7, IL-1β, IL-12 (p70), IL-2, and IL-17 α (CTLA-8), when obefazimod was administered for a longer period. Altogether, our results show the broad anti-inflammatory effect of obefazimod and suggest that it might be a good candidate to treat the chronic inflammation that ART-suppressed PWH exhibit as well as other inflammatory diseases.

Among the study limitations, we should acknowledge the small number of participants that may preclude statistical power of several results; the restricted demographics (all White men), which could hide ethnic- and gender-based differences in response to the intervention; and differences in total time of HIV infection and undergoing ART between the $HIV_{50\text{mg}}/$ $_{12w}$ and HIV_{150mg/4w} groups that might mask differences between the 50- and 150-mg doses of treatment with obefazimod.

It is tempting to speculate that based on the present results, intermediate doses of obefazimod and longer treatment periods than those used in the present study could result in a sustained impact on viral persistence and chronic immune activation and inflammation, while limiting drug-related adverse effects. Of note, obefazimod was not administered to patients with UC at a daily dose >50 mg. Moreover, the present results also support the search of alternative drugs that target viral transcription, processing, and/or maturation of viral transcripts with potential to eliminate or silence viral persistence and simultaneously suppress virus-induced immune activation and inflammation [\[33\]](#page-11-0).

In conclusion, obefazimod impacts chronic immune activation and inflammation in ART-suppressed PWH, further contributing to changes in viral persistence. It would be interesting to better define the drug-associated mechanisms that counteract persistent activation/inflammation, either by decreasing HIV antigen expression and presumably reducing T-cell proliferation, or by selectively upregulating miR-124 through its specific interaction with the CBC. The overall effect of obefazimod suggests a potential role for the drug in context of sustained inflammation during ART, as well as in virus remission strategies involving other compounds that can activate immune cells, such as latency-reversing agents.

Supplementary Data

[Supplementary materials](http://academic.oup.com/jid/article-lookup/doi/10.1093/infdis/jiad251#supplementary-data) are available at *The Journal of Infectious Diseases* online. Consisting of data provided by the authors to benefit the reader, the posted materials are not copyedited and are the sole responsibility of the authors, so questions or comments should be addressed to the corresponding author.

Notes

Author contributions. All authors made substantial contributions to the conception or design of the work, or the data acquisition, analysis, or interpretation of the work. R. D. C., I. M., J. D., D. S., R. P., and J. M.-P. contributed to the study conception and design. J. R. S., B. R., G. S., and B. C. contributed to the inclusion and clinical follow-up of the patients. S. B. contributed to data acquisition, analysis, and interpretation, and drafted the original manuscript including figures and tables. S. B. and M. C. P. contributed to nucleic acid analysis. S. B. and I. E. contributed to flow cytometry analysis; S. B. and S. M.-L. contributed to GALT sample processing. S. B. and M. S. contributed to residual viremia analysis. S. B. and C. G. contributed to measurement of soluble inflammation biomarkers. V. U. contributed to statistical analysis. J. M.-P., S. M.-L., and M. C. P. reviewed the manuscript critically. All authors have approved the final version submitted for publication.

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