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Switching to Bictegravir/Emtricitabine/Tenofovir Alafenamide in Black Americans With HIV-1: A Randomized Phase 3b, Multicenter, Open-Label Study

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Background: With the highest rates of HIV/AIDS in the United States, Black Americans are still underrepresented in HIV medical research.

Setting: BRAAVE (NCT03631732) is a randomized, phase 3b, multicenter, open-label US study.

Methods: Adults identifying as Black or African American and virologically suppressed on 2 nucleoside reverse transcriptase inhibitors (NRTIs) plus third agent were randomized (2:1) to switch to open-label bictegravir/emtricitabine/tenofovir alafenamide (B/F/TAF) once daily or stay on baseline regimen (SBR) for 24 weeks, after which SBR had delayed switch to B/F/TAF. Resistance to non-NRTIs, protease inhibitors, and/or NRTIs was permitted; integrase strand transfer inhibitor resistance was exclusionary. Primary endpoint was proportion of participants with HIV-1 RNA ≥50 copies/mL at week 24 (snapshot algorithm; noninferiority margin of 6%).

Results: Of 558 screened, 495 were randomized/treated (B/F/TAF n = 330; SBR n = 165). Overall, 32% were ciswomen, 2% transwomen, and 10% had an M184V/I mutation. At week 24, 0.6% on B/F/TAF vs 1.8% on SBR had HIV-1 RNA ≥50 copies/mL (difference −1.2%; 95% confidence interval −4.8% to 0.9%), demonstrating noninferiority of B/F/TAF vs SBR. Proportions with HIV-1 RNA <50 copies/mL at week 24 were 96% B/F/TAF and 95% SBR and remained high at week 48. No participant had treatment-emergent resistance to study drug. Treatments were well tolerated. Study drug-related adverse events, mostly grade 1, occurred in 10% of participants on B/F/TAF through week 48 and led to discontinuation in 9 participants through week 48.

Conclusions: For Black Americans with HIV, switching to B/F/TAF was noninferior to continuing a variety of regimens, including those with pre-existing NRTI mutations.

Key Words: bictegravir, HIV, Black Americans, tenofovir alafenamide, INSTI

INTRODUCTION

Among racial and ethnic groups in the United States, Black people have the highest incidence and prevalence of HIV/AIDS.1,2 Black Americans make up about 13% of the US population but account for 42% of HIV infections. In 2018, the rate of diagnosis of new HIV infections among Black Americans was 39.2 per 100,000, as compared with 4.8 among whites and 13.3 in the population as a whole. This
disparity was even greater in women: The rate of new HIV diagnoses was 21.3 per 100,000 for Black women, nearly 13 times higher than the rate of 1.7 in white women. Moreover, the risk of death for Black Americans from HIV/AIDS is approximately 9 times that of their white counterparts.\textsuperscript{1,2} These and other racial disparities show the impacts of structural racism, which contribute to myriad inequities including access to systems of health care, education, and employment. Within the health care system, interpersonal discrimination based on race, gender, and sex also results in disparities in health outcomes.\textsuperscript{3–5} The social factors and structural inequalities that confound racial health disparities in the United States are challenging to disentangle\textsuperscript{6} and likewise may complicate clinical decision-making for patients with HIV despite evidence that neither race nor ethnic origin plays a significant role in HIV treatment outcomes when health care is more equitably distributed.\textsuperscript{7,8} And yet, despite overrepresentation in the US HIV epidemic and clinical need to address health disparities, in clinical trials of HIV treatment regimens, Black people living with HIV (PLWH) are consistently underrepresented.\textsuperscript{9–11} As a result, the findings of clinical trials may not necessarily be generalizable to Black Americans.

The single-tablet regimen of the integrase strand transfer inhibitor (INSTI) bicaptegravir plus the nucleoside reverse transcriptase inhibitors (NRTIs) emtricitabine and tenofovir alafenamide (B/F/TAF) is a recommended initial treatment for most people with HIV.\textsuperscript{12–14} Four randomized international trials have established that switching to B/F/TAF from other combination antiretroviral (ARV) regimens is safe and effective in a general population of virologically suppressed adults with and without resistance to NRTIs.\textsuperscript{15–18} Although the B/F/TAF clinical trials of treatment-naive PLWH enrolled relatively representative proportion of Black participants in the United States, the inclusion of Black PLWH in treatment-experienced studies was low. There is a clear need for clinical data focusing on the safety and efficacy of B/F/TAF switch in virologically suppressed Black American PLWH.

The primary objective of this phase 3b study was to evaluate the efficacy of switching from a regimen of 2 NRTIs and a third agent to a fixed-dose combination of B/F/TAF vs continuing their baseline regimen in virologically suppressed adults self-identifying as African American or Black.

**METHODS**

**Study Design and Participants**

This randomized, open-label, multicenter, active-controlled, phase 3b trial was conducted at 82 outpatient centers in the United States. We enrolled adults (aged 18 years and older) living with HIV who self-described as Black, African American, or mixed race that included Black. Eligible participants had documented plasma HIV-1 RNA levels <$50$ copies/mL for at least 12 months before screening while on a stable ARV regimen other than B/F/TAF that consisted of any 2 NRTIs along with an allowed third agent for at least 6 months. Participants had an estimated glomerular filtration rate (eGFR) $\geq 50$ mL per minute (calculated by the Cockcroft-Gault equation) and no documented history of resistance to INSTIs or tenofovir (TFV) resistance defined as K65R/E/N mutations, $\geq 3$ thymidine analog mutations, or T69 insertions (See Appendix, Supplemental Digital Content, http://links.lww.com/QAI/B674).

Participants were randomized (2:1 ratio) to either switch to B/F/TAF (50 mg/200 mg/25 mg) administered orally once daily without regard to food or stay on baseline regimen (SBR) (which was not provided by the study) from day 1 until week 24. After week 24, the SBR group switched to B/F/TAF for an additional 24 weeks. Participants who completed the study through week 48 and wished to continue B/F/TAF were given the option to receive B/F/TAF for up to an additional 24 weeks in an extension phase and attend study visits every 12 weeks. Randomization was stratified by the baseline third-agent ARV class at entry.

This trial was undertaken in accordance with the Declaration of Helsinki and approved by central or site-specific review boards or ethics committees. All participants provided written informed consent.

A computer-generated allocation sequence (block size of 6) was created by Bracket (San Francisco, CA). Study investigators determined eligibility, obtained a participant number, and received automated treatment assignment based on a randomization sequence.

**Procedures**

Participans had visits on day 1, weeks 4, 12, 24, 36, and 48, with an additional week 28 visit for the SBR group. Laboratory tests included hematological analyses, serum chemistries, cluster of differentiation-4 (CD4) T-cell counts, renal parameters (eGFR) (Covance Laboratories, Indianapolis, IN), and measurement of HIV-1 RNA plasma concentration (Roche TaqMan 2.0; Roche Diagnostics, Indianapolis, IN). Resistance testing by genotyping and phenotyping integrase and reverse transcriptase (Monogram Biosciences, Inc., South San Francisco, CA) was included for any participant with plasma HIV-1 RNA $\geq 200$ copies/mL at confirmed virologic failure and who did not resuppress on study drug.

Safety was assessed by physical examinations, laboratory tests, recording of concomitant drugs, and recording of adverse events, which were coded using the Medical Dictionary for Regulatory Activities (MedDRA, version 22.1). Relatedness of adverse events to study treatment was determined by the investigator.

**Outcomes**

The primary efficacy endpoint was the proportion of participants with HIV-1 RNA $\geq 50$ copies/mL at week 24 as defined by the US FDA-defined snapshot algorithm. The secondary efficacy endpoints were the proportion of participants with HIV-1 RNA $\geq 50$ copies/mL at week 48, the proportion of participants with HIV-1 RNA $<50$ copies/mL at weeks 24 and 48, and the change in CD4 cell count from baseline to weeks 24 and 48.
Statistical Analysis

The study was powered using expected efficacy based on a pooled analysis of Black participants in studies of INSTI-based regimens.\textsuperscript{19} Assuming equal efficacy between the 2 treatment groups, a sample size of 480 participants randomized in a 2:1 ratio (320 in the B/F/TAF group and 160 in the SBR group) would achieve approximately 89% power to detect a prespecified noninferiority margin difference of 6% in the group proportions with HIV-1 RNA $\geq$50 copies/mL at week 24 (B/F/TAF minus SBR) by FDA snapshot algorithm, using a 1-sided alpha level of 0.025%. All randomized participants who received at least one dose of study treatment (either B/F/TAF or baseline regimen) on or after day 1 and did not have pre-existing INSTI resistance-associated mutations based on historical data were included in the primary efficacy analysis.

The proportion with HIV-1 RNA $\geq$50 copies/mL was also assessed at week 48 as a secondary outcome. The proportions of participants with plasma HIV-1 RNA <50 copies/mL at weeks 24 and 48 by FDA snapshot algorithm were prespecified secondary efficacy outcomes, analyzed similarly to the primary efficacy endpoint but using a noninferiority margin of 10%.

Pre-existing drug resistance was assessed by cumulative historical and/or proviral DNA genotypes performed retrospectively from baseline samples. We performed subgroup analyses of the participants with plasma HIV-1 RNA <50 copies/mL at week 24 based on age, sex, baseline NRTI resistance, and specifically baseline M184V/I resistance. A per-protocol analysis was performed as a secondary outcome to assess proportions of participants with plasma HIV-1 RNA <50 copies/mL at week 24 by FDA snapshot algorithm. Weeks 24 and 48 secondary efficacy endpoints were also analyzed with a plasma HIV-1 RNA threshold of <20 copies/mL by FDA snapshot algorithm. We imputed missing as failure (M = F) and missing as excluded (M = E) to estimate the proportion of participants with plasma HIV-1 RNA <50 copies/mL at weeks 24 and 48.

Differences in percentages of participants with HIV-1 RNA <20 copies/mL, < 50 copies/mL or $\geq$50 copies/mL between treatment groups and 95% CI were calculated based on exact methods.

An analysis of variance model with treatment group as a fixed effect was used to analyze CD4 count and percentage. Safety was assessed in all randomized participants who received at least one dose of study treatment. Differences between groups in fasting lipids and body weight were tested using the two-sided Wilcoxon rank-sum test. Safety data were collected from the date the study drug was first given for up to 30 days after the last dose of study drug. Treatment satisfaction questionnaire scores were compared between groups using analysis of covariance adjusting for baseline score.

We used SAS Software Version 9.4 (SAS Institute Inc., Cary, NC) for all analyses. This study was conducted according to protocol and is registered with ClinicalTrials.gov, number NCT03631732.

RESULTS

Between August 28, 2018, and March 6, 2019, 558 individuals were screened for enrollment and 496 were randomized: 331 to B/F/TAF and 165 to SBR. One participant randomized to B/F/TAF never received treatment and was excluded from analysis. Of 330 participants randomized and treated with B/F/TAF, 320 (97%) continued treatment after the week 24 primary endpoint, and 163 of 165 (99%) participants randomized to SBR switched to B/F/TAF at week 24 (Fig. 1). Demographics and baseline characteristics were generally balanced between groups (Table 1). The median age overall was 49 years, 32% were ciswomen, 2% transwomen, and 10% had documented baseline M184V/I mutation. Median time on HIV treatment was 10 years, and median time on baseline ARV regimen was 2 years. Most participants (61%) were receiving an INSTI as a third agent at baseline.

Efficacy

Switching to B/F/TAF was noninferior to SBR for the primary outcome of proportion of participants with plasma HIV-1 RNA $\geq$50 copies/mL at week 24 as defined by the US FDA snapshot algorithm (0.6% (2 of 328 participants) vs 1.8% (3 of 165), difference $-1.2\%$ (95% confidence interval [CI]: $-4.8\%$ to 0.9%)) (Table 2, Fig. 2, panel A). Two participants were found to have baseline INSTI resistance based on historical genotype data and were excluded from the primary analysis because of this protocol violation; 1 participant each with Y143C and Q148K mutations, both were randomized to B/F/TAF and had HIV-1 RNA $<$50 copies/mL at week 24. The secondary efficacy outcome of proportion of participants with plasma HIV-1 RNA $<$50 copies/mL at week 24 was consistent with the primary endpoint and showed high rates of virologic suppression [B/F/TAF 96.3% (316 of 328 participants) vs SBR 94.5% (156 of 165)] (Table 2).

After 48 weeks of receiving B/F/TAF, 95% (310 of 328) in the group randomized to B/F/TAF had HIV-1 RNA $<$50 copies/mL (Fig. 2, panel A; Appendix Table 4, Supplemental Digital Content, http://links.lww.com/QAI/B674). Two had HIV-1 RNA $\geq$50 copies/mL. One individual had an HIV-1 RNA level of 55 copies/mL at week 48 and switched back to the prestudy regimen. The other had an HIV-1 RNA level of 455 copies/mL at week 48, which was suppressed on continued treatment with B/F/TAF; no treatment-emergent resistance was detected. Of 163 participants randomized to SBR who switched to B/F/TAF at week 24, no participants had HIV-1 RNA $\geq$50 copies/mL; 158 (97%) had HIV-1 RNA $<$50 copies/mL at week 48.

Results from the per-protocol, M = F, and M = E analyses conducted at weeks 24 and 48 were consistent with the primary outcome (Table 2). Analysis of the proportion of participants with plasma HIV-1 RNA $<$20 copies/mL showed high and similar rates of virologic suppression at this lower threshold. Efficacy between B/F/TAF and SBR was similar across prespecified subgroups (Fig. 2, panel B), and testing for homogeneity found no significant interactions between treatment and subgroup. We observed no treatment differences in changes from baseline in CD4 cell count and CD4 percentage at week 24 (see Appendix Table 1, Supplemental Digital Content, http://links.lww.com/QAI/B674).
TABLE 1. Baseline Demographics and Disease Characteristics

<table>
<thead>
<tr>
<th></th>
<th>B/F/TAF (n = 330)</th>
<th>SBR (Delayed Switch to B/F/TAF) (n = 165)</th>
</tr>
</thead>
<tbody>
<tr>
<td>Median age, yrs (range)</td>
<td>49 (18–79)</td>
<td>49 (19–70)</td>
</tr>
<tr>
<td>Female sex at birth, n (%)</td>
<td>101 (31)</td>
<td>55 (33)</td>
</tr>
<tr>
<td>Gender identity, n (%)</td>
<td></td>
<td></td>
</tr>
<tr>
<td>Cisgender</td>
<td>317 (96)</td>
<td>159 (96)</td>
</tr>
<tr>
<td>Transgender</td>
<td>6 (2)</td>
<td>6 (4)</td>
</tr>
<tr>
<td>Other gender or prefer not to disclose</td>
<td>7 (2)</td>
<td>0</td>
</tr>
<tr>
<td>Sexual orientation, n (%)</td>
<td></td>
<td></td>
</tr>
<tr>
<td>Heterosexual (female sex at birth)</td>
<td>97 (29)</td>
<td>52 (32)</td>
</tr>
<tr>
<td>Heterosexual (male sex at birth)</td>
<td>62 (19)</td>
<td>41 (25)</td>
</tr>
<tr>
<td>Gay or bisexual (male sex at birth)</td>
<td>162 (49)</td>
<td>67 (41)</td>
</tr>
<tr>
<td>Ethnicity, n (%)</td>
<td></td>
<td></td>
</tr>
<tr>
<td>Hispanic/latinx</td>
<td>15 (5)</td>
<td>3 (3)</td>
</tr>
<tr>
<td>Median CD4 count, cells/μL (Q1, Q3)</td>
<td>747 (570, 922)</td>
<td>758 (494, 969)</td>
</tr>
<tr>
<td>Median eGFR&lt;sub&gt;CG&lt;/sub&gt;, mL/min (Q1, Q3)</td>
<td>110 (88, 132)</td>
<td>107 (86, 132)</td>
</tr>
<tr>
<td>Median weight, kg (Q1, Q3)</td>
<td>88 (79, 103)</td>
<td>89 (76, 104)</td>
</tr>
<tr>
<td>Median BMI, kg/m² (Q1, Q3)</td>
<td>29.2 (25.9, 34.0)</td>
<td>29.3 (25.7, 34.3)</td>
</tr>
<tr>
<td>HBV coinfection, n (%)</td>
<td>15 (5)</td>
<td>3 (2)</td>
</tr>
<tr>
<td>Baseline NRTI backbone,* n (%)</td>
<td></td>
<td></td>
</tr>
<tr>
<td>F/TAF</td>
<td>224 (68)</td>
<td>107 (65)</td>
</tr>
<tr>
<td>F/TDF</td>
<td>56 (17)</td>
<td>34 (21)</td>
</tr>
<tr>
<td>Abacavir/lamivudine</td>
<td>44 (13)</td>
<td>24 (15)</td>
</tr>
<tr>
<td>Other</td>
<td>4 (1)</td>
<td>0</td>
</tr>
<tr>
<td>Baseline 3rd agent,† n (%)</td>
<td></td>
<td></td>
</tr>
<tr>
<td>INSTI</td>
<td>202 (61)</td>
<td>99 (60)</td>
</tr>
<tr>
<td>NNRTI</td>
<td>100 (30)</td>
<td>51 (31)</td>
</tr>
<tr>
<td>PI</td>
<td>30 (9)</td>
<td>14 (9)</td>
</tr>
<tr>
<td>CCR5 antagonist</td>
<td>0</td>
<td>1 (1)</td>
</tr>
<tr>
<td>Baseline ARV resistance,‡ n (%)</td>
<td></td>
<td></td>
</tr>
<tr>
<td>NRTI resistance</td>
<td>44 (13)</td>
<td>26 (16)</td>
</tr>
<tr>
<td>M184V/I</td>
<td>31 (9)</td>
<td>20 (12)</td>
</tr>
<tr>
<td>NNRTI resistance</td>
<td>70 (21)</td>
<td>32 (19)</td>
</tr>
<tr>
<td>PI resistance</td>
<td>36 (11)</td>
<td>26 (16)</td>
</tr>
<tr>
<td>INSTI resistance</td>
<td>8 (2)</td>
<td>3 (2)</td>
</tr>
</tbody>
</table>

*Two participants in the B/F/TAF group did not report a baseline NRTI backbone medication; both were protocol violations.
†One participant in the B/F/TAF group reported both PI and INSTI; 1 subject in the B/F/TAF group reported both INSTI and NNRTI, both were protocol violations.
‡No baseline ARV resistance data B/F/TAF n = 17, SBR n = 7. Resistance assessed by cumulative historical or retrospective baseline proviral DNA genotypes.
BMI, body mass index; CCR5, C-C chemokine receptor type 5; eGFR<sub>CG</sub>, estimated glomerular filtration rate (calculated by Cockcroft-Gault equation); HBV, hepatitis B virus; NNRTI, non-nucleoside reverse transcriptase inhibitor; PI, protease inhibitor; Q, quartile.

At baseline, 13% (44 of 328) in the B/F/TAF group and 16% (26 of 165) in the SBR group had NRTI resistance, including 9% (31 of 328) and 12% (20 of 165) with M184V/I mutations in the B/F/TAF and SBR groups, respectively. Efficacy by snapshot analysis was similar in participants with or without NRTI resistance (Fig. 2, panel C). Among participants with NRTI resistance at baseline 98% (43 of 44) on B/F/TAF and 96% (25 of 26) on SBR had HIV-1 RNA <50 copies/mL at week 24. Among participants with M184V or I mutations at baseline 97% (30 of 31) on B/F/TAF and 95% (19 of 20) on SBR had HIV-1 RNA <50 copies/mL at week 24. One participant with M184V on B/F/TAF discontinued after the baseline visit and did not have virologic data at week 24. In addition, 2% (8 of 328) in the B/F/TAF group and 2% (3 of 165) in the SBR group had primary INSTI resistance detected at baseline, all of whom had HIV-1 RNA <50 copies/mL at week 24.

Altogether, 68 participants with baseline NRTI resistance (including 50 with M184V/I) received B/F/TAF during the study and had postswitch virologic data. At study week 48, all 68 (100%) had HIV-1 RNA <50 copies/mL; 43 randomized to B/F/TAF had 48 weeks of treatment, and 25 with a delayed switch from SBR had 24 weeks of treatment with B/F/TAF. In addition, a total of 11 participants with baseline INSTI resistance received B/F/TAF, and all 11 (100%) had HIV-1 RNA <50 copies/mL at week 48.

One participant in each treatment group met criteria for resistance testing through week 24. No treatment-emergent resistance to any components of the regimens was detected in either group during the study.

Safety

Through week 24, 64% of participants who switched to B/F/TAF (210 of 330) experienced at least one adverse event compared with 52% (85 of 165) of those on SBR (Table 3). The most common adverse events in the B/F/TAF and SBR groups were upper respiratory infection, diarrhea, cough, arthralgia, and bronchitis. The proportion of participants who experienced at least one grade 3 or 4 adverse event was the same in both groups (5%) as was the proportion of participants experiencing a serious adverse event (4%). Investigators attributed events to treatment in 11% (36 participants) in the B/F/TAF group and none in the SBR group. For most participants who had adverse events attributed to treatment, events were either mild (25 of 36) or moderate (8 of 36) and resolved on treatment. Three of these 36 participants experienced grade 3 events: one (somnolence) that resolved on study drug and 2 that led to study drug discontinuation (headache n = 1, agitation, psychomotor hyperactivity, and anxiety n = 1). In total, 7 (2%) of participants in the B/F/TAF group and none in the SBR group had adverse events leading to discontinuation up to week 24. No deaths or pregnancies were reported.

Incidence of grade 3 or 4 laboratory abnormalities through week 24 was comparable between groups: 8% (27 of 329) B/F/TAF and 6% (10 of 165) SBR (see Appendix Table 2, Supplemental Digital Content, http://links.lww.com/QAI/B674). Median [interquartile range (IQR)] change from baseline in eGFR at week 24 was −4.6 (−12.1, 2.6) mL/min for B/F/TAF (n = 318) vs 1.1 (−7.1, 7.8) mL/min for SBR (n = 160); P < 0.001; consistent with the previously described inhibition of tubular secretion of creatinine without affecting renal glomerular function by bictegravir. In the B/F/TAF group, one participant with prostate cancer discontinued study drug because of acute kidney injury secondary to urinary obstruction with

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FIGURE 1. CONSORT diagram. B/F/TAF, bictegravir, emtricitabine, and tenofovir alafenamide; SBR, stay on baseline regimen.

* Because of gunshot.
Hepatitis B virus DNA
HIV-1 RNA
HIV-1 RNA
Discontinued because of other reasons
Discontinued because of AE/death and last available HIV-1 RNA ≥50 copies/mL.
Discontinued because of other reasons† and last available HIV-1 RNA ≥50 copies/mL.
No virologic data in the week 24 window
Discontinued because of AE/death and last available HIV-1 RNA <50 copies/mL.
Discontinued because of other reasons‡ and last available HIV-1 RNA <50 copies/mL.
Missing data but on study drug
HIV-1 RNA <50 copies/mL by per-protocol snapshot analysis
HIV-1 RNA <50 copies/mL by missing = failure
HIV-1 RNA <50 copies/mL by missing = excluded
HIV-1 RNA <20 copies/mL
Hepatitis B virus DNA <29 IU/mL by missing = excluded

 Data are n (%).
 Virology outcomes are based on snapshot algorithm unless otherwise specified.
 The week 24 window is between days 127 and 210 (inclusive).
 Per-protocol analysis excluded patients in full analysis set who were off study drug at week 24 or had low adherence, that is, adherence ≤2.5th percentile among those in study or did not meet the inclusion criteria for baseline ARV medication or met inclusion/exclusion criteria for mutations detected before baseline or after baseline.
 *The difference in percentages of participants between treatment groups and their 95% CIs were calculated based on exact methods.
 †Other reasons include participants who discontinued study drug because of investigator’s discretion, subject decision, lost to follow-up, noncompliance with study drug, protocol violation, pregnancy, and study terminated by sponsor.
 ‡The per-protocol analysis was performed in a similar manner to the primary endpoint but excluded participants with low adherence (below the 2.5th percentile), violation of key entry criteria, or who did not have a plasma HIV-1 RNA value in the week 24 analysis window because of reasons other than lack of efficacy.
 §Difference in percentages and 95% CI were based on a dichotomized response: HIV-1 RNA <50 copies/mL vs HIV-1 RNA ≥50 copies/mL or missing for missing = failure approach and HIV-1 RNA <50 copies/mL vs HIV-1 RNA ≥50 copies/mL for missing = excluded approach.
 ||At baseline, 18 participants had HBV coinfection: 15 (5%) in the B/F/TAF group and 3 (2%) in the SBR group. At baseline, 13 of 15 in the B/F/TAF group and 2 of 3 in the SBR group had HBV DNA <29 IU per mL. The denominator for percentage is the number of participants in the full analysis set with HIV/HBV coinfected and with nonmissig HBV DNA at week 24.

TABLE 2. Virologic Outcomes at Week 24, Full Analysis Set

<table>
<thead>
<tr>
<th>Week 24</th>
<th>B/F/TAF (n = 328)</th>
<th>SBR (No Switch) (n = 165)</th>
<th>B/F/TAF vs SBR Difference in Percentages (95% CI)*</th>
</tr>
</thead>
<tbody>
<tr>
<td>HIV-1 RNA &lt;50 copies/mL</td>
<td>316 (96.3%)</td>
<td>156 (94.5%)</td>
<td>1.8% (−2.0% to 6.8%)</td>
</tr>
<tr>
<td>HIV-1 RNA ≥50 copies/mL</td>
<td>2 (0.6%)</td>
<td>3 (1.8%)</td>
<td>−1.2% (−4.8% to 0.9%)</td>
</tr>
<tr>
<td>HIV-1 RNA ≥50 copies/mL</td>
<td>2 (0.6%)</td>
<td>3 (1.8%)</td>
<td></td>
</tr>
<tr>
<td>Discontinued because of lack of efficacy</td>
<td>0</td>
<td>0</td>
<td></td>
</tr>
<tr>
<td>Discontinued study drug because of AE/death and last available HIV-1 RNA ≥50 copies/mL</td>
<td>0</td>
<td>0</td>
<td></td>
</tr>
<tr>
<td>Discontinued because of other reasons† and last available HIV-1 RNA ≥50 copies/mL</td>
<td>0</td>
<td>0</td>
<td></td>
</tr>
<tr>
<td>No virologic data in the week 24 window</td>
<td>10 (3.0%)</td>
<td>6 (3.6%)</td>
<td></td>
</tr>
<tr>
<td>Discontinued because of AE/death and last available HIV-1 RNA &lt;50 copies/mL</td>
<td>6 (1.8%)</td>
<td>0</td>
<td></td>
</tr>
<tr>
<td>Discontinued because of other reasons‡ and last available HIV-1 RNA &lt;50 copies/mL</td>
<td>4 (2.1%)</td>
<td>1 (0.6%)</td>
<td></td>
</tr>
<tr>
<td>Missing data but on study drug</td>
<td>0</td>
<td>5 (3.0%)</td>
<td>1.4% (−1.0% to 5.3%)</td>
</tr>
<tr>
<td>HIV-1 RNA &lt;50 copies/mL by per-protocol snapshot analysis‡</td>
<td>304/306 (99.3%)</td>
<td>145/148 (98.0%)</td>
<td></td>
</tr>
<tr>
<td>HIV-1 RNA &lt;50 copies/mL by missing = failure§</td>
<td>321/328 (97.9%)</td>
<td>156/165 (94.5%)</td>
<td>3.3% (−0.2% to 8.2%)</td>
</tr>
<tr>
<td>HIV-1 RNA &lt;50 copies/mL by missing = excluded§</td>
<td>321/323 (99.4%)</td>
<td>156/159 (98.1%)</td>
<td>1.3% (−0.8% to 4.9%)</td>
</tr>
<tr>
<td>HIV-1 RNA &lt;20 copies/mL</td>
<td>305/328 (93.0%)</td>
<td>149/165 (90.3%)</td>
<td>2.7% (−2.4% to 8.7%)</td>
</tr>
<tr>
<td>Hepatitis B virus DNA &lt;29 IU/mL by missing = excluded</td>
<td></td>
<td>6/7 (86%)</td>
<td>1/1 (100%)</td>
</tr>
</tbody>
</table>

SBR (P = 0.006 P = 0.02, and P = 0.03, respectively) (see Appendix Table 3, Supplemental Digital Content, http://links.lww.com/QAI/B674). At week 24, treatment differences in changes from baseline were small but statistically significant for fasting total cholesterol (−5-mg/dL B/F/TAF vs −2-mg/dL SBR; P = 0.006), high-density lipoprotein cholesterol (−2-mg/dL B/F/TAF vs −1-mg/dL SBR; P = 0.05), and triglycerides (−5 vs 3 mg/dL; P = 0.01). No significant treatment differences were noted for changes from baseline in low-density lipoprotein cholesterol or total cholesterol: high-density lipoprotein ratio at week 24. A similar proportion of participants in each group were taking lipid-lowering medication at baseline and initiated lipid-lowering agents through week 24. In sensitivity analyses excluding participants on lipid-lowering medications, changes from baseline in fasting lipids were consistent with the full analyses.

Median (IQR) weight at baseline was 88 (79–103) kg in the B/F/TAF group and 89 (76–104) kg in the SBR group (P = 0.66), corresponding to a median (IQR) body mass index of 29 (26–34) in each group. Median (IQR) weight change

Hydrophobia; this adverse event was judged by the investigator as not related to study drug. There were no discontinuations because of renal adverse effects in the SBR group and no reported cases of proximal renal tubulopathy in either group.

A cumulative analysis of safety was performed for all 493 participants treated with B/F/TAF when the last participant reached the week 48 endpoint. At the time of the analysis, the median (IQR) exposure to B/F/TAF was of 61 (56–68) weeks among those randomized to B/F/TAF and 38 (32–44) weeks for those with delayed switch to B/F/TAF; B/F/TAF was well tolerated during the study. Overall, 367 (74%) had at least one adverse event (see Appendix Table 5, Supplemental Digital Content, http://links.lww.com/QAI/B674). Three individuals who received B/F/TAF discontinued treatment after week 24 because of adverse events. Grade 3 or 4 laboratory abnormalities were reported for 60 participants (12%). One death was reported because of a gunshot wound; no pregnancies were reported.

Participants randomized to the B/F/TAF group had higher baseline levels of fasting total and low-density lipoprotein cholesterol and triglycerides compared with those randomized to SBR (P = 0.006 P = 0.02, and P = 0.03, respectively) (see Appendix Table 3, Supplemental Digital Content, http://links.lww.com/QAI/B674). At week 24, treatment differences in changes from baseline were small but statistically significant for fasting total cholesterol (−5-mg/dL B/F/TAF vs −2-mg/dL SBR; P = 0.006), high-density lipoprotein cholesterol (−2-mg/dL B/F/TAF vs −1-mg/dL SBR; P = 0.05), and triglycerides (−5 vs 3 mg/dL; P = 0.01). No significant treatment differences were noted for changes from baseline in low-density lipoprotein cholesterol or total cholesterol: high-density lipoprotein ratio at week 24. A similar proportion of participants in each group were taking lipid-lowering medication at baseline and initiated lipid-lowering agents through week 24. In sensitivity analyses excluding participants on lipid-lowering medications, changes from baseline in fasting lipids were consistent with the full analyses.

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from baseline at week 24 was +0.9 (−1.3, +3.0) kg in the B/F/TAF group and +0.2 (−2.1, +2.0) kg in the SBR group (P = 0.09). At week 48, median (IQR) change in body weight remained at 0.9 kg (−1.5, 4.1) in the B/F/TAF arm. Post hoc exploratory analyses of the change in weight were performed in subgroups by sex at birth for all participants and by baseline NRTIs in those switched to B/F/TAF (see Appendix Fig. 1, Supplemental Digital Content, http://links.lww.com/QAI/B674). In the male sex at birth subanalysis, median change in weight from baseline was similar between groups.

compared with +0.8 (−1.3, 2.5) kg and +0.7 (−2.0, 3.0) kg for those taking TAF- and abacavir (ABC)-containing regimens respectively (P = 0.0235 TDF vs TAF; P = 0.89 ABC vs TAF).

Among the 18 HIV/HBV-coinfected participants, there were no hepatic adverse events, no grade 3 or 4 adverse events, and no participants interrupted or discontinued their study regimen because of an adverse event.

Based on responses using the HIV Treatment Satisfaction Questionnaire, median (IQR) scores were higher with B/F/TAF than SBR at week 24 [29 (24–30) vs 28 (7–30), P < 0.001], indicating a small but significant difference favoring switch to B/F/TAF.

DISCUSSION

In this randomized, open-label, multicenter trial, we found that virologically suppressed Black adults with HIV in the United States can switch from a regimen of 2 NRTIs plus a third agent to B/F/TAF with no loss of virologic efficacy. Few of those who switched to B/F/TAF had HIV RNA ≥50 copies/mL through week 48, and the rate of virologic suppression was high in those with pre-existing resistance to NRTIs. None of the participants experienced treatment-emergent resistance. Treatment satisfaction was higher for participants after switching to B/F/TAF.

Community advisors who participated in the study design and oversight set goals of inclusion based on sex at birth, gender, and sexual orientation to ensure representa-
and safe. Results from this study complement those from other large clinical trials that demonstrated the efficacy, tolerability, and lack of resistance of B/F/TAF. Our results draw us closer to fulfilling a critical need to evaluate infectious disease treatments in a disproportionately affected population.  

**CONCLUSION**

B/F/TAF is safe and effective for Black Americans with HIV switching from a variety of regimens, including those with TFV sensitivity but pre-existing NRTI resistance.

**NOTES**

Community advisors representing HIV advocacy and care organizations across North America reviewed the protocol, provided input into study design, contributed to training for study personnel, and set goals for inclusion across participant groups according to sex at birth, gender identity, and sexual orientation. A study advisory committee composed of 3 community advisors, and 3 clinical site personnel provided ongoing monitoring and input throughout the study.

**ACKNOWLEDGMENTS**

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**REFERENCES**


