

M184V/I does not impact the efficacy of abacavir/lamivudine/dolutegravir use as switch therapy in virologically suppressed patients

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1	M184V/I does not impact the efficacy of abacavir/lamivudine/dolutegravir use as switch
2	in virologically suppressed patients
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20	Short running title: No impact of M184V/I on the efficacy of
21	abacavir/lamivudine/dolutegravir
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Objectives: M184V/I nucleoside reverse transcriptase inhibitor (NRTI) resistance mutations can be selected by either lamivudine/emtricitabine or abacavir. There are controversies about the use of abacavir/lamivudine/dolutegravir combination in HIV-1 treated patients with a fully suppressed HIV viral load and harboring M184V/I. We assessed the efficacy of abacavir/lamivudine/dolutegravir combination when used in HIV pretreated patients with an undetectable viral load (VL) and previously harboring on their genotypic resistance test a M184V/I as unique NRTI resistance and without any resistance to integrase inhibitors.

36 **Patients and methods:** 154 patients with a fully suppressed HIV-1 plasma viral load (< 50 copies/mL) tenofovir/emtricitabine/boosted 37 treated by protease inhibitor or 38 abacavir/lamivudine/boosted inhibitor and switched protease to an 39 abacavir/lamivudine/dolutegravir regimen with M184V/I as unique NRTI resistance mutation 40 in their therapeutic history were retrospectively analyzed up to 12 months after the switch to 41 abacavir/lamivudine/dolutegravir. Assessment of residual viraemia was performed at months 42 1, 3, 6 and 12. Plasma VL with undetected HIV-1 RNA corresponded to an absence of residual viraemia. 43

Results: During the 12 months of follow-up, 3 patients had a blip of VL (53, 62 and 106
copies/mL) at month 3 followed by a subsequent VL < 50 copies/mL. No patient harbored a
virologic failure during the follow-up. Moreover, there was no evolution of residual viraemia
during the follow up.

48 Conclusions: M184V/I as a unique NRTI resistance mutation, regardless of possible selection
49 by regimen containing lamivudine/emtricitabine or abacavir, does not affect the virological
50 response of well controlled patients who switched to abacavir/lamivudine/dolutegravir for at
51 least 12 months.

52 Introduction

Nucleoside/nucleotide reverse transcriptase inhibitors (NRTIs) remain the 'backbone' of 53 antiretroviral therapy. Abacavir was developed alone or in combination with lamivudine, 54 zidovudine and lamivudine + dolutegravir to treat HIV. Lamivudine was developed to be used 55 alone or in combination with zidovudine, abacavir, and abacavir + dolutegravir. 3'-56 thiacytidine inhibitors (lamivudine and emtricitabine) as well as abacavir select rapidly a 57 mutation in vitro in the YMDD region of reverse transcriptase (RT) of HIV-1 (M184I then 58 59 M184V) leading to high level of resistance for lamivudine and emtricitabine and intermediate level of resistance for abacavir. These in vitro M184V/I driven phenotypic resistance 60 mutations to lamivudine, emtricitabine or abacavir were confirmed in clinical trials. ^{1,2,3,4} 61

As M184V/I RT mutation can impact the efficacy of both lamivudine and abacavir, there are controversies about the use of abacavir/lamivudine/dolutegravir combination in HIV-1 treated patients with a fully suppressed viral load and harboring M184V/I as unique NRTI resistance without resistance to integrase inhibitors. The aim of this study was to assess the efficacy of abacavir/lamivudine/dolutegravir (ABC/3TC/DTG) combination when used in pretreated patients with an undetectable viral load and previously harboring on their genotypic resistance testing M184V/I as unique NRTI resistance and without any resistance to integrase inhibitors.

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71 Patients and methods

72 Study population

HIV-1-infected patients were selected from two French university hospitals (Pitié-Salpêtrière
and Bichat-Claude Bernard Hospitals). Inclusion criteria were as follows: being suppressed on
a triple antiretroviral treatment (ART) regimen with tenofovir/emtricitabine/boosted protease
inhibitor or abacavir/lamivudine/boosted protease inhibitor for at least 12 months, having an

historical genotypic resistance test (RNA or DNA) harboring a M184V/I mutation as a unique 77 NRTI mutation, switched to ABC/3TC/DTG between 2015-2018, with at least 12 months of 78 79 follow-up. One hundred fifty four patients fully suppressed for at least 12 months with an HIV-1 plasma viral load (VL < 50 copies/mL) treated by 2 NRTIs + 1 boosted protease 80 inhibitor (PI) (tenofovir/emtricitabine/boosted 81 darunavir (n=68), tenofovir/emtricitabine/boosted atazanavir (n=56), abacavir/lamivudine/boosted darunavir 82 (n=10), abacavir/lamivudine/boosted atazanavir (n=20) and switched to an ABC/3TC/DTG 83 84 regimen were retrospectively analyzed at M1, M3, M6 and M12 after the switch to ABC/3TC/DTG. 85

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87 Ethics

The research was conducted in accordance with the Declaration of Helsinki and national and institutional standards. All the patients gave their written informed consent to have their medical chart recorded in the electronic medical record system Nadis® (www.dataids.org; CNIL number: 770134, 30 October 2001).

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93 Virological and pharmacological assays

94 Plasma HIV-1 RNA was measured using a commercially available PCR assay (Ampliprep 95 COBAS TaqMan V2.0, Roche, Meylan, France). This assay provides quantitative results for HIV-RNA values ≥20 copies/mL. Qualitative results are also given as HIV-RNA detected 96 (but <20 copies/mL) or when PCR target is not detected (VL considered <1 copy/mL).⁵ 97 98 Virological failure was defined as HIV-RNA >50 copies/mL in 2 consecutive determinations. The plasma concentrations of dolutegravir, abacavir and lamivudine were determined using a 99 100 validated LC-MS/MS method (Waters Acquity UPLC-TQD, Milford, MA, USA). Dolutegravir trough levels were interpreted according to different effective cut-offs of 101

1000 ng/mL at 24 h, based on the pharmacokinetic/pharmacodynamic relationship from the
 SAILING trial. ⁶

104 **Results**

Baseline characteristics of the 154 virologically suppressed patients switched to
ABC/3TC/DTG are shown in Table 1. Most patients were male (71.4%), the median age was
44 years (IQR 36–50) and 70% were infected with a subtype B. Median time with VL <50
copies/mL before the switch to ABC/3TC/DTG was 36 months (IQR 14–44) and 12 patients had
received at least 3 different therapeutic regimens before the switch to ABC/3TC/DTG.

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Out of the 154 patients: 16 were infected by a virus harboring M184V, 126 acquired previously a M184V/I after a treatment failure to 2 NRTIs + NNRTI or 2 NRTIs + 1 PI regimen. M184I was detected exclusively in DNA genotype and this represents only 6/154 patients. Patients were selected for having only M184V/I in their historical genotypes, so no other mutations associated with resistance to NRTI were detected. A historical integrase sequence was available for 40 patients (26%) with no evidence of resistance mutations.

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During the 12 months of follow-up, all patients maintained a VL < 50 copies/mL except three 118 119 patients who had a blip (53, 62 and 106 copies/mL) at month 3 followed by a subsequent viral 120 load < 50 copies/mL. Overall, none of the patients harbored a virologic failure (Table 2). 121 Among the 3 blips, at M1, one had no available plasma VL, one had a VL with HIV-RNA 122 detected and the last one a VL with HIV-RNA undetected. The three plasma samples 123 corresponding to the blips were sequenced to search for selection of any new resistance 124 mutations in RT and integrase genes and no emergence of mutation was evidenced. Plasma 125 measurements of abacavir, lamivudine and dolutegravir during blip episodes showed that they 126 were all in therapeutic recommended values.

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128 All plasma samples with VL< 50 copies/mL were assessed for the presence of PCR target or 129 not and no statistical evolution of the percentage of patients harboring a plasma VL below 1 copy/mL was evidenced. The percentage of patients with VL < 1 copy/mL was 70% 130 131 (108/154), 66% (67/101), 69% (85/124), 67% (73/110) and 73% (71/97), respectively. 132 133 134 Discussion 135 This study assessed the efficacy, including the impact on residual viraemia, of a switch to ABC/3TC/DTG in virologically suppressed patients (VL <50 copies/mL) with an history of 136 137 M184V/I from a clinical setting with a follow-up of 1 year. Among these 154 patients, with previous lamivudine or emtricitabine resistance, who received ABC/3TC/DTG, all maintained 138 139 virological suppression at month 12 with no evolution of residual viremia during the follow 140 up. 141 142 To our knowledge, this is the largest study evaluating the impact of M184V/I as a unique 143 NRTI resistance mutation on the virological response after a switch to ABC/3TC/DTG. In 144 addition, virological follow-up using the assessment of residual viraemia in patients switching 145 to ABC/3TC/DTG has not been previously reported. In our study we showed that 70% of 146 patients had no residual viraemia at baseline before switching and that this proportion 147 remained stable during the follow-up: 67% at month 6 and 73% at month 12. 148 149 Interestingly, M184V/I as a unique NRTI resistance mutation, regardless of possible selection 150 by regimens containing lamivudine or emtricitabine or abacavir, did not affect the response of 151 these virologically suppressed patients who switched to ABC/3TC/DTG even when switching

from a TDF containing regimen. This was true both at the 50 copies/mL and 1 copy/mL cut-152 off. Previous studies have addressed the effect of the M184V/I mutation on the efficacy of a 153 154 dolutegravir-containing regimen (dual or triple) in this context of switch and they all described low virological failure rates, irrespective of the presence of the M184V/I mutation. 155 156 In the retrospective study by Gagliardini *et al.*, in patients with a shorter duration of viral 157 suppression before simplification to dolutegravir dual regimen, the group with previous 158 detection of M184V showed higher hazards of virological failure, and the gap of efficacy 159 between the groups increased when reducing the duration of suppression, particularly below 3 years. ⁷ However, in a recent prospective pilot study (n = 41), lamivudine/dolutegravir was 160 161 effective in maintaining virologic control in integrase inhibitor naïve patients with historical lamivudine resistance when baseline proviral DNA Sanger genotype did not detect the 162 persistence of lamivudine resistance-associated mutations.⁸ This has to be confirmed by 163 164 larger prospective studies that could assess the factors associated with virological success of lamivudine/dolutegravir dual regimen. There was another study that showed no impact of 165 166 M184V/I on the risk of virological failure in virologically suppressed patients switching to 167 ABC/3TC/DTG, but in this study they did not analyze clearly the impact of M184V/I as unique NRTI resistance mutation, because at least 2 thymidine analogue mutations were 168 documented in 42.3% of patients with M184V/I.⁹ 169

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The findings of our study can be explained probably by the characteristics (potency and high genetic barrier to resistance) of dolutegravir that is able to control HIV replication even in a context where the total genotypic sensitivity score is less than 3. Indeed, in presence of M184V/I dolutegravir is considered sensitive (1), lamivudine resistant (0) and abacavir possibly resistant (0.5), thus associated with a genotypic sensitivity score of 1.5. Similarly, high rates of tenofovir alafenamide/emtricitabine/bictegravir treatment efficacy have been

observed among patients with pre-existing resistance substitutions, such as M184V/I, 177 baseline tenofovir 178 indicating that genotype did not affect 179 alafenamide/emtricitabine/bictegravir outcomes in suppressed switching participants regimens.¹⁰ Together, these results indicate that three-drug regimens with second generation 180 181 integrase inhibitors, ABC/3TC/DTG and tenofovir alafenamide/emtricitabine/bictegravir, are a treatment option for suppressed patients, including those with evidence of archived 182 183 resistance, such as M184V/I.

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Our study has several limitations. First, the observational nature of the study could have affected the results due to missing or incomplete data (such as HIV DNA). Second, the limited follow-up should be taken into consideration when interpreting the clinical relevance of our findings. Furthermore, we cannot exclude residual confounding given the absence of data on treatment adherence.

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In conclusion, our findings obtained in a clinical cohort confirmed data reported in clinical trials, showing high rate of patients maintaining pVL <50 copies/mL up to 12 months, with a stable proportion of patients (around 70%) without detectable residual viremia during the first year following the switch to ABC/3TC/DTG even in presence of an history of M184V/I as single NRTI mutation.

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206 Transparency declarations

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- 210

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Table 1. Baseline characteristics of the patients

Characteristic	n = 154
Male, n (%)	110 (71.4)
Age (years), median (IQR)	44 (36–50)
Time since HIV diagnosis (years), median (IQR)	15 (12-23)
Duration of ART (years) before switch, median (IQR)	11 (9-13)
Number of previous ART lines before switch, median (IQR)	2 (1-3)
History of INSTI-containing regimen, n (%)	0 (0)
Duration of plasma HIV-1 RNA $<$ 50 copies/mL before switch (months), median	36 (14–44)
(IQR)	
Duration between the last genotype with $M184V/I$ and time of switch (months),	52 (20-62)
median (IQR)	

248 ART, antiretroviral treatment; INSTI, integrase inhibitors

249 Table 2. Characteristics of the patients before and after the switch to250 abacavir/lamivudine/dolutegravir

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	After the switch to ABC/3TC/DTG				
ARV treatment	Type of 184 mutation (n)	Median pretherapeutic VL (cp/mL)	Median pretherapeutic CD4 cell count (mm ³)	Blips	Virologic failure
TDF/FTC+r/DRV (n=68)	M184V (66) M184I (2)	68000	286	1 (62 cp/mL at M3)	None
TDF/FTC+r/ATV (n=56)	M184V (54) M184I (2)	27000	368	2 (106 and 53 cp/mL at M3)	None
ABC/3TC+r/DRV (n=10)	M184V (9) M184I (1)	43000	371	None	None
ABC/3TC+ r/ATV (n=20)	M184V (19) M184I (1)	50000	411	None	None

252 ABC, abacavir ; ARV, antiretroviral ; ATV, atazanavir ; DRV, darunavir ; DTG, dolutegravir ; FTC, emtricitabine ; TDF, tenofovir ; 3TC,

253 lamivudine ; VL, viral load

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