



# Effectiveness of direct-acting antivirals for hepatitis C virus infection in hepatitis C/HIV coinfected individuals

# A multicenter study

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#### **Abstract**

In a hepatitis C virus (HCV)/HIV-positive Brazilian cohort, evaluate the safety and efficacy of HCV DAAs, the frequency of resistance substitutions in the HCV NS5A and NS5B genes and identify predictors of treatment failure.

Retrospective multicenter study of HCV/HIV patients treated with sofosbuvir (SOF)-based regimens at 10 reference centers in Brazil.

Clinical and virological data were collected. Genetic diversity in the NS5A and NS5B genes was assessed by direct nucleotide sequencing. The primary outcome was sustained virological response (SVR) 12 weeks after DAA completion.

Of 643 HCV/HIV patients analyzed, 74.7% were male, median CD4+ T cell count was  $617 \text{ cells/mm}^3$ , 90% had an undetectable HIV viral load. HCV genotype 1 was detected in 80.2%, and 60% were taking at least 1 medication other than antiretroviral drugs during their DAA therapy. Cirrhosis was present in 42%. An SOF/daclatasvir (DCV) regimen was used in most patients (98%). The frequency of NS5A polymorphisms associated with clinically relevant resistance to DCV was 2%; no relevant NS5B variants were identified. The SVR12 rate was 92.8% in an intention to treat (ITT) analysis and 96% in a modified ITT (m-ITT) analysis. AE occurred in 1.6% of patients. By multivariate analysis, therapeutic failure was associated, in the m-ITT analysis, with concomitant use of anticonvulsant drugs (P=.001), age (P=.004), and female gender (P=.004).

SOF/DCV regimens were associated with a high SVR rate in an HCV/HIV population. The use of concurrent anticonvulsant drugs and DAAs decreases the chances of achieving an SVR.

**Abbreviations:** AE = adverse event, DAAs = direct-action antivirals, DCV = daclatasvir, HCV = hepatitis C virus, ITT = intention to treat, SOF = sofosbuvir, SVR = sustained virological response.

Keywords: direct acting antivirals, effectiveness, hepatitis C treatment, hepatitis C-HIV coinfection, safety

# 1. Introduction

Hepatitis C virus (HCV) and HIV coinfection is a major global public health problem. [1,2] According to the World Health Organization (WHO), there are currently around 71 million

individuals chronically infected with HCV, 31 million with HIV, and more than 2 million with HCV-HIV coinfection. [2,3] Since HCV and HIV infections share the same routes of transmission, coinfections are often found with an estimated prevalence of 25%

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The datasets generated during and/or analyzed during the current study are available from the corresponding author on reasonable request.

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to 30% in the general population of people living with HIV/AIDS. These rates are even higher in specific groups, such as in injection drug users. [4,5] Coinfection adversely affects the natural course of hepatitis C infection, accelerating the progression of liver fibrosis and increasing the risk of hepatic decompensation and hepatocellular carcinoma in cirrhotic patients. [6–8] Currently, end-stage liver disease is one of the leading causes of death in the HCV/HIV population. [5,9,10]

The advent of direct-action antivirals (DAAs) has substantially changed the HCV therapy scenario, including in the HIV-infected population. [11–13] In the interferon (IFN) treatment era, a minority of individuals could be treated due to the occurrence of significant adverse effects and low sustained virological response (SVR) rates. [12,14,15] With DAAs, especially IFN-free regimens, HCV therapy has been simplified and coinfected patients have achieved high cure rates. [11,12] Randomized clinical trials and real-life studies in different parts of the world have shown SVR rates greater than 90% in patients with HCV/HIV, similar to that reported for patients without HIV coinfection. [16–21]

However, conflicting results regarding the impact of HIV and immunosuppression on SVR rate in these patients have been reported. [22-24] In some studies, HIV coinfection has been associated with worse clinical outcomes. [22-25] In addition, for this population, adherence to therapy, the high frequency of comorbidities, the concomitant use of medications with potential drug–drug interactions (DDI) with DAAs, and the impact of the presence of resistance-associated substitutions (RASs) in the HCV genome are additional significant challenges that could interfere with the chances of treatment success. [11,26-29]

In Brazil, the public health system has provided free-of-charge treatment with IFN-free DAAs regimens beginning in 2015 to all HCV/HIV coinfected patients, irrespective of the degree of liver fibrosis. [30] We now present results of a real-life study in this population with the following objectives: to estimate the real-life safety and efficacy of sofosbuvir (SOF)-based treatment regimens in an HCV/HIV Brazilian cohort; to investigate the frequency of baseline RASs in HCV NS5A and NS5B genes, and to identify predictors of treatment failure.

#### 2. Methods

# 2.1. Study design and population

We conducted an observational, retrospective, and multicenter study of patients at 10 Brazilian reference centers specialized in the treatment of HCV. These centers are located in 3 Brazilian states: São Paulo, Rio Grande do Sul, and Rio de Janeiro.

Eligible subjects were all patients with HCV/HIV coinfection who started HCV therapy with SOF-based regimens between July 2015 and August 2017, according to the Brazilian Ministry of Health's Therapeutic Guidelines. <sup>[30]</sup> Inclusion criteria were: age ≥ 18 years, having received at least 1 dose of the prescribed treatment regimen, and no history of previous therapy with SOF and/or daclatasvir (DCV) or with any other NS5A antiviral regimen. Patients were not excluded based on the presence of comorbidities.

According to the Brazilian Therapeutic Guidelines, the following treatment regimens were administered for the different HCV genotypes (GT): for HCV GT 1 and GT4: SOF and DCV; for GT2: SOF-ribavirin (RBV); for GT3: SOF-DCV combination or SOF plus pegIFN-RBV. [30] The choice of concomitant

administration of RBV and/or the treatment of HCV GT3 patients was made at the discretion of the attending physician.

This study was approved by the Ethics Committee for the Analysis of Research Projects (Comissão de Ética para Análise de Projetos de Pesquisa—CAPPesq—Hospital das Clínicas da Faculdade de Medicina da Universidade de São Paulo—HC-FMUSP) under protocol no. 46221015.0.0000.0068. All patients provided written informed consent prior to inclusion in the study.

# 2.2. Data collection

Demographic, clinical, and laboratory data were collected. Data was obtained from medical files after the end of treatment, by using a standard clinical questionnaire. For staging of liver disease, the results obtained by liver biopsy were considered according to the classification of the METAVIR Co-operative Study Group. [31] Liver elastography was also used for liver fibrosis staging. The corresponding transient elastography cutoff points (FibroScan, Echosens, France) for Metavir were less than 5.5 kPa = F0, 5.6 to 7.0 = F1, 7.1 to 9.4 kPa = F2, 9.5 to 12.4kPa = F3, and  $\geq 12.5$  kPa = F4. [32] Cirrhosis was defined by liver biopsy or noninvasive method of liver fibrosis, according to the cut-off values mentioned above, or by clinical signs of portal hypertension. When biopsy or elastography were not available, those with APRI score  $\geq 2$  and FIB score  $\geq 3.25$  were also classified as cirrhotic. [33–35] Comorbidity was defined as the presence of any clinical condition that required clinical or pharmacological intervention by the medical team. Known DDI between DAAs and other concomitant medications were identified from the website www.hep-druginteractions.org.

#### 2.3. Outcomes

The primary efficacy endpoint was the proportion of patients with SVR, defined as an HCV RNA level below the limit of quantification (HCV RNA PCR <12 IU/mL; Abbot Molecular, Des Plaines, IL) following 12 weeks of treatment among all those who received at least 1 dose of the study treatment (intention-to-treat analysis (ITT)).

We also analyzed SVR, in a modified intention-to-treat analysis (m-ITT), in which we excluded cases such as loss to clinical follow-up, treatment interruptions, and/or deaths not associated with DAA adverse events.

An adverse event (AE) was defined as any event during treatment that required a dose change or discontinuation of a drug from HCV therapy regimen or other drug intervention. Serious adverse events (SAE) were defined as those that led to discontinuation of DAA therapy or death.

# 2.4. Baseline RASs

We analyzed for RASs in pretreatment samples. HCV RNA extraction was performed from serum using the QIAamp Viral RNA Kit (Qiagen, Hilden, Germany), followed by amplification using the SuperScript III / Platinum Taq High Fidelity One-Step enzyme (InvitrogenTM, Thermo Fisher Brand, Carlsbad, USA), following the manufacturer's protocol. The amplified samples were submitted to a sequencing reaction (bidirectional) derived from the Sanger methodology. The sequences obtained were initially analyzed using the Electropherogram quality analysis program (http://asparagin.cenargen.embrapa.br/phph/). By aligning the amino acid sequences of each sample with reference

Table 1

#### Baseline characteristics of 643 Brazilian HCV-HIV patients.

Characteristics	Overall N=643
Male sex, n (%)	480 (74.7)
Age, median (range)	50 (21; 74)
Injection drug use, n (%)*	217 (49.9)
Comorbidity, n (%)	403 (62.7)
Nonantiretroviral drugs, n (%)	376 (58.5)
Drugs with clinically significant DDI with DAAs	9 (1.4)
CD4+T-cell (cell/mm <sup>3</sup> ) <sup>†</sup>	
median (range)	617 (73; 2321)
< 200, n (%)	31 (4.9)
200-499, n (%)	190 (30.2)
≥500, n (%)	409 (64.9)
Undetectable HIV RNA, n (%) <sup>‡</sup>	551 (89.9)
CDC clinical category C, n (%)§	267 (49.4)
On antiretrovirals, n (%)	636 (98.9)
Prior IFN-based therapy, n (%)	343 (53.3)
HCV Genotype, n (%)	
GT 1	516 (80.2)
1a	322
1b	111
1a/1b	12
Subtype unknown	71
GT 2	9 (1.4)
GT 3	85 (13.2)
GT 4	33 (5.1)
Metavir score, n (%)	
F0-F1	158 (28.1)
F2	100 (17.8)
F3	68 (12.1)
F4/Cirrhosis	238 (42.2)
Child-Pugh A	203 (85.3)
Child-Pugh B	35 (14.7)

Missing data: \* 208; † 13; \* 30; §102; | 79.

CDC = centers for disease control and prevention, DDI = drug-drug interaction, GT = HCV genotype, IFN = interferon.

sequences, NS5A inhibitor RASs were screened at positions 28, 30, 31, 58, 92, and 93; and for NS5B inhibitors at positions 282, 316, and 321. [29,37,38] In addition, all consensus sequences were submitted to Geno2pheno [hcv] 0.92 (https://hcv.geno2pheno.org/) for confirmation of the results. [39]

# 2.5. Statistical analysis

We used bivariate analyses followed by logistic regression to identify factors associated with lack of a SVR. Categorical variables were compared using  $\chi^2$  or Fisher exact tests and

continuous variables were compared using the Student t or Mann–Whitney test. Variables with P < .10 were selected for logistic regression analysis. Variables with P < .05 in the multiple analysis were retained in the final model. Odds ratios (OR) of each variable were estimated with the corresponding confidence interval (95% CI) at the 5% significance level. IBM-SPSS for Windows version 20.0 software was used to perform the analysis.

# 3. Results

#### 3.1. Baseline characteristics

The main baseline characteristics of the 643 HCV/HIV coinfected patients who started DAA therapy are presented in Table 1. The majority were male (74.7%), median age 50, 90% with an undetectable HIV viral load and a median CD4 + T cell count of 617/mm³. Liver cirrhosis was identified in 42% of patients, mostly Child-Pugh A (85.3%) and Child-Pugh B (14.7%). There were no subjects with Child-Pugh C cirrhosis in our cohort. About 60% of patients had at least 1 comorbidity, with an average of 2 medical conditions per patient (range 1–9), including extrahepatic manifestations and hepatitis B virus coinfection. One hundred sixteen patients (18%) had 3 or more comorbidities. The most frequent comorbidities observed were: systemic arterial hypertension (n=150), psychiatric disorders (n=142), diabetes mellitus (n=79), and dyslipidemia (n=72). Twelve patients had hepatitis B virus coinfection (HBsAg positive).

Almost all patients (n=636) were taking antiretroviral medication. Antiretroviral treatment therapy (ART) was done according to Brazilian Guidelines. ART was documented and no combination was excluded. Among these patients, 250 (38.9%) had their medication modified prior to initiating DAA to prevent drug-drug interactions. During HCV treatment with DAAs, 376 patients (58.5%) reported the concomitant use of other medications (an average of 3 drugs per patient) that were unrelated to HIV or HCV treatment. Antihypertensive drugs, psychotropic drugs, and those used to treat diabetes were the most commonly used medication classes. Although anticonvulsant medications are generally contraindicated during SOF and DCV treatment due to the risk of adverse interactions, these drugs were prescribed in 9 (1.4%) cases, as follows: carbamazepine (n=4), phenobarbital (n=2), phenytoin (n=2), and oxcarbazepine (n=1).

# 3.2. Treatment regimens

Table 2 presents the DAAs regimens used in this cohort, according to HCV genotypes.

# Table 2

Treatment regimens and sustained virologic response rate according HCV genotype.

Dogimon n (0/)	%SVR GT1	%SVR GT2	%SVR GT3	%SVR GT4	%SVR total
Regimen, n (%)	(n=516)	(n = 9)	(n = 85)	(n = 33)	(n = 643)
SOF+DCV 12w	97.2 (175/180)	100 (1/1)	85.7 (18/21)	89.5 (17/19)	95.6 (211/221)
SOF+DCV 24w	85.7 (6/7)	0	50 (1/2)	0	77.8 (7/9)
SOF+DCV+RBV 12w	91.4 (257/281)	100 (2/2)	94.3 (50/53)	92.3 (12/13)	92 (321/349)
SOF+DCV+RBV 24w	95.8 (46/48)	0	50 (1/2)	100 (1/1)	94.1 (48/51)
SOF+RBV 12w	0	83.3 (5/6)	0	0	83.3 (5/6)
SOF+pegIFN+RBV 12w	0	0	71.4 (5/7)	0	71.4 (5/7)

<sup>12</sup>w=12 weeks, 24w=24 weeks, DCV=daclatasvir, GT=HCV genotype, pegIFN=pegylated interferon, RBV=ribavirin, SOF=sofosbuvir, SVR=sustained virologic response.

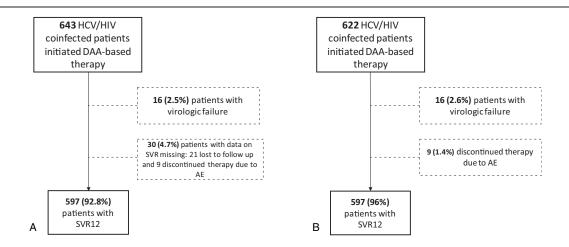


Figure 1. DAA = direct-acting antiviral, SVR = sustained virological response, AE = adverse event. (A) = intention-to-treat approach, (B) = modified intention-to-treat approach.

# 3.3. Response to treatment

The overall rate of SVR in the ITT analysis was 92.8% (n=597; 95% CI 90.9–94.8). Sixteen subjects (2.5%) did not achieve a virological response. For 30 patients (4.7%) no SVR data was available: 9 subjects (1.4%) discontinued DAAs treatment due to SAE and 21 patients (3.3%) were lost to follow-up, interrupted treatment or died due to different causes not associated with HCV or the use of DAAs. In the m-ITT analysis, the SVR rate was 96% (95% CI 94.4%–97.5%) (Fig. 1).

Cirrhotic patients achieved lower SVR rates when compared with noncirrhotic patients, but with no significant difference in both analyses (P=.21 and P=.44 in the ITT and m-ITT approaches, respectively). Child-Pugh B cirrhotic patients (N=35) achieved SVR rates of 88.6% in the ITT analysis and 91.1% in the m-ITT analysis.

SVR rates also did not differ significantly among HCV GTs or between Child-A or Child-B patients. Although cirrhotic patients positive for GT3 had a lower SVR rate (74.1%), this was not significantly different from cirrhotic patients with other HCV genotypes (P = .90).

# 3.4. Safety

One hundred three patients experienced an AE during DAA therapy (AE rate: 16%; 95% CI 13.2–18.9). The most frequent AE was anemia, reported for 90 patients, all of whom were using RBV. The diagnosis of anemia secondary to the use of ribavirin was made by the attending physician, based on a decrease in hemoglobin levels during hepatitis C therapy compared with baseline hemoglobin values. None of the cases with anemia resulted in discontinuation of HCV treatment.

SAE occurred in 10 individuals (1.6%; 95% CI 0.6-2.5), of which 2 died due to liver decompensation; both were Child-Pugh B cirrhotic. The other SAE reported were as follows: decompensation of psychiatric disorder (n=1), elevation of alanine aminotransferase, and aspartate aminotransferase to a level 10 times the upper limit of normal (n=1) and gastrointestinal intolerance (n=6).

Among the 10 patients with SAE, 1 individual with gastrointestinal intolerance completed 8 weeks of therapy and achieved SVR. The other patients did not achieve SVR.

# 3.5. HCV gene polymorphisms

We analyzed for polymorphisms in the HCV NS5B and NS5BA genes in 261 (40.6%) and 254 (39.5%) patients, respectively. In the analysis of the HCV NS5A region, genetic variants were found in 52 (20.5%) patients, In 5 of these individuals (2.0%) the substitutions conferred clinically relevant resistance to DCV. [40] The amino acid variants identified were: L31 M in 1 patient with HCV GT 1a, Y93C in 1 patient with HCV GT 1a, and Y93H in 3 HCV GT 1b-positive patients.

#### 3.6. Predictors of treatment failure

In the final logistic regression model, in both the ITT and m-ITT analyses, the use of drugs with relevant DDI with DAAs was significantly associated with non-SVR (P=.004 and P=.001, respectively). In the ITT analysis, the absence of prior IFN therapy was also associated with therapeutic failure (P=.04), whereas in the m-ITT analysis, female gender and age (> 50 years old) were also predictive of lack of SVR (P=.04 and P=.04, respectively). The m-ITT analysis results are summarized in Table 3.

To evaluate whether characteristics associated with HIV infection could interfere with the likelihood of obtaining SVR, we performed additional analyses on patients with known CD4 + T lymphocyte counts, HIV viral load, and a CDC clinical classification for AIDS. In bivariate and multivariate analysis of these  $520\,\mathrm{HCV/HIV}$  coinfected individuals, only the use of drugs with significant DDI was associated with treatment failure (P=.01). The results are presented in Table 4.

# 4. Discussion

In a cohort of 643 HCV-HIV coinfected patients from 10 centers in Brazil, the SVR rate was 96% following SOF and DCV-based therapy. Using a modified-ITT analysis, therapeutic failures were

Table 3

Results of bivariate and multivariate analyses of factors associated with lack of SVR in a modified intention-to-treat approach—643 HCV-HIV coinfected patients.

Variable	Bivariate analysis			Multivariate analysis		
	OR	95 CI	P	OR	95 CI	Р
Male sex	0.42	0.19-0.95	.03	0.42	0.18-0.99	.04
Age (per year)	1.058	1.004-1.115	.04*	1.06	1.00-1.12	.04
Comorbidity	1.93	0.76-4.90	.16			
Nonantiretroviral drugs	2.33	0.92-5.92	.07	1.62	0.62-4.27	.33
Drugs with clinically significant DDI with DAAs	16.15	3.63-71.87	.003 <sup>†</sup>	12.54	2.68-58.54	.001
Baseline CD4 <sup>+</sup> T-cell/mm <sup>3#</sup>	1.00	0.999-1.001	.86 <sup>‡</sup>			
Detectable baseline HIV RNA##	0,86	0.2-3.76	$>.99^{\dagger}$			
On antiretrovirals	§		>.99			
CDC clinical category C###	2.00	0.78-5.08	.14			
Prior IFN-based therapy	1.04	0.47-2.33	.92			
HCV Genotype 1 (vs GT non-1)	0.50	0.21-1.16	.12 <sup>†</sup>			
HCV Genotype 3 (vs GT non-3)	2.2	0.85-5.68	.12 <sup>†</sup>			
Cirrhosis	1.38	0.61-3.08	.44			
Cirrhosis Child-Pugh B	2.49	0.71-8.77	.15 <sup>†</sup>			
HCV RNA, median	1.00	1.00-1.00	.40 <sup>‡</sup>			
12 wks (vs 24 wks)	1.88	0.62-5.67	>.28 <sup>†</sup>			
Ribavirin use	1.49	0.61-3.62	.38			
RAS associated with resistance to DCV####	§		.99 <sup>†</sup>			

Missing data: (#) 13, (##) 30, (###) 102, (####) 389.

Table 4

Results of bivariate and multivariate analyses of factors associated with lack of SVR for 520 HCV-HIV coinfected patients with complete data on HIV infection.

Variable	Bivariate analysis			Multivariate analysis		
	OR	95 CI	P	OR	95 CI	Р
Male sex	0.58	0.28-1.20	.14			
Age (per year)	1.019	0.975-1.064	.42*			
Comorbidity	1.30	0.62-2.72	.49			
Nonantiretroviral drugs	1.68	0.79-3.59	.18			
Drugs with clinically significant DDI with DAAs	15.58	3.02-80.40	.004 <sup>†</sup>	9.94	1.69-58.37	.01
Baseline CD4 <sup>+</sup> T-cell/mm <sup>3</sup>	1.00	0.999-1.001	.59 <sup>‡</sup>			
Detectable baseline HIV RNA	1.50	0.56-4.06	$.39^{\dagger}$			
On antiretrovirals	§		>.99 <sup>†</sup>			
CDC clinical category C	1.97	0.96-4.08	.06	1.88	0.9-3.95	.09
Prior IFN-based therapy	0.49	0.24-1.00	.05	0.52	0.25-1.08	.08
HCV Genotype 1 (vs GT non-1)	0.61	0.27-1.34	.21			
HCV Genotype 3 (vs GT non-3)	1.44	0.57-3.61	.43			
Cirrhosis	1.64	0.81-3.29	.16			
Cirrhosis Child-Pugh B	2.04	0.58-7.20	.22			
HCV RNA, median	1000	1.00-1.00	.46 <sup>‡</sup>			
12 wks (vs 24 wks)	0.93	0.27-3.15	$>.99^{\dagger}$			
Ribavirin use	1.95	0.83-4.57	.12			
RAS associated with resistance to DCV#	3.9	0.41-37.10	.28 <sup>†</sup>			

Missing data: (#) 266.

CI = confidence interval, DCV = daclatasvir, DDI = drug-drug interaction, GT = genotype, IFN = interferon, OR = odds ratio, RAS = resistance-associated substitutions.

 $<sup>\</sup>chi^2$  test: \* Student t test.

<sup>†</sup> Fisher exact test.

<sup>\*</sup> Mann-Whitney test.

 $<sup>\</sup>S$  There are no cases to estimate.

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<sup>†</sup> Fisher exact test.

<sup>\*</sup> Mann-Whitney test.

 $<sup>\</sup>S$  There are no cases to estimate.

associated with the concomitant use of anticonvulsant drugs (P=.001), age above 50 years old (P=.04), and female gender (P=.04). Regarding drug safety, less than 2% of patients discontinued therapy due to AE. Factors related to HIV immunosuppression were not associated with lower rates of a therapeutic response. Lastly, the frequency of baseline mutations in HCV associated with clinical resistance to DAAs was infrequent, around 2%, and did not impair the SVR rate.

The observed high SVR rate is consistent with results of prior studies on smaller numbers of subjects from outside Brazil that employed the same therapeutic regimen. [40-43] Other studies, however, have associated the presence of HIV coinfection with a higher failure rate of anti-HCV DAA. [20,22-24] Berenguer et al [22] observed in a cohort of more than 2000 coinfected individuals that CD4+ T lymphocyte counts below 200/mm³ and a history of opportunistic disease were associated with therapeutic failure. In contrast, ARV therapy was successful in the majority of patients (98.9%) in our study and less than 5% had a CD4+ T cell count below 200/mm³. It is likely that this effective control of HIV infection contributed to the observed high positive rates of SVR and negated our ability to evaluate the possible impact of variables associated with immunodepression on success rates.

Other factors, such as infection with HCV GT3, liver cirrhosis, or baseline RASs, have also been related to reduced SVR. [19,37,44-<sup>46</sup> RBV use and the duration of therapy have also been reported to interfere with SVR rates in specific populations. [20,40,45] However, these variables were not associated with a lack of SVR in our cohort. In our study, patients who received a regimen of SOF+DVC 24 weeks (n=9; 7 HCV GT1 and 2 HCV GT3) and SOF+pegIFN+RBV (n=7; all HCV GT3; 50% with cirrhosis) had relatively low SVR rates (77.8% and 71.4%, respectively). We did not find a statistical difference in the SVR rates between HCV genotypes 1 and 3 or duration of treatment (12 versus 24 weeks). It is important to highlight that the number of patients receiving these 2 therapy regimens was small (n=16) and according to national guidelines, the choice of concomitant administration of RBV and/or the treatment of HCV GT3 positive patients with IFN-based regimen was made at the discretion of the attending physician.

In our cohort, patients with liver cirrhosis had slightly lower SVR rates than those observed in the noncirrhotic population, but without statistical significance (P=.21 in ITT analysis and P=.44 in m-ITT analysis). When analyzing only cirrhotic patients (N=238), those positive for HCV GT3 achieved the lowest SVR rate (74.1%), but this also was not statistically significant when compared with cirrhotic individuals with other HCV genotypes (P=.90).

Our analysis identified that anticonvulsant medications predicted a reduction in the SVR rate (OR=12.54 in m-ITT analysis). Nine patients (1.4%) were prescribed anticonvulsant medications to control chronic neurological or psychiatric disorders. Anticonvulsant medications are reported to be potent inducers of P-glycoprotein polypeptide and cytochrome P450-CYP3A4 enzymes capable of decreasing sofosbuvir and daclatasvir plasma concentrations, respectively, leading to reduced therapeutic effect of these antivirals. [47–50] It is important to emphasize that in our series, the use of anticonvulsant medications was associated with therapeutic failure in all analyses performed, whether in the multivariate analysis of the 643 patients (by IIT or m-ITT) or in the separate analysis of the 520 coinfected patients, for whom we had all the clinical information related to HIV immunosuppression.

In our study, female gender and age were also significantly associated with a reduction in the SVR (P=.04 and P=.04 with modified-ITT analysis). It is known that with IFN-based anti-HCV therapy, gender (specifically male), and advanced age are predictive of nonresponse to treatment. [51,52] With the advent of DAAs, these variables have become of less concern, as high rates of SVR have been achieved, and most studies have not reported their association with therapeutic failure. [16-19] In the few studies that identified an impact of gender and age on SVR in the DAAs era, the results have been conflicting. Berenguer et al found that males had a higher rate of virological failure in a Spanish HCV/ HIV cohort, while Bischoff reported in a German cohort that males aged over 50 years had a higher SVR rate. [20,22] Our sample consisted predominantly of men (74.7%) and individuals over 40 years of age (89.9%). We found that the prevalence of comorbidities, psychiatric disorders, and the use of non-ARV medications were higher in females than in men in our cohort. These variables, either individually or in combination, may have contributed to the observed female association with therapeutic failure.

The impact of comorbidities on DAAs treatment outcomes has been described previously. [53–56] A recent Brazilian study with mono-infected patients showed that having a higher number of comorbidities was significantly associated with therapeutic failure using SOF-based regimens. [55] In the cohort of HCV/HIV coinfected patients described by Cachay et al, [56] the only factor identified as associated with a lower SVR was the presence of psychiatric disease. Considering the evidence that individuals with HIV infection have a higher prevalence and number of comorbidities, especially psychiatric disorders, it is essential in this population to evaluate all possible comorbidities, medication usage, known drug interactions, and patient adherence to achieve an optimal outcome with HCV therapy. [26,53,56,57]

Regarding substitutions in the HCV genome associated with clinical resistance to DAAs, these were identified in less than 2% of our patients analyzed. The frequency of RASs in HCV NS5A genes in our study was lower than observed in other studies, [38,58–60] and their presence did not impair SVR rates in our population. An explanation for our results is that RASs, especially in the HCV NS5A gene, appear to impact the treatment response mainly in patients infected with HCV GT 1a and GT 3a. [37,38,58] In addition, we highlight that in the present study, RASs were not found at baseline in HCV GT3-positive patients (N=32). This contrasts with results of other Brazilian studies where amino acid substitutions at positions 30 and 93 were identified. [61,62]

Limitations in our study need to be acknowledged. Its retrospective nature did not allow us to obtain totally comprehensive data for all included patients nor was it possible to adequately assess adherence to treatment, relying only on data from medical records. In addition, the high SVR rates observed made it difficult to identify predictors of treatment failure, resulting in a reduction in the statistical power of these analyses. We acknowledge that, although the SOF-DCV regimen is not currently a first-line regimen in the Americas, Europe, and Brazil, it remains one of the preferred regimens according to WHO guidelines. Despite these limitations, this is one of the largest series of subjects to be evaluated using SOF-DCV combination therapy in people living with HIV as well as possibly the largest case series of HCV/HIV coinfected patients in Latin America.

Considering the epidemiological setting and the impact of HCV/HIV coinfection and its outcomes on global public health, the results regarding the HCV treatment response in this

population may contribute to the implementation of improved global strategies for HCV elimination. In addition, it is important to note that HCV treatment in this population goes beyond SVR. Management of the risk for reinfection and the need for hepatocarcinoma surveillance, especially in patients with advanced liver disease, are priorities in the clinical follow-up of this population. It is also necessary to mention that subgroups of coinfected patients continue to have specific barriers to effective care. Comorbidities, especially psychiatric issues, dependency on alcohol, and other drugs, difficulties to access health services and even situations of incarceration reduce access to HCV treatment. [26,28,63] Implementation of health programs involving multidisciplinary teams could improve the overall clinical management of this population.

In summary, the high cure rate of HCV/HIV coinfected individuals after SVR in our multicenter population confirms the efficacy and safety of this protocol and supports its universal distribution.

# **Author contributions**

SMM and MCMC developed the original idea to conduct this study. They were responsible for the data analysis, wrote the first draft and final revision of the manuscript.

AGV, AGL, ACMD, PRAF, DCJ, SBT, CEBM, MPG, FS, KDP, DVN: participated in the study design, in the patient cohort recruitment and data collection and revision of the manuscript.

GLN and RSA were responsible for data analysis JRRP and FMM were responsible for laboratory analysis SSW was responsible for the study concept and revision of the article.

All authors assessed, reviewed, and edited the manuscript and approved the final version for publication.

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