

Impact of Patient and Viral Factors on Virologic Outcome with Long-Acting Cabotegravir and Rilpivirine in Phase 3 Trials (ATLAS, FLAIR, and ATLAS-2M)

Summary

Week 48 Results

- A post hoc multivariable analysis (MVA) showed that 4 factors were associated with confirmed virologic failure (CVF) through Week 48 in FLAIR, ATLAS, and ATLAS-2M:
 - o RPV resistance-associated mutations (RAMs) at baseline
 - Post-hoc RPV C_{min} at Week 8
 - o HIV-1 subtype A6/A1 at baseline
 - Body mass index (BMI) at baseline
- A post hoc baseline factors analysis (BFA) showed that there is an increased risk of CVF if 2 or more baseline factors (among RPV RAMs, HIV-1 subtype A6/A1, and BMI ≥30 kg/m²) are present.
 - The rate of CVF among patients with no or 1 factor present was <0.5%

Beyond Week 48

- Using data through the of study for FLAIR (Week 124), ATLAS (Week 96), and ATLAS-2M (Week 152) the MVA was updated and revealed the following factors associated with CVF:
 - o RPV RAMs at baseline
 - o HIV-1 subtype A6/A1 at baseline
 - o Predicted log₂ cabotegravir and rilpivirine troughs at Week 44
 - o Predicted log₂ cabotegravir trough at Week 4
- The updated BFA confirmed the results of the BFA conducted at Week 48
- Important Safety Information can be found in the <u>Prescribing Information</u> and can also be accessed from the <u>Our HIV Medicines</u> section of viivhealthcare.com/us.

To access additional scientific information related to ViiV Healthcare medicines, visit the ViiV US Medical Portal at viivhcmedinfo.com.



BACKGROUND

Overall in Phase 3 trials reported to date, CAB + RPV LA maintained virologic suppression in 94% (1531/1636) through Week 48.¹ Despite high levels of adherence with the medicines, the rate of CVF (defined as 2 consecutive HIV-1 RNA \geq 200 copies/mL) was 1-2%. For more information about the efficacy and safety of monthly CAB + RPV LA reported in ATLAS and FLAIR please click here. For more information about the efficacy and safety of CAB + RPV LA administered every 2 months in ATLAS-2M please click here.

MULTIVARIABLE ANALYSES

Week 48

The objective of this post hoc MVA was to provide an understanding of potential factors associated with CVF through Week 48 among patients receiving CAB + RPV LA in ATLAS, FLAIR, and ATLAS-2M.¹ A logistic regression model was used to assess the influence of 10 covariables suspected to contribute to virologic outcomes and included:

- CAB trough at Week $8 \le 25^{th}$ percentile ($\le Q1$)
- RPV trough at Week 8 ≤Q1
- HIV-1 subtype A6/A1
- Baseline L74I
- Baseline integrase (IN) mutation
- Baseline RPV RAMs
- Baseline non-nucleoside reverse transcriptase inhibitor (NNRTI) mutation
- Female at birth
- BMI ≥30 kg/m²
- Every-8-week dosing of CAB + RPV LA

Results

Patients were included if they were naïve to CAB + RPV LA (n=1039). Patients (n=597) were excluded from the analysis if they were not naïve to CAB + RPV LA (n=391), had not received CAB + RPV LA (n=22), or had missing data (n=184).

Seventeen patients met the criteria for CVF through Week 48.¹ Of these, 13 patients were included in the MVA. Four patients were excluded from the analysis.

See Table 1 below for a listing of the 10 potential factors among the 13 patients included in the analysis. As can be seen, 12 of the 13 patients had 2 or more factors present.

It should also be noted that there was a high degree of correlation between the presence of L74I (integrase polymorphism) and HIV-1 subtype A6/A1 (see Table 2 below). In this analysis, the presence of L74I only appears to be relevant in the context of HIV-1 subtype A6/A1. Among this group of patients, the rate of CVF was 6.6% (7/106). There were no (0/14) CVFs in patients with HIV-1 subtype A6/A1 and without L74I. Among patients with L74I and HIV-1 subtype B, the most prevalent subtype in North American and Western Europe, there were no CVFs.

Table 1. Per-Patient Accounting of Covariates Included in the Multivariable Analysis¹

Study	Patient ID	CAB C _{min} ≤Q1	RPV C _{min} ≤Q1	HIV-1 Subtype A6/A1	Baseline L74I	Baseline IN Mutation	Baseline RPV Mutation	Baseline NNRTI Mutation	Female at Birth	BMI ≥30 kg/m²	Q8W Dosing
ATLAS-2M	1										
ATLAS-2M	2										
ATLAS	3										
ATLAS	4										
FLAIR	5										
FLAIR	6										
FLAIR	7										
ATLAS-2M	8										
ATLAS-2M	9										
ATLAS	10										
ATLAS-2M	11										
ATLAS-2M	12										
ATLAS-2M	13										

CAB = cabotegravir; C_{min} = trough concentration at prior to Week 8 administration of CAB + RPV LA (4 weeks following first injections); Q1 = 25th percentile; RPV = rilpivirine; IN = integrase; NNRTI = non-nucleoside reverse transcriptase inhibitor; BMI = body mass index; Q8W = every-8-week dosing

Table 2. Proportion of Patients with CVF with and without L74I Across HIV-1 Subtypes¹

Subtype	With	L74I	Without L74I		
	n/N (%)	95% CI	n/N (%)	95% CI	
\6/A1	7/106 (6.6)	2.7 to 13.1	0/14	0.0 to 23.2	
A other	0/4	0.0 to 60.2	0/13	0.0 to 24.7	
	1/7 (14.3)	0.4 to 57.9	0/70	0.0 to 5.1	
3	0/41	0.0 to 8.6	4/714 (0.6)	0.2 to 1.4	
Other	0/5	0.0 to 52.2	1/65 (1.5)	0.0 to 8.3	
Total	8/163 (4.9)	2.1 to 9.4	5/876 (0.6)	0.2 to 1.3	

Apr-23

Four factors were associated with an increased risk of CVF. See Table 3 below for details. Of note, there was no association between female gender at birth or every-8-week dosing and CVF.

It is important to note that the magnitude of the odds ratio in Table 3 below does not show that there is a causal relationship between certain covariables and CVF; it shows the strength of the association.¹

Table 3. Strength of Association of the Covariables and CVF through Week 481

Covariable	Odds Ratio* (95% CI)	P value	
RPV RAMs at baseline [†]	37.24 (8.44 to >99)	<0.001	
Log₂ of post hoc Week 8 RPV C _{min}	4.17 (1.59 to 11.11)	0.004	
Baseline HIV-1 subtype A6/A1	6.59 (1.82 to 25.26)	0.005	
BMI at baseline	1.13 (1.03 to 1.25)	0.014	
Pre-specified IN mutation (excluding L74I non-M mixture) at baseline [‡]	0.11 (0.01 to 0.83)	0.029	
Log ₂ of post hoc Week 8 CAB C _{min}	Not significant		
Female at birth	Not significant		
Q8W dosing	Not significant		
L74I (non-M mixture) polymorphism at baseline	Not significant		
NNRTI RAMs (excluding RPV RAMs) at baseline	Not significant		

^{*}Odds ratios (ORs), 95% penalised profile CIs and penalised likelihood ratio p-values are provided. Covariates with p<0.05 in the final backwards elimination model are presented. CAB and RPV PK parameters were log2-transformed; therefore, the corresponding ORs are per halving of each variable.

CVF = confirmed virologic failure; RPV = rilpivirine; RAMs = resistance associated mutations; C_{min} = trough concentration; BMI = body mass index; CAB = cabotegravir; Q8W = every-8-week dosing; NNRTI = non-nucleoside reverse transcriptase inhibitor

Beyond Week 48

The MVA presented above was updated with data through Week 124 from FLAIR, Week 96 from ATLAS, and Week 152 from ATLAS-2M.² The pooled analysis includes 1651 patients with up to 3 years on study and 1292 patients with non-missing information on selected baseline and post-baseline factors. There were a total of 19 patients with CVF who were exposed to only every-4-week (n=11) or every-8-week (n=8) CAB + RPV LA. Data is only available for patients who received either every-4-week or every-8-week CAB + RPV LA, but not both.

See Table 4 for the rate of CVF by treatment regimen.

[†]Identified per the IAS-USA 2019 list of mutations.

[‡]Identified per the IAS-USA list of mutations associated with resistance to bictegravir, CAB, dolutegravir, elvitegravir or raltegravir and observed mutations during in vitro passage of dolutegravir or seen in a previous dolutegravir study (NCT01328041) in INSTI-experienced subjects.

Table 4. Rates of CVF by Dosing Regimen through End of Study in FLAIR, ATLAS, and ATLAS-2M²

CAB + RPV LA Regimen at CVF				
	Every-4-week Only (N=1129)	Every-8-week Only (N=327)	Switch from every-4-week to every-8-week (N=195)	Overall (N=1651)
CVFs n (%)	11 (1)	8 (2.4)	4 (2.0)	23 (1.4)
Person-years	2621	936	734	4291
Incidence rate/100 person years	0.42	0.85	0.54	0.54
95% CI for incidence rate	(0.21, 0.75)	(0.37, 1.68)	(0.15, 1.40)	(0.34, 0.80)

CAB + RPV LA = long-acting cabotegravir and rilpivirine; CVF = confirmed virologic failure

As seen in the Week 48 MVA, rilpivirine RAMs and HIV-1 subtype A6/A1 remained significant predictors of CVF (see Table 5 below). Predicted cabotegravir and rilpivirine log₂ trough concentrations at Week 44 and predicted cabotegravir log₂ trough at Week 4 were new predictive factors identified. Body mass index and rilpivirine log₂ trough at Week 4 were no longer predictive of CVF.

Table 5. Strength of Association of the Covariables and CVF through End of Study from FLAIR, ATLAS, and ATLAS-2M²

Covariable	Odds Ratio* (95% CI)	<i>P</i> value	
RPV RAMs: Yes/No	25.7 (7.17, 92.2)	<0.0001	
HIV-1 subtype A6/A1: Yes/No	15.5 (4.69, 50.9)	<0.0001	
Predicted log₂ Week 44 CAB trough*	5.99 (1.94, 18.5)	0.0019	
Predicted log₂Week 44 RPV trough*	4.16 (1.04, 16.7)	0.0441	
Predicted log₂ Week 4 CAB trough [†]	2.20 (1.04, 16.7)	0.0100	
BMI at baseline [‡]	Not significant		
Regimen: every-8-week/every-4-week	Not significant		
Integrase L74I: Yes/No [§]	Not significant		
Sex at birth: male/female	Not signifi	cant	
Other NNRTI RAMs: Yes/No [¶]	Not significant		
CAB RAMs: Yes/No	Not significant		
Other INSTI RAMs: Yes/No	Not significant		
Predicted log₂ Week 4 RPV trough [†]	Not signifi	cant	

^{*} After 44 weeks of LA therapy (excludes oral lead-in)

[†]After 4 weeks of LA therapy (excludes oral lead-in)

[‡] BMI was evaluated on a continuous scale

[§]Including mixtures except L74I/M

[¶]Other NNRTI RAMs were also retained in the final selected model, but not considered statistically significant (p=0.0667)

CVF = confirmed virologic failure; RPV = rilpivirine; RAMs = resistance associated mutations; BMI = body mass index; NNRTI = non-nucleoside reverse transcriptase inhibitor; CAB = cabotegravir; INSTI = integrase strand transfer inhibitor

BASELINE FACTOR ANALYSES

Week 48 Results

Of the 4 factors noted above to be associated with CVF through Week 48 in ATLAS, FLAIR, and ATLAS-2M, three (RPV RAMs, HIV-1 subtype A6/A1, and BMI \geq 30 kg/m²) may be present at baseline prior to initiation of CAB + RPV LA.¹ Of note, HIV-1 subtype A is primarily found in eastern Europe, central Asia, and east/central Africa.³

This analysis showed that there is an increased risk of CVF if 2 or more of these baseline factors are present. The rate of CVF among patients no or 1 factor present was <0.5%. See Table 6 below for details.

Table 6. Relationship Between the Presence of Baseline Factors and CVF or HIV-1 RNA <50 copies/mL at Week 48¹

	Proportion of Patients with CVF n/N (%)	Proportion of Patients with HIV-1 RNA <50 copies/mL n/N (%)	
None of the 3 baseline factors	3/732 (0.41)	694/732 (94.8)	
Any 1 of the baseline factors	1/272 (0.37)	261/272 (96)	
HIV-1 Subtype A6/A1 alone	1/95 (1.1)	90/95 (94.7)	
BMI ≥30 kg/m² alone	0/153 (0)	147/153 (96.1)	
RPV RAMs alone	0/24 (0)	24/24 (100)	
At least 2 of the baseline factors	9/35 (25.7)	25/35 (71.4)	
RPV RAMs + Subtype A6/A1	1/3 (33.3)	2/3 (66.7)	
RPV RAMS + BMI ≥30 kg/m²	3/10 (30)	7/10 (70)	
Subtype A6/A1 + BMI ≥30 kg/m²	4/21 (19)	16/21 (76.2)	
All 3 baseline factors	1/1 (100)	0/1 (0)	
Total, n/N (%)	13/1039 (1.25) [95% CI 0.67, 2.13]	980/1039 (94.3) [95% CI 92.74, 95.65]	

Beyond Week 48

The BFA presented above was updated with data through Week 124 from FLAIR, Week 96 from ATLAS, and Week 152 from ATLAS-2M. The pooled analysis includes 1651 patients with up to 3 years on study and 1431 with non-missing information on selected baseline factors. There were a total of 19 patients with CVF who were exposed to only every-4-week (n=11) or every-8-week (n=8) CAB + RPV LA. Data is only available for patients who received either every-4-week or every-8-week CAB + RPV LA, but not both.

The updated analysis confirmed the results of the BFA conducted at Week 48.² See Tables 7 and 8 below.

Table 7. Strength of Association of the Covariables and CVF through End of Study from FLAIR, ATLAS, and ATLAS-2M²

Covariable	Odds Ratio* (95% CI)	P value	
RPV RAMs: Yes/No	21.7 (5.80, 80.8)	<0.0001	
HIV-1 subtype A6/A1: Yes/No	12.9 (4.42, 37.5)	<0.0001	
BMI at baseline	1.09 (1.00, 1.19)	0.0447	
Regimen: every-8-week/every-4-week	Not signific	cant	
Integrase L74I: Yes/No	Not significant		
Sex at birth: male/female	Not significant		
Other NNRTI RAMs: Yes/No	Not signification	cant	
CAB RAMs: Yes/No	Not significant		
Other INSTI RAMs: Yes/No	Not signific	cant	

Table 8. Relationship Between the Presence of Baseline Factors and CVF or HIV-1 RNA <50 copies/mL through End of Study in FLAIR, ATLAS, and ATLAS-2M²

cabotegravir; NNRTI = non-nucleoside reverse transcriptase inhibitor; INSTI = integrase strand transfer inhibitor

Factors at Baseline	CVF, n (%)	HIV-1 RNA <50 copies/mL, n (%)
No baseline factors	4/970 (0.4)	844/970 (87)
Any 1 baseline factor	8/404 (2.0)	343/404 (85)
≥2 factors	11/57 (19)	44/57 (77)
Total, n/N (%)	23/1431 (1.6)	1231/1431 (86)
95% CI	1.0, 2.4	84.1, 88
CVF = confirmed virologic failure		

Some information contained in this response is outside the approved Prescribing Information. This product is not approved for the use described. This response is not intended to offer recommendations for administering this product in a manner inconsistent with its approved labeling.

In order for ViiV Healthcare to monitor the safety of our products, we encourage healthcare professionals to report adverse events or suspected overdoses to the company at 877–844–8872. Please consult the attached Prescribing Information.

This response was developed according to the principles of evidence-based medicine and, therefore, references may not be all-inclusive.

REFERENCES

- 1. Cutrell AG, Schapiro JM, Perno CF, et al. Exploring predictors of HIV-1 virologic failure to long-acting cabotegravir and rilpivirine: a multivariable analysis. *AIDS (London, England)*. 2021;35(9):1333-1342. doi:http://dx.doi.org/10.1097/QAD.000000000002883.
- 2. Orkin C, et al. Expanded multivariable models to assist patient selection for long-acting cabotegravir+rilpivirine treatment: clinical utility of a combination of patient, drug concentration, and viral factors associated with virologic failure over 152 weeks. Presented at HIV Glasgow 2022, October 23-26, 2022, Glasgow, UK and virtually. Oral Presentation.
- Hemelaar J, Elangovan R, Yun J, et al. Global and regional molecular epidemiology of HIV-1, 1990-2015: a systematic review, global survey, and trend analysis. *Lancet Infect Dis.* 2019;19(2):143-155.

doi:http://dx.doi.org/10.1016/S1473-3099(18)30647-9.