LICAVIR Study Synopsis and Table Shells

Background

Data from different studies on the risk of *de novo* hepatocellular carcinoma (HCC) in direct acting antiviral (DAA) treated patients are conflicting and inconsistent. While a potential mechanism may be immune modulation resulting from the rapid decrease of the hepatitis C virus (HCV) viral load ¹, several studies from large cohorts found no evidence of an increased risk of de novo HCC in DAA-treated patients. ²⁻⁵

Given the contemporaneous impact of liver inflammation and fibrosis, cirrhosis, duration of HCV infection, baseline patient demographics including patient age, HCV genotype, and prior treatment response associated with the development of HCC, the evaluation of the causal role of DAAs in the development of incident HCC in HCV infected patients is challenging.

The LICAVIR study was initiated by the ANRS (Agence Française de Recherches Sur Le VIH/Sida et Les Hépatites Virales)/INSERM (Institut National de la Santé et de la Recherche Médicale) and aims to study predisposing factors, clinical, biological, radiological and histological characteristics of HCC at the time of diagnosis from the data obtained from both CIRVIR and HEPATHER cohorts; HCC therapeutic procedures, prognosis and predictive factors of survival will also be analyzed.

AbbVie has engaged with the ANRS-INSERM group to obtain the results of the LICAVIR study and to present them to the Pharmacovigilance Risk Assessment Committee (PRAC) as the response to the following PRAC request: "The MAH should present the complete results of specific analyses from the HEPATHER and CIRVIR cohorts regarding the pattern of presentation of HCC cases (number, size, aggressiveness) in DAA treated and in untreated patients according to the indications presented in the PRAC Assessment Report and include them in the next PSURs."

Study objective

To characterize the de novo HCC cases that occurred with and without DAA treatment

Study methodology

A retrospective data collection from a cohort of patients with chronic HCV infection, who developed HCC up to the end of 2019.

A descriptive analysis of patients treated with any DAA who develop HCC after DAA initiation and patients who develop HCC without DAA treatment. No direct comparisons of treatment groups will be performed.

Table shells / data requests

Notes: for all tables, please provide summary statistics as appropriate. For categorical variables, please provide N (%); for continuous variables please provide N and mean (SD)

For variables that are only available in 1 cohort, please denote that in a footnote and add the N that contributed to that variable.

Patient characteristics should be measured over the period from data start through 1 day prior to HCC diagnosis. For characteristics with multiple measurements or assessments (i.e. labs, fibrosis stage, etc), report the value closest but prior to HCC diagnosis.

Table 1. Characteristics of treatments

| | Frequency | Achieved SVR12, N (%) | | Achieved SVR24, N (%) | | | Days since last treatment, Mean (SD) | |
|-----------------|-----------|--------------------------|----|--------------------------|-----|----|--|--|
| | N, % | Yes | No | Missing | Yes | No | Missing | |
| DAA | | | | | | | | |
| DAA only | | | | | | | | |
| DAA + IFN | | | | | | | | |
| DAA + ribavirin | | | | | | | | |
| DAA + IFN + | | | | | | | | |
| ribavirin | | | | | | | | |
| No DAA | | | | | | | | |
| treatment | | | | | | | | |
| INF | | | | | | | | |
| Ribavirin | | | | | | | | |
| No treatment | | | | | | | | |

Table 2. Frequency of DAA therapy in the study population*

| Treatment | N (%) |
|-------------------------------------|-------|
| DAA | |
| sofosbuvir/ledipasvir | |
| sofosbuvir/daclatasvir | |
| sofosbuvir/simeprevir | |
| elbasvir/grazoprevir | |
| sofosbuvir/velpatasvir | |
| sofosbuvir/velpatasvir/voxilaprevir | |
| glecaprevir/pibrentasvir | |
| dasabuvir, ombitasvir, | |
| paritaprevir, and ritonavir | |
| Other DAA | |

^{*}These groups are not mutually exclusive and may include combination of therapy and/or multiple rounds of therapy

Table 3. Patient characteristics at time of HCC diagnosis, with and without prior DAA treatment, in the LICAVIR study population

| Baseline characteristics | DAA treated (N) | DAA untreated (N) |
|----------------------------------|-----------------|-------------------------|
| Gender | | |
| Male, N (%) | | |
| Female, N (%) | | |
| Age, mean (SD) years | | |
| Ethnicity/Race | | |
| BMI, mean (SD) kg/m ² | | |
| Diabetes, N (%) | | |
| YES | | |
| NO | | |
| Missing | | |
| Dyslipidemia, N (%) | | |
| YES | | |
| NO | | |
| Missing | | |
| Past history of | | |
| cardiovascular events, | | |
| N (%) | | |
| YES | | |
| NO | | |
| Missing | | |
| Arterial Hypertension, | | |
| N (%) | | |
| YES | | |
| NO | | |
| Missing | | |
| HIV co-infection, N (%) | | |
| YES | | |
| NO | | |
| Missing | | |
| HBV co-infection, N (%) | | |
| YES | | |
| NO | | |
| Missing | | |
| Past excessive alcohol | | |
| intake, N (%) | | |
| YES | | |
| NO | | |
| Missing | | |

| Ongoing alcohol | |
|------------------------------|--|
| consumption, N (%) | |
| YES | |
| NO | |
| Missing | |
| Injection drug use, N (%) | |
| Past | |
| | |
| Ongoing Never | |
| | |
| Missing N (9/) | |
| Smoking, N (%) | |
| Past | |
| Ongoing | |
| Never | |
| Missing | |
| Past history of extrahepatic | |
| cancer, N (%) | |
| YES | |
| NO | |
| Missing | |
| Signs of Portal | |
| hypertension, N (%) | |
| Gastroesophageal varices, | |
| Thrombocytopenia | |
| LSE by TE ≥20-25kPa | |
| imaging showing collateral | |
| circulation | |
| missing | |
| Fibrosis stage, N (%) | |
| F0 | |
| F1 | |
| F2 | |
| F3 | |
| F4 | |
| missing | |
| In F4/cirrhotic patients, | |
| Child-Pugh-Turcotte | |
| Classification and MELD | |
| score, N (%) | |
| Child-Pugh A | |
| Child-Pugh B | |
| Child-Pugh C | |
| MELD score (mean, SD) | |
| History of prior hepatic | |
| decompensation, N (%) | |
| YES | |

| NO | |
|---|--|
| Missing | |
| Laboratory testing | |
| HCV genotype, N (%) | |
| Genotype 1 | |
| Genotype 2 | |
| Genotype 3 | |
| Genotype 4 | |
| Genotype 5 | |
| Genotype 6 | |
| Serum albumin g/dL, mean | |
| (SD) | |
| Platelet count 10 ³ /mm ³ , | |
| mean (SD) | |
| Prothrombin time%, mean | |
| (SD) | |
| INR (no units) , mean (SD) | |
| Total bilirubin mg/dL, mean | |
| (SD) | |
| Alpha-fetoprotein ng/mL, | |
| mean (SD) | |
| AST U/L, mean (SD) | |
| ALT U/L, mean (SD) | |
| Anti-HBc antibodies | |
| (HBsAg), N (%) | |
| YES | |
| NO | |
| Missing | |

Table 4: Characteristics of the HCC cases

| Characteristics | DAA treated | DAA untreated |
|------------------------------|-------------|---------------|
| Number of patients with HCC | | |
| Child-Pugh classification | | |
| A | | |
| В | | |
| С | | |
| Missing | | |
| Diagnostic method used, % | | |
| СТ | | |
| MRI | | |
| contrast-enhanced ultrasound | | |
| biopsy | | |
| missing | | |
| Number of nodules, % | | |
| 1 | | |

| 2-3 nodules | | |
|------------------------------------|---|--|
| >3 nodules | | |
| | | |
| missing | | |
| Diameter of largest nodule, mm, n | | |
| (%) | | |
| ≤ 20 | | |
| 21-30 | | |
| 31-50 | | |
| missing | | |
| Portal invasion, n (%) | | |
| Yes | | |
| No | | |
| missing | | |
| Within Milan criteria, n(%) | | |
| 1 nodule ≤ 50 mm | | |
| 2 or 3 nodules ≤ 30 mm | | |
| Outside Milan criteria | | |
| missing | | |
| Days between last imaging | | |
| examination and HCC diagnosis, | | |
| mean (SD) | | |
| Number of screening exams before | | |
| HCC diagnosis, mean (SD) | | |
| Extrahepatic spread, n (%) | | |
| YES | | |
| NO | | |
| Missing | | |
| Intervention for HCC treatment | | |
| Curative treatment, N (%) | | |
| Transplantation | | |
| Resection** | | |
| Ablation | | |
| Pallative treatment, N (%) | | |
| TACE | | |
| Systemic therapy, 1st line | | |
| Systemic therapy, 2nd line | | |
| Best supportive care | | |
| Missing | | |
| AFP levels at HCC diagnosis, mean | | |
| (SD) | | |
| Imaging examinations performed | | |
| during follow-up in HCC, N (%) | | |
| Survival in days/months, mean (SD) | | |
| Overall | | |
| Progression-free | | |
| Days until progression, Mean (SD) | | |
| | 1 | |

| Death, N (%) | |
|-------------------------------|--|
| YES | |
| NO | |
| Missing | |
| Hepatic decompensation, N (%) | |
| YES | |
| NO | |
| Missing | |

References

- 1. Villani R, Vendemiale G, Serviddio G. Molecular Mechanisms Involved in HCC Recurrence after Direct-Acting Antiviral Therapy. *Int J Mol Sci.* 2019; 20: 49.
- 2. Reig M, Mariño Z, Perelló C, Iñarrairaegui M, Ribeiro A, Lens S, et al. Unexpected high rate of early tumor recurrence in patients with HCV-related HCC undergoing interferon-free therapy. *J Hepatol.* 2016;65:719-26.
- 3. Ioannou GN, Green PK, Berry K. HCV eradication induced by direct-acting antiviral agents reduces the risk of hepatocellular carcinoma. *J Hepatol.* 2018;68:25-32.
- 4. Kanwal F, Kramer J, Asch SM, Chayanupatkul M, Cao Y, El-Serag HB. Risk of hepatocellular cancer in HCV patients treated with direct-acting antiviral agents. *Gastroenterology*. 2017;153:996-1005.e1.
- 5. Carrat F. First Prospective Evidence of Decreased Mortality after Direct Acting Antivirals in the French ANRS CO22 HEPATHER Cohort [Poster LB-28]. AASLD Meeting; 2017a; Washington, D.C.