

# Predictors of Undetectable Hepatitis Delta Virus RNA at 48 Weeks After End of Treatment With Bulevirtide Monotherapy in the MYR301 Study

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#### **Disclosures**

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#### **Background**

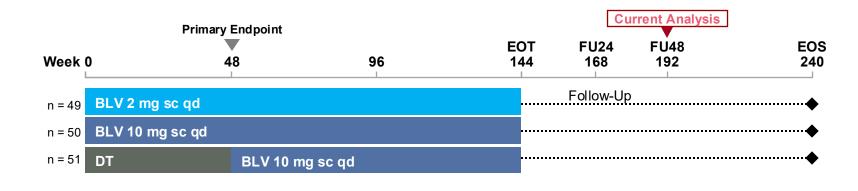
- Hepatitis delta virus (HDV) represents the most severe form of chronic viral hepatitis and is estimated to affect between 9 and 19 million people worldwide<sup>1</sup>
- Bulevirtide (BLV), an entry inhibitor of HDV, is approved in the European Union, Great Britain, Switzerland, the Russian Federation, and Australia (2 mg/day dose) for the treatment of chronic hepatitis delta in patients with compensated liver disease<sup>2,3</sup>
- Monotherapy with BLV 2 mg/day or 10 mg/day has been demonstrated to be effective and safe over 144 weeks of treatment<sup>4-6</sup>

# **Objectives**

- To evaluate predictors of undetectable HDV RNA after 96 or 144 weeks of BLV monotherapy
- To evaluate predictors for sustained HDV RNA undetectability throughout the patient's follow-up period (up to 48 weeks of posttreatment follow-up [FU48]) in those with undetectable HDV RNA at the scheduled end of treatment (EOT)

#### **MYR301 Methods**

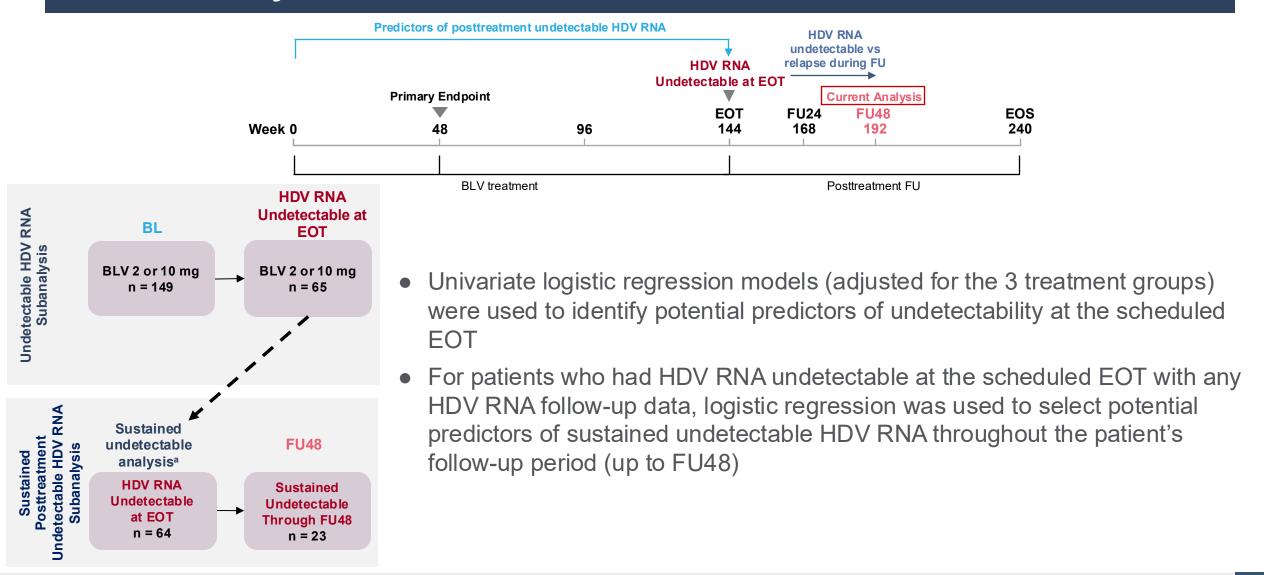
#### **Study Design**



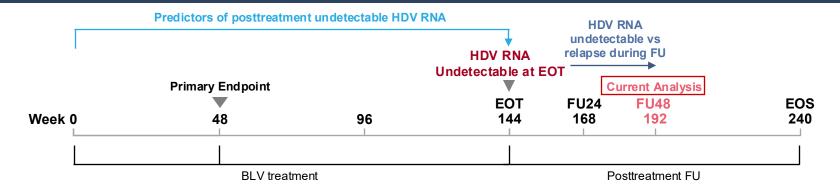
#### **Key Inclusion Criteria**

- CHD with detectable serum HDV RNA
- With or without cirrhosis; CTP score ≤7
- ALT >1 to <10 × ULN; platelets ≥60,000 cells/mm<sup>3</sup>
- MYR301 was a multicentre, open-label, randomised, Phase 3 study (NCT03852719) conducted in 4 countries (Germany, Italy, Russian Federation, and Sweden)
- HDV RNA levels were determined by reverse transcription—quantitative polymerase chain reaction using the RoboGene 2.0 (lower limit of quantitation, 50 IU/mL; lower limit of detection, 6 IU/mL)

## **Predictor Analysis Methods**



#### **Baseline and On-Treatment Characteristics Evaluated as Predictors**



#### Potential Predictors of Undetectable HDV RNA at EOT

#### Baseline<sup>a</sup> Clinical Characteristics

- Treatment (BLV 2 mg vs 10 mg)
- Age, sex, race, weight, BMI
- · Cirrhosis, liver stiffness (kPa)
- · ALT, platelets
- HDV RNA (log<sub>10</sub> IU/mL)
- Previous IFN therapy
- Concomitant HBV treatment, HBsAg, HBV DNA, HBV genotype
- Total bile salt levels (µmol/L)

# Potential Predictors of Sustained Posttreatment Undetectable HDV RNA in Patients Who Achieved Undetectability at EOT

#### **Baseline<sup>a</sup> Clinical Characteristics**

- Treatment (BLV 2 mg vs 10 mg)
- Age, sex, race, weight, BMI
- · Cirrhosis, liver stiffness (kPa)
- · ALT, platelets
- HDV RNA (log<sub>10</sub> IU/mL)
- Previous IFN therapy
- Concomitant HBV treatment, HBsAg, HBV DNA, HBV genotype
- Total bile salt levels (µmol/L)

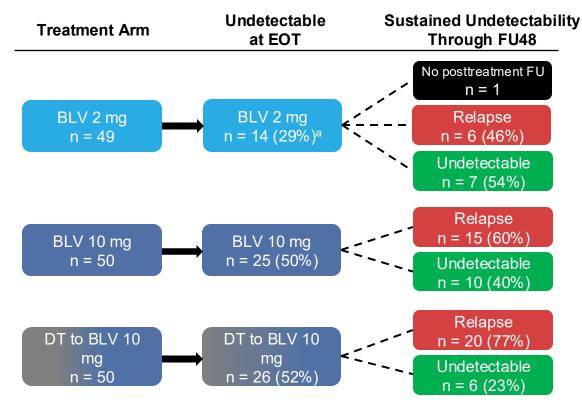
#### Treatment-Related Characteristics

- Duration of HDV RNA continuously undetectable at EOT
- Undetectable HDV RNA at early study timepoints

# **Baseline Demographics and Disease Characteristics**

|   | BLV 2 mg<br>(n = 49) | BLV 10 mg<br>(n = 50) | DT to BLV 10 mg<br>(n = 50) <sup>a</sup> |
|---|----------------------|-----------------------|--|
| Age, years, mean (SD)                       | 44 (9)               | 41 (9)                | 42 (8)                                   |
| Male sex, n (%)                             | 30 (61)              | 30 (60)               | 26 (52)                                  |
| Race, <sup>b</sup> n (%)                    |                      |                       |  |
| White                                       | 41 (84)              | 43 (86)               | 39 (78)                                  |
| Asian                                       | 8 (16)               | 6 (12)                | 11 (22)                                  |
| Cirrhosis present, n (%)                    | 23 (47)              | 24 (48)               | 24 (48)                                  |
| Liver stiffness, kPa, mean (SD)             | 14.0 (8.2)           | 14.8 (9.3)            | 16.1 (11.8)                              |
| ALT, U/L, mean (SD)                         | 108 (63)             | 123 (81)              | 82 (51)                                  |
| HDV RNA, log <sub>10</sub> IU/mL, mean (SD) | 5.10 (1.19)          | 4.96 (1.46)           | 5.03 (1.56)                              |
| Genotype HDV-1, <sup>c</sup> n (%)          | 49 (100)             | 48 (96)               | 50 (100)                                 |
| HBsAg, log <sub>10</sub> IU/mL, mean (SD)   | 3.67 (0.52)          | 3.61 (0.59)           | 3.71 (0.63)                              |
| HBV DNA, log <sub>10</sub> IU/mL, mean (SD) | 1.31 (1.28)          | 1.08 (1.26)           | 0.81 (1.01)                              |
| HBV genotype, n (%)                         |                      |                       |  |
| A   | 2 (4)                | 2 (4)                 | 2 (4)                                    |
| D   | 47 (96)              | 44 (88)               | 44 (88)                                  |
| Other <sup>d</sup> /missing                 | 0                    | 4 (8)                 | 4 (8)                                    |
| Previous IFN therapy, n (%)                 | 26 (53)              | 29 (58)               | 29 (58)                                  |
| Concomitant HBV NA treatment, e n (%)       | 32 (65)              | 27 (54)               | 33 (66)                                  |

## Rates of HDV RNA Undetectability Among Patients Who Received BLV



- Of 64 patients with available follow-up HDV RNA data, 62 had available data through FU48, while 2
  discontinued the study after 24 weeks of posttreatment follow-up (FU24) without experiencing relapse
- Rates of undetectable HDV RNA at EOT were higher in the treatment groups who received BLV 10 mg
- Relapses were more common in patients who received only 2 years of BLV treatment (vs 3 years)

#### **Baseline Predictors of Undetectable HDV RNA at EOT**

**Univariate Logistic Regression Analysis** 

| Baseline; n = 149                 | Undetectable at EOT Test (%) | Undetectable at EOT Reference (%) | Odds Ratio<br>(95% CI) | Favours Test       |                 |
|-----------------------------------|------------------------------|-----------------------------------|------------------------|--------------------|-----------------|
| Treatment arm                     | , ,                          | ,                                 |                        |                    |                 |
| BLV 10 mg (vs BLV 2 mg)           | 50.0%                        | 28.6%                             | 2.5 (1.1, 5.7)         | <b>├──</b>         | <i>P</i> = .03  |
| DT to BLV 10 mg (vs BLV 2 mg)     | 52.0%                        | 28.6%                             | 2.7 (1.2, 6.2)         | <b>├</b>           | <i>P</i> = .02  |
| Cirrhosis at randomisation        |                              |                                   |                        |                    |                 |
| Yes (vs no)                       | 46.5%                        | 41.0%                             | 1.3 (0.6, 2.4)         | <b>├</b>           |                 |
| Liver stiffness                   |                              |                                   |                        |                    |                 |
| ≥15.2 kPa (vs <15.2 kPa)          | 44.7%                        | 43.1%                             | 1.0 (0.5, 2.1)         | <del></del>        |                 |
| ALT                               |                              |                                   |                        |                    |                 |
| ≥1.5 × ULN (vs <1.5 × ULN)        | 44.1%                        | 42.1%                             | 1.2 (0.6, 2.6)         | <b>├</b>           |                 |
| Platelet count (10¹º/L)           |                              |                                   |                        |                    |                 |
| ≥160a (vs <160a)                  | 48.0%                        | 39.2%                             | 1.4 (0.7, 2.8)         | <b>⊢</b>           |                 |
| HDV RNA (log <sub>10</sub> IU/mL) |                              |                                   |                        |                    |                 |
| <5.2a (vs ≥5.2a)                  | 56.8%                        | 30.7%                             | 3.2 (1.6, 6.5)         | <b>├</b>           | <i>P</i> = .001 |
| Previous IFN therapy              |                              |                                   |                        |                    |                 |
| Yes (vs no)                       | 48.8%                        | 36.9%                             | 1.6 (0.8, 3.1)         | ++-                |                 |
| Concomitant HBV treatment         |                              |                                   |                        |                    |                 |
| Yes (vs no)                       | 46.7%                        | 38.6%                             | 1.5 (0.7, 3.0)         | <del>    •  </del> |                 |
| Total bile salts (µmol/L)         |                              |                                   |                        |                    |                 |
| <12.0ª (vs ≥12.0ª)                | 46.6%                        | 41.9%                             | 1.1 (0.6, 2.2)         |                    |                 |
| HBsAg (log <sub>10</sub> IU/mL)   |                              |                                   |                        |                    |                 |
| <3.8a (vs ≥3.8a)                  | 52.9%                        | 35.1%                             | 2.4 (1.2, 4.8)         | <b>├</b>           | <i>P</i> = .02  |
|                                   |                              |                                   | 0.1                    | 1 10               |                 |
|                                   |                              |                                   |                        | dds Ratio (95% CI) |                 |

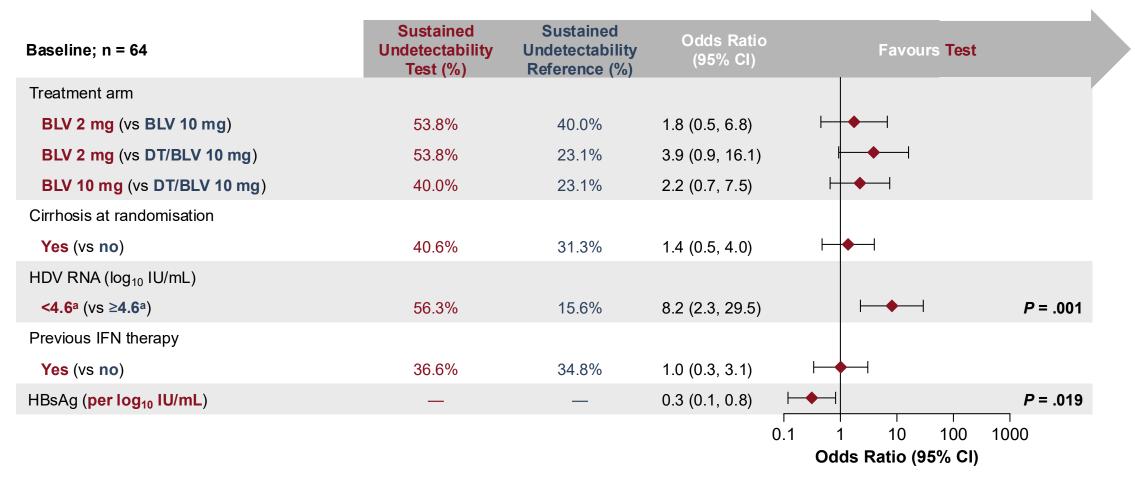
Treatment with BLV 10 mg and lower baseline HDV RNA were predictive of achieving undetectability at EOT

Bold **P-values** indicate significance.

<sup>&</sup>lt;sup>a</sup>Represents the median

## Baseline Predictors of Sustained HDV RNA Undetectability After EOT (FU48)

#### **Univariate Logistic Regression Analysis**



• In patients with undetectable HDV RNA at EOT, lower baseline HDV RNA and lower baseline HBsAg were predictive of sustained undetectability posttreatment

<sup>a</sup>Represents the median.

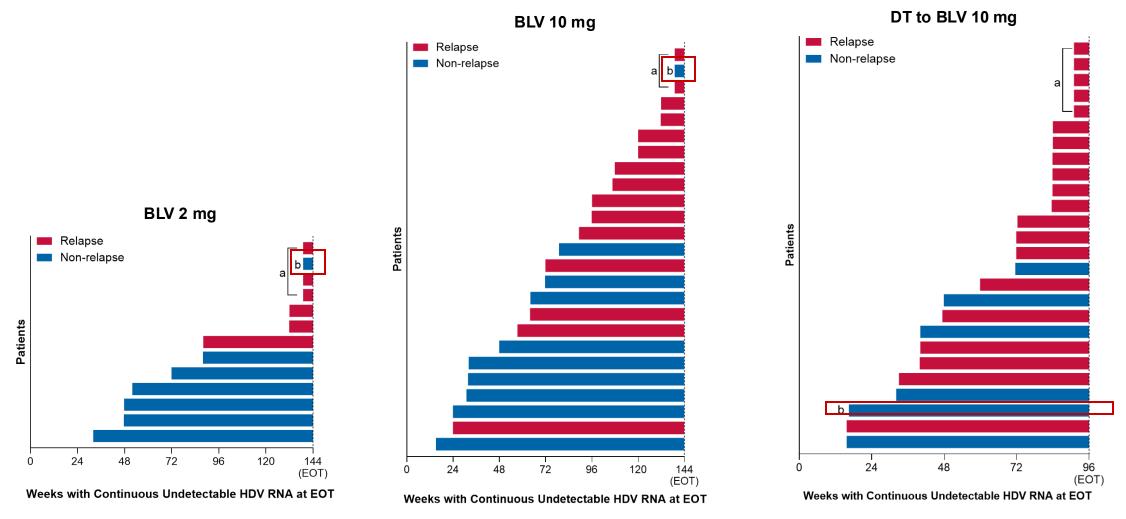
#### On-Treatment Predictors of Sustained HDV RNA Undetectability After EOT (FU48)

#### **Univariate Logistic Regression Analysis**

| On-Treatment; n = 64                     | Sustained<br>Undetectability<br>Test (%) | Sustained<br>Undetectability<br>Reference (%) | Odds Ratio<br>(95% CI) | Favours Test        |                  |
|--|--|---|------------------------|---------------------|------------------|
| Continuously undetectable HDV RNA at EOT |  |   |                        |                     |                  |
| ≥96 weeks (vs <96 weeks)                 | 90.0%                                    | 25.9%   | 26.6 (2.7, 262.4)      | <b>├</b>            | <i>P</i> <.005   |
| ≥48 weeks (vs <48 weeks)                 | 62.5%                                    | 9.4%  | 17.2 (4.0, 75.1)       | <b>├→</b>           | <i>P</i> = .0001 |
| Duration of continuous HDV RNA           |  |   |                        |                     |                  |
| undetectability at EOT (per 24 weeks)    | _  | _   | 2.9 (1.7, 4.8)         | <b>I</b> ◆I         | <i>P</i> <.0001  |
| W16 HDV RNA undetectable                 |  |   |                        |                     |                  |
| Yes (vs no)                              | 80.0%                                    | 32.2%   | 14.5 (1.3, 156.9)      | <b>├</b>            | <i>P</i> = .03   |
| W24 HDV RNA undetectable                 |  |   |                        |                     |                  |
| Yes (vs no)                              | 77.8%                                    | 29.1%   | 10.8 (1.8, 64.8)       | <b>├</b>            | P = .009         |
| W48 HDV RNA undetectable                 |  |   |                        |                     |                  |
| Yes (vs no)                              | 70.8%                                    | 15.0%   | 33.1 (5.9, 186.9)      | <b>├</b>            | <i>P</i> <.0001  |
| W72 HDV RNA undetectable                 |  |   |                        |                     |                  |
| Yes (vs no)                              | 57.1%                                    | 10.3%   | 20.3 (3.9, 107.0)      |                     | <b>P = .0004</b> |
|  |  |   |                        | Odds Ratio (95% CI) |                  |

• In patients with undetectable HDV RNA at EOT, early and longer duration of on-treatment undetectability was predictive of sustained undetectability posttreatment

# Weeks of Continuous On-Treatment Undetectable HDV RNA at EOT and Relapse Status by FU48



- Patients with longer duration of continuous undetectability at EOT were less likely to relapse
- Approximately 93% (38/41) of HDV RNA relapses occurred by the FU24 visit

# Biochemical Responses in Patients With Sustained Undetectable HDV RNA through FU48

| n = 23   | Baseline     | EOT         | FU48                    |  |
|--|--------------|-------------|-------------------------|--|
| ALT normalisation, n (%)                       | N/Aª         | 16 (70%)    | 15 (65%)b               |  |
| <b>ALT</b> , <sup>c</sup> U/L, median (Q1, Q3) | 93 (63, 136) | 32 (23, 44) | 30 (21, 46)<br>(n = 21) |  |

• Biochemical response was maintained in patients with sustained HDV RNA undetectability at FU48

#### **Conclusions**

- Rates of undetectable HDV RNA at EOT were higher in patients treated with BLV 10 mg vs 2 mg monotherapy for 2 to 3 years
  - In addition, rates of undetectable HDV RNA at EOT were associated with lower baseline viral load
- Among patients who had HDV RNA undetectable at the scheduled EOT, a subset of patients were able to sustain undetectable HDV RNA throughout their follow-up period (through FU48)
  - In these patients, biochemical responses were maintained during follow-up
- The majority of HDV RNA relapses (>90%) occurred by the FU24 visit
- The predictors for sustained HDV RNA undetectability through 48 weeks off-therapy were lower baseline HDV RNA levels (<4.6 log<sub>10</sub> IU/mL), lower baseline HBsAg levels, and most strongly, the duration of continuous on-treatment HDV RNA undetectability at EOT

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- Please see corresponding presentations:
  - Zoulim et al, OS-070; Viral Hepatitis B/D Therapy; 09/05/2025 (17:00–18:15)
  - Wedemeyer and Aleman et al, LBO-004; Late Breaker; 10/05/2025 (13:00–14:30)
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