The D-SOLVE Consortium





















Project coordinator/ leader

 Medizinische Hochschule Hannover/ Centre for Individualised Infection Medicine (CiiM), Germany, Prof. Heiner Wedemeyer

Partners

- Karolinska University Hospital, Sweden, Prof. Soo Aleman
- Institut national de la santé et de la recherche médicale (INSERM), France,
 Prof. Thomas Baumert/ Dr. Joachim Lupberger
- Karolinska Institutet, Sweden, Prof. Niklas Björkström
- National Institute for Infectious Diseases
 "Professor Doctor Matei Balş,", Romania,
 Prof. Florin Caruntu
- · Policlinico of Milan, Italy, Prof. Pietro Lampertico
- Helmholz-Zentrum für Infektionsforschung (HZI) /CiiM, Germany, Prof. Yang Li
- Helmholtz-Zentrum für Informationssicherheit (CISPA), Germany, Dr. Yang Zhang



The D-SOLVE Consortium

Understanding the individual host response against Hepatitis D Virus to develop a personalized approach for the management of Hepatitis D

Prof. Dr. Heiner Wedemeyer

Funded by the European Union, Horizon Europe Framework Programme (HORIZON), grant agreement No 101057917.





Images: stock.adobe.com

The pathogen in a nutshell: Hepatitis D virus

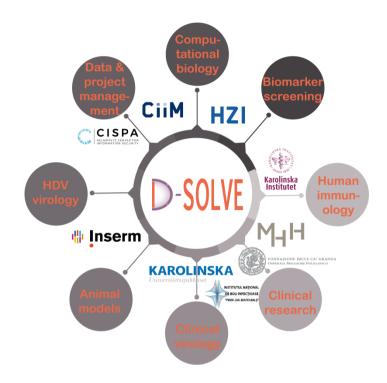
- Hepatitis D is by far the most severe form of chronic viral hepatitis, frequently leading to liver failure, hepatocellular carcinoma and death
- Hepatitis D is caused by coinfection of hepatitis B patients with the hepatitis D virus (HDV)
- There is a large interindividual variability in the course of hepatitis D
- → prototype infection for an individualized infection medicine approach



BUT hepatitis D is an orphan disease

- There is only limited knowledge on disease pathophysiology and host-virus interactions
- No multicenter cohorts of HDV infected patients are available
- urgent need to better understand individual factors determining the outcome of infection and to identify subjects benefiting from currently available treatments

The Consortium: Expertise



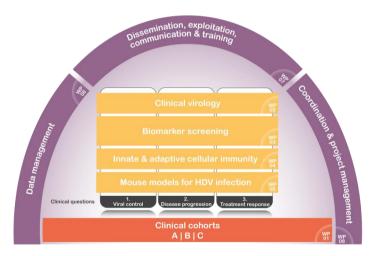
- Several of the leading experts in European HDV research contribute to D-SOLVE
- Clinical centers with an outstanding experience in translational research
- High-level knowledge in performing an unbiased and broad biomarker screening, from host genetics to host transcriptome and proteome analysis
- Mechanistic studies will be performed by renowned specialists in human immunology, HDV immunity and viral hepatitis research

The Aim

- Unbiased screening of a large multicenter cohort of well-defined HDV-infected patients
- Mechanistic studies to determine the functional role of distinct molecules
- Identification of specific parameters and prognostic markers

The Approach

- Key clinical questions on viral control, disease progression and treatment response will be studied across 4 research workpackes
- Similar methodological approaches will be applied for all clinical questions



Clinical cohort A: cross-sectional screening cohort of 750 HDV patients.

Clinical cohort B: retrospective-prospective group of patients with liver biopsies for immunological studies.

Clinical cohort C: prospective clinical trial where controlled stopping of bulevirtide is explored.