Health Data Science

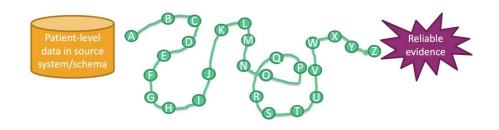
LARGE-SCALE PREDICTIVE MODEL DEVELOPMENT AND VALIDATION USING DATA STANDARDIZED TO THE OMOP COMMON DATA MODEL



Peter R. Rijnbeek
Professor of Medical Informatics
Chair Department of Medical Informatics
Erasmus MC, Rotterdam
The Netherlands
www.healthdatascience.nl



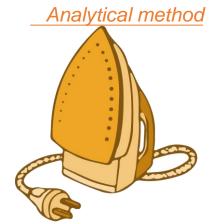
Generating Reliable Evidence at Scale



How can we generate reliable evidence at a large scale, i.e. on many data sources in Europe for many research questions?



The challenges of Real-World Data



The data...















Link to data

















What will it require?







Strong community



Data network

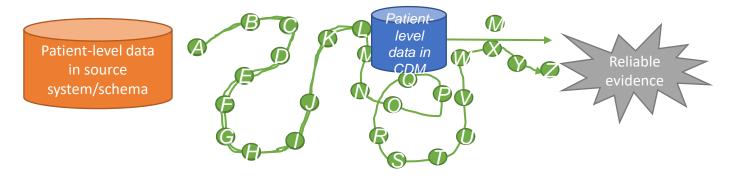


Standardised analytics



Standardization to a common data model





We develop analytical pipelines that can be utilized by data partners across the world.





www.ehden.eu



OHDSI COLLABORATORS

OHDSI COLLABORATORS

Map of Collaborators

The OHDSI community brings together volunteers from around the world to establish open community data standards, develop open-source software, conduct methodological research, and apply scientific best practices to both answer public health questions and generate reliable clinical evidence.

Our community is ALWAYS seeking new collaborators. Do you want to focus on data standards or methodological research? Are you passionate about open-source development or clinical applications? Do you have data that you want to be part of global network studies? Do you want to be part of a global community that truly values the benefits of open science? Add a dot to the map below and JOIN THE JOURNEY!

OHDSI By The Numbers

- · 2,367 collaborators
- 74 countries
- · 21 time zones
- · 6 continents
- 1 community

Large Active Community



















Oncology WG Structure

Observational Cancer Research

CDM & Standardized Vocabularies

Research

Vocab & Development

2nd & 4th Thursday 1-2 pm EST

Genomic

2nd & 4th Tuesday 9-10 am EST

Outreach & Research 2nd & 4th Tuesday 3-4 pm EST

www.ohdsi.org





Vision

The European Health Data & Evidence Network (EHDEN) aspires to be the trusted observational research ecosystem to enable better health decisions, outcomes and care

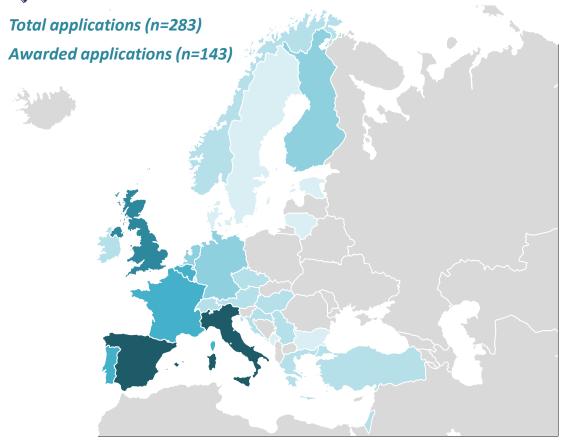
Mission

Our mission is to provide a new paradigm for the discovery and analysis of health data in Europe, by building a large-scale, federated network of data sources standardised to a common data model





EHDEN DATA NETWORK



Italy Spain Belgium Portugal France Germany Finland The Netherlands Switzerland Croatia Serbia Turkey Norway Hungary Greece Ireland Czech Republic Austria Israel Sweden Montenegro Luxembourg Lithuania Estonia Denmark Bulgaria 24

Geographic spread of data partners. The shade of blue indicates the # of data partners in that country (darker = more)









DARWIN EU® Coordination Centre





Search

Human regulatory ✓ Veterinary regulatory ✓

Initiation of DARWIN EU® Coordination Centre advances integration of real-world evidence into assessment of medicines in the EU Share

News 09/02/2022

EMA is initiating today the establishment of the Coordination Centre for the Data Analysis and Real World Interrogation Network (DARWIN EU®).

The role of the Coordination Centre is to develop and manage a network of real-world healthcare data sources across the EU and to conduct scientific studies requested by medicines regulators and, at a later stage, requested by other stakeholders.

The vision of DARWIN EU® is to give EMA and national competent authorities in EU Member States access to valid and trustworthy real-world evidence, for example on diseases, patient populations, and the use, safety and effectiveness of medicines, including vaccines, throughout the lifecycle of a medicinal product.

By supporting decision-making on the development, authorisation and surveillance of medicines, a wide range of stakeholders will benefit, from patients and healthcare professionals to health technology assessment bodies and the pharmaceutical industry. Additionally, DARWIN EU® will provide an invaluable resource to prepare for and respond to future healthcare crises and pandemics.

For example, the availability of timely and reliable real-world evidence can lead to innovative medicines becoming more quickly available to patients. Better evidence also supports more informed regulatory decisionmaking on the safe and effective use by patients of medicines on the market.

EMA will be working with Erasmus University Medical Center Rotterdam to establish the DARWIN EU® Coordination Centre. The contract was awarded to Erasmus University Medical Center Rotterdam following a call for tender for a service provider published in June 2021. The contractor will set up the necessary infrastructure of the Coordination Centre and establish the required business services. It will run scientific studies to answer research questions that come up during the evaluation of medicines in the EU and also maintain a catalogue of real-world data sources and metadata for use in medicine regulatory activities. All studies will be published in the EU catalogue of observational studies.







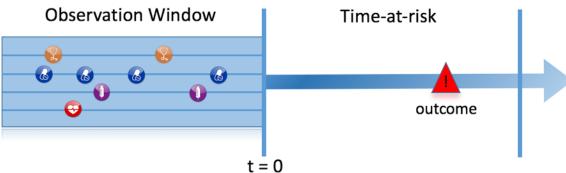






Problem definition





Among a target population (T), we aim to predict which patients at a defined moment in time (t=0) will experience some outcome (O) during a time-at-risk Prediction is done using only information about the patients in an observation window prior to that moment in time.



Model Development Pipeline



Dissemination of study results should follow the minimum requirements as stated in the Transparent Reporting of a multivariable prediction model for Individual Prognosis Or Diagnosis (TRIPOD) statement ¹. Annals of Internal Medicine RESEARCH AND REPORTING METHODS

- Sharing of model development details
- Discrimination and Calibration
- Internal and external validation
- Etc.

Transparent Reporting of a multivariable prediction model for Individual Prognosis Or Diagnosis (TRIPOD): Explanation and Elaboration

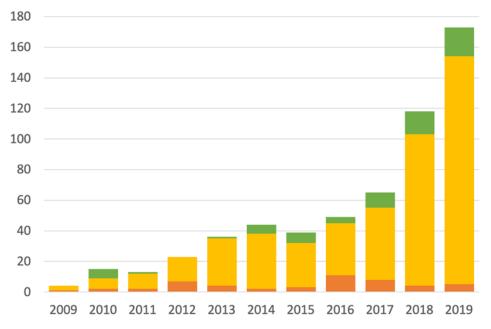
Karel G.M. Moons, PhD; Douglas G. Altman, DSc; Johannes B. Reitsma, MD, PhD; John P.A. Ioannidis, MD, DSc; Petra Macaskill, PhD; Ewout W. Steyerberg, PhD; Andrew J. Vickers, PhD; David F. Ransohoff, MD; and Gary S. Collins, PhD

The TRIPOD (Transparent Reporting of a multivariable prediction model for Individual Prognosis Or Diagnosis) Statement includes a 22-item checklist, which aims to improve the reporting of studies developing, validating, or updating a prediction model, whether for diagnostic or prognostic purposes. The TRIPOD Statement aims to improve the transparency of the reporting of a prediction model study regardless of the study methods used. This explanation and elaboration document describes the rationale; clarifies the meaning of each item; and discusses why transparent reporting is important, with a view to assessing risk of bias and clinical usefulness of the prediction model. Each checklist item of the TRIPOD Statement is explained in detail and accompanied by published examples of good reporting. The document also provides a valuable reference of issues to consider when designing, conducting, and analyzing prediction model studies. To aid the editorial process and help peer reviewers and ultimately readers and systematic reviewers of prediction model studies, it is recommended that authors include a completed checklist in their submission. The TRIPOD checklist can also be downloaded from www.tripod-statement.org.

Ann Intern Med. 2015;162:W1-W73. doi:10.7326/M14-0698 www.annals.org For author affiliations, see end of text. For members of the TRIPOD Group, see the Appendix



Current status of predictive modelling:



> J Am Med Inform Assoc. 2022 Apr 13;29(5):983-989. doi: 10.1093/jamia/ocac002.

Trends in the conduct and reporting of clinical prediction model development and validation: a systematic review

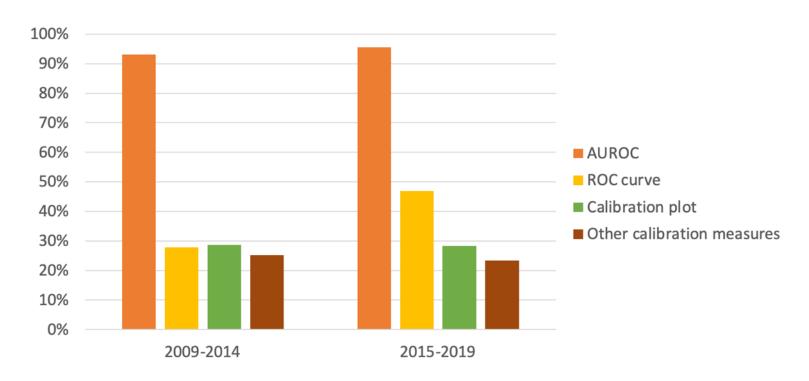
Cynthia Yang ¹, Jan A Kors ¹, Solomon Ioannou ¹, Luis H John ¹, Aniek F Markus ¹, Alexandros Rekkas ¹, Maria A J de Ridder ¹, Tom M Seinen ¹, Ross D Williams ¹, Peter R Rijnbeek ¹

- External validation
- Internal validation
- No validation

Review of 422 papers with 579 models



Current status of predictive modelling:





What is needed?

Full transparency and reproducibility

- 1. Standardised Health Data with respect to structure and terminology
- 2. Standardised Analytical Pipelines that enforce best modelling practices
- 3. Share models and allow extensive external validation across many databases
- 4. Disseminate <u>all</u> performance results



Our mission for Patient-Level Prediction

The Observational Health Data Sciences and Informatics (OHDSI) and the European Health Data and Evidence Network (EHDEN) developed a systematic process to learn and evaluate large-scale patient-level prediction models using observational health data in a large data network

Evidence Generation Evidence Evaluation Evidence Dissemination





www.ehden.eu



Patient-Level Prediction Framework



Design and implementation of a standardized framework to generate and evaluate patient-level prediction models using observational healthcare data 3

Jenna M Reps 🗷, Martijn J Schuemie, Marc A Suchard, Patrick B Ryan, Peter R Rijnbeek

Journal of the American Medical Informatics Association, Volume 25, Issue 8, August 2018, Pages 969–975, https://doi.org/10.1093/jamia/ocy032

Published: 27 April 2018 Article history ▼













Abstract

Objective

To develop a conceptual prediction model framework containing standardized steps and describe the corresponding open-source software developed to consistently implement the framework across computational environments and observational healthcare databases to enable model sharing and reproducibility.

R-package

www.github.com/OHDSI/PatientLevelPrediction

Book-of-OHDSI https://book.ohdsi.org/





EHDEN Academy https://academy.ehden.eu

Study Results

https://data.ohdsi.org



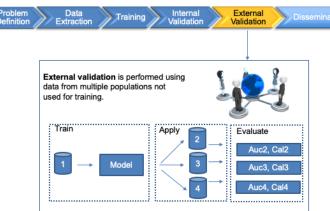
Prediction team in action





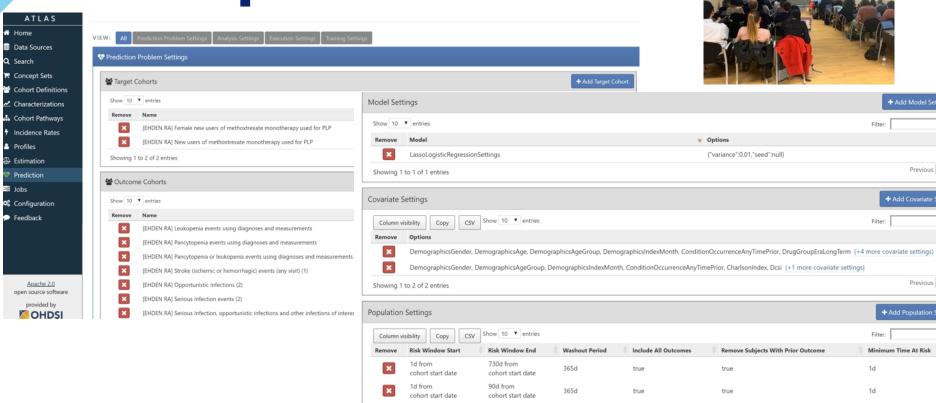








Model Specification



1826d from

cohort start date

cohort start date

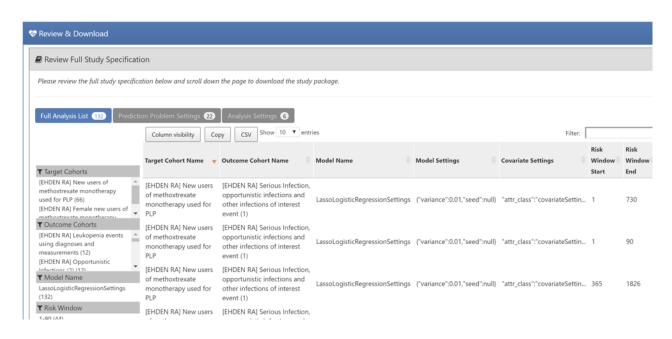
365d

true



true

Generate R-Package and share with the world







Share model performance





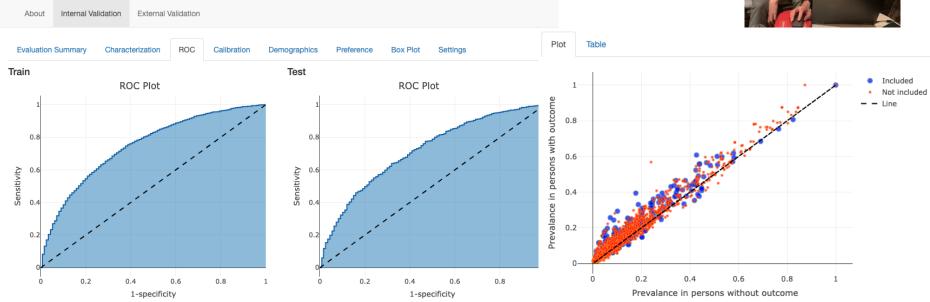


Figure 2. The Receiver Operating Characteristics (ROC) curve shows the ability of the model to discriminate between people with and without the outcome during the time at risk. It is a plot of sensitivity vs 1-specificity at every probability threshold. The higher the area under the ROC plot the higher the discriminative performance of the model. The diagonal refers to a model assigning a class at random (area under de ROC = 0.5).

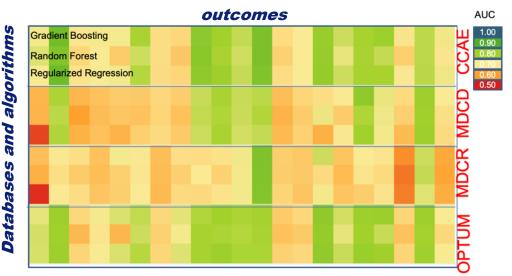
Figure 1. The variable scatter plot shows the mean covariate value for the people with the outcome against the mean covariate value for the people without the outcome. The meaning of the size and color of the dots depends on the settings on the left of the figure.

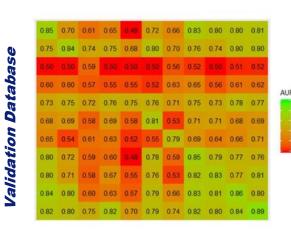
Erasmus MC 2 afms

Large scale validation and dissemination

The tool auto generates a word document containing all the model specifications, internal and external validation results, model details etc. etc. which serves as a kickstart for result dissemination.

Multiple interesting visualisation can be created:





Development Database



Seek COVER: COVID risk prediction

Objective: develop and externally validate **COV**ID-19 **E**stimated **R**isk scores that quantify a patient's risk of hospital admission, hospitalization requiring intensive services or fatality.

> BMC Med Res Methodol. 2022 Jan 30;22(1):35. doi: 10.1186/s12874-022-01505-z.

Seek COVER: using a disease proxy to rapidly develop and validate a personalized risk calculator for COVID-19 outcomes in an international network

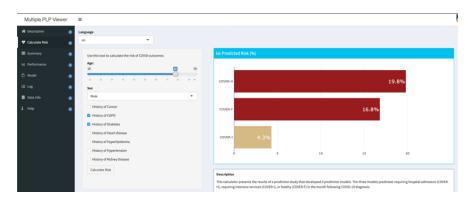
```
Ross D Williams # 1, Aniek F Markus # 1, Cynthia Yang 1, Talita Duarte-Salles 2, Scott L DuVall 3, Thomas Falconer 4, Jitendra Jonnagaddala 5, Chungsoo Kim 6, Yeunsook Rho 7, Andrew E Williams 8, Amanda Alberga Machado 9, Min Ho An 10, María Aragón 2, Carlos Areia 11, Edward Burn 2, 12, Young Hwa Choi 13, Iannis Drakos 14, Maria Tereza Fernandes Abrahão 15, Sergio Fernández-Bertolín 2, George Hripcsak 4, Benjamin Skov Kaas-Hansen 16, 17, Prasanna L Kandukuri 18, Jan A Kors 1, Kristin Kostka 19, Siaw-Teng Liaw 5, Kristine E Lynch 3, Gerardo Machnicki 20, Michael E Matheny 21, 22, Daniel Morales 23, Fredrik Nyberg 24, Rae Woong Park 25, Albert Prats-Uribe 12, Nicole Pratt 26, Gowtham Rao 27, Christian G Reich 19, Marcela Rivera 28, Tom Seinen 1, Azza Shoaibi 27, Matthew E Spotnitz 4, Ewout W Steyerberg 29, 30, Marc A Suchard 31, Seng Chan You 25, Lin Zhang 32, 33, Lili Zhou 18, Patrick B Ryan 27, Daniel Prieto-Alhambra 12, Jenna M Reps # 27, Peter R Rijnbeek # 34.

Affiliations + expand
PMID: 35094685 PMCID: PMC8801189 DOI: 10.1186/s12874-022-01505-z
Free PMC article
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Abstract

Background: We investigated whether we could use influenza data to develop prediction models for COVID-19 to increase the speed at which prediction models can reliably be developed and validated early in a pandemic. We developed COVID-19 Estimated Risk (COVER) scores that quantify a patient's risk of hospital admission with pneumonia (COVER-H), hospitalization with pneumonia requiring intensive services or death (COVER-I), or fatality (COVER-F) in the 30-days following COVID-19 diagnosis using historical data from patients with influenza or flu-like symptoms and tested this in COVID-19 patients.

- · 14 data sources from 6 countries
- Externally validated in 44,507 COVID cases from 5 data sources in South Korea, Spain, USA





Original Article | Published: 20 May 2022

Training prediction models for individual risk assessment of postoperative complications after surgery for colorectal cancer

V. Lin , A. Tsouchnika, E. Allakhverdiiev, A. W. Rosen, M. Gögenur, J. S. R. Clausen, K. B. Bräuner, J. S. Walbech, P. Rijnbeek, I. Drakos & I. Gögenur

Techniques in Coloproctology (2022) | Cite this article

108 Accesses | 1 Altmetric | Metrics

Abstract

Background

The occurrence of postoperative complications and anastomotic leakage are major drivers of mortality in the immediate phase after colorectal cancer surgery. We trained prediction models for calculating patients' individual risk of complications based only on preoperatively available data in a multidisciplinary team setting. Knowing prior to surgery the probability of developing a complication could aid in improving informed decision-making by surgeon and patient and individualize surgical treatment trajectories.

Methods

All patients over 18 years of age undergoing any resection for colorectal cancer between January 1, 2014 and December 31, 2019 from the nationwide Danish Colorectal Cancer Group



We need external validation at scale!

JMIR Med Inform. 2021 Apr; 9(4): e21547. Published online 2021 Apr 5. doi: 10.2196/21547 PMCID: PMC8023380 PMID: 33661754

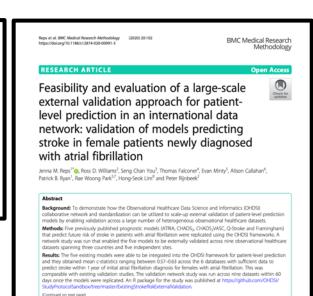
Implementation of the COVID-19 Vulnerability Index Across an International Network of Health Care Data Sets: Collaborative External Validation Study

Monitoring Editor: Christian Lovis

Reviewed by David Maslove, JianLi Wang, and Anoop Austin

Jenna M Reps. BSc, MSc, PhD. ³¹ Chungsoo Kim, PharmD, ² Ross D Williams, MSc, ³ Aniek F Markus, BSc, MSc, ³ Cynthis Yang, BSc, MSc, ³ Talita Duarte-Salles, MPH, PhD. ⁴ Thomas Falconer, BSc, MSc, ⁵ Litendra, Jonnagaddala, MIS, PhD, ⁶ Andrew Williams, PhD, ⁷ Sergio Fernandez-Bertolin, MSc, ⁴ Scott LD/viall, PhD, ⁶ Siristin Koatka, MPH, ⁹ Gowtham Rao, MD, PhD, ¹ Azza Shoaibi, PhD, ¹ Anna Ostropolets, MD, ⁵ Matthew E Spotnitz, MPH, MD, ⁵ Lin Zhang, PhD, ^{10,1} Paula Casajust, BSc, MSc, ¹Z Ewout W Steverberg, MSc, PhD, ^{13,14} Fredrik Nyberg, MPH, MD, PhD, ¹⁵ Seniamin Skov Kaas-Hansen, MSc, MD, ^{16,17} Young Hwa Choi, MD, PhD, ¹⁰ Daniel Morales, PhD, MBChB, ¹⁹ Seniamin Skov Kaas-Hansen, MSc, MD, ^{16,17} Young Hwa Choi, MD, PhD, ²⁰ Carlos Areis, MSc, PT, ²¹ Mishael E Matheny, MD, MPH, MS, ²² Kiristine E Lynch, PhD, ⁸ Maria Aragón, MSc, ⁴ Rae Woong Park, MD, PhD, ²³ George Hricosak, MD, MS, ⁵ Christian G Reich, MD, PhD, ⁹ Marc A Suchard, MD, PhD, ²⁴ Seng Chan You, MD, MS, ²³ Patrick B Ryan, PhD, ¹Daniel Prieto-Alhambra, MD, PhD, ²⁵ and Peter R Rijnbeek, PhD³

Join the network!



O Comment on this paper Using the OHDSI network to develop and externally validate a patient-level prediction model for Heart Failure in Type II Diabetes Mellitus O Ross D. Williams, Jenna M. Reps, Jan A Kors, Patrick B Ryan, Ewout Steyerberg, Katia M. Verhamme, doi: https://doi.org/10.1101/2021.04.06.21254966 This article is a preprint and has not been peer-reviewed [what does this mean?]. It reports new medical research that has yet to be evaluated and so should not be used to guide clinical practice. Preview PDF Abstract Introduction Heart Failure (HF) and Type 2 Diabetes Mellitus (T2DM) frequently coexist and exacerbate symptoms of each other. Treatments are available for T2DM that also provide beneficial treatment effects for HF. Guidelines recommend that patients with HF should be given Sodium-glucose co-transporter-2 inhibitors in preference to other second-line treatments for T2DM. Increasing personalization of treatment means that patients who have or are at risk of HF receive a customised treatment. We aimed to develop and externally validate prediction models to predict the 1-year risk of incident HF in T2DM patients starting second-line treatment.



Acknowledgement

This work would not have been possible without many contributors in the OHDSI community and the EHDEN project







www.ehden.eu



@IMI_EHDEN



IMI_EHDEN



github.com/EHDEN







This project has received funding from the Innovative Medicines Initiative 2 Joint Undertaking (JU) under grant agreement No 806968. The JU receives support from the European Union's Horizon 2020 research and innovation programme and EFPIA.

