

# Better predictions using big(ger) datasets

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## **Risk prediction**

- Risk prediction = foreseeing / foretelling
   ... (probability) of something that is yet unknown
- Turn available information (predictors) into a statement about the probability:

... of having a particular disease -> diagnosis

... of developing a particular event -> **prognosis** 





#### Why do we predict?

- Identification of high risk individuals
  - To inform patients and their families
  - To guide treatment decisions ("precision medicine")
  - To design randomized trials
- Data analysis
  - To deal with missing values
  - To match/subclassifiy patients
  - **–** ...



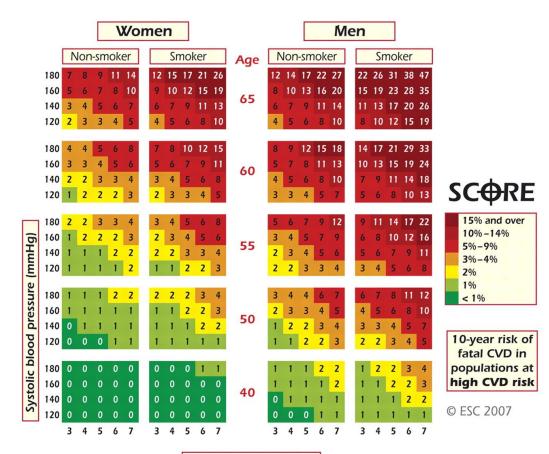


#### How do we predict?

- Combine information from multiple predictors
  - Patient characteristics (e.g. age, gender)
  - History and physical examination results (e.g. blood pressure)
  - Imaging results
  - (Bio)markers (e.g. coronary plaque)
- Develop a multivariable statistical model
  - Need for individual participant data (e.g. from cohort studies)
  - Many strategies available (e.g. logistic regression)

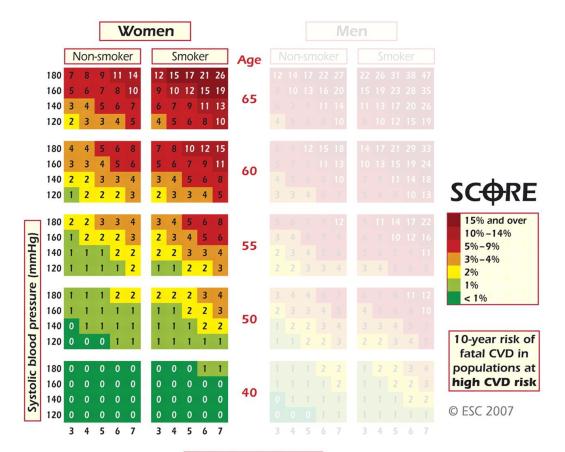






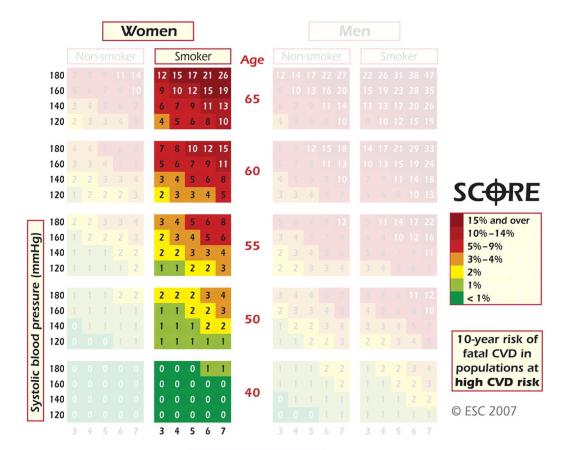






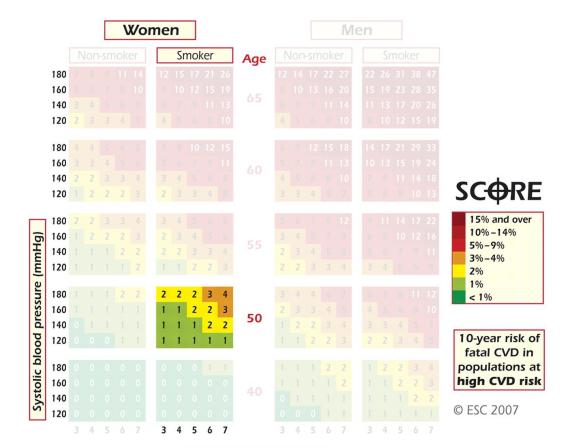






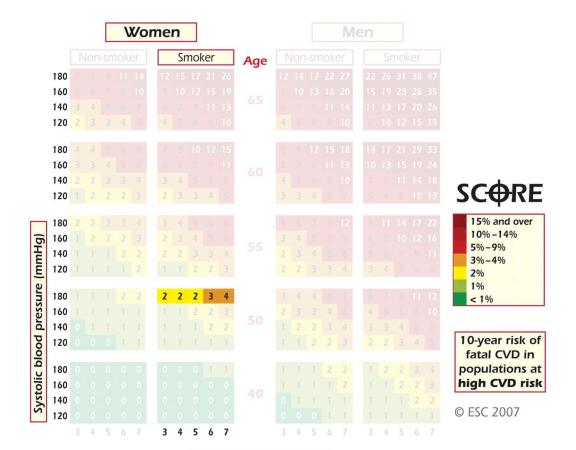






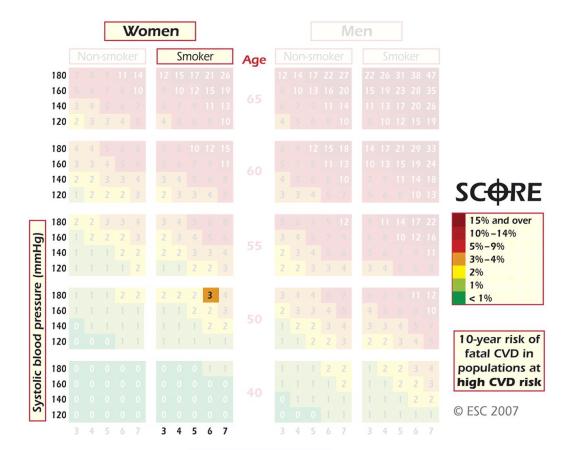














@TPA\_Debray



## **Watson for Oncology**

"Bring personalized, evidence-supported cancer care plans to your patients"



- Interpret cancer patients' clinical information
- Digest doctor's notes, medical studies, and clinical guidelines
- Provide individualized treatment recommendations
- Adopted by more than 150 hospitals and healthcare organizations





#### **Hype meets reality**

- Focus on US clinical practice and demographics
- Reliance on varies among hospitals
- Multiple examples of unsafe and incorrect treatment recommendations
- Lack of validation by independent scientists
- Lack of clinical trials to assess effectiveness



## Most models are not as good as we think

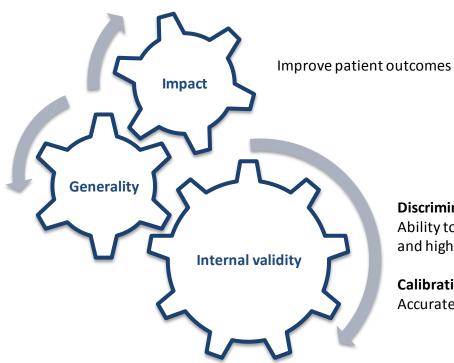


"All models are wrong, but some are useful"

George Box

## What is a "good" prediction model?

Good and consistent performance across different settings and populations



#### Discrimination

Ability to distiguish between low risk and high risk patients

#### Calibration

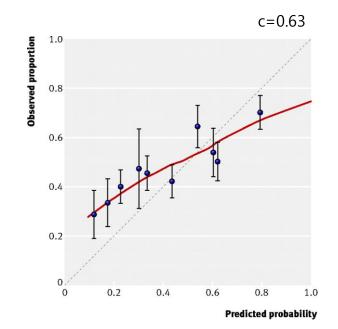
Accurate risk predictions



#### What is a "good" prediction model?

#### Common measures of prediction model performance

- Discrimination
  - Concordance (c-) statistic
  - Area under the ROC curve
- Calibration
  - Calibration intercept (calibration-in-the-large)
  - Calibration slope
  - Ratio of expected versus observed events





## Most models are not as good as we think

How to assess and improve the generalizability of prediction models?





## The rise of "big" data sets





## The rise of "big" data sets

Data increasingly available for thousands or even millions of patients from multiple practices, hospitals, or countries.

- Meta-analysis of individual participant data from multiple studies
  - Observational studies
  - Randomized controlled trials
- Analyses of databases and registry data containing e-health records









#### Validation of existing prediction models

- Assess prediction model performance multiple times
  - In new patients from the same (target) population
  - In new patients from different (but related) populations
  - To evaluate generalizability across different settings and populations
- Meta-analysis methods are needed
  - To summarize prediction model performance
  - To investigate sources of between-study heterogeneity





## Validation of QRISK 2

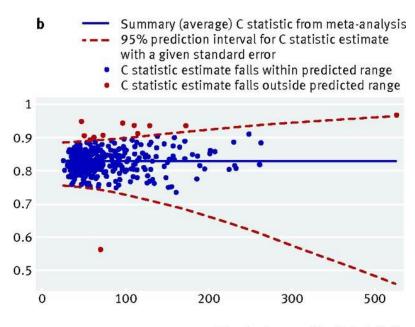
#### Registry data with 1.58 million patients from 364 practices

**Objective** To evaluate the performance of the QRISK2 score for predicting 10-year cardiovascular disease in an independent UK cohort of patients from general practice records and to compare it with the NICE version of the Framingham equation and QRISK1.

**Design** Prospective cohort study to validate a cardiovascular risk score.

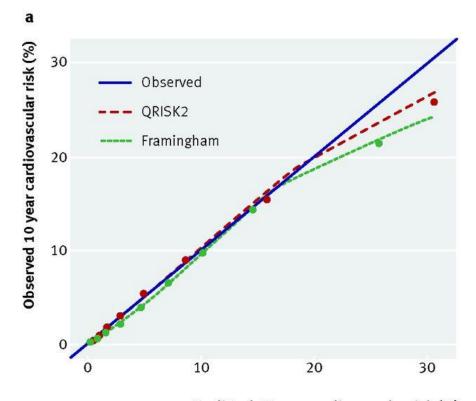
**Setting** 365 practices from United Kingdom contributing to The Health Improvement Network (THIN) database.

**Participants** 1.58 million patients registered with a general practice between 1 January 1993 and 20 June 2008, aged 35-74 years (9.4 million person years) with 71 465 cardiovascular events.



Standard error of logit C statistic

Summary (average) C statistic = 0.83 (95% CI 0.826 to 0.833) 95% prediction interval for true C statistic in a new practice = 0.76 to 0.88



Predicted 10 year cardiovascular risk (%)



### **Key references**

- Debray et al. A framework for meta-analysis of prediction model studies with binary and time-to-event outcomes. Stat Methods Med Res 2018
- Debray et al. A guide to systematic review and meta-analysis of prediction model performance. BMJ 2017
- Riley et al. External validation of clinical prediction models using big datasets from e-health records or IPD meta-analysis: opportunities and challenges.
   BMJ 2016
- Snell et al. Multivariate meta-analysis of individual participant data helped externally validate the performance and implementation of a prediction model. J Clin Epidemiol 2015









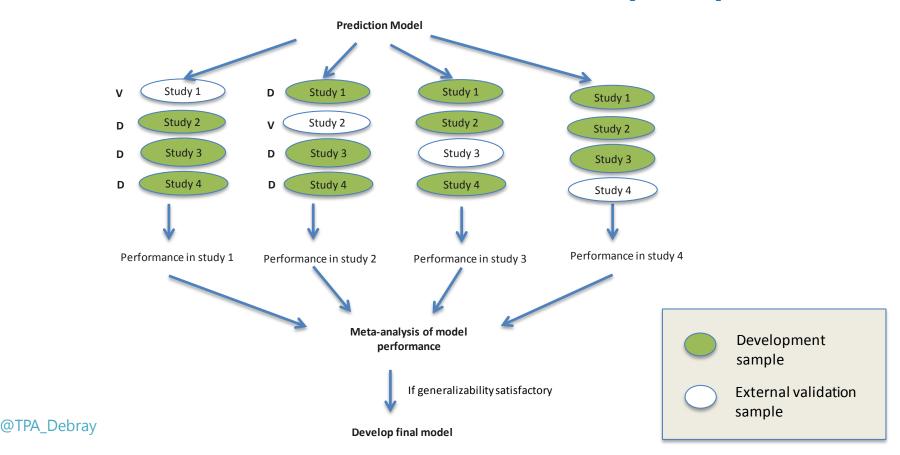
#### **Development of prediction models**

- Access to big(ger) datasets enables to assess model transportability (rather than merely reproduciblity) during its development
- Identify and account for heterogeneity in prediction model performance via <u>internal-external cross-validation</u>
- This allows to optimize prediction model generalizability

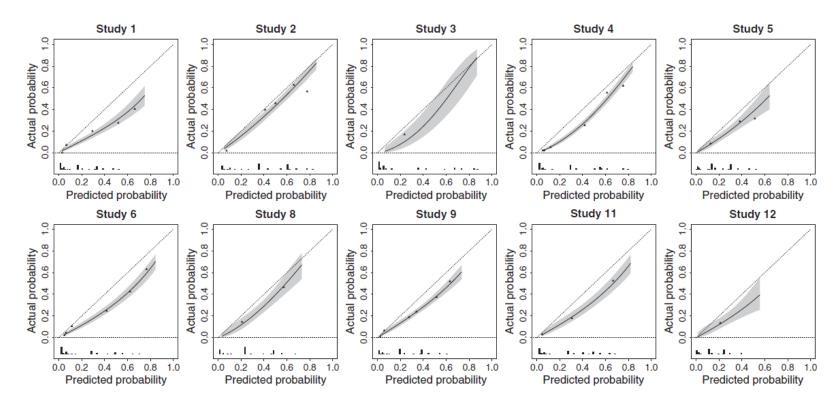




#### **Internal-external cross-validation (IECV)**



#### **IECV** allows for many external validations





#### **Development and validation of ENCALS**

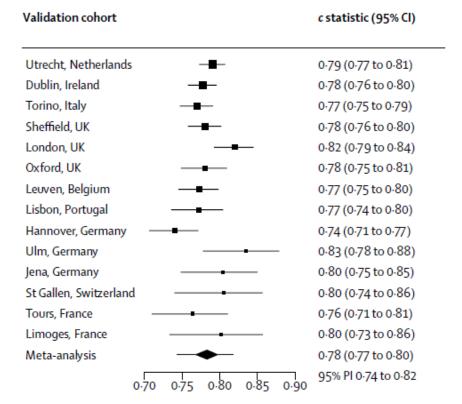
#### Prognosis for patients with amyotrophic lateral sclerosis (ALS)

- Cohort data from 11,475 patients from 14 European ALS centres
- Composite survival outcome (non-invasive ventilation for more than 23 h per day, tracheostomy, or death)
- Development of multivariable Royston-Parmar models
- Assessment of generalizability via IECV





#### **Development and validation of ENCALS**

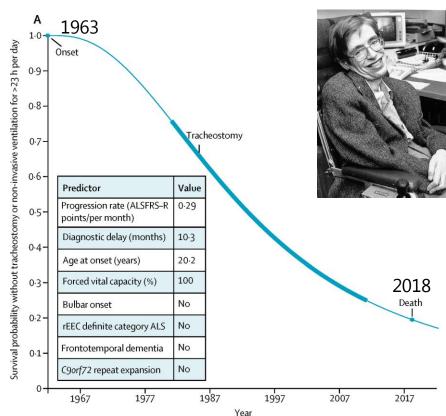


Pr(c>0.70)=100%



#### The life expectancy of Stephen Hawking

- Prediction in 1963: 2 year survival
- Prediction according to ENCALS:
   94% to survive at least 10 years
- Young age of onset was the most important factor for his long survival



#### **Key references**

- Ahmed et al. Developing and validating risk prediction models in an individual participant data meta-analysis. BMC Med Res Methodol 2014
- Debray et al. A framework for developing, implementing, and evaluating clinical prediction models in an individual participant data meta-analysis. Stat Med 2013
- Debray et al. Individual Participant Data (IPD) Meta- analyses of Diagnostic and Prognostic Modeling Studies: Guidance on Their Use. PLoS Med 2015





# **Future developments**



#### **Software**

metamisc: Diagnostic and Prognostic Meta-Analysis

Meta-analysis of diagnostic and prognostic modeling studies. Summarize estimates of prognostic factors, diagnostic test accuracy and prediction model performance. Validate, update and combine published prediction models. Develop new prediction models with data from multiple studies.

Version: 0.1.9

Depends: R (≥ 3.2.0), stats, graphics

Imports: <u>metafor</u> (≥ 2.0.0), <u>mvtnorm</u>, <u>ellipse</u>, <u>lme4</u>, <u>plyr</u>, <u>ggplot2</u>

Suggests: <u>runjags, rjags, testthat</u> (≥ 1.0.2)

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Author: Thomas Debray (5) [aut, cre], Valentijn de Jong [aut]

Maintainer: Thomas Debray <thomas.debray at gmail.com>

License: <u>GPL-3</u>

URL: <a href="http://r-forge.r-project.org/projects/metamisc/">http://r-forge.r-project.org/projects/metamisc/</a>

NeedsCompilation: no

In views: <u>MetaAnalysis</u>
CRAN checks: metamisc results

Downloads:

Reference manual: metamisc.pdf

Package source: metamisc 0.1.9.tar.gz

Windows binaries: r-devel: metamisc 0.1.9.zip, r-release: metamisc 0.1.9.zip, r-oldrel: metamisc 0.1.9.zip

OS X binaries: r-release: metamisc 0.1.9.tgz, r-oldrel: metamisc 0.1.8.tgz

Old sources: metamisc archive

Linking:

R

Please use the canonical form https://CRAN.R-project.org/package=metamisc to link to this page.

#### **Guidance and methods**

- Prognostic Research in Health Care: concepts, methods and impact editors: Richard Riley, Danielle Van der Windt, Peter Croft, Karel Moons
- Evidence synthesis using individual participant data: Concepts, Methods and Guidance for Clinical Research editors: Richard Riley, Jayne Tierney, Lesley Stewart
- Handbook of Meta-analysis
   editors: Christopher Schmid, Theo Stijnen, Ian White