# The slowly changing landscape of predictive modeling in biomedicine

Lara Lusa

Franziska Kappenberg, Willi Sauerbrei, Matthias Schmid and Jörg Rahnenführer

# STRATOS

No conflict of interest



#### Project of the STRATOS initiative on prediction modeling

- Acknowledgements:
  - Project initiated by Willi Sauerbrei, Matthias Schmid and Jörg Rahnenführer
  - Comments from Federico Ambrogi, Riccardo de Bin, Anne-Laure Boulesteix, Ben van Calster, Mitch Gail, Frank Harrell, Marianne Huebner
- Which are the important steps for the <u>development of useful</u> prediction models?
  - Diagnosis, prognosis and treatment selection
- Which are the advantages/weaknesses of machine learning and statistical models?
- In the last few years:
  - considerable interest in similar topics emerged from several groups, due to important changes in the prediction modeling landscape

Suggestions from the editorials

Data are more complex and more machine learning methods are used

"As clinical research catches up with other fields and finds itself immersed in the <u>era</u> <u>of big data</u>, the opportunity to apply <u>more computational and data-driven techniques</u> *increases.*" Goldstein et al., 2018, Health Informatics

• The existing best practice recommendations from the traditional biostatistics and medical statistics literature are no longer sufficient to guide the use of predictive models.

*"while <u>many best practice recommendations</u> for design, conduct, analysis, reporting, impact assessment, and clinical implementation <u>can be borrowed</u> from the traditional biostatistics and medical statistics literature, <u>they are not sufficient</u> <u>to guide the use of ML/AI in research</u>." Vollmer et al., 2019, Machine learning and artificial intelligence research for patient benefit: 20 critical questions on transparency, replicability, ethics, and effectiveness* 



#### How is predictive modeling changing?



Made with PubMed by Year: http://esperr.github.io/pubmed-by-year

The interest in predictive modeling in medicine is growing, and so is the use of machine learning methods

## The "systematic review collectors"



Maarten van Smeden @MaartenvSmeden

#### ...

#### Small update to the prediction modeling landscape

#### Traduci il Tweet

#### The prediction modelin



- 408 models for COPD prognosis (Bellou, 2019)
- 363 models for cardiovascular disease general population (Damen, 20
- · 327 models for toxicity prediction after radiotherapy (Takada 2022)
- 263 prognosis models in obstetrics (Kleinrouweler, 2016)
- 258 models mortality after general trauma (Munter, 2017)
- 232 models related to COVID-19 (Wynants, 2020)
- 160 female-specific models for cardiovascular disease (Baart, 2019)
- 142 models for mortality prediction in preterm infants (van Beek 2021)
- 119 models for critical care prognosis in LMIC (Haniffa, 2018)
- 101 models for primary gastric cancer prognosis (Feng, 2019)
- 99 models for neck pain (Wingbermühle, 2018)
- 81 models for sudden cardiac arrest (Carrick, 2020)
- 74 models for contrast-induced acute kidney injury (Allen, 2017)
- 73 models for 28/30 day hospital readmission (Zhou, 2016)
- 68 models for preeclampsia (De Kat, 2019)
- 68 models for living donor kidney/iver transplant counselling (Haller, 20
- 67 models for traumatic brain injury prognosis (Dijkland, 2019)
- 64 models for suicide / suicide attempt (Belsher, 2019)
- 61 models for dementia (Hou, 2019)
- 58 models for breast cancer prognosis (Phung, 2019)
- 52 models for pre-eclampsia (Townsend, 2019)
- 52 models for colorectal cancer risk (Usher-Smith, 2016)
- 48 models for incident hypertension (Sun, 2017)
- 46 models for melanoma (Kaiser, 2020)
- 46 models for prognosis after carotid revascularisation (Volkers, 2017)
- 43 models for mortality in critically ill (Keuning, 2019)

Gary Collins @ @GSCollins · 22 mar 2022 I've also recently been updating my list @MaartenvSmeden (for a talk). Below are ~260 systematic reviews of clinical prediction models - you might've missed a couple ? - it's incomplete, and probably has some inaccuracies (still updating it).

Number of reviewed models	Number of systematic reviews since 2020
>100	7
50-100	13
25-49	22
10-24	33
<10	7





#### Selection of reviews



887 papers and 1448 models (after excluding pure validations and studies published before 2005)

#### Selected reviews



Review	Populati on	Index model	Outcome	Year of publication of the papers	Mod els	Pap ers
				included in the review		
Wynants et al.	Patient with confirmed COVID-19	All available prognostic models	All outcomes	2020 to 2022	501	368
Li et al.	Patients with vascular conditions	Predictive models that use ML methods	All outcomes (included segmentation)	1991 to 2021	202+	202
Sun et al.	Patient with heart failure	All available prognostic models <u>from 2011</u>	All-cause mortality or all- cause readmission of HF patients	2011 to 2021	176	78
Ndjaboue et al.	People with pre- diabetes and any type of diabetes, except gestational diabetes	All available models for which there was reported internal and/or external validation	Diabetes-related health conditions (complications)	2000 to 2020	175	75
Ogink et al	Surgical orthopaedic population	Prognostic models from studies that included <u>at least one</u> <u>ML-based prediction</u>	Orthopaedic surgical outcomes	1996 to 2020	218	56
He et al.	Patients diagnosed with cervical cancer	All available models	Clinical outcome (recurrence, metastasis, death, etc.)	1987 to 2020	74	52
Haller et al.	Recipients or donors in living organ transplantation	All available models	Any outcome occurring after transplantation donation in the recipient or donor	2004 to 2021	48	35
Gade et al.	Community-dwelling older adults (60+) of the general	All available models	Falls	1994 to 2019	54	21 7

#### Completeness of reviews: Information available at paper level





Some information is missing systematically in some reviews

Number of events and of variables can be difficult to retrieve

**EPV** is problematic



#### The reviews are heterogenous



#### Main findings – time trends (without COVID-19)

**IR**M



From 2005, graphs are mostly on log-2 scale on the y-axes (positively asymmetric distributions)



## Zooming in – trends only

Data are bigger, more variables are used and more models are fitted BUT the changes are smaller than might have been expected





## **Missing values**



#### 3 reviews ignore the information

#### Method

Unclear / No information Other No Need To Report / None Variable omission Indicator methods / Dummy Other imputation Multiple imputation Single imputation Complete Case

Information about missing data is still rarely reported in papers (gray, 54% of papers with no information)

Imputation methods (blue/red) are becoming more common Complete case analysis is still the most common choice

The quality of reporting did not improve substantially in time



## Internal validation









#### Vast heterogeneity across reviews!

Internal validation is becoming more

## Internal validation – if performed, which?



Vast heterogeneity across reviews!

Cross-validation is gaining popularity, split-based methods are common only in some reviews



#### **External validation**



External validation is still uncommon (might be performed in subsequent papers) More common in the review from Njdaboue, due to inclusion criteria



16

#### Time trends in reporting

Year	Internal val.	External val.	Discrimination	Classification	Calibration
2005/09	21/46 (46%)	8/46 (17%)	20/46 (43%)	20/25 (80%)	12/46 (26%)
2010/14	82/148 (55%)	27/141 (19%)	98/148 (66%)	47/72 (65%)	47/141 (33%)
2015/19	142/200 (71%)	26/167 (16%)	149/200 (74%)	62/100 (62%)	50/167 (30%)
2020/21	82/125 (66%)	14/109 (13%)	87/125 (70%)	42/82 (51%)	30/109 (28%)
COVID-19	248/368 (67%)	95/368 (26%)	209/368 (57%)	114/368 (31%)	82/368 (22%)

Increase of reporting of internal validation and discrimination measures, poor reporting of calibration

## Type of model: ML vs statistical methods



Li (using ML models) and Ogrink (at least one ML model) selected models based on the use of ML – but many of their models were classified (by us) as statistical

Time changes are not visible within reviews (the overall summary are influenced <sub>17</sub> by weight of each review)

## Comparison of statistical and ML models

	n	Ме	Mean	Range	IQR	
Number of patients						
Statistical	381	421	11,891	4 to 1,621,149	160 to 1475	
ML	260	347	19,753	8 to 1,567,636	130 to 1071	
Number of events						
Statistical	292	84	591	7 to 28,140	41 to 288	
ML	166	95	689	10 to 46,163	48 to 214	
Number of candidate variables						
Statistical	225	23	33	1 to 1224	14 to 37	
ML	68	33	289	7 to 15,000	24 to 49	
Number of selected variables						
Statistical	370	6	12	1 to 488	4 to 10	
ML	113	6	19	2 to 618	3 to 13	
Number of models						
Statistical	400	1	2	1 to 10	1 to 2	
ML	157	1	1	1 to 8	1 to 1	

The size of the datasets was similar (more extremes for ML methods)

Many more candidate variables for ML, but similar number of selected



# Summary statistics of predictive performance measures by type of model

	Discrimination	Calibration	Classification
Statistical	334/449 (74%)	171/441(39%)	96/255(38%)
ML	124/277 (45%)	24/261(9%)	129/262(49%)
Both	80/99 (81%)	13/69(19%)	41/78(53%)
Unclear	25/62 (40%)	13/60(22%)	19/52(37%)

Very poor reporting of calibration of ML models, poor for discrimination.



#### Conclusions

- Quantitative assessment of changes is important, but it is not straightforward
- Reviews
  - Not many that include many prediction models and have (complete relevant) publicly available data
  - wide heterogeneity across reviews in almost all the aspects
    - Example: COVID/Wynants vs the other 2020/21 papers
    - Truly reflecting differences in the fields or somehow related to the review process?
- The changes in predictive modeling are not as substantial as it might have been anticipated
  - Growth in the 2015-19 period in number of patients and variables, followed by a stabilization, the centers of the distributions are stable, extreme values are more common



## Changes in predictive modeling

- Larger sample sizes (and number of events) and larger mean number of selected variables, more models per paper, but
  - the central part of the distributions are mostly stable in the 2020/21 period
  - similar median number of variables (candidate and selected)
    - Underreported pre-selection?
    - Under-representation of imaging models in our data?
    - Time lag makes the growth not yet observable?
    - Simpler models are still preferred (even if more candidates are available)?
- There might be a trend towards increasingly following guidelines
  - performing/reporting internal validation,
  - using resampling methods instead of sample splitting
  - reporting discrimination
  - more imputation methods for missing values (but still poor reporting for missing data)
- Not a very clear increase of the use of ML methods (with)in reviews and not striking differences in the characteristics of data being used



Beyond our draft...

- ... and back to the original idea of the project
- ML vs statistical models
  - The distinction is difficult. Should we focus on model complexity?
  - Comparison of the performance -> reliable?
- Assessment of bias
  - Larger for ML/complex models
  - how reliable is the assessment?
- Need for new guidelines?
  - All the basic principles apply and are still not always used
  - <u>Comprehensible understanding of methods needed to identify</u> <u>specificities related to "new models"</u>





## Selection of reviews

- Type of prognostic models
  - Multivariate prognostic models (more than 2 candidate predictors)
  - Developmental studies
- Type of review
  - Published in 2020 or after
  - Includes development models (not only validation)
  - Per paper/per model data are available (table format) for most of the information suggested in the CHARMS checklist
  - Includes at least 30 models/papers
  - Source: the lists of the "systematic review collectors" and additional PubMed search
- 8 selected reviews, including 841 papers and 1499 models







