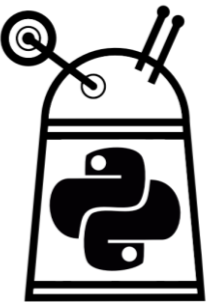
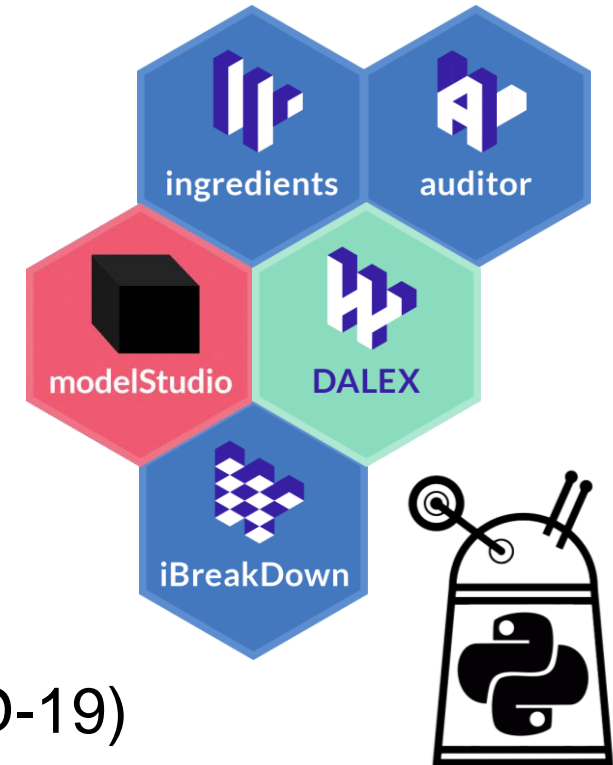
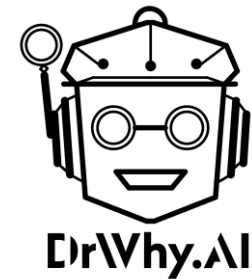


# XAI to support prediction making in COVID-19 pandemic

Hubert Baniecki, Warsaw, 10.2020

# Hi!

- Data Science student at Warsaw University of Technology
- **Research Software Engineer**  
**at MI2 DataLab lead by Przemyslaw Biecek**
- interested in XAI and model-human interaction
- creating & maintaining the DrWhy.AI universe
- **member of DeCoviD project** (grant: IDUB against COVID-19)
- **MI2 DataLab in MOCOS group** (MOdelling COronavirus Spread)



# 1. Prediction Making in COVID-19

literature review & critique

# 2. Real case study

R & Python XAI tools

# 3. MI2 DataLab in MOCOS

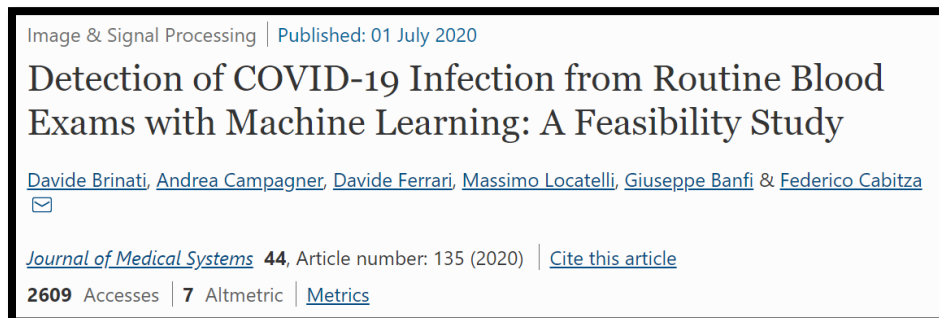
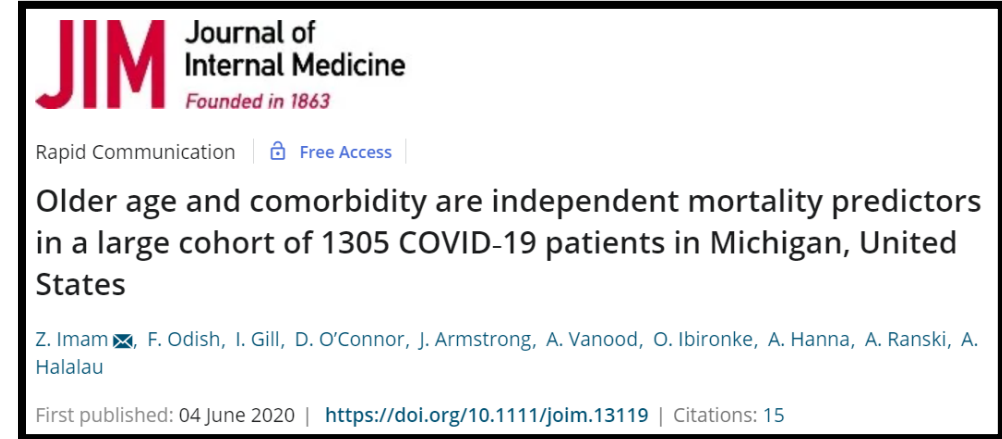
COVID-19 analysis for Poland

# 1. Prediction Making in COVID-19

literature review & critique

# Literature

- diagnosis of COVID-19
- prognosis of patient mortality risk
- prediction of illness severity, hospitalization risk, length of hospital stay
- tabular data (e.g. age, blood test) & image data (medical lung images)
- statistical models / ML models / DL models



# Review

107 studies

91 diagnostic models  
2/3 on medical images

50 prognostic models  
1/2 for mortality  
1/6 for illness severity

mostly preprints

**thebmj** covid-19 Research ▾ Education ▾ News & Views ▾ Campaigns ▾ Jobs ▾

Research

**Prediction models for diagnosis and prognosis of covid-19: systematic review and critical appraisal**

BMJ 2020 ; 369 doi: <https://doi.org/10.1136/bmj.m1328> (Published 07 April 2020)  
Cite this as: BMJ 2020;369:m1328

Article Related content Metrics Responses Peer review

Laure Wynants , assistant professor<sup>1 2</sup>, Ben Van Calster , associate professor<sup>2 3</sup>, Gary S Collins , professor<sup>4 5</sup>, Richard D Riley, professor<sup>6</sup>, Georg Heinze , associate professor<sup>7</sup>, Ewoud Schuit , assistant professor<sup>8 9</sup>, Marc M J Bonten, professor<sup>8 10</sup>, Darren L Dahly, principal statistician<sup>11 12</sup>, Johanna A A Damen, assistant professor<sup>8 9</sup>, Thomas P A Debray, assistant professor<sup>8 9</sup>, Valentijn M T de Jong, assistant professor<sup>8 9</sup>, Maarten De Vos, associate professor<sup>2 13</sup>, Paula Dhiman, research fellow<sup>4 5</sup>, Maria C Haller, medical doctor<sup>7 14</sup>, Michael O Harhay, assistant professor<sup>15 16</sup>, Liesbet Henckaerts, assistant professor<sup>17 18</sup>, Pauline Heus, doctoral candidate<sup>8 9</sup>, Nina Kreuzberger, research associate<sup>19</sup>, Anna Lohmann, researcher in training<sup>20</sup>, Kim Luijken, doctoral candidate<sup>20</sup>, Jie Ma, medical statistician<sup>5</sup>, Glen P Martin, lecturer<sup>21</sup>, Constanza L Andaur Navarro, doctoral student<sup>8 9</sup>, Johannes B Reitsma, associate professor<sup>8 9</sup>, Jamie C Sergeant, senior lecturer<sup>22 23</sup>, Chunhu Shi, research associate<sup>24</sup>, Nicole Skoetz, medical doctor<sup>19</sup>, Luc J M Smits, professor<sup>1</sup>, Kym I E Snell, lecturer<sup>6</sup>, Matthew Sperrin, senior lecturer<sup>25</sup>, René Spijker, information specialist<sup>8 9 26</sup>, Ewout W Steyerberg, professor<sup>3</sup>, Toshihiko Takada, assistant professor<sup>8</sup>, Ioanna Tzoulaki, assistant professor<sup>27 28</sup>, Sander M J van Kuijk, research fellow<sup>29</sup>, Florian S van Royen, research fellow<sup>8</sup>, Jan Y Verbakel, assistant professor<sup>30 31</sup>, Christine Wallisch, research fellow<sup>7 32 33</sup>, Jack Wilkinson, research fellow<sup>22</sup>, Robert Wolff, medical doctor<sup>34</sup>, Lotty Hooft, associate professor<sup>8 9</sup>, Karel G M Moons, professor<sup>8 9</sup>, Maarten van Smeden, assistant professor<sup>8</sup>

Author affiliations ▾

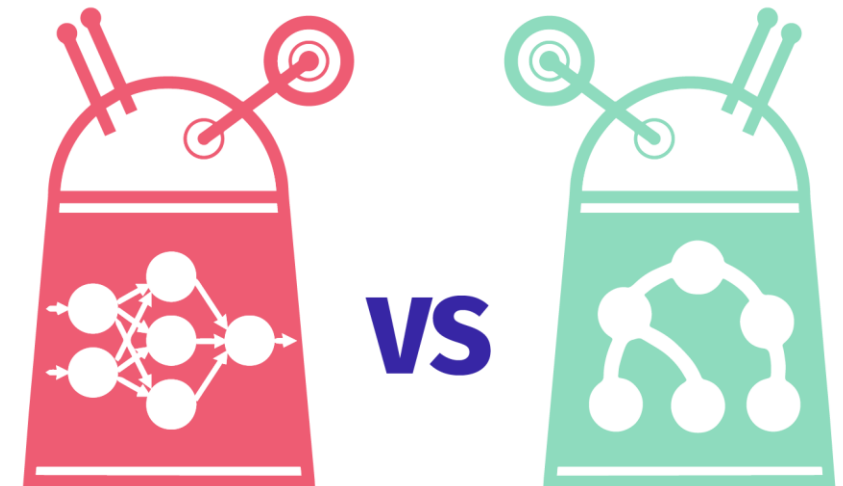
Correspondence to: L Wynants [laure.wynants@maastrichtuniversity.nl](mailto:laure.wynants@maastrichtuniversity.nl)

Accepted 31 March 2020  
Final version accepted 1 July 2020

**+ continuously updated up to 2 years**

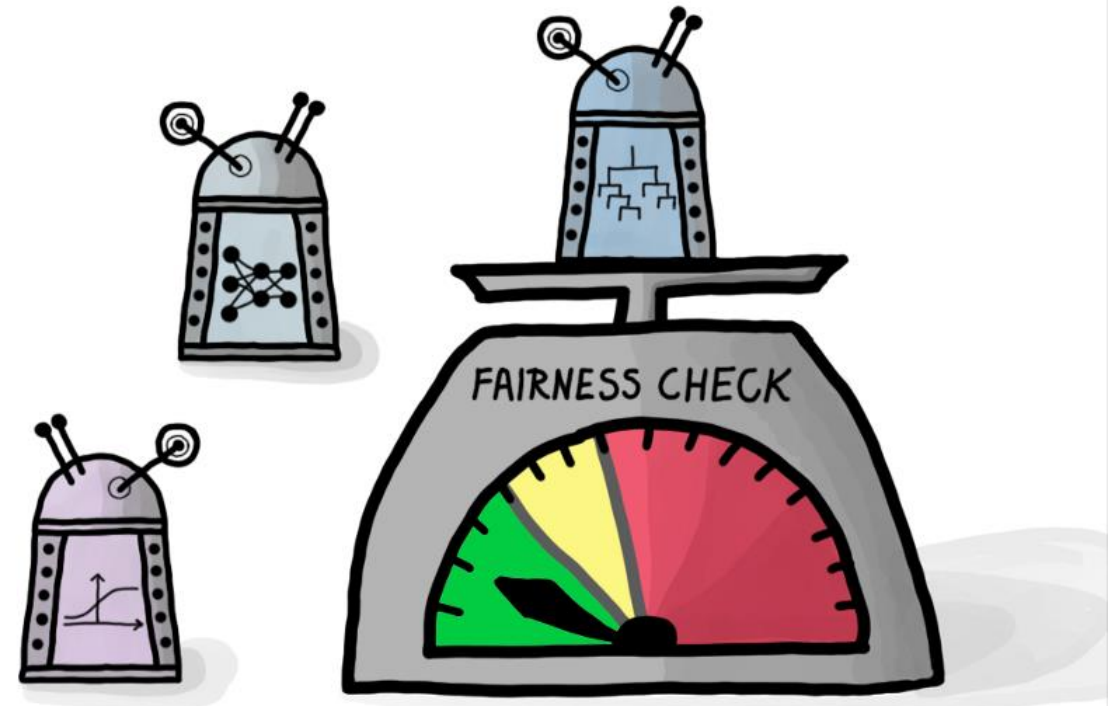
# Critique

- all models biased and/or overfitted
- insufficiently validated
- only 1/3 with code!
- limited, poorly diversified data
- do not use these models in practice
- future: better validation for prediction making



# Responsible AI, ML, Be Responsible!

- Explainable AI, nowadays Responsible AI
- data exploration & model explanation
- fair, unbiased models
- complete model documentation
- performance is not validation





## 2.Real case study

R & Python XAI tools

# Case study

February, Wuhan, China

COVID-19 infected patients  
485 (375 + 110)  
external test dataset

predict mortality from clinical  
variables (e.g. blood test)

XGBoost (0.99 AUC) to  
assess variable importance

interpretable decision tree  
3 variables – 0.96 AUC

## nature machine intelligence


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Article | Published: 14 May 2020

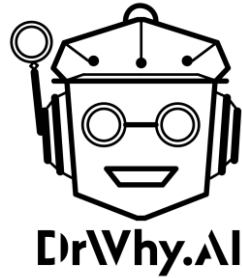
### **An interpretable mortality prediction model for COVID-19 patients**

Li Yan, Hai-Tao Zhang, [...] Ye Yuan 

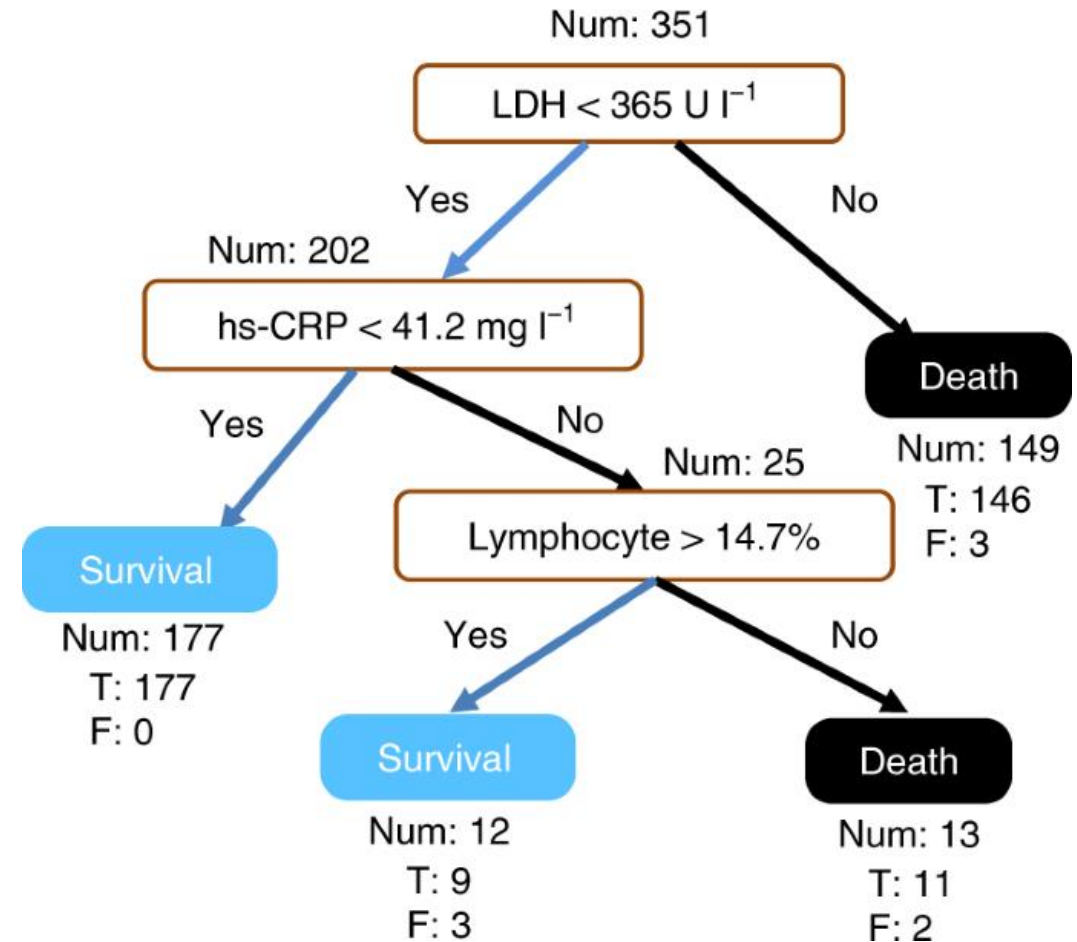
*Nature Machine Intelligence* **2**, 283–288(2020) | [Cite this article](#)

**112k** Accesses | **26** Citations | **1085** Altmetric | [Metrics](#)

# Wh-questions



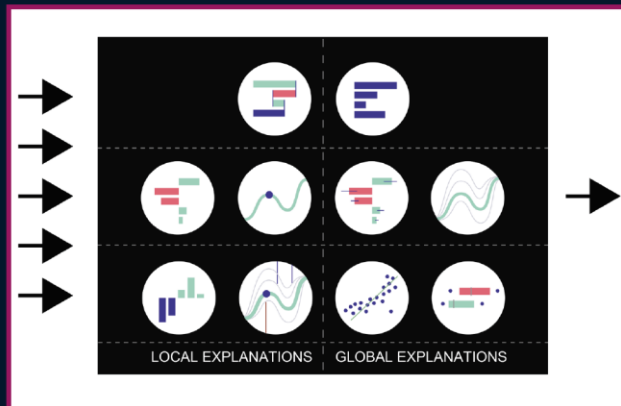
- Why is the performance so good?
- Why 3 variables out of 50 are sufficient?
- Why was age not important?
- What are the continuous relationships between variables and the target in the XGBoost model?
- ...



DATA SCIENCE SERIES

# EXPLANATORY MODEL ANALYSIS

Explore, Explain, and  
Examine Predictive Models



PRZEMYSŁAW BIECEK  
TOMASZ BURZYKOWSKI

 **CRC Press**  
Taylor & Francis Group  
A CHAPMAN & HALL BOOK

<https://pbiecek.github.io/ema>

## Model Exploration Stack

What is the model prediction  
for the selected instance?

$f(x)$  AUC  
RMSE

How good is the model?

ROC curve  
LIFT, Gain charts  
Chapter 15

Which variables contribute to  
the selected prediction?

Break Down  
SHAP, LIME  
Chapters 6, 7, 8, 9

Which variables are important  
to the model?

Permutational  
Variable Importance  
Chapter 16

How does a variable  
affect the prediction?

Ceteris Paribus  
Chapters 10, 11

How does a variable affect  
the average prediction?

Partial Dependence Profile  
Accumulated Local Effects  
Chapters 17, 18

Does the model  
fit well around  
the prediction?

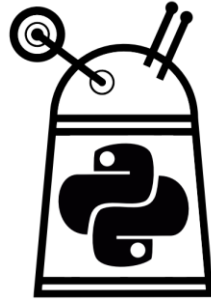
Chapter 12

Does the model  
fit well in  
general?

Chapter 19

PREDICTION LEVEL  
LOCAL EXPLANATIONS

MODEL LEVEL  
GLOBAL EXPLANATIONS



DALEX

moDel Agnostic Language for Exploration and eXplanation

Python 649 97

Python-check passing pypi package 0.2.2 downloads 11k

# MODEL

- scikit-learn
- tensorflow, keras
- xgboost, lightgbm
- ANY

# DATA

- pandas
- numpy

pip install dalex

import dalex as dx

dx.Explainer

# EXPLANATIONS

- **result** attribute (pandas)
- **plot** method (plotly)

# METHODS

predict/model + parts/profile/diagnostics  
/surrogate/performance

```
In [ ]: ▶ import seaborn as sns
import xgboost as xgb
import dalex as dx
```

```
In [ ]: ▶ # paper supplementary code
from utils_features_selection import data_preprocess, col_miss
```

```
In [ ]: ▶ def data_read():
    """
    read paper supplementary data and preprocess
    """

    data_df_unna, data_pre_df = data_preprocess()
    col_miss_data = col_miss(data_df_unna)
    col_miss_data['Missing_part'] = col_miss_data['missing_count'] / len(data_df_unna)
    sel_cols = col_miss_data[col_miss_data['Missing_part'] <= 0.2]['col']
    data_df_sel = data_df_unna[sel_cols].copy()
    data_raw = data_df_sel.fillna(-1)
    name_dict = {'乳酸脱氢酶': 'LDH', '淋巴细胞(%)': 'Lymphocytes(%)', '超敏C反应蛋白': 'hs-CRP',
                '年龄': 'Age', 'Type2': 'Target'}
    data_raw.rename(columns=name_dict, inplace=True)
    data_train = data_raw.loc[:, name_dict.values()]
    return data_train
```

```
In [ ]: ▶ # Load data
data = data_read()

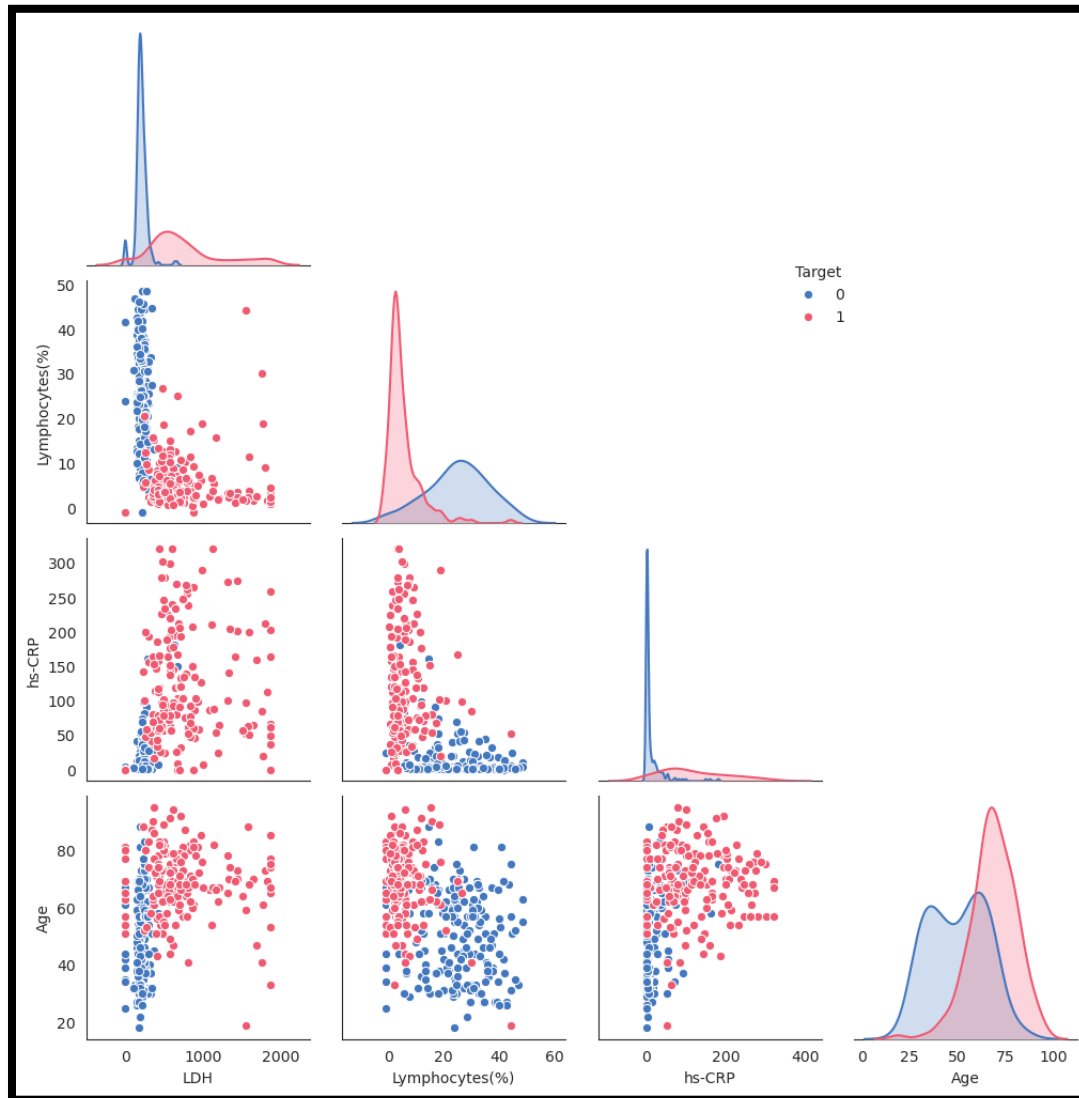
# change Target [0.0, 1.0] to [0, 1]
data.Target = data.Target.astype(int)

# plot data exploration
colors = ["#4378bf", "#f05a71"]
sns.set_palette(sns.color_palette(colors))
g = sns.pairplot(data, hue='Target', corner=True)
g._legend.set_bbox_to_anchor((0.75, 0.75))
g.fig.set_size_inches(10.5, 10)
```

```
In [ ]: ▶ # Load data
data = data_read()
X, y = data.drop('Target', axis=1), data.Target

# make a model
model = xgb.XGBClassifier(max_depth=4, learning_rate=0.2, reg_lambda=1,
                          n_estimators=150, subsample=0.9, colsample_bytree=0.9,
                          random_state=0)
model.fit(X, y)
```

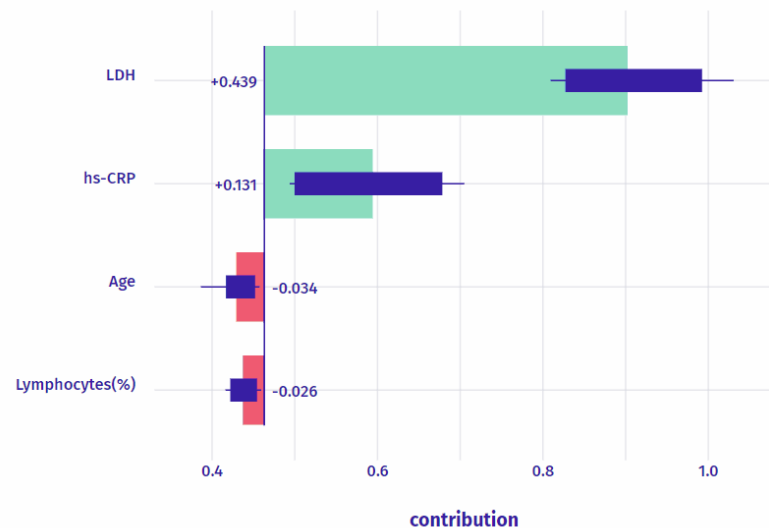
# Data exploration



	Accuracy	Balanced Accuracy	ROC AUC	F1 Score	Time Taken
<b>Model</b>					
<b>LGBMClassifier</b>	1.00	1.00	1.00	1.00	0.03
<b>XGBClassifier</b>	1.00	1.00	1.00	1.00	0.03
<b>RandomForestClassifier</b>	1.00	1.00	1.00	1.00	0.16
<b>ExtraTreesClassifier</b>	0.99	0.99	0.99	0.99	0.15
<b>SVC</b>	0.99	0.99	0.99	0.99	0.01
<b>KNeighborsClassifier</b>	0.99	0.99	0.99	0.99	0.03
<b>GaussianNB</b>	0.99	0.99	0.99	0.99	0.02
<b>AdaBoostClassifier</b>	0.99	0.99	0.99	0.99	0.09
<b>QuadraticDiscriminantAnalysis</b>	0.98	0.98	0.98	0.98	0.02
<b>NuSVC</b>	0.97	0.97	0.97	0.97	0.02
<b>LinearSVC</b>	0.97	0.97	0.97	0.97	0.01
<b>LinearDiscriminantAnalysis</b>	0.97	0.97	0.97	0.97	0.02
<b>BaggingClassifier</b>	0.97	0.97	0.97	0.97	0.04
<b>LogisticRegression</b>	0.97	0.97	0.97	0.97	0.02
<b>CalibratedClassifierCV</b>	0.97	0.97	0.97	0.97	0.13
<b>RidgeClassifier</b>	0.97	0.97	0.97	0.97	0.02
<b>RidgeClassifierCV</b>	0.97	0.97	0.97	0.97	0.01
<b>DecisionTreeClassifier</b>	0.96	0.96	0.96	0.96	0.04
<b>Perceptron</b>	0.96	0.96	0.96	0.96	0.02
<b>SGDClassifier</b>	0.96	0.96	0.96	0.96	0.01
<b>NearestCentroid</b>	0.96	0.96	0.96	0.96	0.02
<b>LabelPropagation</b>	0.95	0.94	0.94	0.95	0.02
<b>LabelSpreading</b>	0.95	0.94	0.94	0.95	0.02
<b>BernoulliNB</b>	0.95	0.94	0.94	0.95	0.02

Shapley Values

D X



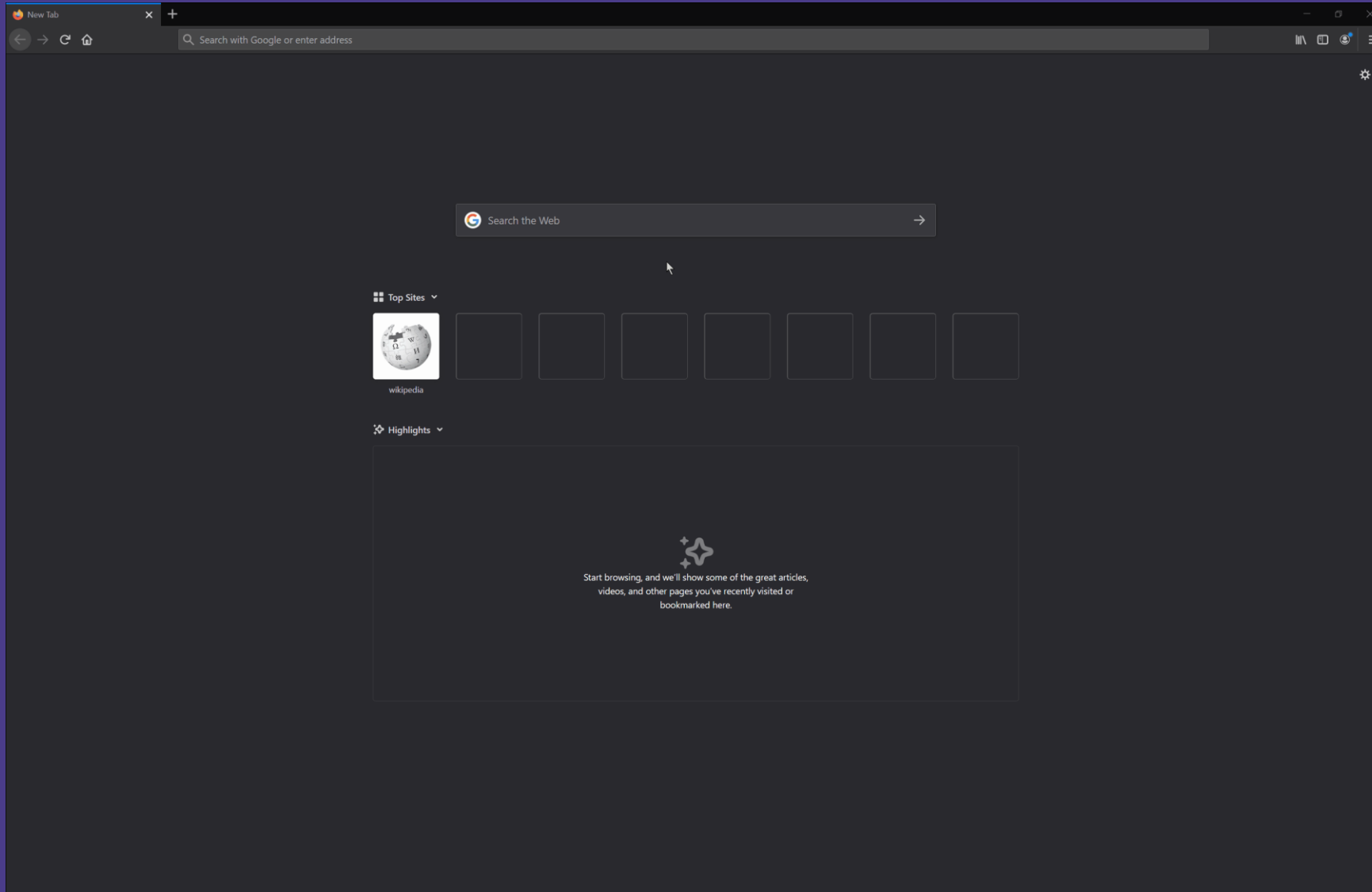
Ceteris Paribus [Local]  
Feature Importance [Global]  
Partial Dependence [Global]  
Accumulated Dependence [Global]  
Residuals vs Feature [Global]  
Feature Distribution [EDA]  
Target vs Feature [EDA]  
Average Target vs Feature [EDA]  
Break Down [Local]

**VS** AUC, ACC  
PREDICTION



# DrWhy.AI blog: Responsible ML

<https://medium.com/responsibleml>

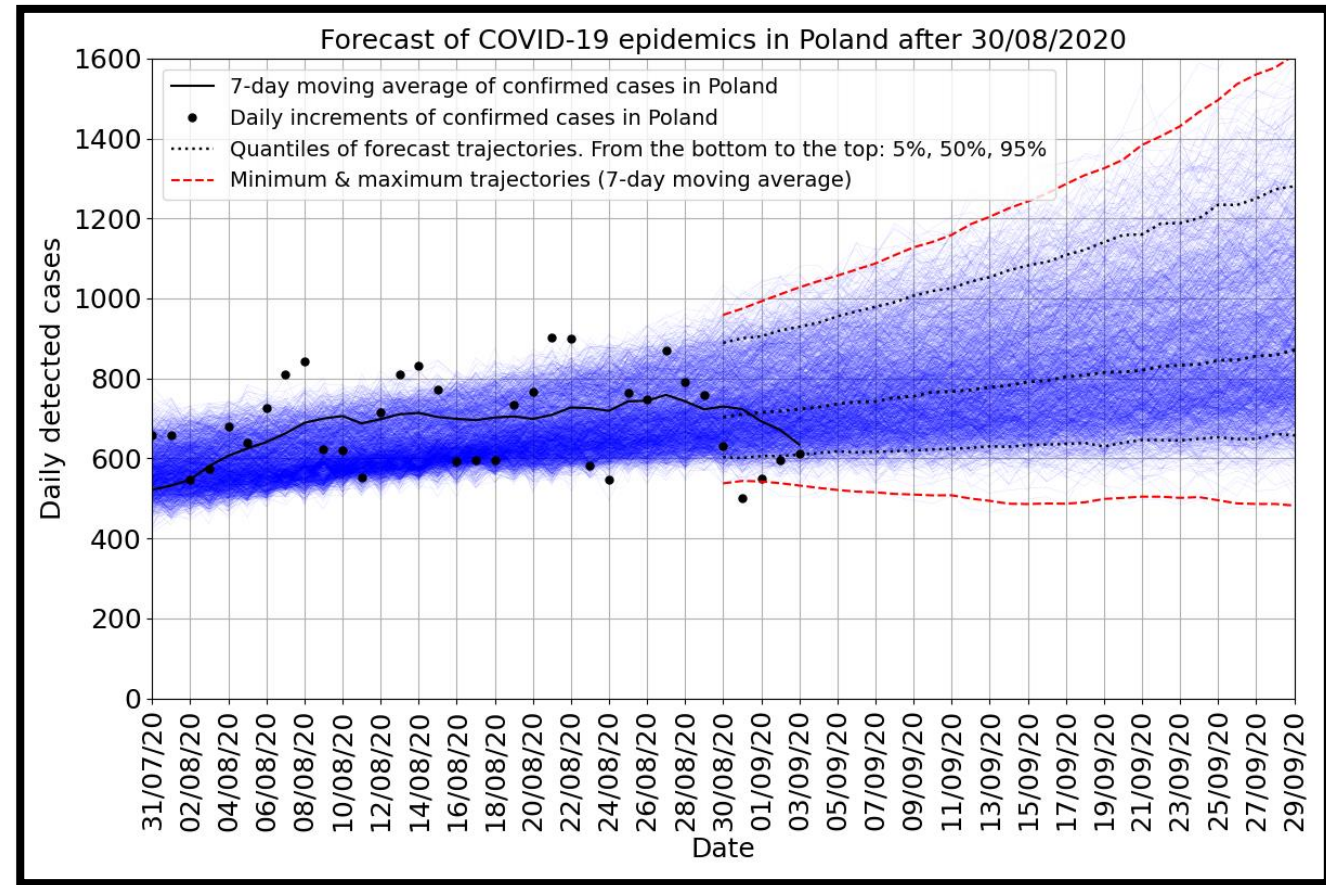


# 3.MI2 DataLab in MOCOS

COVID-19 analysis for Poland

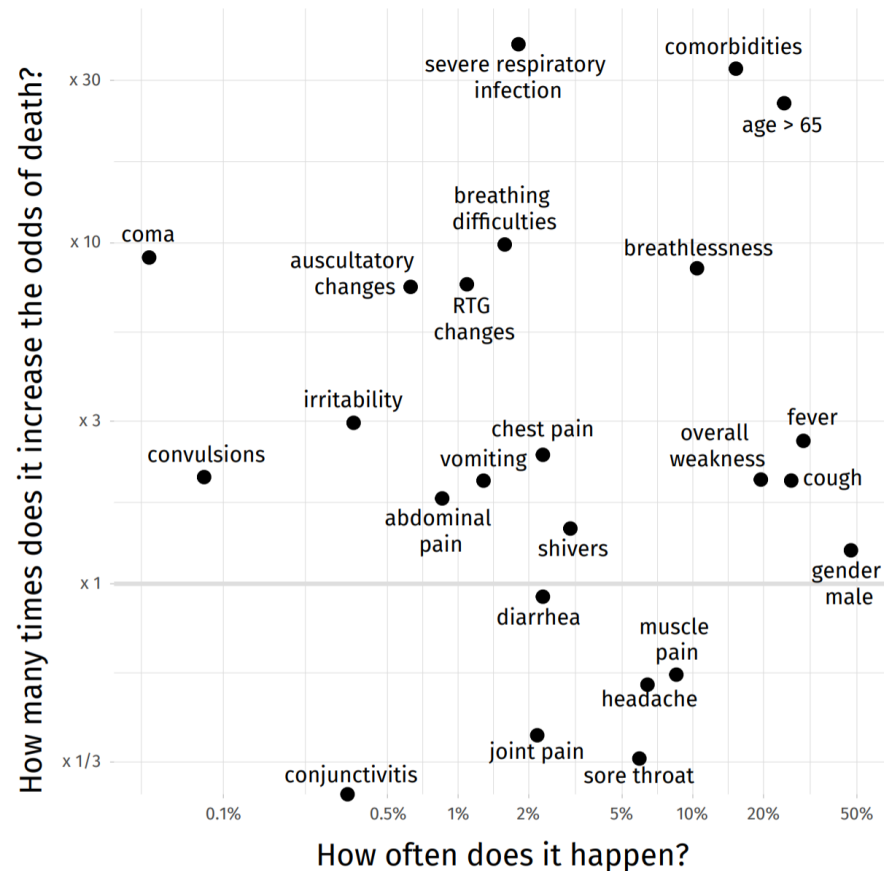
# COVID-19 in Poland

- **MI2 DataLab in MOCOS**
- **MOdelling COronavirus Spread**  
group lead by Tyll Krüger
- specific data of 30k COVID-19  
infected people in Poland
- reports to Ministry of Health
- data exploration, model  
development & explanation



# Data exploration

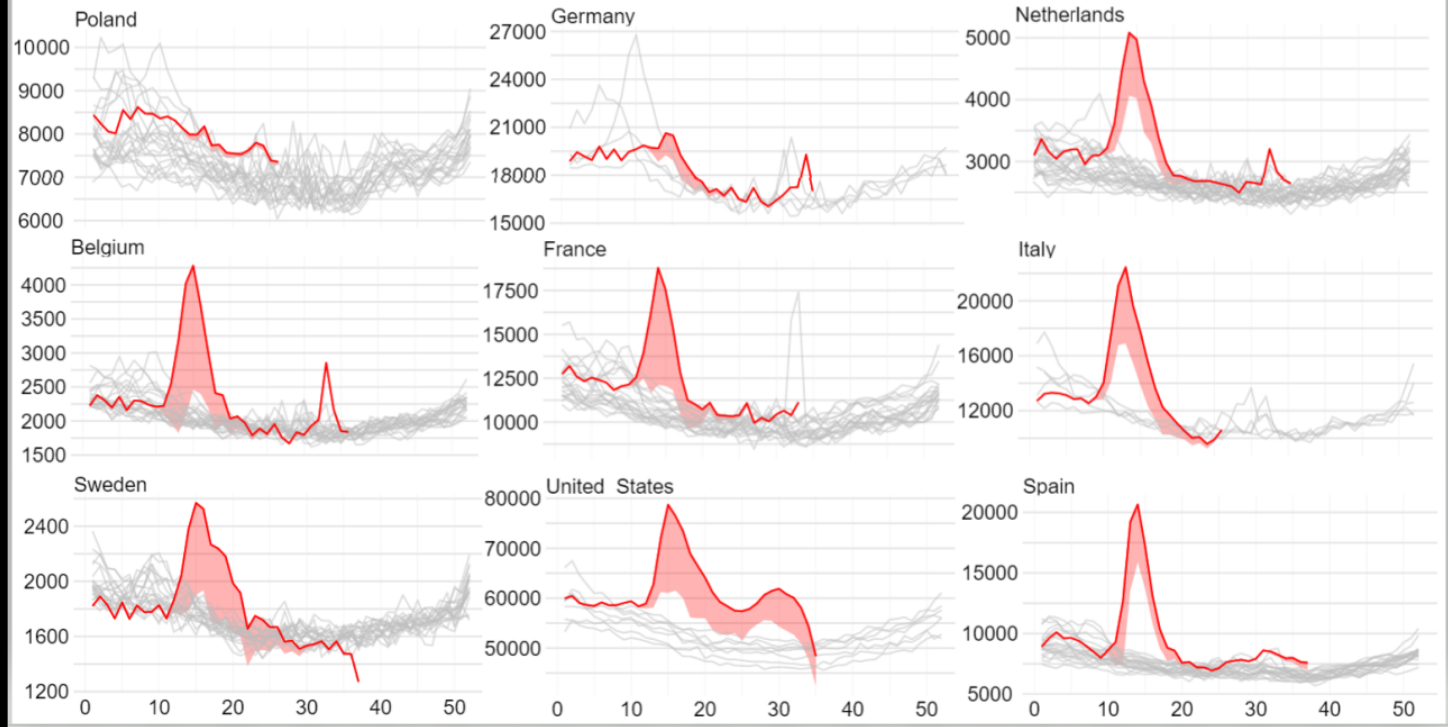
## Frequency and severity of factors associated with Covid19

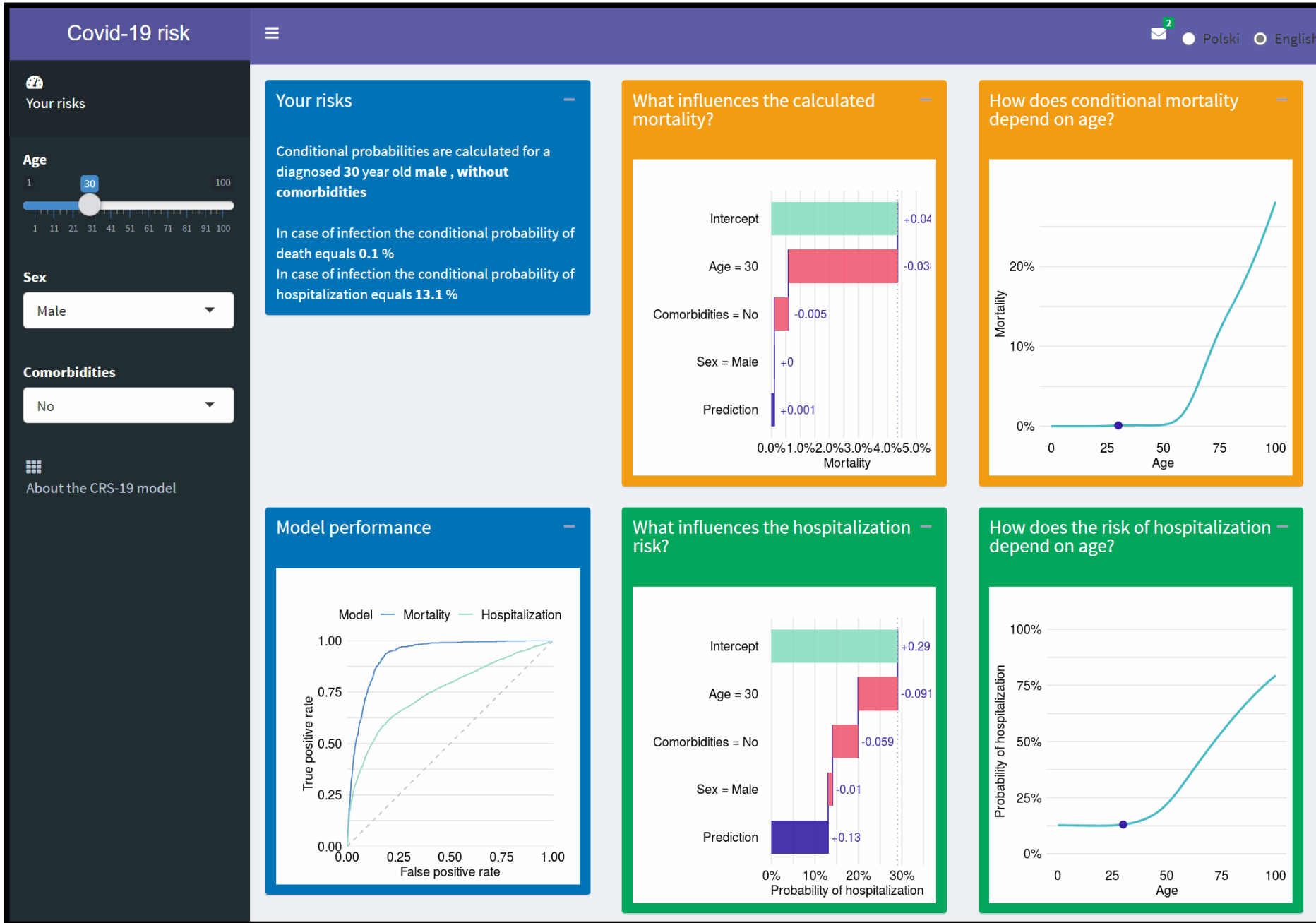


## Number of deaths vs week number in 2000-2020

### Red line - deaths in 2020

### Red field - deaths reported as COVID-19





**VS AUC, ACC PREDICTION**

# Takeaways

Advance the prediction making (not only in COVID-19 pandemic):

- Be Responsible!
- Performance-based model validation is not enough
- Proper results need data exploration & model explanation
- **DrWhy.AI:** R & Python XAI tools, resources for Responsible ML

# Contact me

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DrWhy.AI	<a href="https://drwhy.ai">https://drwhy.ai</a>
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