Interpretable Machine Learning for COVID-19:

An Empirical Study on Severity Prediction Task

- Background
- Dataset & Models
- Interpretation
- Conclusion

Background

Background

Who is responsible for the 1% error, the laptop?

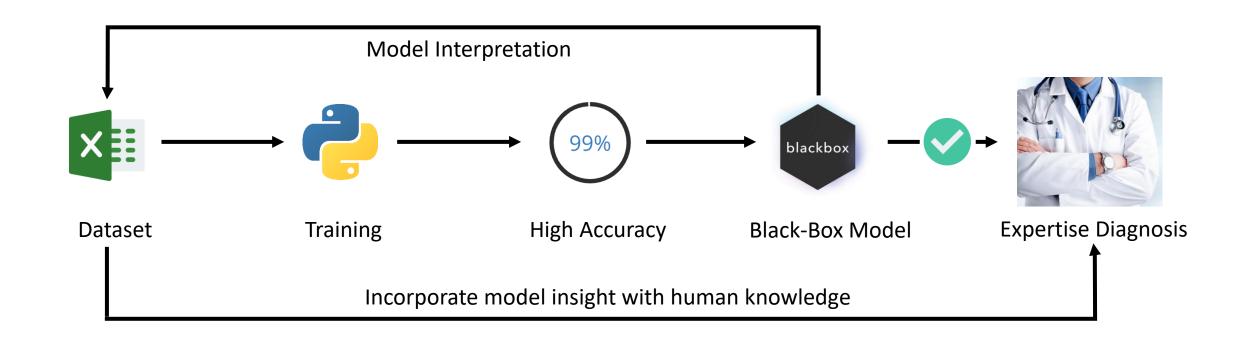
Dataset

Training

High Accuracy

Black-Box Model

Computer Diagnosis



Dataset & Models

Dataset & Models

Dataset Description

Source Code

https://covid.wuhanstudio.cc/hy-features.html

https://github.com/wuhanstudio/interpretable-ml-covid-19

Preprocessing

- 1. Remove clinical irrelevant columns ['MedNum', 'No']
- 2. Remove columns that have no data ['LVEF', 'SO2', 'PO2', 'YHZS', 'RML', 'RUL', 'RLL', 'LUL', 'LLL']
- 3. Remove columns with fewer records ['Onset2Admi', 'Onset2CT1', 'Onset2CTPositive1', 'Onset2CTPeak']
- 4. Remove patients that have incomplete records. [Height == "", cTnl == ""]

Train Test Split (86 patients, 21 severe, 65 normal, with 55 features)

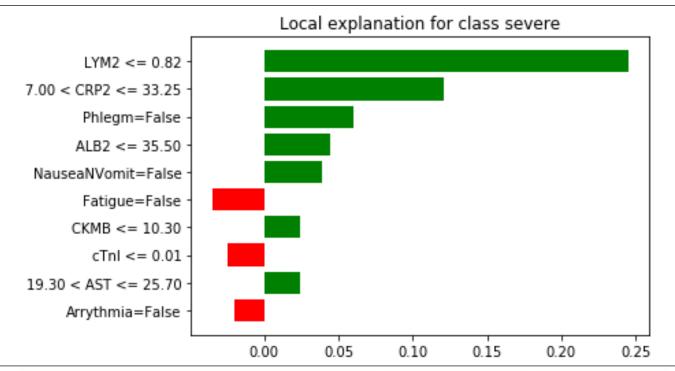
- 1. 90% as Training Set, 10% as Testing Set
- 2. Five folds Cross Validation

In this research, we focus on model **interpretation**. **Interpretation** gives us valuable insights even if these are not 99% perfect models.

Classifier	Cross Validation	Test Set		95% confidence-interval	
	F1	Precision	Recall	F1	
Decision Tree	0.56	0.67	0.50	0.57	0.307
Random Forest	0.64	0.56	0.25	0.33	0.324
XGBoost	0.62	0.78	1.00	0.80	0.271
Neural Network	0.53	0.78	1.00	0.80	0.271

Interpretation

Correct Predictions



Feature	Value
LYM2	0.46
CRP2	22.20
Phlegm=False	True
ALB2	35.30
NauseaNVomit=Fa	alse True
Fatigue=False	True
CKMB	8.30
cTnI	0.01
AST	19.70
Arrythmia=False	True

Prediction probabilities			
normal	0.02		
severe		0.98	

Feature	Patients Data	Normal Range	Conclusion
LYM <= 0.82	0.46	(1.1, 3.2)	\downarrow
CRP (7.00, 33.25)	22.20	(0, 5)	\uparrow
No Phlegm	True	-	No Symptom
ALB <= 35.50	35.30	(35, 55)	Normal
No Nausea and Vomit	True	-	No Symptom
CKMB <= 10.30	8.30	(0, 18)	Normal
AST (19.30, 25.70)	19.70	(29, 35)	\downarrow

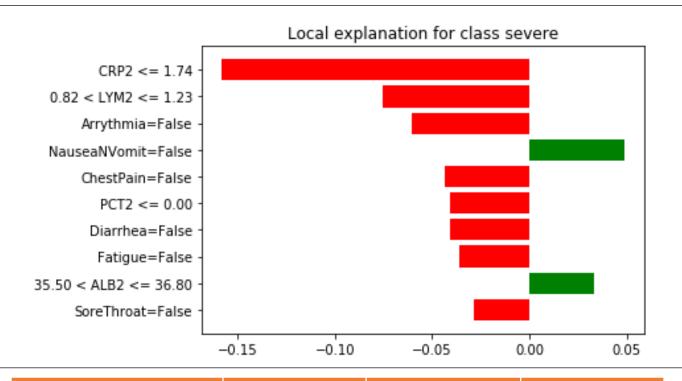
Feature	Patients Data	Normal	Conclusion
No Fatigue	True	-	Normal
cTnl <= 0.1	0.01		
No Arrythmia	True	-	Normal

No Phlegm, No Nausea and Vomit is more likely to turn severe?

This is counterintuitive, but corroborated by medical research months later after the outbreak of the pandemic.

Patient 0 - Normal

Correct



Feature	Value
CRP2	0.46
LYM2	0.84
Arrythmia=False	True
NauseaNVomit=Fal	se True
ChestPain=False	True
PCT2	0.00
Diarrhea=False	True
Fatigue=False	True
ALB2	36.50
SoreThroat=False	True

Prediction p	robabilities	
normal		0.97
severe	0.03	

Feature	Patients Data	Normal Range	Conclusion
No Nausea and Vomit	True	-	No Symptom
ALB (35.50, 36.80)	36.50	(35, 55)	Normal

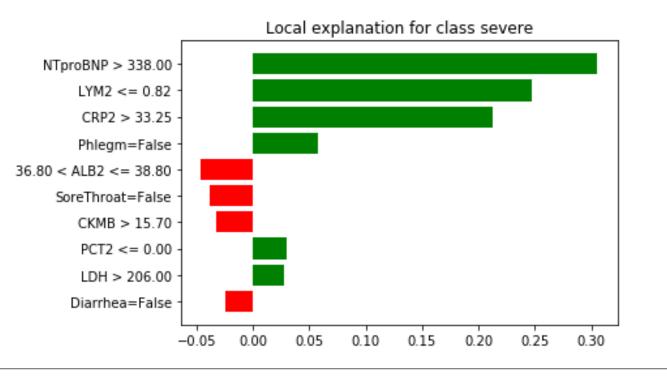
Our models notice that without some specific symptoms may indicate a higher probability of turning severe.

Feature	Patients Data	Normal	Conclusion
CRP <= 1.74	0.46	(0, 5)	Normal
LYM (0.82, 1.23)	0.84	(1.1, 3.2)	\downarrow
No Arrythmia	True	X	
No ChestPain	True	X	
PCT <= 0.00	0.00	(0, 0.5)	Normal
No Diarrhea	True	Χ	
No Fatigue	True	X	
No SoreThroat	True	Χ	

Wrong Predictions

Patient 3 - Severe





Feature	Value
NTproBNP	475.00
LYM2	0.81
CRP2	78.76
Phlegm=False	True
ALB2	37.60
SoreThroat=Fals	e True
CKMB	17.90
PCT2	0.00
LDH	263.00
Diarrhea=False	True

Prediction probabilities			
normal	0.42		
severe	0.58		

Feature	Patients Data	Normal Range	Conclusion
NTproBNP > 338.00	475.00	(0, 300)	\uparrow
LYM <= 0.82	0.81	(1.1, 3.2)	\downarrow
CRP > 33.25	78.76	(0, 5)	\uparrow
No Phlegm	True	-	
PCT	0.00	(0, 0.5)	Normal
LDH	263.00	(109, 245)	\uparrow

Feature	Patients Data	Normal Range	Conclusion
ALB (36.80, 38.80)	37.60	(35, 55)	Normal
No SoreThroat	True	-	Normal
CKMB > 15.70	17.90	(0, 18)	Normal
No Diarrhea	True	-	Normal

A normal patient is diagnosed as sever by our models. But actually he is indeed severe from medical tests.

Conclusion

Conclusion

 \triangleright Elderly people are more likely to turn severe. The average age of severity level [0-3]: 36.8 --> 47.5 --> 54.3 --> 69.4

Interpretation

- Inflammation Indicators: High CRP (Virus Infection)
- Blood Routing Examination: Low LYM
- Biomarker: High LDH, Low ALB

Symptoms: No Phlegm, No diarrhea, No Nausea and Vomit