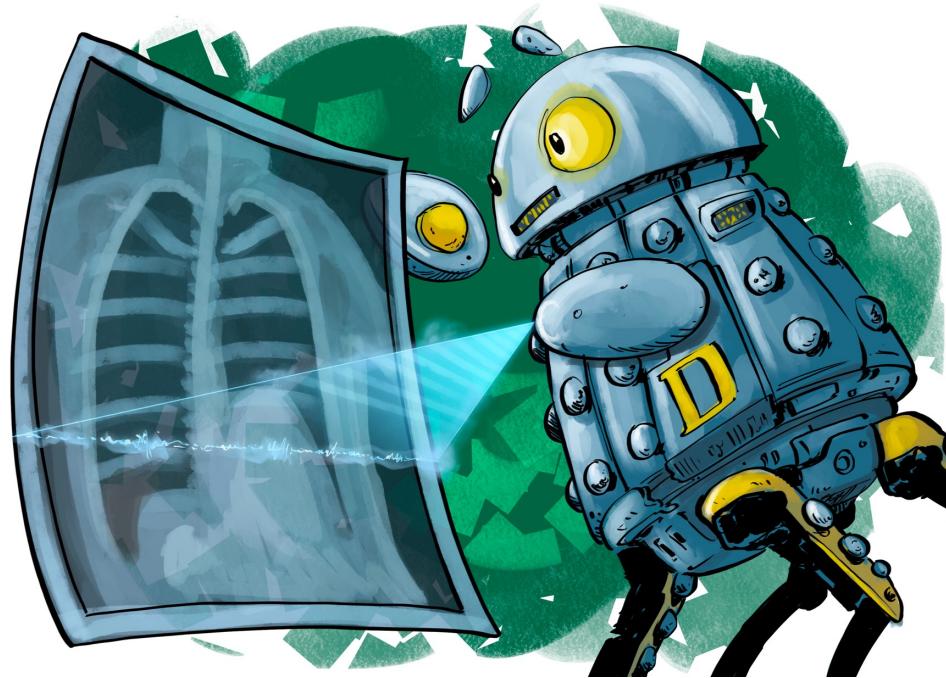


Jak budować skuteczne
i bezpieczne modele uczenia
maszynowego dla zastosowań
medycznych

How to build trustworthy machine
learning models for medical
applications

Przemysław Biecek



About me and MI².ai



Associate professor at University of Warsaw and Warsaw University of Technology
Past: Samsung, Netezza, IBM, Disney, iQor

Research area: **Human Oriented AI**

Research projects:

MLGenSig (NCN), DALEX (NCN), HOMER (NCN), DeCovid (IDUB)

X-LUNGS (NCBiR): Responsible Artificial Intelligence for Lung Diseases



Where we are with AI?

MI × LUNGS



“

Data is the new oil”

Clive Humby



“

AI is the new electricity.

Andrew Ng, Baidu

”



https://www.slideshare.net/jaypod/digitaltransformation50soundbites/19-Data_is_the_new_oilClive

<https://www.newworldai.com/forget-the-hype-what-every-business-leader-needs-to-know-about-artificial-intelligence-now/>

Where we are with AI?

MI × LUNGS

List of failures in AI applications

Read more at: <https://romanlutz.github.io/ResponsibleAI/>

Speech Detection

- Oh dear... AI models used to flag hate speech online are, er, racist against black people
- The Risk of Racial Bias in Hate Speech Detection
- Toxicity and Tone Are Not The Same Thing: analyzing the new Google API on toxicity, PerspectiveAPI.
- Voice Is the Next Big Platform, Unless You Have an Accent
- Google's speech recognition has a gender bias
- Fair Speech report by Stanford Computational Policy Lab, also covered in [Speech recognition algorithms may also have racial bias](#)
- Automated moderation tool from Google rates People of Color and gays as "toxic"
- Someone made an AI that predicted gender from email addresses, usernames. It went about as well as expected

Image Labelling & Face Recognition

- Google Photos identified two black people as 'gorillas'
- When It Comes to Gorillas, Google Photos Remains Blind
- The viral selfie app ImageNet Roulette seemed fun – until it called me a racist slur
- Google Is Investigating Why it Trained Facial Recognition on 'Dark Skinned' Homeless People
- Gender Shades: Intersectional Accuracy Disparities in Commercial Gender Classification
- Machines Taught by Photos Learn a Sexist View of Women
- Tenants sounded the alarm on facial recognition in their buildings. Lawmakers are listening.
- Google apologizes after its Vision AI produced racist results

Public Benefits & Health



Topics

Video

Events

Podcasts

Machine Learning is Creating a Crisis in Science

The adoption of machine-learning techniques is contributing to a worrying number of research findings that cannot be repeated by other researchers.

Kevin McCaney

Wed, 02/27/2019 - 11:28



Photo credit: metamorworks/iStock

Do we have a problem?

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AN EPIC FAILURE: OVERSTATED AI CLAIMS IN MEDICINE

	Epic +	Epic -	Total
Sepsis	843	1,709	2,552
No Sepsis	6,128	29,775	35,903
Total	6,971	31,484	38,455

- Of the 2,552 patients with sepsis, ESM only generated sepsis alerts for 843 (33 percent). They missed 67 percent of the people with sepsis.
- Of the 6,971 ESM sepsis alerts, only 843 (12 percent) were correct; 88 percent of the ESM sepsis alerts were false alarms, creating what the authors called “a large burden of alert fatigue.”

Results: We identified 27 697 patients who had 38 455 hospitalizations (21 904 women [57%]; median age, 56 years [interquartile range, 35–69 years]) meeting inclusion criteria, of whom sepsis occurred in 2552 (7%). The ESM had a hospitalization-level area under the receiver operating characteristic curve of 0.63 (95% CI, 0.62–0.64). The ESM identified 183 of 2552 patients with sepsis (7%) who did not receive timely administration of antibiotics, highlighting the low sensitivity of the ESM in comparison with contemporary clinical practice. The ESM also did not identify 1709 patients with sepsis (67%) despite generating alerts for an ESM score of 6 or higher for 6971 of all 38 455 hospitalized patients (18%), thus creating a large burden of alert fatigue.

Conclusions and relevance: This external validation cohort study suggests that the ESM has poor discrimination and calibration in predicting the onset of sepsis. The widespread adoption of the ESM despite its poor performance raises fundamental concerns about sepsis management on a national level.

<https://pubmed.ncbi.nlm.nih.gov/34152373/>

Do we have a problem?



IBM spent more than \$15 billion on Dr. Watson with no peer-reviewed evidence that it improved patient health outcomes. Watson Health has disappointed so soundly that IBM is now looking for someone to take it off their hands.

“ Five years and \$60 million later, MD Anderson fired Watson after “multiple examples of unsafe and incorrect treatment recommendations.”

Dr. Watson was the most hyped computerized health care system, but it is hardly the only disappointing one. Most recently, a 2021 study looked at 2,212 research papers published during the period from January 1, 2020, to October 3, 2020, that described new machine learning models for diagnosing or prognosing COVID-19 from chest radiographs and chest computed tomography images. Their conclusion: “None of the models identified are of potential clinical use.”

AI act to the rescue!

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EUROPEAN COMMISSION

Brussels, 21.4.2021

COM(2021) 206 final

2021/0106(COD)

Proposal for a

REGULATION OF THE EUROPEAN PARLIAMENT AND OF THE COUNCIL

LAYING DOWN HARMONISED RULES ON ARTIFICIAL INTELLIGENCE (ARTIFICIAL
INTELLIGENCE ACT) AND AMENDING CERTAIN UNION LEGISLATIVE ACTS

{SEC(2021) 167 final} - {SWD(2021) 84 final} - {SWD(2021) 85 final}

EXPLANATORY MEMORANDUM

1. CONTEXT OF THE PROPOSAL

1.1. Reasons for and objectives of the proposal

This explanatory memorandum accompanies the proposal for a Regulation laying down harmonised rules on artificial intelligence (Artificial Intelligence Act). Artificial Intelligence (AI) is a fast evolving family of technologies that can bring a wide array of economic and societal benefits across the entire spectrum of industries and social activities. By improving prediction, optimising operations and resource allocation, and personalising service delivery, the use of artificial intelligence can support socially and environmentally beneficial outcomes and provide key competitive

**The Regulatory Framework
4 levels of risk in AI:**

Unacceptable risk
High risk
Limited risk
Minimal or no risk

Important but overlooked aspects



To build a good AI system it is needed to take care of:

- high quality **data**
- methods of reducing undesired **bias**
- methods of exploring and **explaining** models
- many other topics that I will not talk about today

Analysis | Open Access | Published: 15 March 2021

Common pitfalls and recommendations for using machine learning to detect and prognosticate for COVID-19 using chest radiographs and CT scans

Michael Roberts , Derek Driggs, Matthew Thorpe, Julian Gilbey, Michael Yeung, Stephan Ursprung,

Angelica I. Aviles-Rivero, Christian Etmann, Cathal McCague, Lucian Beer, Jonathan R. Weir-McCall,

Zhongzhao Teng, Effrossyni Gkrania-Klotsas, AIX-COVNET, James  Recommendations for data

Bibiane Schönlieb

Nature Machine Intelligence 3, 199–217 (2021) | [Cite this article](#)

72k Accesses | 201 Citations | 1160 Altmetric | [Metrics](#)

First, we advise caution over the use of public repositories, which can lead to high risks of bias due to source issues and Frankenstein datasets as discussed above. Furthermore, authors should aim to match demographics across cohorts, an often neglected but important potential source of bias; this can be impossible with public datasets that do not include demographic information, and including paediatric images⁸⁶ in the COVID-19 context introduces a strong bias.

Using a public dataset alone without additional new data can lead to community-wide overfitting on this dataset. Even if each individual study observes sufficient precautions to avoid overfitting, the fact that the community is focused on outperforming benchmarks on a single public dataset encourages overfitting. Many public datasets containing images taken from preprints receive these images in low-resolution or compressed formats (for example

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

Laure Wynants,^{1,2} Ben Van Calster,^{2,3} Gary S Collins,^{4,5} Richard D Riley,⁶ Georg Heinze,⁷ Ewoud Schuit,^{8,9} Marc M J Bonten,^{8,10} Darren L Dahly,^{11,12} Johanna A Damen,^{8,9} Thomas P A Debray,^{8,9} Valentijn M T de Jong,^{8,9} Maarten De Vos,^{2,13} Paula Dhiman,^{4,5} Maria C Haller,^{7,14} Michael O Harhay,^{15,16} Liesbet Henckaerts,^{17,18} Pauline Heus,^{8,9} Michael Kammer,^{7,19} Nina Kreuzberger,²⁰ Anna Lohmann,²¹ Kim Luijken,²¹ Jie M...
Glen P Martin,²² David J McLernon,²³ Constanza L Andaur Navarro,^{8,9} Johannes E...
Jamie C Sergeant,^{24,25} Chunhu Shi,²⁶ Nicole Skoetz,¹⁹ Luc J M Smits,¹ Kym I E Snijders,²⁷ Matthew Sperrin,²⁷ René Spijker,^{8,9,28} Ewout W Steyerberg,³ Toshihiko Takada,⁸ Ioanna Tzoulaki,^{29,30} Sander M J van Kuijk,³¹ Bas C T van Bussel,^{1,32} Iwan C C van der Velde,¹ Florien S van Royen,⁸ Jan Y Verbakel,^{33,34} Christine Wallisch,^{7,35,36} Jack Wilkinson,³⁷ Robert Wolff,³⁷ Lotty Hooft,^{8,9} Karel G M Moons,^{8,9} Maarten van Smeden⁸

<https://www.bmjjournals.org/content/bmjj/369/bmj.m1328.full.pdf>

Cite this as: **BMJ** 2020;369:m1328

<http://dx.doi.org/10.1136/bmj.m1328>

Originally accepted:

31 March 2020

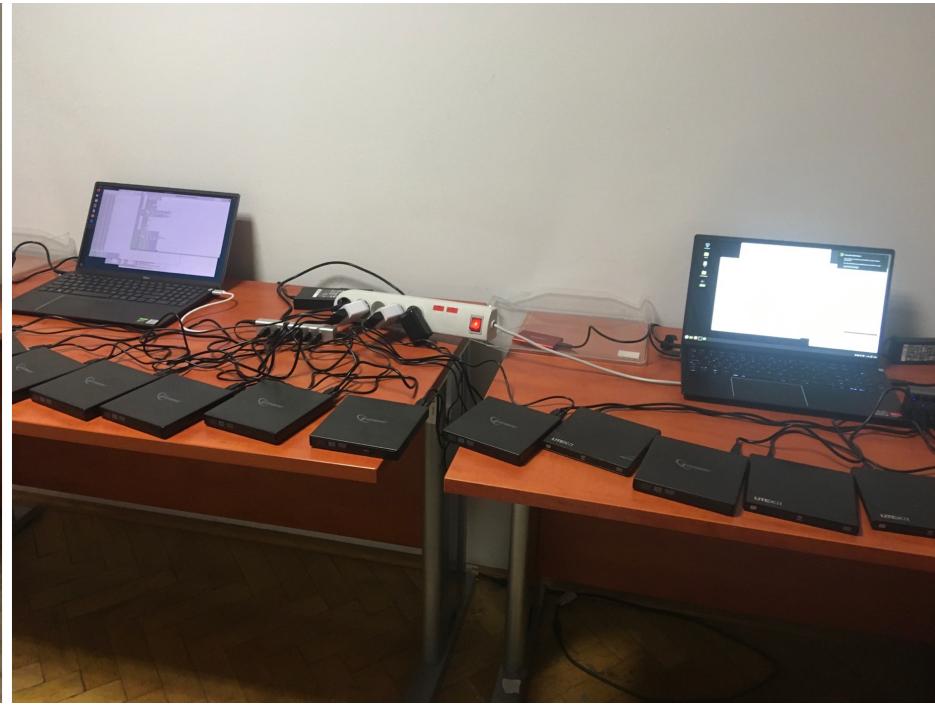
Final version accepted:

12 January 2021

Conclusion

Several diagnostic and prognostic models for covid-19 are currently available and they all report moderate to excellent discrimination. However, these models are all at high or unclear risk of bias, mainly because of model overfitting, inappropriate model evaluation (eg, calibration ignored), use of inappropriate data sources and unclear reporting. Therefore, their performance estimates are probably optimistic and not representative for the target population. The COVID-PRECISE group does not recommend any of the current prediction models to be used in practice, but one

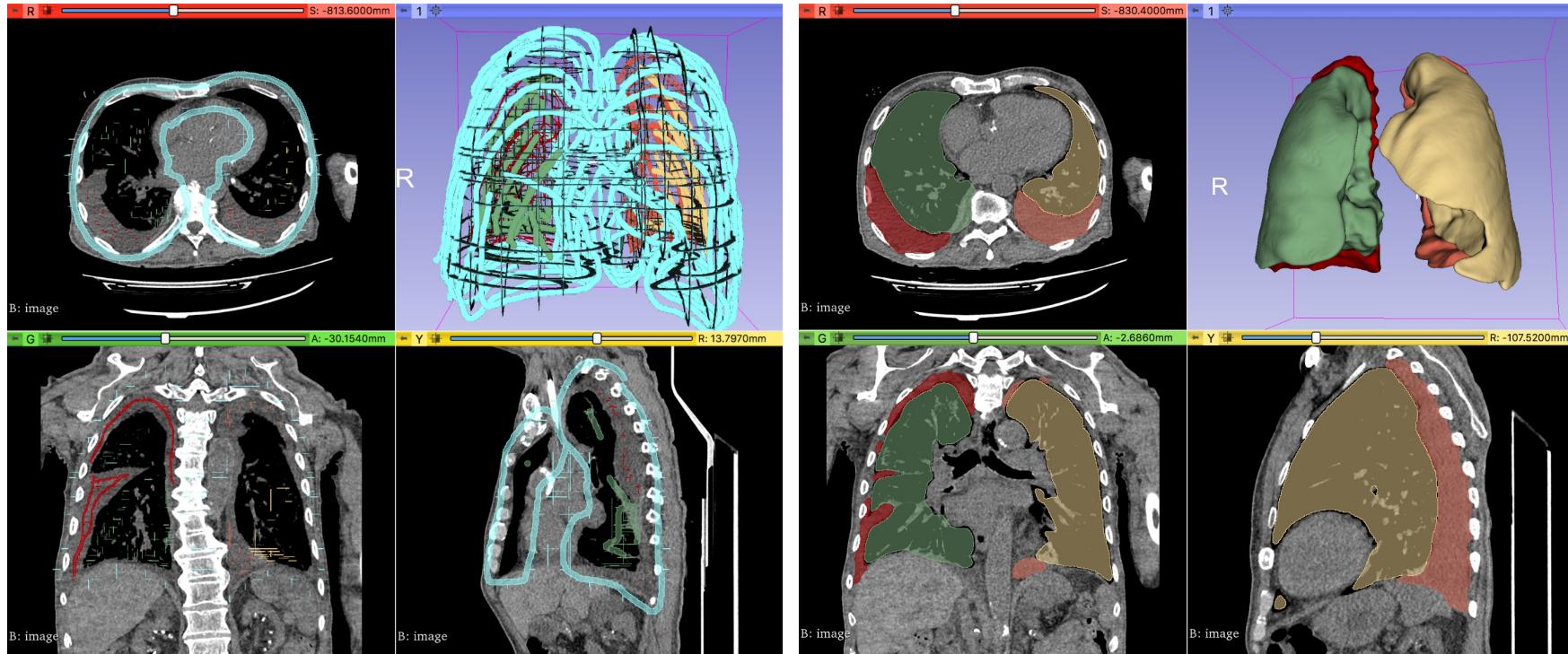
Data is King



- We prepared a dataset with more than 40,000 CT scans (plus xrays).
- The data is representative of the Polish population.
- **Data will be made public** for other teams soon.
- We will also prepare a special benchmark allowing for proper validation of the developed solutions.

Data is King

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- The data have different modalities (series of images + text medical records + tabular data).
- And we have identified over 30 different tasks for them.
- The tasks vary from binary classification to regression to segmentation.

Data is King

RadioTator

The screenshot shows the RadioTator interface. At the top left are 'Filters' and a 'Filter' button. Below are two dropdown menus, both currently set to 'Select...'. To the right are filter options: 'And' (radio button), 'Or' (radio button, selected), 'substring' (radio button), 'word' (radio button), 'positive' (radio button, selected), 'negative' (radio button). Further right are checkboxes for 'With final annotations' (unchecked) and 'Without final annotations' (checked). Below these are fields for 'Text ID:' (empty), 'Text length:' (set to 'All'), and two search terms: 'right lung' and 'inflammation', each with 'And' or 'Or' radio buttons.

Result number: 0-5 / 200

Previous page Next page

Save annotations

Annotations listed:

- Cardiomegaly
- Lesion
- Pleural Effusion
- Interesting case. Must check later.

- Very often, relevant information is stored in plain text descriptions.
- And these descriptions may be very diverse.
- Extracting structures data can be aided by applying advanced NLP techniques
- In the picture you can see the RadioTator that we developed for more efficient extraction of data from text

Important but overlooked aspects

To build a good AI system it is needed to take care of:

- high quality **data**
- methods of reducing undesired **bias**
- methods of exploring and **explaining** models
- many other topics that I will not talk about today

Recurring Nightmares: Bias

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ProPublica

Local Initiatives

Data Store

Facebook Twitter Donate



Graphics & Data Newsletters About

Get the Big Story

Join

Racial Justice

Health Care

Labor

Criminal Justice

More...

Series

Video

Impact

Search



MACHINE BIAS



Facebook Ads Can Still Discriminate Against Women and Older Workers, Despite a Civil Rights Settlement



New research and Facebook's own ad archive show that the company's new system to ensure diverse audiences for housing and employment ads has many of the same problems as its predecessor.

by Ava Kofman and Ariana Tobin, Dec. 13, 2019, 5 a.m. EST

Inactive
Nov 4, 2019 - Nov 14, 2019
ID: 991590397853985



About social issues, elections or politics

Dolese Bros. Co.
Sponsored • Paid for by Dolese Bros. Co.

A local career to keep you close to home. We're hiring CDL drivers! #DoleseDelivers #CareerOpportunities



Now Hiring CDL Drivers
Join our team today!
WWW.DOLESE.COM

Apply Now

See Ad Details

Data About This Ad

Inactive
Nov 4, 2019 - Nov 14, 2019
ID: 991590397853985

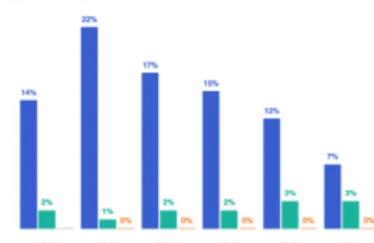
20K - 25K Impressions

\$100 - \$199 Money spent (USD)

Who Was Shown This Ad

Age and Gender

Men Women Unknown



Left: A Facebook ad for Dolese Bros. Co. Right: Facebook's chart shows that 87 percent of the people who saw the ad were men.

Recurring Nightmares: Bias

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Key point

- Discrimination defines a situation where an individual is disadvantaged in some way on the basis of 'one or multiple protected grounds'.

PROTECTED GROUNDS

- Sex
- Gender identity
- Sexual orientation
- Disability
- Age
- Race, ethnicity, colour and membership of a national minority
- Nationality or national origin
- Religion or belief
- Social origin, birth and property
- Language
- Political or other opinion

https://fra.europa.eu/sites/default/files/fra_uploads/fra-2018-handbook-non-discrimination-law-2018_en.pdf

<https://fra.europa.eu/en/publication/2018/handbook-european-non-discrimination-law-2018-edition>

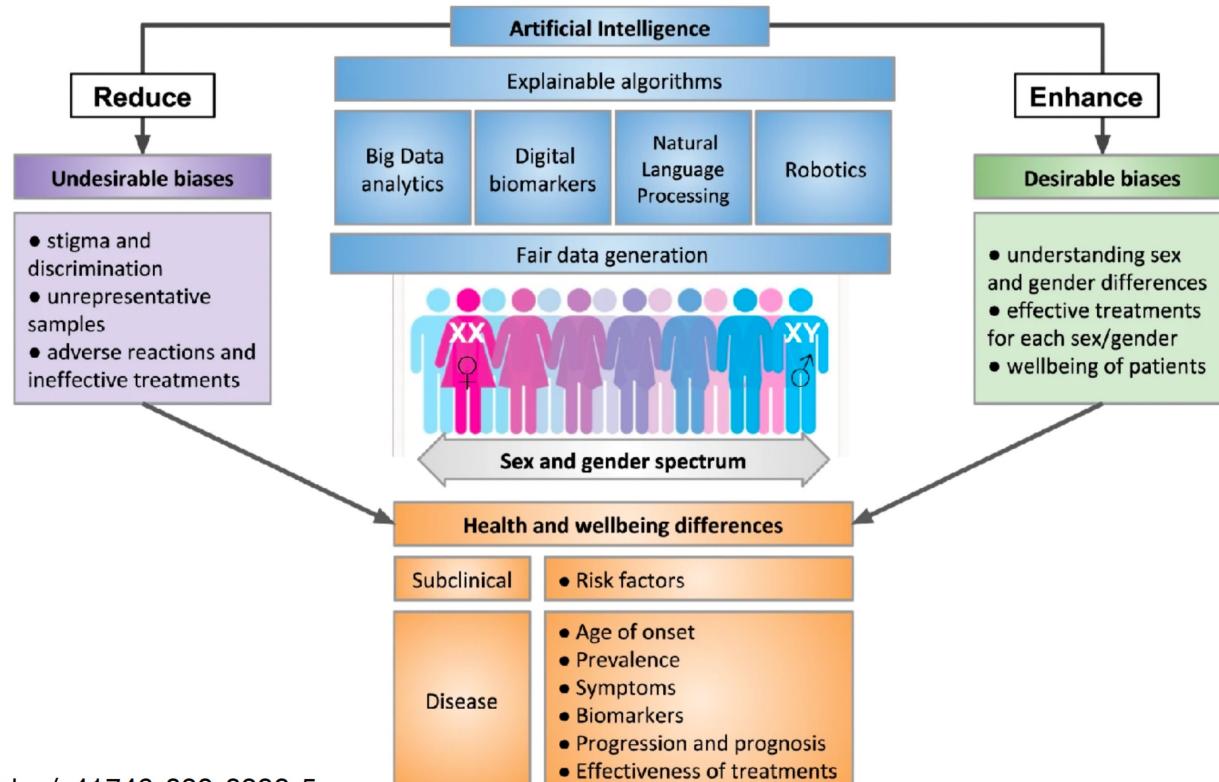


Recurring Nightmares: Bias

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Sex and gender differences and biases in artificial intelligence for biomedicine and healthcare

Cirillo et al 2020



Recurring Nightmares: Bias



Sex and gender differences and biases in artificial intelligence for biomedicine and healthcare

Cirillo et al 2020

Clinical conditions and studies	Current status without the desirable bias	Utility of the desirable bias
Autistic spectrum disorder	<p>There is a current lack of consideration of the demonstrated age-dependent sex differences in the symptomatology related with impairments in social communication and interaction, expressive behaviour, reciprocal conversation, non-verbal gestures for diagnostic purposes¹²³.</p>	Differential diagnostic criteria for males and females could facilitate the identification of the clinical diagnosis leading to appropriate treatment.
Cardiovascular disorders	<p>Although it has been documented that men and women respond differently to many cardiovascular medications such as statins, angiotensin-converting enzyme inhibitors and β-Blockers among others, adopted treatments do not consider sex differences¹²⁴.</p> <p>Despite the fact that Coronary heart disease (CHD) is the leading cause of death among women¹²⁵, the majority (67%) of patients enroled in clinical trials for cardiovascular devices are male¹²⁶.</p>	Making prescriptions according to the sex of the patient could lead to improved health benefits.
Genome-wide association studies (GWAS)	<p>Most of genome-wide association studies (GWAS) focus on white male subjects¹²⁷ and those that explore sex differences in complex traits are scarce¹²⁸.</p>	The application of a desirable bias towards women would lead to a more accurate representation of sex differences in clinical research.
Human immunodeficiency virus (HIV)	<p>The observed lower female representation in HIV clinical trials depends, among other factors, from the disadvantaged awareness about treatment and enrolment options compared with men¹²⁹⁻¹³¹.</p>	The introduction of desirable biases to deliberately include female subjects and other ethnicities in GWAS could lead to better account for potential sex differences in disease that are currently unknown because of being overlooked.
		Promoting empowerment initiatives in those patients with disadvantages will increase their exposure to treatment options and clinical trial enrolment.

Recurring Nightmares: Bias



- **Historical bias.** The data are correctly sampled and correspond well to the observed relationships, but due to different treatment in the past some prejudices are encoded in the data. Think about gender and occupation stereotypes.
- **Representation bias.** The available data is not a representative sample of the population of interest. Think about the available facial images of actors, often white men. Or genetic sequences of covid variants, mostly collected in developed European countries. Or crime statistics in the regions to which the police are directed.
- **Measurement bias.** The variable of interest is not directly observable or is difficult to measure and the way it is measured may be distorted by other factors. Think of the results of the mathematics skills assessment (e.g. PISA) measured by tasks on computers not that widely available in some countries.
- **Evaluation bias.** The evaluation of the algorithm is performed on a population that does not represent all groups. Think of a lung screening algorithm tested primarily on a population of smokers (older men).
- **Proxy bias.** The algorithm uses variables that are proxies for protected attributes. Think of male/female only schools where the gender effect can be hidden under the school effect.

Recurring Nightmares: Bias

Group fairness / statistical parity / independence / demographic parity

$$P(\hat{Y} = 1|A = a) = P(\hat{Y} = 1|A = b) \quad \hat{Y} \perp A$$

Predicted class is independent from protected attribute

	$Y = 1$	$Y = 0$	
$\hat{Y} = 1$	TP	FP	P
$\hat{Y} = 0$	FN	TN	N
	π_1	π_0	

Equalized odds, Separation, Positive Rate Parity

$$P(\hat{Y} = 1 | A = a, Y = 1) = P(\hat{Y} = 1 | A = b, Y = 1)$$

$$\hat{Y} \perp A \mid Y$$

$$P(\hat{Y} = 1 | A = a, Y = 0) = P(\hat{Y} = 1 | A = b, Y = 0)$$

Equal True Positive Rate TPR = TP/(TP+FN) for each subgroup and

equal False Positive Rate FPR = FP/(FP+TN) for each subgroup

Predicted class is independent from protected attribute given true class

	$Y = 1$	$Y = 0$	
$\hat{Y} = 1$	TP	FP	P
$\hat{Y} = 0$	FN	TN	N
	π_1	π_0	

Predictive Rate Parity, Sufficiency

$$P(Y = 1|A = a, \hat{Y} = 1) = P(Y = 1|A = b, \hat{Y} = 1)$$

$$Y \perp A \mid \hat{Y}$$

$$P(Y = 1|A = a, \hat{Y} = 0) = P(Y = 1|A = b, \hat{Y} = 0)$$

Equal Positive Predictive Value $\text{PPV} = \text{TP} / (\text{TP} + \text{FP})$ for each subgroup and

equal Negative Predictive Value $\text{NPV} = \text{NP} / (\text{NP} + \text{FN})$ for each subgroup

True class is independent from protected attribute given predicted class

	$Y = 1$	$Y = 0$	
$\hat{Y} = 1$	TP	FP	P
$\hat{Y} = 0$	FN	TN	N
	π_1	π_0	

Recurring Nightmares: Bias

COMPAS case

Demographic parity:

frequency of convicted by the court equal for each subpopulation

fair from society's perspective

Equal opportunity:

frequency of convicted among innocents equal for all subpopulations

fair from the prisoner's perspective (ProPublica)

Predictive Rate Parity:

frequency of innocents among convicts is equal for subpopulations

fair from the judge's perspective (Northpointe)

	$Y = 1$	$Y = 0$	
$\widehat{Y} = 1$	TP	FP	P
$\widehat{Y} = 0$	FN	TN	N
	π_1	π_0	

	$Y = 1$	$Y = 0$	
$\widehat{Y} = 1$	TP	FP	P
$\widehat{Y} = 0$	FN	TN	N
	π_1	π_0	

	$Y = 1$	$Y = 0$	
$\widehat{Y} = 1$	TP	FP	P
$\widehat{Y} = 0$	FN	TN	N
	π_1	π_0	

Recurring Nightmares: Bias

COMPAS case

Demographic parity:

frequency of convicted by the court equal for each subpopulation

fair from society's perspective

Equal opportunity:

frequency of convicted and released for all subpopulations

fair from the prisoner's perspective

Predictive Rate Parity:

frequency of innocents among convicts is equal for subpopulations

fair from the judge's perspective (Northpointe)

The fairness trade-off (fairml.org)

Except in trivial situations, you cannot control all three criteria.

In most cases, you can't even control two of them.

	$Y = 1$	$Y = 0$	
$\widehat{Y} = 1$	TP	FP	P
$\widehat{Y} = 0$	FN	TN	N
π_0			
$= 0$			
FP			P
TN			N
π_0			
$= 0$			
$\widehat{Y} = 1$	TP	FP	P
$\widehat{Y} = 0$	FN	TN	N
π_1			
π_0			

Recurring Nightmares: Bias



arXiv.org > stat > arXiv:2104.00507

Search...

Help | Advanced

Statistics > Machine Learning

[Submitted on 1 Apr 2021]

fairmodels: A Flexible Tool For Bias Detection, Visualization, And Mitigation

Jakub Wiśniewski, Przemysław Biecek

Machine learning decision systems are getting omnipresent in our lives. From dating apps to rating loan seekers, algorithms affect both our well-being and future. Typically, however, these systems are not infallible. Moreover, complex predictive models are really eager to learn social biases present in historical data that can lead to increasing discrimination. If we want to create models responsibly then we need tools for in-depth validation of models also from the perspective of potential discrimination. This article introduces an R package fairmodels that helps to validate fairness and eliminate bias in classification models in an easy and flexible fashion. The fairmodels package offers a model-agnostic approach to bias detection, visualization and mitigation. The implemented set of functions and fairness metrics enables model fairness validation from different perspectives.

The package is designed not only to examine a single model

Comments: 15 pages, 9 figures

Subjects: Machine Learning (stat.ML); Machine Learning (cs.LG)

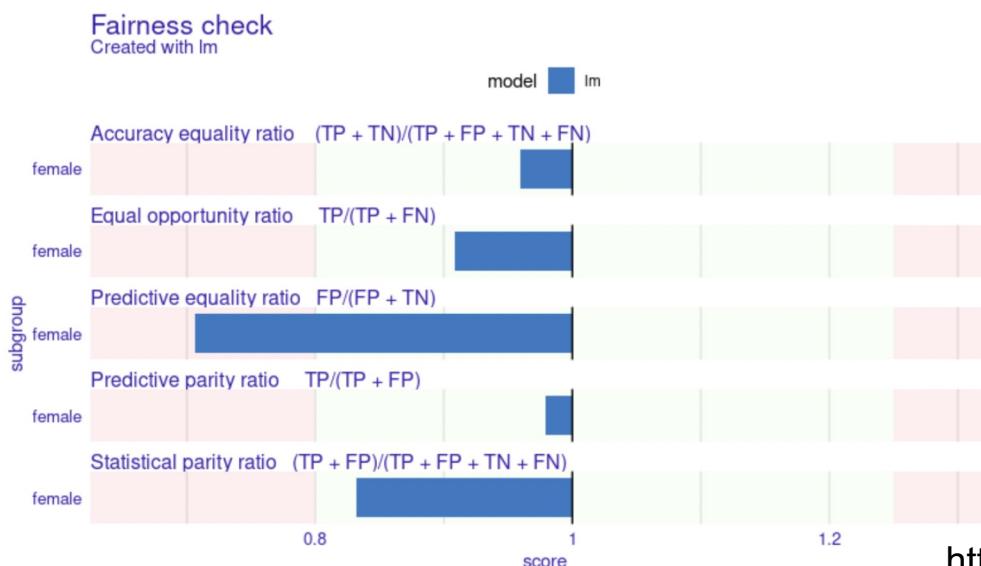
Cite as: arXiv:2104.00507 [stat.ML]

(or arXiv:2104.00507v1 [stat.ML] for this version)

Submission history

From: Przemysław Biecek [view email]

[v1] Thu, 1 Apr 2021 15:06:13 UTC (1,920 KB)



Important but overlooked aspects

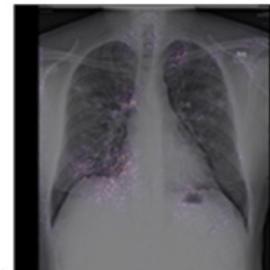
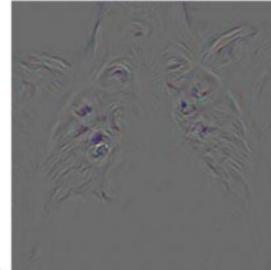
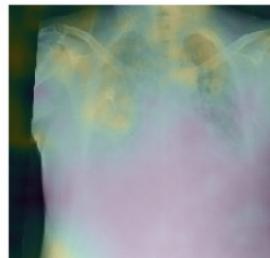
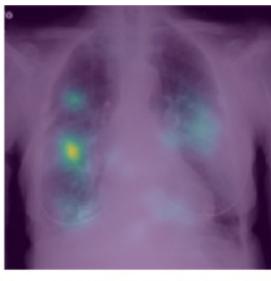
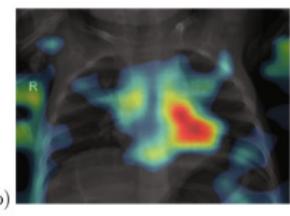
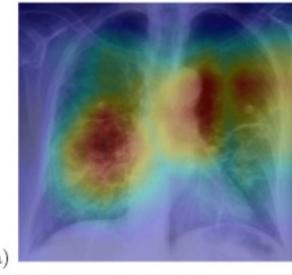
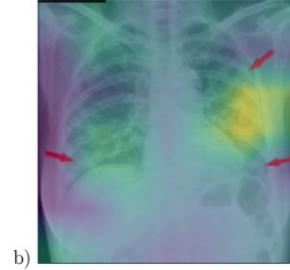
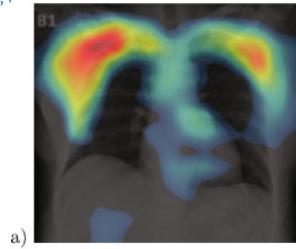
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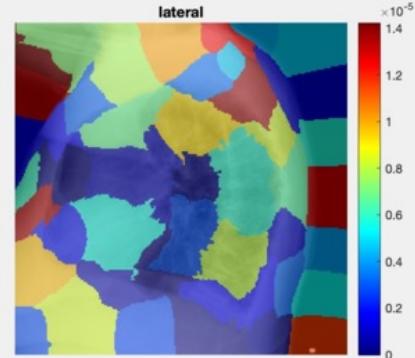
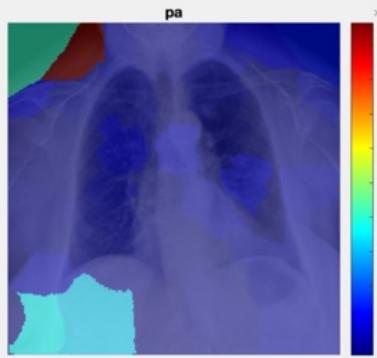
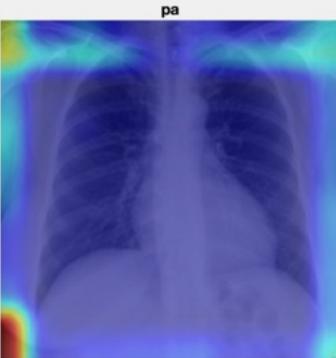
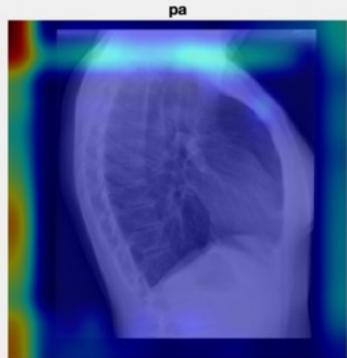
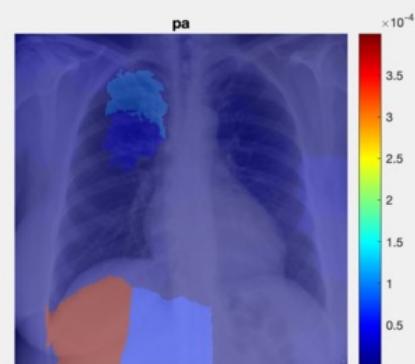
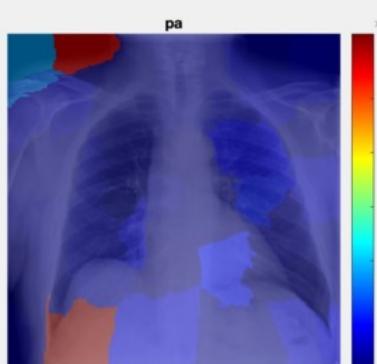
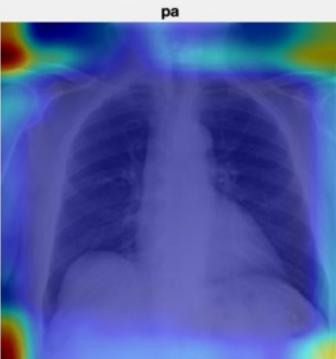
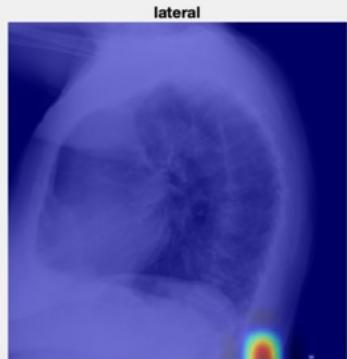
Checklist for responsible deep learning modeling of medical images based on COVID-19 detection studies

Weronika Hryniewska ^a, Przemysław Bombiński ^b, Patryk Szatkowski ^b, Paulina Tomaszewska ^a, Artur Przelaskowski ^a, Przemysław Biecek ^{a, i}



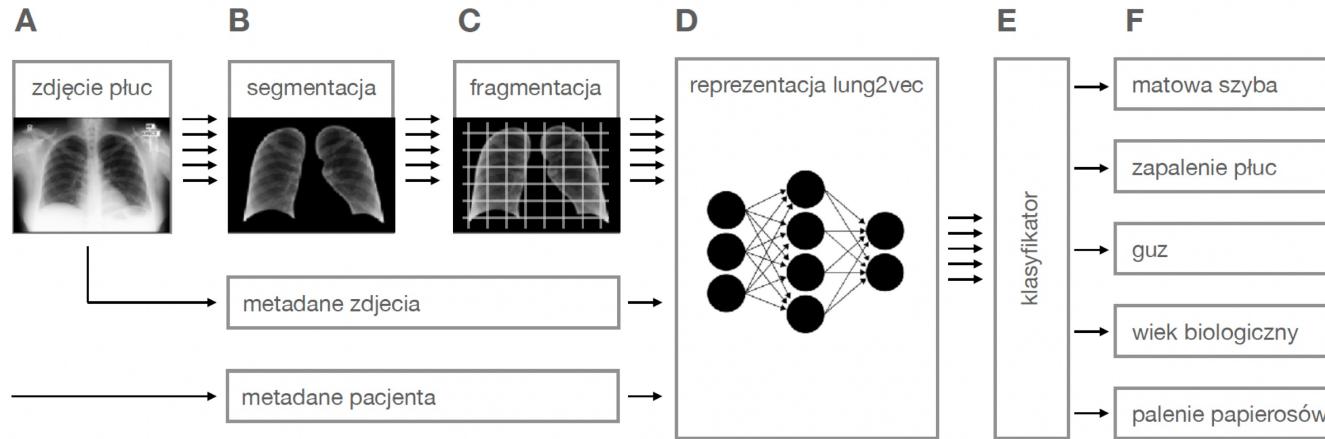
Explain!

MI × LUNGS

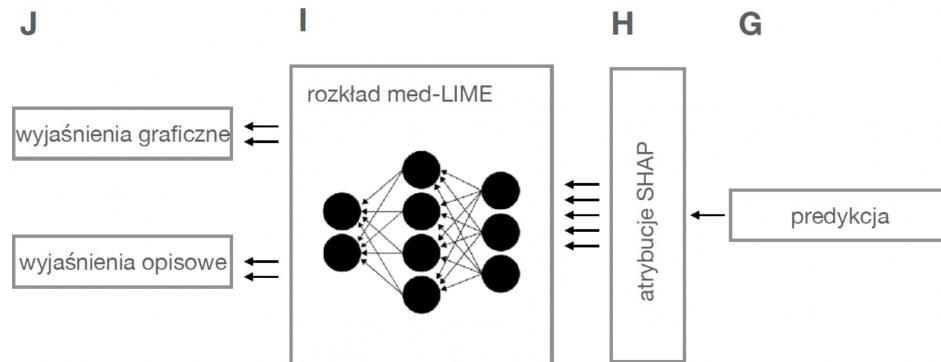


Explain!

Moduł predykcyjny



Moduł wyjaśnienia



Explain!

[Submitted on 15 Nov 2021 (v1), last revised 30 Apr 2022 (this version, v2)]

LIMEcraft: Handcrafted superpixel selection and inspection for Visual Explanations

Weronika Hryniwska, Adrianna Grudzień, Przemysław Biecek

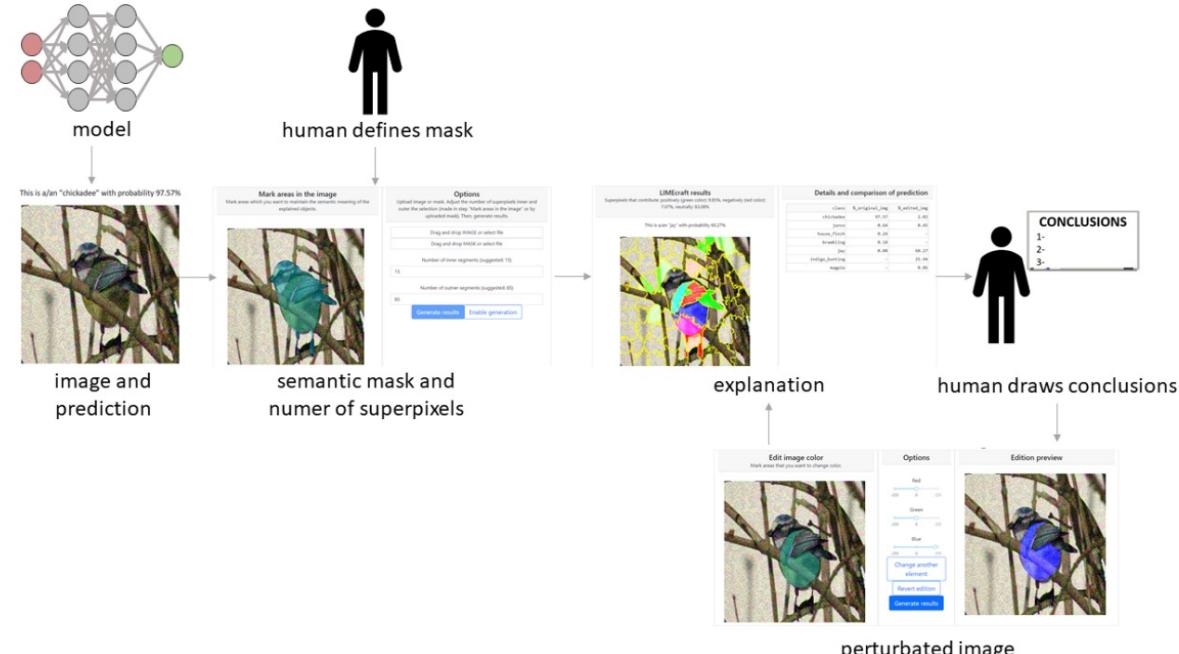
The increased interest in deep learning applications, and their hard-to-detect biases result in the need to validate and explain complex models. However, current explanation methods are limited as far as both the explanation of the reasoning process and prediction results are concerned. They usually only show the location in the lack of possibility to interact with explanations makes it difficult to verify and creates a significant risk when using the model. The risk is compounded by the semantic meaning of the explained objects. To escape from the trap of a tool and a process called LIMEcraft. LIMEcraft enhances the process of explaining semantically consistent areas and thoroughly examine the prediction for the Experiments on several models show that our tool improves model safety by may indicate model bias. The code is available at: [this http URL](#)

Subjects: Computer Vision and Pattern Recognition (cs.CV); Machine Learning (cs.LG)

Cite as: [arXiv:2111.08094 \[cs.CV\]](#)

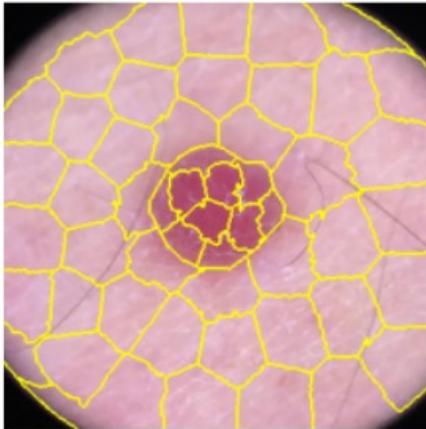
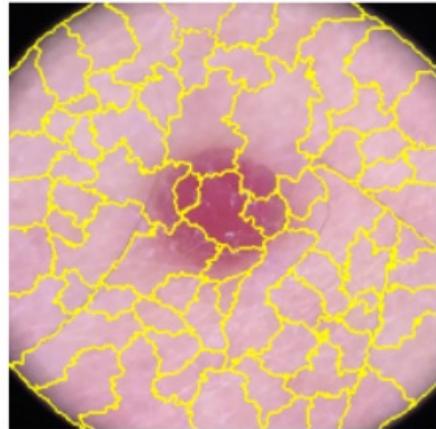
(or [arXiv:2111.08094v2 \[cs.CV\]](#) for this version)

<https://doi.org/10.48550/arXiv.2111.08094> ⓘ



Explain!

MI × LUNGS



Try this yourself
<https://limecraft.mi2.ai/>

Explain!

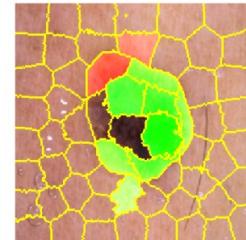
MI × LUNGS



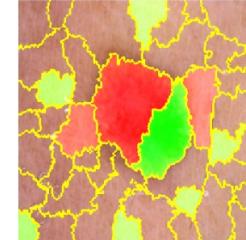
(a) Skin lesion



(b) Selected mask



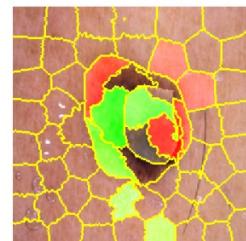
(c) LIMEcraft results
melanocytic nevi 54.79%



(d) LIME results



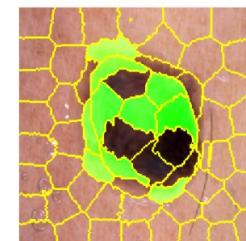
(e) Color edition



(f) LIMEcraft for (e) benign
keratosis-like 99.80%



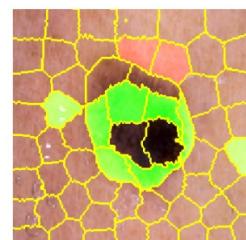
(g) Shape edition



(h) LIMEcraft for (g)
melanoma 52.86%



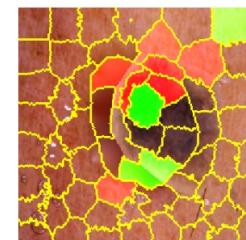
(i) Hair removal



(j) LIMEcraft for (i)
melanoma 66.36%



(k) Color, rotation and loca-
tion edition



(l) LIMEcraft for (k) benign
keratosis-like 98.82%

Try this yourself

<https://limecraft.mi2.ai/>

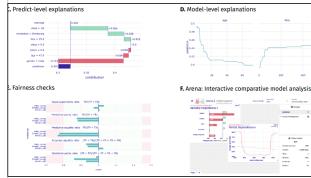
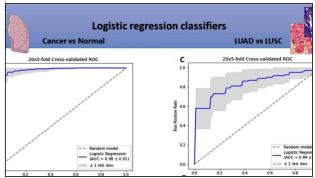
Important but overlooked aspects



To build a good AI system it is needed to take care of:

- high quality **data**
- methods of reducing undesired **bias**
- methods of exploring and **explaining** models
- many other topics that I will not talk about today

↓ Papers



A Signature of 14 Long Non-Coding RNAs (lncRNAs) as a Step towards Precision Diagnosis for NSCLC[†]

Anetta Sulewska, Jacek Niklinski, Radoslaw Charkiewicz, Piotr Karabowicz, Przemyslaw Biecek, Hubert Baniecki, Oksana Kowalczuk, Miroslaw Kozlowski, Patrycja Modzelewska, Piotr Majewski et al.

dalex: Responsible Machine Learning with Interactive Explainability and Fairness in Python[†]

Hubert Baniecki, Wojciech Kretowicz, Piotr Piątyszek, Jakub Wiśniewski, Przemysław Biecek

Journal of Machine Learning Research (2021)

<https://www.mi2.ai/>

RadioTator

Filters

With categories
Without categories
All documents

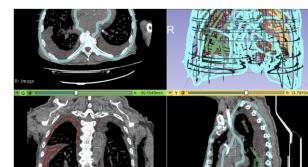
Text ID: right lung Text length: All

Filter: And Or With final annotations
Normal Without final annotations
substring word Regex positive negative

Result number: 0-5/200

Save annotations

Annotations: Cardiomegaly, Lesion, Pleural Effusion, Interesting case. Must check later



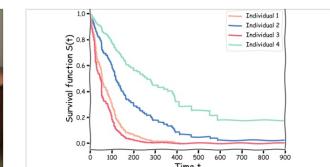
<https://medium.com/responsibleml>

RadioTator: A Tailored Tool for Rapid Medical Text Annotation

Creating an annotation app for text data to use for radiologists.

Jakub Wiśniewski

Jun 14 · 4 min read



Towards the largest database of Polish lung medical images

In healthcare is gaining more at more attention. In MI2DataLab, we work on this area in the project: "xLungs—Responsible Artificial...

Paulina Tomaszewska

May 10 · 2 min read

Responsible Machine Learning for Survival Analysis

A brief introduction to survival analysis and the use of machine learning models in this area

Mateusz Krzyżniński

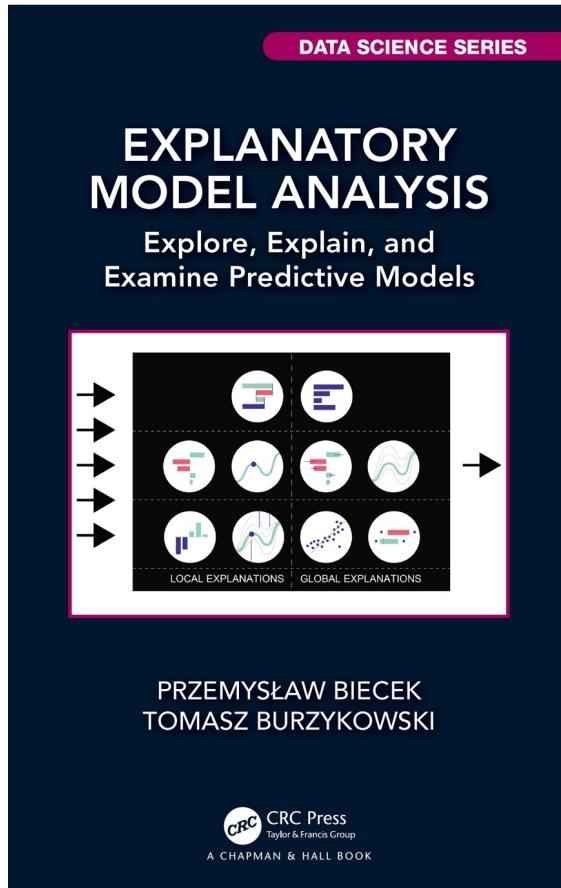
Jan 19 · 9 min read

Paulina Tomaszewska

Apr 14 · 2 min read

More

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<https://ema.drwhy.ai/>



<https://github.com/BetaAndBit/RML>