

Supplementary Material

Explainable Artificial Intelligence Models Using Real-World Electronic Health Record Data: a Systematic Scoping Review

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Appendix Table 1. Inclusion and exclusion criteria for the systematic review

| Type | Criteria | Rationale |
|-----------|--|--|
| Inclusion | XAI in medicine and healthcare | XAI can be applied to many different applications, but the focus of our review is in medicine and healthcare, since this area is one of the most critical applications of AI and XAI. |
| | The AI model is built based on EHR data | In this review, we focus on predictive modeling using EHR data. |
| | They study is not older than 2009 | We are interested in XAI in medicine in the past decade. Also, emergence of new generation of complex AI models like deep learning, makes the XAI challenge more significant. |
| | The study's focus is on XAI, thus a relevant term is used in the title and/or abstract | The goal of the study is on XAI and the relevant terms include: explainable, explainability, interpretable, interpretability, understandable, understandability, comprehensible, comprehensibility, intelligible, machine learning, artificial intelligence, prediction model, predictive model, deep learning, AI, neural network |
| Exclusion | Details of the paper is not available | Abstract papers, or the papers that could not be accessed through the university library or the interlibrary loan, and system demonstrations are not included. |
| | Unpublished | We excluded papers that were uploaded on arXiv or other archiving systems but have not been published in a peer-reviewed venue. |
| | Opinion or other review papers | This systematic review is not a review of reviews, also opinion papers do not fulfill the requirement of delivering an XAI method. |
| | Duplicated | Usually querying multiple databases returns similar papers, thus we removed the duplicates. |

Appendix Table 2. Reproducibility analysis of the articles. The source codes are either available and explicitly referred to in the article (marked as “√”) or not (marked as “-”). The datasets are either proprietary (marked as “-”) or not (marked as “√”). If the provided link to the dataset and/or the source code was broken or there was no dataset and/or source code available in the referred link to the host website, we also marked it as not available “-”.

| Article | Source code is accessible | Source code repository | Dataset is accessible | Dataset |
|------------------------------|---------------------------|---|-----------------------|---|
| Das et.al, 2019 [1] | √ | GitHub | √ | ADNI dataset |
| Knijnenburg et al., 2016 [2] | √ | GitHub | √ | Data provided in the supplementary material of the article, the validation dataset: drug sensitivity dataset from the Cancer Therapeutics Response Portal |
| Xiao et al., 2018 [3] | √ | GitHub | - | A real-world EHR repository of Congestive Heart Failure cohort and simulated EHR data |
| Choi et al., 2016 [4] | √ | GitHub | - | Sutter Health EHR data |
| Hu et al., 2019 [5] | √ | GitHub | √ | Retrovirus Integration Database |
| Hao et al., 2018 [6] | √ | GitHub | √ | The Cancer Genome Atlas (TCGA) |
| Bernardini et al., 2019 [7] | √ | Code Ocean | √ | Federazione Italiana Medici di Medicina Generale (FIMMG) dataset ¹ |
| Zhang et al., 2015 [8] | √ | Downloadable from the Journal’s website | √ | SIDER database |
| Aditya and Pande, 2017 [9] | √ | Downloadable from the Journal’s website | √ | Open Access Series of Imaging Studies (OASIS) |
| Fejza et al., 2018 [10] | - | N/A | √ | EHR data from the Premier healthcare database |
| Yoon et al., 2018 [11] | - | Journal website ² | √ | United Network for Organ Sharing (UNOS) database |

| | | | | |
|----------------------------------|---|-----------------------|---|--|
| Jalali and Pfeifer, 2016 [12] | ✓ | GitHub | ✓ | The Cancer Genome Atlas (TCGA) |
| Kaji et al., 2019 [13] | ✓ | GitHub | ✓ | MIMIC-III |
| Brisimi et al., 2018 [14] | - | N/A | - | Made available by the Boston Medical Center |
| Eck et al., 2017 [15] | ✓ | Sourceforge | ✓ | Data generated by IS-pro technique, made available on SourceForge |
| Settouti et al., 2012 [16] | - | N/A | ✓ | Pima Indian Dataset in UCI repository |
| Davoodi and Moradi, 2018 [17] | - | N/A | ✓ | MIMIC-III, Diabetes (PID), Heart Disease (HD) and Liver datasets from UCI repository |
| Kwon et al., 2018 [18] | - | GitHub | ✓ | Health Insurance Review and Assessment Service (HIRA) National Patients Sample |
| Shickel et al., 2019 [19] | - | N/A | ✓ | MIMIC-III, University of Florida Health Hospital EHR data |
| Pan et al., 2019 [20] | - | N/A | - | Endocrinology Department at Guangzhou Women and Children's Medical Center EHR data |
| Huang, 2009 [21] | - | N/A | - | Research data not available |
| Van den Bulcke et al., 2011 [22] | - | N/A | - | Data from a systematic screening for newborns by the PCMA screening center (Belgium) |
| Bouktif et al., 2014 [23] | ✓ | Institutional website | ✓ | Three heart disease datasets in UCI repository |

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|--|---|---|---|--|
| Ponce and Martinez-Villaseñor, 2017 [24] | - | N/A | ✓ | Wisconsin Breast Cancer Dataset in UCI repository |
| Crielaard and Papapetrou, 2018 [25] | - | N/A | - | Stockholm electronic patient record (EPR) corpus (HealthBank) |
| Ming et al., 2019 [26] | ✓ | GitHub | ✓ | Wisconsin Breast Cancer Dataset and Pima Indian Diabetes Dataset in UCI repository |
| Hajiloo et al., 2013 [27] | - | N/A | ✓ | Prostate cancer dataset and colon cancer dataset in other publications |
| Luo, 2016 [28] | - | N/A | - | Practice Fusion Diabetes Classification |
| Zhang et al., 2018 [29] | ✓ | GitHub | - | De-identified EHR data from the University of Virginia Health System |
| Hartono, 2018 [30] | - | N/A | - | Cancer microarray data sets obtained from Gene Expression Model Selector ³ |
| Zhao and Bolouri, 2016 [31] | - | N/A | ✓ | The Cancer Genome Atlas (TCGA) |
| Kim et al., 2016 [32] | - | N/A | ✓ | NIH Epigenome Roadmap, NHGRI, ENCODE |
| Valdes et al., 2016 [33] | ✓ | Dedicated website (https://www.mediboostml.com/) | ✓ | 13 data sets corresponding to all binary classification problems in the field of life sciences within the UCI Repository |
| Stiglic et al., 2012 [34] | ✓ | Open source Weka package | ✓ | 40 datasets in the UCI Repository, eSol database, 9 Gene Expression Machine Learning Repository (GEMLeR) datasets |

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|---------------------------------|---|-----------------|---|--|
| Zuallaert et al., 2018 [35] | ✓ | Project website | - | Datasets in other publications |
| Ghafouri-Fard et al., 2019 [36] | - | N/A | - | Proprietary genotyping data |
| Paredes et al., 2018 [37] | - | N/A | - | Portuguese dataset of ACS patients |
| Barakat et al., 2010 [38] | - | N/A | - | A diabetes dataset in Oman |
| Park et al., 2018 [39] | - | N/A | - | Seoul National University Bundang Hospital EHR data |
| Ge et al., 2018 [40] | - | N/A | - | Asan Medical Center EHR data |
| Lakkaraju et al., 2016 [41] | - | N/A | - | Medical diagnosis records of about 150K patients collected by a web-based EHR company and two other non-medical datasets |
| Che et al., 2017 [42] | - | N/A | - | Pediatric ICU dataset collected at the Children's Hospital Los Angeles. |

¹ <http://vrai.dii.univpm.it/content/fimmg-dataset>

² Only the pseudo code is available

³ No longer available

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