# REVIEW





# Automated machine learning with interpretation: A systematic review of methodologies and applications in healthcare

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## **Abstract**

Machine learning (ML) has achieved substantial success in performing healthcare tasks in which the configuration of every part of the ML pipeline relies heavily on technical knowledge. To help professionals with borderline expertise to better use ML techniques, Automated ML (AutoML) has emerged as a prospective solution. However, most models generated by AutoML are black boxes that are challenging to comprehend and deploy in healthcare settings. We conducted a systematic review to examine AutoML with interpretation systems for healthcare. We searched four databases (MEDLINE, EMBASE, Web of Science, and Scopus) complemented with seven prestigious ML conferences (AAAI, ACL, ICLR, ICML, IJCAI, KDD, and NeurIPS) that reported AutoML with interpretation for healthcare before September 1, 2023. We included 118 articles related to AutoML with interpretation in healthcare. First, we illustrated AutoML techniques used in the included publications, including automated data preparation, automated feature engineering, and automated model development, accompanied by a real-world case study to

Abbreviations: AAAI, annual AAAI conference on artificial intelligence; ACL, annual meeting of the association for computational linguistics; AutoML, automated machine learning; ANN, artificial neural networks; AUPRC, area under the precision recall curve; AUROC, area under the receiver operating characteristic curve; BMI, brain machine interfaces; CNN, convolutional neural networks; ChIP-seq, chromatin immunoprecipitation sequences; DNA-seq, DNA sequences; DNase-seq, DNase I hypersensitive site sequences; DSC, dice similarity coefficient; EMG, electromyogram; ECG, electrocardiogram; FE, feature engineering; fMRI, functional magnetic resonance imaging; GMM, Gaussian mixture model; GAN, generative adversarial network; GAT, graph attention network; GBM, gradient boosting machine; GNB, Gaussian Naive Bayes; GRU, gated recurrent unit; HD, hausdorff distance; HLAN, hierarchical label-wise attention network; ICD, international classification of diseases; IB, information bottleneck; ICLR, international conference on learning representations; ICML, international conference on machine learning; IJCAI, international joint conference on artificial intelligence; IOU, intersection over union; KDD, ACM SIGKDD conference on knowledge discovery & data mining; LDA, linear discriminant analysis; LSTM, long short-term memory; LR, logistic regression; LASSO, least absolute shrinkage and selection operator; MDL, minimum description length; ML, machine learning; MLP, multilayer perceptron; MSE, mean squared error; MNase-seq, micrococcal nuclease digestion with deep sequencing; NeurIPS, annual conference on neural information processing systems; OCT, optical coherence tomography; PRISMA, preferred reporting items for systematic reviews and meta-analyses; PSD, predictive sparse decomposition; PSNR, peak signal-to-noise ratio; PNN, probabilistic neural networks; R-CNN, region convolutional neural networks; RF, random forest; RMSE, root mean squared error; RNA-seq, RNA sequences; RNN, recurrent neural networks; SVM, support vector machine; SHAP, shapley additive explanations; TPOT, tree-based pipeline optimization tool; VAE, variational autoencoder.

Han Yuan, Kunyu Yu and Feng Xie are contributed equally.

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demonstrate the advantages of AutoML over classic ML. Then, we summarized interpretation methods: feature interaction and importance, data dimensionality reduction, intrinsically interpretable models, and knowledge distillation and rule extraction. Finally, we detailed how AutoML with interpretation has been used for six major data types: image, free text, tabular data, signal, genomic sequences, and multi-modality. To some extent, AutoML with interpretation provides effortless development and improves users' trust in ML in healthcare settings. In future studies, researchers should explore automated data preparation, seamless integration of automation and interpretation, compatibility with multi-modality, and utilization of foundation models.

## KEYWORDS

automated machine learning, explainable machine learning, health informatics

# 1 | INTRODUCTION

The rapid growth of biomedical big data has led to greater opportunities for the deployment of modern data-driven technologies such as machine learning (ML) [1]. ML techniques have achieved substantial success in processing various types of data and performing diverse tasks in the context of healthcare [2–4]. However, the effective exploitation of healthcare data by ML models necessitates the rigorous configuration of every part of the ML pipeline, which relies heavily on specialized technical knowledge and extensive effort.

To help professionals with borderline expertise in data science to better use ML techniques, Automated ML (AutoML) has emerged as a prospective solution. The objective of AutoML, as defined by Yao et al. [5], is to allow computer programs to replace human tuning in the process of determining all or a part of model configurations while maintaining good performance and high computational efficiency. Configurations in this context refer to all factors that are specified prior to model training and affect the final performance, including input data, feature sets, hyperparameters, and model architectures. Therefore, a complete AutoML pipeline encompasses the automation of data preparation, feature engineering, and model development [6].

AutoML techniques in the ML pipeline cater to various levels of coding proficiency. For example, sophisticated AutoML methods such as NASLib [7], which require advanced programming knowledge, aim to provide greater flexibility for experienced ML engineers. AutoML software packages such as auto-sklearn [8] focus primarily on model development, that is, algorithm selection and hyperparameter optimization, targeting users with mediate coding skills. Additionally, commercial AutoML platforms such as Google Cloud's AutoML

system and H2O Driverless artificial intelligence (AI) offer no-coding solutions, featuring user-friendly interfaces and rapid convergence capabilities. Table 1 provides an overview of the toolkits developed by leading companies.

In healthcare, the extensive application of ML significantly amplifies the advantages of implementing AutoML approaches. It enables healthcare professionals with borderline ML knowledge to build high-quality models using a fully automated pipeline [9] and further addresses privacy concerns without sharing data with external ML engineers. AutoML systems effectively fill the gap between the lack of ML expertise among healthcare practitioners and the need for data analytics based on ML models [10]. AutoPrognosis [11] describes an end-to-end diagnosis and prognosis modeling framework that helps healthcare professionals leverage clinical data for risk prediction across diverse clinical settings. Additionally, AutoML improves the efficiency of ML engineers by automating tedious and time-consuming tasks such as data preprocessing [12]. For example, nnU-Net [13] introduces a self-configured biomedical image segmentation method that automates the conversion of raw image data into representative structured features.

Although AutoML systems help both healthcare professionals and ML engineers to process medical data effortlessly, the interpretability of these systems should be improved to boost confidence in the reliability of the generated ML models [14]. Given the potentially serious consequences of medical AI failures, greater demands are being placed on the interpretation of ML models in clinical decision-making to fulfill both medical validation and regulatory requirements. Thus, in contrast to conventional AutoML systems primarily centered on ML development, AutoML with interpretation aligns more closely with the real-world requirements in healthcare settings [15].

TABLE 1 Overview of AutoML toolkits from leading companies.

Company	Toolkit	Modality	Website
Amazon	AutoGluon	Multi- modality	https://auto.gluon.ai/stable/index.html
	SageMaker	Multi- modality	https://aws.amazon.com/sagemaker/canvas/
Apple	Create ML	Multi- modality	https://developer.apple.com/machine-learning/create-ml/
Google	Vertex AI	Multi- modality	https://cloud.google.com/vertex-ai?hl=en
IBM	AutoAI	Tabular data	https://www.ibm.com/products/watson-studio/autoai
	watsonx.ai	Multi- modality	https://www.ibm.com/products/watsonx-ai
Meta	Looper	Multi- modality	https://research.facebook.com/publications/looper-an-end-to-end-ml-platform-for-product-decisions/
Microsoft	Azure machine learning	Multi- modality	https://azure.microsoft.com/en-us/products/machine-learning/automatedml/#overview
NVIDIA	TAO	Multi- modality	https://developer.nvidia.com/tao-toolkit

Note: The companies are listed alphabetically for ease of reference.

Because AutoML systems with interpretation are fundamental to facilitating the clinical adoption of AI technologies, we conducted this review to gain insight into how they empower the health community by lowering the entry barrier and enhancing the credibility of ML algorithms. In recent years, several researchers [6, 9, 10, 16-19] have reviewed the development and application of either AutoML or ML interpretations. However, none have provided a systematic and in-depth summary of AutoML with interpretation, particularly its applications in healthcare. In our review, we aim to integrate existing research practices by categorizing data modalities, AutoML techniques, and interpretation methods to acquire a comprehensive understanding of AutoML with interpretation in healthcare and inspire future research topics. The purpose of the categorization is to provide practitioners with an insight into how various AutoML with interpretation systems have been implemented in different medical tasks.

The promising application of AutoML with interpretation in healthcare necessitates a systematic review of cutting-edge research to bridge the gap between technical innovation and practical application. We envisage that this review will empower healthcare practitioners by providing well-organized and referable information about AutoML with interpretation systems, and further facilitate the real-world deployment of ML systems in diverse healthcare settings.

# **METHODS**

#### Search strategy and data sources 2.1

We conducted a systematic review that encompassed both methodology and application studies on AutoML with interpretation for healthcare. We performed a literature search on four databases: MEDLINE, EMBASE, Web of Science, and Scopus. Given that some of the latest ML research is often presented at conferences and may not be included in these four databases, we also searched for research papers in the proceedings of seven relevant and prestigious ML conferences: AAAI, ACL, ICLR, ICML, IJCAI, KDD, and NeurIPS. The searched terms in the medical domain were ("medical" OR "clinical" OR "health" OR "healthcare" OR "medicine"). We also added the terms ("ML" OR "deep learning" OR "AI") to limit the search to ML-based studies, and ("automated" OR "automatic") AND ("interpretable" OR "explainable" OR "interpretability") to include studies on AutoML with interpretation. We restricted our search to papers published before September 1, 2023.

#### 2.2 Inclusion and exclusion criteria

We followed the Preferred Reporting Items for Systematic reviews and Meta-Analyses guidelines [20] to conduct the

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systematic review. We included all papers published in English that used AutoML with interpretation to perform healthcare tasks. We excluded review articles, workshop papers, duplicate records, and studies not relevant to AutoML with interpretation or healthcare. Each article was independently screened by at least two reviewers, and, if ambiguous, discussed with the corresponding author to reach a consensus.

# 2.3 | Data analysis

Table 2 presents our evaluation and summary of the papers from three aspects: AutoML techniques, interpretation methods, and target data types. For AutoML techniques, we identified three main research directions: automated data preparation, automated feature engineering, and automated model development [9]. For interpretation methods, we summarized from four angles: knowledge distillation and rule extraction, intrinsiinterpretable models, data dimensionality reduction, and feature interaction and importance [18, 19]. For target data types, we classified the included articles into six categories: image, free text, tabular data, signal data, genomic sequence, and multi-modality. Additionally, Table 2 lists specific applications and performance advantages of AutoML for users focused on specific tasks.

## 3 | RESULTS

Figure 1 illustrates the literature selection process for this systematic review. Our initial search yielded 2730 papers. We removed 1378 duplicates; hence, we used 1352 records for title and abstract screening. We excluded 1184 records because they were either not relevant to healthcare (n = 331) or did not use AutoML methods (n = 722); were conference papers that were not from listed conferences (n = 9); were not research articles (n = 121); or were not in English (n = 1). As a result, we included 168 articles for full-text review. Finally, we included 118 papers for systematic review. Figure 2 shows the rising trend of publications in AutoML with interpretation for healthcare and indicates that image and tabular data constituted the major subsets for all included publications. In this section, we first summarize AutoML techniques. Then, we elaborate on the ML interpretations used in the included articles. Finally, we summarize the representative AutoML with interpretation systems for different data modalities.

# 3.1 | AutoML techniques

For AutoML techniques, we followed the previous classification criteria [9] based on three stages of the ML pipeline: automated data preparation (n = 18), automated feature engineering (n = 95), and automated model development (n = 31). Figure 3 provides a comprehensive overview and Table 3 offers a detailed description of the ML components automated by AutoML within the healthcare sector. Specifically, data preparation refers to the process of collecting and processing raw data into a suitable format for downstream ML stages. AutoML has been leveraged to deal with processes such as automatic data collection [96, 106], noise filtering [27, 28, 44, 119], missing value imputation [87, 95, 110, 126, 133], data imbalance compensation [87, 90, 102, 140], data normalization [44], redundant data removal [53], outlier removal [133], sample clustering [135], data pattern shift detection [137], and continuous variable binning [109]. Feature engineering describes the process of creating new features or modifying existing features to enhance ML performance and AutoML has been used to facilitate automatic feature generation [21, 60, 61, 63, 64, 66, 68, 71, 72, 76, 77, 79–81, 103, 120, 122, 123], selection [70, 72, 80, 98, 99, 101, 102, 105, 108, 121, 124, 127, 135, 141], and transformation [67, 78, 107, 138, 142, 143]. Model development refers to the process of creating, training, and optimizing a model based on either the formatted data or modified features. AutoML has also been used for the selection of main backbone models [65, 86, 88, 89, 91-93, 125, 144], the tuning of model-specific parameters [24, 98, 100, 119, 126, 128, 136], and the optimization of modelspecific [21, 24, 40, 59, 74, 86, 89–93, 102, 110, 122, 124, 131, 138, 144] or agnostic hyperparameters [24, 62, 69, 95].

Additionally, we conducted a comparative analysis of commonly used metrics between AutoML and the most competitive baseline in the last column of Table 2, which demonstrated that AutoML outperformed conventional ML solutions across various data types. Specifically, slashes ("/") divide AutoML performance and the most competitive baseline performance. Hyphens ("-") indicate that specific results were not reported in the original papers. Ampersands ("&") separate the same evaluation metrics across different tasks or experimental settings and commas (",") separate different evaluation metrics. We retained all measurement units and decimal digits from the original papers. We did not report results from studies in which visual performance comparisons were made without quantitative data or from studies involving an excessive number of tasks because of content constraints.

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TABLE 2 Information summary of the included studies on AutoML with interpretation for healthcare.

Data	Year	Year Paper	Automated data preparation	Automated feature engineering	Automated model development	Interpretability methods	Model name	Main ML architectures	Healthcare applications	Performance comparison
Image	2023	Alkhalaf et al. [21]		>	>	Feature interaction and importance	AAOXAI-CD	CNN, RNN, GRU, LSTM	Cancer classification	Accuracy: 99.00/97.00 & 99.42/98.43
Image	2023	Berghe et al. [22]		>		Feature interaction and importance		U-net, CNN	Structural lesions detection	Accuracy: 0.89/- & 0.92/-, AUROC: 0.92/- & 0.91/-
Image	2023	Cabon et al. [23]		>		Data dimensionality reduction		LR, RF, SVM	Functional age estimation	MAE: 1.4/- & 1.6/- & 1.3/-
Image	2023	Choi et al. [24]			>	Knowledge distillation and rule extraction	SimpleMind	CNN, U-net	Endotracheal tube assessment, kidney segmentation, prostate segmentation	Accuracy: 89/-, DSC: 0.881/0.878 & 0.842/ 0.818, HD: 46.4/30.7
Image	2023	Custode et al. [25]		>		Intrinsically interpretable models		U-net, CNN, decision tree	Lung status evaluation	
Image	2023	Dai et al. [26]		>		Feature interaction and importance	MS-net	CNN	Lung nodules assessment Accuracy: 92.4/90.0 & 88.5/87.3	Accuracy: 92.4/90.0 & 88.5/87.3
Image	2023	Gerbasi et al. [27]	>	>		Feature interaction and importance	<b>DeepMiC</b> a	CNN, U-net	Microcalcifications detection	Accuracy: 0.83/- & 0.83/-, AUROC: 0.95/- & 0.89/-, AUPRC: 0.78/-, IOU: 0.74/-
Image	2023	Ghassemi et al. [28]	>			Feature interaction and importance		GAN	COVID-19 classification	Accuracy: 89.24/88.05 & 98.25/94.69 & 96.20/94.69 & 99.2/ 89.6, AUROC: 97.22/ 96.71 & 99.79/99.03 & 99.43/99.43 & 99.95/99.60 & 99.95/99.99
Image	2023	2023 Jun et al. [29]		>		Feature interaction and importance		CNN, U-net	Noninvasive meningioma AUROC: 0.770/0.757, triaging DSC: 0.910/0.907 (Contii	AUROC: 0.770/0.757, DSC: 0.910/0.907 (Continues)

TABLE 2 (Continued)

Data type	Year Paper	Automated data preparation	Automated feature engineering	Automated model Interpret development methods	Interpretability methods	Model name	Main ML architectures	Healthcare applications	Performance comparison
Image	2023 Leong et al. [30]		>		Feature interaction and importance		Decision tree	Lung water content evaluation	AUROC: 0.719/- & 0.756/-
Image	2023 Orton et al. [31]		>		Data dimensionality reduction		LASSO	Molecular, histopathology and clinical target prediction	
Image	2023 Pham et al. [32]		>		Feature interaction and importance		U-net, CNN	Human epidermal growth factor receptor-2 classification	F1-score: 0.80/0.81
Image	2023 Saglam et al. [33]		>		Feature interaction and importance		XGBoost, SVM	Early onset schizophrenia classification	Accuracy: 0.80/0.78, AUROC: 0.85/0.83
Image	2023 Taşcı et al. [34]	34]	>		Feature interaction and importance	DGXAINet	CNN, SVM	Brain tumor classification	Accuracy: 98.42/95.75 & 99.96/98.91
Image	2023 Wang et al. [35]		>		Feature interaction and importance		CNN, U-net	Parkinson's disease classification	AUROC: 0.901/0.856
Image	2023 Xiang et al. [36]		>		Feature interaction and importance		CNN, GCN	Prostate cancer classification	Accuracy: 0.677/0.584, AUROC: 0.985/- & 0.986/-
Image	2023 Yoon et al. [37]	[37]	>		Feature interaction and importance		CNN	Anterior disc displacement classification	AUROC: 0.985/0.910 & 0.960/0.861
Image	2022 Yu et al. [38]	_	>		Feature interaction and importance		CNN, RF	Idiopathic pulmonary fibrosis prediction	AUROC: 0.987/-
Image	2022 Basso et al. [39]	[39]	>		Data dimensionality reduction		LDA, LR, RF	Glomerular disorder classification	Accuracy: 77/- & 87/-
Image	2022 Chen et al. [40]	[40]	>	>	Intrinsically interpretable models		R-CNN, U- net, LR	Blunt splenic injury triaging	Accuracy: 92/-, AUROC: 0.83/0.88

TABLE 2 (Continued)

Data type	Year Paper	Automated data preparation	Automated feature engineering	Automated model development	Interpretability methods	Model name	Main ML architectures	Healthcare applications	Performance comparison
Image	2022 Falco et al. [41]	ıl. [41]	>		Intrinsically interpretable models		Fuzzy rules	COVID-19 classification	Accuracy: 80.67/80.28
Image	2022 Kakileti et al. [42]		>		Knowledge distillation and rule extraction		V-net, RF	Early vascularity evaluation	AUROC: 0.85/0.79
Image	2022 Maqsood et al. [43]	_	>		Feature interaction and importance		CNN, SVM	Brain cancer prediction	Accuracy: 97.47/93.85 & 98.92/98.59
Image	2022 McCay et al. [44]	>	>		Knowledge distillation and rule extraction		LR, SVM, decision tree, LDA	Cerebral palsy prediction	Accuracy: 100/100 & 38/ 38 & 97.37/86.84
Image	2022 Mou et al. [45]	l. [45]	>		Feature interaction and importance	DeepGrading	CNN	Corneal confocal microscopy estimation	Accuracy: 84.10/82.40
Image	2022 Nafisah et al. [46]		>		Feature interaction and importance		CNN, U-net	Tuberculosis detection	Accuracy: 0.987/0.980, AUROC: 0.999/0.990
Image	2022 Nijiati et al. [47]	_	>		Feature interaction and importance		CNN, U-net	Active pulmonary tuberculosis classification	Accuracy: 0.910/0.895
Image	2022 Sharma et al. [48]		>		Feature interaction and importance		CNN, U-net	COVID-19 classification	Accuracy: 97.45/98.70, AUROC: 0.998/0.980
Image	2022 Park et al. [49]	l. [49]	>		Feature interaction and importance		U-net, LightGBM	Pilocytic astrocytomas classification	AUROC: 0.930/0.785
Image	2022 Sharma et al. [50]		>		Feature interaction and importance	COVID- MANet	CNN, U-net	COVID-19 classification	Accuracy: 97.37/97.16, IOU: 93.64/91.40, DSC: 96.70/95.49
Image	2022 Suri et al. [51]	. [51]	>		Feature interaction and importance	COVLIAS 2.0- cXAI	CNN, U-net	COVID-19 localization	Accuracy: 98.5/98.2, AUROC: 0.990/0.988

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TABLE 2 (Continued)

u u	98.5/99.4	Accuracy: 0.8961/0.9458	9.85/-	Accuracy: 91.2/- & 95.0/-	-/56	).962/0.986, 978/0.997	1.9460/0.9249,	AUROC: 0.9470/0.9797, DSC: 0.9796/0.9732, HD: 20.2249/41.0517	AUROC: 0.9470/0.9797, DSC: 0.9796/0.9732, HD: 20.2249/41.0517 Accuracy: 88.69/88.55 & 82.29/81.82 & 96.88/94.27	AUROC: 0.9470/0.9797, DSC: 0.9796/0.9732, HD: 20.2249/41.0517 Accuracy: 88.69/88.55 & 82.29/81.82 & 96.88/94.27 Accuracy: 98.13/87.69 & 98.13/87.69 & 98.26/98.18	9470/0.9732, HD: 0517 8.69/88.55 & & 96.88/94.27 8.13/87.69 & & 8 98.26/98.18
Performance comparison	n Accuracy; 98.5/99.4	Accuracy: 0	ion Accuracy: 0		AUROC 0.95/-	Accuracy: 0.962/0.986, AUROC: 0.978/0.997					
Healthcare applications	COVID-19 classification	Brain disease classification	Lung cancer classification Accuracy: 0.85/-	Bacterial pneumonia classification, COVID-19 classification	U-net, Gaussian Placenta health process prediction regression	Diabetic retinopathy triaging		COVID-19 classification	COVID-19 classificatio	COVID-19 classification COVID-19 detection Breast cancer classification	COVID-19 classification COVID-19 detection Breast cancer classification Retinal-vessel caliber measurement
Main ML architectures	GNB, SVM, decision tree, LR, KNN, RF	CNN, GRU	CNN, decision tree	R-CNN, K- means, SVM	U-net, Gaussian process regression	CNN		LR	LR	CNN	CNN CNN CNN
Model name					APPLAUSE				CovidNet3D	CovidNet3D	CovidNet3D
Interpretability methods	Feature interaction and importance	Feature interaction and importance	Intrinsically interpretable models	Knowledge distillation and rule extraction	Feature interaction and importance	Feature interaction and importance		Feature interaction and importance	Feature interaction and importance Feature interaction and importance	Feature interaction and importance Feature interaction and importance Data dimensionality reduction	Feature interaction and importance Feature interaction and importance Data dimensionality reduction Data dimensionality reduction
Automated model Interpret									>	>	>
Automated feature engineering	>	>	>	>	>	>		>	>	>	> >
Automated data preparation		>									
Year Paper	Ullah et al. [52]	Fu et al. [53]	Horry et al. [54]	Myeongkyun et al. [55]	Pietsch et al. [56]	Shorfuzzaman et al. [57]		Zhao et al. [58]	Zhao et al. [58] He et al. [59]	Zhao et al. [58]  He et al. [59]  Boumaraf et al. [60]	Zhao et al. [58]  He et al. [59]  Boumaraf et al. [60]  Cheung et al. [61]
Year	2022	2021	2021	2021	2021	2021	2021		2021	2021	2021 2021 2021
Data type	Image	Image	Image	Image	Image	Image	Image		Image	Image	Image

TABLE 2 (Continued)

										-IV/-LIADV	ANCES
Performance comparison	AUROC: 0.81/0.75	Accuracy: 90.22/88.88	Accuracy: 0.89/0.89 & 0.92/0.89, AUROC: 0.96/ 0.84 & 0.91/0.84	AUROC: 0.93/0.89	Accuracy: 0.946/0.895 & 0.917/0.878, AUROC: 0.918/0.845	Accuracy: 96.7/84.0, AUROC: 0.990/0.926	PSNR: 36.47/37.95 & 44.17/44.09	Accuracy: 72.43/-	AUROC: 0.884/0.721	DSC: 0.84/0.87 & 0.75/0.82	Accuracy: 78.15/75.46 (Continues)
Healthcare applications	Cardiac events prediction AUROC: 0.81/0.75	Leukocyte classification	Glioblastoma diagnosis	Early neoplasia classification	Congenital heart disease interpretation	Bladder cancer diagnosis	fMRI sensing	Breast cancer prediction	Prostate cancer recurrence prediction	Brain tumor segmentation, penumbra estimation	fMRI analysis
Main ML architectures	XGBoost	PSD, RF	CNN	CNN	CNN, RNN	PNN, SVM, LR, Adaboot, RF, MLP	Adaptive smoothing, game encoding	Decision tree	Autoencoder, LASSO, ridge regression, SVM	Boltzmann machine, RF	LASSO, elastic net
Model name		WBC-Profiler	TPOT, Auto- SkLearn								Remurs
Interpretability methods	Feature interaction and importance	Data dimensionality reduction	Data dimensionality reduction	Data dimensionality reduction	Feature interaction and importance	Data dimensionality reduction	Intrinsically interpretable models	Intrinsically interpretable models	Data dimensionality reduction	Feature interaction and importance	Data dimensionality reduction
Automated model			>				>	>			
Automated feature	>	>		>	>	>			>	>	>
Automated data preparation											=
Year Paper	Tamarappoo et al. [63]	l Yan et al. [64]	2020 Rucco et al. [65]	2020 Putten et al. [66]	) Wang et al. [67]	2020 Yin et al. [68]	2020 Lecouat et al. [69]	2019 Wu et al. [70]	Yamamoto et al. [71]	Pereira et al. [72]	7 Song et al. [73]
Year	2021	2021	2020	2020	2020	2020	2020	2019	2019	2018	2017
Data type	Image	Image	Image	Image	Image	Image	Image	Image	Image	Image	Image

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TABLE 2 (Continued)

Data type	Year	Year Paper	Automated data preparation	Automated feature engineering	Automated model Interpret development methods	Interpretability methods	Model name	Main ML architectures	Healthcare applications	Performance comparison
Free text	2021	2021 Diao et al. [74]		>	>	Feature interaction and importance		LightGBM	ICD coding	Accuracy: 95.2/91.3
Free text	2021	Kulshrestha et al. [75]		>		Feature interaction and importance		Elastic net, XGBoost, CNN	Chest injury prediction	AUROC: 0.93/-
Free text	2021	Blanco et al. [76]		>		Feature interaction and importance		GRU	Death cause extraction	AUROC: 53.3/52.1 & 49.4/58.8 & 58.2/62.0
Free text	2021	Dong et al. [77]		>		Feature interaction and importance	HLAN	GRU	Medical coding	AUROC: 88.4/88.3 & 94.5/96.9 & 88.5/90.2
Free text	2020	2020 Yang et al. [78]		>		Feature interaction and importance	AMFF	LSTM	Medical entity tagging	F1-score: 94.48/90.23 & 92.11/88.46 & 68.34/64.61 & 80.51/80.03
Free text	2020	2020 Li et al. [79]		>		Feature interaction and importance	MultiResCNN	CNN	ICD coding	F1-score: 0.073/0.068 & 0.608/0.584
Free text	2019	Atutxa et al. [80]		>		Feature interaction and importance		RNN, transformer	ICD coding	F1-score: 0.838/0.786 & 0.963/0.935 & 0.952/0.895
Free text	2018	Duarte et al. [81]		>		Feature interaction and importance		GRU	ICD coding	Accuracy: 89.320/79.802 & 81.349/70.754 & 76.112/67.404
Tabular data	2023	Li et al. [82]		>		Feature interaction and importance	FETCH	MLP	Hepatitis classification	F1-score: 0.9290/0.8839
Tabular data	2023	Junaid et al. [83]		>		Feature interaction and importance		SVM, RF, LightGBM	Parkinson's disease prediction	
Tabular data	2023	Islam et al. [84]		>		Data dimensionality reduction		LR, MLP, RF, XGBoost	Hypertension prediction	AUROC: 0.894/0.829

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Data	77	Automated data		Automated model	Interpretability		Main ML	Healthcare	Performance
Tabular data	2023 Wang et al. [85]			Knowledg distillation rule extra	Knowledge distillation and rule extraction		LR	Heart failure prediction	ACCUTACY: 0.999/0.995, AUROC: 0.981/0.979
Tabular data	2023 Zhang et al. [86]			>	Feature interaction and importance		LR, RF, GBM, MLP	Severe acute pancreatitis prediction	Accuracy: 0.910/0.920, AUROC: 0.907/0.849
Tabular data	2022 Agüero et al. [87]	>	>		Knowledge distillation and rule extraction		MLP, GRU, LSTM	Antimicrobial multidrug resistance prediction	Accuracy: 65.40/-, AUROC: 66.73/-
Tabular data	2022 Chou et al. [88]		>	>	Feature interaction and importance		XGBoost, RF, LR	Spinal cord injury prediction	AUROC: 0.68/-
Tabular data	2022 Cui et al. [89]			>	Feature interaction and importance		LR, RF, XGBoost, MLP, GBM	Early death prediction	Accuracy: 0.772/-, AUROC: 0.820/-
Tabular data	2022 Danilatou et al. [90]	>	>	>	Feature interaction and importance		LR, RF, SVM, decision tree	Mortality prediction	AUROC: 0.93/0.85 & 0.87/0.79
Tabular data	2022 Thongprayoon et al. [91]			>	Feature interaction and importance		RF, decision tree, XGBoost, MLP	Acute kidney injury prediction	Accuracy: 0.72/0.74, AUROC: 0.79/0.78
Tabular data	2022 Yin et al. [92]			>	Feature interaction and importance		RF, GBM, MLP, LR, XGBoost	Severe acute pancreatitis prediction	Accuracy: 0.953/0.943, AUROC: 0.945/0.898
Tabular data	2022 Yu et al. [93]			>	Feature interaction and importance		XGBoost, LR, GBM, RF, MLP	Mortality prediction	Accuracy: 0.879/0.857, AUROC: 0.888/0.782
Tabular data	2022 Zhang et al. [94]		>		Data dimensionality reduction		XGBoost, CNN	Ischemic stroke classification	Accuracy: 0.6020/0.5671, AUROC: 0.6757/0.6532
Tabular data	2021 Alaa et al. [95]	>	>	>	Knowledge distillation and rule extraction	AutoPrognosis RF, AdaBoost, MLP	RF, AdaBoost, MLP	Breast cancer prediction	AUROC: 0.771/0.773 & 0.823/0.792 & 0.777/0.763 & 0.815/0.784 & 0.790/ 0.778 & 0.803/0.775

TABLE 2 (Continued)

	% %									
Performance comparison	MAE: 5.34/5.94 & 3.80/ 4.05, RMSE: 8.24/9.98 & 6.05/6.68	RMSE: 0.545/0.561 & 0.494/0.588	AUPRC: 0.807/0.736			AUROC: 0.780/0.778		RMSE: 36/- & 104/-	MSE: 0.04/-	Accuracy: 0.88/-, AUROC: 0.92383/-
Healthcare applications	Personalized lifestyle recommendations	Attention-deficit hyperactivity disorder prediction	Mortality prediction	Asthma hospital visit prediction	Asthma hospital visit prediction	Mortality prediction	GBM, XGBoost, Disease classification AdaBoost	Emergency department and outpatient cancer clinic time prediction	Breast cancer prediction	Acute inflammations diagnosis, breast cancer diagnosis, etc.
Main ML architectures	RF	Deep LASSO	GBM, XGBoost	XGBoost	XGBoost	RF	GBM, XGBoost, AdaBoost	Congestion graphs, generalized Jackson networks	RF, LASSO, SVM	Elastic net, linear programming
Model name						AutoScore	mAML			Interval coded scoring
Interpretability methods	Data dimensionality reduction	Data dimensionality reduction	Feature interaction and importance	Knowledge distillation and rule extraction	Knowledge distillation and rule extraction	Intrinsically interpretable models	Data dimensionality reduction	Feature interaction and importance	Knowledge distillation and rule extraction	Intrinsically interpretable models
Automated model development			>		>		>		>	
Automated feature engineering		>	>	>		>	>	>		>
Automated data preparation	· >	[/		_			>			
Year Paper	Chiang et al. [96]	Laria et al. [97]	Ikemura et al. [98]	Luo et al. [99]	Tong et al. [100]	Xie et al. [101]	Yang at al. [102]	Senderovich et al. [103]	Banerjee et al. [104]	Billiet et al. [105]
Year	2021	2021	2021	2021	2020	2020	2020	2019	2018	2018
Data	Tabular data	Tabular data	Tabular data	Tabular data	Tabular data	Tabular data	Tabular data	Tabular data	Tabular data	Tabular data

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		Performance	comparison
			SI
		Healthcare	applications
		Main ML	Model name architectures
			Model name
		Interpretability	methods
	Automated	model	development methods
	Automated	feature	engineering
	Automated	data	preparation
ontinued)			Year Paper
E 2 (C			Year
TABLE 2		Data	type

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Performance comparison	AUROC: 0.836/- & 0.883/ - & 0.916/- & 0.828/- & 0.820/- & 0.781/- & 0.908/ - & 0.845/- & 0.890/- & 0.875/- & 0.910/- & 0.850/ - & 0.924/- & 0.890/-	F1-score 0.820/0.615 & 0.895/0.832	Accuracy: 91/90	Accuracy: 73.21/68.91 & 86.35/84.50 & 70.43/65.12 & 83.60/88.12		Accuracy: 0.866/0.864 & 0.831/0.849 & 0.836/0.815 & 0.867/0.875	AUROC: 0.887/0.703 & 0.739/0.315 & 0.743/0.614 & 0.870/0.896	AUROC: 0.966/0.957	Accuracy: 98.88/97.27 & 93.65/93.77 & 94.13/91.54	Accuracy: 97.3/72	Accuracy: 98.9/- & 98.5/-	(Continues)
Healthcare applications	Mortality prediction	Diabetes prediction, oncology prediction	Cerebral palsy classification	Histamine H1 receptor binding prediction	Prostate cancer classification	Sleep triaging	Respiratory pathology detection	Neonatal seizure detection	Myocardial infarction prediction	Epilepsy detection	Myocardial infarction detection	
Main ML architectures	RF, LASSO, GBM	Transformation graph	Naïve bayes, LR	Decision tree	Neuro-fuzzy system	LR, GBM	CNN, LR	CNN, GAT	CNN, knowledge graph	Autoencoder	CNN	
Model name	Pythia				NEFCLASS		DeepBreath					
Interpretability methods	Data dimensionality reduction	Intrinsically interpretable models	Data dimensionality reduction	Intrinsically interpretable models	Intrinsically interpretable models	Intrinsically interpretable models	Feature interaction and importance	Feature interaction and importance	Knowledge distillation and rule extraction	Feature interaction and importance	Feature interaction and importance	
Automated model development					>							
Automated feature engineering		>	>			>	>	>	>	>	>	
Automated data preparation	>		=	>	>				-			
Year Paper	Corey et al. [106]	Khurana et al. [107]	Laet et al. [108]	Drakakis et al. [109]	Keles et al. [110]	Donckt et al. [111]	Heitmann et al. [112]	Raeisi et al. [113]	Han et al. [114]	Huang et al. [115]	Jahmunah et al. [116]	
Year	2018	2018	2017	2016	2007	2023	2023	2023	2022	2022	2022	
Data type	Tabular data	Tabular data	Tabular data	Tabular data	Tabular data	Signal	Signal	Signal	Signal	Signal	Signal	

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Data type	Year	Year Paper	Automated data preparation	Automated feature engineering	Automated model Interpret development methods	Interpretability methods	Model name	Main ML architectures	Healthcare applications	Performance comparison
Signal	2022	Yang et al. [117]		>		Knowledge distillation and rule extraction		KNN, SVM, RF, MLP	Cardiac abnormalities classification	Accuracy: 99.0/98.7
Signal	2021	Lee et al. [118]		>		Feature interaction and importance		CNN	Arrhythmia classification F1-score: 81.75/82.2	F1-score: 81.75/82.2
Signal	2021	Fuchs et al. [119]	>		>	Intrinsically interpretable models		Fuzzy rules	Tremor severity assessments	MAE: 1.85/6.41 & 2.30/8.65
Signal	2021	2021 Kim et al. [120]		>		Data dimensionality reduction		CNN	BMI channel selection	Accuracy: 76.8/79.6 & 58.3/58.5 & 70.8/71.4
Signal	2019	Saboo et al. [121]		>		Data dimensionality reduction		GMM	Active electrodes selection	AUROC: 0.974/0.752
Signal	2019	Tison et al. [122]		>	>	Feature interaction and importance		CNN	Cardiac disease detection	AUROC: 0.94/- & 0.91/- & 0.86/- & 0.77/-
Genomic	2021	Clauwaert et al. [123]		>		Feature interaction and importance		Transformer	Genome annotation	AUROC: 0.740/0.882 & 0.920/0.961 & 0.976/0.958 & 0.981/0.978 & 0.976/ 0.964, AUPRC: 0.039/ 0.035 & 0.057/0.132 & 0.141/0.098 & 0.128/0.128 & 0.141/0.137
Genomic	2020	2020 Le et al. [124]		>	>	Data dimensionality reduction	TPOT-FSS	TPOT, XGBoost	TPOT, XGBoost TPOT enhancement	
Genomic	2019	Trabelsi et al. [125]			>	Feature interaction and importance	deepRAM	CNN, RNN	DNA/RNA sequence binding specificities prediction	AUROC: 0.930/- & 0.951/-
Genomic	2018	Nagorski et al. [126]	>		>	Data dimensionality reduction	SpaCC	Convex optimization	Cancer epiGenomictics subtype discovery	

TABL	TABLE 2 (Continued)									
		Automated Automate	Automated	d Automated						
Data		data	feature	model	Interpretability		Main ML Healthcare	Healthcare	Performance	ce
type	Year Paper	preparation	n engineering	ring development methods	methods	Model name	Model name architectures applications	applications	comparison	a
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Performance comparison	Accuracy: 82.35/79.41 & 88.24/88.24 & 97.06/97.06		AUROC: 0.81/0.69	C-statistic: 0.72/0.70 & 0.70/0.72	AUROC: 0.849/0.847	Accuracy: 0.79/-, AUROC 0.85/-	Accuracy: 0.849/- & 0.872/- & 0.885/- & 0.926/-	AUROC: 0.845/0.841 & 0.845/0.826, AUPRC: 0.464/0.453 & 0.180/0.125	AUROC: 0.957/0.994 & 0.987/0.910, AUPRC: 0.798/0.975 & 0.299/0.262	AUROC: 0.686/0.612 & 0.726/0.646 & 0.760/0.689 & 0.862/0.800
Healthcare applications	Cancer prediction	Ovarian cancer detection	Prostate cancer detection	Cardiac resynchronization therapy outcome prediction	Visual acuity prediction	Stroke prediction	Alzheimer's disease classification, frontotemporal dementia classification	In-hospital mortality prediction, physiological decompensation prediction, length of stay prediction	Medical condition prediction	Medical image report generation
Main ML architectures	SVM	Bayesian network	CNN, U-net, SVM	VAE, LR, cox regression	XGBoost, CNN, U-net	XGBoost	SVM, KNN, decision tree	Transformer, RF, LSTM	LR	Graph transformer
Model name	OFSSVM						GA-MADRID		PheVis	KERP
Interpretability methods	Data dimensionality reduction	Knowledge distillation and rule extraction	Feature interaction and importance	Feature interaction and importance	Feature interaction and importance	Data dimensionality reduction	Data dimensionality reduction	Feature interaction and importance	Intrinsically interpretable models	Feature interaction and importance
Automated Interpret development methods		>			>					>
Automated feature engineering	>		>	>	>	>	>	>	>	
Automated data preparation		<u>~</u>				>	>		>	
Year Paper	.8 Shen et al. [127]	2007 Yap et al. [128]	3 Roest et al. [129]	:3 Wouters et al. [130]	.2 Abbas et al. [131]	.2 Gerbasi et al. [132]	.2 Gutierrez et al. [133]	2. Zhang et al. [134]	11 Ferté et al. [135]	2019 Li et al. [136]
Data type Yea	Genomic 2018 sequence	Genomic 2000 sequence	Multi- 2023 modality	Multi- 2023 modality	Multi- 2022 modality	Multi- 2022 modality	Multi- 2022 modality	Multi- 2022 modality	Multi- 2021 modality	Multi- 2019 modality

TABLE 2 (Continued)

Data			Automated Automated data feature		Automated model	Interpretability		Main ML	Healthcare	Performance
type	Year	Year Paper	preparation engineering		development methods	methods	Model name	Model name architectures	applications	comparison
Multi- modality	2018 Chen et al.	Chen et al. [137]	>			Data dimensionality reduction	DASSA	IB, MDL	Disease propagation pattern detection	
Multi- modality	2017	2017 Guo et al. [138]		>	>	Knowledge distillation and rule extraction		Hierarchical probabilistic framework	Dermatology image analysis	Accuracy: 75.3/62.9, AUROC: 0.78/0.67

Abbreviations: AUPRC, area under the precision recall curve; AUROC, area under the receiver operating characteristic curve; C-statistic, concordance statistic; DSC, dice similarity coefficient; HD, hausdorff distance; IOU, intersection over union; MAE, mean absolute error; MSE, mean squared error; PSNR, peak signal-to-noise ratio; RMSE, root mean squared error

Furthermore, we implemented a toy example to compare AutoML solutions with classic ML models based on 44 918 de-identified patients from BIDMC critical care units [145]. The prediction target was in-hospital mortality (8.81% across all patients) and the candidate variables were age, temperature, platelet, glucose, sodium, lactate, potassium, bicarbonate, heart rate, respiration rate, hematocrit, creatinine, hemoglobin, chloride, anion gap, white blood cells, blood urea nitrogen, systolic blood pressure, diastolic blood pressure, mean arterial pressure, and peripheral capillary oxygen saturation [146]. We randomly divided the entire dataset using the ratio 6:2:2 for model training, validation, and testing. For traditional ML models, we optimized the hyperparameters using grid search based on the area under the receiver operating characteristic curve (AUROC) evaluated on the validation set. For AutoML solutions, we automatically determined the hyperparameters using their inherent algorithms; therefore, their training data included both the training and validation sets. Figure 4 presents the AUROC results on the unseen test set, which demonstrates that the two AutoML solutions of AutoGluon [147] and TPOT [124] statistically significantly outperformed the conventional ML models random forest [148], gradient boosting machine (GBM) [149], and Knearest neighbor [150]. We made the code open access to enable reproducibility and serve as an exemplary case study [151].

# 3.2 | Interpretation methods

Regarding the ML interpretations, we grouped them into four categories based on the commonly adopted criteria [18, 19]: feature interaction and importance (n = 63), data dimensionality reduction (n = 27), intrinsically interpretable models (n = 14), and knowledge distillation and rule extraction (n = 14).

Feature interaction entails quantifying the effect of one feature on another, considering their mutual influence, whereas feature importance involves discerning the significance of input features in shaping the output targets of ML models [61, 63, 67, 72, 76, 78-81, 98, 103, 108, 122, 125, 136]. In the healthcare domain, the alignment of feature interaction and importance with clinical expertise enhances healthcare professionals' trust in ML outputs [152]. However, when feature interaction and importance diverge from established knowledge, ML models may encounter overfitting issues. Remarkably, such disparities occasionally reveal previously unidentified markers [153].

Data dimension reduction refers to the use of a subset of the most informative raw inputs or modified

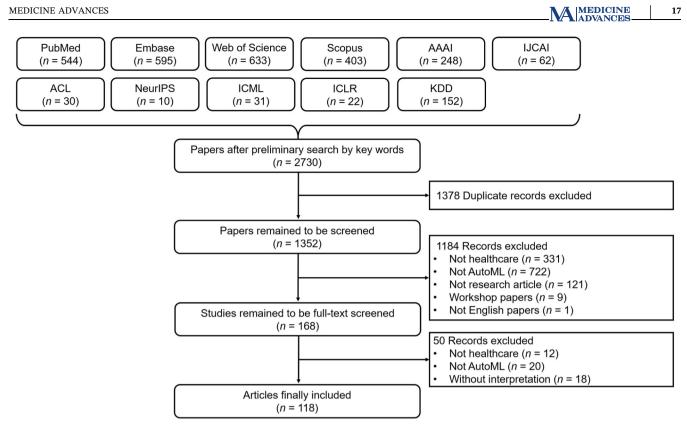
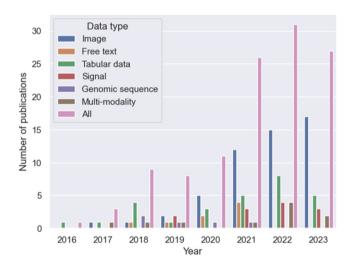


FIGURE 1 Literature selection flow for automated machine learning with interpretation in healthcare.



MEDICINE ADVANCES

FIGURE 2 Timeline of publications on automated machine learning with interpretation for healthcare since 2016. Our search concluded on September 1, 2023, which accounts for the lower number of included publications published in 2023 compared with those published in 2022.

features in model development and subsequent analyses [61, 62, 64–66, 68, 71, 73, 76, 102, 106, 108, 120, 121, 124, 126, 127, 137]. In the context of highdimensional samples, data dimension reduction helps the model to focus on salient features, thereby simplifying model complexity and enhancing its interpretability [154]. Additionally, data dimension reduction enables the effective graphical visualization of data distributions within a low-dimensional space [155]. This visualization reveals latent data patterns that can be integrated into subsequent model development, thereby enhancing both model performance and interpretability [18, 19].

Intrinsically interpretable models represent the application of transparent models to solve prediction problems [18] such as logistic regression [101, 111, 135, 140, 141, 156, 157], decision tree [25, 54, 70, 105, 137], fuzzy rules [41, 110, 119], and mathematical solid decision functions [40, 69, 107]. Intrinsically interpretable models feature simple architectures or algorithms, thereby fostering a clear understanding of the relationship between inputs and outputs [18, 19]. These models may not consistently achieve predictive performance comparable with that of their black box counterparts, but within high-stakes tasks that impact lives, model transparency is substantially more important than marginal performance superiority [158].

Knowledge distillation and rule extraction refer to the processes of simplifying intricate ML models into either streamlined models or human-comprehensible rules, respectively [99, 100, 104, 128, 138]. Knowledge distillation is a technique designed to train simple student models by mirroring the behavior of complex teacher models while preserving model performance [159]. Post distillation, student models demonstrate reduced

**FIGURE 3** Overview of the ML components automated by automated ML within the healthcare sector. This figure is reproduced from [139] with permission. ML, machine learning.

TABLE 3 Description of ML components automated by AutoML in the healthcare sector.

Stages	Operations	Description
Automated data preparation	Automatic data collection	Collecting raw data in an automated manner.
	Noise filtering	Removing inherent noise from the data.
	Missing value imputation	Filling in missing values in the dataset.
	Data imbalance compensation	Addressing and compensating for imbalanced classes in the data.
	Data normalization	Scaling data to a standard range.
	Redundant data removal	Eliminating duplicate or unnecessary data entries.
	Outlier removal	Identifying and removing anomalous data points.
	Samples clustering	Grouping similar data samples together.
	Data pattern shift detection	Detecting changes in data patterns over time.
	Continuous variable binning	Converting continuous variables into discrete bins.
Automated feature engineering	Automatic feature generation	Creating features automatically using algorithms.
	Feature selection	Choosing the most relevant features for modeling.
	Feature transformation	Transforming features to a more suitable form for modeling.
Automated model development	Backbone model selection	Choosing the main model architecture.
	Model tuning	Adjusting model-specific parameters for better performance.
	Hyperparameter optimization	Finding the best hyperparameters for better performance.

complexity, which renders them more comprehensible to humans and potentially bolsters transferability [160]. Rule extraction yields human-understandable rules because each rule inherently provides a logical explanation for its decision [161]. Based on these interpretation methods discussed above, healthcare practitioners can discern potential errors and ascertain the reliability of ML models [162].

# 3.3 Data modalities

In this section, we discuss AutoML with interpretation for different types of healthcare data: image (n=53), free text (n=8), tabular data (n=29), signal (n=12), genomic sequence (n=6), and multi-modality (n=10). Figures 5 and 6 present the summary statistics of AutoML and interpretation techniques in the included

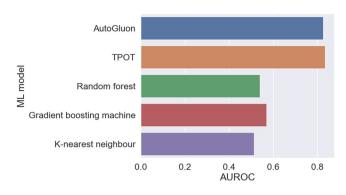


FIGURE 4 AUROC comparison of automated ML solutions versus conventional ML methods for real-world in-hospital mortality prediction. AUROC, area under the receiver operating characteristic curve; ML, machine learning.

publications. For AutoML techniques, automated feature engineering dominated in five out of the six modalities; for the genomic sequence, automated model development was more prevalent. Regarding interpretation methods, feature interaction and importance were widely used in all modalities except the genomic sequence, where data dimensionality reduction was the preferred approach. For each data modality, we focused on the principal tasks addressed by AutoML with interpretation systems and elaborated on them using representative studies.

#### 3.3.1 **Image**

Medical images are essential diagnostic tools for a spectrum of diseases [66, 163]. AutoML with interpretation enables clinicians with little coding experience [164] to perform a spectrum of healthcare tasks, such as retinalvessel caliber measurement [61], breast cancer classification [60], and thoracopathy lesion localization [165]. Based on whether they transform raw pixels into useable features, current systems can be classified into two categories: (1) two-step systems that consist of feature extraction and subsequent modeling [64, 66, 68, 70, 71, 166]; and (2) end-to-end systems without the explicit extraction of intermediate features [67, 69].

Two-step systems first extract image features from raw pixels and build up the subsequent analysis based on the extracted features. Various methods have been proposed to automate the extraction of image features, including both commercial software and homemade models. Yin et al. [68] applied the commercial software CellProfiler [167] and ImageJ [168] to extract individual and textual features, and then integrated domain knowledge from pathologists to shortlist useful features. PDE [66] has also demonstrated its effectiveness in automatic feature extraction. By contrast, Yan et al. [64]

developed a feature extraction tool and demonstrated the effectiveness of their homemade model through a comparison with human clinicians. With diverse off-the-shelf solutions, multiple tools have been combined to improve the robustness of extracted features [169]. In these systems, the most common interpretation is feature interaction and importance that results from mapping the extracted features back to the original images and highlighting relevant pixels or patches [64, 71]. Additionally, in some systems, inherently interpretable models are applied based on the extracted features to improve model interpretations [70, 166]. For instance, Wu et al. [70] implemented a decision tree to mimic how radiologists interpret the extracted features. Moreover, knowledge distilled from an inherently interpretable model, such as a decision tree, can serve as diagnostic guidelines in the future [166].

Different from two-step methods, end-to-end systems process image inputs without the implicit extraction of intermediate features and output predictions of interest in addition to useful interpretations [170]. In the task of compressed sensing for functional magnetic resonance imaging (fMRI), Lecouat et al. [69] automated the architecture design and parameter training of artificial neural networks (ANN) based on convex optimization and non-cooperative games [171]. To enhance interpretability, they introduced a decision function with sparse parameters and clear mathematical formulas. Wang et al. [67] developed a classic end-to-end system for congenital heart disease classification, including automatic data clustering and model parameter tuning. Similar to twostep systems, their system highlighted important areas on the input image toward ML predictions and used these sub-areas as an interpretation. Although end-to-end systems provide more ceaseless automation and are thus more user-friendly, users should choose the appropriate systems based on whether they need the intermediate features for further modeling and interpretation [68].

#### 3.3.2 Free text

Medical text records various patients' information, such as hospitalization descriptions, diagnoses, and treatments [172]. Accurate mining of such information can summarize patients' former health conditions and guide subsequent interventions [173]. A fundamental task addressed by AutoML with interpretation is the coding of unstructured raw clinical notes into structured medical codes, such as the international classification of diseases (ICD). Similar to the two-step systems adopted in medical image analysis, this process extracts standard intermediate features from text records, and these intermediate

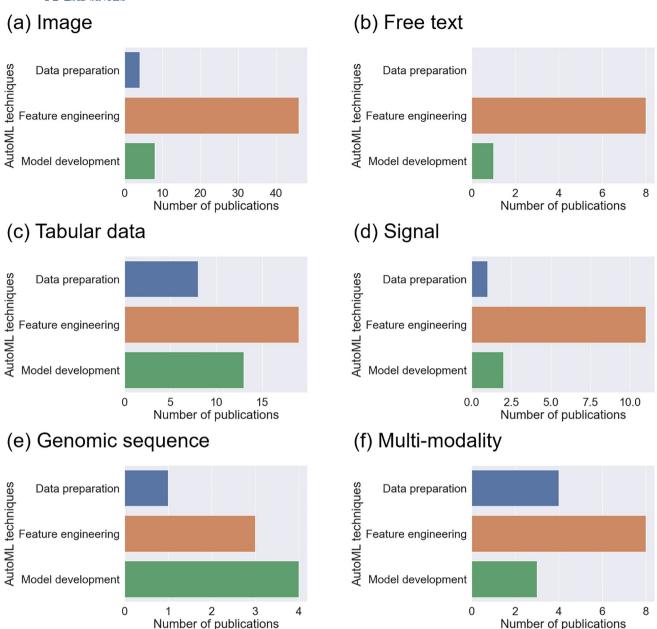


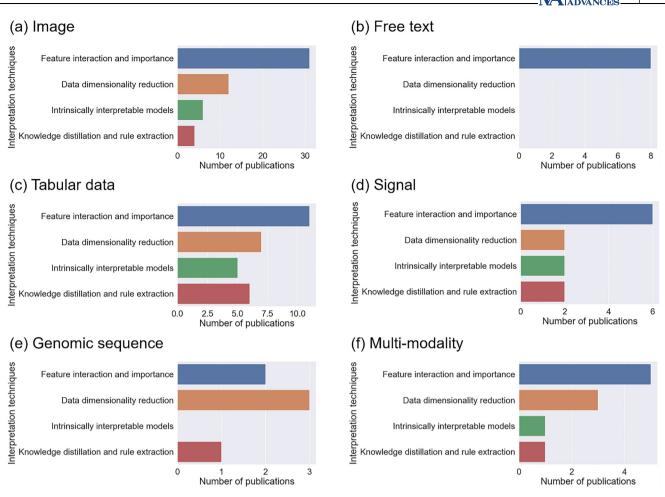
FIGURE 5 Summary statistics of automated machine learning techniques for the included publications targeting each data modality: (a) image; (b) free text; (c) tabular data; (d) signal; (e) genomic sequence; and (f) multi-modality.

features facilitate various subsequent analyses [80]. Conventionally, such a transformation was conducted manually [79, 80], but has been gradually replaced by either commercial software or home-made models to save time and eliminate errors. For example, commercial software called clinical text analysis and knowledge extraction system has been demonstrated to be an effective method for mapping trauma encounter text to structured medical concepts [75]. Additionally, researchers have demonstrated that homemade models are useful for generating informative feature vectors from free text and subsequently projecting these vectors to medical codes [74, 76, 77, 80, 81]. Duarte et al. proposed a

framework similar to residual learning, wherein word embeddings are processed using a gated recurrent unit (GRU) to generate representations [81]. These representations are then concatenated with the initial embeddings to prevent information loss and enhance model accuracy. Additionally, Atutxa et al. demonstrated that beyond classic recurrent neural networks (RNN) such as GRU, convolutional neural networks (CNN) and transformers are also effective for mapping diagnostic text to ICD codes [80].

Across all analytical tasks that use medical text, the attention mechanism is the most important backbone. It is valued not only because of its superior performance in

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**FIGURE 6** Summary statistics of interpretation techniques for the included publications targeting each data modality: (a) image; (b) free text; (c) tabular data; (d) signal; (e) genomic sequence; and (f) multi-modality.

the attention-based transformer [174] but also because of its inherent weights that provide feature interaction and the importance of each part in the input text [76, 78, 81]. For instance, in the sentence "He should undergo chemotherapy when he is diagnosed before his cancer cells metastasize," attention detects that "his cancer cells metastasize" is a crucial component in the automatic determination of the patient's cause of death [76]. In recent studies, researchers introduced the hierarchical attention mechanism, which uses the various types of attention and interprets feature representations on different levels. The hierarchical label-wise attention network [77] applies two-level attention mechanisms at the word-level and sentence-level for selecting important words and sentences in each paragraph, respectively.

# 3.3.3 | Tabular data

Tabular data, the most common data format in healthcare, includes structured demographic data, vital signs, lab tests, diagnoses, treatments, and procedures [1]. Unlike pixels in images and words in free text, raw features, such as gender in tabular data, typically have clinically explainable meanings, therefore feature engineering becomes the focal point of automation and interpretation for AutoML systems. It should be noted that the proposed methods for tabular data in the included studies can also be applied to structured information derived from unstructured healthcare data, as illustrated in the two-step methods above. In this section, we focus on studies in which researchers explored raw inputs in a structured tabular format.

Traditional feature engineering for tabular data is labor-consuming and costly. It requires ML engineers' intuition and domain knowledge [107]. By contrast, automatic and interpretable feature engineering automatically performs transformation and aggregation across candidate features in a transparent manner. For example, Khurana et al. [107] proposed automatic feature selection and transformation based on intrinsically interpretable transformation graphs, and found that the

modified features reduced ML errors. Their work demonstrated the utility of intrinsically interpretable models in feature engineering. AutoScore [141, 156] further exploits the full potential of the intrinsically interpretable clinical score as the backbone for predicting parameters such as the in-hospital mortality rate [141, 156], survival time [157], and rare event occurrence [140]. Although complicated ML models have dominated the analysis of high-dimensional data, for tabular data with a limited number of features, transparent features and intrinsically interpretable models are still preferred in practice [175, 176].

In addition to feature engineering, data preparation (pre-processing) [106] and model development [98] have been automated using AutoML with interpretation systems. Ikemura et al. [98] automated the entire ML lifecycle using commercial software [177] and interpreted models through feature interaction and importance generated by Shapley additive explanations (SHAP) [178]. In addition to commercial software such as H2O.ai, researchers have also developed comprehensive homemade systems for mining clinical tabular data. mAML [102] is an example that includes automated imbalance compensation [179], feature selection [180], and hyperparameter optimization [181]. Specifically, imbalance compensation is addressed using RandomOverSampler [179], SMOTE [182], and ADASYN [183]. Feature selection methods include the distal DBA method [184], HFE [180], and mRMR [185]. Hyperparameter optimization is performed using a grid search [181].

# 3.3.4 | Signal data

Signal data refers to electrical or mechanical signals collected from physiological sensors to monitor the functioning of the human body and make informed intervention decisions [186]. ML has been applied to identify the sophisticated relationships between various signal inputs and clinical events. AutoML with interpretation further automates and improves the reliability of this analytical process. A promising research direction involves transforming signal data into two-dimensional representations and subsequently applying imagerelated methods [118]. However, in this section, we focus on these techniques specifically designed for signal data to avoid confusion. Specifically, Fuchs et al. [119] used an intrinsically interpretable fuzzy model to analyze tremor signals, in which the wrapper approach [187] and pyFUME [188] automate feature selection and model development, respectively. Kim et al. [120] proposed an automated channel selection method based on CNN for electroencephalograms. analyzing They further

elucidated neurophysiological feature interaction and importance by correlating the selected channels with specific brain regions. In addition to the end-to-end architecture, Tison et al. [122] devised a two-step framework for predicting distinct heart diseases. Initially, the system autonomously generated features using a CNNhidden Markov model from electrocardiograms (ECG). Subsequently, these features were input into a GBM for predicting the target diseases. Finally, the system calculated the interaction and importance of segments within ECG as the model interpretation. A similar strategy was implemented by Jahmunah et al. [116] in which ECG beats were first extracted using an off-the-shelf algorithm and then input into the downstream DenseNet [189] for myocardial infarction detection. Han et al. [114] conducted an extensive investigation into the use of AutoML for diagnosing myocardial infarction. On top of clinical standards, diagnostic guidelines, and DenseNet-based signal morphology, they developed an interpretable diagnostic system based on production rules.

# 3.3.5 | Genomic sequence

Genomic sequence data [190, 191] indicate the precise order and arrangement of fundamental genetic elements, such as nucleotides (adenine, thymine, cytosine, and guanine), within DNA sequences (DNA-seq). In addition to DNA-seq, other common genomic sequences include RNA sequences (RNA-seq), Deoxyribonuclease I hypersensitive site sequences (DNase-seq), micrococcal nuclease digestion with deep sequencing (MNase-seq), and chromatin immunoprecipitation sequences (ChIPseq). These sequences encapsulate the detailed composition of genetic material, thereby offering fundamental information that is essential for comprehending potential associations between genetic patterns and diseases [192]. The principal application of AutoML with interpretation in genomic sequence data mining is to identify genomic sites of interest from the entire genomic sequence. Trabelsi et al. [125] proposed deepRAM for identifying protein binding sites in DNA and RNA-seq based on a hybrid architecture of CNN and RNN. The hyperparameters were automatically tuned through a combination of random search and cross-validation. Sequence motifs, which represent patterns with biological significance, were extracted from the initial CNN layer to improve interpretability [125]. In addition to genomic sites, AutoML has been applied to the data mining of gene expression data. Shen et al. [127] introduced elastic netbased [193] automatic feature selection to a support vector machine (SVM), which demonstrated that feature selection boosted both model performance

interpretability. In addition to classic ML models such as SVM, the transformer has gradually gained popularity in genomic sequence analyses, such as automatic prokaryotic genome annotation [123]. In addition to inherent attention in the transformer for acquiring feature interaction and importance, data dimensionality reduction [124] and rule extraction [128] are used to improve model interpretability.

# 3.3.6 | Multi-modality

Multi-modality refers to the simultaneous use of more than one data type discussed above to gain a comprehensive understanding of a patient's condition [194]. The integration of these complementary modalities enhances the overall diagnostic accuracy of ML models [195]. AutoML with interpretation is highly valued for processing complex data that involve multiple modalities [196]. PheVis [135] uses a dictionary-based named entity recognition tool to extract medical concepts from free text and then fuses these features with diagnosis codes to predict rheumatoid arthritis and tuberculosis. The SAFE algorithm [197] is used for automatic feature selection, and logistic regression is used for the transparent modeling of the relationship between shortlisted features and medical conditions of interest. Similarly, Zhang et al. [134] combined phenotypical features from free text and clinical features from tabular data to predict in-hospital mortality, physiological decompensation, and length of stay in intensive care units. Compared with features from a single modality of either free text or tabular data, multimodal features have led to statistically significant improvements in performance across most evaluated settings. For analogous frameworks within the field of image modality and signal modality, readers can refer to Abbas et al. [131] and Wouters et al. [130], respectively. They used different tools to extract features from image or signal data and combined them with tabular features, which achieved state-of-the-art performance. In addition to integrating different data modalities for predicting events of clinical interest, the aligned data of different modalities facilitates the translation of high-dimensional data into human-understandable formats, such as human language. KERP [136] was proposed to automatically generate free text reports for medical images, where feature interaction and importance, derived from attention weights, are leveraged to connect generated reports with original image regions, mimicking the inference process of a human radiologist.

In addition to the six detailed data categories above, healthcare data can also be generally classified as spatial or sequential data. Image data primarily encompasses spatial information, whereas temporal tabular data, free text, signal data, and genomic sequence data fall into the sequential data category. Medical videos represent an integration of both spatial and sequential data. The shared characteristics across different modalities pave the way for a unified architecture that is capable of handling various data types. Chen et al. [137] designed DASSA for automatic pattern change detection within any sequential data and demonstrated its potential for analyzing the aforementioned sequential data within a unified framework.

# 4 | DISCUSSION

As a fundamental component for the successful implementation of ML in healthcare, AutoML with interpretation reduces the barriers to the full lifecycle of ML analyses and provides interpretations for healthcare professionals [198]. Through a systematic literature review, we discussed the methodologies and applications of AutoML with interpretation for six data types: image, free text, tabular data, signals, genomic sequence, and multimodality. We identified three components that have been automated in ML analyses: data preparation, feature engineering, and model development. We summarized four major interpretation methods: feature interaction and importance, data dimensionality reduction, intrinsically interpretable models, and knowledge distillation and rule extraction. Using Table 2, readers can easily identify papers in which AutoML with interpretation and model performance are discussed for their tasks of interest. Despite the promising performance achieved by AutoML with interpretation systems, several challenges persist, including the absence of automatic data preparation, the loose integration of automation and interpretation, and the unmet compatibility with multi-modality. Additionally, the latest advancements in foundation models have the potential to revolutionize AutoML with interpretation.

The first challenge of current AutoML with interpretation systems is the absence of automatic data preparation, as highlighted by the finding that automatic data preparation was integrated into AutoML with interpretation systems in only 18 out of 118 studies [199]. Realworld healthcare records contain issues such as missing values, outliers, inconsistencies, duplicates, and non-standardization [200]. These issues constitute almost 50%–80% of the overall workload in the complete lifecycle of ML analyses, underscoring the necessity for automated data preparation within the infrastructure of future AutoML systems [201]. Additionally, we suggest that ML engineers should frequently communicate with

healthcare professionals during the system design phase to align their work with real-world demands [202]. For instance, although complex ANNs have become the primary choice in some application domains, such as reinforcement learning [203], intrinsically interpretable models are favored in healthcare settings, such as emergency departments [204]. Hence, ML engineers should ensure the inclusion of common intrinsically interpretable models in their systems rather than exclusively incorporating various ANN architectures.

The second challenge identified in the included papers is the loose integration of automation and interpretation. In all the included studies, the researchers addressed interpretation issues to some extent. Researchers should leverage the insights gained from interpretation to enhance their model automation rather than merely adding post hoc explanations as the last module in their frameworks. A good demonstration was provided by Ikemura et al. [98]. They applied SHAP and PD plots to analyze the decision processes of their AutoML models, indicated potential medical knowledge from their studies, and further reused these findings to enhance their models. The interaction between model development and model interpretation can be achieved by automated feature selection, which reveals feature importance, offers model interpretation, simplifies model structure, and potentially enhances model performance [205]. In addition to automated feature selection, for future AutoML with interpretation, researchers should explore the research direction of developing the tightly knit integration of AutoML and ML interpretations.

Furthermore, the expanding collection of multimodalities presents an opportunity for ML engineers to develop an AutoML with interpretation system that emulates a human clinician's inference process based on various types of healthcare data [177]. Specifically, when patients visit a hospital, clinicians and nurses investigate their former medical records, which are in the form of text and tabular data. Then, some tests may be conducted on the feedback image and signal data. Some advanced treatments involve genome sequencing, which introduces genetic data into the consultation and diagnosis. Handling such abundant and complex information requires a great deal of domain knowledge. The scenario becomes even more intricate when healthcare professionals seek to leverage ML, and this is an exact application scenario for AutoML with interpretation systems. Given the recent versatile application of the transformer for the data types image [206], free text [207], signal data [208], and genomic sequence [209], future researchers can explore the development of comprehensive AI doctors that use multi-modal healthcare data as inputs, automate the entire pipeline of data analyses, and

generate results along with interpretations based on a unified backbone architecture.

Recent advancements in foundation models for text, image, and multi-modality have the potential to significantly enhance all three stages of ML: data preparation, feature engineering, and model development [210]. These models excel in zero-shot learning, which enables them to perform tasks without additional training on specific datasets. For example, large language models, such as ChatGPT, can perform a range of tasks from ICD code extraction [211] to risk triage prediction [212] based on prompts provided by healthcare professionals. This zeroshot capability elevates ML to an unprecedented level of automation, potentially obviating the need for tedious data preparation and computationally intensive model development in certain tasks [213]. By contrast, in tasks in which foundation models exhibit suboptimal performance, they can serve as effective tools for feature engineering. The representations within their architectures can be extracted to enhance downstream models [214]; in previous studies, researchers validated that downstream models embedded with these representations outperformed powerful baseline models [215].

Our study had certain limitations that warrant refinement in future work. First, we sought to provide an overview of current AutoML with interpretation systems in healthcare settings. Hence, we did not consider the technical details of AutoML and interpretation techniques. For readers interested in these technical intricacies, we recommend referring to the original papers for a more in-depth exploration. In future work, we may conduct a detailed review of areas such as the underlying algorithms, methodologies, and implementation frameworks. Second, for a given data modality, various commercial software and homemade solutions are readily available, as illustrated above. Although Figure 4 exemplifies the effectiveness of AutoML in predicting inhospital mortality for a real-world application, we refrained from suggesting a one-size-fits-all solution because of the heterogeneous properties of datasets across different scenarios. In future endeavors, we could undertake a thorough benchmarking analysis to delineate guidelines. An exemplary precedent is in the investigation conducted by Gijsbers et al. [216], wherein they meticulously scrutinized 9 AutoML frameworks across 71 classification and 33 regression tasks. Finally, to ensure that all the reviewed papers underwent peer review, we excluded preprints, which may have resulted in the latest developments in the field being overlooked. In future studies, we could explore the integration of bibliometric methodologies to discern high-quality preprints from a broader pool, thereby enhancing the comprehensiveness of paper inclusion [217].

# 5 | CONCLUSION

AutoML with interpretation is essential for the successful uptake of ML by healthcare professionals. This review provides a comprehensive summary of the current state of AutoML with interpretation systems in the context of healthcare. To some extent, the proposed systems facilitate effortless development and improve users' trust in ML in healthcare settings. In future studies, researchers should focus on automated data preparation, the seamless integration of automation and interpretation, compatibility with multi-modalities, and the utilization of foundation models to expedite clinical implementation.

# **AUTHOR CONTRIBUTIONS**

Han Yuan: Conceptualization (equal); data curation (equal); formal analysis (equal); investigation (equal); methodology (equal); visualization (equal); writing—original draft (lead); writing—review & editing (lead). Kunyu Yu: Conceptualization (equal); data curation (equal); formal analysis (equal); investigation (equal); methodology (equal); writing—original draft (equal); writing—review & editing (equal). Feng Xie: Conceptualization (equal); data curation (equal); investigation (equal); methodology (equal); writing—original draft (equal); writing—review & editing (equal). Mingxuan Liu: Formal analysis (supporting); investigation (supporting); writing—original draft (supporting); writing—review & editing (supporting), Shenghuan Sun: Formal analysis; investigation; writing—original draft.

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# CONFLICT OF INTEREST STATEMENT

All authors declare that they have no conflicts of interest.

# DATA AVAILABILITY STATEMENT

Data sharing is not applicable to this article as no new data were created or analyzed in this study.

# ETHICS STATEMENT

This study is exempt from review by the ethics committee because it does not involve human participants, animal subjects, or sensitive data collection.

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