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REVIEW ARTICLE

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JOURNAL OF

Spin Practices and Low Reporting Quality in Studies on Prediction Model of Diagnosis Based on TCM Information: A Systematic Review and Evidence Mapping

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Abstract

Background: Despite the abundance of published studies on prediction models for diagnosing Traditional Chinese Medicine (TCM), there remains a lack of comprehensive assessment regarding reporting and methodological quality, as well as an absence of examination into the objectivity of linguistic aspects within these studies.

Methods: The PubMed, Cochrane Library, Web of Science, CNKI, VIP, and WANFANG databases were systematically searched from inception to October 30th, 2023, to identify studies developing and/or validating diagnostic and prognostic TCM prediction models using supervised machine learning. PROBAST and TRIPOD were employed to assess the reporting and methodological quality of identified studies. A previous article about spin in prognostic factor studies already identified several practices, which we modified for our data extraction the present study was registered on PROSPERO with the registration number CRD42023450907.

Results: 35 and 19 eligible studies published in Chinese and English were identified respectively from 1746 records. The clinical diseases with the most publications were diabetes (n = 7, 14.8%), coronary heart disease (n = 6, 11.1%), and lung cancer (n = 5, 9.26%). Primary analysis and sensitivity analysis confirmed that the reporting and methodological quality of included studies were correlated ($r_s = 0.504$, p < 0.001). The quality of the CM prediction model requires improvement by including a structured title, participants and predictor's selection, statistical analysis methods, model performance and interpretation. Two studies (4.55%) recommended the model to be used in daily practice lacked any external validation of the developed models. Six studies (13.63%) made recommendations for clinical use in their main text without any external validation. Reporting guidelines were only cited in one study (1.85%).

Conclusion: The available evidence indicated TCM information can provide predict information for different diseases, but the scientific quality of published studies needs to be improved.

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Keywords

- Traditional Chinese medicine
- Prediction model
- Reporting standards
- Methodological quality
- Spin practice

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Introduction

The clinical method of four diagnoses, involving inspection, auscultation and olfaction, inquiry, and palpation, is emphasized in Traditional Chinese Medicine (TCM) [1]. Collecting TCM clinical indices holds important guiding significance for subsequent diagnosis and treatment conducted by TCM doctors. In general, a variety of information is required to arrive at an accurate clinical diagnosis including the patient's medical history, physical examination, symptoms, signs, etc. With the rapid development of information technology and the popularization of big data, the modernization of diagnostic approaches has become integral to TCM [2]. In order to improve the accuracy and reliability of TCM diagnosis, so as to decrease some challenges, such as high subjectivity, strong empirical dependence, and inconsistent diagnostic outcomes, researchers have delved into harnessing machine learning and data mining technology to establish TCM prediction models that can assist in the diagnosis of various diseases in clinical practice. For instance, Wang, et al. [3], stated that the machine learning classification model based on tongue and pulse data exhibits glorious performance and can adequately predict PCOS risk; Shi, et al. [4] investigated the tongue image features among patients with lung cancer and constructed a lung cancer risk warning model using machine learning methods.

Since the publication of the Transparency Report on Multivariate Prediction Models for Individual Prognosis or Diagnosis (TRIPOD) statement [5] and the Predictive model Risk of Bias Assessment Tool (PROBAST) [6], the publication of machine learning clinical prediction methods rapidly increases. Nonetheless, studies using machine learning techniques often face questions about their actual effectiveness within the clinical workflow. In recent years, a large number of prediction models have been published, but often with inadequate reporting and methodological quality. For example, Li et al. [7], used PROBAST to assess the methodological quality of published systematic reviews of predictive models for motor function in patients 3-6 months after a stroke. Their assessment highlighted limitations, particularly the failure to report key information during predictive model development and validation. Despite the seeming objectivity and straightforwardness of prediction models based on readily available clinical information, their potential to provide risk estimates for diverse outcomes is

hindered by a lack of comprehensive reporting. This limitation restricts the utility of study findings, impeding subsequent validation endeavors, evidence synthesis, or in daily practice, leading to research waste.

In addition, a methodology study focusing on predictive models has highlighted that the misuse of lingual, whether intentional or unintentional, affects the interpretation of findings and has been described as "rotation" [8]. Spin is sometimes referred to as "scientific hype," a situation where scientific findings are inappropriately exaggerated and widely understood as a biased presentation [9]. In the scientific literature, spin refers to the specific reporting practice of distorting the interpretation of a result, thereby misleading the reader into perceiving the outcome more favorably. Evidence suggests that "spin" or over-exaggeration of scientific findings is prevalent across various study types, including randomized therapeutic interventions, observational studies, biomarker analyses, diagnostic test accuracy studies, prognostic factor investigations, and systematic reviews. This phenomenon significantly influences reader interpretation and decisionmaking [10].

There is no comprehensive summary of the evidence reported in TCM diagnostic prediction models, along with an evaluation of the scientific quality of published studies. Thus, we present the outcomes by a systematic review to summarize both the reporting and methodological quality of TCM diagnosis prediction model studies, in addition to examining spin practices. Moreover, we provide an evidence gap map based on Area Under the ROC Curve (AUC) results for each included study, addressing the knowledge gaps identified in this realm.

Methods

For the reporting of this study, we adhered to the Preferred Reporting Items for Systematic Reviews and Meta-Analyses (PRISMA) 2020 statement [11]. To increase the transparency of the research process and avoid selective reporting of results, we registered the systematic review on the International prospective register of systematic review (PROSPERO, https:// www.crd.york.ac.uk/PROSPERO/), with registration number CRD42023450907. Ethical approval was waived as the study design is an overview of published systematic reviews with meta-analyses and, as such, did not involve additional human participation.

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Literature search

A comprehensive and systematic search was performed to identify primary studies on machine learning-based prediction models of TCM diagnostic. Six large electronic databases, PubMed (https:// pubmed.ncbi.nlm.nih.gov/), Web of Science (https:// www.webofscience.com/), the Cochrane Library (https://www.cochranelibrary.com/), VIP (http:// www.cqvip.com/), WANFANG (https://www. wanfangdata.com.cn/), and CNKI (https://www.cnki. net/), were searched from inception to October 30th, 2023, with Medical Subject Heading and keyword terms that included "Machine Learning", "Deep learning", " Predict*", " Validat*", "Medicine, Chinese Traditional", "Complem entary Therapies", "diagnostic", and "Prognostic". The reference lists of all selected systematic reviews with meta-analyses were manually searched to identify additional articles potentially meeting the inclusion criteria. The details of all search strategies used in this review are presented in Supplementary Material.

Literature selection

We included studies that met any of the following criteria: 1) described the development or validation of one or more CM multivariable prediction models using any machine learning technique aiming for individualized predictions, or 2) reported on the incremental value or model extension aiming to develop a CM indicator-based prediction model. A multivariable prediction model was defined as a model aiming to predict a health outcome by using two or more predictor variables. We excluded studies that: 1) investigated single predictor, test, or biomarker or its causality with an outcome; 2) employed machine learning to enhance the images or signals interpretation; 3) utilized predictors solely for genetic traits or molecular markers; 4). utilized predictors solely for the effectiveness of specific CM interventions. The search was restricted to human subjects, English and Chinese language articles, and those accessible through our institution's resources. As meta-analysis essentially involves the statistical synthesis of findings from individual studies [12], preclinical studies, qualitative systematic reviews, narrative reviews, protocols, duplicate records, and publications of non-target intervention were excluded. All records identified from electronic databases were imported into Endnote (Version X9, Clarivate Analytics) to assess eligibility. Firstly, two independent reviewers (JK.Lim and XA.Xiao)

screened records based on their titles and abstracts, after deduplication. Subsequently, the full text of potentially eligible records was then downloaded for further scrutiny. Any disagreement during the selection process was solved through discussion or consultation with the third author (ZX. Xu).

Quality assessment

The TRIPOD Statement consists of a comprehensive checklist comprising 22 items that are deemed crucial for the comprehensive reporting of studies focused on the development or validation of multivariable prediction models. These items encompass various aspects, spanning from the title and abstract (items 1 and 2), background and objectives (item 3), methods (items 4 through 12), results (items 13 through 17), discussion (items 18 through 20), and other information (items 21 and 22). TRIPOD explicitly covers the development and validation of prediction models for both diagnosis and prognosis, spanning all medical domains and all types of predictors [5]. PROBAST, on the other hand, consists of 4 domains (participants, predictors, outcome, and analysis), containing 20 signaling questions to facilitate Risk of Bias (ROB) assessment. The tool is now widely used as a general instrument for appraising the methodological quality of primary studies that develop, validate, or update (for example, extend) multivariable prediction models for diagnosis or prognosis [6].

'Spin practice' is defined as any issue that could make the clinical utility of the developed or validated prediction model look more favorable than the study design and results can underpin. A previous article about spin in prognostic factor identified several practices, which we considered for use in our data extraction [13]. For a detailed description of extracted items.

Two investigators (JK.Lim and XA.Xiao) independently evaluated the methodological and reporting quality of the studies included in the overview. Specifically, "Yes" (Y), "Partial Yes" (PY), and "No" (N) were used to answer the questions related to items in the TRIPOD. "Low ROB", "Unclear ROB", and "High ROB" were used to answer the questions related to items in PROBAST tools. The results of the assessment were cross-checked, and consensus was reached among the investigators. Any disagreement between them arising from discussions was resolved by consultation with the third author (ZX. Xu).

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Data extraction

The general information encompassed in the extracted data including the title of publication, first author's name, year of publication, writing language, journal title, study type (diagnosis vs prognosis), total sample size, the aim of the study (development only or development with external validation), clinical diseases, CM diagnosis methods, the mention of CM syndrome differentiation, the use of machine learning, and the reference to reporting guideline. The extraction form was pilot tested on five articles and was subsequently implemented using Excel 2019 (Microsoft Corporation, WA, USA). Two investigators (JK.Lim and XA.Xiao) independently extracted key outcome information and the study characteristics from each document using a predesigned Excel spreadsheet. Any disagreement was resolved through discussion or by consultation with the third author (ZX.Xu).

Statistical analysis

Data relating to prediction outcomes of various machine learning based on CM indicators for clinical diseases in the included studies were descriptively summarized. The compliance rate of TRIPOD statement items was calculated for each study and reported the numbers and their percentages of "Y", "PY", or "N" responses. Meanwhile, the numbers and percentages of "Low ROB", "unclear", or "High ROB" responses were calculated for PROBAST items. Several evidence mapping methods were used to visualize the results of quality assessment [14,15]. A radar plot was used to present the assessment of reporting quality, while a bar chart was used to display the results of methodological quality. Meanwhile, a bubble plot was used to visualize the results of the overall methodological quality of each included study. Each bubble represented a primary study, the size of each bubble proportional to sample size (ie., total number of participants), and the color of the bubble represented the disease specialty classified based on ICD-11 (International classification of diseases-11) [16]. The X-axis reflected the best AUC score for internal validation of the prediction model study, whereas the Y-axis presented the overall methodological quality assessed by the PROBAST. The overall confidence of each study was classified as "Critically Low" (more than one critical flaw with or without noncritical weaknesses), "Low" (one critical flaw with or without non-critical weaknesses), "Moderate" (more than one non-critical weakness), or "High"

(no or one non-critical weakness). Two reviewers (JY.Li and M.Zhou) with a background in evidencebased medicine, employed this tool to independently evaluate the methodological quality of the included studies, while any further disagreement was resolved through detailed discussions.

In addition, Spearman's rank (r_s) test was employed to explore the correlation between reporting quality and methodological quality using the number of responses in TRIPOD and PROBAST items for each study. To assess the robustness of the primary analysis, sensitivity analysis was conducted after excluding the "PY" and "unclear" responses. The strength of correlation was rated as low $(r_s <$ 0.4), moderate $(r_s: 0.4~0.7)$, and high $(r_s > 0.7)$ [17]. Excel 2019 and Stata 16.0 (StataCorp, College Station, TX, USA) were used to analyze and visualize the data. Statistical significance was set as two-sided p < 0.05.

Results

A total of 1746 records were identified from six databases (PubMed = 307, Web of Science = 101, The Cochrane Library =10, CNKI = 656, VIP = 494, WANFANG =178). 1637 records were screened after removing the 109 duplications. Through the meticulous evaluation of the titles and abstracts, a further 1523 records were excluded. Ultimately, 54 studies [3,4,18–69] focusing on machine learningbased prediction models for CM diagnostic were selected for final inclusion in the research. A flow diagram of the study selection process is presented in figure 1.

General characteristics of included studies

Table 1 shows the characteristics of the included articles. All 54 articles originated from Chinese institutions, of which 19 (35.19%) were written in English and 35 (64.81%) in Chinese. 32 studies (59.26%) were published in 16 various journals, 15 studies (27.78%) were published in master dissertations, and 7 (12.96%) for doctoral dissertations. The broad spectrum of total sample sizes spanned from 12,000 as the largest to 98 as the smallest. Three studies (5.5%) focused on prognostic models [45,53,56] and 51 (94.44%) on diagnostic models. Most studies reported the development of prediction models, including internal validation (n = 44, 81.5%), and 10 (18.5%) performed external validation. Among the clinical diseases under investigation, the most prevalent were diabetes (n



Table 1	Table 1: General characteristics of included articles ($n = 54$).										
Code	Author	Language	Publication	Study type	Sample Size	Study aim	Clinical disease	CM predictive factor	CM syn- drome differen- tiation	Machine learning	Refer- ence to reporting guideline
1	[66]	Chinese	J of CM	Diagnosis	1021	Develop- ment only	Meno- pause	Conventional CM information	NR	RF; SVM; ANN	NR
2	[58]	Chinese	J of Zhejiang Chinese Medical University	Diagnosis	305	Develop- ment with external validation	Stroke	Conventional CM information	yes	LR	NR
3	[54]	Chinese	J of Hunan University of CM	Diagnosis	300	Develop- ment only	Stroke	Conventional CM information	yes	SVM; KNN; RF; ExtraTree; XG- Boost; LightGBM	NR
4	[4]	English	Front Physiol	Diagnosis	862	Develop- ment only	Lung cancer	Tongue instrument parameters	NR	DT; SVM; RF; NN; NB; LR	NR
5	[3]	English	Digit Health	Diagnosis	486	Develop- ment only	PCOS	Pulse-taking instru- ment parameters; Tongue instrument parameters	NR	AdaBoost; SVM	NR
6	[18]	English	Comput Math Meth- ods Med	Diagnosis	238	Develop- ment only	CHD	Laboratory test in- dicators	yes	ANN	NR
7	[69]	Chinese	J of Yan- shan Uni- versity	Diagnosis	906	Develop- ment only	Corpore- ity	Tongue instrument parameters	NR	LR; RF; XGBoost; AdaBoost	NR
8	[64]	Chinese	Intelligent computers and appli- cations	Diagnosis	2520	Develop- ment only	Corpore- ity	Tongue instrument parameters	NR	LR; ANN; SVM; NB	NR

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) ICINE	9	[61]	Chinese	Masteral disserta- tion	Diagnosis	415	Develop- ment only	COPD	Conventional CM information	yes	NN	TRIPOD
VE MEI	10	[39]	Chinese	Masteral disserta- tion	Diagnosis	350	Develop- ment only	Lung cancer	Conventional CM information	NR	CNN	NR
TERNATIV	11	[61]	Chinese	Masteral disserta- tion	Diagnosis	235	Develop- ment with external validation	MCI	Conventional CM information	yes	LR; XGBoost	NR
AL	12	[68]	Chinese	Masteral disserta- tion	Diagnosis	845	Develop- ment only	Eczema	Pulse-taking instru- ment parameters	yes	SVM; RF; ANN	NR
EMIOLOGY	13	[50]	Chinese	Tianjin CM	Diagnosis	149	Develop- ment only	Hyper- tension	Pulse-taking instrument parameters;Tongue instrument param- eters	NR	LR	NR
rea(s): EPID	14	[67]	Chinese	Doctoral disserta- tion	Diagnosis	1676	Develop- ment with external validation	Cardiac failure	Conventional tongue information	yes	RF; DT	NR
Subject A	15	[49]	Chinese	Masteral disserta- tion	Diagnosis	500	Develop- ment with external validation	Hyper- tension	Conventional CM information	yes	LR	NR
	16	[50]	Chinese	J of CM	Diagnosis	4723	Develop- ment with external validation	Diabetes	Conventional tongue information	NR	NB	NR
	17	[42]	Chinese	Masteral disserta- tion	Diagnosis	1087	Develop- ment only	Metror- rhagia	Conventional CM information	yes	RF; SVM; ANN; DT	NR
	18	[14]	English	J Biomed Inform	Diagnosis	570	Develop- ment only	Diabetes	Tongue instrument parameters	NR	GA_XGBT; DT; KNN; LR; SVM; ANN; RF	NR
	19	[14]	English	Int J Med Inform	Diagnosis	1512	Develop- ment with external validation	Diabetes	Conventional CM information	NR	NB; LR; RF; SVM; XGB; ANN; KNN; DT	NR
	20	[25]	English	Comput Math Meth- ods Med	Diagnosis	342	Develop- ment with external validation	CHD	Hand image param- eters	NR	MTIALM	NR
	21	[25]	English	J Integr Med	Diagnosis	10060	Develop- ment only	Liver cancer	Conventional CM information	yes	PSO-ELM; SVM; BN	NR
	22	[32]	English	Biomed Pharmaco- ther	Diagnosis	586	Develop- ment only	MS	Conventional CM information	NR	DT; SVM; RF	NR
	23	[59]	Chinese	Doctoral disserta- tion	Diagnosis	316	Develop- ment with external validation	Hyper- tension	Pulse-taking instru- ment parameters	yes	RF; SVM; XG- Boost; LGB;ANN	NR

EUICINE	24	[56]	Chinese	Masteral disserta- tion	Prognosis	988	Develop- ment with external validation	Liver cancer	Conventional CM information	yes	AdaBoost; NB; ANN; RF; SVM	NR
VALIVE	25	[47]	Chinese	Doctoral disserta- tion	Diagnosis	1427	Develop- ment only	Depres- sion	Pulse-taking instru- ment parameters; Tongue instrument parameters	yes	DT;KNN;RF;Adabo ost;GDBT;Bootstr ap;NB;SVM	NR
ALIEKN	26	[50]	Chinese	Masteral disserta- tion	Diagnosis	610	Develop- ment only	CHD	Pulse-taking instru- ment parameters; Tongue instrument parameters	yes	LR	NR
<u>U</u> bY	27	[53]	Chinese	Masteral disserta- tion	Prognosis	103	Develop- ment only	Lung cancer	Conventional tongue information	NR	LR	NR
EFIDEMIUL	28	[14]	Chinese	Chinese J of Basic Medicine of Chinese Medicine	Diagnosis	852	Develop- ment only	Diabetes	Conventional tongue information	NR	LR; AN; SVM; NB	NR
bject Area(s):	29	[65]	Chinese	Chinese J of CM	Diagnosis	300	Develop- ment with external validation	Infertility	Conventional CM information	yes	RF; SVM; KNN; ANN	NR
Su	30	[55]	Chinese	Masteral disserta- tion	Diagnosis	804	Develop- ment only	Insom- nia	Conventional CM information	yes	RF	NR
	31	[4]	English	Frontiers in physiology	Diagnosis	736	Develop- ment only	Fatigue	Pulse-taking instru- ment parameters; Tongue instrument parameters	NR	LR; SVM; RF; ANN	NR
	32	[21]	English	Comput Biol Med	Diagnosis	1778	Develop- ment with external validation	NAFLD	Conventional CM information	NR	AdaBoost; GBDT; NB; ANN; RF; SVM	NR
	33	[4]	English	Biomed Res Int	Diagnosis	522	Develop- ment only	Lung cancer	Pulse-taking instru- ment parameters; Tongue instrument parameters	yes	RF; LR; SVM; ANN	NR
	34	[27]	English	BMC Med Inform De- cis Mak	Diagnosis	12,000	Develop- ment only	AIDS	Conventional CM information	yes	KNN; SVM; ANN; RF	NR
	35	[24]	English	J Healthc Eng	Diagnosis	950	Develop- ment with external validation	Corpore- ity	Conventional CM information	yes	BPNN	NR
	36	[21]	English	Medicine (Baltimore)	Diagnosis	2436	Develop- ment only	Gastrop- athy	Conventional CM information	yes	DT; SVM; RF	NR
	37	[45]	Chinese	Masteral disserta- tion	Prognosis	1713	Develop- ment only	RA	Pulse-taking instru- ment parameters; Tongue instrument parameters	yes	KNN; SVM; DF; RF; ANN; Ada- Boost	NR
	38	[62]	Chinese	Chinese J of CM	Diagnosis	622	Develop- ment only	Diabetic Neph- ropathy	Modern CM instru- ment parameters	yes	ANN	NR

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JICINE	39	[30]	Chinese	Doctoral disserta- tion	Diagnosis	1385	Develop- ment only	CHD	Modern CM instru- ment parameters	yes	DF	NR
VE ME	40	[48]	Chinese	Masteral disserta- tion	Diagnosis	802	Develop- ment only	Diabetes	Conventional CM information	NR	LR	NR
KNA	41	[38]	Chinese	Doctoral disserta- tion	Diagnosis	708	Develop- ment only	NAFLD	Tongue instrument parameters	NR	SVM	NR
ALIE	42	[52]	Chinese	Masteral disserta- tion	Diagnosis	340	Develop- ment only	Stroke	Conventional CM information	yes	LR	NR
JLUGY	43	[28]	English	Evid Based Comple- ment Alter- nat Med	Diagnosis	523	Develop- ment only	Hyper- tension	Pulse-taking instru- ment parameters	NR	LR	NR
DFM	44	[20]	English	JMIR Mhealth Uhealth	Diagnosis	467	Develop- ment only	Diabetes	Pulse-taking instru- ment parameters	NR	LR; RF; SVM	NR
rea(s): E P	45	[43]	Chinese	Masteral disserta- tion	Diagnosis	733	Develop- ment only	CHD	Conventional CM information	yes	SVM; ANN	NR
Subject A	46	[36]	Chinese	Doctoral disserta- tion	Diagnosis	515	Develop- ment only	Lung cancer	Conventional tongue information	yes	AdaBoost; NB; ANN; RF; SVM	NR
	47	[40]	Chinese	Doctoral disserta- tion	Diagnosis	520	Develop- ment only	Hyper- tension	Pulse-taking instru- ment parameters; Tongue instrument parameters	yes	Xgboost	NR
	48	[35]	Chinese	J of Beijing University of CM	Diagnosis	98	Develop- ment only	Gastrop- athy	Conventional CM information	yes	RF; SVM	NR
	49	[25]	English	Biomed Res Int	Diagnosis	929	Develop- ment only	Hyper- tension	Pulse-taking instru- ment parameters	NR	KNN; RF; Ada- Boost; Gradient Boosting; SVM	NR
	50	[41]	Chinese	Masteral disserta- tion	Diagnosis	397	Develop- ment only	Urticaria	Conventional tongue information	yes	ANN; DF	NR
	51	[63]	Chinese	Informa- tion J of Chinese Medicine	Diagnosis	919	Develop- ment only	Gastrop- athy	Conventional CM information	yes	RF	NR
	52	[34]	English	Biomed Res Int	Diagnosis	827	Develop- ment only	Diabetes	Tongue instrument parameters	NR	SVM; k-NN; NB; BPNN; GASVM	NR
	53	[51]	Chinese	J China J of CM and Pharmacy	Diagnosis	136	Develop- ment only	Hepatic failure	Conventional CM information	NR	LR	NR
	54	[26]	English	IEEE Trans Biomed Eng	Diagnosis	525	Develop- ment only	Corpore- ity	Conventional CM information	NR	RF; SVM; XG- Boost; LGB; ANN	NR

AIDS: Acquired Immune Deficiency Syndromes; CHD: Coronary Heart Disease; MCI: Mild Cognitive Impairment; MS: Metabolic Syndrome; NAFLD: Non-Alcoholic Fatty Liver Disease; PCOS: Polycystic Ovarian Syndrome; RA: Rheumatoid Arthritis; ANN: Artificial Neutral Network; DT: Decision Tree; BPNN: Back Propagation Neural Network; DF; Extremely Randomized Trees, Extra trees; GDBT : Gradient Boosting; KNN: K Nearest Neighbor; LR: Logistic Regression; NB: Naive Bayes; SVM: Support Vector Machine; RF: Random Forest

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Subject Area(s):

= 7, 14.8%) [20,22,23,34,44,57,62], Coronary Heart Disease (CHD) (n = 6, 11.1%) [18,31,37,43,50,67], and lung cancer (*n* = 5, 9.26%) [4,29,36,39,53]. 31 studies (57.41%) used conventional CM fourdiagnosis information, including inquiry scale, tongue diagnosis and pulse diagnosis, which binary data were used for prediction. Moreover, 9 studies (16.67%) used objectified parameters collected by tongue instrument and 12 studies (22.22%) by pulsetaking instrument. 29 studies (53.7%) predicted the outcome of different CM syndrome types. Support Vector Machine (SVM) and Random Forest (RF) were the most widely used machine learning, utilized in 28 (51.85%) and 22 (40.74%) instances, respectively. Remarkably, the citation of reporting guidelines was observed in only one study (1.86%) [61], which mentioned TRIPOD.

Reporting and methodological quality assessment

Based on the TRIPOD checklist⁵, the reporting quality of the CM diagnostic prediction studies was suboptimal. The results of the assessment are presented in figure 2 and further elucidated in **Supplementary File 2-Data** extraction. All primary studies (n = 54, 100%) reported sufficient details on item 10d (statistical analysis methods: specify all measures used to assess model performance), item 13b (participants: describe the characteristics of the participants), and item 14a (model development: specify the number of participants and outcome events in each analysis). Given the potential variation in abstract formats across different journals, the contents reported in the abstracts were evaluated rather than rigidly assessing item 2 against the "structured format" requirement. Then 50 (81.81%) studies reported sufficient details for this item. 14 studies (25.92%) reported sufficient details on item 4a (source of data: describe the study design or source of data) and (specify the key study dates). None of the studies achieved a "Y" rating for item 6b (Report any actions to blind assessment of the outcome to be predicted), and similarly, no study reported details on item 7b (report any actions to blind assessment of predictors for the outcome and other predictors). In terms of study size and missing data handling, 23 studies (42.59%) reported sufficient details on item 8 (explain how the study size was arrived at for predictors), and 15 (27.78%) described how missing data were handled (e.g., complete-case analysis, single imputation, multiple imputation). 10 included studies (18.52%) furnished performance measures with 95%Cls for the prediction model in item 16. Only 2 studies (3.7%) reported the results from model updating in item 17. Most of the studies reported study limitations in item 18 (n = 51, 94.44%), but



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only three studies (5.56%) discussed the results with regard to performance in the development data. 21 studies (38.89%) provided information about the availability of **supplementary** resources in item 21 (e.g., study protocol, web calculator, and data sets). 26 studies (48.15%) reported funding sources detailly in the study in item 22.

The methodological quality of the studies of CM diagnosis prediction model including four domains were evaluated by the PROBAST [6]. For domain 1: Participants, 13 studies (24.07%) reported study designs as a cohort design, nested case-control or case-cohort design (with proper adjustment of the baseline risk/hazard in the analysis), while the remains were unclear. 32 studies (59.26%) reported the inclusion and exclusion criteria of patients in the prediction model in detail and referred to the disease diagnostic criteria of Chinese and Western medicine, respectively. Six studies (11.11%) did not report CM diagnostic criteria in detail. There is no information on whether inappropriate inclusions or exclusions took place in 16 included studies (29.63%). For domain 2: Predictors, there is no information on how predictors were defined or assessed for most of the studies (n = n)33, 61.11%). Meanwhile, most studies (*n* = 35, 64.81%) did not report detailed outcome information to evaluate the predictors and showed no information on whether predictors would be available at the time the model is intended to be used for prediction (n = n)36, 66.67%). For domain 3: Outcome, the outcome determination has been used which is considered optimal or acceptable for all included studies (n = n)54, 100%), while 46 of them mentioned but did not prespecified these methods (85.19%). 33 studies (61.11%) did not report information about whether the outcome was determined without knowing information about predictors. 49 studies (90.74%) did not provide information on time intervals to illustrate the relationship between predictor assessment and outcome ascertainment. For domain 4: Analysis, all studies reported information on the number of candidate predictor parameters or number of participants with the outcome, such that the Events Per Variable (EPV) can be calculated, while 10 of the studies them were not reported the reasonable number of participants (EPV < 10). Continuous and categorical predictors were handled appropriately in all primary studies (n = 54, 100%). 43 studies (79.63%) reported data analysis for all participants. The remains of 11 studies (20.37%) were with no information, in which five of the studies (9.26%) reported participants with missing data are omitted from the analysis. Although most studies (n = 47, 87.04%) appropriately evaluated relevant model performance metrics (e.g., calibration and discrimination) and had detailed internal validation techniques (e.g., bootstrapping and crossvalidation). 45 studies (83.33%) did not provide on whether complexities (e.g., censoring, competing risks, sampling of control participants) in the data are present or accounted for appropriately. The details of the PROBAST assessment are presented in figure 3 and **Supplementary File 2-Data** extraction.

In addition, the correlation between the reporting and methodological quality of the included studies based on the number of "Y" and "low ROB" responses was moderate (r_s = 0.504, p < 0.001). Sensitivity analysis did not decrease the correlation (r_s = 0.568, p<0.001).

Extent of spin practices across sections

A thorough assessment of spin practices was conducted, categorizing 15 distinct spin practices that occurred across various sections: title, results, discussion, and conclusion in both the abstract and main text. Remarkably, no spin practices were identified in titles concerning inconsistency with study results or use of leading words (n = 0, 0%). Within the results section, three spin practices (n = n)3, 5.56%) were observed, involving the use of leading words such as "Excellent" "significant", "accurate", and "improve". Similar practices were also found in the discussion and conclusion section (n = 3, 5.56%). 27 studies (50%) and 43 studies (79.63%) used strong statements to describe the model in the abstract and the main text respectively. However, there is no spin practice in reporting differences between performance measures pre-specified in methods and results. In the main text, leading words were used as spin practices in 17 studies (31.48%) for results, 28 studies (51.85%) for discussion, and 28 studies (51.85%) for conclusion. Finally, qualifiers (such as "very" and "may") were used frequently to describe findings in the main text (n = 6, 11.11%). In the both of abstract and the main text, the diagnostic prediction model based on CM indicators, neither emphasized model relevance nor recommended application of the model in a different setting or population without external validation in the same study. Detailed information is presented in table 2 for further insights.

Evidence mapping of CM diagnostic prediction model

As shown in figure 4, the evaluation of overall methodological quality revealed that 36 CM diagnosis prediction model publications were judged as

Subject Area(s): EPIDEMIOLOGY | ALTERNATIVE MEDICINE



Figure 3 Methodological quality based on the PROBAST tool.

Table 2: Frequency of 'spin' practices in abstract, and main tex	t.				
	No. (%)				
	Abstract (n = 54)	Main text (n = 54)			
Title					
Title is inconsistent with the study results	0 (0)	NA			
Use of leading words	0 (0)	NA			
Results section					
Differences between performance measures pre-specified	2 (2)	2 (2)			
in Methods and reported in Results section	0(0)	0(0)			
Use of leading words	3 (5.56)	17 (31.48)			
Novel	0 (0)	3 (5.56)			
Excellent	1 (1.85)	4 (7.41)			
Accurate	2 (3.7)	0 (0)			
Significant	3 (5.56)	16 (29.62)			
Improved	1(1.85)	11 (20.37)			
Use of strong statements to describe the model and/or	07 (50)	42 (70 (2))			
model performance / accuracy / effectiveness	27 (50)	43 (79.63)			
Spin in tables or figures	NA	10 (18.52)			
Non relevant models are not discussed	NA	45 (83.33)			
Discussion and conclusion section					
Use of strong statements to describe the model and/or	0 (0)	12 (24 07)			
model performance / accuracy / effectiveness	0(0)	13 (24.07)			
Use of leading words	3 (5.56)	28 (51.85%)			
Novel	0 (0)	3 (5.56)			
Excellent	1(1.85)	4 (7.41)			
Accurate	2 (3.7)	0 (0)			
Significant	3 (5.56)	17 (31.48)			
Improved	1(1.85)	23 (42.59)			
Invalid comparison of results to previous development and/	NA	10 (10 50)			
or validation studies is given	NA	10 (18.52)			

	Nonrelevant models are not discussed	NA	0 (0)
	Emphasis on model relevance in the abstract while the	0 (0)	0 (0)
	results reported previously do not support such position	0(0)	0(0)
	Discrepancy between full-text and abstract explanation of	0 (0)	0 (0)
	the study findings	0(0)	0(0)
	Recommendation to use the model in clinical practice	2 (4 55)	6 (12,62)
	without external validation in same study	2 (4.55)	0 (13.03)
ç	Recommendation to use the model in a different setting or	0 (0)	7 (12.06)
	population without external validation in the same study	0(0)	7 (12.96)
	Qualifiers are used	6 (11.11)	25 (42.30)
1	Very	3 (5.56)	19 (35.19)
ļ	May/Might	6 (11.11)	13 (24.07)
	Other benefits not prespecified in Methods are addressed	NA	0 (0)
5	Conclusions are inconsistent with the reported study results	0 (0)	0 (0)
5	Conclusion focuses solely on significant results	6 (11.11)	10 (18.52)
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"Critically Low" quality, 14 studies were judged as "Low" quality, while only four were considered "Moderate" quality.

Based on ICD-11 categorization, the 54 included studies could be divided into 12 categories: Circulatory disorders (CHD: n = 5; hypertension: n = 6; cardiac failure: n = 1), Endocrine, nutritional, or metabolic diseases (PCOS: n = 1; diabetes: n=8), Neoplasm (lung cancer: n = 5; liver cancer: n = 2), Urogenital Diseases (metrorrhagia: n = 1; infertility: n = 1), Neurological disorders (Stroke: n = 3), Dermatologic Diseases (eczema: n = 1; urticaria: n = 1), Musculoskeletal system or connective tissue disorders (RA: n = 1), Infectious or parasitic diseases (AIDS: n = 1), Sleepwake disorder (insomnia: n = 1), Mental, behavioral, or neurodevelopmental disorders (depression: n=1; fatigue: n = 1; MCI: n = 1), Digestive system disease (hepatic failure: n = 1; gastropathy: n = 2; NAFLD: n = 2), and corporeity (n = 4). Four studies with "Moderate" methodological quality were observed to predict COPD (AUC=0.814) [61], cardiac failure (AUC=0.871) [67], stroke (AUC = 0.95) [54], and MCI (AUC = 0.83) [60]. 15 disease prediction models diagnosed by CM yielded AUC values greater than 0.9. Among them, J Li, et al. [22] tested diabetics with the GA_XGBT model had reported the strongest predictive power, the AUC is 0.984 with the methodological quality judged as "Critically Low". SJ Yao, et al. [66] evaluated menopause utilizing an RF model, achieving AUC



Subject Area(s):

of 0.98 with the methodological quality judged as "Low". In circulatory disorders studies, LL Hu reported the best decision tree predictive result for CHD (AUC=0.917) [43], with the quality of evidence was evaluated as "Critically Low". YL Shi, et al. [30] demonstrated the Neural Network model provided the strongest predictive power for Non-Small-Cell Lung Cancer (NSCLC) based on objective CM indicators, achieving an AUC of 0.94, which also judged as "Critically Low" quality in terms of methodological quality.

Discussion

Principal findings

This systematic review is the first overview summarizing the clinical evidence for machine learning-based prediction models of CM diagnostic. Included studies mostly published in journals related to complementary medicine, intelligent computing, and biomed pharmacology from 2004 to 2023. The overall quality of reporting and methodological quality of this systematic review included in the present study were suboptimal, highlighting a pressing need for higher-quality publications in the realm of CM diagnostic prediction models. Content requiring marked improvement in reporting and method quality in terms of Title, details of participants, calculate of sample size, details of statistical analysis, model validation, details of model performance, interpretation, supplementary information, and funding report. For spin practice, attention needs to be paid for rigor of expression. And it is not recommended to apply models to different settings or populations without external validation in the same study.

Over recent years, a large number of CM prediction models have been published in the field of CM diagnostics, which usually provide risk estimates based on readily available clinical information, laboratory indicators, objective tongue and pulse parameters, etc. The integration of contemporary data science technology with traditional Chinese medicine diagnostic principles has yielded predictive CM diagnostic models capable of forecasting disease development and treatment outcomes for patients. With the reproducibility issues and the waste of research resources in biomedical studies have attracted considerable attention in the science community [70,71]. Researchers begin to know that the comprehensive and transparent reporting of the study design, the research process, and the final results are key safeguards to avoid the problems described above [72, 73]. As an existing review have observed, studies describing the development and validation (including updating) of prediction models often fail to report critical information, adherence to the TRIPOD checklist is strongly advocated [8]. Through open and transparent reporting of the evaluation process and results, the scientific validity and verifiability of CM diagnostics prediction model research can identify the problems and deficiencies of CM diagnostics prediction model [70]. We believed that the confidence of researchers in relative model research can be increased, the basis for further research can be provided. In addition, it should be noted that the assessment of methodological quality is based on the main content, so clear and comprehensive reporting is essential for the assessment of methodological quality⁶. As demonstrated in our study, we assessed the correlation between the reporting and methodological quality of the studies included in this study, similar to the article by Chapman, et al. [17], assessing the quality of systematic reviews in highimpact surgical journals. According to the results presented in this study, diagnostic criteria for CM and Western medicine need to be provided in detail to reduce selection bias in relevant studies of CM prediction models. The calculation of the sample size is essential. If the sample size cannot meet the calculated results, the algorithm of the predictive model needs to be improved to reduce the overfitting, which is the main effect due to the insufficient sample size. The selection of predictive models should be based on previously published studies, which can reduce the waste of resources. The detailed machine algorithm needs to be described in detail in the supplementary material. Most notably, external validation of prediction models is essential. Because external validation can show not only the stability of the predictive power, but also its generalization and generalization ability. This is the most important support for selecting to use for clinical.

According to the spin practice problems, we know the reward system within academia and the increasing amount of published research make spin in research, to some extent, necessary and therefore more frequent. As authors, our natural inclination is to ensure the publication of our work, leading us consciously and subconsciously to use the language to increase the credibility and readability of our findings [9]. A systematic review including 35 publications assessing misleading practices showed that spin evaluation varies per study design In the moment of the moment of

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[10]. Navarro, et.al. reported that spin practices and poor reporting standards are also present in 152 studies, on prediction models using machine learning techniques⁹. We believe that a tailored framework for the identification of spin will enhance the sound reporting of prediction model studies. Therefore, this research not only added the evidence for CM model prediction of spin practice, as well as suggested that authors should make every effort to avoid distortion and hype in the further research.

The combination of four diagnostic methods in CM is partly used to realize the idea of "preventive treatment of diseases". Our results showed that there is a limited number of quantitative included models based on CM syndrome differentiation, as same as seldomly mentioned the content of complex traditional Chinese medicine diagnostic thinking process. Therefore, we hope that the incorporation of machine learning algorithms in CM diagnosis presents a viable approach to enhance the model performance for more discriminative classification of CM syndromes.

The evidence mapping results of this study showed that many of the predictive models based on CM diagnostic parameters had satisfactory test results, especially for diabetes, CHD, and lung cancer. However, the specific construction and optimization of the model still need the support of clinical data, and the clinical application value also needs further scientific evaluation. A number of studies have shown that the CM diagnosis prediction model based on tongue and pulse-taking instruments can accurately predict the development of patients' diseases and treatment effects [4,29,30]. However, this study has not yet further compared the prediction differences between the indicators based on conventional CM diagnosis and the objective prediction indicators of tongue diagnosis and pulse diagnosis parameters, as it has little relevance to the research topic of this review.

Strengths and limitations

To our knowledge, there has been no systematic review and evidence-mapping about spin practices and reporting quality in CM diagnosis studies and, particularly not in studies on machine learningbased prediction models. In terms of its advantages, the results of this study filled gaps in the knowledge and can be used to guide future research and clinical decision-making. Secondly, we prospectively registered the protocol on a widely recognized website, thus guaranteeing the transparency of our research process and avoiding the possibility of selection bias. Thirdly, we explored the relationship between the reporting and methodological quality of the included studies, and sensitivity analysis confirmed the robustness of the primary analysis. However, several limitations are worth highlighting. Firstly, we did not perform new meta-analyses by re-estimation of predicted value, because doing so would have been beyond the research aims of this research review. Secondly, we focused on the use of leading words in spin practices rather than allowing a certain degree of rhetoric and evaluating it within its specific context. Similarly, we could not determine if the use of qualifiers was detrimental because we only counted the occurrence rather than evaluating its use to show uncertainty. The assessment of spin practice inherently involves a degree of subjectivity, as reviewers' judgments play a pivotal role. While we diligently addressed any disagreements through discussion to minimize reading bias, it is plausible that others might interpret authors' statements differently, particularly in cases of linguistic spin. Thirdly, the absence of a comparison group limits our ability to establish causal relationships or infer associations between study characteristics, spin practices, and reporting standards. Our aim was to spotlight spin-indicative practices through a descriptive analysis, rather than delve into causal relationships. Despite these limitations, we still provided exploratory evidence about the presence of spin and reporting quality in CM diagnosis prediction model studies.

Conclusion

The available evidence indicates that predictive models based on CM indicators to predict diseases are worthy of consideration and can provide predictions for different diseases, but the scientific quality of published studies needs to be improved. Moreover, the predictive performance of CM diagnosis prediction models needs to be confirmed through high-quality external validation across multiple countries with large sample sizes. These concerted efforts are essential to solidify the credibility and applicability of predictive CM diagnosis models in real-world clinical scenarios.

Authors' Contributions

Conceptualization, ZXX and JYL; methodology, JYL; software, JKL; validation, YM X and JKL; formal

analysis, JYL; investigation, ZXX and JYL; resources, JYL; data curation, JKL and JYL; writing—original draft preparation, JYL; writing—review and editing, JYL and ZXX; visualization, JKL and JYL; supervision, ZXX, XXA and MZ; All authors have read and agreed to the published version of the manuscript.

Data availability

The Ethics Committee of Shanghai University of Traditional Chinese Medicine limited the measurement data used to support the results of this study in order to protect the privacy of patients. For researchers who meet the criteria for obtaining confidential data, the data of this study are available from the corresponding author upon request.

Conflicts of interest

The authors declare that there are no conflicts of interest.

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