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Predicting population health with machine learning: a scoping review

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PREDICTING POPULATION HEALTH WITH MACHINE LEARNING: A SCOPING REVIEW

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ABSTRACT

Objective To determine how machine learning has been applied to prediction applications in population health contexts, including which outcomes were studied, which data sources were used, and how models were developed.

Design A scoping review.

Data Sources MEDLINE, EMBASE, CINAHL, ProQuest, Scopus, Web of Science, Cochrane Library, INSPEC, and ACM Digital Library were searched on July 18th, 2018.

Eligibility criteria We included English articles published since 1980 that used machine learning to predict population health-related outcomes. We excluded studies that only used logistic regression or were restricted to a clinical context.

Data extraction and synthesis We summarized findings extracted from published reports, which included general study characteristics, aspects of model development, reporting of results, and model discussion items.

Results Of 22 618 articles found by our search, 231 were included in the review. The United States (n=71, 30.74%) and China (n=40, 17.75%) produced the most studies and cardiovascular disease (n=22, 9.52%) was the most studied outcome. The median number of observations was 5414 (interquartile range (IQR)=16 543·5) and the median number of features was 17 (IQR=31). The most commonly used data sources were health records (n=126, 54.5%) and investigator-generated (n=86, 37.2%). Many studies did not incorporate recommended guidelines on machine learning and predictive modeling. Predictive discrimination was commonly assessed using area under the receiver operator curve (n=98, 42.42%) and calibration was rarely assessed (n=22, 9.52%).

Conclusions Machine learning applications in population health have concentrated on regions and diseases well-represented in traditional data sources, infrequently using big data. Additionally, important aspects of model development were under-reported. Greater use of big data and uptake of guidelines for predictive modeling could improve the yield from machine learning applications in population health.

Registration Registered on the Open Science Framework on July 17th, 2018 (available at: <https://osf.io/rnqe6/>).

Strengths and limitations of this study

- Our review is one of the first syntheses of machine learning applications in population and public health.
- We used a comprehensive search strategy, including nine peer-reviewed databases, grey literature, and reference searching.
- We extracted a wide array of study characteristics, including important elements of predictive modeling reporting guidelines.
- Since both machine learning and population health have broad definitions, there may be some relevant articles that were not included.
- Given our focus on prediction, we could not address many other important intersections of machine learning and population health, such as surveillance and health promotion.

INTRODUCTION

Predictive models have a long history in clinical medicine. One well-known example is the Framingham risk score, which was first developed in 1967.[1] Such models have proliferated throughout clinical practice to inform management and interventions, including preventive approaches. More recently, researchers have developed prediction models beyond individual clinical applications, for population health uses.[2,3] While there is no universal definition of population health, it includes “the health outcomes of a group of individuals, including the distribution of such outcomes within the group.”[4] Similarly to clinical medicine, population-level models can be used to identify high-risk groups, directing the implementation of preventive

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3 interventions. Additionally, population health prediction models can inform policymakers about
4 future disease burden and help to assess the impact of public health actions. Thus far, most
5 predictive modeling in both medicine and population health has used parametric statistical
6 regression models. More recently, there has been increasing interest in the use of a broader range
7 of machine learning methods for prediction tasks.[5–7]
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17 Machine learning can be loosely defined as the study and development of algorithms that learn
18 from data with little or no human assistance.[8] These approaches have been increasingly applied
19 in the past two decades as a result of the enabling growth of big data reserves and computational
20 power.[9] Recent machine learning applications to prediction in population health contexts
21 include forecasting childhood lead poisoning,[10] yellow fever incidence,[11] and the onset of
22 suicidal ideation.[12]
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33 The distinction between machine learning algorithms and parametric regression models is
34 debated.[13] Regression models tend to impose more structure on the data, requiring greater
35 human input for the verification of distributional assumptions and incorporation of domain
36 knowledge in choosing the input parameters.[14] Algorithms employed in machine learning
37 often derive more structure directly from the data, making fewer distributional assumptions
38 about the data or variables. The literature remains divided on the relative advantages of more
39 traditional approaches compared to newer methods;[15] however, given the wide variation in
40 applications and the data used in these examples, broad assessments of superiority are often not
41 appropriate. Also, there are debates regarding the differences in developing and validating
42 machine learning approaches for health applications.[15,16]
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Population health applications of prediction models are relatively new compared to clinical applications; correspondingly, the role of machine learning in these applications has been far less studied and discussed in the health literature. The goals of our review are to determine how machine learning has been applied to prediction in population health, the nature of the models and data used, and how the models have been developed. We hope that our results will help to inform future research in this area, including the development of guidelines for machine learning applications in population health.

METHODS

We based our scoping review on the framework proposed by Arksey and O'Malley[17] and refined by the Joanna Briggs Institute.[18] We also followed the more recent Preferred Reporting Items for Systematic Reviews and Meta-analysis Extension for Scoping Reviews.[19] Our study protocol was registered on the Open Science Framework on July 17th, 2018 (available at: <https://osf.io/rnqe6/>).

Our initial goal was to scope out all machine learning applications in population health. However, the screening process identified a much larger number of publications than anticipated. Consequently, to describe the subject area comprehensively, we restricted our scope to articles predicting future outcomes.

Search Strategy

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3 Our search strategy consisted of peer-reviewed literature databases, grey literature, and reference
4 searches. First, we searched nine interdisciplinary, indexed databases (MEDLINE, EMBASE,
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6 CINAHL, ProQuest, Scopus, Web of Science, Cochrane Library, INSPEC, and ACM Digital
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8 Library) on July 18th, 2018. Our search was informed by consultation with a health science
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10 librarian, a machine learning textbook,[20] and a similar registered review.[15] Supplementary
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12 table A includes an example search query.
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19 Our grey literature search included Google Scholar and Google. We developed a Google Scholar
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21 search based on terms related to ‘machine learning’ and ‘population health’, which was refined
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23 based on the relevance of initial results. The first 200 results were included in screening. A
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25 similar approach was used for the general Google search, which we restricted to the first 30
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27 results. We examined relevant websites for publications. Results were limited to articles
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29 published on or before the date of the peer-reviewed literature search.
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35 Finally, we searched the references of relevant reviews for additional articles. Most of these
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37 reviews were identified during screening.
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41 **Eligibility Criteria**

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46 We included articles if they used machine learning to develop a predictive model that could be
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48 applied in a population health context. Therefore, we excluded articles where the model was
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50 trained primarily on people with a pre-existing disease. We also excluded articles that were only
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52 indirectly related to population health; for example, traffic accident models that did not predict a
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54 health outcome. Studies predicting individual outcomes were included if the approach was
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3 determined to be scalable to a population level. Finally, articles using only logistic regression
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5 were excluded. See appendix A for the full eligibility criteria.
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10 In order to manage the scope, articles were excluded if their full text could not be retrieved with
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12 our institutional licenses and if they were not written in English. Finally, articles published prior
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14 to 1980 were excluded as earlier machine learning investigators lacked comparable amounts of
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16 digitized data, software, and computational resources.
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21 **Screening Process**

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26 Initially, individual reviewers screened titles for obvious irrelevance to the review topic (JDM
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28 and EB). Examples of articles removed at this stage are outlined in appendix B. Then, we
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30 imported remaining references into Covidence systematic review management software.[21]
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33 Two reviewers screened the abstracts of remaining articles (JDM, EB, MO, and DF). Prior to
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35 evaluating full texts using all eligibility criteria, we then screened out articles that did not focus
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37 on a prediction application (JDM, EB, MO). Finally, two reviewers screened the full text of
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39 remaining articles (JDM, EB, MO). Conflicts were resolved by discussion between at least two
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41 reviewers.
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46 **Data Extraction and Synthesis**

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51 Individual authors extracted article data (JDM, EB, MO, and DF). We based our extraction items
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53 on important aspects of machine learning identified in a recent biomedical guideline[16] and on
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55 the transparent reporting of a multivariable prediction model for individual prognosis or
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3 diagnosis (TRIPOD) statement.[22] Major extraction categories included general study
4 characteristics (e.g. geographic location and sample size), model development (e.g. algorithms
5 used and type of validation), results (e.g. discrimination and calibration measures), and model
6 discussion (e.g. practical costs of errors and implementation). See supplementary table B for a
7 description of each extraction item.
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17 We computed descriptive statistics for all extraction items. We also completed a narrative
18 synthesis of discussion elements.
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23 **Patient and Public Involvement Statement**

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28 There was no patient or public involvement in this study.
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32 **RESULTS**

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37 We initially retrieved 16 172 articles, after removing duplicates (figure 1). We excluded 6494
38 articles after title screening, 7860 after abstract screening, 1453 when screening out non-
39 prediction articles, and 121 after full-text screening. This resulted in 231 articles being included
40 in the final review (appendix C).
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49 **General Study Characteristics**

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53 The number of articles published in the population health prediction area that used machine
54 learning increased dramatically after 2007 (supplementary figure A). Studies were undertaken
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worldwide, with the largest representation from the United States (US) (n=71, 30.74%) and China (n=40, 17.75%) (table 1). Relatively few articles came from Oceania (n=2, 0.87%), Africa (n=5, 2.16%), and the Americas outside of the US (n=13, 5.63%).

Characteristic*	Number or Median	Percent or Interquartile Range
Region		
United States	71	30.74%
Asia Excluding China	41	17.75%
China	40	17.32%
Europe	36	15.58%
Americas Excluding United States	13	5.63%
Africa	5	2.16%
Oceania	2	0.87%
Multi-region	15	6.49%
Not Reported	8	3.46%
Year published		
before 1990	1	0.4%
1990-1999	3	1.3%
2000-2004	13	5.6%
2005-2009	18	7.8%
2010-2014	70	30.3%
2015-2018	126	54.5%
Outcome level†		
Individual Risk Prediction	139	60.17%
Population Risk prediction	92	39.83%
Number of observations	5414	16 543.5

Not reported	72	31.2%
Number of features	17	31
Not reported	59	25.5%
Used any unstructured text		
Yes	24	10.4%
No	207	89.6%
Machine learning model was compared with other statistical methods	111	48.1%
Reported data pre-processing*		
Yes	160	69.3%
No	71	30.7%
Reported method of feature selection		
Yes	164	71.0%
No	67	29.0%
Reported hyper-parameter search		
Yes	114	49.4%
No	117	50.6%
Method of Validation		
Holdout	112	48.5%
Cross-validation or bootstrap	84	36.4%
External	15	6.5%
Not reported	32	13.9%
Reported descriptive statistics[§]		
Yes	140	60.6%
No	91	39.4%
Discussed the practical costs of prediction errors[¶]		
Yes	36	15.6%

No	195	84.4%
Stated rationale for using machine learning		
Yes	179	77.5%
No	52	22.5%
Discussed model usability		
Yes	91	39.4%
No	140	60.6%
Stated model limitations		
Yes	161	69.7%
No	70	30.3%
Discussed model implementation		
Yes	184	79.7%
No	47	20.3%
Dataset Availability by Study		
Closed	149	64.5%
Public	42	18.2%
Closed and Public	38	16.5%
Unknown	1	0.4%

^{*}Refer to supplementary table A for a description of each characteristic and rationales for including some elements.

[†]Individual risk prediction refers to studies that developed models to predict the health outcomes of individuals, while population risk prediction refers to studies that developed models to predict aggregated population-level health outcomes.

^{*}Whether any aspects of data cleaning or pre-processing were reported. Examples include how missing data was handled, whether log transformations were done, and if derived variables were generated.

[§]Included a broad array of descriptive statistics such as sample population demographics, feature distributions, and outcome distributions.

^{*}Whether the article discussed the relative risks of false negative and false positive results based on their predictive model in contexts where it might be used.

^{||}Closed refers to datasets that were not immediately available in the public domain or were not identifiable as such.

Table 1: Summary statistics of included articles

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3 The median number of observations in each article was 5414 (interquartile range (IQR)=16
4 543.5) and the median number of features (i.e. independent variables) used was 17 (IQR=31)
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6 (table 1). Seventy-two studies (31.2%) did not report the number of observations. These studies
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8 often used data from reportable disease databases, which do not necessarily have a firm sampling
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10 frame, making ascertainment of the number of observations difficult.
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16 **Algorithms**

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21 The most frequently used machine learning algorithms were neural networks (n=95, 41.13%),
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23 followed by support vector machines (n=59, 25.54%), single tree-based methods (n=52,
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25 22.51%), and random forests (n=48, 20.78%) (supplementary table C). About half of the articles
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27 made a comparison with statistical methods (n=111, 48.1%), which were generally logistic
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29 regression or autoregressive integrated moving average models (table 1).
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35 **Outcomes**

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39 Non-communicable disease outcomes were assessed by many articles (n=95, 41.13%), with
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41 communicable diseases (n=76, 32.90%) and non-disease outcomes (n=60, 25.97%) studied
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43 somewhat less often. The outcome most frequently predicted was cardiovascular disease (n=22,
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45 9.52%) (figure 2). Other commonly forecasted non-communicable disease outcomes were
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47 suicidality (n=13, 5.63%), cancer (n=12, 5.19%), and perinatal health (n=12, 5.19%). Influenza
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49 (n=15, 6.49%) and dengue fever (n = 14, 6.06%) were the most predicted communicable disease
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51 outcomes. Aside from non-communicable and communicable disease, mortality (n=13, 5.63%)
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53 and healthcare utilization (n=14, 6.06%) were also frequently predicted.
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Data

Data sources were usually structured (n=207, 89.6%) and closed, i.e. not publicly available (n=189, 81.8%) (table 1). The most frequently reported data sources were health records (n=126, 54.5%) and investigator-generated (e.g. cohort studies) (n=86, 37.2%) (table 2). A large proportion of studies (n=42, 18.2%) used an environmental data source (e.g. satellite imagery), mostly for prediction of infectious disease. Government databases (n=32, 13.9%) and internet-based data (n=21, 9.1%) were less frequently used. Among studies from China and the US, 80.0% and 67.6% respectively used health records data, whereas 54.5% of studies overall used these data sources (supplementary figure B).

Sources of Data Used*	Number	Percent
Environmental	42	18.2%
Geographical Information Database	12	5.2%
Meteorological/Air Quality Datasets	32	13.9%
Satellite Imagery	21	9.1%
Health Records Database	126	54.5%
Clinical Record Database†	46	19.9%
Disease Registry	2	0.9%
Population Health Survey	15	6.5%
Reportable Disease Database	42	18.2%
Other Health Records Database	30	13.0%
Government Database	32	13.9%
Census	11	4.8%

Vital Statistics	13	5.6%
Other Government Database	14	6.1%
HealthMap	3	1.3%
Private Insurance Data	9	3.9%
Private Insurance Claims	9	3.9%
Private Insurance Questionnaire	3	1.3%
Internet-based	21	9.1%
Search engine	12	5.2%
Social Media	12	5.2%
Investigator-generated[‡]	86	37.2%
Public Repositories[§]	19	8.2%
Health Organization Reports[¶]	5	2.2%
Not Reported	6	2.6%

*Categories are not mutually exclusive.

[†]Any dataset produced primarily for the purpose of delivering clinical care, such as electronic medical records and administrative healthcare databases produced by hospitals.

[‡]Any datasets resulting from researcher-driven studies such as randomized controlled trials, cohort studies, and case-control studies.

[§]Any freely available datasets such as MIMIC or the UC Irvine Machine Learning Repository.

[¶]Health-related reports, typically including disease burden estimates, produced by non-governmental or governmental organizations such as the World Health Organization.

Table 2: Data sources

Features

Biomedical and sociodemographic features were frequently used (supplementary figure C). Of these, the most commonly used were disease history (43.3%), age (48.5%), and sex/gender (41.1%). Among lifestyle features, smoking was the most frequently used (25.1%) and of

environmental features, meteorology was common (17.3%). Social media posts (5.2%) and web search queries (5.2%) were not often used. See supplementary table D for more details.

Model Development and Validation

The majority of articles reported how data pre-processing (n=160, 69.3%) and feature selection (n=164, 71%) were done (table 1). Fewer authors reported how hyperparameters were selected (n=114, 49.4%). Most studies used a holdout method of validation (n=112, 48.5%), fifteen (6.5%) externally validated their models, and thirty-two (13.9%) did not report how models were validated.

Performance Metrics

Most articles reported a prediction discrimination metric (n=172, 74.46%), with fewer reporting a measure of overall model fit (n=77, 33.33%), and few reporting a measure of calibration (n=21, 9.09%) (table 3). The most common discrimination metrics employed were area under the receiver operator curve (n=98, 42.42%), accuracy (n=76, 32.90%), and recall (n=68, 29.44%). Calibration was mostly assessed with graphing methods (n=9, 3.90%) and Hosmer-Lemeshow statistics (n=8, 3.46%). Overall performance was usually measured with a form of mean error, such as root mean squared error (n=35, 15.15%).

Prediction Performance Metrics Used	Number	Percent
Any overall performance metric	77	33.33%
RMSE	35	15.15%

MSE	26	11.26%
MAE	24	10.39%
MAPE	23	9.96%
R2*	19	8.23%
Correlation	8	3.46%
AIC or BIC	8	3.46%
Other performance metric [†]	21	9.09%
Any discrimination metric	172	74.46%
Area under the curve [‡]	98	42.42%
Accuracy [§]	76	32.90%
Recall [¶]	68	29.44%
Precision	39	16.88%
F statistics	10	4.33%
Likelihood Ratio**	4	1.73%
Youden Index	3	1.30%
Manual or visual comparison	3	1.30%
Other discrimination metric ^{††}	4	1.73%
Any calibration metric	21	9.09%
Manual or visual comparison ^{‡‡}	9	3.90%
Hosmer-Lemeshow	8	3.46%
Observed/Expected	5	2.16%
Other calibration metric ^{§§}	3	1.30%
Any reclassification metric	6	2.60%
Net Reclassification Index	5	2.16%
Integrated Discrimination Improvement	3	1.30%

RMSE = Root Mean Squared Error; MSE = Mean Squared Error; MAE = Mean Absolute Error; MAPE = Mean Absolute Percentage Error; AIC = Akaike Information Criterion; BIC = Bayesian Information Criterion.

*Includes R2 and pseudo-R2 metrics.

[†]Includes penalty error, Total Sum of Squares, proportional reduction in error, overall prediction error, specific prediction error, Nash-Sutcliffe, Root Mean Squared Percentage Error (2), mean relative absolute error, Analysis of Variance F-stat, 2LogLikelihood, relative efficiency, deviance, Ljung-Box test, mean absolute deviation, standard error, Mean Percentage Error, Brier score, and log score.

[‡]Includes c-statistic, s-index, area under ROC / AUC.

[§]Includes accuracy, misclassification, and error rate.

[¶]Includes sensitivity, specificity, true/false positive, and true/false negative.

^{||}Includes positive predictive value, negative predictive value, and precision.

^{**}Includes positive/negative LR.

^{**}Includes G-means (2), k-statistic, Matthews correlation coefficient.

^{**}Includes calibration plots.

^{§§}Includes mean bias (from Bland-Altman plot), calibration factoring, and Calibration statistic.

Table 3: Prediction Performance Metrics

Study Discussion

Most articles included some discussion of their rationale for using machine learning (n=179, 77.5%), limitations of their study (n=161, 69.7%), and how the model might be implemented (n=184, 79.7%) (table 1). Few discussed model usability (n=91, 39.4%) and only a small number discussed the costs of prediction errors in real-world contexts (n=36, 15.6%). See appendix D for a narrative synthesis of discussion reporting items.

DISCUSSION

Our results show that machine learning is increasingly being applied to make predictions related to population health. Nearly half of the included studies were conducted in the US or China.

Both countries produce the greatest number of scientific publications in general;^[23] however, they also likely benefited from robust health data infrastructures. The US has rapidly digitized

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3 much of its healthcare system, resulting in large electronic medical records (EMRs) linked with
4 government data through public-private partnerships, including processes to make these data
5 available to researchers.[24,25] Both the US and China made greater use of health records and
6 less use of investigator-generated data relative to other regions, which may have made machine
7 learning projects more tractable. They also used more internet-based data, which typically
8 includes many observations and is high-dimensional, making it amenable to machine learning
9 methods. Other countries with substantial EMR-use and government database linkage such as
10 Finland, Singapore, and Denmark[26] likely have untapped potential for machine learning
11 research. We noted that studies from Oceania, Africa, and the Americas (outside of the US) were
12 limited. This may be partly due to less availability of traditional sources of structured health data.
13 However, given that machine learning methods can incorporate non-traditional data sources,
14 there is the potential to expand use of these methods even when structured health data is
15 unavailable.

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35 We found that a wide range of population health outcomes have been the focus of machine
36 learning prediction models. However, relative to morbidity and mortality, multiple outcome
37 categories like cancer, human immunodeficiency virus, dementia, gastroenteritis, pneumococcal
38 disease, perinatal health, tuberculosis, and malaria appear understudied.[27] Many of these
39 conditions are most prevalent in regions with decreased access to traditional health data, perhaps
40 stymieing research. If machine learning methods are used to leverage novel data sources for
41 research in these regions, it could enable greater study of neglected diseases.

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3 Most investigators did not analyze a large number of observations and features. We observed a
4 high reliance on investigator-generated data, which likely made it difficult to achieve high
5 sample sizes or high dimensional data. The use of smaller datasets may affect the performance of
6 studied models, as machine learning algorithms generally require a high number of observations
7 relative to features.[28] Additionally, most studies focused on features typical of clinical
8 prediction models, such as biomedical factors and limited aspects of broader socioeconomic or
9 environmental determinants of health. We also observed infrequent use of unstructured data and
10 wearable data for prediction purposes. A reliance on small datasets and traditional numbers and
11 types of features is unlikely to fully leverage any benefits of machine learning. This may be
12 contributing to the small differences frequently seen between parametric regression and machine
13 learning model performance. Greater use of linked population-level databases, large EMRs,
14 internet data, and unstructured features would likely improve these approaches.
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33 Based on the elements of model development that we studied, adherence to existing machine
34 learning[16] and prediction model[22] guidelines appears limited. Most articles did not report
35 their method of hyper-parameter selection, discuss practical costs of prediction errors, or
36 consider model usability, which are needed for transparency and model assessment. Many
37 studies did not report the number of features included, method of validation, method of feature
38 selection, or any performance metric. Given these issues, it would be difficult or impossible to
39 compare many of these machine learning models with existing approaches. However, we
40 acknowledge that existing guidelines were not available when many included studies were
41 published. Future work should apply existing guidance,[16] including from TRIPOD,[22] and
42 anticipate the forthcoming TRIPOD-ML statement.[29]
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Lastly, we noted that included studies rarely assessed predictive performance in terms of calibration, which refers to a model's ability to accurately predict the absolute probability of outcomes.[30] In contrast, discrimination measures of predictive performance quantify a model's ability to correctly rank-order individuals. Many traditional machine learning tasks, such as image recognition, often have a high signal to noise ratio. In these cases, discrimination may be a suitable lone performance metric, as the algorithm can achieve near perfect performance.

Conversely, health outcomes tend to be more stochastic. As a result, accurate prediction of probabilities is more important.[30] Models can have good predictive discrimination, but poor calibration, making them less useful in practice, particularly for population health applications. A further issue is that many measures of discrimination, such as accuracy and recall, artificially impose a threshold for calling events. Thresholds should ideally be ascertained by decision-makers based on their cost-utility curves.[30] Overall, applications of machine learning in population health would benefit from greater use of calibration performance metrics.

A strength of our study is that we addressed an understudied area, the intersection of machine learning and population health. Additionally, prediction is an application with untapped potential in population health, and where machine learning has the potential to make significant improvements. Our study also employed a comprehensive search strategy, including numerous multidisciplinary peer-reviewed databases, alongside a grey literature search. Furthermore, we applied insights from the field of clinical prediction modeling to population health and machine learning. Finally, given the focus on prediction, we were able to take a comprehensive approach to data extraction and synthesis.

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6 In terms of limitations, concentrating on prediction prevented us from exploring applications of
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8 machine learning to other important aspects of population health, such as disease surveillance.
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10 These should be the focus of future research. Our review was also limited by including only
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12 English articles and articles with available full text, which may have introduced selection bias.
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14 Lastly, the two main concepts underlying our review, machine learning and population health,
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16 are not universally defined. As a result, we may have excluded articles that may be relevant to
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18 these fields.
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24 This was the first scoping review specifically focused on machine learning prediction in
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26 population health applications. Predictive modeling in population health can help to inform
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28 preventive interventions, anticipate future disease burden, and assess the impact of health
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30 policies and programs. Advances in machine learning offer opportunities to improve these
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32 models, particularly when incorporating big data. This is still a nascent field, but based on our
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34 findings more research in Oceania, Africa, and South America would be particularly beneficial.
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36 Diseases such as malaria, tuberculosis, and dementia should also be further studied. Additionally,
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38 future machine learning projects could incorporate larger datasets and more non-traditional
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40 features. Greater use of resources such as HealthMap, social media, web search patterns, remote
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42 sensing, and WHO reports would enable more work in regions without formal data sources and
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44 enrich research in others. Another largely untapped prospect is using machine learning and high-
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46 dimensional data to incorporate richer representations of the social determinants of health.
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48 Opportunities should continue to grow as governments increasingly digitize their health service
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50 records and link databases to both health and non-health data. Overall, as applications of
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3 machine learning in population health develop, adherence to existing guidance[16,22,29] will
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5 improve our ability to assess and advance machine learning applications. Finally, it will be
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7 important to evaluate the impact of prediction models on decisions made in population health
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9 and the practice of public health.
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14 **CONTRIBUTORS**

15
16 JDM contributed to the design of the study and led the literature search, article screening, data extraction, analysis,
17
18 and writing of the manuscript. EB contributed to the design of the study, the literature search, article screening, data
19
20 extraction, and analysis. MO contributed to article screening, data extraction, analysis, and writing of the
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22 manuscript. DF contributed to article screening and data extraction. KK contributed to the design of the study. LCR
23
24 led the design of the study. All authors interpreted study results and contributed to drafting of the manuscript.
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29
30 We are grateful to Catherine Bornbaum for her assistance with the initial design of the study.
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35
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37
38 design, data collection, data analysis, data interpretation, or writing of the report. The corresponding author had full
39
40 access to all the data in the study and had final responsibility for the decision to submit for publication.
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44 **COMPETING INTERESTS**

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46 None declared.
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50 **PATIENT CONSENT FOR PUBLICATION**

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52 Not required.
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ETHICS APPROVAL

Not required as only prior published research was included in the review.

DATA AVAILABILITY

The full data extraction table used for this review will be made publicly available after publication with no end date on Mendeley Data (DOI: 10.17632/7rrz9xrp2j.1).

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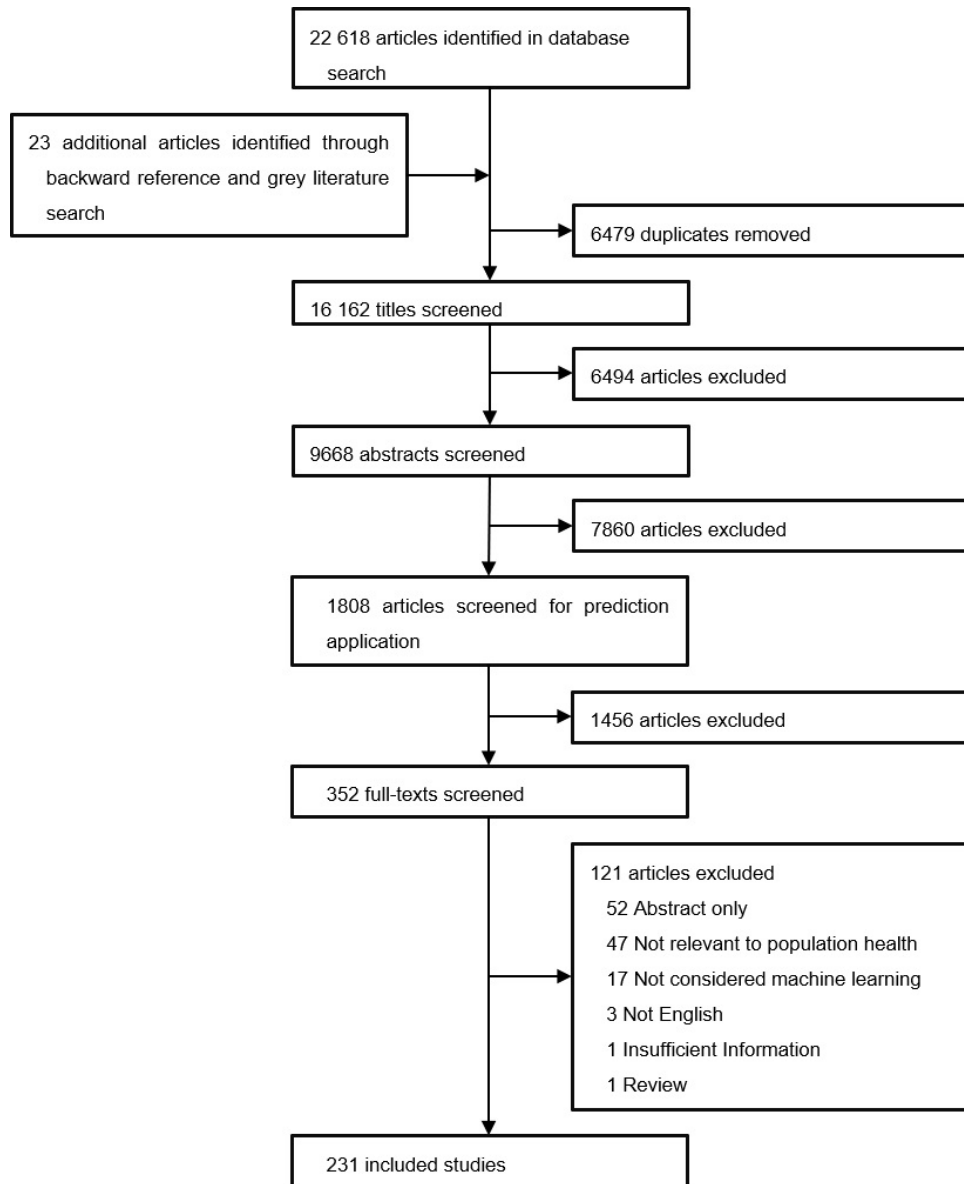
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12 13 14 15 16 **FIGURES LEGENDS**

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18 **Figure 1:** PRISMA flowchart of article screening process.

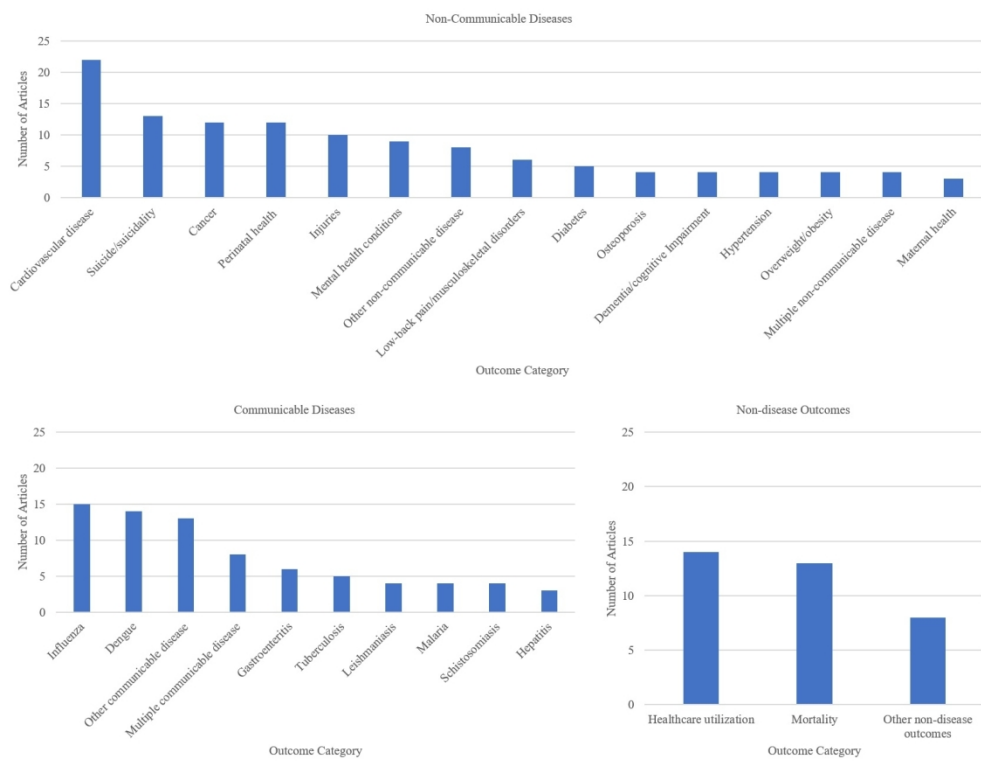
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21 **Figure 2:** Number of articles by outcome.



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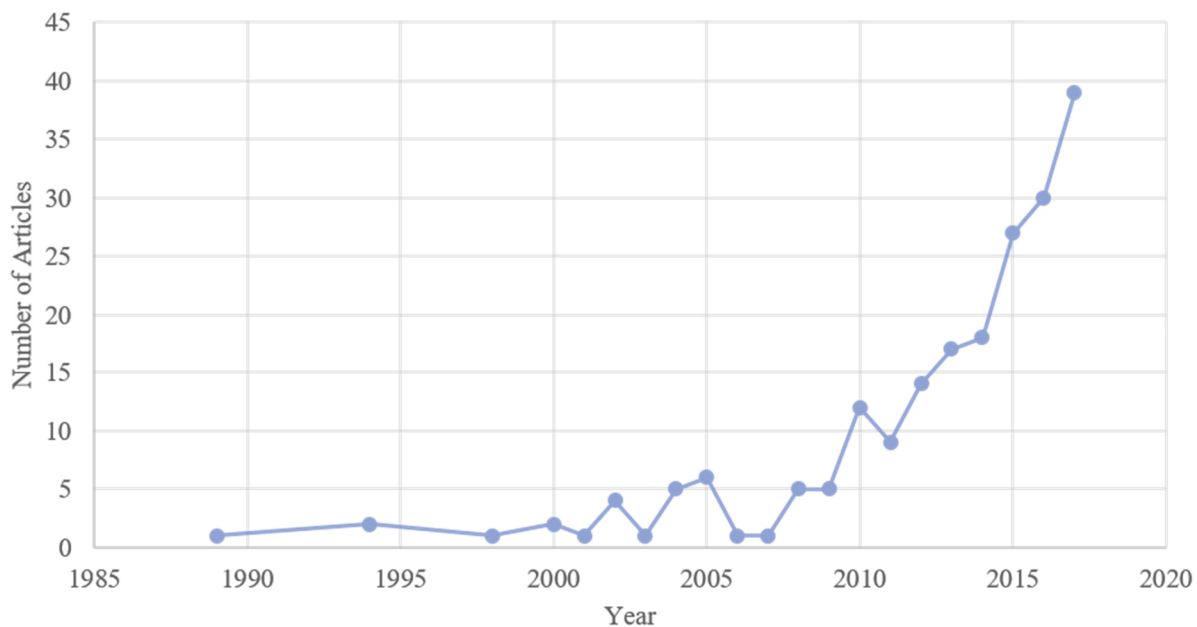
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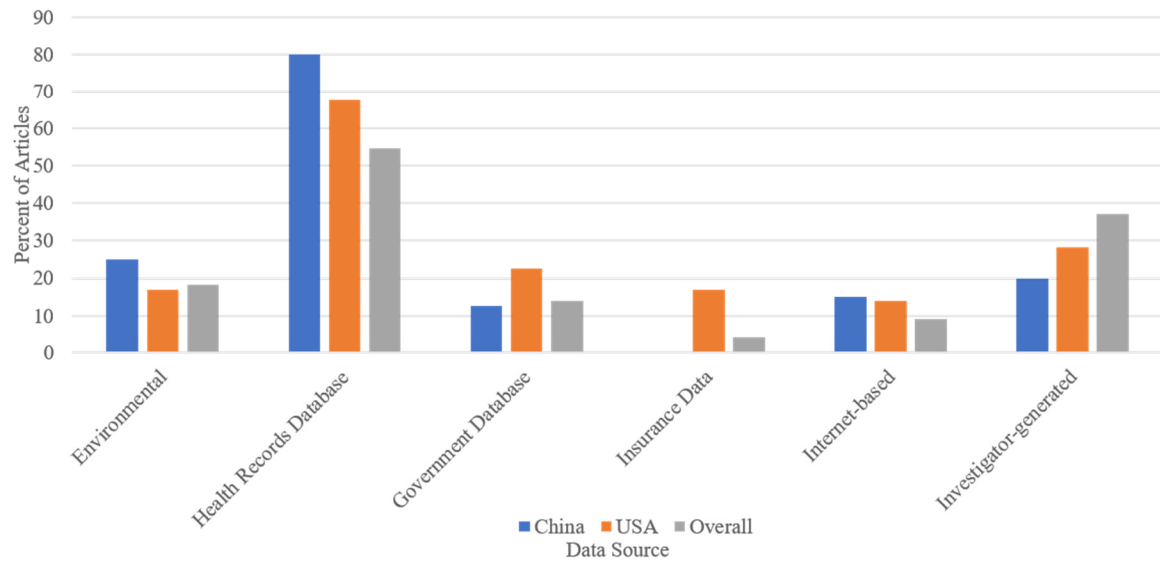


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Supplementary Figure A: Number of articles by publication year. Articles from 2018 are not plotted in this figure because the review does not include all studies published that year.

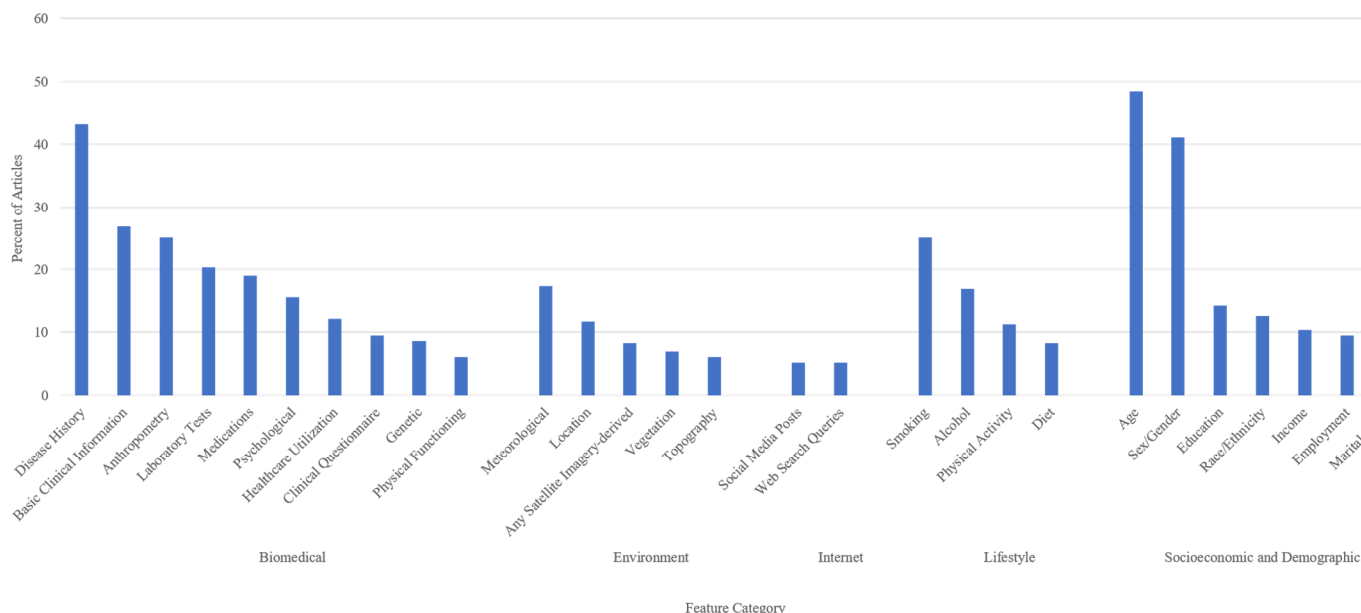


Supplementary Figure B: Data Source by selected region.



review only

Supplementary Figure C: Most commonly used feature categories.



review only

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Supplementary Table A: MEDLINE search query¹

Machine Learning Terms	Population Health Terms
1. Exp Artificial Intelligence/	24. Exp Population Health/
2. Exp "neural networks (computer)"/	25. Exp Population Surveillance/
3. Support vector machine*.kf,tw	26. Exp Health Equity/
4. Neural net*.kf,tw	27. Health status/
5. Perceptron*.kf,tw	28. Health status disparities/
6. Deep learning.kf,tw	29. Public health systems research/
7. Random forest*.kf,tw	30. "Social determinants of health"/
8. Lasso*.kf,tw	31. Health surveys/
9. Gaussian mixture*.kf,tw	32. Health status indicators/
10. Bayesian network*.kf,tw	33. "global burden of disease"/
11. Classification tree*.kf,tw	34. Global health/
12. Regression tree*.kf,tw	35. Environmental health/
13. Relevance vector machine*.kf,tw	36. Harm reduction/
14. Nearest neighbo*.kf,tw	37. Public health informatics/
15. Probability estimation tree*.kf,tw	38. Community medicine/
16. Elastic net*.kf,tw	39. Public health/
18. Naive bayes.kf,tw	40. Epidemiology/
19. Genetic algorithm*.kf,tw	41. Preventive medicine/
20. Artificial intelligence.kf,tw	42. Occupational medicine/
21. Machine learning.kf,tw	43. Environmental medicine/
22. Statistical learning.kf,tw	44. Public health practice/
23. /or 1-22	45. Preventive health services/
	46. Health promotion/
	47. public health.kf,tw
	48. population health.kf,tw
	49. health promot*.kf,tw
	50. population surveillance.kf,tw
	51. health surveillance.kf,tw
	52. health equity.kf,tw
	53. preventive medicine.kf,tw
	54. health protection.kf,tw
	55. disease prevention.kf,tw
	56. social determinant* of health.kf,tw
	57. health determinant*.kf,tw
	58. determinant* of health.kf,tw
	59. occupational medicine.kf,tw
	60. community medicine.kf,tw
	61. epidemiolog*.kf,tw
	62. health status*.kf,tw
	63. global health.kf,tw

	64. environmental health.kf,tw
	65. harm reduction.kf,tw
	66. environmental medicine.kf,tw
	67. /or 24-66
	68. 23 and 67

¹Limited to articles published in 1980 or after.

For peer review only

Supplementary Table B: Data Extraction Field Descriptions

Data Extraction Field	Description
Title	The article titles.
First Author	The last name and first initial of the first listed author of each article
Year of Publication	The year of publication noted for each article.
Outcome level	One of two categories: <ol style="list-style-type: none"> <i>Population risk prediction</i>: the aggregated outcome of a whole population was predicted <i>Individual risk prediction</i>: outcomes of individual participants were predicted
Outcome	<p>Selected from the following, which are not mutually exclusive, as some articles predicted multiple outcomes:</p> <p>Non-communicable Disease</p> <ol style="list-style-type: none"> <i>Cardiovascular disease</i>: any disease characterized by atherosclerosis and resulting ischemia, including myocardial infarction and stroke <i>Suicide/suicidality</i> <i>Cancer</i> <i>Perinatal health</i>: including pre-term birth, fetal alcohol spectrum disorder, congenital heart disease, growth failure, and neural tube defects <i>Mental health conditions</i> <i>Osteoporosis</i> <i>Low-back pain and other musculoskeletal disorders</i> <i>Diabetes</i> <i>Dementia and cognitive Impairment</i> <i>Hypertension</i> <i>Injuries</i>: including fractures, falls, traffic injury, and foreign body injuries <i>Overweight and obesity</i> <i>Maternal health</i>: including fertility, pregnancy risk, and severe maternal morbidity <i>Multiple non-communicable disease</i> <i>Other non-communicable disease</i>: including liver disorders, Crohn's disease, glaucoma, dental caries, and lead poisoning <p>Communicable Disease</p> <ol style="list-style-type: none"> <i>Influenza</i> <i>Dengue</i> <i>Gastroenteritis</i> <i>Tuberculosis</i> <i>Leishmaniasis</i> <i>Malaria</i> <i>Schistosomiasis</i> <i>Hepatitis</i>: of viral origin <i>Multiple communicable disease</i> <i>Other communicable disease</i>: including zika, hand food and mouth disease, leptospirosis, yellow fever, West Nile, and typhoid fever <p>Non-disease Outcomes</p> <ol style="list-style-type: none"> <i>Mortality</i> <i>Healthcare utilization</i> <i>Other non-disease outcomes</i>: including health behaviours, vitamin d status, and wellness score
Region	<p>Categorized based on Organisation for Economic Cooperation and Development (OECD) region except for the United States and China, which were given their own categories due to the high number of publications. One of the following:</p> <ol style="list-style-type: none"> <i>Africa</i> <i>Americas except for the United States</i> <i>Asia except for China</i> <i>China</i> <i>Europe</i> <i>Oceania</i> <i>United States</i> <i>Multi-region</i> <i>Other/Unknown</i>
Study Setting	One of two categories:

	<ol style="list-style-type: none"> 1. <i>Clinical</i>: when data was collected in any type of clinical setting 2. <i>Community</i>: when data was collected in a community setting
Data Source Categories	<p>Selected from the following categories, which were not mutually exclusive, and often more than one was used:</p> <ol style="list-style-type: none"> 1. <i>Geographical Information Database</i>: any dataset containing basic map-based spatial information such as distances and topography 2. <i>Meteorological/Air Quality Datasets</i> 3. <i>Satellite Imagery</i>: examples include the moderate resolution imaging spectroradiometer (MODIS) and the Shuttle Radar Topography Mission (SRTM) 4. <i>Clinical Record Database</i>: any dataset produced primarily for the purpose of delivering clinical care, such as electronic medical records and administrative healthcare databases produced by hospitals 5. <i>Disease Registry</i>: a dataset maintained to monitor and/or provide care for a specific disease 6. <i>Population Health Survey</i>: a regular epidemiological survey administered periodically to assess the health of populations 7. <i>Reportable Disease Database</i>: a dataset containing reports of diseases for which it is mandatory for healthcare providers to report 8. <i>Other Health Records Database</i>: any other health records dataset not encompassed in other categories, including various surveillance systems 9. <i>Census</i> 10. <i>Vital Statistics</i>: information regularly collected by governments regarding births and deaths 11. <i>Other Government Database</i>: other governmental datasets including socioeconomic and demographic information 12. <i>HealthMap</i>: a public health surveillance system using natural language processing to analyze informal data sources such as online news, individual reports, expert-curated discussions 13. <i>Private Insurance Claims</i>: including medical, hospital, and prescription drug claims 14. <i>Private Insurance Questionnaires</i> 15. <i>Internet Search</i>: including the number of searches of certain key terms and meta data such as the location of the searches 16. <i>Social Media</i>: both posts and metadata 17. <i>Investigator-generated</i>: any datasets resulting from researcher-driven studies such as randomized controlled trials, cohort studies, and case-control studies 18. <i>Public Repositories</i>: any freely available datasets such as MIMIC 19. <i>Health Organization Reports</i>: health-related reports, typically including disease burden estimates, produced by non-governmental or governmental organizations such as the World Health Organization 20. <i>Not Reported</i>
Feature Categories	<p>Selected from the following categories, which were not mutually exclusive, as often more than one category was used (if more than one instance of a feature category was found in an article it was only counted once):</p> <p>Biomedical</p> <ol style="list-style-type: none"> 1. <i>Anthropometry</i>: measurements of the human body such as height and weight 2. <i>Basic Clinical Information</i>: information typically collected during a brief physician encounter such as a focused medical history and physical examination, including blood pressure 3. <i>Basic Medical Tests</i>: any test requiring somewhat specialized equipment such as an electrocardiogram 4. <i>Clinical Questionnaire</i>: a standardized questionnaire administered in a clinical context such as the Montreal Cognitive Assessment or Patient Health Questionnaire-9 5. <i>Disease History</i>: information regarding present and/or past diagnoses of an individual 6. <i>Genetic</i> 7. <i>Healthcare Utilization</i> 8. <i>Instrumental Activities of Daily Living</i>: features relating to an individual's daily functioning in areas such as cooking and shopping 9. <i>Laboratory Tests</i>: any features derived from human specimens requiring specialized equipment for analysis, such as hematological and microbiological results 10. <i>Medical Imaging</i> 11. <i>Medications</i> 12. <i>Physical Functioning</i>: features including the presence of any physical disabilities or the status of activities of daily living 13. <i>Prenatal</i>: relevant aspects of the period before birth such as the use of prenatal vitamins or the results of routine lab results

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4	14. <i>Psychological</i> : features including mood or anxiety symptoms
5	15. <i>Self-Reported Health Status</i>
6	Internet-based
7	16. <i>Social Media Images</i>
8	17. <i>Social Media Location</i> : either aggregated or individual
9	18. <i>Social Media Metadata</i> : any information other than the content of social media posts, such as the frequency of general posts and time of posting
10	19. <i>Social Media Posts</i> : social media post content
11	20. <i>Social Network</i> : the interconnections among individuals in a social media platform
12	21. <i>Web Search Metadata</i> : any aspects of web searches other than their content
13	22. <i>Web Search Queries</i> : the content of web search queries either individual or aggregated
14	Lifestyle
15	23. <i>Alcohol</i>
16	24. <i>Diet</i>
17	25. <i>Physical Activity</i>
18	26. <i>Sleep</i>
19	27. <i>Smoking</i>
20	28. <i>Unspecified</i>
21	29. <i>Other Substance-use</i>
22	30. <i>Other Lifestyle</i>
23	Environment
24	31. <i>Air Quality</i>
25	32. <i>Any Satellite Imagery-derived</i>
26	33. <i>Biodiversity and Domestic Animals</i>
27	34. <i>Satellite-based Built Environment</i>
28	35. <i>Other Built Environment</i>
29	36. <i>Connectivity</i> : the ease of access to large urban centers and/or general services
30	37. <i>Electrical Lighting (satellite-based)</i>
31	38. <i>General Environmental Exposures (not included in other categories)</i>
32	39. <i>Hazard</i> : characteristics of an external hazard such as the presence of lighting on a roadway
33	40. <i>Satellite-based Land-use</i>
34	41. <i>Other Land-use</i>
35	42. <i>Location</i>
36	43. <i>Meteorological</i>
37	44. <i>Surface Water Distribution/Flooding (satellite-based)</i>
38	45. <i>Satellite-based Topography</i>
39	46. <i>Other Topography</i>
40	47. <i>Vector/Reservoir Characteristics</i> : including mosquito surveillance numbers and the population of non-human primates in the case of yellow fever
41	48. <i>Vegetation (satellite-based)</i> : such as the normalized difference vegetation index (NDVI)
42	49. <i>Water Composition</i>
43	50. <i>Other Satellite Imagery-derived</i>
44	51. Population Disease or Healthcare Statistics
45	Socioeconomic and Demographic
46	52. <i>Adverse Adult Experiences/Trauma</i>
47	53. <i>Adverse Childhood Experiences</i>
48	54. <i>Age</i>
49	55. <i>Antisocial Behaviour</i>
50	56. <i>Economy Makeup</i> : such as the number of individuals working in various types of occupations
51	57. <i>Education</i>
52	58. <i>Electricity</i>
53	59. <i>Employment</i>
54	60. <i>Garbage Collection</i>
55	61. <i>Healthcare System</i> : such as the availability of universal, public healthcare
56	62. <i>Household Characteristics</i> : the number of individuals in the household and their ages
57	63. <i>Housing Structure</i> : aspects of the physical structure of housing such as the number of units and age of the building
58	64. <i>Human Development Index</i>
59	65. <i>Immigration Status</i>
60	66. <i>Income</i>
	67. <i>Income Inequality</i>
	68. <i>Language</i>
	69. <i>Legal System</i>
	70. <i>Literacy</i>

	<p>71. <i>Marital Status</i></p> <p>72. <i>Occupational Risk</i>: including risk factors for low-back pain such as prolonged sitting or injury from repetitive movements</p> <p>73. <i>Parental</i>: including disciplinary styles and the amount of time spent at home and number of parent-child activities</p> <p>74. <i>Peer Group</i>: behaviours of peer group</p> <p>75. <i>Political Stability</i></p> <p>76. <i>Population and Population Density</i></p> <p>77. <i>Population Growth</i></p> <p>78. <i>Race/Ethnicity</i></p> <p>79. <i>Religion</i></p> <p>80. <i>Sanitation</i>: availability of sewage systems</p> <p>81. <i>Sex/Gender</i></p> <p>82. <i>Social Support</i></p> <p>83. <i>Unspecified</i></p> <p>84. <i>Vehicle Ownership</i>: at population level</p> <p>85. <i>Water Supply Quality</i></p> <p>86. <i>Wealth</i></p> <p>87. <i>Other Socioeconomic and Demographic</i></p> <p>88. Other Features</p> <p>89. Not Reported</p>
Number of Datasets Used	The number of distinct datasets used regardless of the number of sources.
Dataset Availability	<p>Selected from the following categories:</p> <ol style="list-style-type: none"> 1. <i>Public</i>: all the datasets used by article authors were publicly available 2. <i>Closed</i>: all the datasets were not publicly available or appeared not to be available 3. <i>Closed and Public</i>: the datasets used were a mix of available and not available
Any Unstructured Text Used	Natural human language was included in the model as a feature with no initial ordinal/nominal structure imposed.
Number of Observations	The number of individuals or other units of observations (such as countries) included in the predictive model. If multiple subsets of the data and/or distinct datasets were used for different models, the largest number was used.
Machine Learning Algorithm Type	<p>The algorithm type used to build the predictive model, with multiple types often used in the same article. Algorithms were only counted once when used in each article, even if used to build multiple different models in the same article. Selected from the following categories:</p> <ol style="list-style-type: none"> 1. <i>Neural Networks</i>: includes deep learning/deep neural networks as well as other simpler neural networks 2. <i>Support Vector Machine</i> 3. <i>Single Tree-based Methods</i>: includes classification trees, regression trees, and decision trees 4. <i>Random Forest</i> 5. <i>Least Absolute Shrinkage and Selection Operator (LASSO)</i> 6. <i>Bayesian Networks</i>: includes naïve bayes 7. <i>Feature Selection Methods</i>: includes k-means clustering and genetic algorithms; these were often used as a pre-processing step and in a few cases this was the only use of machine learning (i.e. a machine learning model was not used to build the predictive model itself) 8. <i>Boosted Tree-based Methods</i>: includes gradient boosting and boosted trees 9. <i>K-Nearest Neighbour</i> 10. <i>Elastic Net</i> 11. <i>Ridge Regression</i> 12. <i>Other</i>: includes association rule learning, single task learning, multitask learning, rough set classifier, associative classification, bagging, partial least squares discriminant analysis, Just-Add-Data Bio Tool, super learner, particle swarm optimization, ant colony optimization, Isomap, PCA, Disease State Index, Stacking, kernel conditional density estimation, stepwise deletion, conditional random fields, contrast mining, grammatical evolution, Learning from Examples Using ROugh Sets, AUtoregression with exogenous outputs, and natural language processing
Compared with Other Statistical Methods	Whether the machine learning method's predictive performance was compared with a traditional parametric statistical regression model such as logistic regression (yes/no).
Reported Data Pre-processing	Whether any aspects of data cleaning or pre-processing were reported (yes/no). Examples include how missing data was handled, whether log transformations were done, and if derived variables were generated. Missing data and all model development processes have been identified as important to report by TRIPOD.[1]
Reported Method of Feature Selection	Whether the method of feature selection was reported (yes/no). When there is a high number of features initially, this is usually done using algorithmic, domain knowledge-

	informed, or mixed approaches. Feature selection is an important element of reporting as identified by TRIPOD.[1]
Number of Features	The number of features included in the final prediction model after feature selection. If multiple models were used in one article, the largest number of features was chosen.
Reported Hyper-parameter Search	Whether the process for determining the hyper-parameters of the machine learning model, such as the number of features used to build each tree in a random forest, was reported (yes/no). This is an important aspect of model development[2], and thus considered an important element to report by the TRIPOD statement.[1]
Method of Validation	How the authors validated the predictive performance of their model, selected from one of the following categories: <ol style="list-style-type: none"> 1. <i>Holdout</i>: the dataset was divided into two parts; one part was used to train the model and the other was used to test the model 2. <i>Cross-validation and bootstrap</i>: the dataset was either divided into more than two parts and repeatedly trained and tested on different parts of the dataset or random sampling with replacement was used to train the model 3. <i>External</i>: the model was tested on a completely separate dataset
Reported Descriptive Statistics	Whether the article reported any descriptive statistics regarding their sample (yes/no). We considered a broad array of descriptive statistics including sample population demographics, feature distributions, and outcome distributions. These are all important reporting elements according to TRIPOD.[1]
Calibration Metrics	The types of calibration predictive performance metrics used to evaluate models, which could be more than one. Calibration refers to a model's ability to accurately predict absolute probabilities of the outcome occurring.[3] One or more of the following categories was selected if a calibration metric was used: <ol style="list-style-type: none"> 1. <i>Manual or visual comparison</i>: includes calibration plots 2. <i>Hosmer-Lemeshow</i> 3. <i>Observed/Expected</i>: is a ratio or comparison of observed and predicted/expected probabilities 4. <i>Other calibration metric</i>: includes mean bias (from Bland-Altman plot), calibration factoring, calibration statistic
Discrimination Metrics	The types of discrimination predictive performance metrics used to evaluate models, which could be more than one. Discrimination refers to a model's ability to correctly rank-order individuals according to their likelihood of developing the outcome.[3] One or more of the following categories was selected if a discrimination metric was used: <ol style="list-style-type: none"> 1. <i>Area under the curve</i>: meaning receiver operator curve; also includes c-statistic and s-index 2. <i>Accuracy</i>: includes accuracy, misclassification, and error rate 3. <i>Recall</i>: includes sensitivity, specificity, true/false positive, and true/false negative 4. <i>Precision</i>: includes positive predictive value, negative predictive value, and precision 5. <i>F statistics</i> 6. <i>Likelihood Ratio</i>: includes both positive and negative likelihood ratios 7. <i>Youden Index</i> 8. <i>Manual or visual comparison</i> 9. <i>Other discrimination metric</i>: includes G-means, k-statistic, and Matthews correlation coefficient
Overall Goodness of Fit Metrics	The types of overall goodness of fit performance metrics used to evaluate models, which could be more than one. Overall goodness of fit refers to a model's predictions' concordance with observed outcomes. One or more of the following categories was selected if an overall performance metric was used: <ol style="list-style-type: none"> 1. <i>Root mean squared error</i> 2. <i>Mean squared error</i> 3. <i>Mean absolute error</i> 4. <i>Mean absolute percentage error</i> 5. <i>R²</i>: includes pseudo-R2s 6. <i>Correlation</i> 7. <i>Akaike Information Criterion or Bayesian Information Criterion</i> 8. <i>Other performance metric</i>: includes penalty error, total sum of squares, proportional reduction in error, overall prediction error, specific prediction error, Nash-Sutcliffe, root mean squared percentage error, mean relative absolute error, analysis of variance F-stat, -2LogLikelihood, relative efficiency, deviance, Ljung-Box test, mean absolute deviation, standard error, Brier score, log score, and mean percentage error
Did Machine Learning Models Outperform Traditional Methods?	Whether the machine learning-based predictive models outperformed the statistical parametric regression models based on the performance metrics supplied by the authors (yes/no). However, this should not be taken to mean that the difference in model performance was reliable or valid. Often, important performance metrics and essential aspects of model development were not reported, making accurate comparisons difficult.

Discussed the Practical Costs of Prediction Errors	Whether the article discussed the relative risks of false negative and false positive results based on their predictive model in contexts where it might be used (yes/no). These costs are important for determining the usefulness and application of predictive models.[3]
Stated Rationale for Using Machine Learning	Whether the article stated any reasons for using a machine learning approach instead of a statistical parametric regression approach (yes/no).
Rationale for Using Machine Learning - Free Text	Reviewers included article quotations and summaries in this section to capture different rationales for using machine learning. Reviewers attempted to only extract free text regarding each specific type of rationale once
Discussed Model Usability	Whether the article discussed any aspect of how the model could be practically used in a relevant context (yes/no).
Stated Model Limitations	Whether the article discussed any potential limitations of the research (yes/no).
Model limitations - Free Text	Reviewers included article quotations and summaries in this section to capture different reported limitations. Reviewers attempted to only extract free text regarding each specific type of limitation once.
Discussed Model Implementation	Whether the article included discussion of any consequences of model implementation such as potential clinical, population-health, and policy-level impacts (yes/no).
Model Implementation - Free Text	Reviewers included article quotations and summaries in this section to capture different reported consequences of model implementation. Reviewers attempted to only extract free text regarding each specific type of implementation impact once.

Supplementary Table C: Types of machine learning algorithms used.

Types of Algorithms	Number	Percent
Neural Networks*	95	41.13%
Support Vector Machine	59	25.54%
Single tree-based methods [†]	52	22.51%
Random Forest	48	20.78%
LASSO	25	10.82%
Bayesian Networks [‡]	23	9.96%
Feature selection methods [§]	20	8.66%
Boosted tree-based methods	19	8.23%
K-Nearest Neighbour	19	8.23%
Elastic Net	9	3.90%
Ridge regression	5	2.16%
Other	22	9.52%

*Includes deep neural networks.

[†]Includes CART, decision trees.

[‡]Includes naive bayes.

[§]Includes cluster methods (e.g. k-means clustering) and genetic algorithms.

^{||}Includes gradient boosting and boosted trees.

^{||}Including (all algorithms used once unless otherwise specified) association rule learning (n=3), single task learning, multitask learning, rough set classifier, associative classification, bagging, partial least squares discriminant analysis, Just-Add-Data Bio Tool, super learner, particle swarm optimization, ant colony optimization, isomap, principal components analysis, disease state Index, stacking, kernel conditional density estimation, stepwise deletion, conditional random fields, contrast mining, grammatical evolution, Learning from Examples Using ROugh Sets, AUtoregression with exogenous outputs, and natural language processing (n=2).

Supplementary Table D: Detailed feature categories included in studies.

Feature Category	Number of Articles	Percent
Biomedical	141	61.04
Anthropometry	58	25.11
Basic Clinical Information	62	26.84
Basic Medical Tests	10	4.33
Clinical Questionnaire	22	9.52
Disease History	100	43.29
Genetic	20	8.66
Healthcare Utilization	28	12.12
Instrumental Activities of Daily Living	6	2.60
Laboratory Tests	47	20.35
Medical Imaging	10	4.33
Medications	44	19.05
Physical Functioning	14	6.06
Prenatal	10	4.33
Psychological	36	15.58
Self-Reported Health Status	7	3.03
Internet-based	21	9.09
Social Media Images	1	0.43
Social Media Location	5	2.16
Social Media Metadata	4	1.73
Social Media Posts	12	5.19
Social Network	3	1.30
Web Search Metadata	1	0.43
Web Search Queries	12	5.19
Lifestyle	81	35.06
Alcohol	39	16.88
Diet	19	8.23
Physical Activity	26	11.26
Sleep	11	4.76
Smoking	58	25.11
Unspecified	4	1.73
Other Substance-use	13	5.63
Other Lifestyle	13	5.63
Environment	82	35.50
Air Quality	5	2.16
Any Satellite Imagery-derived	19	8.23
Biodiversity and Domestic Animals	2	0.87
Built Environment	8	3.46
Satellite	4	1.73
Other	4	1.73

Connectivity	4	1.73
Electrical Lighting ¹	1	0.43
General Environmental Exposures (not included in other categories)	9	3.90
Hazard	10	4.33
Land-use	2	0.87
Satellite	1	0.43
Other	1	0.43
Location	27	11.69
Meteorological	40	17.32
Surface Water Distribution/Flooding ¹	6	2.60
Topography	14	6.06
Satellite	12	5.19
Other	2	0.87
Vector/Reservoir Characteristics	9	3.90
Vegetation ¹	16	6.93
Water Composition	1	0.43
Other Satellite Imagery-derived	7	3.03
Population-level Disease or Healthcare Statistics	38	16.45
Socioeconomic and Demographic Factors	150	64.94
Adverse Adult Experiences/Trauma	5	2.16
Adverse Childhood Experiences	4	1.73
Age	112	48.48
Antisocial Behaviour	2	0.87
Economy Makeup	1	0.43
Education	33	14.29
Electricity	2	0.87
Employment	22	9.52
Garbage Collection	1	0.43
Healthcare System	5	2.16
Household Characteristics	10	4.33
Housing Structure	4	1.73
Human Development Index	1	0.43
Immigration Status	5	2.16
Income	24	10.39
Income Inequality	3	1.30
Language	2	0.87
Legal System	1	0.43
Literacy	2	0.87
Marital Status	21	9.09
Occupational Risk	10	4.33
Parental	3	1.30
Peer Group	1	0.43
Political Stability	1	0.43

Population and Population Density	11	4.76
Population Growth	2	0.87
Race/Ethnicity	29	12.55
Religion	3	1.30
Sanitation	5	2.16
Sex/Gender	95	41.13
Social Support	10	4.33
Unspecified	6	2.60
Vehicle Ownership	2	0.87
Water Supply Quality	5	2.16
Wealth	2	0.87
Other Socioeconomic and Demographic	29	12.55
Other Features	17	7.36
Not Reported	1	0.43

¹See supplementary table B for greater detail regarding feature categories

²Satellite-derived

Appendix A: Eligibility Criteria

The following types of articles were excluded:

- Reviews;
- Focused on a methodological development;
- Only included an abstract;
- Only used linear regression, logistic regression, generalized additive models, or other approaches not considered machine learning for the purpose of this review;
- Only applied models to diagnosis, treatment decisions, or prognosis of individuals who already had a disease;
- Only related to logistics, human resources, finance, or management involved in provision of public health services;
- Focused on occupational health, traffic accidents, or environmental monitoring, with no direct link to population health outcomes;
- Used smart home or home monitoring systems;
- Used advanced imaging or other expensive predictors that would be difficult or unsafe to scale to a population level;
- Focused on clinical decision support systems;
- Predicted adverse drug effects, except vaccines.

Appendix B: Examples of article titles removed during title screening

1. Improved classification of mangroves health status using hyperspectral remote sensing data
2. Diesel engine and propulsion diagnostics of a mini-cruise ship by using artificial neural networks
3. Relationship between benthic macroinvertebrate bio-indices and physicochemical parameters of water: A tool for water resources managers
4. Adaptive one-switch row-column scanning
5. Development of a distributed bearing health monitoring and assessing system
6. Neural networks based sensor validation and recovery methodology for advanced aircraft engines
7. Mining images in publicly-available cameras for homeland security
8. The human pulvinar and attentional processing of visual distractors
9. Text classification techniques in oil industry applications
10. Research on acoustic mechanical fault diagnosis method of high voltage circuit breaker based on improved EEMD

Appendix C: All studies included in the review.

- 1 Achrekar H, Gandhe A, Lazarus R, *et al.* Predicting Flu Trends using Twitter data. 2011. doi:10.1109/INFCOMW.2011.5928903
- 2 Adamou M, Antoniou G, Greasidou E, *et al.* Mining Free-Text Medical Notes for Suicide Risk Assessment. Proc. 10th Hell. Conf. Artif. Intell. 2018. doi:10.1145/3200947.3201020
- 3 Adams LJ, Bello G, Dumancas GG. Development and Application of a Genetic Algorithm for Variable Optimization and Predictive Modeling of Five-Year Mortality Using Questionnaire Data. *Bioinform Biol Insights* 2015;**9**:31–41. doi:<https://dx.doi.org/10.4137/BBI.S29469>
- 4 Agarwal A, Baechle C, Behara RS, *et al.* Multi-method approach to wellness predictive modeling. *J Big Data* 2016;**3**:1–23. doi:<http://dx.doi.org/10.1186/s40537-016-0049-0>
- 5 Agarwal V, Zhang L, Zhu J, *et al.* Impact of Predicting Health Care Utilization Via Web Search Behavior: A Data-Driven Analysis. *J Med Internet Res* 2016;**18**:e251.<http://ovidsp.ovid.com/ovidweb.cgi?T=JS&PAGE=reference&D=medp&NEWS=N&AN=27655225>
- 6 Agopian AJ, Lupo PJ, Tinker SC, *et al.* Working towards a risk prediction model for neural tube defects. *Birth Defects Res A Clin Mol Teratol* 2012;**94**:141–6. doi:<https://dx.doi.org/10.1002/bdra.22883>
- 7 Ahn C, Hwang Y, Park SK. Predictors of all-cause mortality among 514,866 participants from the Korean National Health Screening Cohort. *PLoS One* 2017;**12**. doi:<http://dx.doi.org/10.1371/journal.pone.0185458>
- 8 Aichele S, Rabbitt P, Ghisletta P. Illness and intelligence are comparatively strong predictors of individual differences in depressive symptoms following middle age. *Aging Ment Health* 2017;**;**1–10. doi:<https://dx.doi.org/10.1080/13607863.2017.1394440>
- 9 Akbulut A, Ertugrul E, Topcu V. Fetal health status prediction based on maternal clinical history using machine learning techniques. *Comput Methods Programs Biomed* 2018;**163**:87–100. doi:<http://dx.doi.org/10.1016/j.cmpb.2018.06.010>
- 10 Akhavan P, Karimi M, Pahlavani P, *et al.* Risk mapping of Cutaneous Leishmaniasis via a fuzzy C Means-based Neuro-Fuzzy inference system. 2014;**40**:19–23. doi:10.5194/isprsarchives-XL-2-W3-19-2014
- 11 Alby S, Shivakumar BL. A prediction model for type 2 diabetes risk among Indian women. *ARPN J Eng Appl Sci* 2016;**11**:2037–43.<https://www.scopus.com/inward/record.uri?eid=2-s2.0-84959387072&partnerID=40&md5=0fde9764a6290488b1c3472e2bbb5f7c> NS -
- 12 Allen T, Murray KA, Zambrana-Torrel C, *et al.* Global hotspots and correlates of emerging zoonotic diseases. *Nat Commun* 2017;**8**:1124. doi:<https://dx.doi.org/10.1038/s41467-017-00923-8>
- 13 Allore H, Tinetti ME, Araujo KLB, *et al.* A case study found that a regression tree outperformed multiple linear regression in predicting the relationship between impairments and Social and Productive Activities scores. *J Clin Epidemiol* 2005;**58**:154–61.<http://ovidsp.ovid.com/ovidweb.cgi?T=JS&PAGE=reference&D=med5&NEWS=N&AN=15680749>
- 14 Al-Mallah MH, Elshawi R, Ahmed AM, *et al.* Using Machine Learning to Define the Association between Cardiorespiratory Fitness and All-Cause Mortality (from the Henry Ford Exercise Testing Project). *Am J Cardiol* 2017;**120**:2078–84. doi:10.1016/j.amjcard.2017.08.029
- 15 Almeida AS, Werneck GL. Prediction of high-risk areas for visceral leishmaniasis using socioeconomic indicators and remote sensing data. *Int J Health Geogr* 2014;**13**. doi:10.1186/1476-072X-13-13
- 16 Alves EB, Costa CHN, de Carvalho FAA, *et al.* Risk Profiles for Leishmania infantum Infection in Brazil. *Am J Trop Med Hyg* 2016;**94**:1276–81. doi:10.4269/ajtmh.15-0513
- 17 Amini P, Ahmadiania H, Poorolajal J, *et al.* Evaluating the High Risk Groups for Suicide: A Comparison of Logistic Regression, Support Vector Machine, Decision Tree and Artificial Neural Network. *Iran J Public Health* 2016;**45**:1179–87.<https://www.ncbi.nlm.nih.gov/pmc/articles/PMC5149472/pdf/IJPH-45-1179.pdf> NS -
- 18 Amini P, Maroufizadeh S, Samani RO, *et al.* Factors Associated with Macrosomia among Singleton Live-births: A Comparison between Logistic Regression, Random Forest and Artificial Neural Network Methods. *Epidemiol Biostat Public Heal* 2016;**13**. doi:10.2427/11985
- 19 Anand A, Shakti D. Prediction of diabetes based on personal lifestyle indicators. 2015;**;**673–6. doi:10.1109/NGCT.2015.7375206
- 20 Anderson RT, Balkrishnan R, Camacho F. Risk classification of Medicare HMO enrollee cost levels using a decision-tree approach. *Am J Manag Care* 2004;**10**:89–98.<http://ovidsp.ovid.com/ovidweb.cgi?T=JS&PAGE=reference&D=med5&NEWS=N&AN=15011809>
- 21 Asensio-Cuesta S, Diego-Mas JA, Alcaide-Marzal J. Applying generalised feedforward neural networks to

- classifying industrial jobs in terms of risk of low back disorders. *Int J Ind Ergon* 2010;**40**:629–35. doi:10.1016/j.ergon.2010.04.007
- 22 Ayyagari R, Vekeman F, Lefebvre P, *et al.* Pulse pressure and stroke risk: development and validation of a new stroke risk model. *Curr Med Res Opin* 2014;**30**:2453–60. doi:10.1185/03007995.2014.971357
- 23 Azeez A, Obaromi D, Odeyemi A, *et al.* Seasonality and Trend Forecasting of Tuberculosis Prevalence Data in Eastern Cape, South Africa, Using a Hybrid Model. *Int J Environ Res Public Health* 2016;**13**. doi:10.3390/ijerph13080757
- 24 Bakar AA, Kefli Z, Abdullah S, *et al.* Predictive models for dengue outbreak using multiple rulebase classifiers. 2011;;5 pp. doi:10.1109/ICEEI.2011.6021830
- 25 Balaraman S, Schafer JJ, Tseng AM, *et al.* Plasma miRNA Profiles in Pregnant Women Predict Infant Outcomes following Prenatal Alcohol Exposure. *PLoS One* 2016;**11**. doi:http://dx.doi.org/10.1371/journal.pone.0165081
- 26 Bandyopadhyay S, Wolfson J, Vock DM, *et al.* Data mining for censored time-to-event data: a Bayesian network model for predicting cardiovascular risk from electronic health record data. *Data Min Knowl Discov* 2015;**29**:1033–69. doi:10.1007/s10618-014-0386-6
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Appendix D: Narrative Synthesis of Aspects of Discussion

Rationale for applying machine learning approaches mainly centered around it being “state of the art” or better suited to modeling complex data than regression. Machine learning was thought to be “state of the art” due to improved accuracy and deeper insights. Discussions of complex modeling focused on capturing non-linear relationships, interactions, and high-dimensionality.

When authors discussed model limitations, frequent concerns were an inadequate sample size, too few features, questionable generalizability, and a lack of interpretability. Aspects of the data other than sample size and feature number, such as potential measurement error or selection bias, were infrequently mentioned.

When discussing model implementation, many articles stated that predictive accuracy would be improved; but they did not frequently discuss how this could be translated to specific health-related policies or actions. Additionally, they rarely mentioned organizations and knowledge users that would be best suited to leverage the model.

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Predicting population health with machine learning: a scoping review

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PREDICTING POPULATION HEALTH WITH MACHINE LEARNING: A SCOPING REVIEW

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ABSTRACT (300 / 300 words)

Objective. To determine how machine learning has been applied to prediction applications in population health contexts. Specifically, to describe which outcomes have been studied, the data sources most widely used, and whether reporting of machine learning predictive models aligns with established reporting guidelines.

Design. A scoping review.

Data Sources. MEDLINE, EMBASE, CINAHL, ProQuest, Scopus, Web of Science, Cochrane Library, INSPEC, and ACM Digital Library were searched on July 18th, 2018.

Eligibility criteria. We included English articles published between 1980 and 2018 that used machine learning to predict population health-related outcomes. We excluded studies that only used logistic regression or were restricted to a clinical context.

Data extraction and synthesis. We summarized findings extracted from published reports, which included general study characteristics, aspects of model development, reporting of results, and model discussion items.

Results. Of 22 618 articles found by our search, 231 were included in the review. The United States (n=71, 30.74%) and China (n=40, 17.75%) produced the most studies. Cardiovascular disease (n=22, 9.52%) was the most studied outcome. The median number of observations was 5414 (interquartile range (IQR)=16543.5) and the median number of features was 17 (IQR=31). Health records (n=126, 54.5%) and investigator-generated data (n=86, 37.2%) were the most common data sources. Many studies did not incorporate recommended guidelines on machine learning and predictive modeling. Predictive discrimination was commonly assessed using area under the receiver operator curve (n=98, 42.42%) and calibration was rarely assessed (n=22, 9.52%).

Conclusions. Machine learning applications in population health have concentrated on regions and diseases well-represented in traditional data sources, infrequently using big data. Important aspects of model development were under-reported. Greater use of big data and reporting guidelines for predictive modeling could improve machine learning applications in population health.

Registration. Registered on the Open Science Framework on July 17th, 2018 (available at: <https://osf.io/rnqe6/>).

Strengths and limitations of this study

- Our review is one of the first syntheses of machine learning applications in population and public health.
- We used a robust search strategy, including nine peer-reviewed databases, grey literature, and reference searching, to comprehensively describe the literature.
- We compared reported study characteristics to established predictive modeling reporting guidelines, which provide an objective measure of the quality of reporting.
- Since both machine learning and population health have broad definitions, there may be some relevant articles that were not included.
- Given our focus on prediction, we could not address many other important intersections of machine learning and population health, such as surveillance and health promotion.

Word Count: 3663

INTRODUCTION

Predictive models have a long history in clinical medicine. One well-known example is the Framingham risk score, which was first developed in 1967.[1] Such models have proliferated throughout clinical practice to inform management and interventions, including preventive approaches. More recently, researchers have developed prediction models beyond individual clinical applications, for population health uses.[2,3] While there is no universal definition of population health, it generally encompasses “the health outcomes of a group of individuals, including the distribution of such outcomes within the group.”[4] Similarly to clinical medicine, population-level models can be used to identify high-risk groups, directing the implementation of preventive interventions. Additionally, population health prediction models can inform policymakers about future disease burden and help to assess the impact of public health actions. Thus far, most predictive modeling in both medicine and population health has used parametric statistical regression models. More recently, there has been increasing interest in the use of a broader range of machine learning methods for prediction tasks.[5–7]

Machine learning can be loosely defined as the study and development of algorithms that learn from data with little or no human assistance.[8] These approaches have been increasingly applied in the past two decades as a result of the enabling growth of big data reserves and computational power.[9] Recent machine learning applications to prediction in population health contexts include forecasting childhood lead poisoning,[10] yellow fever incidence,[11] and the onset of suicidal ideation.[12]

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3 The distinction between machine learning algorithms and parametric regression models is
4 debated.[13] Regression models tend to impose more structure on the data, requiring greater
5
6 human input for the verification of distributional assumptions and incorporation of domain
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8 knowledge in choosing the input parameters.[14] Algorithms employed in machine learning
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10 often derive more structure directly from the data, making fewer distributional assumptions
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12 about the data or variables. The literature remains divided on the relative advantages of more
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14 traditional approaches compared to newer methods;[15] however, given the wide variation in
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16 applications and the data used in these examples, broad assessments of superiority are often not
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18 appropriate. Also, there are debates regarding the differences in developing and validating
19
20 machine learning approaches for health applications.[15,16]
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28 Population health applications of prediction models are relatively new compared to clinical
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30 applications; correspondingly, the role of machine learning in these applications has been far less
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32 studied and discussed in the health literature. The goals of our review are to determine how
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34 machine learning has been applied to prediction in population health, the nature of the models
35
36 and data used, and how the models have been developed. We also sought to assess how well the
37
38 published literature aligns with recommended guidelines for reporting of predictive models and
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40 machine learning, by extracting features related to model development and performance that are
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42 highlighted by two such guidelines.[16,17]
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49 **METHODS**

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3 We based our scoping review on the framework proposed by Arksey and O'Malley[18] and
4 refined by the Joanna Briggs Institute.[19] We also followed the more recent Preferred Reporting
5 Items for Systematic Reviews and Meta-analysis Extension for Scoping Reviews.[20] Our study
6 protocol was registered on the Open Science Framework on July 17th, 2018 (available at:
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12 <https://osf.io/rnqe6/>).

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17 Our initial goal was to scope out all machine learning applications in population health.
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19 However, the screening process identified a much larger number of publications than anticipated.
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21 Consequently, to describe the subject area comprehensively, we restricted our scope to articles
22 predicting future outcomes.
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28 **Search Strategy**

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32 Our search strategy consisted of peer-reviewed literature databases, grey literature, and reference
33 searches. First, we searched nine interdisciplinary, indexed databases (MEDLINE, EMBASE,
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CINAHL, ProQuest, Scopus, Web of Science, Cochrane Library, INSPEC, and ACM Digital
Library) on July 18th, 2018 for papers published between 1980 and 2018. Our search was
informed by consultation with a health science librarian, a machine learning textbook,[21] and a
similar registered review.[15] Supplementary Table A includes the full MEDLINE search
strategy and filters, and serves an example search query for all database searches.

Our grey literature search included Google Scholar and Google. We developed a Google Scholar
search based on terms related to 'machine learning' and 'population health', which was refined
based on the relevance of initial results. The first 200 results were included in screening. A

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2
3 similar approach was used for the general Google search, which we restricted to the first 30
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5 results. We examined relevant websites for publications. Results were limited to articles
6
7 published on or before the date of the peer-reviewed literature search. Finally, we searched the
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9 references of relevant reviews for additional articles. Most of these reviews were identified
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11 during screening.
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14 15 16 **Eligibility Criteria**

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21 We included articles if they used machine learning to develop a predictive model that could be
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23 applied in a population health context. Therefore, we excluded articles where the model was
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25 trained primarily on people with a pre-existing disease. We also excluded articles that were only
26
27 indirectly related to population health; for example, traffic accident models that did not predict a
28
29 health outcome. Studies predicting individual outcomes were included if the approach was
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31 determined to be scalable to a population level. Finally, articles using only logistic regression
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33 were excluded. See Appendix A for the full eligibility criteria.
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40 In order to manage the scope, articles were excluded if their full text could not be retrieved with
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42 our institutional licenses and if they were not written in English. Finally, articles published prior
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44 to 1980 were excluded as earlier machine learning investigators lacked comparable amounts of
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46 digitized data, software, and computational resources.
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50 51 **Screening Process**

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3 Initially, individual reviewers screened titles for obvious irrelevance to the review topic (JDM
4 and EB). An example of an obviously irrelevant topic would be a paper describing the *machine*
5
6 *health* lifespan of a piece of industrial equipment; specific examples of articles removed at this
7
8 stage are listed in Appendix B. Then, we imported remaining references into Covidence
9
10 systematic review management software.[22] Two reviewers screened the abstracts of remaining
11
12 articles (JDM, EB, MO, and DF). Prior to evaluating full texts using all eligibility criteria, we
13
14 then screened out articles that did not focus on a prediction application (JDM, EB, MO). Finally,
15
16 two reviewers screened the full text of remaining articles (JDM, EB, MO). Conflicts were
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18 resolved by discussion between at least two reviewers.
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26 **Data Extraction and Synthesis**

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30 Individual authors extracted article data (JDM, EB, MO, and DF). We based our extraction items
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32 on features identified in a recent biomedical guideline for reporting of machine learning
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34 predictive models [16] and on the transparent reporting of a multivariable prediction model for
35
36 individual prognosis or diagnosis (TRIPOD) statement.[17] Major extraction categories
37
38 identified from these guidelines included general study characteristics (e.g. geographic location
39
40 and sample size), model development (e.g. algorithms used and type of validation), results (e.g.
41
42 discrimination and calibration measures), and model discussion (e.g. practical costs of errors and
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44 implementation). See Supplementary Table B for a description of each extraction item.
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51 We computed descriptive statistics for all extraction items. For categorical extracted features
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53 (e.g. whether or not unstructured text was used, the method of validation used), we calculated the
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55 total number and percent of all studies in a particular category. For continuous extracted features
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(e.g. number of observations in the study sample), we calculated the median value and the interquartile range (range between quartile 1 and quartile 3 in the value distribution). We also completed a narrative synthesis of discussion elements based on the text of included manuscripts.

Patient and Public Involvement Statement

There was no patient or public involvement in this study.

RESULTS

We initially retrieved 16 172 articles, after removing duplicates (Figure 1). We excluded 6494 articles after title screening, 7860 after abstract screening, 1453 when screening out non-prediction articles, and 121 after full-text screening. This resulted in 231 articles being included in the final review (Appendix C).

General Study Characteristics

The number of articles published in the population health prediction area that used machine learning increased dramatically after 2007 (Supplementary Figure A). Studies were undertaken worldwide, with the largest representation from the United States (US) (n=71, 30.74%) and China (n=40, 17.75%) (Table 1). Relatively few articles came from Oceania (n=2, 0.87%), Africa (n=5, 2.16%), and the Americas outside of the US (n=13, 5.63%).

Characteristic*	Number of Articles [#]	Percent of Articles**
Region		
United States	71	30.74%
Asia Excluding China	41	17.75%
China	40	17.32%
Europe	36	15.58%
Americas Excluding United States	13	5.63%
Africa	5	2.16%
Oceania	2	0.87%
Multi-region	15	6.49%
Not Reported	8	3.46%
Year published		
before 1990	1	0.4%
1990-1999	3	1.3%
2000-2004	13	5.6%
2005-2009	18	7.8%
2010-2014	70	30.3%
2015-2018	126	54.5%
Outcome level[†]		
Individual Risk Prediction	139	60.17%
Population Risk prediction	92	39.83%
Number of observations	Median = 5414 [#]	IQR = 16543.5 ^{**}
Not reported	72	31.2%
Number of features	Median = 17 [#]	IQR = 31 ^{**}
Not reported	59	25.5%
Used any unstructured text		
Yes	24	10.4%

Characteristic*	Number of Articles [#]	Percent of Articles**
No	207	89.6%
Machine learning model was compared with other statistical methods	111	48.1%
Reported data pre-processing[‡]		
Yes	160	69.3%
No	71	30.7%
Reported method of feature selection		
Yes	164	71.0%
No	67	29.0%
Reported hyper-parameter search		
Yes	114	49.4%
No	117	50.6%
Method of Validation		
Holdout	112	48.5%
Cross-validation or bootstrap	84	36.4%
External	15	6.5%
Not reported	32	13.9%
Reported descriptive statistics[§]		
Yes	140	60.6%
No	91	39.4%
Discussed the practical costs of prediction errors[¶]		
Yes	36	15.6%
No	195	84.4%
Stated rationale for using machine learning		
Yes	179	77.5%
No	52	22.5%

Characteristic*	Number of Articles [#]	Percent of Articles**
Discussed model usability		
Yes	91	39.4%
No	140	60.6%
Stated model limitations		
Yes	161	69.7%
No	70	30.3%
Discussed model implementation		
Yes	184	79.7%
No	47	20.3%
Dataset Availability by Study		
Closed	149	64.5%
Public	42	18.2%
Closed and Public	38	16.5%
Unknown	1	0.4%

*Refer to Supplementary Table A for a description of each characteristic and rationales for including some elements.

[†]Individual risk prediction refers to studies that developed models to predict the health outcomes of individuals, while population risk prediction refers to studies that developed models to predict aggregated population-level health outcomes.

[‡]Whether any aspects of data cleaning or pre-processing were reported. Examples include how missing data was handled, whether log transformations were done, and if derived variables were generated.

[§]Included a broad array of descriptive statistics such as sample population demographics, feature distributions, and outcome distributions.

[¶]Whether the article discussed the relative risks of false negative and false positive results based on their predictive model in contexts where it might be used.

^{||}Closed refers to datasets that were not immediately available in the public domain or were not identifiable as such.

[#]In rows where the characteristic being measured is an integer count (e.g. number of features), this column refers to the median value.

^{**}In rows where the characteristic being measured is an integer count (e.g. number of features), this column refers to the interquartile range (IQR; quartile 3 – quartile 1).

Table 1: Summary statistics of included articles

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3 The median number of observations in each article was 5414 (interquartile range (IQR)=16
4 543.5) and the median number of features (i.e. independent variables) used was 17 (IQR=31)
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6 (Table 1). Seventy-two studies (31.2%) did not report the number of observations. These studies
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8 often used data from reportable disease databases, which do not necessarily have a firm sampling
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10 frame, making ascertainment of the number of observations difficult.
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16 **Algorithms**

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21 The most frequently used machine learning algorithms were neural networks (n=95, 41.13%),
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23 followed by support vector machines (n=59, 25.54%), single tree-based methods (n=52,
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25 22.51%), and random forests (n=48, 20.78%) (Supplementary Table C). About half of the
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27 articles made a comparison with statistical methods (n=111, 48.1%), which were generally
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29 logistic regression or autoregressive integrated moving average models (Table 1).
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35 **Outcomes**

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39 Non-communicable disease outcomes were assessed by many articles (n=95, 41.13%), with
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41 communicable diseases (n=76, 32.90%) and non-disease outcomes (n=60, 25.97%) studied
42
43 somewhat less often. The outcome most frequently predicted was cardiovascular disease (n=22,
44
45 9.52%) (Figure 2). Other commonly forecasted non-communicable disease outcomes were
46
47 suicidality (n=13, 5.63%), cancer (n=12, 5.19%), and perinatal health (n=12, 5.19%). Influenza
48
49 (n=15, 6.49%) and dengue fever (n = 14, 6.06%) were the most predicted communicable disease
50
51 outcomes. Aside from non-communicable and communicable disease, mortality (n=13, 5.63%)
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53 and healthcare utilization (n=14, 6.06%) were also frequently predicted.
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Data

Data sources were usually structured (n=207, 89.6%) and closed, i.e. not publicly available (n=189, 81.8%) (Table 1). In general, high-dimensional data with many observations, such as multi-linked electronic medical records (EMRs) or internet-based data, may offer the most value for machine learning applications. These data types were represented in some of the articles captured, for which the most frequently reported data sources were health records (n=126, 54.5%) and investigator-generated (e.g. cohort studies) (n=86, 37.2%) (Table 2). A large proportion of studies (n=42, 18.2%) used an environmental data source (e.g. satellite imagery), mostly for prediction of infectious disease. Government databases (n=32, 13.9%) and internet-based data (n=21, 9.1%) were less frequently used. Among studies from China and the US, 80.0% and 67.6% respectively used health records data, whereas 54.5% of studies overall used these data sources (Supplementary Figure B).

Sources of Data Used*	Number	Percent
Environmental	42	18.2%
Geographical Information Database	12	5.2%
Meteorological/Air Quality Datasets	32	13.9%
Satellite Imagery	21	9.1%
Health Records Database	126	54.5%
Clinical Record Database [†]	46	19.9%
Disease Registry	2	0.9%
Population Health Survey	15	6.5%
Reportable Disease Database	42	18.2%

Other Health Records Database	30	13.0%
Government Database	32	13.9%
Census	11	4.8%
Vital Statistics	13	5.6%
Other Government Database	14	6.1%
HealthMap	3	1.3%
Private Insurance Data	9	3.9%
Private Insurance Claims	9	3.9%
Private Insurance Questionnaire	3	1.3%
Internet-based	21	9.1%
Search engine	12	5.2%
Social Media	12	5.2%
Investigator-generated[‡]	86	37.2%
Public Repositories[§]	19	8.2%
Health Organization Reports[¶]	5	2.2%
Not Reported	6	2.6%

^{*}Categories are not mutually exclusive.

[†]Any dataset produced primarily for the purpose of delivering clinical care, such as electronic medical records and administrative healthcare databases produced by hospitals.

^{**}Any datasets resulting from researcher-driven studies such as randomized controlled trials, cohort studies, and case-control studies.

[§]Any freely available datasets such as MIMIC or the UC Irvine Machine Learning Repository.

[¶]Health-related reports, typically including disease burden estimates, produced by non-governmental or governmental organizations such as the World Health Organization.

Table 2: Data sources

Features

The median number of features used in a machine learning algorithm was 17 (IQR = 31; Table 1). The frequency of specific feature categories used are shown in Supplementary Figure C and

Supplementary Table D. Biomedical and sociodemographic features were frequently used (Supplementary Figure C). Of these, the most commonly used were disease history (43.3%), age (48.5%), and sex/gender (41.1%). Among lifestyle features, smoking was the most frequently used (25.1%) and of environmental features, meteorology was common (17.3%). Social media posts (5.2%) and web search queries (5.2%) were not often used. In general, most studies focused on features typical of clinical prediction models, such as subject demographics, behaviours, and medical histories. We observed limited use of other data, such as unstructured text or image-based features, which are difficult to parse using traditional statistical approaches and could benefit more from machine learning applications

Model Development and Validation

The majority of articles reported how data pre-processing (n=160, 69.3%) and feature selection (n=164, 71%) were done (Table 1). Fewer authors reported how hyperparameters were selected (n=114, 49.4%). Most studies used a holdout method of validation (n=112, 48.5%), fifteen (6.5%) externally validated their models, and thirty-two (13.9%) did not report how models were validated.

Performance Metrics

Most articles reported a prediction discrimination metric (n=172, 74.46%), which quantifies a model's ability to correctly rank-order individuals (Table 3).[23] Discrimination is a useful performance metric in cases where classification is the primary goal, including many machine learning-relevant tasks such as image recognition. The most common discrimination metrics

employed were area under the receiver operator curve (n=98, 42.42%), accuracy (n=76, 32.90%), and recall (n=68, 29.44%).

In clinical and public health settings, accurate prediction of outcome probabilities is important to the practical utility of a tool, so assessing model calibration is very important.

Few articles in our study reported a measure of calibration (n=21, 9.09%), which describes how well a model predicts the absolute probability of outcomes (table 3).^[23] Calibration was mostly assessed with graphing methods (n=9, 3.90%) and Hosmer-Lemeshow statistics (n=8, 3.46%).

Some articles also reported a measure of overall model fit (n=77, 33.33%). Overall performance was usually measured with a form of mean error, such as root mean squared error (n=35, 15.15%).

Prediction Performance Metrics Used	Number	Percent
Any overall performance metric	77	33.33%
RMSE	35	15.15%
MSE	26	11.26%
MAE	24	10.39%
MAPE	23	9.96%
R2*	19	8.23%
Correlation	8	3.46%
AIC or BIC	8	3.46%
Other performance metric [†]	21	9.09%
Any discrimination metric	172	74.46%
Area under the curve [‡]	98	42.42%
Accuracy [§]	76	32.90%
Recall [¶]	68	29.44%
Precision	39	16.88%

F statistics	10	4.33%
Likelihood Ratio**	4	1.73%
Youden Index	3	1.30%
Manual or visual comparison	3	1.30%
Other discrimination metric††	4	1.73%
Any calibration metric	21	9.09%
Manual or visual comparison‡‡	9	3.90%
Hosmer-Lemeshow	8	3.46%
Observed/Expected	5	2.16%
Other calibration metric§§	3	1.30%
Any reclassification metric	6	2.60%
Net Reclassification Index	5	2.16%
Integrated Discrimination Improvement	3	1.30%

RMSE = Root Mean Squared Error; MSE = Mean Squared Error; MAE = Mean Absolute Error; MAPE = Mean Absolute Percentage Error; AIC = Akaike Information Criterion; BIC = Bayesian Information Criterion.

*Includes R2 and pseudo-R2 metrics.

†Includes penalty error, Total Sum of Squares, proportional reduction in error, overall prediction error, specific prediction error, Nash-Sutcliffe, Root Mean Squared Percentage Error (2), mean relative absolute error, Analysis of Variance F-stat, 2LogLikelihood, relative efficiency, deviance, Ljung-Box test, mean absolute deviation, standard error, Mean Percentage Error, Brier score, and log score.

‡Includes c-statistic, s-index, area under ROC / AUC.

§Includes accuracy, misclassification, and error rate.

¶Includes sensitivity, specificity, true/false positive, and true/false negative.

||Includes positive predictive value, negative predictive value, and precision.

**Includes positive/negative LR.

††Includes G-means (2), k-statistic, Matthews correlation coefficient.

‡‡Includes calibration plots.

§§Includes mean bias (from Bland-Altman plot), calibration factoring, and Calibration statistic.

Table 3: Prediction Performance Metrics

Study Discussion and Narrative Synthesis

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3 Most articles included some discussion of their rationale for using machine learning (n=179,
4 77.5%), although some articles did not mention or explain their rationale (n = 52, 22.5%) (Table
5
6
7 1). Rationale for applying machine learning approaches mainly focused on being “state of the
8
9 art” or better suited to modeling complex data than regression.
10

11
12 Most articles also had some discussion of the limitations of their study (n=161, 69.7%), and how
13
14 the model might be implemented (n=184, 79.7%) (Table 1). Frequent concerns were an
15
16 inadequate sample size, too few features, questionable generalizability, and a lack of
17
18 interpretability. When discussing model implementation, many articles stated that predictive
19
20 accuracy would be improved; but they did not frequently discuss how this could be translated to
21
22 specific health-related policies or actions.
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25
26 Less than half of the articles discussed model usability (n=91, 39.4%); that is, whether and how
27
28 the model could practically be used in a relevant context. This is an important reporting
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30 component of the TRIPOD statement (“Discuss the potential clinical use of the model and
31
32 implications for future research”) and is relevant for understanding real-world applications of
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34 prediction models.[17] Also, only a small number discussed the costs of prediction errors in real-
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36 world contexts (n=36, 15.6%).
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40 See Appendix D for further narrative synthesis of discussion reporting items.
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44 **DISCUSSION**

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49 Our results show that machine learning is increasingly being applied to make predictions related
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51 to population health. However, applications of machine learning to population health prediction
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53 tasks have not capitalized fully on the opportunities presented by emerging big data resources
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3 and efficient machine learning algorithms. Furthermore, reporting of these models often does not
4 align with established guidelines for reporting of prediction models, which limits their ability to
5 be critically appraised, compared with existing statistical models, or implemented in clinical or
6 public health practice.
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10 *Applications of Machine Learning Prediction Models*

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12 Nearly half of the included studies were conducted in the US or China. Both countries produce
13 the greatest number of scientific publications in general;^[24] however, they also likely benefited
14 from robust health data infrastructures. The US has rapidly digitized much of its healthcare
15 system, resulting in large EMRs linked with government data through public-private
16 partnerships, including processes to make these data available to researchers.^[25,26] Both the US
17 and China made greater use of health records and less use of investigator-generated data relative
18 to other regions, which may have made machine learning projects more tractable. They also used
19 more internet-based data, which typically includes many observations and is high-dimensional,
20 making it amenable to machine learning methods. We noted that studies from Oceania, Africa,
21 and the Americas (outside of the US) were limited. This may be partly due to less availability of
22 traditional sources of structured health data. However, given that machine learning methods can
23 incorporate non-traditional data sources, there is the potential to expand use of these methods
24 even when structured health data is unavailable.
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47 We found that a wide range of population health outcomes have been the focus of machine
48 learning prediction models. However, relative to morbidity and mortality, multiple outcome
49 categories like cancer, human immunodeficiency virus, dementia, gastroenteritis, pneumococcal
50 disease, perinatal health, tuberculosis, and malaria appear understudied.^[27] Many of these
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3 conditions are most prevalent in regions with decreased access to traditional health data, perhaps
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5 stymieing research. If machine learning methods are used to leverage novel data sources for
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7 research in these regions, it could enable greater study of neglected diseases.
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10 Most investigators did not analyze a large number of observations and features. We observed a
11
12 high reliance on electronic health records and investigator-generated data, including the use of
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14 relatively small study cohorts. Small study sample sizes or narrow data collection associated with
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16 these data sources can make it difficult to achieve high sample sizes or high dimensional data,
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18 which may impact machine learning algorithm performance. Specifically, the use of smaller
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20 investigator-generated datasets may affect the performance of studied models, as machine
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22 learning algorithms generally require a high number of observations relative to features.[28]
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26 Additionally, most studies focused on features typical of clinical prediction models, such as
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28 biomedical factors and limited aspects of broader socioeconomic or environmental determinants
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30 of health. We also observed infrequent use of unstructured data and wearable data for prediction
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32 purposes. A reliance on small datasets and traditional numbers and types of features is unlikely
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34 to fully leverage any benefits of machine learning. This may be contributing to the small
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36 differences frequently seen between parametric regression and machine learning model
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38 performance. Greater use of linked population-level databases, large EMRs, internet data, and
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40 unstructured features would likely improve these approaches.
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47 *Reporting of Machine Learning Prediction Models*

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49 Based on the elements of model development that we studied, adherence to existing machine
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51 learning[16] and prediction model[17] guidelines appears limited. Most articles did not report
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53 their method of hyper-parameter selection, discuss practical costs of prediction errors, or
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3 consider model usability, which are needed for transparency and model assessment. Many
4 studies did not report the number of features included, method of validation, method of feature
5 selection, or any performance metric. Given these issues, it would be difficult or impossible to
6 compare many of these machine learning models with existing approaches. However, we
7 acknowledge that existing guidelines were not available when many included studies were
8 published. Future work should apply existing guidance,[16] including from TRIPOD,[17] and
9 anticipate the forthcoming TRIPOD-ML statement.[29]
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22 Lastly, we noted that included studies rarely assessed predictive performance in terms of
23 calibration, which refers to a model's ability to accurately predict the absolute probability of
24 outcomes.[23] In contrast, discrimination measures of predictive performance quantify a model's
25 ability to correctly rank-order individuals. Many traditional machine learning tasks, such as
26 image recognition, often have a high signal to noise ratio. In these cases, discrimination may be a
27 suitable lone performance metric, as the algorithm can achieve near perfect performance.
28 Conversely, health outcomes tend to be more stochastic. As a result, accurate prediction of
29 probabilities is more important.[23] Models can have good predictive discrimination, but poor
30 calibration, making them less useful in practice, particularly for population health applications. A
31 further issue is that many measures of discrimination, such as accuracy and recall, artificially
32 impose a threshold for calling events. Thresholds should ideally be ascertained by decision-
33 makers based on their cost-utility curves.[23] Overall, applications of machine learning in
34 population health would benefit from greater use of calibration performance metrics.
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51 *Strengths and Limitations of this Review*

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3 A strength of our study is that we addressed an understudied area, the intersection of machine
4 learning and population health. Additionally, prediction is an application with untapped potential
5 in population health, and where machine learning has the potential to make significant
6 improvements. Our study also employed a comprehensive search strategy, including numerous
7 multidisciplinary peer-reviewed databases, alongside a grey literature search. Furthermore, we
8 applied insights from the field of clinical prediction modeling to population health and machine
9 learning. Finally, given the focus on prediction, we were able to take a comprehensive approach
10 to data extraction and synthesis.
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24 In terms of limitations, concentrating on prediction prevented us from exploring applications of
25 machine learning to other important aspects of population health, such as disease surveillance.
26 These should be the focus of future research. Our review was also limited by including only
27 English articles and articles with available full text, which may have introduced selection bias.
28 Because of the broad scope of this review, and inconsistent reporting of model development and
29 validation in reviewed articles, we were unable to carry out a critical appraisal of the literature
30 and are unable to comment significantly on the overall performance of published machine
31 learning population health prediction tools. This would be of great value for understanding the
32 clinical and population health relevance of machine learning prediction tools. Lastly, the two
33 main concepts underlying our review, machine learning and population health, are not
34 universally defined. As a result, we may have excluded articles that may be relevant to these
35 fields.
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51 *Research Recommendations and Conclusion*

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3 This was the first scoping review specifically focused on machine learning prediction in
4 population health applications. Predictive modeling in population health can help to inform
5 preventive interventions, anticipate future disease burden, and assess the impact of health
6 policies and programs. Advances in machine learning offer opportunities to improve these
7 models, particularly when incorporating big data. Countries with substantial EMR-use and
8 government database linkage such as Finland, Singapore, and Denmark[30] likely have untapped
9 potential for machine learning research. This is still a nascent field, but based on our findings
10 more research in Oceania, Africa, and South America would also be particularly beneficial.
11 Diseases with a high global burden of disease that were underrepresented in our findings include
12 malaria, tuberculosis, and dementia, which may be opportune for further study.[31] Additionally,
13 future machine learning projects could incorporate larger datasets and more non-traditional
14 features. Greater use of resources such as HealthMap, social media, web search patterns, remote
15 sensing, and WHO reports would enable more work in regions without formal data sources and
16 enrich research in others. Another largely untapped prospect is using machine learning and high-
17 dimensional data to incorporate richer representations of the social determinants of health.
18 Opportunities should continue to grow as governments increasingly digitize their health service
19 records and link databases to both health and non-health data. Overall, as applications of
20 machine learning in population health develop, adherence to existing guidance[16,17,29] will
21 improve our ability to assess and advance machine learning applications. We hope that our
22 results will help to inform future research in this area, including the development of guidelines
23 for machine learning applications in population health. Finally, it will be important to evaluate
24 the impact of prediction models on decisions made in population health and the practice of
25 public health.
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CONTRIBUTORS

JDM contributed to the design of the study and led the literature search, article screening, data extraction, analysis, and writing of the manuscript. EB contributed to the design of the study, the literature search, article screening, data extraction, and analysis. MO contributed to article screening, data extraction, analysis, and writing of the manuscript. TP and VG contributed to the design of the study and supervised work. DF contributed to article screening and data extraction. KK contributed to the design of the study. LCR led the design of the study. All authors interpreted study results and contributed to drafting of the manuscript.

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COMPETING INTERESTS

None declared.

PATIENT CONSENT FOR PUBLICATION

Not required.

ETHICS APPROVAL

Not required as only prior published research was included in the review.

DATA AVAILABILITY

The full data extraction table used for this review will be made publicly available after publication with no end date on Mendeley Data (DOI: 10.17632/7rrz9xrp2j.1).

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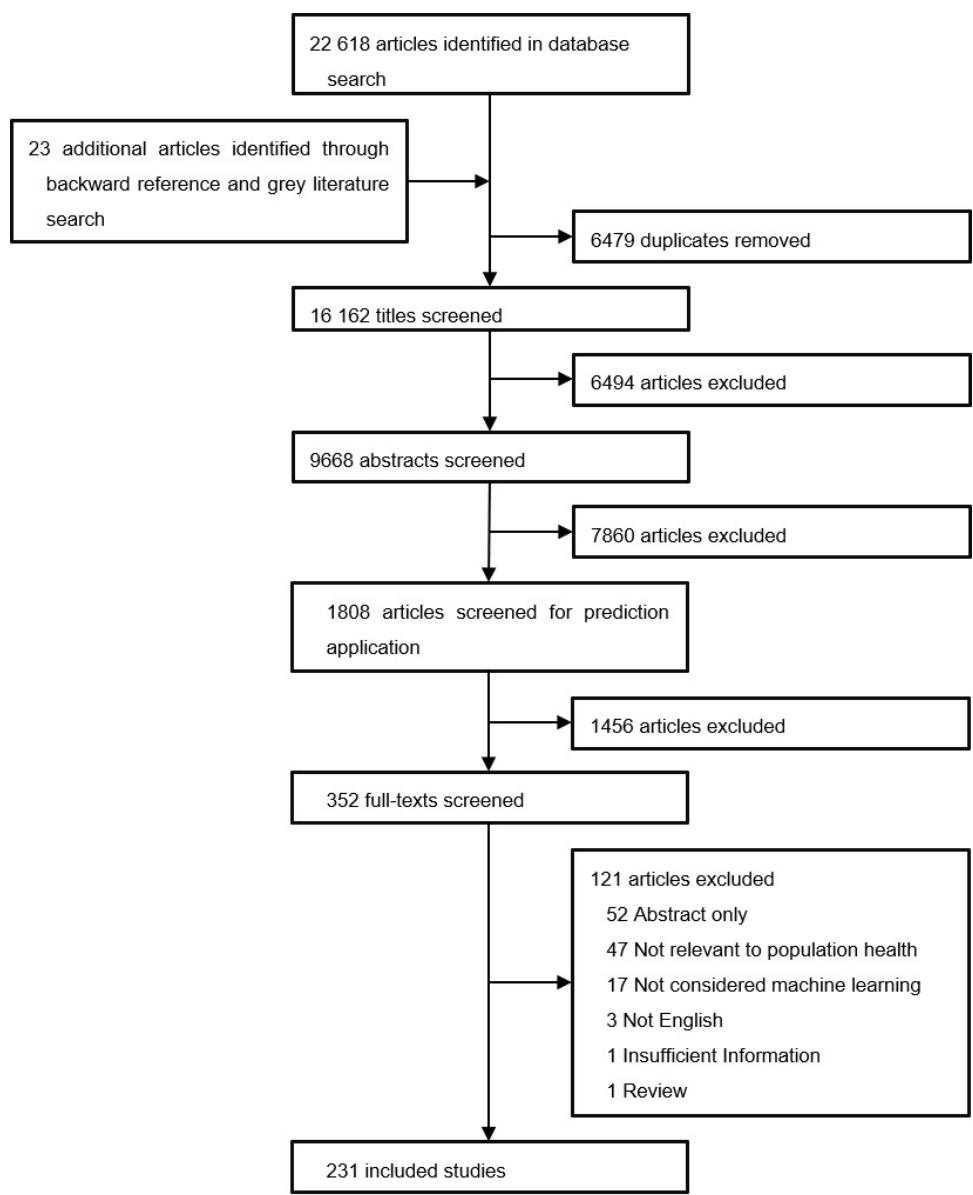
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20 **FIGURES LEGENDS**

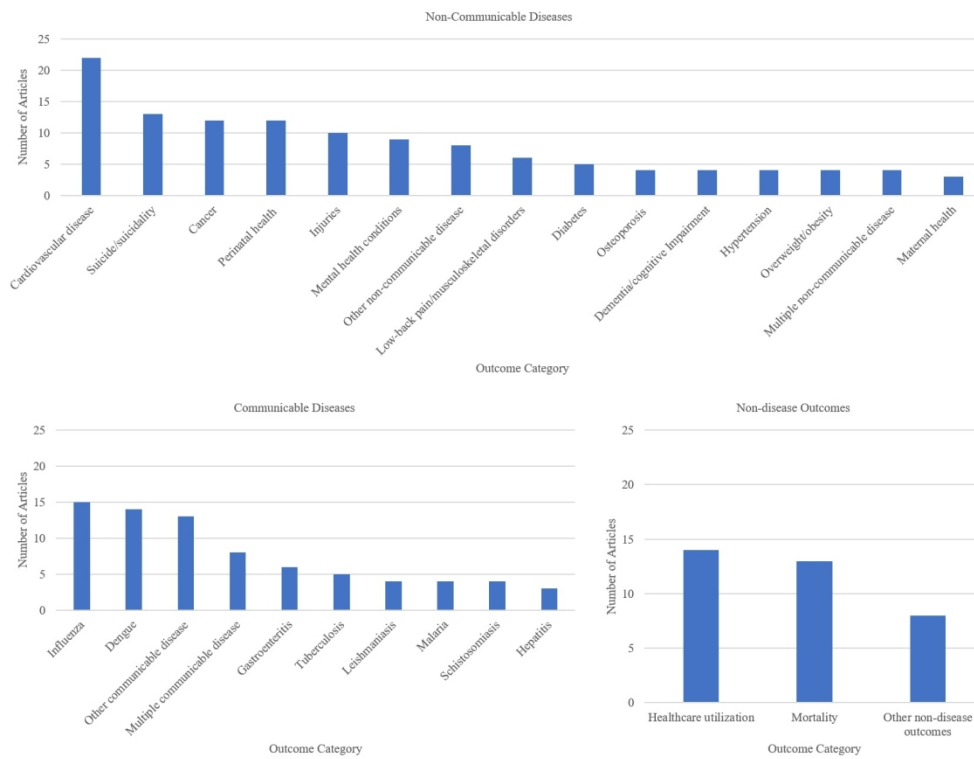
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22 **Figure 1:** PRISMA flowchart of article screening process.
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25 **Figure 2:** Number of articles by outcome.
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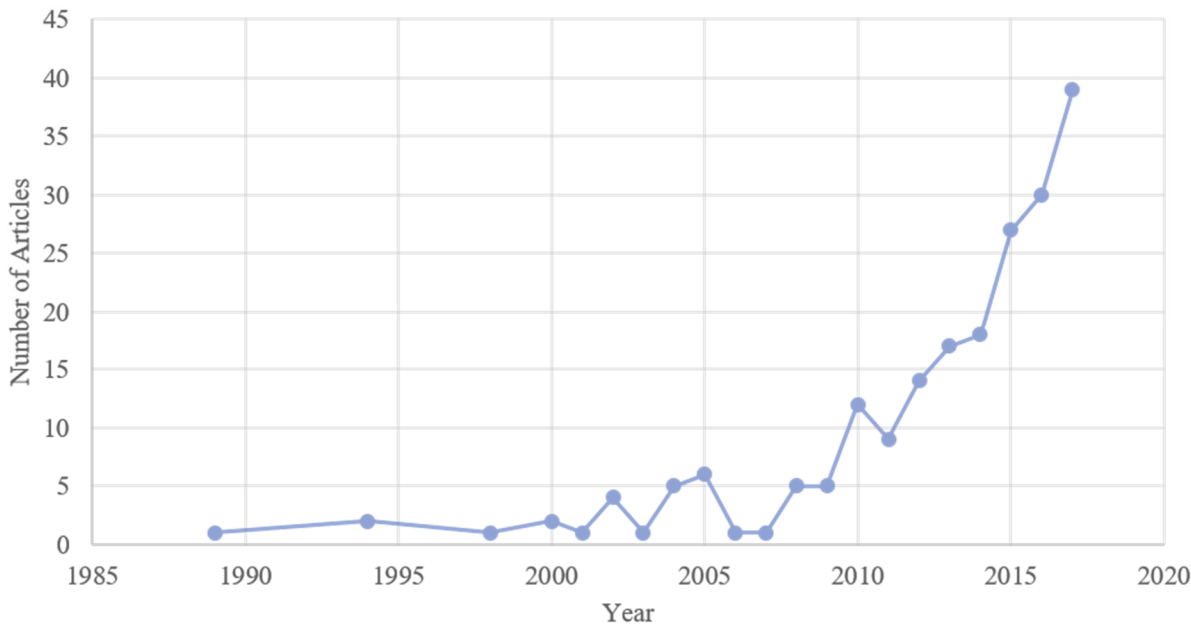


144x180mm (144 x 144 DPI)



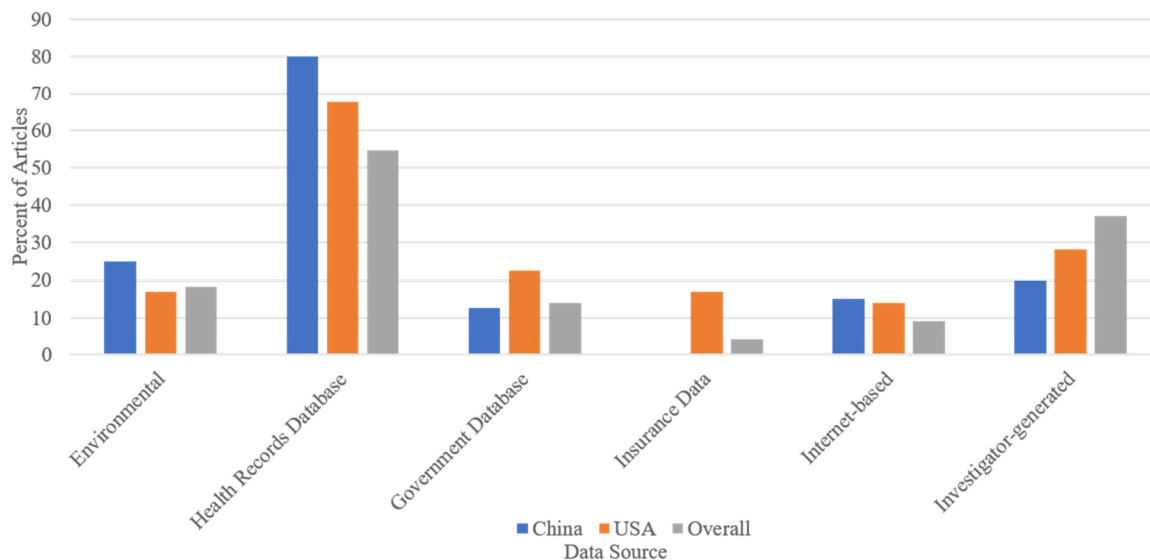
298x230mm (144 x 144 DPI)

Supplementary Figure A: Number of articles by publication year. Articles from 2018 are not plotted in this figure because the review does not include all studies published that year.



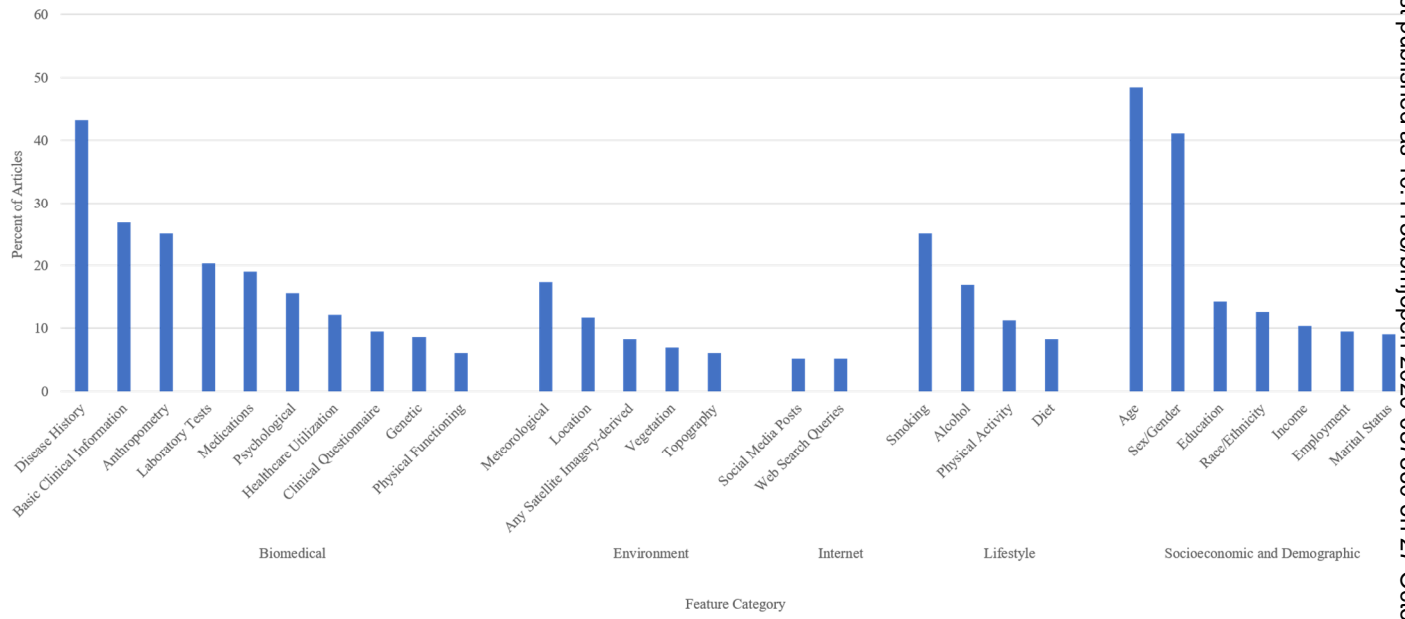
review only

Supplementary Figure B: Data Source by selected region.



review only

Supplementary Figure C: Most commonly used feature categories.



review only

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Supplementary Table A: MEDLINE search query¹

Machine Learning Terms	Population Health Terms
1. Exp Artificial Intelligence/	24. Exp Population Health/
2. Exp "neural networks (computer)"/	25. Exp Population Surveillance/
3. Support vector machine*.kf,tw	26. Exp Health Equity/
4. Neural net*.kf,tw	27. Health status/
5. Perceptron*.kf,tw	28. Health status disparities/
6. Deep learning.kf,tw	29. Public health systems research/
7. Random forest*.kf,tw	30. "Social determinants of health"/
8. Lasso*.kf,tw	31. Health surveys/
9. Gaussian mixture*.kf,tw	32. Health status indicators/
10. Bayesian network*.kf,tw	33. "global burden of disease"/
11. Classification tree*.kf,tw	34. Global health/
12. Regression tree*.kf,tw	35. Environmental health/
13. Relevance vector machine*.kf,tw	36. Harm reduction/
14. Nearest neighbo*.kf,tw	37. Public health informatics/
15. Probability estimation tree*.kf,tw	38. Community medicine/
16. Elastic net*.kf,tw	39. Public health/
18. Naive bayes.kf,tw	40. Epidemiology/
19. Genetic algorithm*.kf,tw	41. Preventive medicine/
20. Artificial intelligence.kf,tw	42. Occupational medicine/
21. Machine learning.kf,tw	43. Environmental medicine/
22. Statistical learning.kf,tw	44. Public health practice/
23. /or 1-22	45. Preventive health services/
	46. Health promotion/
	47. public health.kf,tw
	48. population health.kf,tw
	49. health promot*.kf,tw
	50. population surveillance.kf,tw
	51. health surveillance.kf,tw
	52. health equity.kf,tw
	53. preventive medicine.kf,tw
	54. health protection.kf,tw
	55. disease prevention.kf,tw
	56. social determinant* of health.kf,tw
	57. health determinant*.kf,tw
	58. determinant* of health.kf,tw
	59. occupational medicine.kf,tw
	60. community medicine.kf,tw
	61. epidemiolog*.kf,tw
	62. health status*.kf,tw
	63. global health.kf,tw

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	64. environmental health.kf,tw
	65. harm reduction.kf,tw
	66. environmental medicine.kf,tw
	67. /or 24-66
	68. 23 and 67

¹Limited to articles published in 1980 or after.

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Supplementary Table B: Data Extraction Field Descriptions

Data Extraction Field	Description
Title	The article titles.
First Author	The last name and first initial of the first listed author of each article
Year of Publication	The year of publication noted for each article.
Outcome level	One of two categories: <ol style="list-style-type: none"> <i>Population risk prediction</i>: the aggregated outcome of a whole population was predicted <i>Individual risk prediction</i>: outcomes of individual participants were predicted
Outcome	<p>Selected from the following, which are not mutually exclusive, as some articles predicted multiple outcomes:</p> <p>Non-communicable Disease</p> <ol style="list-style-type: none"> <i>Cardiovascular disease</i>: any disease characterized by atherosclerosis and resulting ischemia, including myocardial infarction and stroke <i>Suicide/suicidality</i> <i>Cancer</i> <i>Perinatal health</i>: including pre-term birth, fetal alcohol spectrum disorder, congenital heart disease, growth failure, and neural tube defects <i>Mental health conditions</i> <i>Osteoporosis</i> <i>Low-back pain and other musculoskeletal disorders</i> <i>Diabetes</i> <i>Dementia and cognitive Impairment</i> <i>Hypertension</i> <i>Injuries</i>: including fractures, falls, traffic injury, and foreign body injuries <i>Overweight and obesity</i> <i>Maternal health</i>: including fertility, pregnancy risk, and severe maternal morbidity <i>Multiple non-communicable disease</i> <i>Other non-communicable disease</i>: including liver disorders, Crohn's disease, glaucoma, dental caries, and lead poisoning <p>Communicable Disease</p> <ol style="list-style-type: none"> <i>Influenza</i> <i>Dengue</i> <i>Gastroenteritis</i> <i>Tuberculosis</i> <i>Leishmaniasis</i> <i>Malaria</i> <i>Schistosomiasis</i> <i>Hepatitis</i>: of viral origin <i>Multiple communicable disease</i> <i>Other communicable disease</i>: including zika, hand food and mouth disease, leptospirosis, yellow fever, West Nile, and typhoid fever <p>Non-disease Outcomes</p> <ol style="list-style-type: none"> <i>Mortality</i> <i>Healthcare utilization</i> <i>Other non-disease outcomes</i>: including health behaviours, vitamin d status, and wellness score
Region	<p>Categorized based on Organisation for Economic Cooperation and Development (OECD) region except for the United States and China, which were given their own categories due to the high number of publications. One of the following:</p> <ol style="list-style-type: none"> <i>Africa</i> <i>Americas except for the United States</i> <i>Asia except for China</i> <i>China</i> <i>Europe</i> <i>Oceania</i> <i>United States</i> <i>Multi-region</i> <i>Other/Unknown</i>
Study Setting	One of two categories:

	<ol style="list-style-type: none"> 1. <i>Clinical</i>: when data was collected in any type of clinical setting 2. <i>Community</i>: when data was collected in a community setting
Data Source Categories	<p>Selected from the following categories, which were not mutually exclusive, and often more than one was used:</p> <ol style="list-style-type: none"> 1. <i>Geographical Information Database</i>: any dataset containing basic map-based spatial information such as distances and topography 2. <i>Meteorological/Air Quality Datasets</i> 3. <i>Satellite Imagery</i>: examples include the moderate resolution imaging spectroradiometer (MODIS) and the Shuttle Radar Topography Mission (SRTM) 4. <i>Clinical Record Database</i>: any dataset produced primarily for the purpose of delivering clinical care, such as electronic medical records and administrative healthcare databases produced by hospitals 5. <i>Disease Registry</i>: a dataset maintained to monitor and/or provide care for a specific disease 6. <i>Population Health Survey</i>: a regular epidemiological survey administered periodically to assess the health of populations 7. <i>Reportable Disease Database</i>: a dataset containing reports of diseases for which it is mandatory for healthcare providers to report 8. <i>Other Health Records Database</i>: any other health records dataset not encompassed in other categories, including various surveillance systems 9. <i>Census</i> 10. <i>Vital Statistics</i>: information regularly collected by governments regarding births and deaths 11. <i>Other Government Database</i>: other governmental datasets including socioeconomic and demographic information 12. <i>HealthMap</i>: a public health surveillance system using natural language processing to analyze informal data sources such as online news, individual reports, expert-curated discussions 13. <i>Private Insurance Claims</i>: including medical, hospital, and prescription drug claims 14. <i>Private Insurance Questionnaires</i> 15. <i>Internet Search</i>: including the number of searches of certain key terms and meta data such as the location of the searches 16. <i>Social Media</i>: both posts and metadata 17. <i>Investigator-generated</i>: any datasets resulting from researcher-driven studies such as randomized controlled trials, cohort studies, and case-control studies 18. <i>Public Repositories</i>: any freely available datasets such as MIMIC 19. <i>Health Organization Reports</i>: health-related reports, typically including disease burden estimates, produced by non-governmental or governmental organizations such as the World Health Organization 20. <i>Not Reported</i>
Feature Categories	<p>Selected from the following categories, which were not mutually exclusive, as often more than one category was used (if more than one instance of a feature category was found in an article it was only counted once):</p> <p>Biomedical</p> <ol style="list-style-type: none"> 1. <i>Anthropometry</i>: measurements of the human body such as height and weight 2. <i>Basic Clinical Information</i>: information typically collected during a brief physician encounter such as a focused medical history and physical examination, including blood pressure 3. <i>Basic Medical Tests</i>: any test requiring somewhat specialized equipment such as an electrocardiogram 4. <i>Clinical Questionnaire</i>: a standardized questionnaire administered in a clinical context such as the Montreal Cognitive Assessment or Patient Health Questionnaire-9 5. <i>Disease History</i>: information regarding present and/or past diagnoses of an individual 6. <i>Genetic</i> 7. <i>Healthcare Utilization</i> 8. <i>Instrumental Activities of Daily Living</i>: features relating to an individual's daily functioning in areas such as cooking and shopping 9. <i>Laboratory Tests</i>: any features derived from human specimens requiring specialized equipment for analysis, such as hematological and microbiological results 10. <i>Medical Imaging</i> 11. <i>Medications</i> 12. <i>Physical Functioning</i>: features including the presence of any physical disabilities or the status of activities of daily living 13. <i>Prenatal</i>: relevant aspects of the period before birth such as the use of prenatal vitamins or the results of routine lab results

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4	14. <i>Psychological</i> : features including mood or anxiety symptoms
5	15. <i>Self-Reported Health Status</i>
6	Internet-based
7	16. <i>Social Media Images</i>
8	17. <i>Social Media Location</i> : either aggregated or individual
9	18. <i>Social Media Metadata</i> : any information other than the content of social media posts, such as the frequency of general posts and time of posting
10	19. <i>Social Media Posts</i> : social media post content
11	20. <i>Social Network</i> : the interconnections among individuals in a social media platform
12	21. <i>Web Search Metadata</i> : any aspects of web searches other than their content
13	22. <i>Web Search Queries</i> : the content of web search queries either individual or aggregated
14	Lifestyle
15	23. <i>Alcohol</i>
16	24. <i>Diet</i>
17	25. <i>Physical Activity</i>
18	26. <i>Sleep</i>
19	27. <i>Smoking</i>
20	28. <i>Unspecified</i>
21	29. <i>Other Substance-use</i>
22	30. <i>Other Lifestyle</i>
23	Environment
24	31. <i>Air Quality</i>
25	32. <i>Any Satellite Imagery-derived</i>
26	33. <i>Biodiversity and Domestic Animals</i>
27	34. <i>Satellite-based Built Environment</i>
28	35. <i>Other Built Environment</i>
29	36. <i>Connectivity</i> : the ease of access to large urban centers and/or general services
30	37. <i>Electrical Lighting (satellite-based)</i>
31	38. <i>General Environmental Exposures (not included in other categories)</i>
32	39. <i>Hazard</i> : characteristics of an external hazard such as the presence of lighting on a roadway
33	40. <i>Satellite-based Land-use</i>
34	41. <i>Other Land-use</i>
35	42. <i>Location</i>
36	43. <i>Meteorological</i>
37	44. <i>Surface Water Distribution/Flooding (satellite-based)</i>
38	45. <i>Satellite-based Topography</i>
39	46. <i>Other Topography</i>
40	47. <i>Vector/Reservoir Characteristics</i> : including mosquito surveillance numbers and the population of non-human primates in the case of yellow fever
41	48. <i>Vegetation (satellite-based)</i> : such as the normalized difference vegetation index (NDVI)
42	49. <i>Water Composition</i>
43	50. <i>Other Satellite Imagery-derived</i>
44	51. Population Disease or Healthcare Statistics
45	Socioeconomic and Demographic
46	52. <i>Adverse Adult Experiences/Trauma</i>
47	53. <i>Adverse Childhood Experiences</i>
48	54. <i>Age</i>
49	55. <i>Antisocial Behaviour</i>
50	56. <i>Economy Makeup</i> : such as the number of individuals working in various types of occupations
51	57. <i>Education</i>
52	58. <i>Electricity</i>
53	59. <i>Employment</i>
54	60. <i>Garbage Collection</i>
55	61. <i>Healthcare System</i> : such as the availability of universal, public healthcare
56	62. <i>Household Characteristics</i> : the number of individuals in the household and their ages
57	63. <i>Housing Structure</i> : aspects of the physical structure of housing such as the number of units and age of the building
58	64. <i>Human Development Index</i>
59	65. <i>Immigration Status</i>
60	66. <i>Income</i>
	67. <i>Income Inequality</i>
	68. <i>Language</i>
	69. <i>Legal System</i>
	70. <i>Literacy</i>

	<p>71. <i>Marital Status</i></p> <p>72. <i>Occupational Risk</i>: including risk factors for low-back pain such as prolonged sitting or injury from repetitive movements</p> <p>73. <i>Parental</i>: including disciplinary styles and the amount of time spent at home and number of parent-child activities</p> <p>74. <i>Peer Group</i>: behaviours of peer group</p> <p>75. <i>Political Stability</i></p> <p>76. <i>Population and Population Density</i></p> <p>77. <i>Population Growth</i></p> <p>78. <i>Race/Ethnicity</i></p> <p>79. <i>Religion</i></p> <p>80. <i>Sanitation</i>: availability of sewage systems</p> <p>81. <i>Sex/Gender</i></p> <p>82. <i>Social Support</i></p> <p>83. <i>Unspecified</i></p> <p>84. <i>Vehicle Ownership</i>: at population level</p> <p>85. <i>Water Supply Quality</i></p> <p>86. <i>Wealth</i></p> <p>87. <i>Other Socioeconomic and Demographic</i></p> <p>88. Other Features</p> <p>89. Not Reported</p>
Number of Datasets Used	The number of distinct datasets used regardless of the number of sources.
Dataset Availability	<p>Selected from the following categories:</p> <ol style="list-style-type: none"> 1. <i>Public</i>: all the datasets used by article authors were publicly available 2. <i>Closed</i>: all the datasets were not publicly available or appeared not to be available 3. <i>Closed and Public</i>: the datasets used were a mix of available and not available
Any Unstructured Text Used	Natural human language was included in the model as a feature with no initial ordinal/nominal structure imposed.
Number of Observations	The number of individuals or other units of observations (such as countries) included in the predictive model. If multiple subsets of the data and/or distinct datasets were used for different models, the largest number was used.
Machine Learning Algorithm Type	<p>The algorithm type used to build the predictive model, with multiple types often used in the same article. Algorithms were only counted once when used in each article, even if used to build multiple different models in the same article. Selected from the following categories:</p> <ol style="list-style-type: none"> 1. <i>Neural Networks</i>: includes deep learning/deep neural networks as well as other simpler neural networks 2. <i>Support Vector Machine</i> 3. <i>Single Tree-based Methods</i>: includes classification trees, regression trees, and decision trees 4. <i>Random Forest</i> 5. <i>Least Absolute Shrinkage and Selection Operator (LASSO)</i> 6. <i>Bayesian Networks</i>: includes naïve bayes 7. <i>Feature Selection Methods</i>: includes k-means clustering and genetic algorithms; these were often used as a pre-processing step and in a few cases this was the only use of machine learning (i.e. a machine learning model was not used to build the predictive model itself) 8. <i>Boosted Tree-based Methods</i>: includes gradient boosting and boosted trees 9. <i>K-Nearest Neighbour</i> 10. <i>Elastic Net</i> 11. <i>Ridge Regression</i> 12. <i>Other</i>: includes association rule learning, single task learning, multitask learning, rough set classifier, associative classification, bagging, partial least squares discriminant analysis, Just-Add-Data Bio Tool, super learner, particle swarm optimization, ant colony optimization, Isomap, PCA, Disease State Index, Stacking, kernel conditional density estimation, stepwise deletion, conditional random fields, contrast mining, grammatical evolution, Learning from Examples Using ROugh Sets, AUtoregression with exogenous outputs, and natural language processing
Compared with Other Statistical Methods	Whether the machine learning method's predictive performance was compared with a traditional parametric statistical regression model such as logistic regression (yes/no).
Reported Data Pre-processing	Whether any aspects of data cleaning or pre-processing were reported (yes/no). Examples include how missing data was handled, whether log transformations were done, and if derived variables were generated. Missing data and all model development processes have been identified as important to report by TRIPOD.[1]
Reported Method of Feature Selection	Whether the method of feature selection was reported (yes/no). When there is a high number of features initially, this is usually done using algorithmic, domain knowledge-

	informed, or mixed approaches. Feature selection is an important element of reporting as identified by TRIPOD.[1]
Number of Features	The number of features included in the final prediction model after feature selection. If multiple models were used in one article, the largest number of features was chosen.
Reported Hyper-parameter Search	Whether the process for determining the hyper-parameters of the machine learning model, such as the number of features used to build each tree in a random forest, was reported (yes/no). This is an important aspect of model development[2], and thus considered an important element to report by the TRIPOD statement.[1]
Method of Validation	How the authors validated the predictive performance of their model, selected from one of the following categories: <ol style="list-style-type: none"> 1. <i>Holdout</i>: the dataset was divided into two parts; one part was used to train the model and the other was used to test the model 2. <i>Cross-validation and bootstrap</i>: the dataset was either divided into more than two parts and repeatedly trained and tested on different parts of the dataset or random sampling with replacement was used to train the model 3. <i>External</i>: the model was tested on a completely separate dataset
Reported Descriptive Statistics	Whether the article reported any descriptive statistics regarding their sample (yes/no). We considered a broad array of descriptive statistics including sample population demographics, feature distributions, and outcome distributions. These are all important reporting elements according to TRIPOD.[1]
Calibration Metrics	The types of calibration predictive performance metrics used to evaluate models, which could be more than one. Calibration refers to a model's ability to accurately predict absolute probabilities of the outcome occurring.[3] One or more of the following categories was selected if a calibration metric was used: <ol style="list-style-type: none"> 1. <i>Manual or visual comparison</i>: includes calibration plots 2. <i>Hosmer-Lemeshow</i> 3. <i>Observed/Expected</i>: is a ratio or comparison of observed and predicted/expected probabilities 4. <i>Other calibration metric</i>: includes mean bias (from Bland-Altman plot), calibration factoring, calibration statistic
Discrimination Metrics	The types of discrimination predictive performance metrics used to evaluate models, which could be more than one. Discrimination refers to a model's ability to correctly rank-order individuals according to their likelihood of developing the outcome.[3] One or more of the following categories was selected if a discrimination metric was used: <ol style="list-style-type: none"> 1. <i>Area under the curve</i>: meaning receiver operator curve; also includes c-statistic and s-index 2. <i>Accuracy</i>: includes accuracy, misclassification, and error rate 3. <i>Recall</i>: includes sensitivity, specificity, true/false positive, and true/false negative 4. <i>Precision</i>: includes positive predictive value, negative predictive value, and precision 5. <i>F statistics</i> 6. <i>Likelihood Ratio</i>: includes both positive and negative likelihood ratios 7. <i>Youden Index</i> 8. <i>Manual or visual comparison</i> 9. <i>Other discrimination metric</i>: includes G-means, k-statistic, and Matthews correlation coefficient
Overall Goodness of Fit Metrics	The types of overall goodness of fit performance metrics used to evaluate models, which could be more than one. Overall goodness of fit refers to a model's predictions' concordance with observed outcomes. One or more of the following categories was selected if an overall performance metric was used: <ol style="list-style-type: none"> 1. <i>Root mean squared error</i> 2. <i>Mean squared error</i> 3. <i>Mean absolute error</i> 4. <i>Mean absolute percentage error</i> 5. R^2: includes pseudo-R2s 6. <i>Correlation</i> 7. <i>Akaike Information Criterion or Bayesian Information Criterion</i> 8. <i>Other performance metric</i>: includes penalty error, total sum of squares, proportional reduction in error, overall prediction error, specific prediction error, Nash-Sutcliffe, root mean squared percentage error, mean relative absolute error, analysis of variance F-stat, -2LogLikelihood, relative efficiency, deviance, Ljung-Box test, mean absolute deviation, standard error, Brier score, log score, and mean percentage error
Did Machine Learning Models Outperform Traditional Methods?	Whether the machine learning-based predictive models outperformed the statistical parametric regression models based on the performance metrics supplied by the authors (yes/no). However, this should not be taken to mean that the difference in model performance was reliable or valid. Often, important performance metrics and essential aspects of model development were not reported, making accurate comparisons difficult.

Discussed the Practical Costs of Prediction Errors	Whether the article discussed the relative risks of false negative and false positive results based on their predictive model in contexts where it might be used (yes/no). These costs are important for determining the usefulness and application of predictive models.[3]
Stated Rationale for Using Machine Learning	Whether the article stated any reasons for using a machine learning approach instead of a statistical parametric regression approach (yes/no).
Rationale for Using Machine Learning - Free Text	Reviewers included article quotations and summaries in this section to capture different rationales for using machine learning. Reviewers attempted to only extract free text regarding each specific type of rationale once
Discussed Model Usability	Whether the article discussed any aspect of how the model could be practically used in a relevant context (yes/no).
Stated Model Limitations	Whether the article discussed any potential limitations of the research (yes/no).
Model limitations - Free Text	Reviewers included article quotations and summaries in this section to capture different reported limitations. Reviewers attempted to only extract free text regarding each specific type of limitation once.
Discussed Model Implementation	Whether the article included discussion of any consequences of model implementation such as potential clinical, population-health, and policy-level impacts (yes/no).
Model Implementation - Free Text	Reviewers included article quotations and summaries in this section to capture different reported consequences of model implementation. Reviewers attempted to only extract free text regarding each specific type of implementation impact once.

Supplementary Table C: Types of machine learning algorithms used.

Types of Algorithms	Number	Percent
Neural Networks*	95	41.13%
Support Vector Machine	59	25.54%
Single tree-based methods [†]	52	22.51%
Random Forest	48	20.78%
LASSO	25	10.82%
Bayesian Networks [‡]	23	9.96%
Feature selection methods [§]	20	8.66%
Boosted tree-based methods	19	8.23%
K-Nearest Neighbour	19	8.23%
Elastic Net	9	3.90%
Ridge regression	5	2.16%
Other	22	9.52%

*Includes deep neural networks.

[†]Includes CART, decision trees.

[‡]Includes naive bayes.

[§]Includes cluster methods (e.g. k-means clustering) and genetic algorithms.

^{||}Includes gradient boosting and boosted trees.

^{||}Including (all algorithms used once unless otherwise specified) association rule learning (n=3), single task learning, multitask learning, rough set classifier, associative classification, bagging, partial least squares discriminant analysis, Just-Add-Data Bio Tool, super learner, particle swarm optimization, ant colony optimization, isomap, principal components analysis, disease state Index, stacking, kernel conditional density estimation, stepwise deletion, conditional random fields, contrast mining, grammatical evolution, Learning from Examples Using ROugh Sets, AUtoregression with exogenous outputs, and natural language processing (n=2).

Supplementary Table D: Detailed feature categories included in studies.

Feature Category	Number of Articles	Percent
Biomedical	141	61.04
Anthropometry	58	25.11
Basic Clinical Information	62	26.84
Basic Medical Tests	10	4.33
Clinical Questionnaire	22	9.52
Disease History	100	43.29
Genetic	20	8.66
Healthcare Utilization	28	12.12
Instrumental Activities of Daily Living	6	2.60
Laboratory Tests	47	20.35
Medical Imaging	10	4.33
Medications	44	19.05
Physical Functioning	14	6.06
Prenatal	10	4.33
Psychological	36	15.58
Self-Reported Health Status	7	3.03
Internet-based	21	9.09
Social Media Images	1	0.43
Social Media Location	5	2.16
Social Media Metadata	4	1.73
Social Media Posts	12	5.19
Social Network	3	1.30
Web Search Metadata	1	0.43
Web Search Queries	12	5.19
Lifestyle	81	35.06
Alcohol	39	16.88
Diet	19	8.23
Physical Activity	26	11.26
Sleep	11	4.76
Smoking	58	25.11
Unspecified	4	1.73
Other Substance-use	13	5.63
Other Lifestyle	13	5.63
Environment	82	35.50
Air Quality	5	2.16
Any Satellite Imagery-derived	19	8.23
Biodiversity and Domestic Animals	2	0.87
Built Environment	8	3.46
Satellite	4	1.73
Other	4	1.73

Connectivity	4	1.73
Electrical Lighting ¹	1	0.43
General Environmental Exposures (not included in other categories)	9	3.90
Hazard	10	4.33
Land-use	2	0.87
Satellite	1	0.43
Other	1	0.43
Location	27	11.69
Meteorological	40	17.32
Surface Water Distribution/Flooding ¹	6	2.60
Topography	14	6.06
Satellite	12	5.19
Other	2	0.87
Vector/Reservoir Characteristics	9	3.90
Vegetation ¹	16	6.93
Water Composition	1	0.43
Other Satellite Imagery-derived	7	3.03
Population-level Disease or Healthcare Statistics	38	16.45
Socioeconomic and Demographic Factors	150	64.94
Adverse Adult Experiences/Trauma	5	2.16
Adverse Childhood Experiences	4	1.73
Age	112	48.48
Antisocial Behaviour	2	0.87
Economy Makeup	1	0.43
Education	33	14.29
Electricity	2	0.87
Employment	22	9.52
Garbage Collection	1	0.43
Healthcare System	5	2.16
Household Characteristics	10	4.33
Housing Structure	4	1.73
Human Development Index	1	0.43
Immigration Status	5	2.16
Income	24	10.39
Income Inequality	3	1.30
Language	2	0.87
Legal System	1	0.43
Literacy	2	0.87
Marital Status	21	9.09
Occupational Risk	10	4.33
Parental	3	1.30
Peer Group	1	0.43
Political Stability	1	0.43

Population and Population Density	11	4.76
Population Growth	2	0.87
Race/Ethnicity	29	12.55
Religion	3	1.30
Sanitation	5	2.16
Sex/Gender	95	41.13
Social Support	10	4.33
Unspecified	6	2.60
Vehicle Ownership	2	0.87
Water Supply Quality	5	2.16
Wealth	2	0.87
Other Socioeconomic and Demographic	29	12.55
Other Features	17	7.36
Not Reported	1	0.43

¹See supplementary table B for greater detail regarding feature categories

²Satellite-derived

Appendix A: Eligibility Criteria

The following types of articles were excluded:

- Reviews;
- Focused on a methodological development;
- Only included an abstract;
- Only used linear regression, logistic regression, generalized additive models, or other approaches not considered machine learning for the purpose of this review;
- Only applied models to diagnosis, treatment decisions, or prognosis of individuals who already had a disease;
- Only related to logistics, human resources, finance, or management involved in provision of public health services;
- Focused on occupational health, traffic accidents, or environmental monitoring, with no direct link to population health outcomes;
- Used smart home or home monitoring systems;
- Used advanced imaging or other expensive predictors that would be difficult or unsafe to scale to a population level;
- Focused on clinical decision support systems;
- Predicted adverse drug effects, except vaccines.

Appendix B: Examples of article titles removed during title screening

1. Improved classification of mangroves health status using hyperspectral remote sensing data
2. Diesel engine and propulsion diagnostics of a mini-cruise ship by using artificial neural networks
3. Relationship between benthic macroinvertebrate bio-indices and physicochemical parameters of water: A tool for water resources managers
4. Adaptive one-switch row-column scanning
5. Development of a distributed bearing health monitoring and assessing system
6. Neural networks based sensor validation and recovery methodology for advanced aircraft engines
7. Mining images in publicly-available cameras for homeland security
8. The human pulvinar and attentional processing of visual distractors
9. Text classification techniques in oil industry applications
10. Research on acoustic mechanical fault diagnosis method of high voltage circuit breaker based on improved EEMD

Appendix C: All studies included in the review.

- 1 Achrekar H, Gandhe A, Lazarus R, *et al.* Predicting Flu Trends using Twitter data. 2011. doi:10.1109/INFCOMW.2011.5928903
- 2 Adamou M, Antoniou G, Greasidou E, *et al.* Mining Free-Text Medical Notes for Suicide Risk Assessment. Proc. 10th Hell. Conf. Artif. Intell. 2018. doi:10.1145/3200947.3201020
- 3 Adams LJ, Bello G, Dumancas GG. Development and Application of a Genetic Algorithm for Variable Optimization and Predictive Modeling of Five-Year Mortality Using Questionnaire Data. *Bioinform Biol Insights* 2015;**9**:31–41. doi:https://dx.doi.org/10.4137/BBI.S29469
- 4 Agarwal A, Baechle C, Behara RS, *et al.* Multi-method approach to wellness predictive modeling. *J Big Data* 2016;**3**:1–23. doi:http://dx.doi.org/10.1186/s40537-016-0049-0
- 5 Agarwal V, Zhang L, Zhu J, *et al.* Impact of Predicting Health Care Utilization Via Web Search Behavior: A Data-Driven Analysis. *J Med Internet Res* 2016;**18**:e251. http://ovidsp.ovid.com/ovidweb.cgi?T=JS&PAGE=reference&D=medp&NEWS=N&AN=27655225
- 6 Agopian AJ, Lupo PJ, Tinker SC, *et al.* Working towards a risk prediction model for neural tube defects. *Birth Defects Res A Clin Mol Teratol* 2012;**94**:141–6. doi:https://dx.doi.org/10.1002/bdra.22883
- 7 Ahn C, Hwang Y, Park SK. Predictors of all-cause mortality among 514,866 participants from the Korean National Health Screening Cohort. *PLoS One* 2017;**12**. doi:http://dx.doi.org/10.1371/journal.pone.0185458
- 8 Aichele S, Rabbitt P, Ghisletta P. Illness and intelligence are comparatively strong predictors of individual differences in depressive symptoms following middle age. *Aging Ment Health* 2017;**;**1–10. doi:https://dx.doi.org/10.1080/13607863.2017.1394440
- 9 Akbulut A, Ertugrul E, Topcu V. Fetal health status prediction based on maternal clinical history using machine learning techniques. *Comput Methods Programs Biomed* 2018;**163**:87–100. doi:http://dx.doi.org/10.1016/j.cmpb.2018.06.010
- 10 Akhavan P, Karimi M, Pahlavani P, *et al.* Risk mapping of Cutaneous Leishmaniasis via a fuzzy C Means-based Neuro-Fuzzy inference system. 2014;**40**:19–23. doi:10.5194/isprsarchives-XL-2-W3-19-2014
- 11 Alby S, Shivakumar BL. A prediction model for type 2 diabetes risk among Indian women. *ARPN J Eng Appl Sci* 2016;**11**:2037–43. https://www.scopus.com/inward/record.uri?eid=2-s2.0-84959387072&partnerID=40&md5=0fde9764a6290488b1c3472e2bbb5f7c NS -
- 12 Allen T, Murray KA, Zambrana-Torrel C, *et al.* Global hotspots and correlates of emerging zoonotic diseases. *Nat Commun* 2017;**8**:1124. doi:https://dx.doi.org/10.1038/s41467-017-00923-8
- 13 Allore H, Tinetti ME, Araujo KLB, *et al.* A case study found that a regression tree outperformed multiple linear regression in predicting the relationship between impairments and Social and Productive Activities scores. *J Clin Epidemiol* 2005;**58**:154–61. http://ovidsp.ovid.com/ovidweb.cgi?T=JS&PAGE=reference&D=med5&NEWS=N&AN=15680749
- 14 Al-Mallah MH, Elshawi R, Ahmed AM, *et al.* Using Machine Learning to Define the Association between Cardiorespiratory Fitness and All-Cause Mortality (from the Henry Ford Exercise Testing Project). *Am J Cardiol* 2017;**120**:2078–84. doi:10.1016/j.amjcard.2017.08.029
- 15 Almeida AS, Werneck GL. Prediction of high-risk areas for visceral leishmaniasis using socioeconomic indicators and remote sensing data. *Int J Health Geogr* 2014;**13**. doi:10.1186/1476-072X-13-13
- 16 Alves EB, Costa CHN, de Carvalho FAA, *et al.* Risk Profiles for Leishmania infantum Infection in Brazil. *Am J Trop Med Hyg* 2016;**94**:1276–81. doi:10.4269/ajtmh.15-0513
- 17 Amini P, Ahmadinia H, Poorolajal J, *et al.* Evaluating the High Risk Groups for Suicide: A Comparison of Logistic Regression, Support Vector Machine, Decision Tree and Artificial Neural Network. *Iran J Public Health* 2016;**45**:1179–87. https://www.ncbi.nlm.nih.gov/pmc/articles/PMC5149472/pdf/IJPH-45-1179.pdf NS -
- 18 Amini P, Maroufizadeh S, Samani RO, *et al.* Factors Associated with Macrosomia among Singleton Live-births: A Comparison between Logistic Regression, Random Forest and Artificial Neural Network Methods. *Epidemiol Biostat Public Heal* 2016;**13**. doi:10.2427/11985
- 19 Anand A, Shakti D. Prediction of diabetes based on personal lifestyle indicators. 2015;**;**673–6. doi:10.1109/NGCT.2015.7375206
- 20 Anderson RT, Balkrishnan R, Camacho F. Risk classification of Medicare HMO enrollee cost levels using a decision-tree approach. *Am J Manag Care* 2004;**10**:89–98. http://ovidsp.ovid.com/ovidweb.cgi?T=JS&PAGE=reference&D=med5&NEWS=N&AN=15011809
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Appendix D: Narrative Synthesis of Aspects of Discussion

Rationale for applying machine learning approaches mainly centered around it being “state of the art” or better suited to modeling complex data than regression. Machine learning was thought to be “state of the art” due to improved accuracy and deeper insights. Discussions of complex modeling focused on capturing non-linear relationships, interactions, and high-dimensionality.

When authors discussed model limitations, frequent concerns were an inadequate sample size, too few features, questionable generalizability, and a lack of interpretability. Aspects of the data other than sample size and feature number, such as potential measurement error or selection bias, were infrequently mentioned.

When discussing model implementation, many articles stated that predictive accuracy would be improved; but they did not frequently discuss how this could be translated to specific health-related policies or actions. Additionally, they rarely mentioned organizations and knowledge users that would be best suited to leverage the model.