

REVIEW ARTICLE



A Review on the Methods and Clinical Hurdles for the Use of Artificial Intelligence in Forecasting Cardiovascular Risk

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Abstract: Cardiovascular diseases are the leading contributors to global mortality, calling for more sensitive, patient-specific risk evaluation to guide early interventions. While traditional risk calculators rely on a small set of linear clinical metrics and generate static, point-in-time assessments, modern computational models offer a paradigm shift. This review discusses how machine learning and deep learning algorithms process high-dimensional clinical data, raw electrocardiographic waveforms, continuous physiological streams from wearable devices, multi-modal diagnostic imaging, and comprehensive genomic assays to provide precise and individualized risk assessment. Deep neural networks automatically extract subclinical risk markers and discover novel phenotypes, frequently outperforming conventional calculators. However, the translation of these algorithms into clinical environments is hindered by persistent methodological limitations, including inadequate external validation, poor model calibration, inconsistent reporting, data heterogeneity, and a lack of transparency in "black-box" models. Ethical concerns regarding algorithmic bias and clinical utility must be addressed to ensure safe clinical deployment. Resolving these technical barriers requires establishing standardized reporting, prospective multi-center validation, and transparent interpretability frameworks. Shifting from traditional calculators to dynamic, continuously updated risk estimation platforms utilizing privacy-preserving collaborative frameworks such as federated learning represents the future of cardiovascular risk assessment.

Keywords: Artificial Intelligence; Cardiovascular Disease; Machine Learning; Clinical Translation; Precision Medicine.

1. Introduction

The prevention and management of cardiovascular diseases require patient-specific risk estimation to guide therapeutic interventions. For decades, clinical practice has relied on risk calculators derived from large cohort studies, such as the Framingham Risk Score and the Pooled Cohort Equations [1]. While these tools have provided a foundational framework for primary prevention, they are constrained by significant methodological limitations. Traditional calculators rely on a small, predetermined set of classical risk factors, including age, biological sex, systolic blood pressure, smoking status, and lipid profiles [1, 2]. This narrow focus fails to capture the intricate, multi-layered physiological processes that contribute to atherosclerosis and myocardial dysfunction. Standard equations operate on linear assumptions that cannot model the complex, non-linear interactions among cardiovascular variables [1].

A critical drawback of legacy risk calculators is their static nature. These tools generate a singular, point-in-time risk estimate that is updated infrequently, typically every five to ten years [2, 3]. This approach ignores the dynamic evolution of cardiovascular health, which changes continuously in response to lifestyle modifications, therapeutic interventions, aging, and the onset of comorbidities [2]. Additionally, traditional models are prone to geographic and demographic miscalibration. Because they were developed using predominantly historical cohorts with limited racial and socioeconomic diversity, they frequently over-estimate or under-estimate risk when applied to contemporary, ethnically diverse populations [1, 3].

The rapid digitization of healthcare has resulted in an abundance of high-dimensional clinical data, creating new pathways for cardiovascular risk assessment. Advanced computational techniques can process vast datasets that extend far beyond traditional risk markers [4, 5]. Modern clinical repositories contain complex variables, including high-resolution electrocardiographic waveforms, continuous physiological streams from wearable devices, multi-modal diagnostic imaging, and comprehensive genomic assays [4]. Traditional statistical approaches, such as Cox proportional hazards models, are poorly suited for extracting meaningful insights from such massive, unstructured, and highly collinear data streams.

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Computational algorithms, particularly machine learning and deep learning, are uniquely capable of analyzing high-dimensional datasets. These systems can identify subtle, non-linear patterns and multi-variable interactions without requiring rigid a priori assumptions [5]. Deep neural networks can discover novel cardiovascular phenotypes and latent risk indicators that are invisible to human experts by utilizing unsupervised representation learning. Consequently, data-driven modeling offers an opportunity to transition from generic, population-averaged risk estimations to continuous, highly personalized cardiovascular forecasting.

Despite the growing number of high-performing algorithms developed in academic settings, their integration into routine clinical practice remains highly limited. This translational gap is driven by several unresolved technical and methodological challenges. A primary limitation is the retrospective and single-center design of most predictive models [6]. Algorithms often achieve exceptional performance on internal validation datasets but experience substantial performance degradation, known as algorithmic drift, when tested on external, independent cohorts [6, 7]. This lack of generalizability arises from overfitting to institutional clinical practices, specific imaging protocols, and local patient demographics.

A lack of methodological transparency severely limits clinician trust. Many state-of-the-art predictive frameworks, especially deep neural networks, operate as "black boxes," providing highly accurate predictions without clear clinical explanations for their outputs [8]. This lack of explainability makes it difficult for healthcare providers to verify the clinical safety of an algorithmic decision, identify potential confounding variables, or explain the diagnostic reasoning to patients [8, 9]. Additionally, inconsistent reporting practices regarding data preprocessing, handling of missing values, hyperparameter optimization, and validation strategies prevent the replication of published models, slowing down scientific progress in the field [6].

This review provides a discussion of the current state of artificial intelligence in cardiovascular risk prediction, focusing on the transition from technical development to real-world clinical application. The objective is to analyze the computational methodologies applied to diverse clinical modalities, evaluate the clinical challenges of these technologies, and identify future research directions.

2. Artificial Intelligence in Cardiovascular Prediction

2.1. Machine Learning on Structured Clinical Data

2.1.1. Supervised Classifiers and Ensemble Architectures

Supervised machine learning algorithms have been widely used to predict cardiovascular outcomes using structured data from electronic health records, such as laboratory measurements, demographic details, medication histories, and diagnostic codes [8]. Traditional classifiers, including regularized logistic regression, support vector machines, and Naive Bayes, serve as essential baselines for risk modeling. However, ensemble learning methods, specifically tree-based architectures such as Random Forests, Gradient Boosting Machines, Extreme Gradient Boosting (XGBoost), and LightGBM, routinely outperform single-classifier approaches in clinical prediction tasks [8, 9].

These ensemble models work by combining multiple weak decision trees to construct a highly robust predictive consensus. Tree-based systems are particularly effective at managing tabular clinical data, which often contain non-linear interactions, highly correlated variables, and missing data points [9]. For instance, in predicting heart failure hospitalization or major adverse cardiovascular events (MACE), XGBoost models can capture complex relationships between subtle declines in renal function, variations in blood pressure, and medication adherence patterns that traditional linear models overlook.

2.1.2. Model Calibration, Reclassification, and Decision Metrics

Evaluating model performance in clinical settings requires looking beyond standard discrimination metrics like the Area Under the Receiver Operating Characteristic curve (AUROC) [10]. While AUROC measures a model's ability to distinguish between patients who will experience an event and those who will not, it does not evaluate the clinical utility of the predictions. Therefore, models must also be assessed using calibration curves, which evaluate how closely the predicted probabilities align with actual observed event rates across the risk spectrum [10, 11]. A model with high discrimination but poor calibration can lead to clinical errors by over-estimating risk in low-risk individuals or under-estimating risk in high-risk patients.

To quantify the clinical value added by a new algorithmic model over established risk calculators, researchers utilize Net Reclassification Improvement (NRI) and Integrated Discrimination Improvement (IDI) [10]. The NRI metric calculates whether the new model correctly reclassifies patients into higher or lower risk categories, allowing for more appropriate therapeutic interventions, such as initiating statin therapy or scheduling advanced diagnostic tests. Decision curve analysis is also employed to evaluate the net clinical benefit of algorithmic interventions across varying decision thresholds, ensuring that model adoption improves patient outcomes without introducing excessive diagnostic burden.

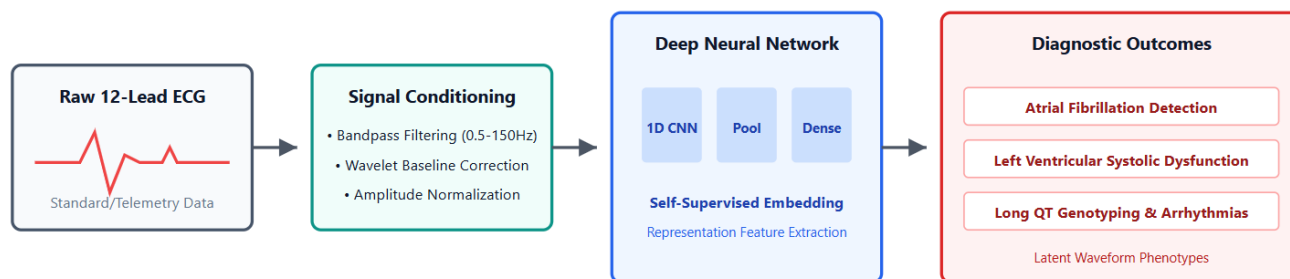


Figure 1. Operational workflow of deep learning-enabled electrocardiography.

Raw waveforms obtained from standard 12-lead arrays or mobile point-of-care devices undergo automated high-pass/low-pass filtering to remove physical noise. They are subsequently processed by convolutional neural architectures (using one-dimensional convolutions and self-supervised weights) to segment temporal features, yielding high-accuracy predictions of clinical conditions that may remain occult or subclinical on standard visual evaluation.

2.2. Deep Learning Architectures in Electrocardiography

2.2.1. Waveform Analysis and Convolutional Networks

Electrocardiography (ECG) is a fundamental tool in cardiovascular medicine, but manual interpretation is limited by human visual perception and established diagnostic criteria. Deep learning models, particularly Convolutional Neural Networks (CNNs), have transformed ECG analysis by processing raw, digitized voltage-time waveforms directly [11, 12]. These networks utilize layers of mathematical convolutions to automatically extract hierarchical features from the ECG signal, capturing subtle morphological variations and temporal dependencies that escape human detection.

CNNs can detect subclinical structural heart diseases, such as left ventricular systolic dysfunction, even during normal sinus rhythm by analyzing raw 12-lead ECG recordings [11, 14]. The algorithms identify minute alterations in depolarization and repolarization patterns, reflecting underlying myocardial remodeling. Additionally, specialized deep learning architectures, such as Temporal Convolutional Networks (TCNs) combined with Graph Neural Networks (GNNs), are used to analyze dynamic temporal variations across multiple leads, enabling highly accurate classification of complex arrhythmias [13, 15].

2.2.2. Self-Supervised Representation Learning and Foundation Models

A major challenge in training deep learning models for ECG analysis is the scarcity of high-quality, expert-labeled datasets. Manual labeling of millions of electrocardiograms is highly time-consuming and prone to inter-observer variability. To address this limitation, self-supervised learning (SSL) has emerged as a powerful paradigm [14]. Under this approach, neural networks are pre-trained on massive datasets of unlabeled ECG waveforms by solving pretext tasks, such as predicting masked signal segments or identifying contrastive representations of the same patient's cardiac cycles.

This pre-training phase allows the network to learn a rich, generalized representation of cardiac electrophysiology. These pre-trained systems, or foundation models, can then be fine-tuned on small, annotated datasets for specific clinical tasks, such as predicting long-term atrial fibrillation risk or identifying silent myocardial ischemia [14]. Foundation models developed through SSL demonstrate exceptional generalizability, maintaining high predictive performance across diverse clinical settings, patient demographics, and recording devices.

2.3. Deep Learning Applications in Cardiovascular Imaging

2.3.1. Echocardiography and Magnetic Resonance Imaging Automation

Cardiovascular imaging generates massive volumes of high-resolution, multi-dimensional data, making it highly suitable for deep learning applications. In echocardiography and Cardiac Magnetic Resonance Imaging (CMR), deep learning algorithms automate repetitive, time-consuming quantitative tasks [11]. Fully automated CNN pipelines segment cardiac chambers, delineate myocardial borders, and calculate clinical parameters such as left ventricular ejection fraction (LVEF), end-diastolic volume, and global longitudinal strain [11].

Deep learning reduces inter-institutional variability and improves diagnostic efficiency by eliminating manual tracing. Beyond automation, deep learning models analyze subtle textural features and deformation patterns in imaging studies to identify subclinical pathologies, such as early-stage hypertrophic cardiomyopathy or cardiac amyloidosis, before severe clinical symptoms manifest [11,

16]. In CMR analysis, deep learning systems evaluate late gadolinium enhancement patterns to precisely quantify myocardial scar burden, providing essential prognostic information for patients at risk of sudden cardiac death.

2.3.2. Multimodal Data Fusion and Latent Representation

While single-modality models achieve high accuracy, cardiovascular diseases are multi-faceted, and relying on a single data source can limit prognostic performance. Multimodal deep learning architectures address this by fusing imaging datasets with tabular clinical records, ECG waveforms, and genomic markers [16]. This integration can occur at different stages of the modeling process. Early fusion combines raw data features into a single input vector, while late fusion trains separate neural networks for each modality and aggregates their final outputs to make a prediction.

Intermediate fusion, where deep representations from different modalities are projected into a shared latent space, has proven highly effective [16]. For instance, multimodal architectures that integrate myocardial strain imaging from echocardiograms with longitudinal electronic health records and baseline biomarkers show superior performance in predicting heart failure exacerbations compared to single-modality models. This multi-modal approach ensures that predictions are based on a holistic view of the patient's cardiovascular profile.

Table 1. Characterization of Multimodal Data Inputs and Preprocessing

Cardiovascular Data Modality	Typical Raw Formats	Common Artifacts & Noise	Preprocessing & Feature Extraction Pipelines	Common Fusion Strategies	Computational & Storage Demands
Digitized 12-Lead ECG	XML, DICOM-wave, CSV (typically sampled at 250 Hz to 1000 Hz)	Baseline wander, powerline interference (50/60 Hz), electromyographic noise	Bandpass filtering (0.5 Hz - 150 Hz), wavelet denoising, pan-tompkins QRS detection	Intermediate fusion via 1D convolutional latent embeddings	Moderate (requires specialized signal processing libraries)
Cardiac Imaging (CT / MRI)	DICOM slices, NIfTI volumes	Motion artifacts, respiratory blur, metallic hardware streak artifacts	Hounsfield Unit windowing, bias-field correction, spatial resampling, automated U-Net segmentation	Late fusion of convolutional feature maps or tabular extraction	High (requires high-RAM GPUs and extensive disk arrays)
Structured EHR & Clinical Tables	SQL databases, HL7/FHIR JSON resources	Missing values, inconsistent ICD coding, temporal irregularity	MICE (Multivariate Imputation by Chained Equations), Z-score normalization, one-hot encoding	Early fusion via concatenation; late fusion of classifier logits	Low (readily processed on standard CPU instances)
Wearable PPG & Telemetry	Raw photoplethysmography infrared channels, accelerometer streams	Motion artifacts, ambient light leakage, skin-contact fluctuations	Bandpass filtering (0.5 Hz - 4 Hz), derivative-based peak detection, signal-quality index (SQI) gating	Early fusion of temporal features or late fusion of alarm alerts	Low to Moderate (frequently optimized for edge processors)
Genomic & Transcriptomic Assays	PLINK files, VCF (Variant Call Format), FASTQ	Sequencing errors, batch effects, population stratification	Quality control filtering (minor allele frequency > 0.01, Hardy-Weinberg equilibrium $p > 10^{-6}$), PCA for ancestry	Early fusion of polygenic risk scores (PRS) as clinical features	High (primarily in the initial alignment and variant-calling phases)

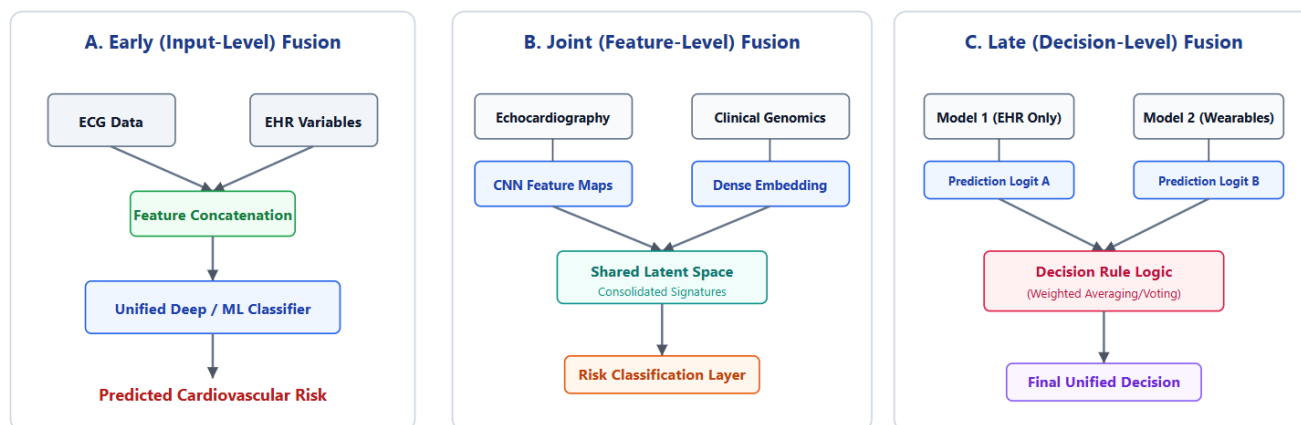


Figure 2. Classification of data integration methods in multimodal systems.

2.4. Continuous Monitoring and Wearable Diagnostics

2.4.1. Real-Time Signal Processing and Edge Computing

The widespread adoption of consumer smartwatches, patch monitors, and smart rings has enabled continuous, ambulatory monitoring of physiological signals, including photoplethysmography (PPG) and single-lead ECG [11, 17]. Processing these continuous, high-volume data streams requires efficient computational frameworks. Traditional approaches that transmit raw physiological signals to centralized cloud servers face challenges related to network bandwidth, high latency, and patient privacy concerns.

To overcome these issues, edge computing paradigm deploys optimized, lightweight machine learning models directly onto wearable hardware. These edge algorithms perform real-time signal preprocessing, filter out motion artifacts, and run continuous anomaly detection [17]. Edge-enabled wearables can instantly notify patients and healthcare providers of potentially life-threatening events by processing data locally, such as acute myocardial ischemia or malignant arrhythmias, while minimizing data transmission and battery consumption.

2.4.2. Diagnostic Performance and Quality-Assessment Protocols

Ambulatory monitoring devices play a critical role in detecting paroxysmal arrhythmias, such as silent atrial fibrillation, which are often missed during brief clinical examinations [17]. Large-scale clinical trials have demonstrated that wearable devices, when paired with deep learning detection models, can successfully identify atrial fibrillation in real-world cohorts. However, translating these systems to routine clinical practice presents challenges.

Physiological signals collected during daily activities are often degraded by motion artifacts, poor sensor contact, and environmental noise, leading to false-positive detections. These false positives can cause patient anxiety and place an unnecessary burden on healthcare systems through excess clinic visits. Therefore, wearable-based AI systems must utilize robust quality-assessment algorithms that automatically discard low-quality data segments and output confidence scores alongside clinical predictions to guide follow-up care [17].

2.5. Analysis of Representative Literature

To provide a consolidated overview of the current research landscape and resolve redundancies identified in earlier literature evaluations, the table below synthesizes the methodological approaches, clinical outcomes, and limitations of representative cardiovascular artificial intelligence studies.

3. Translational Perspectives

3.1. Clinical Implications and Challenges

The transition of artificial intelligence models from computational benchmarks to bedside cardiology is complicated by fundamental differences in performance evaluation. Many published predictive algorithms demonstrate high statistical discrimination, typically

reported as a high area under the receiver operating characteristic curve (AUROC) [11]. However, a model with strong statistical discrimination may offer minimal clinical utility if it does not change therapeutic pathways or improve patient outcomes [11, 25]. The central clinical question is whether a fractional increase in predictive accuracy justifies the operational cost of model deployment, the disruption of clinical workflows, and the potential risk of false positives.

Table 2. Methods, Findings, and Translational Gaps of Cardiovascular AI Studies

Study	AI Approach & Data Type	Main Objective	Findings	Limitations & Gaps	Future Directions
Andritsch & Dike (2026) [18]	Explainable AI (XAI), ML/DL models using SHAP and LIME on structured clinical datasets.	Evaluate explainability in cardiovascular AI prediction models.	Explainable models improved interpretability while maintaining comparable prediction performance.	Limited external validation and lack of interpretability standardization across deep learning architectures.	Development of standardized, universally applicable, and clinically interpretable XAI frameworks.
Kavila et al. (2022) [19]	Artificial Neural Network (ANN) and model-agnostic XAI on clinical datasets.	Assess model-agnostic XAI for cardiovascular prediction.	An ANN multi-level model achieved 87% prediction accuracy and improved classification performance.	Preprocessing methodologies, validation strategies, and reproducibility metrics were insufficiently reported.	Integration of heterogeneous multimodal cardiovascular data within interpretable architectures.
Alshraideh et al. (2025) [20]	Hybrid CNN and generative language model framework on clinical and multimodal datasets.	Develop a hybrid AI framework for early heart failure prediction.	Hybrid framework achieved high predictive performance with strong Area Under the Curve and F1-scores.	Lacked prospective clinical validation and performance evaluation in multicenter environments.	Expansion of hybrid frameworks into real-time clinical decision-support systems.
Joshi et al. (2023) [21]	AI-assisted cardiac computed tomography (CT) imaging analysis.	Assess clinical applications of AI in cardiac CT imaging.	AI automated coronary artery plaque quantification, calcium scoring, and long-term event prediction.	Limited standardization across different imaging platforms and scarcity of clinical implementation studies.	Broader clinical automation, standardization, and prospective multicenter validation trials.
Chitrapathy et al. (2025) [22]	Machine learning integrated with IoT-based monitoring on the UCI dataset and health telemetry data.	Improve cardiovascular prediction using AI and internet-of-things (IoT) platforms.	High classification accuracy was reported for ambulatory cardiovascular prediction.	Significant concerns regarding training dataset diversity, potential biases, and overall generalizability.	Deployment and validation of models using large-scale, diverse, and wearable-derived cohorts.
Umar et al. (2023) [23]	Hybrid ANN-Genetic Algorithm (ANN-GA) model on structured cardiovascular datasets.	Compare hybrid machine learning approaches for cardiovascular disease prediction.	The hybrid ANN-GA approach outperformed traditional machine learning classifiers, including SVMs and decision trees.	Constrained by a small sample size, limited cohort diversity, and lack of external validation.	Extensive validation of hybrid models across geographically and demographically diverse cohorts.
Li et al. (2022) [24]	Machine learning and biomechanics-integrated modeling.	Explore the interplay between artificial intelligence and biomechanical variables.	Machine learning reduced manual computational steps and improved hemodynamic assessment efficiency.	Data integration challenges and poor model scalability across complex vascular networks.	Integration of automated computational biomechanics with advanced multimodal AI architectures.

Evaluating the clinical value of predictive tools requires measuring how effectively they risk-stratify patients to guide specific clinical choices. For instance, a model predicting coronary event risk must align with actionable decisions, such as the initiation of intensive lipid-lowering therapies or referral for invasive coronary angiography [25, 26]. Clinical efficacy must be demonstrated in diverse, resource-limited healthcare environments, where the deployment of complex computational architectures must be balanced against basic patient-care needs [11].

3.2. Validation and Reliability

While discrimination measures a model's ability to separate high-risk from low-risk patients, clinical safety is heavily dependent on model calibration. Calibration evaluates how closely the predicted risk probabilities align with observed historical event rates [10]. A poorly calibrated model may systematically over-estimate risk, leading to unnecessary diagnostic testing and patient anxiety, or under-estimate risk, resulting in missed therapeutic opportunities [11, 26]. In preventive cardiology, evaluating model performance must incorporate calibration metrics, including Hosmer-Lemeshow goodness-of-fit statistics, calibration slopes, intercepts, and Brier scores [10].

In addition to calibration, metrics such as Net Reclassification Improvement (NRI) and Integrated Discrimination Improvement (IDI) are essential for quantifying the incremental clinical benefit of a novel model over existing standard risk calculators [10]. Evaluating generalizability is another challenge. Most cardiovascular predictive models are developed and validated using retrospective data from single institutions [6]. When these models are applied to external cohorts, they frequently experience a substantial drop in performance [7]. This performance degradation, or algorithmic drift, occurs due to variations in local patient demographics, clinical practice patterns, coding practices, and imaging hardware [6, 7]. To ensure reliability, models must undergo prospective external validation across diverse geographical, socioeconomic, and ethnic groups [11, 25].

3.3. Bias and Fairness Issues

Artificial intelligence models are highly susceptible to reproducing and magnifying systemic biases present in their training datasets [1, 26]. Historical clinical data often reflect disparities in healthcare access, treatment patterns, and diagnostic strategies across different demographic groups. For example, traditional cardiovascular cohort studies, such as the original Framingham cohorts, relied on predominantly white, middle-class populations, leading to miscalibrated risk estimation when applied to contemporary, multi-ethnic patient groups [1].

If an algorithm is trained on data with socioeconomic or racial disparities, it may perpetuate these inequities. Bias can manifest in several ways, including lower predictive performance in marginalized groups or systematic underestimation of cardiovascular risk in specific demographic cohorts [26]. Addressing these issues requires systematic fairness assessments during model development, utilizing quantitative fairness metrics and algorithmic debiasing techniques. Integrating diverse, multi-center datasets is essential to ensure that predictive tools are both technically accurate and clinically equitable [1, 25].

Table 3. Clinical Implications and Diagnostic Relevance of AI Modalities in Cardiovascular Care

Clinical Domain	AI Application	Translational Relevance	Clinical Impact
Electrocardiography	Automated arrhythmia classification and subclinical waveform feature extraction.	Identifies latent cardiovascular phenotypes and subtle structural anomalies.	Facilitates early diagnostic detection and improves risk stratification.
Heart Failure Prediction	Hybrid machine learning and deep learning risk estimation systems.	Continuous longitudinal monitoring and clinical deterioration modeling.	Aids in reducing hospital admission rates and mitigating mortality risk.
Cardiac Imaging	Automated computer-assisted CT, MRI, and echocardiography workflows.	Standardizes plaque characterization and ventricular chamber segmentation.	Optimizes clinical throughput and enhances diagnostic reproducibility.
Wearable Monitoring	Streamlined photoplethysmography and single-lead ECG analysis.	Ambulatory physiological tracking and remote event surveillance.	Supports real-time intervention and outpatient monitoring.
Explainable AI	Feature-importance mapping utilizing SHAP and LIME architectures.	Explains algorithmic decisions, improving clinician transparency.	Elevates trust among healthcare providers and supports decision-making.
Multimodal AI	Data fusion integrating imaging, ECG, genomic, and laboratory variables.	Provides a unified, multidimensional view of patient health.	Enables highly personalized, precision-guided cardiovascular care.

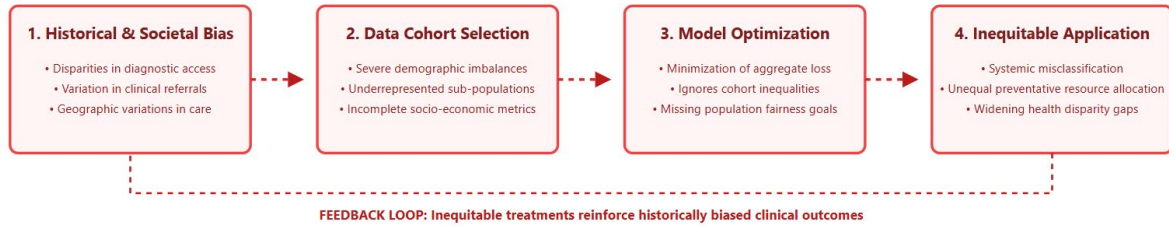


Figure 3. Flow of algorithmic bias propagation across the cardiology translational lifecycle

3.4. Clinical Adoption Barriers

The integration of predictive algorithms into routine clinical workflows is a major challenge for translation [11]. Healthcare professionals already manage high cognitive loads and complex electronic health record (EHR) systems. Introducing independent computational tools that require manual data entry, generate excessive alerts, or operate outside existing software frameworks can lead to user frustration and "alert fatigue" [27]. For successful adoption, predictive systems must be integrated directly into local EHR environments, delivering automated, real-time risk assessments at the point of care [11, 27].

Table 4. Algorithmic Fairness and Bias Mitigation in Predictive Cardiology

Metric / Mitigation Class	Specific Strategy / Formalism	Mathematical Definition / Algorithmic Mechanism	Clinical Application & Implications in Cardiology
Group Fairness Metric	Demographic Parity (Statistical Parity)	$P(\hat{Y} = 1 A = a) = P(\hat{Y} = 1 A = b)$ <p>(where \hat{Y} is the model prediction, and A represents a protected attribute like biological sex or race)</p>	Ensures that the rate of high-risk stratification (e.g., referral for coronary angiography) is equal across all demographic groups, independent of baseline prevalence.
Group Fairness Metric	Equal Opportunity	$P(\hat{Y} = 1 A = a, Y = 1) = P(\hat{Y} = 1 A = b, Y = 1)$ <p>(where $Y=1$ is the true clinical event, such as myocardial infarction)</p>	Ensures that the true positive rate (sensitivity) is equal across demographic groups, preventing underdiagnosis or missed preventive therapies in marginalized patient populations.
Group Fairness Metric	Predictive Equality	$P(\hat{Y} = 1 A = a, Y = 0) = P(\hat{Y} = 1 A = b, Y = 0)$	Standardizes the false positive rate across groups, mitigating the risk of disproportionate anxiety and unnecessary diagnostic testing (e.g., cardiac catheterization) in lower-risk cohorts.
Bias Mitigation	Pre-processing (e.g., Reweighting)	$W(x) = \frac{P(A=a) \cdot P(Y=y)}{P(A=a, Y=y)}$ <p>(modifying sample weights $W(x)$ before training)</p>	Adjusts historical datasets to correct for past treatment disparities, ensuring that minority populations who received suboptimal care do not bias the baseline model.
Bias Mitigation	In-processing (e.g., Adversarial Debiasing)	$\min_{\theta_G} \max_{\theta_D} [\mathcal{L}_{\text{predictor}}(\theta_G) - \lambda \mathcal{L}_{\text{adversary}}(\theta_G, \theta_D)]$ <p>(where the adversary attempts to predict the protected attribute from the latent features)</p>	Trains predictive networks in a game-theoretic framework to ensure that deep feature extractors cannot store or utilize demographic proxies when estimating heart failure mortality.
Bias Mitigation	Post-processing (e.g., Reject Option Classification)	Adjust decision thresholds τ_a and τ_b dynamically for different demographic groups close to the decision boundary.	Minimizes disparities in clinical decision-making post-hoc without requiring model retraining, particularly useful in locked, commercially deployed diagnostic systems.

Another barrier is the opaque, "black-box" nature of advanced deep learning architectures [11]. Clinicians are hesitant to trust automated recommendations that lack clear explanatory reasoning, especially when those recommendations contradict clinical intuition or standard guidelines. Explainable AI (XAI) methodologies, including SHapley Additive exPlanations (SHAP) and Local Interpretable Model-agnostic Explanations (LIME), help address this issue by identifying the specific clinical features that drive a model's prediction [18]. However, further work is required to standardize interpretability frameworks and ensure that algorithmic explanations are clinically meaningful, reproducible, and easy to understand [11, 18].

4. Future Trends

4.1. Temporal Trajectory Modeling

The future of cardiovascular prediction lies in transitioning from static, point-in-time risk estimates to dynamic, continuous risk trajectories [8, 28]. Traditional models generate static risk assessments that are infrequently updated, ignoring the progressive nature of cardiovascular disease. Modern computational systems can process continuous streams of physiological data from wearables and home monitoring devices alongside longitudinal health records to build adaptive, real-time risk profiles [8, 29].

These dynamic models can generate individualized survival curves, illustrating how a patient's risk profile evolves in response to lifestyle modifications, therapeutic changes, and clinical events [28]. This shift enables proactive, preventive interventions, allowing clinicians to adjust therapies before acute clinical events occur. For example, continuous monitoring systems can track subtle changes in heart rate variability, arterial stiffness, and physical activity to predict impending heart failure exacerbations, enabling early medication adjustments and reducing hospitalization rates [8, 28, 29].

4.2. Reporting Standards and Lifecycle Governance

Realizing the potential of dynamic cardiovascular prediction requires addressing challenges in reporting and validation [29, 30]. The lack of standardization in data preprocessing, validation methods, and performance reporting makes it difficult to compare and replicate models. The scientific community must adopt standardized reporting frameworks, such as TRIPOD-AI and DECIDE-AI, to improve transparency and reproducibility [30].

Table 5. Comparison of Reporting Guidelines and Lifecycle Governance

Guideline Standard /	Developmental Phase	Stakeholders	Core Reporting	Translational Benefit in Cardiovascular AI
TRIPOD-AI (Transparent Reporting of a multivariable prediction model of Individual Prognosis Or Diagnosis-AI)	Model Development & Validation	Machine learning developers, clinical statisticians, journal editors	Explicit description of AI architecture, hyperparameter tuning protocols, handling of missing data, and external validation cohort selection.	Mitigates reporting inconsistencies and poor reproducibility in retrospective ML studies based on structured electronic health records.
DECIDE-AI (Development and Assessment of Robotic and Artificial Intelligence Devices in Clinical Trials)	Early-Stage Clinical Evaluation	Clinical investigators, regulatory officers, system implementers	Documentation of human-AI interaction, pilot safety profiles, user-centered design feedback, and institutional workflow adaptation.	Guides the initial translational transition of deep learning tools from silent computational profiling to live bedside decision support.
MINIMAR (Minimum Information for Medical AI Reporting)	Raw Data and Cohort Curation	Database administrators, data engineers, epidemiologists	Strict cataloging of demographic representation, clinical characteristics, data source origins, and potential selection biases in the training pool.	Exposes latent cohort biases in large historical cardiovascular databases before model optimization begins.
FDA SaMD Action Plan (Software as a Medical Device)	Post-Market Surveillance & Deployment	Medical device manufacturers, federal regulators, healthcare administrators	Implementation of a Predetermined Change Control Plan (PCCP), real-world performance monitoring, and structured algorithm update protocols.	Establishes a safe, legal framework for managing algorithmic drift and continuous learning without violating active diagnostic approvals

Clinical deployment requires robust regulatory pathways that treat software as a medical device (SaMD). Regulated clinical systems must feature structured post-market surveillance and continuous auditing to detect and mitigate algorithmic drift over time [29, 30]. Establishing these governance standards is essential to maintain model safety, clinical efficacy, and user trust throughout the software lifecycle.

4.3. Federated Learning and Data Privacy

Developing generalizable predictive models requires training on large, diverse, multi-institutional datasets [30]. However, consolidating sensitive patient data into centralized repositories is challenging due to patient privacy regulations, data security concerns, and institutional data-sharing restrictions. Federated learning (FL) offers a solution by enabling multi-center collaboration without sharing raw patient data [31].

Under a federated framework, institutions train local models on their own clinical data and share only the resulting mathematical updates with a centralized coordinator. The coordinator aggregates these updates to refine a global model, which is then shared back with the participating centers [31]. This process is repeated iteratively. Federated learning preserves patient privacy, complies with strict data protection laws, and helps reduce algorithmic bias by enabling models to learn from highly diverse, multi-national cohorts [30, 31, 32].

4.4. Limitations

Several key limitations affect the current state of computational prediction in cardiovascular medicine [1]. As a narrative appraisal, the present literature survey is subject to potential selection bias, and the absence of primary meta-analytical pooling limits the generalizability of some findings. The vast majority of published predictive models are based on retrospective, single-center datasets [6]. These models frequently overfit to local practices and imaging parameters, leading to poor generalizability when applied to external populations.

Electronic health record data are often incomplete, noisy, and inconsistently coded across healthcare networks, which introduces noise and compromises training quality [1]. The opaque nature of complex deep learning models limits clinical interpretability and clinician trust [8]. The clinical deployment of these advanced technologies requires significant digital infrastructure, technical expertise, and financial resources, which are often unavailable in low-resource settings, potentially widening existing disparities in healthcare delivery [1].

5. Conclusion

Artificial intelligence is a major advance in cardiovascular medicine, offering tools to transition from legacy, population-averaged risk estimators to precise, individualized risk assessment. These methods can capture complex, non-linear relationships and identify subclinical risk patterns that are invisible to human clinicians by utilizing machine learning algorithms on tabular clinical records, deep learning architectures on physiological waveforms, and automated quantitative pipelines in cardiovascular imaging. However, converting these computational models into safe, effective clinical tools requires overcoming persistent challenges in model calibration, external validation, workflow integration, and algorithmic fairness. Resolving these hurdles requires adopting standardized reporting frameworks, implementing rigorous external validation protocols, developing transparent explainability tools, and utilizing privacy-preserving collaborative learning models. Overcoming these methodological and translational barriers is essential to establish artificial intelligence as a cornerstone of precision cardiovascular care.

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