

# It doesn't matter what they say in the papers... It's still ROC and roll to me

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*Comment on:* Chong SL, Niu C, Piragasam R, *et al.* Adding heart rate n-variability (HRnV) to clinical assessment potentially improves prediction of serious bacterial infections in young febrile infants at the emergency department: a prospective observational study. Ann Transl Med 2023;11:6.

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Heart rate variability (HRV) is a quantitative method for calculating the variance in the length of time between successive heartbeats. Although healthy humans at rest are generally considered to have steady heart rates, there are minute fluctuations in the length of time between successive R waves, on the order of milliseconds. For more than half a century, a loss of HRV has been associated with autonomic dysfunction in the setting of illness. Perhaps the earliest description of HRV was by Hon and Lee in 1963 (1) with the observation that loss of HRV was associated with fetal distress and death, which formed the foundation for modern interpretative standards of cardiotocography (CTG) during labor.

In the decades that followed this observation, a wide range of HRV measures have been developed, evolving with advancements in computer technology to allow faster, more sophisticated calculations on increasingly large datasets. Alterations in HRV have been associated with increased risk of death after myocardial infarction (2), diabetes (3), depression (4), sepsis in adults (5) and neonates (6), and outcomes in neonatal hypoxic-ischemic encephalopathy (7). While many research tools for calculating HRV measures exist and the association of depressed HRV with illness is unequivocal, HRV monitoring is not a standard of care in the intensive care unit (ICU) or non-ICU setting.

Early warning of impending patient deterioration is an important and universal goal in health care. In contrast to doctors of yesteryear who had only a keen eye and a "sixth sense" at their disposal, the modern healthcare provider has vast sums of organized, digitized data at their fingertips. Big data predictive analytics hold great potential to improve patient health (8). However, this bounty has created a new problem-identifying salient and robustly predictive factors across heterogeneous and nuanced patient populations. A number of different systems have been developed to assess illness severity and predict morbidity or mortality. However, despite the initial promise of many of these systems, significant gaps have led to uneven adoption and failure to improve outcomes.

The Acute Physiology and Chronic Health Evaluation (APACHE) score was first proposed in 1985 by Knaus *et al.* (9) and encompasses lab values, limited vital signs, and acute and chronic medical conditions to provide a quantitative measure of illness severity and a prediction of mortality in the first 24 hours after admission to the ICU. Although currently in the fourth version (APACHE-IV), the

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APACHE-II score remains widely used due to ease of use and availability of online calculators. While performance is high for the first 24 hours, the APACHE score has been criticized for its static nature, failure to account for subsequent treatment or the hospital course after the first 24 hours, and interrater reliability concerns about certain variables (10,11).

The Sequential Organ Failure Assessment (SOFA) is a similar illness severity score designed with the explicit intent of quantifying the degree of organ failure in critical illness (such as sepsis) across cardiovascular, respiratory, neurologic, renal, hematologic, and hepatic domains (12). Like APACHE-II, SOFA is widely utilized in clinical practice. Pediatric (pSOFA) and neonatal (nSOFA) variants were developed more recently (13,14). Unlike APACHE-II, SOFA can be recalculated at regular intervals, accounting for longitudinal changes in the patient's course and treatment. The latest revision of consensus definitions of sepsis in adult patients uses SOFA as key criteria (15), yet significant concerns about SOFA's reliability and predictive performance have been raised (16,17).

Despite their utility for benchmarking ICU performance, research, or characterizing a septic patient, many argue that severity of illness scores like APACHE and SOFA should not be used for individual patient decision-making (18,19). To date, the literature lacks evidence demonstrating that clinical usage of SOFA leads to any change in patient outcome, despite widespread adoption.

Complimentary to early warning scores, has been the development and deployment of so-called "Rapid Response" teams (RRT), predesignated medical teams which are available to respond at short notice to medical wards when a decompensating patient is recognized. The RRT concept is based on several observations: (I) most inhospital cardiac arrest occurs outside the ICU, (II) signs of decompensation are present 6–8 hours before arrest, and (III) similar dedicated teams responding to traumas led to improved morbidity and mortality outcomes. Although the implementation of RRT has been widespread, owing to recommendations from accreditation agencies, the quality of evidence supporting this practice remains low and inconsistent, as shown in more than one meta-analysis covering hundreds of studies (20,21).

While the application of Big Data analysis techniques may be technically possible and yield statistically impressive results, if it is not done with an understanding of the framework in which medicine is practiced, it is likely to struggle with real-world deployment, face significant adoption hurdles, and ultimately fail to positively alter patient outcomes.

Quantitative early warning algorithms should seek to make good on the promises of outcome calculators and rapid response teams by providing accurate, timely warning of an impending event where mitigation is likely to prevent damage or adverse outcomes. The challenge lies in integrating predictive analytics with clinician decisions (22). In doing so, such technology should reduce mortality and morbidity, but only if it performs as intended in the population to which it is applied. Failure to critically evaluate a predictive algorithm has resulted in many reportedly accurate algorithms falling short in the clinical setting. The TRIPOD (Transparent Reporting of a multivariable prediction model for Individual Prognosis or Diagnosis) guidelines were developed as a standardized reporting format to facilitate interpretation of results across studies (23). Here, we review six key principles to consider when evaluating and reporting the results of predictive analytics.

(I) Multiple metrics of accuracy should be evaluated and reported (24); for models developed using existing datasets where the outcome is known, the Area Under the Curve of the Receiver Operating Characteristic (Abbreviated AUC, AUROC, or ROC) is the most commonly used and an easily compared metric for performance evaluation. But AUC has drawbacks. First, it is calculated against every possible threshold of model output, and the tradeoff between sensitivity and specificity is calculated at each of these thresholds and summed. Yet, in clinical practice, the output of the model will result in distinct action thresholds (often only one action threshold, in fact). So, while AUC gives the modeler a wide picture of overall model performance, it does not measure the performance at the action threshold(s). Second, because AUC is derived from sensitivity and specificity, it is a backward-looking metric. Sensitivity and specificity start with a patient's known diagnosis and then look back to determine whether the test was positive for patients with the diagnosis (sensitivity) or negative for patients without the diagnosis (specificity). However, in clinical practice, the outcome is not yet known; thus, forward-looking metrics such as predicted risk, relative risk, odds ratio, positive predictive value, and negative predictive value better match the clinical picture confronting the provider. Each of these metrics is calculated at a single threshold, which holds more clinically relevant information, but limits the range of predictive information conveyed. Calibration plots or measures provide information on how the accuracy of the predicted risk compared to the observed risk of an event or outcome. Importantly, a model with excellent discrimination (AUC >0.9) is rarely well-calibrated in the clinical setting, especially for rare outcomes.

- A second principle for consideration is the method (II)and timing of action. Algorithms and early warning systems (EWSs) do not impact patients directlythey impact patients by prompting a change in clinical actions or decisions. The evaluator must ask two questions: (i) what will the predicted risk generated by the algorithm prompt the clinician to do that will intuitively affect patient outcome positively? And (ii) what is the desired timing relative to the event that an action (i.e., intervention) needs to occur in order to achieve the improved outcome? As a simple example that ties these concepts together, if we know that each hour of delay in starting antibiotics to treat sepsis is measurable in terms of increased mortality, then if we can improve the initiation of antibiotics by 3 hours with an EWS, we should be able to decrease mortality. Here we have defined an action (start antibiotics) that addresses the clinical event, and a defined time (3 hours) prior to the event at which to evaluate model performance.
- (III) As alluded to above, a critical element of designing any EWS is implementation. A busy ICU is full of constantly alarming monitors, the vast majority of which are false positives and contribute to alarm fatigue. A new EWS should not further exacerbate this problem. Display of predictive analytics without alerts avoids this issue and allows users to interpret trends and risk in the context of the full clinical picture, but requires the clinician to seek out the information rather than having it automatically presented as an alert. Selecting an alert threshold that minimizes false positives will inherently decrease true positives, and may also alter the timing of the "early" warning to such a degree that there is no longer any action that can positively alter the outcome. Hence, the metrics of model performance must be calculated at the

defined point in time relative to the clinical event (as outlined in the second principle). Continuing the sepsis prediction example, if we believe that antibiotics are over-utilized at a ratio of 10:1 in our patient population (which implies that standard care has a 9% positive predictive value), then we might choose a threshold that results in a 15% or 20% positive predictive value for early warning of sepsis, which would represent a dramatic improvement relative to the current standard of care.

- (IV) Again, our previous principle has become a segue into the next: for any new predictive paradigm, quantitative comparison to the existing paradigm is necessary for determining if the newer, likely more complex, model has sufficiently greater performance to justify the increased computational and/or data input demands. Metrics of model performance reported for a new algorithm are contextual—an AUC of 0.70 or a PPV of 15% can benefit the patient if they represent improvements to the existing paradigm. Statistical tools such as the net reclassification index (25) quantify the degree to which performance shifts.
- (V) Orthogonality provides a more complete picture of the patient. For example, in preterm infants, birth weight and gestational age provide nearly identical information, as opposed to gestational age and sex, which capture distinctly different elements of risk. Additionally, physiologic monitoring data stand to tell clinicians more about what they don't already know than other clinical risk factors. Many predictive algorithms use information that the provider is already aware of, in order to quantitate a risk prediction. This approach can only benefit the patient to the extent that the clinician is unaware or unable to process these data on their own. In contrast, algorithms that recognize subtle patterns in vital signs before the change is recognized at the bedside can alter the timing of interventions and change the course of illness. In practice, clinicians are remarkably adept at assimilating relevant clinical information, and through years of training and practice, intuiting the status of the patient. An algorithm that regurgitates back to the clinician something that they already know might provide decision support once the condition is recognized, but will not be useful for early warning. The patient will only benefit to the extent the model

is providing information that is either new (i.e., orthogonal) or beyond the capacity of the clinician to process in absence of the model.

(VI) Costs must be quantified. The deployment of any clinical decision support system involves costs, which are often far greater than the researcher initially conceives. Data acquisition interfaces, user interface hardware and software, regulatory and quality system burdens, installation and training costs, 24/7 technical support, and post-market surveillance are some of the many costs that must be borne. Cost-effectiveness for improving outcomes (e.g., reduction in length of stay, avoidance of complications or comorbidities) can be assessed against the capital or operating costs of the system or devices, where commercial options are available, or projected costs where such a system is envisioned. Costs should also be assessed along a different dimension-when predictions are inaccurate, what is the cost (financial, pain, increased length of stay) of additional testing and treatment for false positives? What is the cost (death, increased morbidities) for false negatives?

In a manuscript by Chong et al., published of Annals of Translational Medicine (26), the authors describe a quantitative algorithm that uses conventional HRV, a novel second-order derivative of HRV (termed HRnV), demographics, vital signs, and laboratory results to identify which infants presenting to a hospital emergency ward were at risk for serious bacterial infection (SBI). In building this model, the authors utilized several of the previously outlined evaluation principles-the inclusion of orthogonal data elements and sequential comparison of more complex models using multiple accuracy metrics. The reported model shows significant promise for identifying infants at risk for SBI, permitting more targeted allocation of resources, and improving antibiotic stewardship, all of which are likely to increase the cost-effectiveness of care. We look forward to seeing future studies of this promising technology that evaluate thresholds for clinical action and external validation in a wider patient population. Given the financial and time costs of clinical trials, it is imperative that new models be rigorously evaluated and validated before deployment in a clinical setting or randomized clinical trials.

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