Deep-learning-based real-time prediction of acute kidney injury outperforms human predictive performance

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Acute kidney injury (AKI) is a major complication after cardiothoracic surgery. Early prediction of AKI could prompt preventive measures, but is challenging in the clinical routine. One important reason is that the amount of postoperative data is too massive and too high-dimensional to be effectively processed by the human operator. We therefore sought to develop a deep-learning-based algorithm that is able to predict postoperative AKI prior to the onset of symptoms and complications. Based on 96 routinely collected parameters we built a recurrent neural network (RNN) for real-time prediction of AKI after cardiothoracic surgery. From the data of 15,564 admissions we constructed a balanced training set (2224 admissions) for the development of the RNN. The model was then evaluated on an independent test set (350 admissions) and yielded an area under curve (AUC) (95% confidence interval) of 0.893 (0.862–0.924). We compared the performance of our model against that of experienced clinicians. The RNN significantly outperformed clinicians (AUC = 0.901 vs. 0.745, p < 0.001) and was overall well calibrated. This was not the case for the physicians, who systematically underestimated the risk (p < 0.001). In conclusion, the RNN was superior to physicians in the prediction of AKI after cardiothoracic surgery. It could potentially be integrated into hospitals’ electronic health records for real-time patient monitoring and may help to detect early AKI and hence modify the treatment in perioperative care.

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In summary, we retrospectively analysed EHR time series data with high temporal resolution (up to 1 min) generated at a tertiary care center for cardiovascular diseases. Based on \( n = 2224 \) admissions, we developed an RNN that continuously (every 15 min) predicted the probability of developing AKI defined as KDIGO stage 2 or 3 within the first 7 days after cardiothoracic surgery.

Supplementary Tables 1–4 show a comparison of baseline characteristics between AKI- and non-AKI cases in the training, balanced and imbalanced test set and the whole study population before matching AKI- and non-AKI cases.

Table 1 shows the performance metrics of our RNN evaluated on an independent test set with \( n = 350 \) patients. The model achieved an area under curve (AUC) (95% confidence interval (CI)) of 0.893 (0.862–0.924). In addition, we trained a model with only serum creatinine as input and yielded an AUC of 0.805 (0.768–0.842). Thus, the addition of further parameters led to an absolute increase of around 10 percentage points in the AUC. However, a model using all features but creatinine and glomerular filtration rate (GFR) (the GFR is calculated from creatinine) performed almost as good as the full model with an AUC of 0.887 (0.855–0.919)—probably due to high correlation between creatinine and other features, e.g., urea. For further performance metrics of these reduced models see Supplementary Tables 5 and 6.

A table with the model performance metrics derived from an imbalanced test set with incidence rate of 10% AKI (see Supplementary Results 1) can be found in Supplementary Table 7. In addition, we analysed some examples of the predictions of individual patients including false-positive and false-negative predictions. These can be found in Supplementary Figs. 1–3.

### RNN vs. human-level performance—experimental design

We set up an experiment to compare our ML model against experienced physicians (Fig. 1). For each of the \( n = 350 \) patients of our balanced test set a quasi-random point in time in their observation period was chosen, further denoted as ‘prediction point’ (For more information about quasi-random samples see the ‘Methods’ section.).

At the chosen prediction point, seven experienced physicians and the ML model each had to make a prediction (between 0 and 100%) of how likely the patient was to develop AKI within the first 7 days after surgery.

All time series information up to the ‘prediction point’ was graphically displayed for the physicians to mimic the electronic patient chart.

### Performance of RNN and physicians

The performance of our RNN and the physicians’ assessment can be found in Table 2 (Note that the metrics of the RNN are slightly different from those in section ‘Performance of the RNN based prediction’. The reason is that in the RNN vs. human experiment only one prediction point per patient was evaluated, whereas for the complete evaluation of the RNN all predictions of the whole observation periods for all patients were evaluated.)

The median (interquartile range (IQR)) prediction value for the physicians was 0.36 (0.15–0.70) vs. 0.51 (0.12–0.86) for the RNN.

Across all metrics, the RNN outperformed the physicians. We obtained an AUC of 0.901 for the RNN vs. 0.745 for the physicians (\( p < 0.001, Z = 6.85, \) DeLong’s test). The receiver operating characteristic (ROC) curves and the precision-recall curves are displayed in Fig. 2a and Fig. 2b, respectively.

The mean of our predictive quality score \( S (S = r, \text{if the patient developed AKI and } S = 1−r, \text{if the patient did not develop AKI}) \) was significantly higher for the RNN than for the experienced physicians (0.754 vs 0.639, \( p < 0.001, \) t-statistic = 8.47, df = 349, paired t-test).

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**Table 1. Model performance metrics for balanced test set.**

| Metric       | Threshold-independent metrics, (95% CI) | Threshold-dependent metrics, (95% CI) |
|--------------|----------------------------------------|---------------------------------------|
| AUC          | 0.903                                  | 0.893 (0.862–0.924)                   |
| PR_AUC       | 0.124                                  | 0.124 (0.094–0.159)                   |
| Acc          | 0.825                                  | 0.825 (0.786–0.863)                   |
| Sens         | 0.746                                  | 0.746 (0.707–0.785)                   |
| Spec         | 0.836                                  | 0.836 (0.786–0.885)                   |
| F1           |                                        | 0.852 (0.802–0.904)                   |
| FPR          |                                        | 0.083 (0.073–0.093)                   |
| NPV          |                                        | 0.070 (0.050–0.090)                   |
| PPV          |                                        | 0.745 (0.707–0.785)                   |

Note: The median (interquartile range (IQR)) prediction value for the physicians was 0.36 (0.15–0.70) vs. 0.51 (0.12–0.86) for the RNN.
In addition, we investigated the calibration of the RNN’s and physicians’ predictions. Calibration describes how close the predicted probabilities are to the observed frequencies. A perfectly calibrated model would have one point at (0,0) and one at (1,1) in a calibration plot (it would always predict 0 for negatives and 1 for positives). For a well-calibrated model, the points lie on the diagonal between (0,0) and (1,1). Figure 2c illustrates that in the intervals of high prediction values of physicians, the predicted frequencies of AKI largely correspond to the observed frequencies (upper right part of the calibration curve). However, for several patients that developed AKI, physicians predicted low AKI probabilities (false-negative predictions, lower left part of the calibration curve). This is also reflected in the observation that the physicians’ median (IQR) prediction value was lower than the RNN’s (Physicians: 0.36 (0.15–0.70) and RNN: 0.51 (0.12–0.86)). Overall the physicians’ predictions were not well calibrated ($p < 0.001$, $X^2 = 165.5$, df = 8, Hosmer-Lemeshow-test).

In contrast, Fig. 2d displays a very well calibration ($p = 0.37$, $X^2 = 8.67$, df = 8, Hosmer-Lemeshow-test) for the RNN, with most of the points lying very close to the diagonal, even in intervals of low prediction values.

We investigated the performance of our RNN and physicians at different points in time before the event (AKI or non-AKI/discharge) (see Table 3). Not-surprisingly, both, humans and RNN, performed worse when the event was further away in time. However, low sensitivity rates could also be observed when the event was very close ($\leq 2$ h). In this group the median total observation length was very short, meaning that patients who developed AKI, developed it rapidly after surgery. Thus, there was probably not enough information available before the event to reliably predict AKI. However, even in this interval, the RNN reached a sensitivity of 0.789.

**DISCUSSION**

We developed an RNN for real-time prediction of postoperative AKI within 7 days after cardiothoracic surgery—based on routinely collected features during the hospital stay and then retrospectively validated it on an independent test set.

To test the clinical significance, we performed a side-by-side comparison of our model against experienced physicians. Such direct comparisons are highly needed, but hardly ever performed in clinical ML studies. We had expected our model to perform nearly as well as the physicians, and had designed our study as a non-inferiority-experiment. Surprisingly, our RNN significantly outperformed experienced clinicians in terms of the mean of our performance metric S. (S indicates how close a prediction is to the observed outcome). In addition, the model reached a significantly higher AUC than the physicians ($0.901$ vs. $0.745$, $p < 0.001$, DeLong’s test) and was overall well calibrated (Hosmer-Lemeshow-Test: $p = 0.37$ vs. $p < 0.001$ for physicians).

Physicians showed an overall low sensitivity of 0.594 at AKI prediction. They predicted lower risk probabilities in general. They reached a maximum sensitivity of 0.793 for the 2–6 h interval before the event and a minimum sensitivity of 0.387 for the...
Table 2.

| Threshold-independent metrics, (95% CI) | Metrics based on a threshold of 0.5 for positive/negative classification, (95% CI) |
|---------------------------------------|-----------------------------------------------------------------------------------|
|                                       | AUC  | PR_AUC | Brier | Acc  | Sens  | Spec  | F1    | FPR  | NPV   | PPV   |
| RNN                                   | 0.901 | 0.907  | 0.891 | 0.845 | 0.891 | 0.797 | 0.851 | 0.160 | 0.842 | 0.850 |
| Physicians                            | 0.745 | 0.747  | 0.746 | 0.594 | 0.521 | 0.667 | 0.829 | 0.171 | 0.671 | 0.741 |

24–48 h interval before the event. Thus, they systematically underestimated the risk of AKI. This suggests that physicians mainly recognize AKI stage 3 or dialysis and that lower AKI stages are erroneously considered unproblematic. It has been demonstrated, however, that even minor increases in serum creatinine after cardiac surgery are associated with an increased mortality risk.

The participating physicians each had at least one year working experience on a cardiothoracic intensive care unit (ICU), but were no specialists in nephrology. This reflects a realistic clinical setting on an ICU, where nephrologists are usually not available around the clock.

In contrast to the physicians, our RNN yielded an overall high sensitivity of 0.851 with a maximum sensitivity of 0.971 in the 2–6 h interval before the event and a minimum sensitivity of even 0.750 in the 48–168 h interval before the event. In summary, our RNN was superior to experienced physicians in the prediction of AKI after cardiothoracic surgery.

From a modeling point of view, our RNN could easily be integrated into an EHR system. It does not require any additional human input as all data transformation is implemented programmatically. Allowing for personalized predictions, it may enable earlier identification and intervention in high-risk patients and thus contribute to an improvement of patient care and safety. However, the transfer of such a retrospective model from research to real implementation raises additional challenges. Technical barriers, data security when exporting personal data to external software systems, and business considerations may be diverse and can conflict with each other.

Our model achieved highly accurate results with an overall AUC of 0.893 in our internal validation. It outperformed existing classical prediction models that are based on logistic regression from static pre- and intraoperative variables, as well as a dynamic model that predicted AKI at three points in time (pre-operative, at ICU admittance and 24 h after ICU admittance). These models reached AUCs ranging from 0.72–0.85 in their respective internal validation cohorts and used slightly different definitions of AKI (see Table 4). The proposed model does not create additional workload for physicians, as it only used routinely collected data of the EHR. As such, it only employs data that is available at the time of prediction and all data transformations are implemented programmatically. It is worth noting that the model performed very well, although it was built on a relatively small sample size of 2224 admissions.

Previous studies have demonstrated the benefits of using ML for AKI prediction. Thottakkara et al. applied different ML approaches to forecast postoperative AKI and observed promising performances in their internal validation cohort (AUC between 0.797 and 0.858). Bihorac et al. used an ML algorithm to assess the risk of 8 postoperative complications including AKI and reported an AUC of 0.80 (0.79–0.80) for AKI prediction. The approach of both studies, however, relied exclusively on static, mostly preoperative features.

A multi-center ward-based AKI prediction model was developed by Koyner et al. using a discrete time survival model with an AUC (95% CI) of 0.76 (0.76–0.77) for AKI of at least stage 2.

In 2018, Koyner et al. published another study using EHR data for AKI risk prediction and reached an AUC (95% CI) of 0.90 (0.90–0.90) for predicting stage 2 AKI within the next 24 h and 0.87 (0.87–0.87) within the next 48 h. Cheng et al. built ML models to forecast AKI over various time horizons and obtained an AUC of 0.765 (prediction one day before the event). In these studies, however, the urine output criterion of AKI, a central component in the KDIGO definition was not integrated, which can lead to a false-negative classification of AKI cases. In our training and test cohort around 30% of the AKI cases were defined by the urine criteria of KDIGO (see Supplementary Table 8). We can assume that a substantial proportion of the patients in the above studies would
Table 3. Performance metrics of recurrent neural network (RNN) and physicians in temporal dependence to the event.

| Predictor | Time to event | patients | AKI | MOL | AUC | PR_AUC | Brier | Acc | Sens | Spec | F1 | FPR | NPV | PPV |
|-----------|---------------|----------|-----|-----|-----|--------|-------|-----|------|------|----|-----|-----|-----|
| RNN       | 0 h to 2 h    | 54       | 19  | 8.3 h | 0.913 | 0.837 | 0.113 | 0.870 | 0.789 | 0.914 | 0.811 | 0.086 | 0.899 | 0.833 |
| Physicians| 0 h to 2 h    | 54       | 19  | 8.3 h | 0.709 | 0.552 | 0.199 | 0.759 | 0.632 | 0.829 | 0.649 | 0.171 | 0.806 | 0.667 |
| RNN       | 2 h to 6 h    | 63       | 29  | 12.5 h | 0.881 | 0.88  | 0.13  | 0.825 | 0.862 | 0.794 | 0.820 | 0.206 | 0.871 | 0.781 |
| Physicians| 2 h to 6 h    | 63       | 29  | 12.5 h | 0.853 | 0.861 | 0.152 | 0.794 | 0.793 | 0.794 | 0.780 | 0.206 | 0.818 | 0.767 |
| RNN       | 6 h to 12 h   | 63       | 34  | 17.8 h | 0.942 | 0.948 | 0.088 | 0.921 | 0.971 | 0.862 | 0.930 | 0.138 | 0.962 | 0.892 |
| Physicians| 6 h to 12 h   | 63       | 34  | 17.8 h | 0.811 | 0.798 | 0.19  | 0.746 | 0.618 | 0.897 | 0.724 | 0.103 | 0.667 | 0.875 |
| RNN       | 12 h to 24 h  | 74       | 42  | 36.4 h | 0.888 | 0.921 | 0.128 | 0.824 | 0.881 | 0.750 | 0.851 | 0.250 | 0.828 | 0.822 |
| Physicians| 12 h to 24 h  | 74       | 42  | 36.4 h | 0.693 | 0.706 | 0.257 | 0.689 | 0.667 | 0.719 | 0.709 | 0.281 | 0.622 | 0.757 |
| RNN       | 24 h to 48 h  | 60       | 31  | 46.4 h | 0.890 | 0.899 | 0.142 | 0.817 | 0.774 | 0.862 | 0.814 | 0.138 | 0.781 | 0.857 |
| Physicians| 24 h to 48 h  | 60       | 31  | 46.4 h | 0.718 | 0.774 | 0.246 | 0.633 | 0.387 | 0.897 | 0.522 | 0.103 | 0.578 | 0.800 |
| RNN       | 48 h to 168 h | 36       | 20  | 99.0 h | 0.875 | 0.929 | 0.132 | 0.806 | 0.750 | 0.875 | 0.811 | 0.125 | 0.737 | 0.882 |
| Physicians| 48 h to 168 h | 36       | 20  | 99.0 h | 0.647 | 0.741 | 0.274 | 0.611 | 0.400 | 0.875 | 0.533 | 0.125 | 0.538 | 0.800 |

AKI number of patients with acute kidney injury, MOL median total observation length, AUC area under curve, PR_AUC precision-recall AUC, Brier Brier score, Acc accuracy, Sens sensitivity, Spec specificity, F1 F1-score, FPR false-positive rate, NPV negative predictive value, PPV positive predictive value.

Fig. 2 Discrimination and calibration of the predictions of recurrent neural network (RNN) and physicians. a receiver operating characteristics (ROC), b precision-recall curve, c calibration of physicians, d calibration of RNN. AUC area under curve. H-L Hosmer-Lemeshow-Test, PR_AUC precision-recall AUC. The RNN outperformed clinical physicians regarding AUC (a) and PR_AUC (b). Physicians systematically underestimated the risk of acute kidney injury (predicted risks < observed risks, c). In contrast, the RNN was overall well calibrated (d).
also have met the urine criteria first. Probably not all of them have been classified as false-negative, as they might have met the creatinine criterion at a later stage. In our population, 11% of the AKI-cases in the training set and 12% in the test set exclusively fulfilled the urine criterion and would have been diagnosed false-negatively without this criterion. The median (IQR) diagnosis delay of patients who met both criteria within 7 postoperative days was 14.0 h (6.3–27.3 h) in the training set and 13.3 h (5.3–22.4 h) in the test set. Especially in models with short prediction horizons, there is a high risk that the prediction of imminent AKI and consequently initiation of preventive measurements is delayed when not integrating the urine criterion.

In addition, these previous models were restricted to patients with a serum creatinine of <3 mg/dl (Koyner et al.) or even normal when not integrating the urine criterion. Consequently initiation of preventive measurements is delayed. It has been shown that early intervention can prevent AKI or its progression to higher stages. Therefore, the prediction of our model was not limited to AKI requiring dialysis, but rather AKI at any stage.

Another study for continuous AKI prediction on a large data set was performed by Tomashev et al.34. The developed RNN predicted AKI stage 2 or 3 with an AUC of 0.971 24 h before onset. Also in these studies the urine output criterion of AKI was not incorporated. In addition, in the study of Tomashev et al. only patients were included for whom at least one year of EHR data were available before admission. They added aggregate features of up to five years of historical information of each individual patient. This approach requires that patients are already known in the admitting hospital, which is often not the case. It is unclear how their algorithm would perform on patients without any prior medical history. In contrast, we used a real uncurated data stream in our model that only contained information generated after admission.

Meyer et al.35 used an RNN to predict AKI requiring dialysis, mortality and postoperative bleeding after cardiac surgery using routinely collected parameters within the first 24 hours after surgery. The deep-learning model provided very accurate predictions (positive predictive value (PPV)/sensitivity for AKI: 0.87/0.94) that outperformed usual clinical risk scores.

Our model predicted AKI at a time frame up to 7 days after cardiothoracic surgery. Compared to the observation windows of the studies mentioned above, this is a much longer time period. Events in the near future are usually easier to predict than those in the more distant future. To intervene early when the kidneys are merely at risk of injury, a longer prediction window might be necessary. It has been shown that early intervention can prevent AKI or its progression to higher stages. Therefore, the prediction of our model was not limited to AKI requiring dialysis,
but included the prediction of AKI stages 2 or 3 according to the KDIGO definition.

To conclude, based on a relatively small sample size, we developed a highly accurate model for the prediction of AKI after cardiac surgery that significantly outperformed experienced physicians, could potentially be integrated into EHR systems and might prevent severe complications following AKI through real-time patient surveillance. In a long-term perspective, an extension of the application from a simple risk prediction model to treatment decision support tool is also conceivable.

This study has several shortcomings. The observation periods of the included patients varied widely in length. For most patients it ended in <3 days while some outliers lasted for up to 7 days. We only used the start of nephrotoxic drug administration as a feature. Consideration of exact dose, administration route (e.g., i.v., p.o, ...), and administration length could reflect the underlying pharmacodynamics better and improve the prognostic performance.

Our RNN is currently cohort specific for cardiothoracic surgery patients that most likely have different characteristics and risk factors than, e.g., neurosurgical patients. Implementing the same approach on other patient cohorts could give a deeper insight into the generalizability of our method.

Our study is retrospective. Thus, in our RNN vs. physicians head-to-head comparison, physicians only received EHR data and could not clinically evaluate patients. Information such as volume status (except for weight), general condition, etc. or additional examinations (e.g., ultrasound) were not available to them and to the RNN. This deviation from the physicians’ usual workflow in clinical practice may explain some of the observed performance deficits. Real clinical data can be very noisy, leading to reduced performance and greater burden of deploying completely automated systems. This stresses once again the fact that artificial intelligence should be utilised in support systems for physicians and not as their replacement.

External validation trials should be performed on prospective data. In addition, they should focus on usage and acceptance of a system such as the one described here in a real clinical setting.

METHODS
Ethics and reporting guideline
This study was approved by the institutional data protection officer and ethics committee of Charité – Universitätsmedizin Berlin (EA2/180/17). The approval included the collection of data on implied consent: We only used retrospective data and the patients were not actively involved in the study. The requirement of informed consent of the participating physicians was waived by the Institutional Review Board (IRB) of Charité – Universitätsmedizin Berlin due to anonymized data acquisition. Reporting of development and validation of the prediction model follows widely the guideline of the TRIPOD statement.

Patient selection process
We retrospectively analysed EHR time series data generated between October 2012 and February 2018 at a tertiary care center for cardiovascular diseases.

We included adult patients (18+) that were admitted at least once to the operating theatre for cardiothoracic surgery (15,564 admissions/13,895 patients). We excluded patients without any creatinine or urine flow values, patients receiving hemodialysis before the end of the operation or having a baseline creatinine level \( \geq 4.0 \text{ mg/dl} \) (2322 admissions/1487 patients).

Within this collection of 12,978 admissions, 1308 cases were identified with severe postoperative AKI defined as stage 2 or 3 according to KDIGO AKI guidelines—briefly, an increase in serum creatinine to at least twice the baseline value or a decrease in urine flow \(< 0.5 \text{ ml/kg/h} \) for \( \geq 12 \text{ h} \). As AKI can develop over multiple days, we defined a study period of 7 days after cardiothoracic surgery. The global AKI label of a patient was set positive when the KDIGO criteria stage 2 or 3 was fulfilled at any point within these 7 postoperative days.

The observation time of each patient started when the patient was transferred to the ICU or recovery room. It ended when the patient was either discharged, or when the KDIGO criteria for AKI stage 2 or 3 were fulfilled, or after 7 days after the end of the first surgery.

Each AKI-case was assigned a control out of the non-AKI pool (11,670 admissions/11,046 patients). The controls were matched to the cases on

![Flow chart of patient selection process](image_url)
observation length. Thus, we generated a balanced data set that we then randomly split into a training set (85%, 2224 admissions/2180 patients) and the remaining set (15%, 392 admissions/patients) while keeping the cases with their respective controls.

For the 392 patients of the remaining set we manually checked physicians' notes in the EHR data and consequently excluded 28 patients. Exclusion criteria were primarily insufficient documentation of the type of surgery, false recording of surgery times or notion of end-stage kidney disease in the patients’ history that was not detected by automated filtering.

Out of this set, we randomly selected 350 patients that formed the final test set for model evaluation and comparison with human-level performance. A detailed flow chart of the patient selection process is shown in Fig. 3.

The baseline characteristics were well balanced between the training and the test and are summarized in Supplementary Table 8.

The density distribution and a histogram of the observation periods for patients in the training and test sets is shown in Fig. 4. Most patients were either discharged or diagnosed with AKI within the first 3 days after the first surgery.

### Feature selection and preprocessing

We developed our model based on 96 routinely collected clinical parameters. Table 5 gives an overview of all considered features. They can be grouped into static features (e.g., patient and surgery characteristics, 25 features) that do not change over the observation period and frequently measured dynamic features (e.g., lab values, vital signs, blood gas values and fluid output, 49 features). In addition, we included a variety of widely administered agents that have been reported to potentially cause nephrotoxic effects42 (47 features).

The last creatinine/urea value before surgery was used as a baseline. If there was none available in the five days before surgery, we used the first postoperative value.

We observed that urine output was sometimes incompletely documented on normal wards. As this could lead to false-positive AKI diagnoses we considered urine values reliable only when they were recorded in the operation theatre, the recovery room or the ICU. Thus, on normal wards AKI was only defined by the creatinine criterion whereas in the recovery room or the ICU both AKI criteria (creatinine and urine) were used.

EHR systems are often designed with billing and revision purposes in mind, making certain retrospective therapeutic analyses difficult to conduct due to missing information28. In our case, the type of operation that patients underwent was available partly in unstructured textual and partly in categorical form. To access both types of data, we developed a separate set of bag-of-words logistic regression models that predicted the type of operation based on unstructured text describing the operation procedures. As explanatory variables we used all single words or abbreviations that occurred in the pool of text information in its training set. The probability of a specific surgery type \(Y_j\) \((j = 1, 2, \ldots, 17)\) was given by:

\[
P(Y_j = 1) = \frac{\exp(\beta_0 + \beta_1 x_1 + \beta_2 x_2 + \ldots)}{1 + \exp(\beta_0 + \beta_1 x_1 + \beta_2 x_2 + \ldots)}
\]

where \(x_j\) denotes a count variable indicating how often word/abbreviation \(j\) occurred in a patient’s surgery procedure description \((j = 1, 2, \ldots, no.\text{ distinct words/abbreviations})\). For further information see Supplementary Note 1, Supplementary Tables 9 and 10.

Time sequences with 15-min intervals of all features served as input to our model.

Except for the nephrotoxic agents, missing values were filled by forward imputation. If no precedent value was available, static default values defined by a clinical expert were imputed (one value per feature). The same default values were used for all patients and they were imputed programatically. They are shown in Supplementary Table 11.

It is extremely difficult to determine the exact effect duration of a drug due to varying excipients, dosages, drug combinations, application types and patient conditions. Therefore, the administration of a drug was considered as an event. For each nephrotoxic agent class in Table 5 a binary feature was created and its value was set to 1 only at the single time slice immediately following the administration of the drug.

Except for the operation types all continuous features were then scaled as follows49:

\[
X_{\text{scaled}} = \frac{X - \mu(X_{\text{train}})}{\text{IQR}(X_{\text{train}})}
\]

where \(\mu(X_{\text{train}})\) denotes the median and IQR\((X_{\text{train}})\) the IQR of the feature \(X\) in the training set. In total, the model was built on a data matrix of 36,244,608 single data points.

For patient selection, preprocessing of features and imputation of missing data, we used R v3.3.3 (R Core Team (2017), R: A language and environment for statistical computing, R Foundation for Statistical Computing, Vienna, Austria. URL https://www.R-project.org/) and Python v3.6.7 (The Python Software Foundation, Beaverton, OR) with modules IPython50 (v7.5.0), Matplotlib51 (v3.1.0), Scikit-learn52 (v0.19.1), Pandas53 (v0.24.2) and Numpy54 (v1.16.2).

### Modeling

In contrast to classical prediction models such as logistic regression, RNNs are able to capture the temporal development of features in a truly sequential fashion as they incorporate information about preceding time steps, links between single timesteps and a direct indicator of the current position in the timeline (see Fig. 5).

We constructed a set of RNNs with different architectures (preceding convolutional layer, different cell types) which allow to process dynamic features in a sequential fashion and are able to capture the temporal development of features in a truly sequential fashion as they incorporate information about preceding time steps, links between single timesteps and a direct indicator of the current position in the timeline (see Fig. 5).

As the parameters of an RNN depend on their initialization and the order in which the training instances are presented, 10 final models with the same hyperparameters but different initializations were trained on the training set. Our final model comprised a uniform ensemble of the 10 constituent models.

For the modeling process we used Python v3.6.7 (The Python Software Foundation, Beaverton, OR) with modules TensorFlow55 (v1.13.1), TensorFlow 56 (v1.13.1), Scikit-learn57 (v0.19.1), Pandas58 (v0.24.2) and Numpy59 (v1.16.2).
where $t_0$ is the number of timesteps, $y_i$ the prediction at time step $i$ and $y_{ij}$ the true label of patient $j$.

The $\text{MSE}_{\text{pat}}$ ranges from 0 to 1, with value 0 meaning perfect prediction and 1 meaning worst prediction. Random guessing (always predicting 0) would result in a $\text{MSE}_{\text{pat}}$ of 0.25. In contrast to the metrics mentioned above, the $\text{MSE}_{\text{pat}}$ is independent of the individual observation length of a patient and the resulting number of predictions per patient.

We adjusted the threshold for positive class prediction until a fixed sensitivity of 0.85 on cross-validation folds in the training set was reached (threshold $= 0.41$).

Our model predicted the risk of developing AKI every 15 min after the initial surgery. The predictions of an individual patient can be regarded as a cluster of usually highly correlated data. We therefore had to adjust the CIs of our model’s metrics. We calculated the 95% CI of each metric $X$ as follows:

$$X + 1.96\sigma(X)$$

with a standard error $\sigma(X)$ of variable $X$ of

$$\sigma(X) = \sqrt{\frac{X(1 - X)}{n_{\text{eff}}}}$$

To account for intracluster correlation, our sample size $n$ was adjusted, resulting in an effective sample size of

$$n_{\text{eff}} = \frac{n}{\text{DE}} = \frac{\sum_{i=1}^{k} \sum_{j=1}^{m_i} 1}{\text{DE}}$$

where $k$ is the number of patients and $m_i$ the number of time steps of patient $i$. DE denotes the design effect, also called variance inflation factor, and can be calculated as follows:

$$\text{DE} = \frac{mk}{\sum_{i=1}^{k} \frac{m_i}{1 + m_i - 1}\text{ICC}}$$

where ICC is the intracluster correlation coefficient. The ICC was calculated using the R package ICC61 (v2.3.0).

Comparing RNN vs. human performance

We set up an experiment to compare the performance of our RNN against that of experienced physicians (see Fig. 1). For each patient in the test set, a quasi-random point in time in their observation period was chosen, further denoted as the ‘prediction point’. In contrast to real uniform random samples, which tend to form clusters and contain regions without any points at all, quasi-random sequences reduce the probability of cluster formation while still being uniformly distributed62,63. This method prevented us from accidentally exclusively sampling prediction points from e.g. the first half of the patients’ observation periods.

### Table 5. Input feature overview.

| Feature Group (no. features) | Features                                                                 |
|-----------------------------|--------------------------------------------------------------------------|
| Patient characteristics (4) | Age, sex, weight, height                                                 |
| Laboratory results (25)     | Phosphate, total bilirubin, baseline creatinine, creatinine, baseline urea, urea, glomerular filtration rate, creatine kinase (CK), CK-MB, red blood count, white blood count, platelets, C-reactive protein, gamma-glutamyltransferase, glutamic oxaloacetic transaminase, hemoglobin, international normalized ratio, lactate dehydrogenase, magnesium, hematocrit, prothrombin time, partial thromboplastin time, mean corpuscular hemoglobin, mean corpuscular volume, mean corpuscular hemoglobin concentration |
| Surgery characteristics (20)| Aortic cross-clamp time, cardiopulmonary bypass time, time in operation theatre, surgery procedure (from logistic regression text model, see Supplementary Note 1) |
| Vital signs (8)             | Systolic, mean and diastolic arterial pressure, central venous pressure, heart frequency, pulse, body temperature, oxygen saturation |
| Arterial blood gas values (BGA) (15) | Base excess, bicarbonate, glucose, hemoglobin, oxygen saturation, partial pressure of carbon dioxide and oxygen, total carbon dioxide, pH level, potassium, sodium, calcium, lactate, carboxyhemoglobin, oxyhemoglobin |
| Fluid output (2)            | Bleeding Rate, urine flow rate                                           |
| Nephrotoxic agents (22)     | Allopurinol, Aminoglycosides, Amphotericin B, Antiplatelet agents (clopidogrel, ticlopidine), Benzodiazepines, Cephalosporins, Cyclosporine, Haloperidol, Ketamine, Nonsteroidal anti-inflammatory drugs, Paracetamol, Penicillines, Proton pump inhibitors, Pyrazolone derivatives, Quinolones, Ranitidine, Rifampin, Sulfonamides, Tacrolimus, (Val-)/Ganciclovir, Aciclovir, Vancomycin |

Red Blood Cell Transfusions

Fig. 5 Architecture of a recurrent neural network (RNN). At each time step, the model receives the current time slice data as input as well as the own output from the preceding time step. The features are captured in a truly sequential fashion.
At each prediction point, a physician and the RNN had to predict whether a patient would develop AKI within the first 7 days after surgery. All time series information up to the ‘prediction point’ was graphically displayed for the physicians to mimic the electronic patient chart—although here not in 15-min intervals but in the originally recorded time resolution (up to 1 min).

To create a realistic setting, physicians not only received information about nephrotoxic agents, but of all administered drugs. In addition, the surgery type was given to them as unstructured text manually extracted from physicians’ notes. This information was not available to the RNN model. Physicians were explicitly informed about the incidence rate of 50% AKI in our test set.

A physician as well as the RNN made a probability prediction r of the development of AKI for each patient at the respective prediction point. In addition, the physicians made a binary decision (development of AKI: yes/no).

We asked 14 physicians to participate in our study, 10 of whom agreed (response rate = 0.71). All had to meet the selection criteria of ≥5 years of clinical experience and ≥1 year of work experience on a cardiothoracic ICU. From the 10 volunteers we selected seven physicians with different levels of expertise (senior resident up to senior consultant) to create a most realistic setting. Their working experience on a cardiothoracic ICU ranged from at least one year up to several years. None of the participating physicians were specialists in nephrology as nephrologists are usually not constantly available on an ICU. Each physician made predictions for 50 different patients.

Statistical analysis

The initial aim of our study was to show that the RNN is not inferior to experienced physicians in the prediction of AKI. For both, RNN and physicians, the predictive quality of each probability prediction r was measured by a score S as follows:

\[ S = r, \text{ if the patient developed AKI} \]

\[ S = 1 - r, \text{ if the patient did not develop AKI} \]

A prior investigation of the RNN’s predictions had shown that S was normally distributed. Thus, for sample size calculation and power analysis we considered the transformed score \( X \), which was approximately normally distributed:

\[ X = -\log(-\log(S)) \]

We assumed that \( X \) of the physicians’ predictions would also be normally distributed.

Based on a significance level of \( \alpha = 0.025 \), a power of at least 80% and a non-inferiority margin of \( \delta = 0.3 \) (this corresponds to a non-inferiority margin of 5.5% for sensitivity + specificity), we obtained a sample size of \( N = 350 \).

Both for RNN and physicians, we calculated AUC, PR, AUC, brier score, accuracy, sensitivity, specificity, PPV, NPV, FPR, and F1-score. We set the threshold for positive class prediction to 0.5 as this was also the threshold in the physicians’ predictions that corresponded to the ‘yes/no’-classification. We calculated CIs for all metrics as described in Section ‘Measuring RNN Performance’ whereas the effective sample size was \( n_{eff} = n = 350 \) as there was no clustering.

For the statistical comparison of S between RNN and physicians we applied a paired t-test. We used DeLong’s method to compare the two correlated ROC curves using the R package pROC \(^6^5 \) (v1.9.1). In addition, we investigated the calibration of both, physicians’ and RNN’s predictions, with the Hosmer-Lemeshow-Test using the R package ResourceSelection \(^6^6 \) (v0.3-2). All three comparisons mentioned above were tested on a significance level of \( \alpha = 0.05 \).

Reporting summary

Further information on research design is available in the Nature Research Reporting Summary linked to this article.

DATA AVAILABILITY

The EHR data used in this study contain protected health information (PHI) and cannot be published for reasons of data protection. The dataset may be available from the German Heart Center Berlin subject to ethical approvals.

CODE AVAILABILITY

The source code of this publication may be obtained from the corresponding author upon request. To enable independent replication, we describe the experiments and implementation in detail in the ‘Methods’ section and in the Supplementary Information.

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