

Machine Learning–Aided Chronic Kidney Disease Diagnosis Based on Ultrasound Imaging Integrated with Computer‑Extracted Measurable Features

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Abstract

Although ultrasound plays an important role in the diagnosis of chronic kidney disease (CKD), image interpretation requires extensive training. High operator variability and limited image quality control of ultrasound images have made the application of computer-aided diagnosis (CAD) challenging**.** This study assessed the efect of integrating computer-extracted measurable features with the convolutional neural network (CNN) on the ultrasound image CAD accuracy of CKD. Ultrasound images from patients who visited Severance Hospital and Gangnam Severance Hospital in South Korea between 2011 and 2018 were used. A Mask regional CNN model was used for organ segmentation and measurable feature extraction. Data on kidney length and kidney-to-liver echogenicity ratio were extracted. The ResNet18 model classifed kidney ultrasound images into CKD and non-CKD. Experiments were conducted with and without the input of the measurable feature data. The performance of each model was evaluated using the area under the receiver operating characteristic curve (AUROC). A total of 909 patients (mean age, 51.4 ± 19.3 years; 414 [49.5%] men and 495 [54.5%] women) were included in the study. The average AUROC from the model trained using ultrasound images achieved a level of 0.81. Image training with the integration of automatically extracted kidney length and echogenicity features revealed an improved average AUROC of 0.88. This value further increased to 0.91 when the clinical information of underlying diabetes was also included in the model trained with CNN and measurable features. The automated step-wise machine learning–aided model segmented, measured, and classifed the kidney ultrasound images with high performance. The integration of computer-extracted measurable features into the machine learning model may improve CKD classifcation.

Keywords Chronic kidney disease · Computer-aided diagnosis · Convolutional neural network · Deep learning · Kidney ultrasound · Machine learning

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Introduction

Diferentiating chronic kidney disease (CKD) from acute kidney injury (AKI) is the most fundamental step in the management of impaired kidney function. AKI refers to a sudden loss of kidney function, whereas in CKD, kidney damage usually persists for some amount of time. When the cause of kidney damage is properly identifed and removed, kidney function eventually recovers in AKI [\[1](#page-8-0)]. However, in patients with CKD, kidney damage is irreversible and often progressive, leading to end-stage renal disease (ESRD) [\[2](#page-8-1)]. Periodic blood test results that refect kidney function are useful in distinguishing CKD from AKI. However, when such data are not obtainable, making an accurate diagnosis of CKD versus AKI is often challenging.

Imaging studies are key diagnostic modalities for identifying CKD [\[3,](#page-8-2) [4\]](#page-8-3). In particular, ultrasound imaging is the most important scheme used to detect CKD in clinical practice [\[5](#page-8-4)–[7\]](#page-8-5). The well-known advantages of ultrasound imaging, including its low cost and noninvasive nature and lack of ionizing radiation, have made it an attractive option for detecting kidney diseases [\[8](#page-8-6)]. The characteristic changes in kidney length, echogenicity, and corticomedullary differentiation enable a CKD diagnosis without the need for invasive procedures such as kidney biopsy [[9\]](#page-8-7). However, kidney ultrasound image acquisition and interpretation are prone to operator dependency [[10–](#page-8-8)[12](#page-8-9)]. Ultrasound-based diagnosis is highly dependent on the ability and experience of the person performing ultrasonography, and therefore requires extensive training and experience for optimal use. This limits its widespread application in facilities or regions lacking well-experienced specialists.

The application of machine learning in medical image analysis has been widely studied [[13,](#page-8-10) [14\]](#page-8-11). Machine learning algorithms are helpful for reducing false-positive results in computed tomography (CT) image lung nodule detection $[15, 16]$ $[15, 16]$ $[15, 16]$ $[15, 16]$ $[15, 16]$. In addition, the ability to identify the malignant potential of breast nodules on CT and magnetic resonance imaging (MRI) images was greatly improved with machine learning–based computer-aided diagnosis (CAD) [[17,](#page-8-14) [18](#page-8-15)]. Accumulating studies have also attempted to apply machine learning techniques to improve the detection rates of ultrasound imaging [[19,](#page-8-16) [20](#page-8-17)]. However, machine learning–obtained diagnostic accuracy in ultrasound was often inferior to that performed by experienced radiologist [[21](#page-8-18), [22\]](#page-8-19). This would be due to the ultrasound-specifc characteristics such as low image quality and high inter- and intra-observer variability [[10\]](#page-8-8).

In this study, assessments were performed to evaluate whether integrating data on measurable features into a convolutional neural network (CNN) could improve the CAD accuracy of CKD in ultrasound imaging. Segmentation of the kidney and extraction of measurable features, such as kidney length and echogenicity, were performed using deep learning methods.

Material and Methods

Study Design and Subjects

(1) age $<$ 18 years; (2) ESRD with dialysis at the time of ultrasound imaging; (3) previous kidney transplantation; (4) previous nephrectomy; (5) known kidney cysts, solid mass, or hydronephrosis; and (6) missing liver or kidney images on ultrasound. A total of 909 patients were included in the fnal analysis. This study was conducted in accordance with the Declaration of Helsinki and approved by the Institutional Review Board of Yonsei University Health System Clinical Trial Center (1–2018-0039). The need for informed consent was waived by the institutional review board owing to the study's retrospective design.

Patient Classification

Serum creatinine levels were determined using an isotope dilution mass spectrometry–traceable method at a central laboratory. The eGFR was calculated using the Chronic Kidney Disease Epidemiology Collaboration (CKD-EPI) creatinine equation [[25](#page-8-22)]. Laboratory results within 3 months before kidney ultrasound imaging were used for patient classification. Those whose eGFR was \geq 90 mL/min/1.73 m² within the prior 3 months were labeled the non-CKD group. Those whose eGFR was $< 60 \text{ mL/min}/1.73 \text{ m}^2$ throughout the prior 3 months were classifed as the CKD group (Fig. [1](#page-2-0)).

Data Collection and De‑Identification

Demographic and laboratory data were retrieved from electronic medical records. Data on the presence of comorbidities at the time of ultrasound imaging, including hypertension, diabetes, and coronary artery disease, were also collected. All kidney ultrasound images were obtained by board-certifed radiologists. Ultrasound machines (Philips Healthcare, [Amsterdam, Netherlands], Siemens Healthineers [Erlangen, Germany], and GE Healthcare [Chicago, Illinois, USA]) were used for the ultrasound image acquisition. For training and testing, ultrasound images containing both kidney and liver were used. The number of ultrasound images including both the kidney and liver difered for each patient. An average of 3.7 images were employed for evaluation. Although the number of ultrasound images varied for each patient, a single prediction for each patient was derived using average voting. Therefore, the weights of patients were equally used for evaluation. The ultrasound images were transferred as Digital Imaging and Communications in Medicine (DICOM) fles, and pixel data were converted to portable network graphics fles. Images that did not include annotations for segmentation or length measurements were selected for the analysis. In addition, patients' personal information was deleted from the image fles before the evaluation. This de-identifcation program was newly developed for the study. It blacked out the upper corners of the images with personal data to mask the identifying

Fig. 1 Flow diagram of the included participants

information. Since this program only modifed the upper corners of the images, it did not fabricate the ultrasound images of the organs that were used for analysis.

Kidney and Liver Segmentation

A Mask regional convolutional neural network (R-CNN) model, which adds an additional head to the Faster R-CNN model, was used for the kidney and liver segmentation [\[26,](#page-8-23) [27](#page-8-24)]. The Mask R-CNN is commonly used for instance segmentation where each pixel is assigned an object class and instance identity. This instance segmentation method performs segmentation in a two-step manner: initial detection of the object region followed by object segmentation within the detected area. The U-Net and fully convolutional network, which is widely used in semantic segmentation techniques in medical image segmentation, requires post-processing to flter out small segments and remove holes within the segmentation [[28](#page-9-0)–[30](#page-9-1)]. However, the instance segmentation method requires no additional processing for incorrect segmentation removal. In addition, fewer holes were created compared to one-step methods. The Mask R-CNN models used ImageNet-pretrained ResNeXt101 as the backbone [\[26,](#page-8-23) [31](#page-9-2)]. For feature extraction, ROIAlign and Feature Pyramid Networks (FPN) were used [[32](#page-9-3)]. The ultrasound images were input into the backbone, and the inference and recombined features were extracted from each layer through the FPN. The extracted features were resampled at diferent scales and ratios at each anchor using ROIAlign and inputted into the Mask R-CNN head for detection, classifcation, and segmentation. Multi-task loss was used during training. Softmax cross-entropy for classifcation, smooth L1 for bounding box regression, and binary cross-entropy were used for segmentation. To obtain the trained segmentation model, the combined loss was minimized using a gradient descent (Fig. [2A](#page-3-0)).

Segmentation, Training, and Validation

For segmentation training and validation, 9407 ultrasound images were acquired from 456 patients. A trained technician manually depicted the two-dimension regions of the kidney and liver under nephrologist supervision. A total of 3917 kidney segmentations and 7234 liver segmentations were collected (some images contained both kidney and liver); of them, 993 images were randomly selected for validation. The stochastic gradient descent was utilized with a batch size of 48, and the learning rate started at 0.001 and decreased by 0.1 at 30,000 steps and 40,000 steps. The model was trained in 50,000 steps. The segmentation performance was validated on the validation dataset every 2500 steps using dice coefficient, and the most accurate model was selected.

Fig. 2 The schematic diagram of the algorithm used. A Mask R-CNN model was used to perform segmentation of the kidney and liver (**A**) and extract measurable features (**B**). ResNet18 model was used

to classify the kidney ultrasound images into CKD or non-CKD (**C**). CKD, chronic kidney disease; R-CNN, regional convolutional neural network; ROI, region of interest; RPN, region proposal network

Extraction of Measurable Features

Measurable features were extracted from the segmented ultrasound images using the Mask R-CNN model. The kidney length was derived by calculating the longest horizontal length of the segmented kidney. To derive the actual length (cm) of the kidney, the kidney length in the image space was calibrated using the machine information stored in the DICOM fle. Measuring the absolute level of echogenicity is difficult since echogenicity varies with ultrasound machine operating settings and patient's state of hydration. Therefore, kidney cortical echogenicity is generally graded relative to the liver parenchyma [[33\]](#page-9-4). Liver echogenicity was obtained by averaging the pixels of the segmented liver area. For calculation of the kidney cortical echogenicity, the segmented kidney was first divided into 3×4 grid cells along the longest horizontal axis. The top middle cell was considered to represent cortical echogenicity, and the pixels in this area were averaged (Fig. [2B](#page-3-0)).

Classification of Kidney Ultrasound Image

For the classification model, the ResNet18 model was used [[34\]](#page-9-5). This model classifed kidney ultrasound images into CKD or non-CKD. To input meaningful areas in the classifcation model, the kidney region was cropped into a rectangular shape aligned with the longest horizontal axis of the kidney segmentation. ResNet uses a skip connection to the convolutional layer, where the input features are added to convoluted features. After the convolutional stage was fnished, global average pooling was applied to the convoluted output feature, and a fully connected layer was used to classify the images as CKD or non-CKD. For classifcations considering both the image feature and the extracted measurable feature, the features were pre-processed via standardization (mean of 0 and standard deviation of 1) and combined through channel-wise concatenation to the penultimate layer and input to the following fnal fully connected layer. Softmax cross-entropy was used to calculate the error between the true label and the model prediction (Fig. [2](#page-3-0)C).

Classifcation, Training, and Testing

Using the model trained in the segmentation, a total of 3495 ultrasound images that contained both the kidney and the liver were acquired from 909 patients. These images were used for the classifcation training and testing. For the training dataset, 839 patients (3001 images, 92%) were randomly selected. The remaining 70 patients (247 images, 8%) were used as the testing dataset. For the validation dataset, the patients in the training dataset were divided into 10 groups. Consequently, 10 validation datasets were constructed by selecting one group for validation (approximately 249 images [8%]) and the others for training. To reduce the diference in brightness between ultrasound manufacturers, images that contained both kidney and liver segments were selected, and the kidney echogenicity was normalized against that of the liver. The adaptive moment estimation optimizer was used with a batch size of 16 and a learning rate of 0.001, and the learning rate was reduced to 90 epochs following the cosine annealing learning rate scheduler [[35,](#page-9-6) [36](#page-9-7)]. In addition, the augmentation techniques (random fip, translate, rotate, scale, contrast, noise, dropout, and blur) were used for training. To ensure reliability, 10 training sessions were conducted on one validation dataset (tenfold cross-validation), resulting in a total of 100 training processes (10 training sessions with diferent training seeds for each of the 10 diferent validation datasets). For each training process, the most accurate model was chosen, and the best model was evaluated in the test dataset. Three types of experiments were conducted: "image training without any measurable features," "image training with extracted measurable features," and "image training with extracted measurable features and diabetes information." Each of the three experiments was trained with a randomly initialized model and an ImageNet-pretrained model, resulting in a total of 600 experiments. The fnal classifcation results were then calculated for each subject. In cases where multiple ultrasound images included both kidney and liver per subject, each network probability output was averaged to yield one diagnostic result. These evaluations were performed on Intel i7-8700 K and Nvidia GeForce GTX 1080 Ti.

Statistical Methods

Variables were expressed as mean and standard deviation (SD) for continuous variables and frequency and percentage (%) for categorical variables. Intergroup comparisons were performed using Student's *t*-test for continuous variables and Pearson's chi-squared test for categorical variables. The CNN-based CKD classifcation was compared with the diagnosis of CKD based on eGFR, while the model performance was evaluated using the area under the receiver operating characteristic curve (AUROC). All statistical analyses were performed using SPSS Statistics v.23 (IBM Corporation, Armonk, NY, USA) and STATA v.15.1 (Stata Corporation, College Station, TX, USA). Statistical signifcance was set at two-sided values of $p < 0.05$.

Results

Baseline Characteristics

The mean participant age was 51.4 ± 19.3 years; 414 (49.5%) of them were male (Table [1\)](#page-4-0). The non-CKD group comprised 524 (57.6%) participants, while 385 (42.4%)

Table 1 Patients' baseline characteristics

participants were allocated to the CKD group. The CKD group was older and had a higher prevalence of hypertension, diabetes, and coronary artery disease. The average eGFR was 115.5 ± 29.1 and 27.3 ± 11.0 mL/min/1.73 m² in the non-CKD and CKD groups, respectively.

Kidney Segmentation

To evaluate the performance of the segmentation implemented by the Mask R-CNN model, its segmented area was compared to the area depicted by trained technicians. In the validation dataset, the best model achieved a mean intersection on union (mIoU) of 81.62, while the frequencyweighted intersection on union (fwIoU) was 89.35 (Fig. [3](#page-5-0)A and B).

Measurable Feature Extraction

Measurable features, including kidney length and cortical echogenicity, were also obtained using the Mask R-CNN model. The average kidney length was 9.49 ± 1.11 cm in the non-CKD group and 8.90 ± 0.94 cm in the CKD group $(p<0.001;$ Fig. [4A](#page-5-1)). When the kidney-to-liver echogenicity ratio was compared between the non-CKD and CKD groups, the echogenicity was signifcantly higher in the CKD group than in the non-CKD group $(1.08 \pm 0.44 \text{ vs. } 1.42 \pm 0.53,$ *p*<0.001; Fig. [4](#page-5-1)B).

CKD Detection Performance Using Ultrasound Image Information Only

The model performance was evaluated using a test dataset that had been excluded from the training beforehand. Three experiments were completed, and 10 training sessions were performed for each of the 10 validation set experiments. The average AUROC from the model trained with ultrasound images achieved only a level of 0.81 (sensitivity 78.2%, specificity 71.5% , and accuracy 74.4%) (Fig. [5\)](#page-6-0).

Values are expressed as mean \pm SD for continuous variables and frequency (%) for categorical variables *CAD* coronary artery disease, *CKD* chronic kidney disease, *eGFR* estimated glomerular fltration rate

Fig. 3 Kidney segmentation using Mask R-CNN model. The best model achieved an mIoU of 81.62 and an fwIoU of 89.35. **A** and **B** are representative images of kidney and liver segmentation that were

CKD Detection Performance After Integrating Deep Learning–Extracted Measurable Features

The performance of CKD detection from the model trained with ultrasound images only was compared with models that integrated deep learning–extracted measurable feature data and information on diabetes. When achieved by the Mask R-CNN model. fwIoU, frequency-weighted intersection on union; mIoU, mean intersection on union; R-CNN, regional convolutional neural network

data on measurable features, including kidney length and kidney-to-liver echogenicity ratio, were extracted through deep learning algorithms, the average AUROC for CKD detection improved to 0.88 (sensitivity 86.1%, specificity 77.5%, and accuracy 81.2%). Adding information on the presence of diabetes among the participants into the training models resulted in a CKD detection AUROC of

Fig. 4 Extracted measurable features by the Mask R-CNN model. **A** The average lengths of the kidney were 9.49 ± 1.11 cm and 8.90±0.94 cm in non-CKD and CKD groups, respectively. **B** The

ratios of echogenicity of the kidney to liver were 1.08 ± 0.44 and 1.42±0.53 in non-CKD and CKD groups, respectively. CKD, chronic kidney disease; R-CNN, regional convolutional neural network

Fig. 5 ROC curve of CKD detection performance. Three types of experiments were conducted. There were 10 validation datasets for each experiment, and the AUROC was obtained for each dataset (**A**– **C**). The average AUROC was 0.81 in training without any measurable features, 0.88 in image training with extracted measurable fea-

0.91 (sensitivity 89.4%, specificity 82.9%, and accuracy 85.9%) (Fig. [5\)](#page-6-0).

Discussion

In this study, CKD was detected using kidney ultrasound image analysis using deep learning methods. A sequential step-by-step process was employed for the image analyses. First, kidney and liver segmentations were performed using a Mask R-CNN model. Next, the measurable features were extracted from the segmented kidney images. Features such as kidney length and relative cortical echogenicity compared to that of the liver were extracted from the images. Concomitantly, the kidney images were analyzed using the ResNet18 model for CKD detection. Finally, three diferent models, the image-only model, the model including extracted measurement data, and the model also including diabetes history information, were used for CKD detection. Through this systematic process, the accuracy of detecting CKD was signifcantly increased when computer-extracted measurable features were added to the machine learning models.

tures, and 0.91 in image training with extracted measurable features and diabetes information, respectively (**D**). AUROC, area under the receiver operating characteristic curve; ROC, receiver operating characteristic

Recently, machine learning algorithms have been applied to increase the accuracy and practicality of medical image analyses. [[15](#page-8-12)[–18\]](#page-8-15) Using CNN methods, the detection of breast cancer in screening mammograms revealed an AUC of 0.98 [\[37](#page-9-8)]. Functional magnetic resonance imaging (fMRI) data were analyzed using LeNet-5 to detect Alzheimer's disease, which resulted in a mean accuracy of 96.86% [[38](#page-9-9)]. Although employing machine learning algorithms to medical image data has resulted in high accuracy, the number of systems utilizing ultrasound images is limited compared to CT or MRI [\[39](#page-9-10), [40\]](#page-9-11)**.** Several investigations have achieved relatively high accuracy, such as the study that applied a threelayer deep belief network in time-intensive curves extracted from contrast-enhanced US (CEUS) video sequences to classify malignant and benign focal liver lesions, which revealed an accuracy of 86.36% [\[41](#page-9-12)]. However, most of the attempts at applying machine learning to ultrasound images have resulted in lower accuracies than other imaging modalities. In this study, the AUROC for CKD classifcation was signifcantly improved by the imputing of measurable feature data that were automatically extracted through a machine learning process in addition to the CNN-based analysis. This

improvement in accuracy shows that imputing supplementary data that could be obtained from the images themselves may compensate for the low accuracy of the CNN-based methods for evaluating ultrasound images.

The CKD-detecting machine learning process developed in this study features an end-to-end characteristic. Each step of segmentation of the organs, extraction of measurable features, and CNN-based classifcation proceeds automatically without the need for manual intervention. Machine learning methods were assigned to each of these steps. An overall CKD classifcation accuracy of 85.6% was recently obtained by training kidney ultrasound images through a ResNet model [[42](#page-9-13)]. However, in that study, the boundaries of the kidneys were cropped based on kidney length annotations that were manually performed by sonographers. Although the accuracy of CKD classifcation (85.9%) was comparable, it should be noted that a fully automated process from segmentation to classifcation was applied in the current investigation. Compared to previous machine learning–based classifcation systems, this end-to-end automated workfow would be a noteworthy advantage for practical application in the clinical feld.

The accessibility to medical ultrasound imaging is relatively higher than that of other imaging modalities, such as CT and MRI. In addition to the relatively lower cost associated with equipment installation, ultrasound machines require much less space and are portable $[6, 7]$ $[6, 7]$ $[6, 7]$ $[6, 7]$. These characteristics allow ultrasound to be the primary choice for organ imaging in developing countries and in regions that have poor medical access. However, the comparatively low image quality and inter-operator variability were the main limitations of ultrasound, which has prevented it from being more widely used $[10]$ $[10]$. In this regard, the presented automated machine learning–based CKD classifcation model may help overcome the shortcomings of ultrasound diagnosis. Considering the recent increase in CKD incidence in developing countries, the clinical usefulness of automated high-precision models is high. In addition, the strategy of combining measurable data with CNN to improve ultrasound classifcation accuracy could also be applied to diagnose other diseases of various organs. Since high-prevalence diseases such as liver diseases [\[41](#page-9-12)] and breast [[43\]](#page-9-14) and thyroid cancer [\[44](#page-9-15)] are initially screened through ultrasound imaging, methods for improving machine learning–based classifcation accuracy are imperative.

This study has several limitations. First, kidney ultrasound images with large cysts, solid masses, and hydronephrosis were not included in the analysis. The decision to exclude these disease states was made so that a working model solely focused on CKD classifcation could be developed. However, real-world ultrasound images would include various fndings, and a system capable of diferentiating the concomitant abnormalities would be needed for actual

clinical use. Second, in order to compare the images of normal eGFR patients with those with chronic kidney disease, images from patients whose eGFRs were between 60 and 90 were not collected. eGFR higher than 90 is considered normal kidney function, while the presence of chronic kidney disease is suspected when eGFR is lower than 60 [[23,](#page-8-20) 24]. This selection criterion would affect the real-world efficiency of the developed system. Nonetheless, this conceptual study shows that the integration of computer-extracted measurable features improves the classifcation accuracy of image machine learning models. Third, although a fully automated process was applied to classify the images for CKD detection, the ultrasound images were initially acquired by clinicians. This would have resulted in inter-operator variability, which could have introduced bias to the analysis. Forth, the fact that the validation sets used for segmentation and classifcation were not identical could result in overftting the training set on the segmentation task. This may lead to an overestimation of overall performance. In addition, since the pixel normalization for values of kidney to liver was performed per dataset rather than per-image basis, a possibility of data leak which may have resulted in performance augmentation should also be considered. Finally, the ultrasound images were not standardized because of the study's retrospective nature. Ultrasound images were uploaded at various angles, and the ultrasound machines used also varied. However, the variability of ultrasound machines may enable view-invariant properties, which may increase the practicality of real-world applications.

Conclusion

The automated step-wise machine learning–aided model successfully segmented the kidney and liver on ultrasound images. It was also effective at extracting measurable features, such as kidney size and echogenicity. The integration of computer-extracted measurable features into the machine learning model resulted in a signifcantly improved CKD classifcation. Further investigations are needed to apply this model in the real world.

Author Contribution S. L. and J. T. P. designed the study; S. L. obtained the patient's clinical information; Y. A. K. contributed to obtaining ultrasound images; M. K., K. B., and I. H. L. performed machine learning–aided analysis; and S. L. and J. T. P. analyzed the data; S. L. and M. K. made the fgures; S. L., M. K., and J. T. P. drafted the manuscript; and S. E. L., I. H. L., S. W. K., and J. T. P. revised the paper. All authors contributed important intellectual content during manuscript drafting or revision, and accepted accountability for the overall work by ensuring that questions pertaining to the accuracy or integrity of any portion of the work were appropriately investigated and resolved. And Sangmi Lee and Myeongkyun Kang contributed equally to this study as co-frst authors.

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Code Availability Not applicable.

Declarations

Ethics Approval This study was conducted in accordance with the Declaration of Helsinki and approved by the Institutional Review Board of Yonsei University Health System Clinical Trial Center (1–2018-0039).

Consent to Participate The need for informed consent was waived by the institutional review board owing to the study's retrospective design.

Conflict of Interest The authors declare no competing interests.

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