Real-time use of artificial intelligence in the evaluation of cancer in Barrett’s oesophagus

Alanna Ebigbo,1 Robert Mendel,2,3 Andreas Probst,1 Johannes Manzener,1 Friederike Prinz,1 Luis A de Souza Jr.,4 Joao Papa,5 Christoph Palm,2,3 Helmut Messmann1

MESSAGE
Based on previous work by our group with manual annotation of visible Barrett oesophagus (BE) cancer images, a real-time deep learning artificial intelligence (AI) system was developed. While an expert endoscopist conducts the endoscopic assessment of BE, our AI system captures random images from the real-time camera livestream and provides a global prediction (classification), as well as a dense prediction (segmentation) differentiating accurately between normal BE and early oesophageal adenocarcinoma (EAC). The AI system showed an accuracy of 89.9% on 14 cases with neoplastic BE.

IN MORE DETAIL
This paper follows up on our prior publication on the application of AI and deep learning in the evaluation of BE.1 2 In our initial publications, we developed a computer-aided diagnosis (CAD) model and demonstrated promising performance scores in the classification and segmentation domains during BE assessment.1 2 However, these results were achieved on optimal endoscopic images, which may not mirror the real-life situation sufficiently. To enable the seamless integration of AI-based image classification into the clinical workflow, our previous system was developed further to increase the speed of image analysis for classification and the resolution of the dense prediction, which shows the color-coded spatial distribution of cancer probabilities.3 4 Still based on deep convolutional neural nets (CNNs) and a residual net (ResNet) architecture with DeepLab V.3+, a state-of-the-art encoder–decoder network was adapted.1 2 To transfer the endoscopic livestream to our AI system, a capture card (Avermedia, Taiwan) was plugged to the endoscopic monitor.

Online supplementary video 1 shows the setting of AI-based BE evaluation in the endoscopy room of the University Hospital Augsburg (figure 1). The AI prediction can be started at any time using either a button on the keyboard or a foot switch. The video clip shows examples of three different patients and the AI predictions at different locations. Note that image freeze is not necessary for the AI analysis, which is randomly done by the system, but was done for evaluation purposes of this study to ensure the same basis for human and AI predictions. As can be seen, a dynamic blue bar at the bottom of the screen provides a continuous prediction of the probability of cancer in real time.

In conformity to our previous AI system,2 an image is classified as EAC if the global cancer prediction exceeds 90%.

Online supplementary video 2 shows the user interface (UI) and the application features. The left part of the UI consists of a preview window of the livestream of the endoscopic examination (or optionally a video) and a list of patients examined. The main window shows the probability-based dense prediction. A real-time partial prediction is shown by the dynamic blue bar at the bottom of the screen. Three different visualisations of the results are provided: the Barrett view, the cancer view and the combined view. Additionally, the global prediction of cancer for a particular region of interest is located as a thick blue bar on the lower part of the UI. Finally, online supplementary video 2 sketches briefly the AI method used for global and dense prediction (figure 2). The method is based on DeepLab V.3+, an encoder–decoder artificial neural network.2 Similar to our previous CAD system,2 the encoder is ImageNet-pretrained and based on a ResNet with 101 layers. It successively reduces the spatial resolution of the feature maps in each layer to learn semantic information at a higher abstraction level. The last ResNet layer uses the concept of dilated convolutions followed by a spatial pyramid pooling to allow the integration of larger contexts into the classification.5 For global prediction, two fully connected layers are linked to the encoder path.3 For the dense prediction, the decoder uses the semantic information of the encoder and distributes it to full spatial resolution. Therefore, the encoder neural network is used for classification (global prediction) as well as for segmentation (dense prediction).

In our present study, the AI system was trained using a total of 129 endoscopic images from our image database at the University Hospital Augsburg. To validate the system in real time, additional images (36 of early EAC and 26 of normal BE) from 14 patients (4 female and 10 male) were assessed by the AI system concurrently with the endoscopic examination by an expert BE endoscopist. All images were validated by pathological examination of resection specimens (EAC), as well as forceps biopsies (normal BE). The AI system had excellent performance scores in the classification task with a sensitivity and specificity of 83.7% and 100.0%, respectively, and an overall accuracy of 89.9%.

Find more details on Methods and Results online.

1Department of Gastroenterology, Universitätsklinikum Augsburg, Augsburg, Germany
2Regensburg Medical Image Computing (ReMIC), Ostbayerische Technische Hochschule Regensburg (OTH Regensburg), Regensburg, Germany
3Regensburg Center of Health Sciences and Technology, OTH Regensburg, Regensburg, Germany
4Department of Computing, Federal University of São Carlos, São Carlos, Brazil
5Department of Computing, São Paulo State University, Bauru, Brazil

Correspondence to
Dr Alanna Ebigbo, Department of Gastroenterology, Universitätsklinikum Augsburg, Augsburg 86156, Germany; alanna.ebigbo@uk-augsburg.de and Professor Christoph Palm, Ostbayerische Technische Hochschule Regensburg, Regensburg 93053, Germany; christoph.palm@oth-regensburg.de

AE and RM contributed equally.


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To our knowledge, this is the first real-time application of a deep learning AI system in the evaluation and diagnosis of early EAC in BE in a real-life setting. The video demonstrations show the difficulties encountered by expert and non-expert endoscopists during BE assessment. Since the AI system showed a very similar performance compared with the experienced endoscopist (the AI system and the expert disagreed in only one of the examined regions), the AI-based CAD has the potential to improve the quality of BE assessment, especially for non-expert endoscopists. Study limitations include the low number of patients included in the present study. Furthermore, elevated lesions delineated by the subjective impression of the endoscopist were included in our initial study. However, further optimisation with evaluations will follow.

**Contributors**

AE: study concept and design, drafting of the manuscript, analysis and interpretation of data, and critical revision of the manuscript. RM: study concept and design, software implementation, drafting of the manuscript, analysis and interpretation of data, and critical revision of the manuscript. AP: study concept and design, acquisition of data, drafting of the manuscript, critical revision of the manuscript and study supervision. JM: study concept and design, critical revision of the manuscript. P: study concept and design, acquisition of data and critical revision of the manuscript. JD: statistical analysis and critical revision of the manuscript. PK: statistical analysis; critical revision of the manuscript and study supervision. CP: study concept and design, drafting of the manuscript, analysis and interpretation of data, statistical analysis, critical revision of the manuscript, administrative and technical support, and study supervision. HM: study concept and design, acquisition of data, drafting of the manuscript, critical revision of the manuscript, administrative and technical support, and study supervision. AE and RM contributed equally.

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**REFERENCES**