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Hybrid deep learning models OPEN for the screening of Diabetic Macular Edema in optical coherence tomography volumes

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Several studies published so far used highly selective image datasets from unclear sources to train computer vision models and that may lead to overestimated results, while those studies conducted in real-life remain scarce. To avoid image selection bias, we stacked convolutional and recurrent neural networks (CNN-RNN) to analyze complete optical coherence tomography (OCT) cubes in a row and predict diabetic macular edema (DME), in a real-world diabetic retinopathy screening program. A retrospective cohort study was carried out. Throughout 4-years, 5314 OCT cubes from 4408 subjects who attended to the diabetic retinopathy (DR) screening program were included. We arranged twentytwo (22) pre-trained CNNs in parallel with a bidirectional RNN layer stacked at the bottom, allowing the model to make a prediction for the whole OCT cube. The staf of retina experts built a ground truth of DME later used to train a set of these CNN-RNN models with diferent confgurations. For each trained CNN-RNN model, we performed threshold tuning to fnd the optimal cut-of point for binary classifcation of DME. Finally, the best models were selected according to sensitivity, specifcity, and area under the receiver operating characteristics curve (AUROC) with their 95% confdence intervals (95%CI). An ensemble of the best models was also explored. 5188 cubes were non-DME and 126 were DME. Three models achieved an AUROC of 0.94. Among these, sensitivity, and specifcity (95%CI) ranged from 84.1–90.5 and 89.7–93.3, respectively, at threshold 1, from 89.7–92.1 and 80–83.1 at threshold 2, and from 80.2–81 and 93.8–97, at threshold 3. The ensemble model improved these results, and lower specifcity was observed among subjects with sight-threatening DR. Analysis by age, gender, or grade of DME did not vary the performance of the models. CNN-RNN models showed high diagnostic accuracy for detecting DME in a real-world setting. This engine allowed us to detect extra-foveal DMEs commonly overlooked in other studies, and showed potential for application as the frst flter of non-referable patients in an outpatient center within a population-based DR screening program, otherwise ended up in specialized care.

Keywords Deep learning, Diabetic Macular Edema, Diabetic Retinopathy, Optical Coherence Tomography, Screening, Telemedicine

Abbreviations

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Diabetes mellitus (DM), specially type II, remains a challenge for public health as the incidence continues to rise^{1,[2](#page-8-1)}, and with it the burden of associated comorbidities as diabetic eye disease^{[3](#page-8-2)}. Among the latter, diabetic macular edema (DME) is the most prevalent sight-threatening (ST) condition although treatable while macu-lar changes are not yet permanent^{[4](#page-8-3)}. Several countries have implemented successful diabetic retinopathy (DR) screening programs, mainly based in fundus retinography (FR), that lowered the overall incidence of blindness⁵. Nevertheless, recent reports advocated for the potential beneft of adding optical coherence tomography (OCT) to FR, as it may detect indirect early DME signs often missed by FR in a non-negligible proportion,⁶ improving the cost-effectiveness of screening models⁷. As a counterpart, the deployment of such program faces new resourcing challenges that could jeopardize the sustainability of any public health system; adding OCT to FR raises the instrumentation costs, and highly skilled specialists are needed to read the OCT cubes to achieve the best results^{6[,8](#page-8-7)}.

To address some of these challenges, artifcial intelligence, especially deep convolutional neural networks (CNNs), have been widely tested for the automated classifcation of macular pathologies in FR and/or OCT scans, as well as to assist the retina specialists to speed-up their decision-making process^{[9](#page-8-8)}. In this sense, several studies published using CNNs for the classifcation of DME showed specialist-level or even outperformed the results achieved by retina specialists, although their generalizability raises important concerns^{[10,](#page-8-9)11}. In some studies, authors used 2D-OCT scans (B-scans) and that introduced a selection bias as training images must be previously selected from the OCT cube^{12,13}. Moreover, the majority used central fovea B-scans that could potentially miss both non-foveal involving anatomical signs and DMEs^{[10](#page-8-9)-13}. Some authors relied in FR alone to detect DME, thus based on indirect signs that likely indicate progression which in turn may be easier to detect by the models^{10[,14](#page-8-13)}. Furthermore, in most of them the authors did not state the criteria followed to assess DME^{[10](#page-8-9)}. In consequence, most of the datasets used in these studies may not represent the actual distribution of the disease in the general population, so that may lead to overestimated performances and poor generalizability in turn.

Recurrent neural networks (RNN) are deep learning models suitable to analyze time-series data and could be stacked at the output of several CNNs to predict time-dependent outcomes in images[15](#page-9-0),[16](#page-9-1). Hence, such hybrid model (CNN-RNN) applied to OCT B-scans would allow to analyze the OCT cube in a row and sequentially in time.

Thus, the aim of the present study was to build hybrid deep learning models (CNN-RNN) with different settings and trained with an *"*ad hoc*"* ground truth of DME, then to test them for the detection of DME in an unselected dataset from a real-world DR screening program.

Methods

Model development and training with a ground truth of DME

A detailed description of all models settings, training and testing, as well as image processing was shown in Supplementary Material 1. Briefy, we pre-trained a backbone custom CNN using a publicly available dataset comprised of B-scans of DME, drusen, choroidal neovascularization, and normal macula (Supplementary Material $1.1-1.2$ ¹². Then, twenty-two (22) of these pre-trained CNNs were stacked in parallel, initialized with the pre-training weights, and fed with 22 unselected B-scans extracted from the OCT cube. Researchers did not intervene in the selection of images for training or testing, so that avoided image selection bias. To this end, we divided each 128-slice cube into those that mainly comprised the foveal zone and those which mainly comprised the parafoveal zone. The foveal zone was captured between slices 60 to 85 from where we automatically extracted every two (12 B-scans), being the remaining the parafoveal zone from where we automatically extracted every ten (10 B-scans; Supplementary Material 1.3). Every pre-trained CNN outputs an embedding of image features from the fatten layer. Ten, all were concatenated into a sequence which was forwarded to the bidirectional recurrent layer. The bidirectional wrapper moves a cell forward and another backwards along the sequence to learn dependencies between time-dependent features. Finally, the output from the RNN layer was fully-connected to a sigmoid layer to predict the probability of DME for the 22-slice OCT cube (Supplementary Fig. 1).

We used a ground truth of DME and normal macula, described elsewhere^{[6](#page-8-5)} but enriched with additional OCT cubes from a second DR screening program (Hospital Clínic of Barcelona, Spain). Additional samples were graded following the same criteria^{[6](#page-8-5)}. All images were acquired using the Topcon 3D OCT-Maestro 1. The ground truth dataset was split into training, validation, and test, ensuring a similar proportion of DME between folds, and avoiding data leakage by creating splits of unique subjects. The binary cross-entropy was used as the loss function, although accuracy, the area under the receiver operating characteristic (ROC) curve (AUROC),

and area under the precision-recall curve (AUPRC) were also computed and compared. From the pool of trained models, we selected those with the best generalizable metrics in the test set.

Study cohort and data collection

A retrospective cohort study nested in a teleophthalmology real-world DR screening program was conducted. From November 2015 to March 2019, we included all diabetic patients (either type), of any gender, and aged≥18 years. We included one eye per subject, which was the afected in case of unilateral DME, or a random sample if both eyes had the same diagnosis (DME or non-DME).

The characteristics of the screening program were described elsewhere^{[6](#page-8-5)}. In short, screening visits were conducted by a technician in an outpatient center and collected health data of interest, measured the best-corrected visual acuity (BCVA), and acquired a 3-feld FR (frst centered on the macula, second on the disc, and third supero-temporal)¹⁷, and a 6 \times 6 mm OCT macular cube scan. Patients were re-imaged under pupil dilation in case of low quality of images. Then, the technician forwarded all the abovementioned data, also including the ETDRS average thicknesses, the ETDRS topographic map, and the macular volume, to a retina specialist in the Hospital who acts as the gatekeeper to specialized care. There, the retina specialist makes the initial diagnosis, and decides whether to refer the patient. The retina specialist assessed DME based on the presence of macular thickening≥300 µm with anatomical signs of DME (cysts, microaneurysms, exudates, neurosensory detach-ment, and hyperreflective dots), without signs of another macular disease in the FR^{18[,19](#page-9-4)}. The severity of DME was also assessed depending on the distance to central fovea as proposed by the International Clinical Diabetic Retinopathy and Diabetic Macular Edema Disease Severity Scales (ICDRSS), into mild, moderate and severe²⁰. Subjects whose OCT and FR were missing or ungradable were excluded from the study cohort.

Model evaluation in the study cohort

All 22-slice OCT cubes from the study cohort were extracted and pre-processed, then fed to the CNN-RNN models to predict the probability of DME. Model predictions were evaluated against the diagnosis made by the retina specialist in the screening program, as described above. ROC curves, AUROC, and the partial AUROC (pAUROC) at a range of false positive rate (FPR) < 0.05 and < 0.1, along with their 95% confdence intervals (95%CI) were computed. Sensitivity, specifcity, positive predictive value (PPV), negative predictive value (NPV), and the cumulative incidence of DME were calculated at a range of 1000 prospectively tuned thresholds set to predict the best parameters for screening: (1) Youden index (threshold 1), (2) the highest sensitivity (threshold 2), and (3) the highest specifcity (threshold 3). For the latter two, we established a baseline specifcity and sensitivity of at least 0.8 or greater. Only the models that yielded the best AUROC with a pAUROC over 0.80 were considered onwards. The selected models were ensembled to build a voting classifier based on the mode of class predictions obtained at each model threshold.

Additionally, the characteristics of false positives and false negatives obtained were reviewed by a third retina specialist.

Finally, as an exploratory analysis, we tested the potential generalizability of the models for classifcation of referable DR. In this case, we graded DR as mild non-proliferative, moderate non-proliferative, and referable DR, which included severe non-proliferative and proliferative plus moderate and severe DME^{20} DME^{20} DME^{20} .

Statistical analysis

Baseline characteristics of the population were expressed as median and interquartile range (IQR) for quantitative variables, and as frequencies and percentages for qualitative variables. Diferences between two medians were tested using the Mann–Whitney's U, and the test on the equality of proportions was used to compare two qualitative variables. A p-value < 0.05 was set as statistically significant.

ROC curves and AUROC (95%CI) were computed to test the model performance as it is independent of the prevalence of the disease. Partial AUROC (pAUROC; 95%CI) was also computed to obtain more nuanced models²¹. Youden (J) index was computed to represent the maximum sensitivity and specificity given for a single point on the ROC curve. Diagnostic accuracy for binary outcomes was assessed using the sensitivity, specifcity, PPV, and NPV, afer probability thresholding. Diagnostic accuracy was also stratifed by gender, age, laterality, BCVA, grade of DR, OCT quality of image, and grade of DME. For the latter, due to low numbers, we collapsed moderate and severe. Incidence of DME predicted by the models was calculated as the number of true positives divided by the total number of subjects. Intervals at 95% of confdence (95%CI) were calculated using the standard normal distribution or the binomial distribution for proportions²².

To test the assumption of no missing cases due to lower resolution of images (22-slice cubes) we carried out a sensitivity analysis by severity of DME, as well as by other covariates as grade of DR, BCVA, quality of image, among others.

Models were developed in GPU-enabled Tensorfow v.2.4, and diagnostic accuracy was computed with Scikitlearn v. 1.2.2, for Python. The remaining analyses were run with STATA/MP v.17 (Stata Corp LLC, College Station, TX, USA).

Ethics approval and consent to participate

The study protocol was approved by the Ethics Committee of the University Hospital "Príncipe de Asturias" on March 2, 2020. The need for informed consent was waived due to the retrospective nature of the study. This study complied with the provisions of Spanish and European laws on personal data as well as with the Declaration of Helsinki (Fortaleza 2013).

Results

Baseline and clinical characteristics of the study cohort

The characteristics of the study population were detailed in Table [1](#page-4-0). We included 5314 screens from 4408 subjects. Of them, 126 (2.37%) were DMEs, and 5188 (97.6%) had no record of DME throughout the study period. The majority of DMEs were mild (77; 61.1%), while moderate and severe contributed equally (23; 18.3%, each). Compared to subjects without, those with DME had fewer screening visits, were older, predominantly males, with longer duration of DM, and had a greater prevalence of acute myocardial infarction (Table [1](#page-4-0)). Regarding vision, subjects with DME had lower BCVA, greater prevalence of moderate and severe non-proliferative DR, as well as proliferative DR, and had greater proportion of epiretinal membranes, pupil dilation or previous macular scars. In OCT, subjects with DME showed greater average macular thicknesses and volume across the ETDRS grid (Table [1\)](#page-4-0).

Training of hybrid models with a ground truth dataset of DME

The ground truth dataset comprised 650 OCT cubes from unique subjects, being 433 (66.6%) of them in the training set (127 -29.3%- DMEs, and 306 -70.7%- non-DMEs), 100 (15.4%) in the validation set (20 -20%- DMEs, and 80 -80%- non-DMEs), and 117 (18%) in the test set (40 -34.2%- DMEs, and 77 -65.8%- non-DMEs). Seven (7) diferent hybrid models yielded the best predictive and generalizable results in the test set (Supplementary material 1.4–1.6).

Model selection and diagnostic accuracy in the study cohort

The abovementioned models were then tested in the study cohort and three showed the best performance. Their architecture characteristics and results were detailed in Supplementary material 1.5–1.6. AUROC (95%CI) was 0.94 (0.92–0.97) for model 1, 0.94 (0.91–0.97) for model 2, and 0.94 (0.91–0.97) for model 3. For the region at FPR < 0.05, pAUROC (95%CI) was 0.83 (0.79-0.87), 0.84 (0.80-0.89), and 0.81 (0.76-0.85) for models 1, 2, and 3, respectively, while at FPR<0.10, pAUROC (95%CI) was 0.87 (0.83–0.91), 0.88 (0.84–0.92), and 0.85 (0.81–0.89) for models 1, 2, and 3, respectively (Fig. [1](#page-5-0)).

Sensitivity and specifcity were calculated at the three thresholds. Model 2 showed the most balanced results at threshold 1: sensitivity (95%CI) was 90.5 (84.0–95.0), and specifcity (95%CI) was 90.8 (90.0–91.6). Similar results but with higher specifcity was reached by the ensemble model: sensitivity (95%CI) was 88.9 (82.1–93.8), and specificity (95%CI) was 93.3 (92.6–94.0). The ensemble model showed the best results regarding sensitivity and specifcity at the remaining thresholds: sensitivity (95%CI) was 92.1 (82.1–93.8), and specifcity (95%CI) was 83.1 (82.1–84.1) at threshold 2, while sensitivity (95%CI) was 81.0 (73.0–87.4), and specifcity (95%CI) was 97.0 (96.5–97.4) at threshold 3 (Table [2\)](#page-5-1).

Negative predictive values (NPV; 95%CI) were high and similar in all scenarios (above 99.5). As expected, positive predictive values (PPV; 95%CI) were low (as it drastically depends on the prevalence of the disease but increases with higher specifcity); with the ensemble model at threshold 3, PPV was 39.5 (33.5–45.8). Cumulative incidence of DME for the study period was similar across all thresholds and for all models (Table [2](#page-5-1)).

Diagnostic accuracy of the ensemble model by diferent covariates

The stratified analysis showed a similar diagnostic accuracy across all strata, excepting for moderate and severe DME which resulted in higher sensitivity and specifcity compared to mild DME and overall; the highest sensitivity (95%CI) was 100 (92.9–100) at threshold 2, and the highest specifcity was reached at threshold 3 (specifcity; 95%CI = 97.0; 96.5–97.4). In contrast, the specifcity observed among subjects with ST-DR (including severe non-proliferative and proliferative) was lower as compared to overall; at threshold 3 it was 62.9 (44.9–78.5), and even lower at thresholds 1 and 2 (Table [3\)](#page-6-0).

Characteristics of false positives and false negatives

The characteristics of false positives and negatives obtained with the ensemble model at threshold 3 were explored. In this confguration, we obtained 24 false negatives and 156 false positives. According to the review made by the third retina specialist, 87.8% of all false negatives predicted by the model were correctly classifed indeed. They were mainly characterized by epiretinal membranes, macular thickening without signs of DME, and anatomical signs of DME without macular thickening, and just 4 DMEs (12.1%) were missed by the model. On the other hand, anatomical signs of DME without macular thickening, images with artifacts (eye blinking, cropped images) or skewed, and age-related macular degeneration accounted for 54% of false positives, and 17 (10%) DMEs were misdiagnosed by the retina specialist, thus correctly classifed by the model (Fig. [2\)](#page-7-0).

Generalizability of models for classifcation of referable DR

The three selected models yielded almost chance prediction in the classification of mild non-proliferative DR. Conversely, an increasing trend to a high performance was observed with severity. Overall, AUROC for referable DR was equal or above 0.89 (Fig. [3](#page-7-1)).

Discussion

Results from the present study showed that our hybrid (CNN-RNN) deep learning models achieved high diagnostic accuracy in the identifcation of DME, overall and by severity, in a dataset of OCT cubes generated in a real-world DR screening program as a product of routine clinical care.

Computer vision models based on CNNs have been widely tested for the detection of DR or DME so far²³. Nevertheless, very few have been designed for direct clinical application in real-life, with prospectively collected

Table 1. Baseline characteristics of the study cohort. *DME* diabetic macular edema, *IQR* interquartile range, *CV* cardiovascular, *BCVA* best corrected visual acuity, *OCT* optical coherence tomography, *CST* central subfeld thickness. *Includes proliferative and photocoagulated proliferative DR.

Figure 1. AUROC and pAUROC (95%CI) for the classifcation of DME, with the best three models. ROC: receiver operating characteristics; AUROC: area under the ROC curve; DME: diabetic macular edema. Vertical dotted and colored lines represented pAUROC at FPR (1-specificity) < 0.05 and FPR < 0.10.

Table 2. Diagnostic accuracy of the three best models after threshold tuning. N_{DME} : 126; $N_{non\text{-DME}}$: 5188. *DME* diabetic macular edema, *CI* confdence interval, *PPV* positive predictive value, *NPV* negative predictive value. *Treshold 1: Youden index. **†**Treshold 2: highest sensitivity. **‡**Treshold 3: highest specifcity.

datasets, and well-defined diseases^{10[,13](#page-8-12)[,14](#page-8-13),24}. CNNs require significant amounts of manually expert-labeled data for training and validation. At present, some publicly available datasets comprised of OCT B-scans or FR have been used in numerous studies, however, using flat images as those implies selective sampling^{10,12}. The screening of DR in FR alone leads to low sensitivity in the diagnosis of DME, as the criteria to assess DME is based on the presence of indirect signs that are indeed markers of progression, and a non-negligible proportion of false positives and negatives (almost 50% of DME without indirect signs are missed)⁶ Likewise, selective sampling also occurs in B-scans as they must be previously selected or, in other cases, only central fovea B-scans were extracted from the OCT cube^{10,12}. For these reasons, DMEs with clear signs of the disease are overrepresented in existing datasets, and may introduce a significant bias with misleadingly outstanding results¹⁰. Otherwise, when transported to real-world datasets, much more realistic results have been reported^{[13](#page-8-12),[14](#page-8-13),[24](#page-9-9)-[26](#page-9-10)}.

To our knowledge, this is the frst study that applied hybrid (CNN-RNN) models to analyze OCT cubes with the aim to detect DMEs of either grade. Hybrid models have been proposed to predict time-dependent outcomes as treatment response in colorectal and lung cancer^{15[,16](#page-9-1)}, although we took advantage of this feature to analyze OCT cubes with just one expert annotation per volume, and also to avoid misalignments when re-assembling the cube if 3D-CNNs were used^{[11](#page-8-10)}. Furthermore, another potential benefit of our approach is the management of time, making our models transferable to predict time-to-event outcomes as treatment response (e.g. vascular

Table 3. Sensitivity and specificity of the ensemble model at all thresholds, by different covariates. N_{DME} : 126; *Nnon-*DME: 5188. *DME* diabetic macular edema, *CI* confdence interval, *BCVA* best corrected visual acuity, *ST* sight-threatening, *OCT* optical coherence tomography. *ST-DR included severe non-proliferative and proliferative DR.

endothelial growth factor inhibitors) or risk of progression. At present, other authors have also developed a model for segmentation and identifcation of several referable retinal diseases using OCT cubes, so that precluded the comparability with our results 27 27 27 .

Regarding the screening of DME, Liu et al*.* [13](#page-8-12) carried out a study nested in a real-word DR screening program and reported a sensitivity, specifcity, and AUROC of 91.3, 97.5, and 0.944, respectively, using a combination of FR and OCT. Taking the Youden index as reference, our ensemble model yielded a sensitivity (95%CI) and specifcity (95CI%) of 88.9 (82.1–93.8) and 93.3 (92.6–94.0), while AUROC for the best three models was 0.94. Our results were slightly lower, but some considerations should be made to adequately compare the results: Liu et al. used B-scans and FR but without assessment by grade of DME[13](#page-8-12). Of note, using a modest population of 600 diabetics, they reported a PPV of 75.4% strongly infuenced by a high prevalence of DME of almost fourfold the observed in our population. In practical terms, since PPV depends on prevalence, with the sensitivity and specifcity reported by them we would barely reach a PPV of 50% in our population, and suggesting PPV alone as inadequate metric for screening in imbalanced datasets. Moreover, baseline and clinical characteristics of the study population were not reported, so that prevented an extensive interpretation of results 13 .

Tis pioneer study evaluated the ability of the models to detect diferent grades of DME, and showed no diferences in performance to detect mild DMEs, but outperformed the overall results in moderate and severe DMEs, so this trend may support, in part, our concerns about the misleading results observed in studies using selected datasets^{[10](#page-8-9),[28](#page-9-12)}. It is important to stress that, in our population, the majority of DMEs were mild (61.1%) as a consequence of a well-established screening program where patients are detected earlier^{[6](#page-8-5)}. Nevertheless, is also relevant to detect early DMEs to personalize screening visits, adjust antidiabetic treatments, and manage other cardiovascular risk factors, to improve their visual prognosis $29,30$ $29,30$ $29,30$.

The design of the best strategy for DME screening in the community must consider multiple dimensions, as well as the impact of potential misdiagnoses, the number of false positives that specialized care could undertake or the cost-efectiveness, among others. Population-based screening of a disease with such low prevalence would benefit most from higher specificity as increases PPV to a greater extent than higher sensitivity 31 , and would rule out most subjects screened in a frst step, reducing drastically the workload of the specialist. Based on our

Figure 3. ROC and AUROC (95%CI) for different grades of diabetic retinopathy with the best three models. ROC: receiver operating characteristics; AUROC: area under the ROC curve; DR: diabetic retinopathy; DME: diabetic macular edema. * Referable DR included severe non-proliferative DR, proliferative DR, moderate DME, and severe DM.

results, that would cost 3–6% (156–311 subjects) false positives, and almost 20% false negatives (24–25 DMEs missed), throughout the study period. Conversely, screening based on sensitivity would result more efficient although highly challenging in turn since advanced stages of DR share indirect markers with DME and both entities ofen concur. In this scenario, we would miss 7.9% (10) DMEs at the cost of 16.9% (877) false positives, at best. Alternatively, the Youden index would return more balanced metrics overall; 7–9% false positives, and 9–11% false negatives. Of note, afer exploring the characteristics of false negatives, only 4 DMEs were incorrectly predicted by the model, so re-calculated sensitivity would rise to 94% with specifcity unchanged. On the other hand, the characteristics of false positives revealed that the models may learn anatomical signs of DME that could be shared with ST-DR, explaining the low specifcity of the models among these subjects. Regarding the latter, our models showed potential transferability to referable DR, although specifc training would be needed specially for mild and moderate stages³².

Our study has strengths and limitations that need to be discussed. We trained our models based on a ground truth of DME constructed by retinal experts with high expertise and low inter-individual variability, and then evaluated the models within a well-defned population extracted from the target population where these models could be further applied. In addition, our models were able to analyze OCT cubes, avoiding selective sampling and allowing to detect extra-foveal DMEs, which in turn were the majority in our population and ofen missed due to subclinical presentations. Finally, the ensemble model improved the results, but its complexity must be limited to be deployed efectively within a teleophthalmology screening program. By contrast, an external validation of the models, using datasets from other imaging sources and populations would be necessary. Despite we achieved good results indeed, we did not analyze the 128 B-scans from the OCT cube, so further approaches to build more complete samples would be needed. In this line, the combination of image-related data as thicknesses from the ETDRS grid may reduce the number of cases where anatomical signs were present but without reaching 300 µm. Finally, artifcial intelligence is usually considered as black box models due to their complexity and lack of interpretability, and that reduces the trust of specialists in their predictions³³. For this reason, medical researchers in this feld must make eforts towards explainable artifcial intelligence that support retina specialists to make meaningful clinical decisions.

Conclusions

Our model was able to detect DME in complete OCT cubes from a real-world screening program, making unnecessary to select training images and avoiding selection bias. Tis approach showed potential applicability to discard in a frst step the majority of non-DMEs otherwise referred to specialized care for re-evaluation, then reducing the workload and needs of retina specialists, then speeding-up the access to the teleophthalmology screening program as a consequence.

Data availability

In line with Spanish and European laws, data share must be authorized by the Ethics Committee of the University Hospital "Príncipe de Asturias", so authors are not allowed to openly make them available. However, data could be available by reasonable request from any organization or researcher to the corresponding author, provided that the Ethics Committee authorize specifcally the data transfer.

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ARM: Conceptualization, Methodology, Sofware, Formal Analysis, Investigation, Resources, Data Curation, Writing-Original Draf, Visualization, Supervision, Project Administration, Funding Acquisition; CA: Conceptualization, Validation, Resources, Data Curation, Writing-Review & Editing, Supervision, Project Administration, Funding Acquisition; GA: Validation, Data Curation, Writing-Review & Editing; MO: Writing-Review & Editing; JZV: Writing-Review & Editing; MAT: Resources, Writing-Review & Editing.

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Additional information

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