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# A digital biomarker of diabetes from smartphone-based vascular signals

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#### **Abstract**

The global burden of diabetes is rapidly increasing, from 451 million people in 2019 to 693 million by 2045<sup>1</sup>. The insidious onset of type 2 diabetes delays diagnosis and increases morbidity<sup>2</sup>. Given the multifactorial vascular effects of diabetes, we hypothesized that smartphone-based photoplethysmography (PPG) could provide a widely-accessible digital biomarker for diabetes. Here, we developed a deep neural network (DNN) to detect prevalent diabetes using smartphone-based PPG from an initial cohort of 53,870 individuals (the "Primary Cohort"), which was then validated in a separate cohort of 7,806 individuals (the "Contemporary

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Cohort"), and a cohort of 181 prospectively-enrolled individuals from three clinics (the "Clinic Cohort"). The DNN achieved an area under the curve (AUC) for prevalent diabetes of 0.766 in the Primary Cohort (95% confidence interval (CI): 0.750–0.782; sensitivity 75%, specificity 65%) and 0.740 in the Contemporary Cohort (95% CI: 0.723–0.758; sensitivity 81%, specificity 54%). When the output of the DNN, called the DNN Score, was included in a regression analysis alongside age, gender, race/ethnicity, and body mass index, the AUC was 0.830 and the DNN Score remained independently predictive of diabetes. The performance of the DNN in the Clinic Cohort was similar to that in other validation datasets. There was a significant and positive association between the continuous DNN Score and hemoglobin A1c (HbA1c) (*p* 0.001) among those with HbA1c. These findings demonstrate that smartphone-based PPG provides a readily attainable, noninvasive digital biomarker of prevalent diabetes.

#### **Keywords**

Diabetes mellitus; Machine learning; deep neural network (DNN); artificial intelligence; remote sensors; smartphone; photoplethysmography (PPG)

Globally, half of all people living with diabetes are undiagnosed (~224 million), and 79% live in low and middle-income countries<sup>1</sup>. Diabetes causes both macrovascular and microvascular multi-organ disease, including coronary heart disease, stroke, neuropathy and kidney disease, among others<sup>3</sup>. A readily attainable, non-invasive digital biomarker of diabetes could facilitate disease detection by making it easier to identify at-risk individuals who would benefit from confirmatory diagnostic testing using HbA1c. Such a tool would have particular impact in underserved populations and those out of reach from traditional medical care.

PPG is a noninvasive optical technique which detects blood flow changes through a vascular bed<sup>4</sup>. It is measured by shining light into tissue, such as the fingertip or wrist, and quantifying the backscattered light that corresponds with changes in blood volume<sup>4</sup>. PPG has long been used clinically to measure heart rate (HR) and peripheral blood oxygen saturation<sup>4</sup>, and research applications have ranged from detection of hypertension<sup>5</sup> to various cardiovascular abnormalities<sup>6,7</sup>. Until recently, PPG recording required specialized equipment, however technological developments have enabled PPG measurement from sensors on smart-devices, like smartphones and fitness trackers. The rapid worldwide adoption of smart-devices over the past decade<sup>8</sup> provides an opportunity to develop non-invasive, widely-scalable digital biomarkers for diseases like diabetes<sup>9</sup>.

PPG is uniquely positioned to capture the multifactorial sequelae of diabetes resulting from a variety of pathophysiologic mechanisms. PPG readily captures sequential heartbeats, enabling not only its longstanding use for HR measurement, but also heart rate variability, which is impacted by diabetic autonomic and neural-regulatory effects<sup>10-12</sup>. Recently, a shared genetic etiology between resting HR and diabetes was identified, implicating mechanisms ranging from metabolism to endothelial aging<sup>13</sup>. Indeed, endothelial dysfunction is an early hallmark of diabetic vascular disease, and is readily detectable in the PPG waveform<sup>14</sup>. Similarly, diabetes-related microvascular arteriosclerosis<sup>6,15</sup> and neuropathy can affect PPG<sup>16</sup>. Given the multitude of mechanisms by which diabetes impacts

PPG, algorithmic analysis of PPG should ideally leverage the complete PPG recording and all the morphologic and temporal information contained therein. DNNs are a class of algorithms<sup>17</sup> which have successfully achieved complex pattern recognition for various medical tasks<sup>18-20</sup>. DNNs provide the advantage of being agnostic to specific sets of predetermined PPG features suspected to predict diabetes, and instead detect patterns using the full PPG record. We therefore hypothesized that PPG obtained from commercially available smartphones and analyzed using a DNN could identify individuals with and without diabetes.

In this study, we first developed and validated a DNN to detect prevalent diabetes in a "Primary Cohort," comprised of 53,870 Health eHeart study<sup>21</sup> participants who contributed 2,589,448 PPG recordings between April 1, 2014 and April 30, 2018 (Figure 1a). Participants self-reported diabetes status and measured PPG by placing an index fingertip on the smartphone camera using the Azumio Instant Heart Rate iOS application (Azumio, Inc; Figure 1b). The Primary Cohort was randomly split into training (70%, n=37,709) and development (10%, n=4,848) datasets—used to train and tune the DNN, respectively—and a Test Dataset (20%, n=11,313), used for DNN validation. The DNN outputs a "DNN Score" between 0 and 1, with higher scores suggesting greater likelihood of diabetes (see Methods). Since many participants contributed >1 recording, we reported DNN performance using AUC<sup>22</sup> at both the 'recording-level,' which treats each recording independently, and the 'user-level,' which averages the DNN Score for all recordings provided by a user; user-level assessment was preferred when possible since clinical application calls for classifying a user as having diabetes or not.

In the Primary Cohort, 3,564 participants (6.6%) had self-reported diabetes and 50,306 (93.3%) did not (Extended Data Figures 1-2). Compared to those without diabetes, those with diabetes were older, male, had higher HR and body mass index (BMI), and were less likely to be non-Hispanic white. In the hold-out Test Dataset, the DNN's AUC to detect diabetes was 0.766 at the user-level (95% CI: 0.750–0.782; recording-level AUC=0.680, 95% CI: 0.678–0.683; Table 1 and Figure 2a). At the chosen cut-off threshold (DNN Score=0.427), user-level sensitivity was 75% and specificity was 65%. Due in part to the low prevalence of diabetes in our cohort (6.6%), the positive predictive value (PPV) of the DNN Score at the user-level and recording-level was 13% and 10%, while the negative predictive value (NPV) was 97% and 96%, respectively (Table 1). DNN performance in the development dataset was not significantly different from the Test Dataset (user-level AUC=0.766, 95% CI: 0.740–0.792; recording-level AUC=0.694, 95% CI: 0.691–0.698).

In addition to validating DNN performance in the Primary Cohort Test Dataset, we employed 2 additional validation cohorts (Figure 1a), providing 3 total examples of algorithm generalizability to datasets distinct from the training dataset<sup>23</sup>. The first was the "Contemporary Cohort," composed of PPG recordings from 7,806 participants newly-enrolled into Health eHeart from May 1 to December 31, 2018. This temporally-distinct validation cohort exhibits the DNN's robustness to secular changes, such as new smartphone models and cameras, that could affect PPG recording. Then, to validate our approach in a real-world clinical setting, we prospectively enrolled an in-person "Clinic Cohort" comprised of 181 consecutive patients referred to 3 cardiovascular prevention clinics

(2 in San Francisco, one in Montreal) between November 1, 2018 and July 30, 2019 (Figure 1a, Extended Data Figure 3). The DNN's user-level AUC to detect diabetes in the Contemporary Cohort was similar to that in the Primary Cohort: 0.740 (95% CI: 0.723–0.758; recording-level AUC=0.661, 95% CI: 0.664–0.667); the DNN had higher sensitivity, but lower specificity, versus the Primary Cohort (Table 1).

In the prospectively enrolled in-person Clinic Cohort, 38 patients (21.0%) had medical record-confirmed diabetes (Extended Data Figure 3). Compared with the Primary Cohort, the Clinic Cohort was substantially older, more male, and had more comorbidity. The Clinic Cohort recording-level AUC (0.682, 95% CI 0.605–0.755) was similar to the recording-level AUC in the Test Dataset and Contemporary Cohorts (Table 1). Compared with the Test Dataset, there was higher sensitivity and PPV, but lower specificity and NPV. When Clinic Cohort patients with a prior diabetes diagnosis were excluded (n=17), 21 patients remained who were newly diagnosed by HbA1c during the clinic visit. In this subset of patients with newly diagnosed diabetes, the DNN AUC was 0.644 (95% CI: 0.546–0.744; Table 1); the DNN correctly identified 16 out of 21 patients with newly diagnosed diabetes (Extended Data Figure 4f).

To investigate whether PPG was predictive of diabetes independently of other predictors and comorbidities, we built nested logistic regression (LogReg) models in the Test Dataset with and without the inclusion of the DNN Score (Table 2). After adjustment for age, gender, race and BMI, the DNN Score remained independently and significantly predictive of prevalent diabetes (Table 2, Supplemental Table 1); the AUC for this prediction model was 0.830 (95% CI: 0.787–0.873; Figure 2a). The DNN Score was also strongly predictive of diabetes independently of all examined comorbidities, including hypertension, hypercholesterolemia and coronary artery disease, among others (Table 2; LogReg-Model 5); the AUC for this prediction model was 0.830 (95% CI: 0.815–0.844; Figure 2a). In all models, the DNN Score was a strong diabetes predictor and was only slightly attenuated after adjustment (Table 2, Supplemental Table 1). Heart rate variability was no longer a significant predictor of diabetes after the DNN Score was added, while HR was attenuated (Table 2, LogReg-Model 4). Compared to participants with a DNN Score below the cutoff (<0.427), those with a DNN Score above the cutoff differed demographically and were nearly twice as likely to have any medical condition (69.4% vs 37.3%; p<0.001; Supplemental Table 2).

We performed several sensitivity analyses for hypertension specifically, since it is co-morbid with diabetes and may directly cause PPG-measurable vascular changes. A subset of Test Dataset participants provided Bluetooth-linked, home-measured blood pressures within 3 months of a PPG recording, totaling 13,007 PPG-blood pressure recording pairs (55 patients with diabetes, 527 patients without diabetes). Though the systolic (but not diastolic) value was a significant univariate predictor of diabetes, after the DNN Score and other (non-hypertension) comorbidities were added into a multivariable model, systolic blood pressure was no longer a significant diabetes predictor; the DNN Score, however, remained strongly independent (odds-ratio: 3.53, 95% CI: 2.20–5.67; p<0.001). Furthermore, after excluding those with self-reported hypertension from the Test Dataset, DNN performance remained similar to that in the full Test Dataset at both user and recording-levels.

Due to the limitations of relying on self-reported diabetes in our primary analysis, we performed additional sensitivity analyses aimed at addressing this. We identified Health eHeart participants who had laboratory-confirmed diabetes based on fasting glucose or HbA1c drawn within 180 days of diabetes self-report (n=12,073). In this subset, the PPV of self-reported diabetes was 81.8% (1,816/2,220) and the NPV was 88.9% (8,767/9,853). We additionally examined the performance of the DNN amongst participants who had laboratory-confirmed diabetes within 180 days of a PPG measurement in the Test Dataset (n=152 users; 9,327 measurements) and Contemporary Cohort (n=94 users; 3,659 measurements). Sampling up to 5 measurements per participant, the DNN's recording-level AUCs were similar when using laboratory-confirmed diabetes or self-reported diabetes in both the Test Dataset (0.670, 95% CI: 0.629–0.710; vs 0.650, 95% CI: 0.606–0.694) and the Contemporary Cohort (0.669, 95% CI 0.618–0.719; vs 0.705, 95% CI: 0.657–0.754).

In these laboratory-confirmed diabetes subsets (n=246), there was also evidence for a significant linear association between the continuous DNN Score and both HbA1c and fasting glucose: one standard-deviation increase in DNN Score was associated with 0.32% increase in HbA1c (beta-coefficient=2.28, 95% CI: 1.27-3.29; p 0.001) and 0.11 mmol/L increase in fasting glucose (beta-coefficient=0.82, 95% CI: 0.30-1.34; p 0.001). Similarly, among Clinic Cohort patients with an HbA1c measured within 7 days of the visit (n=93), there was a positive, borderline association between the DNN Score and HbA1c values (beta-coefficient=1.58, 95% CI: -0.021-3.187; p=0.053). Since longstanding poor glycemic control can adversely affect the vasculature and therefore PPG, we also performed a sensitivity analysis comparing DNN performance between HbA1c strata. Among Test Dataset participants with an HbA1c 7.0-8.0% within 6 months of a PPG measurement, we observed similar recording-level AUC=0.636 (95% CI: 0.587-0.686) to that in those with an HbA1c >8.0%, AUC=0.632 (95% CI: 0.585 – 0.679), suggesting similar DNN performance regardless of glycemic control. We also examined the diagnostic odds-ratio for a positive DNN prediction across different Test Dataset strata of gender, age, time of day, recording length and HR (Figure 2b, Extended Data Figure 5). DNN performance was the highest in those with >6 recordings and HR <100 BPM.

Finally, we performed several analyses to help illuminate the mechanisms by which PPG may capture diabetes-related information. We plotted activation maps from inner DNN layers that illustrate how it encodes input PPG recordings, and its behavior in the presence of artifact (Extended Data Figures 6-7). To investigate the role of PPG morphology to predict diabetes in isolation, we trained a separate DNN using a single cardiac cycle PPG waveform as the sole input; user-level AUC=0.691 (95% CI: 0.680–0.700) and recording-level AUC=0.605 (95% CI: 0.600–0.610). To investigate the role of HR and its derivatives in isolation, we trained a separate DNN using only peak-to-peak PPG intervals as the sole input (which removes all PPG morphology information); user-level AUC=0.721 (95% CI: 0.703–0.740) and recording-level AUC=0.645 (95% CI: 0.642–0.647).

#### **Discussion**

In this large-scale study and validation across 3 distinct cohorts, we show that smartphone-measured PPG, analyzed with deep learning, can serve as an independent, noninvasive

digital biomarker of prevalent diabetes. Importantly, the ability of this PPG biomarker to predict diabetes was independent of standard risk factors and comorbidities, and discrimination further improved when adding easily obtainable covariates such as age, gender, race/ethnicity and BMI. Our validation of this digital biomarker in 3 cohorts demonstrated that the DNN generalizes to prospectively enrolled and real-world clinical populations. This digital biomarker of diabetes could serve as a readily attainable complement to other established tools, providing novel information about vascular and autonomic sequelae of diabetes for clinical applications ranging from screening to therapeutic monitoring. However additional research will be needed to determine its utility in these scenarios.

Our work effectively helps to expand the clinical utility of the PPG modality, since physicians do not currently interpret PPG in the context of diabetes. Prior work has reported associations between individually derived PPG features and diabetes-related physiologic changes, mostly using clinic-based pulse oximeters. The physiologic changes most commonly invoked include heart rate variability<sup>24</sup>, endothelial dysfunction<sup>14</sup>, arterial stiffness<sup>15</sup> and combinations thereof<sup>24,25</sup>, providing important early indications that aspects of the PPG waveform contain diabetes-related information. Our study extends these findings, demonstrating that it is not necessary to derive (and be limited to) particular predefined PPG features; rather, the complete PPG recording—containing all the physiologic information—can be analyzed using a DNN to detect diabetes with strong predictive performance. This PPG-derived DNN biomarker is independent of comorbidities and can be augmented with clinical data, when available, to further improve performance. One of the real-world challenges of using remote sensor data to identify disease biomarkers in ambulatory patients is the multiple potential sources of environmental noise, user error and demographic heterogeneity. Our study makes this crucial translational step by using remotely-measured PPGs from commercially available smartphones in a free-living population.

There are various potential applications for a PPG-based digital biomarker of diabetes. Diabetes has numerous characteristics that make it an ideal candidate for screening, such as a prolonged asymptomatic period and the availability of disease modifying therapy. But since population-wide screening is not currently recommended, a widely accessible smart-device-based tool could be used to identify and encourage individuals at higher-risk of having prevalent diabetes to seek medical care and obtain a low-cost confirmatory diagnostic test like HbA1c<sup>26-28</sup>. Leveraging smart-devices to perform diabetes risk prediction without requiring clinic visits would significantly lower barriers to access given the widespread ownership of smartphones, facilitating measurement amongst many of the 224 million people living globally with undiagnosed diabetes<sup>1</sup>. The discriminative performance of our PPG biomarker is comparable to other commonly used tests such as mammography for breast cancer (AUC range 0.67–0.74)<sup>29</sup> or cervical cytology for cervical cancer (AUC range 0.81–0.86)<sup>30</sup>. It compares favorably to existing diabetes-specific risk scores that have AUCs between 0.74-0.85, some of which require serum glucose measurement and none of which is in common clinical use<sup>23</sup>. Reported AUCs of serum-based diagnostic tests like HbA1c or fasting plasma glucose depend on the gold-standard comparator used, but for prevalent microvascular complications range from 0.82–0.96<sup>31</sup>. Comparatively, the ease and

noninvasiveness of PPG make it widely scalable, and its painlessness makes it attractive for repeated testing. Furthermore, since the PPG biomarker is predictive independently of the demographic and comorbidity components comprising most risk scores, it could also be used to supplement existing scores by capturing complementary vascular and autonomic information.

Of the various mechanisms by which PPG may detect diabetes, PPG likely captures the majority of the HR and heart rate variability information as relates to diabetes<sup>10-13,32</sup>. Both predictors were attenuated in the presence of the DNN Score, and peak-to-peak PPG interbeat intervals had only modestly lower AUC (0.721) than the full PPG record (0.766). While interbeat intervals likely contain the predominant predictive information for diabetes, waveform morphologies likely additionally capture information on diabetic vascular changes ranging from endothelial dysfunction<sup>14</sup> to arterial stiffening<sup>15</sup>.

Our study has several limitations. Participants elected to download the iOS smartphone app and therefore may have higher socioeconomic status, technological competence, or health literacy relative to the general population. Our reliance on self-reported diabetes is another limitation. However, our results generalized to the unselected Clinic Cohort, which had medical-record confirmed diabetes, and sensitivity analysis suggested high PPV/NPV against laboratory-confirmed diabetes. Also, misclassification due to self-report at the algorithm training stage would be expected to bias DNN performance toward the null during validation. In analyses that employed laboratory or blood pressure measurements, the time-windows we employed were large and mainly informative as sensitivity analyses. Future studies are needed to confirm this, and whether PPG signals from other sources, like smartwatches, or obtained from anatomic locations like the toe or ear would perform similarly. Given the lower overall prevalence of diabetes, the PPV of our PPG biomarker ranged from 10-32%, which is similar to existing diabetes risk scores whose PPVs mostly range between 10–25% depending on the population and threshold used<sup>23,33</sup>. While false positives are a concern, confirmatory HbA1c is relatively cost-effective; and since individuals with positive DNN predictions were also more likely to have cardiometabolic conditions, they would likely benefit from medical contact. Depending on the intended use of the biomarker, the DNN Score threshold can also be altered to maximize sensitivity or specificity for the intended application. The cross-sectional nature of our study design limited direct investigation of PPG as a diabetes screening tool, or prediction of incident diabetes. We also did not have sufficient data in the Primary Cohort on the type, severity, or medication use for diabetes. The DNN Score did perform similarly, however, in Clinic Cohort subsets with newly diagnosed diabetes and between HbA1c strata. Finally, we were limited in our attempts to compare our approach against standard diabetes risk scores due to the lack of necessary variables in our cohort. These data availability limitations, however, serve to illuminate the difficulty providers also encounter when deploying existing questionnaire-based prediction models, underscoring a strength of noninvasive, objective PPG-based diabetes detection.

In summary, we demonstrate that PPG recorded using consumer-owned smartphones can provide a readily attainable digital biomarker of prevalent diabetes that is independent of standard risk factors and comorbidities. Remote capture of diabetes-predictive PPG

information from ambulatory users is feasible and provides an easily scalable, noninvasive complement to diabetes risk prediction. The linear association of the DNN Score with HbA1c suggests that PPG may additionally capture information about diabetes severity and control, but this requires further investigation. Though this study leverages a large dataset, additional research is needed ideally in targeted intended-use populations to determine how to best incorporate this digital biomarker into existing practice recommendations for diabetes screening and care—particularly in light of the potential for its wide deployment using existing smart-devices outside of the purview of traditional medical care.

#### **Methods**

#### Data sources and study population

The Primary Cohort: The Primary Cohort was derived from 55,433 Health eHeart participants age 18 years, who self-reported a diabetes diagnosis by a healthcare provider and made at least one PPG recording between April 1, 2014 and April 30, 2018 (Figure 1a). Health eHeart is a worldwide, internet-based, longitudinal electronic cohort of English-speaking adults<sup>21</sup>. PPG waveforms were obtained by placing an index fingertip on the smartphone camera (Figure 1b, Extended Data Figure 8). To assess self-reported diabetes status, participants were asked, "Have you ever been told by a doctor, nurse or other healthcare provider that you have diabetes?" and provided the answer options of "Yes," "No" or "Don't know/prefer not to state." Participants who answered "Don't know/prefer not to state" were excluded from our analysis. Participants completed additional surveys regarding demographics, anthropometrics and medical history to varying degrees. We have demonstrated previously<sup>21</sup> that self-reported past medical history in Health eHeart is strongly correlated with the medical record.

The Primary Cohort was randomly split into training (70%, n=37,709), development (10%, n=4,848) and Test (20%, n=11,313) datasets (Extended Data Figure 2; Life Sciences Reporting Summary). The training dataset was used for DNN development and training, and DNN hyperparameters were tuned in the development dataset. Final model performance is reported in the Test Dataset, which was kept completely separate until the final evaluation step.

**Two Additional Validation Cohorts:** In addition to validating the performance of the DNN algorithm in the Primary Cohort Test Dataset, we additionally reported DNN performance in 2 validation cohorts (Figure 1a), providing 3 examples of validation in datasets separate from training data<sup>23</sup>. The first was the "Contemporary Cohort" which was composed of PPG recordings from 7,806 participants newly enrolled into Health eHeart between May 1, 2018 and December 31, 2018 (Extended Data Figure 3, Supplemental Table 3). This temporally distinct validation cohort helps to account for secular changes, such as changes in smartphone models and cameras, that could affect PPG recording. Then, to test the validity of our approach in a real-world clinical setting, we prospectively enrolled an in-person "Clinic Cohort" comprised of 181 consecutive patients referred to 3 cardiovascular prevention clinics (2 in San Francisco, one in Montreal) between November 1, 2018 and July 30, 2019 (Extended Data Figure 3, Supplemental Table 4). Clinic Cohort participants were

consented, assessed for height, weight, body mass index (BMI), and a trained coordinator obtained at least 15 seconds of a single PPG recording using an iPhone and determined diabetes status by medical chart review. For the subset of Clinic Cohort patients who also had fasting glucose and HbA1c obtained within 7 days of the in-clinic visit, we used the American Diabetes Association diagnosis criteria to classify participants as having/not having diabetes<sup>34</sup>.

The UCSF Institutional Review Board approved the study and all participants gave informed consent.

#### **PPG Waveform Acquisition and Preprocessing**

PPG waveforms were obtained by placing the index fingertip<sup>35</sup> on the smartphone camera using the Azumio Instant Heart Rate iOS smartphone application (Azumio, Inc). Though the app is available for Android and iOS operating systems, data was limited to iOS app versions in this study due to data availability in Health eHeart. Changes in reflected light intensity recorded by the smartphone camera are interpreted as pulsatile blood volume change. The waveforms were pre-processed by the Azumio algorithm for camera artifact removal, utilizing standard de-trending and low pass filter techniques (Figure 1). A Low pass ~0.4Hz, 2nd order, zero phase shift IIR filter is used to find the trend, the trend is subtracted to get the detrended signal. Another Low pass ~10Hz, 2nd order, zero phase shift IIR filter is used to remove high frequency noise. Individual beats corresponding to cardiac cycles were identified using the rising-edge of the PPG signal. If the recording does not have at least 5 seconds of continuous discernable peak-to-peak intervals, it was removed. Waveforms with a length under 5 seconds or with an amplitude of "0", indicating a null signal, were also removed. We excluded outlier PPG measurements defined as HR values of outside of the biologically plausible range of 20–220 bpm. We limited waveforms in our dataset to those collected at either 100Hz or 120Hz, and upsampled recordings of 100 Hz to 120 Hz using the standard polyphase method<sup>36</sup> to minimize variance due to sampling frequency. We derived the onset of each cardiac cycle by identifying the rising-edge of the waveform, used to determine HR and heart rate variability (using Root Mean Square of Successive peak-to-peak interval differences).

#### **DNN Development and Performance**

We built a 39-layer convolutional DNN to detect prevalent diabetes (Extended Data Figure 9). The DNN takes the PPG-waveform as the sole input, which consists of 2,560 samples equivalent to ~21.3 seconds (approximately the mean signal duration), and outputs a DNN Score between 0 and 1 per signal; higher scores suggest greater likelihood of diabetes. Shorter signals were zero-padded up to the fixed-length and longer examples were cropped. All PPG waveforms were standardized using the mean and standard deviation values of the entire training dataset. The network architecture had 39 layers organized in a block structure, consisting of convolutional layers with an initial filter size of 15 and filter number of 16. The size of the filters decreased, and the number of filters increased, as network depth increased. After each convolutional layer, we applied batch normalization<sup>37</sup>, rectified linear activation<sup>38</sup> and dropout <sup>39</sup> with a probability of 0.2. The final flattened and fully connected

softmax layer produced a distribution across the classes of diabetes/no diabetes<sup>40</sup>. Weights were initialized randomly as described by He et al<sup>41</sup>.

We used grid-search to tune the network hyperparameters by searching over the best optimizer, best initializer, number of convolutional layers, the stride size, the filter length, the number of filters, the class weight, the learning rate, the input length of the signal, the batch size, the dropout, the early stopping criteria and the amount of cropping of the start/end of the signal, based on the recording-level development dataset performance. The best performance was achieved by cropping 2 beats from the beginning and 1 beat from the end of the signal; this was applied to all PPG records. For all the models presented, we used the Rectified ADAM optimizer with the default parameters<sup>42</sup>, and a mini-batch size of 512. The learning rate was initialized at  $1 \times 10^{-3}$  and was adjusted based on the effects of variance and momentum during training 42. We halted training after an absence of improvement in the loss within the development set for 8 consecutive epochs. A class weight of 10:1 for diabetes to non-diabetes recordings was applied to our loss function. The best performing model was chosen based on the development dataset recording-level AUC performance and was then applied to all validation sets. We explored different architectures involving recurrent layers, such as long-short-term memory cells and residual blocks (ResNet), and with age or hour of the day added as additional inputs to the DNN, but found no improvement in AUC despite substantial increases in model complexity and runtime. The DNN was trained for 18 epochs.

#### **Grid-search of hyperparameters**

We performed systematic search of hyperparameters among these values:

- Model architecture: Convolutional neural network, ResNet, LSTM
- Number of convolutional layers: 7, 15, 19, 25, 29, 35, 39
- Filter length: 5, 7, 9, 11, 13, 15
- Number of filters to start: 8, 16, 32, 64
- Optimizer: Adam, Rectified Adam
- Class weight for 'diabetes': 5, 10, 15, 20
- Initializer: Glorot, He
- Learning Rate: 10e-1 10e-2, 10e-3, 10e-4, 10e-5
- Input shape [2560, 1]; [2048, 1]
- Batch size: 64, 128, 256, 512
- Dropout: 0.2, 0.4, 0.6
- Early stopping criteria: 6, 8, 12, 20
- (Preprocessing) Number of beats cropped at the start of the signal: 0, 1, 2, 3
- (Preprocessing) Number of beats cropped at the end of the signal: 0, 1, 2, 3

We reported DNN performance using the area under the receiver operating characteristic curve (AUC)<sup>22</sup> in 3 separate test datasets: 1) the Primary Cohort Test Dataset, 2) the Contemporary Cohort, and 3) the in-person Clinic Cohort. Since many participants contributed >1 recording, we assessed model performance both at the 'recording-level,' which treats each recording independently, and at the 'user-level,' which averages the DNN Score for all recordings provided by an individual user. Our primary aim was to evaluate the user-level DNN Score, since the clinical goal would be to classify a patient as having diabetes or not. Clinic cohort patients have only recording-level performance since only a single recording was obtained per patient during their visit. We also plotted the activation maps of several hidden convolutional layers of the trained DNN<sup>43</sup> from an example PPG record to help illuminate some of the higher-level PPG features derived by the DNN (Extended Figures 8-9).

#### **Sensitivity Analyses**

To better ascertain the reliability of self-reported diabetes in the Primary Cohort, we described the PPV and NPV of self-reported diabetes in the larger Health eHeart study<sup>21</sup> using fasting glucose or HbA1c drawn within 180 days of self-reported diabetes; if >1 lab value was available the value closest in time to self-report was used. Laboratory-confirmed diabetes was defined according to the ADA guidelines: HbA1c 7.0%, fasting glucose (fasting glucose 126 mg/dl or 7.0 mmol/L)<sup>34</sup> or non-diabetic range of HbA1c/fasting glucose but self-report of taking diabetes medications. We also examined DNN performance among the subset of Test Dataset and Contemporary Cohort participants who had laboratoryconfirmed diabetes using laboratory values drawn within 180 days of a PPG measurement. For those with multiple measurements, we randomly sampled up to 5 measurements. To understand the performance of the DNN according to glycemic control in the laboratoryconfirmed diabetes cohort, we examined DNN performance in strata of HbA1c above and below 8.0%. Additionally, in the Clinic Cohort, we examined DNN performance after excluding those with a prior diagnosis of diabetes. Linear regression models were fit with the DNN Score as the predictor and either HbA1c or glucose value as the dependent variable in the Test Dataset and Contemporary Cohort. To investigate the role of HR in isolation, we trained a separate DNN to detect diabetes using only peak-to-peak intervals as input and the same architecture and training data as the primary DNN. To investigate the role of the PPG waveform in isolation, we trained a separate DNN using the PPG waveform from a single cardiac cycle, removing the time-domain contribution from consecutive cardiac cycles. In the Clinic Cohort, we also modeled the DNN Score against HbA1c as the dependent variable with linear regression.

#### Statistical Analysis

Basic demographics and previous medical conditions are presented for each dataset, and continuous data are presented as mean  $\pm$  standard deviation. The "DNN Score" is the final layer of the DNN, which is an output distribution for diabetes based on the PPG input. We identified a discrimination threshold for the DNN Score that maximized the macro average sensitivity between the 'diabetes' and 'no-diabetes' classes in the training dataset<sup>44</sup>; this threshold is applied to all relevant performance metrics. We present sensitivity, specificity, positive predictive value (PPV) and negative predictive value (NPV) for each of our test

datasets<sup>22</sup>. Confidence intervals for these metrics were derived by bootstrapping 80% of the test data over 1000 iterations to obtain 5th and 95th percentile values. The diagnostic odds-ratio (odds-ratio) is a measure of the effectiveness of a diagnostic test and is defined as the ratio of the odds of the DNN Score being positive for diabetes if the subject has diabetes, relative to the odds of the DNN Score being positive if the subject does not have diabetes<sup>45</sup>. Odds-ratios, 2-sided p-values for interaction (between the DNN Score, the covariates and diabetes, calculated by the Wald test), sensitivity and specificity were calculated separately between different strata of age, gender and PPG recording characteristics.

To understand the incremental contribution of PPG-based predictions alongside commonly available demographic and clinical predictors of diabetes, we built nested logistic regression (LogReg) models for prevalent diabetes both with and without the inclusion of the standardized DNN Score. LogReg-Model-1 included age as a covariate; LogReg-Model-2 additionally included gender and race/ethnicity; LogReg-Model-3 additionally included BMI. Since HR<sup>12,13</sup> and heart rate variability<sup>46</sup> are known independent predictors of diabetes and can be derived from the PPG signal, we examined the specific role of HR in the PPG-based prediction of diabetes by including the per-record average HR and heart rate variability (calculated using the root mean square of the successive peakto-peak differences) as covariates in LogReg-Model-4. Finally, in LogReg-Model-5 we included clinical comorbidities commonly known to co-occur with diabetes, to ascertain the independent value of the PPG DNN Score for identifying diabetes. All continuous logistic regression variables were standardized using the Z-score to allow comparison between oddsratios and we used complete-case analysis, excluding individuals with missing covariates. Tests for normality were performed and met by all continuous predictors, and there were no adjustments made for multiple comparisons.

A 2-sided p-value <0.05 was considered significant. The convolutional neural network was built in Python 2.7 using Keras (version 2.0.3) and TensorFlow (version 1.13.2). The LogReg models and AUC were derived in SPSS v24.0 (IBM).

**Code Availability:** The code that supports this work is copyright of the Regents of the University of California and can be made available through license.

#### **Extended Data**

	With Diabetes (n = 3,564)	Without Diabetes (n = 50,306)	p-value
Age, years, mean ± SD Gender	54.6±14.7 <i>N=2162</i>	45.0±15.1 <i>N=35,281</i>	<0.001
Females, n (%) Males, n (%)	881 (40.7%) 1,281 (59.3%)	16,664 (47.2%) 18,617 (52.8%)	<0.001
Race and ethnic group	N=2,162	N=35,281	< 0.001
Non-Hispanic White, n (%)	1,564 (72.3%)	26,657 (75.6%)	
Black or African American, n (%)	87 (4.0%)	709 (2.0%)	
Hispanic, Latino or Spanish origin/ancestry, n (%)	219 (10.1%)	3,757 (10.6%)	
Asian, n (%)	167 (7.7%)	2,062 (5.8%)	
Multi-ethnic, n (%)	80 (3.7%)	1,296 (3.7%)	
Other, n (%)	45 (2.1%)	800 (2.3%)	
Waveform data	N=3,564	N=50,306	
Total number of recordings	182,912	2,406,536	
Number of recordings per user, mean ± SD	51.3±101.3	47.8±94.9	0.046
Duration of waveform, seconds, mean ± SD	21.2±10.2	22.1±10.9	<0.001
Heart rate, bpm, mean ± SD	83.8±14.5	79.9±15.1	<0.001
Anthropometric data	N=369	N=5,267	
Height, meters, mean ± SD	1.73±0.11	1.73±0.10	0.98
Weight, kg, mean ± SD	96.5±23.6	81.8±19.7	<0.001
BMI, mean ± SD	32.1±7.0	27.3±5.9	<0.001
Medical conditions	N=3.564	N=50.306	101001
No reported medical	0 (0%)	26,782 (53.2%)	< 0.001
conditions, n (%)	- ()	,,	
Diabetes mellitus, n (%)	3,564 (100%)	0 (0%)	< 0.001
Hypertension, n (%)	2,342 (65.7%)	11,802 (23.5%)	< 0.001
Hypercholesterolemia, n (%)	2,238 (62.8%)	13,120 (26.1%)	< 0.001
Coronary artery disease, n	792 (22.2%)	2,974 (5.9%)	< 0.001
(%)			
Prior MI, n (%)	478 (13.4%)	1,336 (2.7%)	< 0.001
CHF, n (%)	384 (10.8%)	817 (1.6%)	< 0.001
PVD, n (%)	345 (9.7%)	689 (1.4%)	< 0.001
Prior Stroke, n (%)	357 (10.0%)	1,111 (2.2%)	< 0.001
Sleep apnea n (%)	1,198 (33.6%)	5,803 (11.5%)	<0.001

#### Extended Data Fig. 1. Baseline characteristics of the Primary Cohort by Diabetes Status

Primary Cohort sample size was 53,870 individual people. Where data was only available for subgroups of the full cohort, subgroup sample size is denoted by N. Differences in means of continuous variables between 2 groups were compared using the two-sample t-test. Differences in proportions of categorical variables between 2 groups were compared using the Chi-Squared test. Tests of significance were 2 sided. Abbreviations: bpm: beats per minute; CAD: Coronary artery disease; CHF: Congestive heart failure; COPD: Chronic obstructive pulmonary disease; HR: Heart rate, MI: Myocardial Infarction; PVD: Peripheral Vascular Disease.

	Training (n = 37,709)	Development (n = 4,848)	Test (n = 11,313)	p-value
Age, years, mean ± SD	44.9±14.7 <sub>a</sub>	44.7±14.8 a	44.9±14.6 <sub>a</sub>	0.430
Gender	N=26,244	N=3,348	N=7,851	
Females, n (%)	12,459 (47.4%) <sub>a</sub>	1,573 (47.0%) <sub>a,b</sub>	3,513 (44.7%) <sub>b</sub>	0.540
Males, n (%)	13,785 (52.5%) <sub>a</sub>	1,775 (53.0%) <sub>a,b</sub>	4,338 (55.3%) <sub>b</sub>	0.540
Race and ethnic group	N=26,244	N=3,348	N=7,851	
Non-Hispanic White, n (%)	19,772 (75.3%) <sub>a</sub>	2,506 (74.9%) <sub>a</sub>	5,943 (75.7%) <sub>a</sub>	
Black or African American, n (%)	561 (2.1%) a	61 (1.8%) <sub>a</sub>	174 (2.2%) <sub>a</sub>	
Hispanic, Latino or Spanish origin/ancestry, n (%)	2,776 (10.6%) <sub>a</sub>	384 (11.4%) <sub>a</sub>	816 (10.4%) <sub>a</sub>	0.310
Asian, n (%)	1,562 (5.9%) a	208 (6.2%) a	459 (5.8%) a	
Multi-ethnic, n (%)	985 (3.8%) a	119 (3.6%) a	272 (3.5%) a	
Other, n (%)	588 (2.2%) a	70 (2.1%) <sub>a</sub>	187 (2.4%) a	
Waveform data	N=37,709	N=4,848	N=11,313	
Duration of waveform, mean	22.1±11.0 <sub>a</sub>	21.7±10.7 <sub>b</sub>	21.8±10.5 <sub>b</sub>	0.003
± SD				
Heart rate, mean ± SD	80.1±15.1 <sub>a</sub>	80.4±15.2 <sub>a</sub>	80.4±15.1 <sub>a</sub>	0.110
Anthropometric data	N=3,956	N=515	N=1,165	
Height, meters, mean ± SD	1.73±0.10 <sub>a</sub>	1.73±0.10 <sub>a</sub>	1.73±0.10 <sub>a</sub>	0.165
Weight, kg, mean ± SD	$82.3\pm20.02_a$	82.6±20.0 <sub>a</sub>	84.2±21.0 <sub>b</sub>	0.025
BMI, mean ± SD	27.5±6.0 <sub>a</sub>	27.8±6.2 <sub>a</sub>	28.0±6.3 a	0.165
Medical conditions	N=37,709	N=4,848	N=11,313	
No reported medical conditions, n (%)	16,444 (43.6%) a	2,109 (43.5%) a	4,917 (43.5%) <sub>a</sub>	0.990
Diabetes mellitus, n (%)	2,516 (6.7%) a	298 (6.1%) a	750 (6.6%) a	0.382
Hypertension, n (%)	9,851 (26.1%) a	1,293 (26.8%) a	3,000 (26.7%) a	0.562
Hypercholesterolemia, n (%)	10,668 (28.3%) a	1,347 (27.8%) <sub>a</sub>	3,189 (28.1%) a	0.235
Coronary artery disease, n	2,658 (7.1%) a	308 (6.4%) a	800 (7.1%) a	0.186
(%)				
Prior MI, n (%)	1,285 (3.4%) <sub>a</sub>	152 (3.1%) <sub>a</sub>	377 (3.4%) <sub>a</sub>	0.594
CHF, n (%)	852 (2.3%) <sub>a</sub>	115 (2.4%) <sub>a</sub>	234 (2.1%) <sub>a</sub>	0.366
PVD, n (%)	740 (2.0%) <sub>a</sub>	82 (1.7%) <sub>a</sub>	212 (1.9%) <sub>a</sub>	0.395
Prior Stroke, n (%)	1,019 (2.7%) <sub>a</sub>	124 (2.6%) <sub>a</sub>	325 (2.9%) <sub>a</sub>	0.482
Sleep apnea n (%)	4,914 (13.1%) a	620 (12.8%) <sub>a</sub>	1,467 (13.0%) a	0.919

# Extended Data Fig. 2. Baseline Characteristics in the Primary Cohort Training, Development and Test Datasets

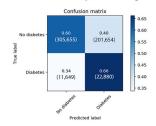
Primary cohort sample size was 53,870 individual people. Where data was only available for subgroups of the full cohort, subgroup sample size is denoted by N. Differences in means of continuous variables between 2 groups were compared using two-sample t-test. Differences in means of continuous variables between 3+ groups were compared using one-way ANOVA. Differences in proportions of categorical variables between the 2+ groups were compared using Chi-Squared. Tests of significance were 2 sided. a, b, c: Each subscript letter denotes a subset of dataset categories whose column proportions do not differ significantly from each other at the 0.05 level. Post-hoc analysis was performed using Fisher's least significant differences to compare means of continuous variables between groups. Abbreviations: SD: Standard deviation; CAD: Coronary artery disease; CHF: Congestive heart failure; COPD: Chronic obstructive pulmonary disease; HR: Heart rate, MI: Myocardial Infarction; PVD: Peripheral Vascular Disease.

	Primary Cohort (n = 53,870)	Contemporary Cohort (n = 7,806)	Clinic Cohort (n = 181)	p-value
Age, years, mean ± SD Gender Females, n (%) Males, n (%)	45.6±16.2 a N=37,443 17,545 (46.9%) a 19,898 (53.1%) a	44.5±16.3 <sub>b</sub> N=3,936 2,237 (56.8%) <sub>b</sub> 1,699 (43.2%) <sub>b</sub>	63.1±14.7 c N=181 66 (36.4%) c 115 (63.5%) c	<0.001
Wales, 11 (76)	19,090 (33.1%) a	1,099 (43.276)6	115 (05.5%) c	<0.001
Race and ethnic group	N=37,443	N=3,936		
Non-Hispanic White, n (%)	28,221 (75.3%) a	2,960 (75.2%) <sub>a</sub>	-	
Black or African American, n (%)	796 (2.1%) <sub>a</sub>	107 (2.7%) <sub>b</sub>	•	
Hispanic, Latino or Spanish origin/ancestry, n (%)	3,976 (10.6%) <sub>a</sub>	456 (11.5%) <sub>b</sub>	-	<0.001
Asian, n (%)	2,229 (5.9%) a	183 (4.6%) ь		
Multi-ethnic, n (%)	1,376 (3.7%) a	119 (3.0%) b	-	
Other, n (%) Waveform data	845 (2.3%) <sub>a</sub> N=53.870	111 (2.8%) <sub>b</sub> <i>N=7.806</i>	N=181	
Duration of waveform, mean ± SD	22.0±10.9 <sub>a</sub>	20.6±9.4 bå	30.1±18.3 c	<0.001
Heart rate, mean ± SD	80.2±15.1 <sub>a</sub>	80.8±14.0 <sub>b</sub>	76.2±24.0 <sub>c</sub>	< 0.001
No. of recordings, mean ± SD	48.1±95.3 <sub>a</sub>	31.3±64.5 <sub>b</sub>	1±0 <sub>c</sub>	<0.001
Anthropometric data	N=5,636	N=650	N=181	
Height, meters, mean ± SD	1.72±0.10 a	1.71±0.01 <sub>b</sub>	1.68±0.26 c	<0.001
Weight, kg, mean ± SD	82.9±20.3 a	82.7±21.8 a	84.4±24.2 <sub>a</sub>	0.643
BMI, mean ± SD	27.7±6.1 a	28.2±6.6 b	28.9±6.6 <sub>в</sub>	0.008
Medical conditions No reported medical	N=53,870 26,782 (49.7%) <sub>a</sub>	<i>N=7,806</i> 3,517 (45.0%) <sub>b</sub>	<i>N=181</i> 49 (27.1%) ₀	<0.001
conditions	0.504 (0.004)	070 (0 70/)	00 (04 00/)	0.004
Diabetes mellitus, n (%)	3,564 (6.6%) a	679 (8.7%) <sub>b</sub>	38 (21.0%) c	<0.001
Hypertension, n (%) Hypercholesterolemia, n (%)	14,144 (26.3%) a 15,349 (28.5%) a	2,452 (31.4%) <sub>b</sub> 2,381 (30.5%) <sub>b</sub>	99 (54.7%) c 76 (42.0%) c	<0.001 <0.001
Coronary artery disease, n (%)	3,766 (7.0%) a	655 (8.4%) <sub>b</sub>	11 (6.1%) a	<0.001
Prior MI, n (%)	1,814 (3.4%) <sub>a</sub>	301 (3.9%) <sub>b</sub>	11 (6.1%) a, b	0.013
CHF, n (%)	1,201 (2.2%) a	232 (3.0%) b	11 (6.1%) c	<0.001
PVD, n (%)	1,034 (1.9%) a	179 (2.3%) <sub>b</sub>	6 (3.3%) <sub>c</sub>	<0.001
Prior Stroke, n (%)	1,468 (2.7%) a	289 (3.7%) <sub>b</sub>	5 (2.8%) a, b	< 0.001
Sleep apnea n (%)	7,001 (12.9%) <sub>a</sub>	1,263 (16.2%) b	13 (7.2%) a. b	<0.001

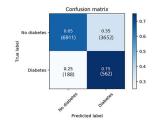
### Extended Data Fig. 3. Baseline Characteristics of the Primary, Contemporary and Clinic Cohorts

Where data was only available for subgroups of the full cohorts, subgroup sample size is denoted by N. Differences in means of continuous variables between 2 groups were compared using two-sample t-test. Differences in means of continuous variables between 3+ groups were compared using one-way ANOVA. Differences in proportions of categorical variables between the 2+ groups were compared using Chi-Squared. Tests of significance were 2 sided. <sup>a, b, c</sup>: Each subscript letter denotes a subset of dataset categories whose column proportions do not differ significantly from each other at the 0.05 level. Post-hoc analysis was performed using Fisher's least significant differences to compare means of continuous variables between groups. Abbreviations: SD: Standard deviation; CAD: Coronary artery disease; CHF: Congestive heart failure; COPD: Chronic obstructive pulmonary disease; HR: Heart rate, MI: Myocardial Infarction; PVD: Peripheral Vascular Disease.

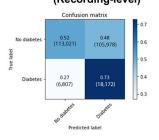
#### a. Test Dataset (Recording-level)



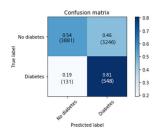
#### b. Test Dataset (User-level)



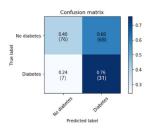
# c. Contemporary Cohort (Recording-level)



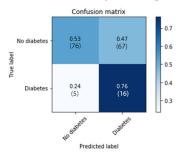
d. Contemporary Cohort (User-level)



#### e. Clinic Cohort (Recording-level)



# f. Clinic Cohort, Newly Diagnosed Diabetes Subset (Recording-Level)



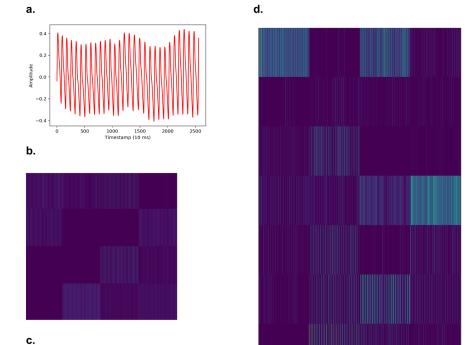
## Extended Data Fig. 4. Data Figure 4. Confusion matrices for DNN performance in 3 validation datasets.

Confusion matrices for the predictions of the DNN in the Test Dataset (a-b), Contemporary Cohort (c-d), and Clinic Cohort (e-f), at both the recording and user-level. Total number of patients are presented in parentheses. The DNN Score cutoff used was 0.427.

Group	Sensitivity	Specificity	OR (95% CI)	Interaction p-value	AUC	
Time of the recording						
0-4 AM	60% (2,463/4,107)	63% (42,167/66,862)	2.56 (2.40-2.73)	0.01	0.66	•
4-8 AM	64% (4,745/7,390)	59% (54,320/91,565)	2.62 (2.49-2.75)	0.04	0.66	
8 AM-12 PM	67% (3,824/5,668)	58% (52,752/90,664)	2.89 (2.72-3.06)	0.78	0.68	
12-4 PM	65% (3,992/6,124)	60% (58,524/97,096)	2.84 (2.69-3.00)	0.37	0.67	•
4-8 PM	70% (4,487/6,446)	58% (49,804/85,206)	3.22 (3.05-3.40)	0.26	0.68	•
3 PM-12 AM	71% (2,566/3,591)	63% (34,878/54,633)	4.42 (4.10-4.76)	< 0.001	0.73	
ength of the recording						
5-10 seconds	62% (1,252/2,010)	58% (14,835/25,429)	2.32 (2.11-2.54)	0.01	0.64	•
0-15 seconds	67% (12,219/18,347)	62% (165,903/268,934)	3.21 (3.11-3.31)	0.04	0.69	
15-20 seconds	67% (5,117/7,638)	60% (67,179/112,338)	3.02 (2.87-3.17)	0.03	0.68	
20 seconds	65% (4,067/6,227)	57% (53,441/93,205)	2.53 (2.40-2.67)	< 0.001	0.66	
leart rate						
60 BPM	43% (1,426/3,343)	83% (76,562/92,029)	3.68 (3.43-3.95)	0.05	0.73	
0-80 BPM	64% (9,168/14,327)	63% (133,211/211,761)	3.01 (2.91-3.12)	0.53	0.68	
80-100 BPM	77% (8,531/11,009)	44% (53,470/121,646)	2.70 (2.58-2.83)	0.53	0.65	
00-120 BPM	73% (2,701/3,695)	38% (16,455/42,985)	1.69 (1.53-1.56)	0.01	0.59	
120 BPM	45% (829/1,848)	68% (21,660/31,485)	1.79 (1.63-1.97)	0.02	0.61	
Overall (Recording-level)	66% (22,880/34,529)	60% (305,655/507,309)	2.98 (2.91-3.05)		0.68	

# Extended Data Fig. 5. DNN performance to predict diabetes according to time of day, recording length and heart rate in the Test dataset.

DNN sensitivity, specificity, diagnostic odds-ratio and AUC to detect prevalent diabetes are presented across strata of age, gender and number of recordings. The Test Dataset sample size is 11,313 individuals. Counts are provided in parentheses for all subgroup metrics. The diagnostic odds-ratio is the ratio of positive likelihood ratio (sensitivity / (1–specificity)) to the negative likelihood ratio ((1–sensitivity)/specificity). The diagnostic odds-ratio is presented at the recording-level with the associated 95% confidence interval. Interaction p-values are two-sided Wald tests for interaction between the DNN Score and the respective covariates for diabetes. Abbreviations: DNN: deep neural network; OR: diagnostic odds ratio; AUC: area under the curve; CI: confidence interval; BPM: beats per minute.

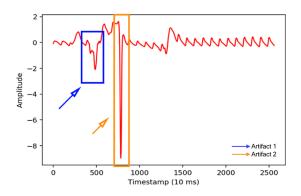


 $\label{lem:extended} \begin{tabular}{ll} Extended Data Fig. 6. Activation maps from several hidden convolutional layers of the trained Deep Neural Network (DNN) for one photoplethysmography (PPG) record. \\ \end{tabular}$ 

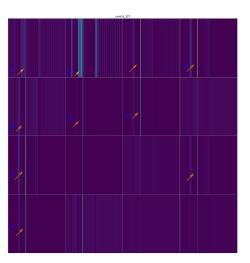
**a.** An example of a PPG recording which serves as the input into the DNN. **b.** The activation map of one example filter (out of 16) from the first convolutional layer of the neural network. This activation map is obtained after the example PPG recording is fed into the trained DNN. Each lighter colored band illustrates "activation" of a model parameter. At this early layer of the neural network, the lighter colored bands correspond directly to each cardiac cycle of the PPG waveform. Thicker lines likely indicate morphological features of the waveform. **c.** Visualization of the activation maps of the 16 filters from the first convolutional layer of the neural network, obtained after the input PPG is fed into the trained DNN. Each of the 16 filters can learn different sets of "features" from the input PPG recording. Filters with more purple bands have more inactive neurons, as compared to those with lighter colors (green being the strongest activation and dark purple being the weakest activation). Six filters appear completely inactivated (all purple), suggesting that the features these filters focus on are not present in this example input PPG. **d.** Visualization of the activation maps of the 7<sup>th</sup> convolutional layer of the DNN, comprised of 32 filters. Broadly, these activation maps from the 7<sup>th</sup> layer of the DNN are more complex compared to those

from the 1<sup>st</sup> layer (b-c), demonstrating how deeper layers of the DNN encode increasingly abstract information representing higher level interactions and complex features.

а



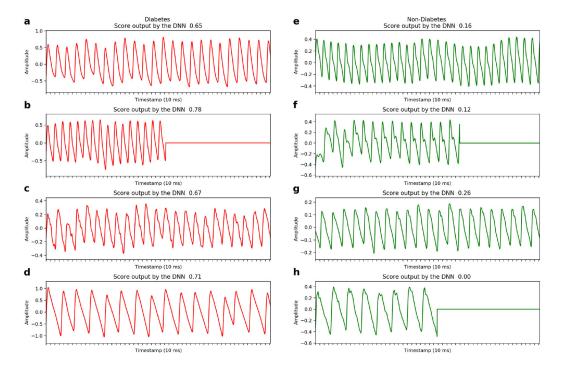
b.



Extended Data Fig. 7. Activation maps from hidden convolutional layers of the trained Deep Neural Network (DNN) for an example photoplethysmography (PPG) recording with artifacts. a. An example PPG recording with 2 artifacts (blue and orange rectangles) which serves as the input into the DNN. b. Activation maps of the 16 filters from the first convolutional layer of the DNN. Each lighter colored band illustrates "activation" of a model parameter. Orange and blue arrow are placed on filters denoting the location of artifacts, highlighted by orange and blue rectangles (a), respectively. Some filters, such as the 4<sup>th</sup> image in the top row, seem to not have activation at the location of the artifactual beats (hollow orange and blue arrows), suggesting that the DNN is "ignoring" data from these artifact locations. Whereas other filters are have activation, suggested by lighter color bars, in the locations of the artifacts (full orange and blue arrows), such as the 2<sup>nd</sup> filter from the left in the top row, suggesting that the DNN is using data from these artifact locations. Some filters, such as the 2<sup>nd</sup> from the left in the bottom row "ignore" the artifactual beats by having uniform

activation throughout the signal length (except where there are artifacts) likely representing

the cardiac cycle. These findings suggest that the DNN is able to identify artifactual beats and differentiate them from good quality waveforms.



#### Extended Data Fig. 8. Example photoplethysmography (PPG) waveforms.

**a**. Examples of raw PPG recordings from individuals with and without diabetes (red/green recordings, respectively), which serve as inputs to the deep neural network. DNN Scores predicted for each recording are shown. PPG recordings are either cropped or zero-padded to the same fixed length (~20.3 seconds) before being input into the DNN. The "flat line" in three examples is a demonstration of zero-padding shorter records to the fixed length. DNN: Deep Neural Network; ms: milliseconds.

### **PPG** signal input PPG Waveform truncated or zero-padded to 2560 Datapoints sampled at 120 Hz (21.3 secs) Convolution **Conv Layers** N filters of stride S, shaped as 1xK **Batch Normalization** K: 15, 15, 15, 13, 13, 13, 11, 11, 11, 9, 9, 9, 7, 7, 18 7, 5, 5, 5 Relu S: 1, 2, 1, 2, 1, 2, 1, 2, 1, 2, 1, 2, 1, 2, 1, 2, 1, 2 N: 16, 16, 16, 16, 16, 16, 32, 32, 32, 32, 32, 32, Dropout 64, 64, 64, 64, 64, 64 Flatten **Fully Connected** Softmax

#### Extended Data Fig. 9. Deep Neural Network architecture.

Diabetes

The neural network had 39 layers organized in a block structure, consisting of convolutional layers with an initial filter size of 15 and filter number (N) of 16. The size of the filters decreased, and the number of filters increased as network depth increased, as shown. After each convolutional layer, we applied batch normalization, rectified linear activation and dropout with a probability of 0.2. The final flattened and fully connected softmax layer produced an output distribution across the classes of diabetes/no diabetes. This output distribution is referred to as the DNN Score. PPG: photoplethysmography; DNN: Deep Neural Network; Hz: Hertz.

No diabetes

### **Supplementary Material**

Refer to Web version on PubMed Central for supplementary material.

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### **Data Availability:**

The data that support the findings of this study are available from the authors and Azumio, but restrictions apply to the availability of these data, which were used under license for the current study, and so are not publicly available. Data are however available from the authors upon reasonable request and with permission of Azumio.

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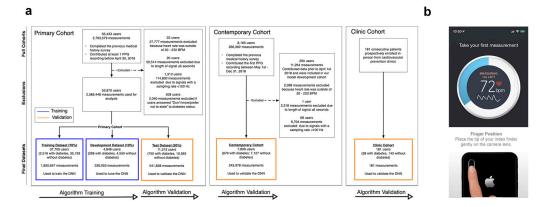


Figure 1: Consort diagram describing the study cohorts and screenshots from the smartphone app used for PPG acquisition

**a.** Description of the datasets used for algorithm development and validation. The deep neural network (DNN) was trained using the training and development dataset of the Primary Cohort (left), and validated using the test dataset of the Primary Cohort. We additionally validated the DNN in the temporally-distinct Contemporary Cohort (middle) and the prospectively enrolled, in-person Clinic Cohort (right). Blue outlines indicate datasets used for model development and training. Yellow outlines indicate datasets used for model validation. All datasets are completely separate and do not contain overlapping participants. **b.** Screenshots from the smartphone app used to acquire user-measured PPG recordings using a smartphone app and camera. PPG: photoplethysmography; BPM: beats per minute. DNN: Deep Neural Network; Hz: Hertz.

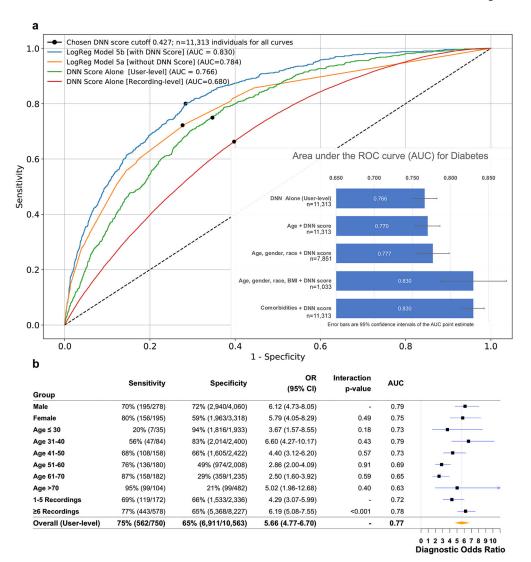


Figure 2: Comparison of model performance to detect diabetes in the Test Dataset.

a. Receiver operating characteristic curves for detection of diabetes, as assessed for the DNN score alone or for the output of LogReg Model 5, which includes comorbidities, with and without the DNN Score. This is calculated at either the recording-level, which treats each recording independently, or at the user-level, which is averaged across all recordings of an individual user. The DNN Score cutoff used (0.427) is indicated by a black dot on each curve. Inset: Bar chart showing the area under the receiver operating characteristic curve (AUC) point estimate values for diabetes in the test dataset by the indicated models; 95% confidence intervals are shown as error bars. b. DNN sensitivity, specificity, diagnostic oddsratio and AUC to detect prevalent diabetes in the Test Dataset, as reported across ranges of age, gender and number of recordings. The Test Dataset sample size is 11,313 individuals. Counts are provided in parentheses for all subgroup metrics. The diagnostic odds-ratio was quantified as the ratio of positive likelihood ratio (sensitivity / (1–specificity)) to the negative likelihood ratio ((1–sensitivity)/specificity), with the associated 95% CI. The diagnostic odds-ratio is presented at the user-level for strata of age, gender and number of recordings. Interaction p-values are two-sided Wald tests between the DNN Score and the respective

covariates for diabetes. Abbreviations: DNN: deep neural network; AUC: area under the receiver operating characteristic curve; OR: diagnostic odds ratio; CI: confidence interval.

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Table 1. Performance of the DNN to detect diabetes using PPG in three validation datasets.

Sample sizes reported indicate individual people. User-level performance metrics are reported based on the average DNN score for all recordings from an individual user. Recording-level performance metrics are calculated treating each recording independently. Since Clinic Cohort participants only received one measurement, only the recording-level metric is reported for this cohort. Samples sizes shown indicate numbers of individual people.

	AUC (95% CI)	Sensitivity*	Specificity*	$\mathrm{PPV}^*$	NPV*
Test Dataset, n=11,313					
User-level	0.766 (0.750-0.782)	75.0% (72.0–77.8%)	65.4% (64.6–66.3%)	$0.766 \ (0.750 - 0.782) \qquad 75.0\% \ (72.0 - 77.8\%) \qquad 65.4\% \ (64.6 - 66.3\%) \qquad 13.3\% \ (12.3 - 14.3\%) \qquad 97.4\% \ (97.0 - 97.7\%)$	97.4% (97.0–97.7%)
Recording-level	0.680 (0.678-0.683)	66.2% (65.8–66.7%)	60.2% (60.1–60.3%)	$0.680 \ (0.678 - 0.683) \\ 66.2\% \ (65.8 - 66.7\%) \\ 60.2\% \ (60.1 - 60.3\%) \\ 10.2\% \ (10.0 - 10.3\%) \\ 96.3\% \ (96.3 - 96.4\%) \\ $	96.3% (96.3 – 96.4%)
Contemporary Cohort, n=7,806					
User level	0.740 (0.722-0.756)	80.7% (77.7–83.6%)	54.4% (53.2–55.5%)	$0.740 \ (0.722 - 0.756) \\ 80.7\% \ (77.7 - 83.6\%) \\ 54.4\% \ (53.2 - 55.5\%) \\ 14.5\% \ (13.3 - 15.5\%) \\ 96.7\% \ (96.2 - 97.2\%) \\ $	96.7% (96.2–97.2%)
Recording-level	0.664 (0.661-0.667)	72.8% (72.2–73.3%)	51.6% (51.4–51.8%)	$0.664 \ (0.661 - 0.667)  72.8\% \ (72.2 - 73.3\%)  51.6\% \ (51.4 - 51.8\%)  14.6\% \ (14.5 - 14.8\%)  94.3\% \ (94.2 - 94.4\%)$	94.3% (94.2–94.4%)
Clinic Cohort, n=181					
Recording-level	0.682 (0.605-0.755)	81.7% (69.2–93.1%)	53.4% (45.8–61.1%)	$0.682 \ (0.605 - 0.755)  81.7\% \ (69.2 - 93.1\%)  53.4\% \ (45.8 - 61.1\%)  31.9\% \ (22.9 - 40.7\%)  91.6\% \ (85.7 - 97.0\%)$	91.6% (85.7-97.0%)
Newly Diagnosed Diabetes, Recording-level, (n=164) 0.644 (0.546-0.744) 75.9% (56.3-92.9%) 53.0% (45.2-61.2%) 19.1% (11.2-28.3%) 93.8% (88.2-98.4%)	0.644 (0.546–0.744)	75.9% (56.3–92.9%)	53.0% (45.2–61.2%)	19.1% (11.2–28.3%)	93.8% (88.2–98.4%)

\*
Metrics are reported at a threshold of DNN score=0.427; this threshold can be altered to optimize DNN performance on specific metrics as suitable for future applications. Abbreviations: CI: Confidence Interval; PPV: Positive predictive value; NPV: Negative predictive value; AUC: Area under the ROC curve.

Table 2. Performance of logistic regression models for prediction of prevalent diabetes with and without the DNN score in the Test Dataset.

All models are shown without ("a") and with ("b") inclusion of the DNN score as a predictor. Models 1-3 are nested models, containing incrementally more demographic predictors and BMI. Model 4 adjusts for heart rate and heart rate variability. Model 5 adjusts for common cardiovascular comorbidities. Sample sizes reported indicate individual people.

Predictor	Multivariable-adjusted OR without DNN Score (95% CI)	<i>p-</i> value <sup>†</sup>	Multivariable- adjusted OR with DNN Score (95% CI)	<i>p-v</i> alue <sup>†</sup>
LogReg Model 1a: Age: AUC=0.691	(95% CI: 0.672–0.710) (n=11,3	13)	LogReg Model 1b: 1a + I (95% CI: 0.754-0.786) (n	
Age, years	1.04 (1.04–1.05)	<0.001	1.01 (1.01–1.02)	<0.001
DNN Score, per SD	-	-	2.69 (2.41-2.99)	< 0.001
<b>LogReg Model 2a: Age, gender and 1</b> (n=7,851)	race: AUC=0.698 (95% CI: 0.6	74 – 0.722)	<b>LogReg Model 2b: 2a + I</b> (0.757 – 0.798) (n=7,851)	ONN score: AUC=0.777
Age, years	1.04 (1.04–1.05)	<0.001	1.01 (1.00–1.02)	0.013
Gender				
Males	Ref.	-	Ref.	-
Females	0.99 (0.82-1.21)	0.996	0.65 (0.53-0.79)	<0.001
Race/Ethnicity		0.003		0.17
Non-Hispanic White, n (%)	Ref.	-	Ref.	-
Black or African American, n (%)	1.87 (1.11-3.15)	0.001	1.40 (0.82 - 2.38)	0.213
Hispanic, Latino or Spanish origin/ ancestry, n (%)	0.73 (0.49-1.07)	0.106	0.69 (0.46 – 1.01)	0.058
Asian, n (%)	1.86 (1.30-2.67)	0.001	1.46 (1.01 – 2.12)	0.047
Multi-ethnic, n (%)	1.27 (0.78-2.07)	0.344	1.29 (0.78 – 2.13)	0.314
Other, n (%)	0.974 (0.49-1.95)	0.941	0.86 (0.42 - 1.74)	0.674
DNN Score, per SD	-	-	2.88 (2.51-3.31)	<0.001
<b>LogReg Model 3a: Age, gender, race</b> (n=1,033)	and BMI: AUC 0.801 (95% C	I: 0.752 – 0.850)	LogReg Model 3b: 3a + I (95% CI: 0.787 – 0.873) (	
Age, years	1.04 (1.02 - 1.06)	<0.001	1.01 (0.99 – 1.04)	0.189
Males	Ref.	-	Ref.	-
Females	0.67 (0.39–1.13)	0.130	0.51 (0.30 – 0.88)	0.015
Race		0.232		0.415
Non-Hispanic White	Ref.	-	Ref.	-
Black or African American	0.33 (0.04 –2.63)	0.294	0.32 (0.04 – 2.63)	0.291
Hispanic, Latino or Spanish origin or ancestry	1.22 (0.45 –3.35)	0.696	1.08 (0.38 – 3.05)	0.884
Asian or Pacific Islander	2.82 (0.97-8.22)	0.058	2.36 (0.77 – 7.24)	0.135
Multi-ethnic	0.416(0.12-1.45)	0.168	0.46 (0.13 – 1.59)	0.218
Other/prefer not to disclose	0 (0)	0.999	0 (0)	0.999
BMI	1.15 (1.11–1.19)	<0.001	1.08 (1.04-1.12)	< 0.001
DNN Score, per SD	-	-	2.12 (1.53 – 2.94)	<0.001

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Predictor Multivariable-adjusted Multivariable*p-v*alue<sup>†</sup> OR without DNN Score adjusted OR with value<sup>†</sup> (95% CI) **DNN Score** (95% CI) LogReg Model 4a: Heart Rate and Heart Rate Variability: AUC 0.586 (95% CI: LogReg Model 4b: 4a + DNN score: AUC: 0.765 (**0.748** – **0.782**) (n=11,313) **0.565** – **0.606**) (n=11,313) 1.02 (1.01-1.02) HR, bpm < 0.001 1.01 (1.00-1.01) 0.024 HRV - RMSSD, per 10 ms 0.97 (0.94-0.99) 0.027 1.02(1.00 - 1.05)0.068 DNN Score, per SD 2.92 (2.65 - 3.21) < 0.001 **LogReg Model 5a: Comorbidities: AUC 0.784 (0.766 – 0.802)** (n=11,313) LogReg Model 5b: 5a + DNN score: AUC 0.830 (0.815 - 0.844) (n=11,313) 3.49 (2.93-4.16) < 0.001 < 0.001 Hypertension, n (%) 2.57(2.15 - 3.07)Hypercholesterolemia, n (%) 2.44 (2.05-2.89) < 0.001 1.97(1.66 - 2.34)< 0.001 0.024 1.22 (0.94 - 1.59) 0.144 Coronary artery disease, n (%) 1.35 (1.04-1.76) Prior MI, n (%) 1.04 (0.74-1.48) 0.815 1.06(0.74 - 1.50)0.765 CHF, n (%) 2.39 (1.67-3.42) < 0.001 2.09 (1.46 - 2.98) < 0.001 PVD, n (%) 1.49 (1.00 -2.21) 0.051 1.43(0.97 - 2.11)0.075 Prior Stroke, n (%) 1.91 (1.39-2.61) < 0.001 1.74 (1.27 - 2.38) 0.001 Sleep apnea n (%) 2.06 (1.72-2.46) < 0.001 1.85(1.54 - 2.22)< 0.001 < 0.001 DNN Score, per SD 2.22(2.00 - 2.46)

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Abbreviations: LogReg: logistic regression; OR: diagnostic odds-ratio; BMI: body mass index; CI: Confidence Interval; DNN: Deep neural network; SD: standard deviation; AUC: area under the receiver operating characteristic curve; HR: Heart Rate; HRV: Heart Rate Variability; RMSSD: Root Mean Square of Successive RR interval differences; y: years. Independent variables were standardized using the Z-score.

 $<sup>\</sup>dot{p}$ -value is calculated using the Wald test for the multivariable adjusted odds ratio (two-sided).