

Blood Glucose Determination by Fourier Transform near Infrared Spectroscopy

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Abstract: Diabetes is a metabolic disorder that is caused by unregulated blood glucose and therefore requires regular and intensive monitoring. Currently, blood sugar monitoring is mostly done invasively by withdrawing blood through a needle or piercing of the fingertips. This method can cause trauma and an infection. However, there is the potential for using a non-invasive measurement of blood glucose levels with near-infrared spectroscopy (NIRS) combined with partial least-square regression. As a pathway to actualize it, the spectrum of whole blood was measured with different glucose levels. A total of 72 NIR spectrum from 8 whole blood samples with different types of glucose levels were measured. A principal component analysis (PCA) and partial least square regression (PLSR) were applied to the spectral data matrix. The results showed that PCA is successfully classified as spectral data based on the glucose content and PLSR model within the clinically accurate region of the Clarke error grid. These results indicate that NIRS has an immense potential to be applied in measuring blood glucose non-invasively.

1 INTRODUCTION

Diabetes is a disease caused by a deficiency of insulin in the body (American Diabetes Association (2004). This disease can increase or decrease blood glucose levels. Under normal conditions, blood sugar levels vary from 80-130 mg/dL. Insulin is created by the pancreas to mediate metabolic reactions with blood and maintain glucose levels in a normal range (Torpy et al. 2014; Center for Disease Control and Prevention, 2016. Uncontrolled diabetes may result in various medical conditions such as a stroke, kidney failure, heart disease, and blindness (Center for Disease Control and Prevention, 2016). Recently, the number of diabetics in the world continues to increase. This increase makes it necessary to be able to detect blood glucose levels. This detection is important not only for those with diabetes but also for non-diabetes people as a part of their routine clinical monitoring. This monitoring often requires fast, painless, non-invasive, and self-

measurement methods (Ferrante et al. 2008; Kurasawa et al. 2017)

Various studies and developments of detection models have been done, such as a glucometer design to non-invasively check blood glucose by applying NIR at a single wavelength (Saleh et al. 2018). This is a promising method. Nevertheless, an assortment of other organic compounds present on the tissue can have an effect on the accuracy. Therefore, spectrum-based measurements are significant to boost the absorbance dynamics for a more thorough analysis. Yano et al. (2001) discussed the possibility of using NIR spectroscopy to simultaneously estimate glucose and citric acid in an aqueous solution of a blood anticoagulant. Zhang et al. (2014) utilized two-dimensional correlation spectroscopy (2DCS) to make the data analysis better. Essential investigations into the NIR spectrum of different organic samples have been carried out since the 1970s. The NIR glucose spectrum was also studied by Simeone et al. (2017) and Yano et al. (2001). In addition, a number of

measurement techniques have been devised like NIR Raman spectroscopy Zhang et al., 2013; Lam et al., 2010, direct diagnostics that utilize an NIR detector chip implanted under the skin Saleh et al., 2018; Uwadaira et al., 2016) and wireless long-term constant observations (Dingari et al. 2011). The NIRS approach with an exploration into different spectral ranges and additional measurement methods has also been reviewed (Pandey et al. 2017) and studied using chemometrics.

Near-infrared spectroscopy (NIRS) facilitates the workflow analysis and enables measurements of a large number of samples in a reasonably quick amount of time. It can gauge the concentration of several constituents. In numerous instances, a specific spectrum of a constituent can be connected to its fingerprint. Samples containing functional groups such as OH, CH, and NH are susceptible to NIR due to the overtones of their fundamental vibrations (R-H) in the IR region which match with the NIR absorptions. Even though the C=C and C-C bonds are not visible in the NIR region, their C-H vibrational frequencies can reveal the C=C and C-C bonds. The NIR absorption is commonly more comprehensive compared to the IR absorption because of the overlapping overtones and combination bands discussed above. Consequently, NIR analyses are complex and necessitate more detailed processes. Fortunately, the development of chemometrics enables NIR data to be utilized in appropriate processes.

This research inspected the application of NIRS to decide the glucose content in whole blood from a healthy volunteer between a range of 80 and 130 mg/dL. This study strove to elaborate on the previous measurements of glucose in an aqueous solution and examine the likelihood of using NIRS and PLSR as a substitute method to devise a non-invasive blood glucose apparatus.

2 METHODS

2.1 Sample Preparations

In this experiment, lifeblood samples were retrieved from a healthy volunteer with the intention of only focusing on the effects of glucose. All the blood drawings were completed within 120 minutes after the volunteer had finished eating and drinking sugary drinks. Blood drawings were taken in 15-minute intervals using a lancet that punctured the individual's fingertips. The drawing procedures followed standard measures using a portable

glucometer. The amount of blood drawn each time was about one drop. A small amount of blood was used to measure the blood sugar levels with a glucometer while the rest was used for scanning by NIRS. There was a total of 8 blood drawings with the glucose levels indicated by a glucometer at 84, 86, 98, 100, 111, 116, 119, and 121 mg/dL. Within 2 hours, the blood sugar levels then returned to their initial levels.

2.2 Data Acquisition

Each blood sample was put on a metal reflector covered by optical glass. The space between the glass and metal reflector was 0.2 mm (thus a 0.4 mm path length with a double pass). A Fourier transform near-infrared spectrometer (Buchi NIRFLEX 500 solid) with a spectral region of 4000-10000 cm^{-1} was applied in a reflectance mode using fiber optics to test each of the sample spectra. Each spectrum had 4 cm^{-1} intervals (thus, each spectrum consisted of 1250 data points) and averaged over 32 measurements. The sample temperatures were sustained at 26°C during the spectral acquisitions. A total of 73 spectra was collected with 9, 8, 9, 9, 11, 9, 8, and 10 spectra for the blood samples with 84, 86, 98, 100, 111, 116, 119, and 121 mg/dL of glucose, respectively.

2.3 Data Analysis

A PCA analysis was applied for the 73 spectra after the smoothing, normalizations, and derivatives. The smoothing procedure was implemented using the Savitzky-Golay method employing a third order polynomial at a frame size of 21. Spectral normalizations were applied to eliminate multiplicative scattering and baseline variations. The details for PLSR have been clarified elsewhere. A total of 73 spectra were divided into two groups, 37 spectra for the calibration set, and 36 spectra for the validation set. The calibration spectra were utilized to devise a prediction model using the partial least square regression (PLSR) method. PLSR attempts to show the relationships between groups of observed variables and latent variables. Validation spectra were applied to cross-validate them by using the PLSR parameters to estimate the concentrations of the validation samples. Both PCA and PLSR procedures were coded using Matlab version 2017b.

3 RESULTS AND DISCUSSION

Red blood cells (erythrocytes), white blood cells (leukocytes), platelets (thrombocytes), and plasma are the main constituents of whole blood (Basu and Kulkarni 2014). Nearly half of whole blood is plasma, and about 90% of it is actually water, while the remaining 10% is protein. Therefore, the NIR spectrum of whole blood resembles the NIR spectrum of water. Figure 1 shows the NIR spectra

of 27 samples, the spectrum of glucose (red) and water (blue). Water and glucose spectra are presented to indicate their bands' positions and widths to the whole blood spectra. The whole blood spectra are characterized by the well-recognized two strong water absorptions that appear at around 5200 cm^{-1} and 7000 cm^{-1} . The absorption of glucose is much weaker when compared to that of water. However, a small skirt at around 4800 cm^{-1} strongly suggests that glucose is present in the whole blood.

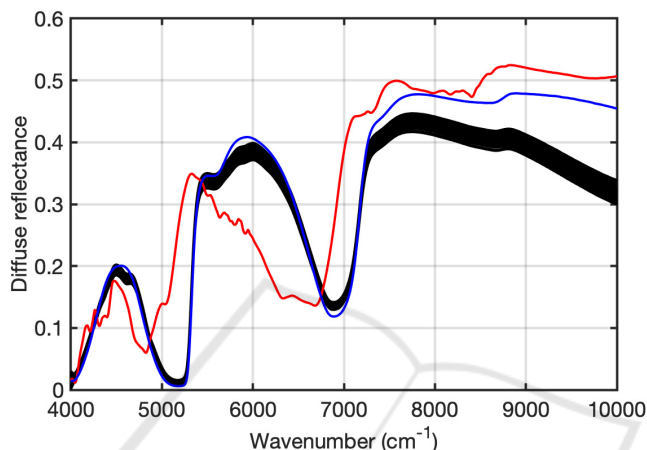


Figure 1. Near-infrared diffuse-reflectance spectra of whole blood (black). For comparison, the diffuse-reflectance spectra of powder glucose (red) and pure water (blue) are also shown. In this figure, the intensity of diffuse-reflectance spectra for powder glucose and water are unscaled.

Although the whole blood spectral patterns are similar to each other, the baseline and intensity are relatively different. They do not entirely overlap due to the baseline variations and multiplicative scattering. For PCA and PLSR analyses, each spectrum was normalized to avoid a multiplicative scattering effect and then the first derivative was

taken to correct the baseline variations. Figure 2 shows the first derivative spectra. The box with the broken line shown at 4200-5000 cm^{-1} indicates this study's target region for analyses (*vide infra*). The inset at the top right corner of the figure enlarges the spectral structure around the target region.

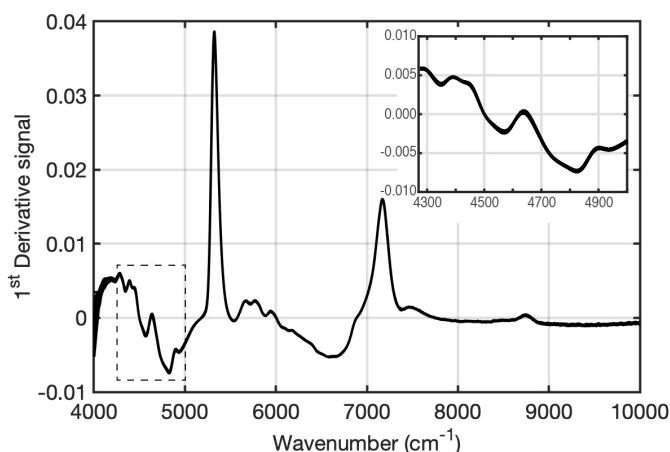


Figure 2. First derivative spectra of the whole blood. The target region for the analyses is indicated by a box with a broken line and enlarged in the inset for clarity.

A singular value decomposition (SVD) analysis of the data matrix was applied to determine the spectral region that effectively contributed to the PCA and PLSR models. It shows that the effective spectral region is 4200 - 500 cm^{-1} . The details of the PCA target spectral region of 4200 - 5000 cm^{-1} . The specified target region was chosen based on the minimum predicted residual error sum of square (PRESS) at the optimum number of latent variables in PLSR calculations. Figure 3 depicts the PCA plot

and PLSR calculations used in this paper are published elsewhere (Abdi, 2010; Tharwat (2016). The PCA and PLSR models were then calculated using a

of the 73 spectral data. Each of the eight groups has different glucose clustering at their unique pc1- pc2 -pc3 space coordinates except one spectrum of 116 mg/dL located at the 84mg/dL group coordinate.

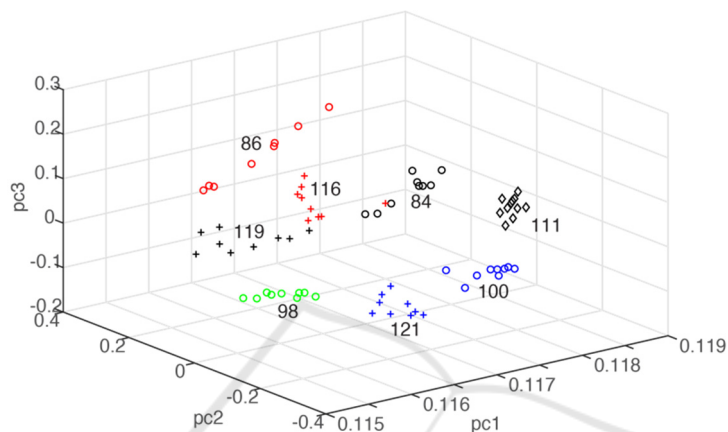


Figure 3. PCA results of seventy-three spectral data. The spectra are grouped into eight clusters based on their glucose content in whole blood. The value indicated in each group designates the glucose content in mg/dL.

In the PLSR analysis, the 73 spectral data were divided into two parts; 37 spectra were used for the calibration data set, while the remaining 36 were used for the validation data set. It was ensured that in each set, the spectrum from samples containing glucose of 84, 86, 98, 100, 111, 116, 119, and 121 mg/dL were represented in almost equal numbers. Initially, the regression parameters were calculated using a calibration data set. Finally, these parameters were employed to predict the glucose levels through the corresponding spectrum in the validation data set. The region for analysis remained the same as that used in the PCA analysis. The number of latent variables was $N=8$, by which the PRESS value reached a minimum. The latent variable obtained in this analysis was the same as that obtained for cases of glucose in an aqueous solution. Figure 4 shows the results of the NIR predictions compared to the ones measured by electrode strips (reference). The blue circles indicate the results of the NIR predictions of the glucose content for samples in the calibration data set, while the red ones represent NIR predictions for the validation data set. The coefficients of the determination (R^2) were 0.97 and 0.75 for the calibration and validation, respectively. For PRESS, the obtained values were 70.7 and 181.4

for the calibration and validation, respectively. Normally, both the R^2 and PRESS for the validation data will be smaller than that of the calibration data set, as in the present case.

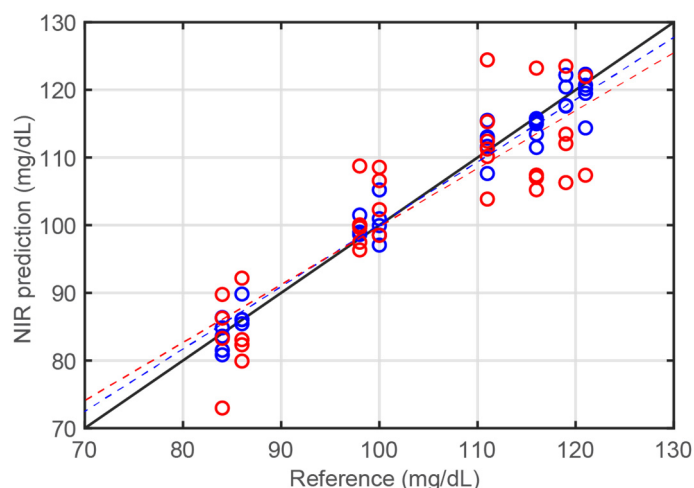


Figure 4. Comparisons between the NIR predictions and the reference values of the glucose content in whole blood. The blue and red circles represent the calibration and validation sets, respectively. The broken blue line shows the linear fitting for the calibration data set, while the broken red line shows the linear fitting for the validation data set. The solid black line is displayed for guidance purposes.

The results of the NIR prediction of glucose in the present study appear relatively scattered as though less precise when compared to the NIR predictions obtained in previous studies Rondonuwu et al., F (2019). In the previous studies, however, the glucose contents were systematically prepared at a certain level so that the reference values were highly accurate. In this study, the reference values solely relied on the strip electrode measurements using a glucometer that has relatively large random deviation values. To evaluate the NIR predictions of the validation set in terms of clinical accuracy, we applied the Clarke error grid analysis. All of the NIR prediction data points in the validation data set were

then transferred into the Clarke diagram, as indicated in Figure 5. In the Clarke diagram, region A was estimated to be clinically accurate, while region B was considered clinically acceptable. In this model, all of the 36 NIR predictions fall into region A, which means they are clinically accurate. Note that the examination range in this study is limited within 80 to 130 mg/dL, which is the range of healthy subjects. A more extensive range is necessary but requires diabetic volunteers. In this study, diabetic volunteers were not employed since they need specialized medical attention, and it is relatively challenging to promote them with extra food and beverages with high calories.

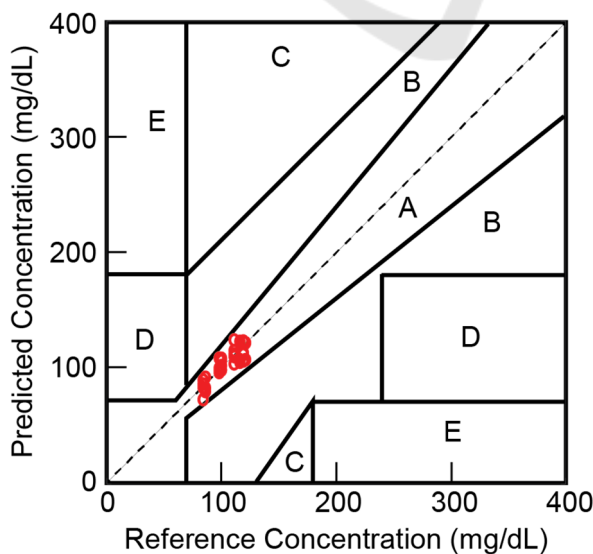


Figure 5. Clarke error grid diagram. The red circles indicate the NIR predictions of the validation data set.

In this study, we only employed a single volunteer for a short time, which means the only differential factor affecting the measurement is glucose. The rest of the constituents, including lipids, proteins, and the physical parameters such as viscosity and temperature, are practically unaltered. In a more realistic model, different volunteers having a variety of whole blood conditions must be included in a database of PLSR calculations to increase the validity and accuracy of the method.

4 CONCLUSION

The glucose content in whole blood can be determined by employing near-infrared spectroscopy followed by the partial least squares regression model in the 4200-5000 cm^{-1} spectral region. Based on the Clarke error grid, the NIR spectroscopy technique followed by a PLSR analysis from a single volunteer in the 80-130 mg/dL was successfully predicted to be clinically accurate. These results shed light on the NIRS technique followed by the PLSR calculation to provide an effective and non-invasive approach to measure blood glucose levels.

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