**Monitoring glucose levels in urine using FTIR spectroscopy combined with univariate and multivariate statistical methods**

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**Highlights**

* Urinary spectral of FTIR with univariate and multivariate statistical can use for diabetes surveillance.
* Spectral of urine analysis exhibited consistent changes related to urea, creatinine, and glucose.
* The 100% accuracy of glucose (1074 cm-1) and strong correlation with glycosuria can discriminate the urine of diabetics
* MCR-ALS analysis suggested that glucose prediction could be helpful in the screening and monitoring of diabetes

**ABSTRACT**

The development of novel platforms for non-invasive continuous glucose monitoring applied in the screening and monitoring of diabetes is crucial to improve diabetes surveillance systems. Attenuated total reflection-Fourier transform infrared (ATR-FTIR) spectroscopy of urine can be an alternative as a sustainable, label-free, fast, non-invasive, and highly sensitive analysis to detect changes in urine promoted by diabetes and insulin treatment. In this study, we used ATR-FTIR to evaluate the urinary components of non-diabetic (ND), diabetic (D), and diabetic insulin-treated (D+I) rats. As expected, insulin treatment was capable to revert changes in glycemia, 24-h urine collection volume, urine creatinine, urea, and glucose excretion promoted by diabetes. Several differences in the urine spectra of ND, D, and D+I were observed, with urea, creatinine, and glucose analytes being related to these changes. Principal components analysis (PCA) scores plots allowed for the discrimination of ND and D+I from D with an accuracy of ~99%. The PCA loadings associated with PC1 confirmed the importance of urea and glucose vibrational modes for this discrimination. Univariate analysis of second derivative spectra showed a high correlation (r: 0.865, p < 0.0001) between the height of 1074 cm-1 vibrational mode with urinary glucose concentration. In order to estimate the amount of glucose present in the infrared spectra from urine, multivariate curve resolution-alternating least square (MCR-ALS) was applied and a higher predicted concentration of glucose in the urine was observed with a correlation of 78.9% compared to urinary glucose concentration assessed using enzyme assays. In summary, ATR-FTIR combined with univariate and multivariate chemometric analyses provides an innovative, non-invasive, and sustainable approach to diabetes surveillance.

**Keywords:** Diabetes, ATR-FTIR, urinary biomarkers, chemometrics analysis

**INTRODUCTION**

Diabetes mellitus (DM) is a chronic metabolic disorder characterized by hyperglycemia related to damage in insulin secretion by pancreatic β-cells and/or reduced insulin action on target tissues [1]. It is estimated that around 537 million people worldwide live with DM [2]. Approximately 90% are affected by type 2 diabetes and 10% by type 1 diabetes [2]. This endocrine disorder is a global health issue and presents higher health costs for surveillance and treatment. Tight control of glycemic levels is critical to reduce morbidity and mortality in diabetic patients. In this context, alternative methods to assess metabolic control are critical to prevent long-term complications and to improve the quality of life in individuals with diabetes [3, 4].

Diabetes affects multiple organs, and it is the leading cause of kidney dysfunctions 6. Polyuria and changes in urinary composition including glucose, creatinine, and urea are present in people with diabetes. Diabetes treatment with optimal glycemic control is capable to suppress these diabetic urinary parameters [5, 6]. Urine is an attractive non-invasive fluid that has been frequently used to support clinical decisions for diabetic patients based on monitoring of renal function [5].

Attenuated total reflection-Fourier transform infrared (ATR-FTIR) spectroscopy is a powerful bioanalytical tool suitable to provide a sustainable, label-free, rapid, non-destructive, and relatively low-cost molecular component analysis in biofluids [7-9]. Over the past decades, research has been conducted in search of novel techniques that can monitor glucose levels in diabetic patients in a rapid, cost-effective, and non-invasive manner. Furthermore, FTIR technology is also available as portable devices, which is ideal for point-of-care medicine. ATR-FTIR is capable of measuring the specific vibrational modes resulting from carbohydrates, proteins, lipids, electrolytes, DNA, and RNA simultaneously and without extensive sample preparation [7, 9]. The discrimination of healthy and pathological conditions, as well as the efficacy of treatments, has been widely performed by ATR-FTIR [10]. Urine-based spectroscopy infrared (IR) bands have already been applied successfully in several studies, for microproteinuria [11], bacteria causing urinary tract infections [12], endometrial and ovarian cancers [13, 14], prostate and urogenital cancer [15, 16], drug-induced crystalluria [17], primary hyperoxaluria [18], and many other diseases [19].

In order to open new avenues for the use of infrared spectroscopy on urine as a novel approach for personalized medicine, we hypothesized that ATR-FTIR is capable of identifying changes in urinary molecules altered by diabetes and thus reverted by appropriate insulin treatment. Therefore, the present study aims to demonstrate the ability of ATR-FTIR spectroscopy combined with univariate and multivariate curve resolution-alternating least square (MCR-ALS) analysis to monitor urinary parameters of non-diabetic and diabetic rats, as well as insulin-treated diabetic animals.

**MATERIAL AND METHODS**

**Animal experiment**

All experiments were conducted following recommendations of the Brazilian Society of Laboratory Animals Science (SBCAL) in the guide for the care and use of laboratory animals. Experimental procedures for the handling, use, and euthanasia were approved by the Ethics Committee for Animal Research of the Federal University of Uberlandia (UFU) (License #CEUA-UFU No. 013/2016) according to Ethical Principles adopted by the Brazilian College of Animal Experimentation (COBEA) and conformed to ARRIVE guidelines.

Male Wistar rats (~260 g) were provided from the Center for Bioterism and Experimentation (REBIR) at the Federal University of Uberlandia. The animals were maintained under standard conditions (12-hour light/dark cycles, light on at 7 a.m.; 22 ± 2 °C; humidity ~60%) and were allowed free access to water and a standard diet within this rodent facility.

Overnight-fasted animals received a single intraperitoneal injection (60 mg/kg) of streptozotocin (STZ) (Sigma-Aldrich, St. Louis, MO. USA) dissolved in 0.1 M citrate buffer (pH 4.5) to induce diabetes. Animals with hyperglycemia (>250 mg/dL) 48 h later were considered diabetics (D). Non-diabetic (ND) animals received injection of 0.9% NaCl (physiological saline) in similar volumes (n = 8). Thus, diabetic animals were divided in placebo-treated Diabetic (D, n = 10) and Diabetic treated with 6 U Insulin (D+I, n = 10). Subsequently, 21 days after diabetes induction, diabetic rats received treatment with vehicle (NaCl 0.9% in D rats) or with 6 U of insulin [NPH insulin, Biobrás, MG, Brazil] (D+I) per day (2 U at 8:30 a.m. and 4U at 5:30 p.m.) subcutaneously in D+I rats. ND animals also received vehicle (0.9% NaCl) [20-22].

**Urine sample collection and preparation**

On day 6 of treatment, the rats were maintained for 24-hours in metabolic cages to allow for 24 h-urine collection from each animal separately. The samples were collected, measured, processed, and stored at -80ºC until further analysis. Creatinine, urea, and glucose concentration in the urine were measured using enzymatic assays following the manufacturer's instructions (Labtest Diagnostica SA, Brazil). Blood glucose levels obtained from the tail vein were measured in overnight-fasted animals using reactive strips for glucometer (Accu-Chek Performa, Roche Diagnostic Systems, Basel, Switzerland). Besides that, the variation of gain/loss body weight (Δ body weight) after STZ or placebo administration was analysed. Animals were euthanized with an excessive anesthetic dose immediately after sample collections.

After that, urine samples were lyophilized (L101, Liotop®) and the samples were analysed after drying.

**FTIR spectroscopy and data analysis**

Infrared spectra were acquired using an FTIR system (Nicolet 6700, Thermo Scientific, Waltham, MA, USA) in Attenuated Total Reflectance (ATR) mode over the range 4000-400 cm-1. Spectra were recorded with a spectral resolution of 4 cm-1 and 100 scans (co-adds) per spectrum. FTIR spectra were smoothed via Savitzky-Golay filter (polynomial of second order in an eleven-point window), followed by rubberband baseline correction, and vector normalized prior to spectra data analysis. The fingerprint region (1800-900 cm-1) was used as the input to PCA and MCR-ALS techniques [23]. All pre-processing steps and spectral analyses were performed using Matlab® R2019b (MathWorks, Natick, MA, USA) and all computational algorithms are available in Github (https://github.com/Biospec/). In addition, second derivative spectra were obtained by applying Savitzky-Golay algorithm with polynomial order 5 and 20 points of the window.

**Glucose, creatinine, and urea and sample preparation**

Glucose, creatinine and urea (>99.0%) were purchased from Sigma-Aldrich (Sigma-Aldrich Dorset, UK). The experiments to analyze the infrared spectra of these molecules, were conducted in a similar protocol as described to urine. Aqueous solutions of glucose (800 mM), creatinine (500 mM), and urea (850 mM) were transferred onto ATR crystal and left to dry at room temperature before data collection.

**Statistical Analysis**

The normality of data distribution was analyzed using the Kolmogorov-Smirnov test. The amplitude of second derivatives were analyzed using one-way analysis of variance (ANOVA), followed by Tukey Multiple Comparison as a post-hoc test. Receiver Operating Characteristic (ROC) curve and Pearson correlation test were applied for all second derivative peaks intensity. All these analyses were performed using the software GraphPad Prism (GraphPad Prism version 7.00 for Windows, GraphPad Software, San Diego, CA, USA). Only values of p < 0.05 were considered significant and the results were expressed as mean ± S.D.

**RESULTS and DISCUSSION**

The DM and insulin treatment effects on animals were characterized by several parameters, which are shown in Table 1. As expected, a reduction in weight gain (p < 0.05), an increase in plasmatic glycemic levels (p < 0.05) and higher 24h-urinary volume (p < 0.05) were observed in D compared with ND rats. By contrast, no change in the concentration of urinary creatinine was observed; however, 24 h urinary creatinine excretion was increased (p < 0.05) in D compared to ND. In addition, the urinary urea concentration decreased (p < 0.05) in D compared to ND, and its level in 24 h urinary urea excretion increased (p < 0.05) in D compared to ND rats. Both glucose concentration in urine and 24 h urinary glucose excretion were also higher (p < 0.05) in D compared to ND.

Insulin treatment resulted in weight gain (p < 0.05) with concomitant reduction in plasma glucose levels (p < 0.05) and urinary volume (p < 0.05) compared to D rats. Creatinine levels in urine did not change during insulin treatment; however, 24 h urinary creatinine excretion decreased (p < 0.05) in D+I compared to D rats. Regarding urea concentration in urine, insulin treatment was not efficient to restore (p > 0.05) this parameter. On the other hand, insulin treatment reduced (p < 0.05) 24 h urinary urea excretion compared to D rats. As expected, the urinary glucose concentration and 24 h urinary glucose excretion were lower (p < 0.05) in D+I compared to D rats (Table 1). Bearing in mind the classical effects of insulin treatment on urinary parameters, diabetic animals treated with insulin or placebo showed similar metabolic profiles as described in previous studies, in which urinary parameters were changed by diabetes induction, and those parameters were restored to normoglycemic profile with insulin treatment [20, 24-26].

Infrared spectra from urine collected using ATR-FTIR are complex signatures and these biochemical fingerprints are related to the many infrared-active molecules within this complex biological mixture. Water is the main component of urine (over 95%) and due to its strong absorption bands in the mid-infrared region, the samples must be dry before ATR-FTIR analysis to generate signals from the signatures of molecules in lower concentrations [9]. Figure 1 shows the fingerprint region (1800-900 cm-1) of spectra acquired from lyophilized urine of non-diabetic and diabetic rats, as well as, the spectra of pure glucose, creatinine, and urea.

The spectral signatures observed in both urinary spectra in non-diabetic and diabetic conditions are linear combinations of the individual spectra from the individual urinary constituents. Urea is an important component in urine (Table 1) and other constituents include creatinine, uric acid, organic and inorganic salts, proteins, hormones, and other metabolites. Spectra of urea and creatinine are also shown in Figure 1 due to their abundance in urine as well as their clinical relevance. Urea is a relatively small molecule with the molecular formula of NH2CONH2, it is therefore not surprising that the IR spectrum of urea is relatively simple and has distinct bands at 1149 cm-1 (NH rocking vibrations), 1459 cm-1 (antisymmetric CN stretch), 1587 cm-1 (CO stretch), 1620 cm-1 (antisymmetric NH bend), and 1671 cm-1 (symmetric NH bend) [27]. The IR spectrum of creatinine (C4H7N3O) depicts bands peaking at 1490 cm-1 (C=N stretch, CN stretch, NCH bend) and three bands between 1500-1700 cm-1 [28]. It is important to notice that urea and creatinine molecules strongly interact with water due to hydrogen bonding with N–H, therefore the IR spectra of both compounds are different in aqueous solution compared to fully dried states. Oliver, Maréchal and Rich [29] demonstrated that the antisymmetric C–N stretch (νas(CN)) vibration from urea, which is found at 1468 cm-1 in aqueous solution, shifts to 1464 cm-1 when urea is completely dry. However, two additional states of urea with νas(CN) peaks at 1454 cm-1 or 1443 cm-1 were also observed in partially hydrated samples [29]. Thus, IR spectra interpretation of such molecules must be performed with caution since the intermediate forms due to different degrees of hydration may be present when whole urine is dried. According to Figure 1, bands from 1700 cm-1 to 1400 cm-1 in spectra collected from urine of both healthy and diabetic rats are mainly due to urea and creatinine.

The exact constitution of urine changes with lifestyle and health status for each individual. Hence, some diseases can result in elevated levels of certain urine components. It is well known that diabetic patients present elevated glucose levels in urine (glycosuria). IR signatures of glucose are also shown in Figure 1 and these vibrations spectral features are mainly dominated by C-O and C-O-C stretching resulting in peaks in the low wavenumber region (900-1200 cm-1) [30, 31]. The low wavenumber region of spectra from diabetic rats is dominated by vibrations related to glucose, which indicates elevated glycosuria. These features are very small in infrared spectra collected from healthy non-diabetic rats (Figure 1).

Spectra from healthy and non-diabetic animals were subjected to PCA in order to evaluate any specific separation related to the development of diabetes. The resultant PCA scores plot (Figure 2A) displayed a clear separation between spectral data acquired from healthy and diabetic animals. Scores related to healthy rats were grouped on the negative side of the first principal component (PC-1) axis, while diabetic animals were clustered on the positive side. PC-1 which accounts for 98% of the total explained variance (TEV), retains information responsible for the clustering pattern obtained in the scores plot. The corresponding PC-1 loadings are illustrated in Figure 2B. Positive loadings on bands peaking at the low wavenumber region indicate the high glucose levels in urine of diabetic rats. On the other hand, negative loadings were observed for bands associated with other urinary constituents, indicating higher urea and creatinine content in urine of non-diabetic animals.

Figure 3 illustrates spectra depicted for each D+I animal (n = 10), as well as averaged spectra of signatures acquired from all D and ND rats. Most spectra from insulin-treated diabetic animals (90 %) presented glucose bands at levels comparable to healthy animals, indicating a positive response to the treatment (D+I in green color). However, the spectrum from one D+I animal with reduced response to insulin treatment presented bands indicating high glucose content (D+I in orange color), indicating a profile similar to urine from non-treated diabetic animals. Similar findings were evidenced by PCA applied to all datasets in ND, D, and D+I rats. PCA scores related to D+I animals were grouped on the same cluster of ND animals, indicating a positive response to insulin (Figure 2C). On the other hand, PCA scores associated with the animal with high urinary glucose content were grouped within the cluster containing scores from untreated diabetic animals. PC-1 loadings are shown in Figure 2D and illustrate similar findings to PC-1 loadings depicted in Figure 2C. These findings indicate the ability of urinary ATR-FTIR spectroscopy as a rapid screening platform for diabetes as well as a tool to evaluate the metabolic response of diabetic patients to insulin treatment.

We also calculated the second derivative of ATR-FTIR spectrum to amplify the spectral variations of the urinary components glucose, creatinine, and urea (Figure 4A) and compared these with ND, D, and D+I rats (Figure 4B).

The second derivative analysis is capable to enhance the separation of overlapping vibrational modes. Relative changes in the amplitude of second derivative are related to the presence of each component. In Figure4.A the amplitude of second derivatives for urea band peaking at 1452 cm-1 were reduced (p < 0.05) in the urine of D compared to ND rats, and the treatment with insulin was effective and caused increase (p < 0.05) in this amplitude compared to D rats. Considering that sensitivity and specificity are basic characteristics to determine the accuracy of diagnostic, triage, and monitoring tests [32-34], receiver operating characteristic (ROC) curve analyses were used to evaluate the potential of these spectral vibrational modes. In this context, we compared hyperglycemic (D) rats to normoglycemic ND and D+I, thus removing the sample of D+I due to the glycemia of 439 mg/dL representing a diabetic metabolic profile. The area under the curve (AUC) of ROC analysis based on amplitude of second derivative of the band peaking at 1452 cm-1 was 0.905 (p < 0.0005), and the best discrimination threshold value was 0.05282 with a sensitivity of 100% and specificity of 88.2%. However, no correlation was observed between these amplitude values and urinary urea concentrations (r = -0.051, p = 0.798) (Figure 5A). In figure 5B the second derivative amplitudes of creatinine at 1490 cm-1 were reduced (p < 0.05) in the urine of D compared to ND rats, and the treatment with insulin increased (p < 0.05) this amplitude compared to D rats. The AUC of ROC analysis based on amplitude at 1490 cm-1 in the second derivative was 0.958 (p < 0.0001), and the cut-off value was 0.01105 with a sensitivity of 100% and specificity of 88.2%. Again, no correlation was observed between the amplitudes at 1490 cm-1 in second derivative with urinary creatinine (r = -0.167, p = 0.405) (Figure 5B). Finally, in Figure 5C the urinary amplitude of glucose at 1074 cm-1 was increased (p < 0.05) in D compared to ND rats, and the treatment with insulin effectively reduced (p < 0.05) this amplitude compared to D rats. The area under the curve (AUC) of ROC analysis based on amplitudes at 1490 cm-1 in the second derivative was 1.0 (p < 0.0001), and the selected discrimination threshold value was 0.125 with both sensitivity and specificity of 100%. In this instance, the analysis of 1490 cm-1 amplitudes with glycosuria showed a strong and positive correlation (r=0.865; p<0.0001) (Figure 5.C) indicating that the main change in the urine between ND and R rats was the level of glucose, which could be recovered in D+I when the rats were treated with insulin.

MCR-ALS is a popular modelling method used to deconvolve spectra from mixtures into their individual components. Thus, MCR-ALS was applied to infrared urinary spectra of ND, D, and D+I animals in order to estimate the relative concentrations of creatinine, urea, and glucose from the whole urine spectrum. MCR-ALS assumes that the recorded spectra are the weighted sum of pure spectra of the components present in the investigated mixture and returns the resolved spectrum of each compound as well as its concentration in the mixture spectrum. Figure 6A shows a good correlation of FTIR spectrum of laboratory-grade commercial glucose (blue line) to the resolved (pure) spectrum obtained through MCR-ALS. Thus, this approach can be applied to estimate the relative concentration of glucose in urine spectra. Figure 6B shows a scatter plot of glucose concentration estimated by MCR-ALS plotted against glucose levels in urine obtained via enzymatic kits. Linear regression showed a positive correlation between glucose levels obtained through MCR-ALS and enzymatic kits (R2 = 0.7893). By contrast, MCR-ALS was not able to reproduce the signatures of urea and creatinine (data not shown), which may have occurred due to the different hydration degree of spectral signatures obtained from pure compounds compared to signatures of both molecules in urine.

Glucose and other monosaccharides absorb are detected at 1200–900 cm -1 and the absorption bands for many of these compounds are C–H stretches and bonds, O–H bonds, and C–C stretches which can overlap [35, 36]. Although the spectra of these monosaccharides are very similar with glucose, some differences in the absorption frequency occur between glucose and with fructose, galactose, mannose, as previously described [36]. Therefore, our data therefore suggest that the vibrational modes at 1074 cm-1 may correspond to the more specific absorption of glucose compared to some model compound spectra.

Although the present study identified novel urinary infrared spectral biomarkers for screening and monitoring metabolic control of diabetes, further studies are needed to confirm these potential urinary biomarkers in humans and to evaluate the suitableness of ATR-FTIR platforms for the screening and monitoring diabetes in human urine. Another potential limitation of the present study is the use of diabetic animals which display higher levels of blood glucose concentration, which is a typical outcome in streptozotocin-induced diabetic rats. Another limitation of urine samples are the changes in urinary components during the day due to changes in hydric behavior and glycemic variation, which can restrict the detection of acute metabolic modulation. We believe that these variations can be captured within the multivariate models and compensate the changes in urinary components.

FTIR spectroscopy is an important analytical tool for sample classification and biomarker detection and identification, however the applicability of this platform to the analysis of biological materials with highly complex chemical constituents has been improved with novel analysis. However, recent advances in the ATR-FTIR coupled with univariate, multivariate and learning machine analysis have greatly enhanced the capability of this platform to discriminate different types of biological samples with or without diseases [7-9, 37]. As a result, an increasing number of studies are investigating the application of infrared spectroscopy in the screening and diagnosis of several diseases [7, 10, 15, 19]. The lack of sensitive and specific methods for early detection and detection of various diseases has led to the search for new approaches. FTIR spectroscopy is simpler, faster, more accurate, less expensive and non-destructive compared to existing screening, diagnostic and monitoring methods for detecting biochemical changes [19]. Therefore, could improve clinical decision-making and patient outcomes at the molecular level in a near future.

**CONCLUSION**

In this study, ATR-FTIR spectroscopy was used to evaluate the urinary spectral differences between non-diabetics and insulin-treated diabetic than diabetic rats. The ATR-FTIR mean spectral analysis exhibited consistent changes related to urea, creatinine, and glucose from ND and D+I than D rats. PCA demonstrated discrimination between the ATR-FTIR spectra of urine with 99% of the cumulative variance of ND compared to D, and 98.3% comparing all groups, including D+I. PCA loadings associated to PC1 indicated urine glucose as one of the components for this discrimination between groups. It was reinforced by the 100% of accuracy to discriminate ND and D+I than D rats using the amplitude of second derivatives of glucose at 1074 cm-1 as well as by the strong correlation (r = 0.865; p < 0.0001) between this vibrational mode with glycosuria. MCR-ALS analyses were also efficient in predicting glucose in the urine with a high relation between glucose in urine by enzymatic kit.

In summary, these results indicate that ATR-FTIR spectroscopy with univariate or multivariate analysis has the potential to provide information about the status of glucose in urine as a novel noninvasive approach to diabetes screening and monitoring.

**ACKNOWLEDGEMENTS and FUNDING**

We would like to thank our collaborators at the Rodent Vivarium Network (REBIR-UFU). The funders had no role in study design, data collection, and analysis, decision to publish, or preparation of the manuscript. This research was supported by a grant from CAPES/CNPq (#458143/2014), CNPq INCT 465763/2014-6, FAPEMIG (#APQ-02872-16), Federal University of Uberlandia and National Institute of Science and Technology in Theranostics and Nanobiotechnology (CNPq Process N.: 465669/2014-0). Caixeta, D.C. received a fellowship from FAPEMIG, Sabino-Silva, R and Lima, C.A. received a fellowship from PrInt CAPES/UFU and CNPq 141629/2015-0, respectively. Zezell, D.M. and Sabino-Silva, R also received a productivity fellowship CNPq.

**CONFLICTS OF INTEREST**

The authors declare that they have no conflicts of interest.

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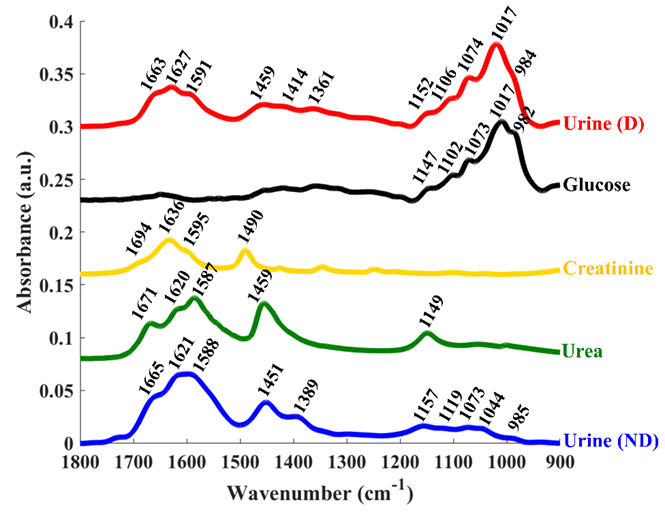
**TABLE**

**Table 1.** Effect of diabetes and insulin on body weight, glycemia, 24 h urinary volume, urinary creatinine concentration, 24 h urinary creatinine excretion, urinary urea concentration, 24 h urinary urea excretion, glycosuria and 24 h urinary glucose excretion.

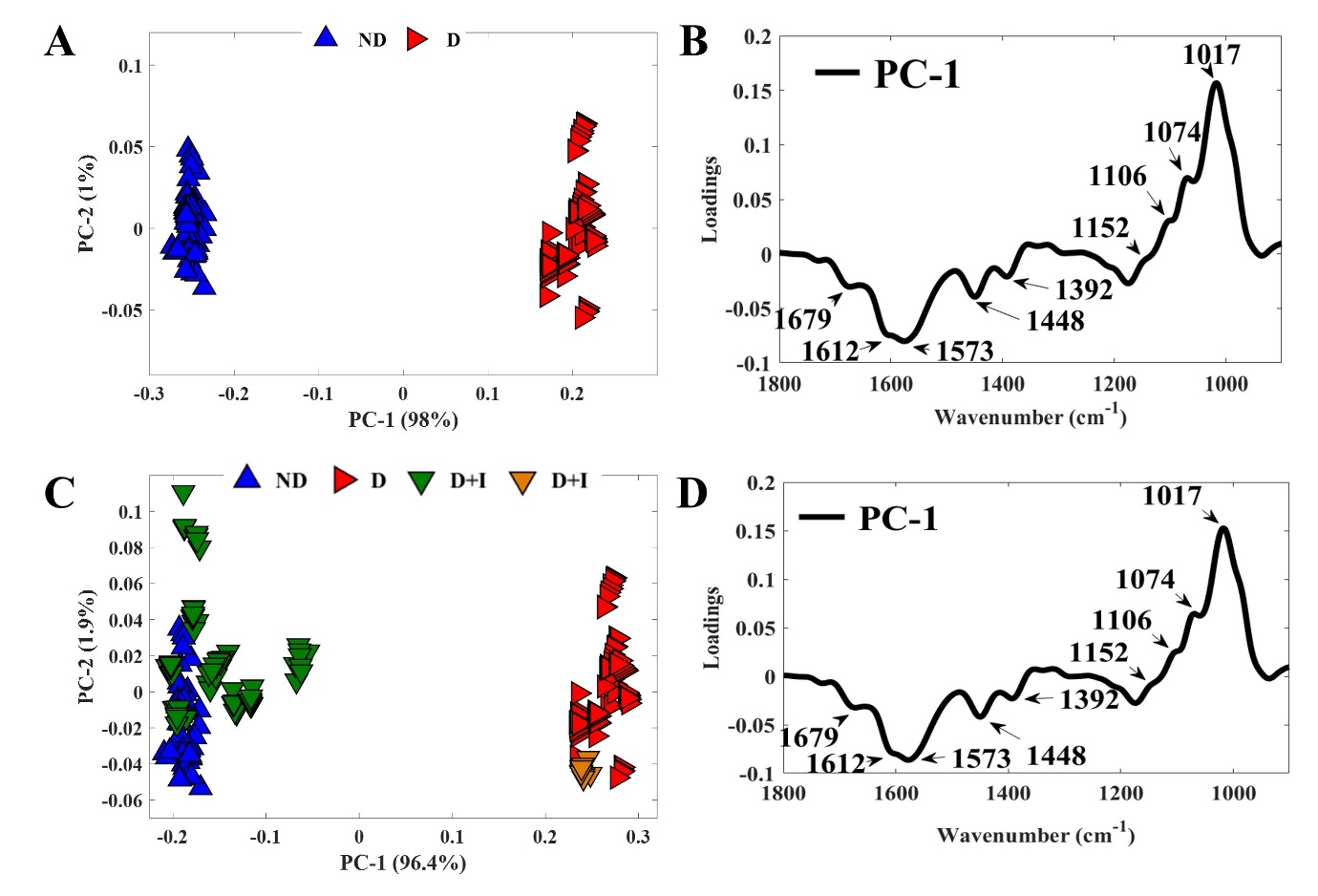
|  |  |  |  |
| --- | --- | --- | --- |
| **Parameters** | **ND** | **D** | **D+I** |
| **Δ Body weight (g)** | 51.6±24.9 | -20.1±37.4\* | 23.2±31.5# |
| **Glycemia (mg/dL)** | 83.7±13.4 | 483.8±59.9\* | 102.4±100.8# |
| **24 h urinary volume (mL)** | 23.2±10.4 | 124.2±31.0\* | 35.1±12.7# |
| **Urinary creatinine concentration (md/dL)** | 0.52±0.15 | 0.62±0.28 | 0.65±0.33 |
| **24 h urinary creatinine excretion (mg/24h)** | 12.6±8.2 | 76.7±40.0\* | 20.7±8.1# |
| **Urinary urea concentration (md/dL)** | 94.6±56.0 | 35.1±9.0\* | 38.4±15.5\* |
| **24 h urinary urea excretion (mg/24h)** | 20.9±14.9 | 44.0±16.6\* | 12.5±5.4# |
| **Glycosuria (mg/dL)** | 24.7±20.3 | 367.8±62.8\* | 165.5±181.3\*# |
| **24 h urinary glucose excretion (mg/24h)** | 4.6±2.6 | 461.3±173.4\* | 51.9±58.3# |

Values are expressed as mean ± S.D. \**p* < 0. 05 vs ND rats; #*p* < 0. 05 vs. D rats. Non-diabetic rats (ND), Diabetic rats (D) and Diabetic rats with insulin treatment (D+I).

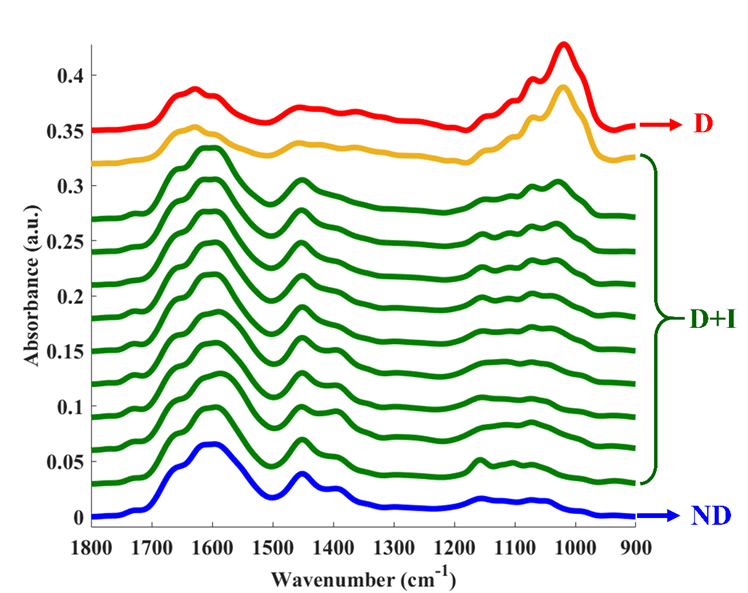
**FIGURES**



**Figure 1.** Fingerprint region (1800-900 cm-1) of typical infrared spectra of urine collected from healthy (ND, blue), diabetic (D, red), along with the spectra from pure urea (green), creatinine (yellow), and glucose (black). Peaks are highlighted which are discussed in the text.



**Figure 2.** PCA scores plots (A) and PC-1 loadings plots (C) obtained by applying PCA to FTIR spectra of urine acquired from healthy (ND) and diabetic (D) animals. PCA scores (C) and loadings (D) results obtained for FTIR spectra of urine acquired from healthy (ND), diabetic (D), and insulin-treated diabetic (D+I) rats. One D+I animal with glycemia of 439 mg/dL is highlighted in orange due to reduced effectiveness of insulin treatment on the diabetic metabolic profile.



**Figure 3.** Averaged FTIR spectra collected from the urine of each insulin-treated diabetic animal, as well as averaged spectra of signatures acquired from all diabetic (red) and non-diabetic rats (blue). Green lines relate to animals with positive response to insulin-treatment, while the orange line represents the animal that displayed no response to insulin treatment.



**1490 cm-1**

**1459 cm-1**

**1073 cm-1**

**1452 cm-1**

**1490 cm-1**

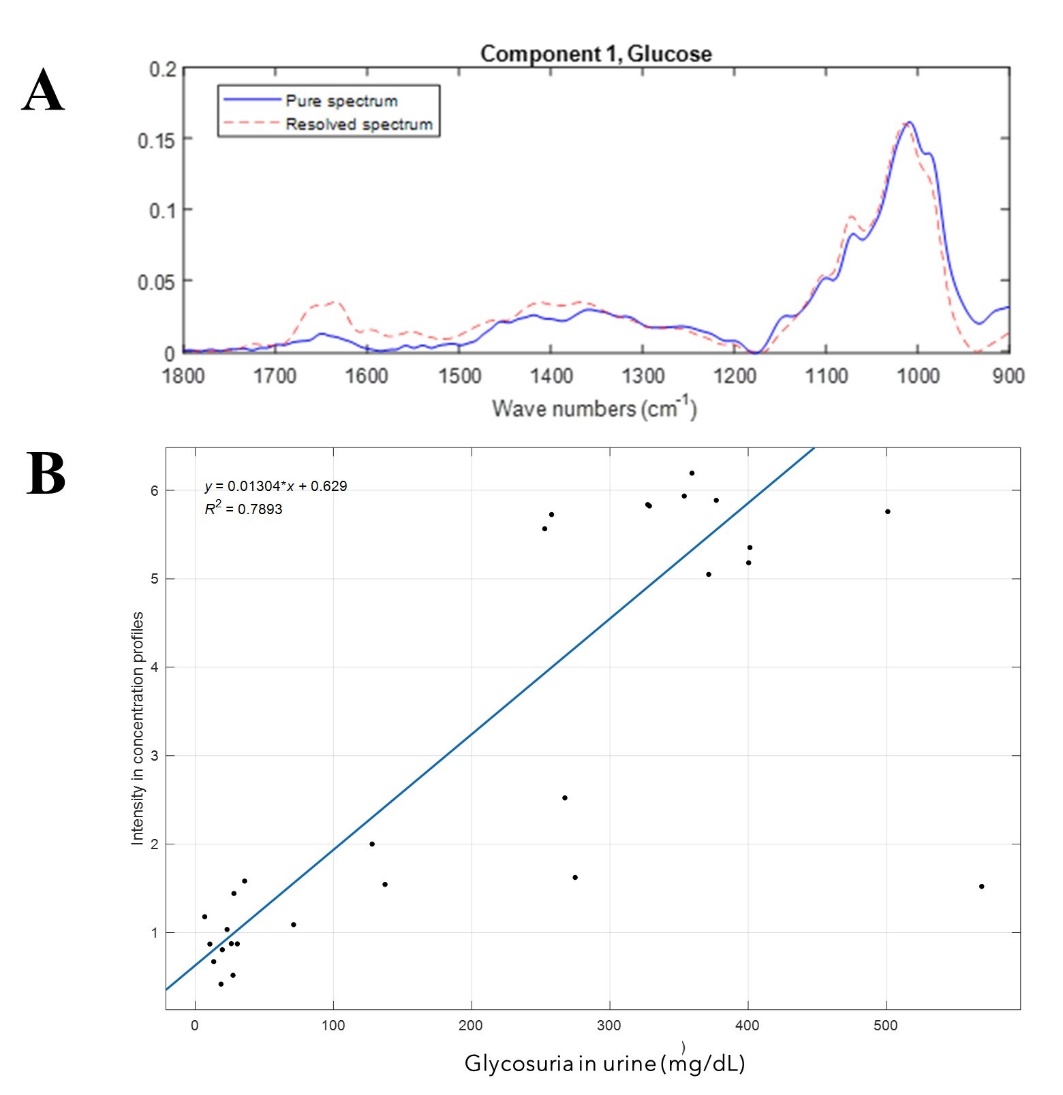
**1074 cm-1**

**A**

**B**

**Figure 4.** Fingerprint region (1800-900 cm-1) of urinary spectra from glucose (black), creatinine (yellow) and urea (green) (A) and non-diabetic (ND, blue), diabetic (D, red) and insulin-treated diabetic (D+I) rats (B).

**Figure 5.** Analyses of three key peaks seen in urea (1452 cm-1), creatinine (1490 cm-1) and glucose (1074 cm-1). For each set the amplitude of second derivative vibrational mode is plotted for 1452 cm-1 (A), 1490 cm-1 (B) and 1074 cm-1(C), along with the ROC curve analysis to discriminate ND and D+I from D rats, and Pearson correlations between: urinary creatinine and amplitude of 1452 cm-1 (A); urinary creatinine and amplitude of 1490 cm-1 (B); and glycosuria and amplitude of 1490 cm-1.



**Figure 6.** FTIR spectra of laboratory-grade commercial glucose (blue line) and resolved spectrum obtained through MCR-ALS (dashed red line) (A). Scatter plot of glucose concentration estimated by MCR-ALS plotted against glycosuria obtained via enzymatic kits (B).