The Clinical Utility of Salivary Biomarkers in the Identification of Type 2 Diabetes Risk and Metabolic Syndrome

Abstract: Type 2 diabetes is traditionally diagnosed by the use of an oral glucose tolerance test and/or HbA1c, both of which require serum collection. Various biomarkers, which are measurable biological substances that provide clinical insight on disease state, have also been effective in the early identification and risk prediction of inflammatory diseases. Measuring biomarker concentrations has traditionally been obtained through serum collection as well. However, numerous biomarkers are detectable in saliva. Salivary analysis has more recently been introduced into research as a potential non-invasive, cost-effective diagnostic for the early identification of type 2 diabetes risk in adults and youth. Therefore, the purpose of this review was to compare 6 established inflammatory biomarkers of type 2 diabetes, in serum and saliva, and determine if similar diagnostic effectiveness is seen in saliva. A lack of standardized salivary analysis, processing, and collection accounts for errors and inconsistencies in conclusive data amongst studies. Proposing a national standardization in salivary analysis, coupled with increased data and research on the utility of saliva as a diagnostic, poses the potential for salivary analysis to be used in diagnostic settings.

Keywords: saliva, inflammation, type 2 diabetes, biomarkers, metabolic syndrome, oxidative stress

Introduction

Type 2 diabetes is a chronic metabolic disorder characterized by a progressive loss of β-cell function relative to insulin resistance, resulting in persistent hyperglycemia.1–4 A complex interplay between genetics and environment attribute to the development of type 2 diabetes, with increased adiposity, energy-dense diet, physical inactivity, and low socioeconomic status being risk factors.5 Development of insulin resistance is associated with low-grade inflammation, which can be induced by oxidative stress mediators and pro-inflammatory mediators such as cytokines and adipokines.6,7 Furthermore, altered inflammatory cytokine profiles are commonly reported in diabetic patients, suggesting that heightened inflammation plays a vital role in the pathophysiology of type 2 diabetes.7–9

The global and national prevalence of type 2 diabetes is substantial. In 2019, 463 million cases of diabetes were estimated globally,10 with studies suggesting that by the year 2025–2030 incidence of type 2 diabetes cases will increase an additional 130–200 million cases globally.11–13 According to the CDC, more than 34 million Americans have diabetes, with 90–95% being type 2 diabetes.14 Moreover, the CDC’s most recent data show approximately 71.6% of adults over...
20 years old in the United States are overweight or obese, and 20.6% of youth age 12–19 are obese, putting them at higher risk for type 2 diabetes (https://www.cdc.gov/obesity/data/index.html). In parallel with the widespread pediatric obesity epidemic, type 2 diabetes, once thought to be an adult disease, is increasing evidently in obese youth. In recent years, as many as 5000 new cases of youth type 2 diabetes are reported per year in the United States, with disproportionate burden on racial and ethnic minorities. Additionally, mortality is a major cause of concern for individuals diagnosed with diabetes. Per the CDC, type 2 diabetes was the seventh leading cause of death in the United States in 2019, with 83,564 total deaths (https://www.cdc.gov/nchs/data/nvsr/nvsr68/nvsr68_09-508.pdf). Likewise, the economic burden of diabetes is massive. In 2017, diagnosed diabetes of all ages, undiagnosed diabetes and prediabetes in adults, and gestational diabetes treatment reached a combined $404 billion in the United States, totaling an annual burden of $13,240 per diagnosed patient. Given its high mortality, cost, and prevalence, type 2 diabetes carries a significant burden on the healthcare system, with an urgent need for expanded treatment and early identification of individuals at high-risk.

Interestingly, biomarkers provide great contributions in understanding the etiology of diseases, monitoring treatment, and providing information with respect to risk prediction, disease prevention, and early intervention. In recent years, biomarkers have been a prospective area of research, specifically, novel plasma and serum biomarkers of type 2 diabetes. Numerous biomarkers have displayed strong, positive correlations to type 2 diabetes, and are now recognized as risk factors. As such, these parameters have been obtained exclusively through drawing blood. Given the invasive nature of needles, along with a potential blood-injection-injury phobia in patients, and especially children, this introduces the question of whether there are less-invasive measures that can be introduced in clinical settings in order to obtain biomarker concentrations.

More recently, saliva sampling has been introduced into research as a potential indicator of type 2 diabetes risk. Numerous molecular components are detectable in saliva; however, its effectiveness and utility in clinical and diagnostic settings is not well known. Thus, the purpose of this review was to compare several established biomarkers of type 2 diabetes, in serum and saliva, and determine if similar effectiveness is seen in saliva, including adiponectin, leptin, C-reactive protein (CRP), interleukin-6 (IL-6), tumor necrosis factor-alpha (TNF-α), and 1,5-anhydroglucitol (1,5-AG).

Methods

Search Strategy

This scientific literature review conducted until May 2020 was executed using the following databases: Pubmed, Google Scholar, Science Direct, Proquest, Elsevier, Scopus, and Embase. The literature search was limited to papers in English language with defined data collection and analysis. The reference section of the articles included were further screened for relevant additional sources. A total of six biomarkers were selected based on relevance in research related to type 2 diabetes and metabolic syndrome (Figure 1). The primary search strategy was conducted using the following terms: saliva, biomarkers, type 2 diabetes, adiponectin, leptin, C-reactive protein, interleukin-6, tumor necrosis factor-alpha, 1,5-anhydroglucitol, and inflammation.

Saliva

Saliva is an antimicrobial oral fluid produced and secreted by salivary glands that contains a variety of molecular components, such as water, proteins, enzymes, hormones, growth factors, metabolites, immunoglobulins, vitamins, and electrolytes. Saliva has a variety of functions, including lubrication of the oral cavity, binding of masticated food, initiation of digestion through starch breakdown by salivary amylase, and oral hygiene. To date, standardization of saliva collection and analysis has not been introduced. Generally, unstimulated whole saliva (UWS) is collected, centrifuged, and stored at −80°C until analyzation using an enzyme-linked immunosorbent assay (ELISA) test or commercial enzyme immunoassay kit.

Prior to our review of six inflammatory biomarkers, we have acknowledged any possibilities regarding the limitations of salivary biomarkers that could be affected by dental health. A total of 10 previous literatures that compared salivary samples and blood biomarkers of inflammation were mainly included in this review (Table 1); however, only two studies addressed oral health by including participants with periodontitis while the remaining studies did not. It is noted that saliva plays a role in the maintenance of dental health to prevent dental caries. In other words, poor dental health often results from
reduced salivary flow. Collectively, poor dental health may potentially affect salivary composition, thereby suggesting that any future research involving salivary collection should address the dental health status of the participants.

Review of Six Inflammatory Biomarkers

Adiponectin

Adiponectin is an adipose-derived plasma protein strictly secreted by adipocytes. It is an anti-inflammatory adipokine that functions in sensitizing insulin action and regulating glucose and lipid metabolism. In healthy individuals, plasma adiponectin is abundant in circulation, accounting for approximately 0.01% of total plasma protein. Adiponectin is anti-diabetic, as it increases insulin sensitivity by inhibiting hepatic glucose production and promoting fatty acid oxidation. Additionally, it is considered an anti-atherogenic hormone, due to its suppression of atherosclerotic events and promotion of vasodilation. Low plasma concentrations of adiponectin have notably been reported in overweight and obese individuals. Adiponectin is inversely correlated with visceral adiposity, a component of metabolic syndrome and an important determinant of insulin sensitivity, and

Figure 1 Linkage between obesity, metabolic syndrome, inflammation, and type 2 diabetes.
Table 1 Review of Six Biomarkers

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<th>Biomarkers</th>
<th>Reference</th>
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| Adiponectin | Thanakun et al (2014) | Healthy (n=46; age 44.5 [median] years; BMI 21.8 kg/m²) vs patients with metabolic syndrome (n=82; age 48 years; BMI 27.3 kg/m²) | • Adiponectin in plasma and saliva were correlated (r=0.211, P=0.018) after controlling for age, gender and BMI.  
• Adiponectin in plasma was higher than those in saliva.  
• Salivary adiponectin levels were not significantly different between the two groups (2.92 vs 2.78 µg/mL). |
| Mamali et al (2012) | 50 healthy adults (age 34 years [mean]; BMI 22.4 kg/m²) | • Adiponectin in serum and saliva correlated (r=0.347, P<0.05) after control for age, BMI and body fat.  
• Mean value of salivary to serum adiponectin ratio was 0.001 (interquartile range 0.0004–0.0025).  
• Serum adiponectin significantly negatively correlated with age (r=−0.290) and BMI (r=−0.339), while salivary adiponectin correlated with neither age nor BMI. |
| Goodson et al (2014) | 744 (4th to 5th graders; age 11 years) youth n=186 for each group: underweight vs normal weight vs overweight vs obese | • Salivary adiponectin decreased by 30% with increasing obesity: normal healthy weight (median 4083 pg/mL) vs overweight and obese (2981 and 2798 pg/mL, respectively).  
• Blood samples are not available. |
| Leptin | Thanakun et al (2014) | Healthy vs patients with metabolic syndrome  
*Description above | • Salivary and plasma leptin had a bivariate relationship (r=−0.161, P=0.069).  
• Leptin in plasma was higher than in saliva.  
• Plasma leptin increased twofold in patients with metabolic syndrome (P<0.001).  
• No differences in salivary leptin in healthy vs metabolic syndrome (P=0.523). |
| Tvarijonavičiute et al (2017) | Adults with type 2 diabetes (n=31; age 49.8 [mean] years; BMI 26.4 kg/m²) vs Control (n=34; age 49.3 years; BMI 24.8 kg/m²) | • 2.1 times higher salivary leptin levels in adults with type 2 diabetes vs non-diabetic (55.5 vs 26.8 pg/mL, P<0.001).  
• Blood samples are not available. |
| Goodson et al (2014) | 744 youth  
*Description above | • 3 times higher salivary leptin in obese children vs healthy counterparts.  
• Blood samples are not available. |
| IL-6 | Monea et al (2014) | Adults with type 2 diabetes and periodontal disease (n=20; age 56.4 [mean] years; BMI 28.9 kg/m²) vs non-diabetic with periodontal disease (n=21; age 52.4 years; BMI 22.0 kg/m²) vs control (n=21; age 53.1 years; BMI 25.1 kg/m²) | • A tendency toward a significant correlation between IL-6 in serum and saliva (r=0.500, P=0.057).  
• A tendency toward a significant correlation between IL-6 and (a) age (r=0.491), (b) BMI (r=0.456), and (c) HbA1c (r=0.504). |
| Costa et al (2010) | Adults with untreated chronic periodontitis (n=24; age 45.4 years) vs type 2 diabetes (n=20; age 53.3 years) vs type 2 diabetes and clinically diagnosed untreated chronic periodontitis (n=24; age 47.8 years) vs control (n=22; age 43.7 [mean] years) | • The group with untreated periodontitis, and the group with type 2 diabetes with untreated periodontitis displayed higher salivary IL-6 concentrations than the control and diabetic groups (P<0.01).  
• Salivary IL-6 was positively correlated with HbA1c levels in the type 2 diabetes and untreated periodontitis group (r=0.60, P<0.01).  
• A correlation was not observed in the type 2 diabetes group (r=0.43, P>0.05). |

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| 1,5-AG     | Mook-Kanamori et al (2014)64 | Adults with type 2 diabetes (n=188; age 53.8 [median] years; BMI 29.5 kg/m²) vs Controls (n=181; age 38.5 [median] years; BMI 27.6 kg/m²) | • 1,5-AG is detectable in saliva 9P = 306×10⁻¹³  
• Salivary 1,5-AG observed in association with type 2 diabetes across all three ethnicity groups (P<0.005, for each group).  
• Salivary 1,5-AG correlates with serum 1,5-AG (r= 0.74).  
• Salivary 1,5-AG correlates with (a) serum glucose (r= -0.51) and (b) HbA1c (r= -0.59). |
| CRP        | Naidoo et al (2014)87 | Black South African children (n=170; age 10 ± 2 years; boys, n=70; girls, n=100); obese children (n=53; boys=24; girls=29; mean BMI=26.2 ± 5 kg/m²) vs overweight (n=24; boys=8; girls=16; mean BMI=19.64 ± 1.49) vs normal weight control group (n=93; boys=38; girls=55; mean BMI=16.88 ± 1.50) | • It was determined that salivary CRP can be used as an alternative marker of chronic disease risk in children.  
• There is a positive relationship between BMI and levels of CRP.  
• Obese children had higher levels of CRP than normal-weight children (P<0.05). |
|            | Goodson et al (2014)97 | 744 (4th to 5th graders; age 11 years) youth n=186 for each group: underweight vs normal weight vs overweight vs obese | • Salivary CRP was 6 times higher in obese than normal healthy weight: normal healthy weight (median 435 pg/mL) vs obese (76 pg/mL, respectively).  
• Blood samples are not available. |
| TNF-α      | Tvarjonaviciute et al (2017)29 | Non-diabetic adults (n=34, males=11, females=2, age=49.3 ± 17.2 years and BMI 24.8 ± 3.5) vs Adults with type 2 diabetes (n=31, males=14, females=16, age 49.8 ± 20.9 and BMI 26.4 ± 5.4) | • TNF-α was 1.9-fold higher in type 2 diabetics.  
• Adults in control group had 4.7 pg/mL of TNF-α in their saliva.  
• Adults with type 2 diabetes had 8.7 pg/mL in their saliva. |
|            | Srinivasan et al (2017)26 | Healthy individuals (n=20, males=10, females=10, age=48 ± 6) vs Type 2 Diabetics (n=20, males=12, females=8, age=56.5 ± 11) | • There were no significant differences between healthy individuals and type 2 diabetics, about 100 pg/mL difference between the two groups. |

positively correlated to high-density lipoprotein (HDL).35,41,42 Furthermore, individuals with metabolic syndrome reported a 15–44% decreased plasma adiponectin level than in controls, with each study reporting the metabolic syndrome patients having a higher BMI and waist circumference than the control group.43,44 A number of studies reported hypoadiponectinemia resulting in insulin resistance and type 2 diabetes.45,46 Likewise, low plasma adiponectin concentration increases the risk of type 2 diabetes, suggesting adiponectin as a potential biomarker for type 2 diabetes (Figure 1).

To date, few studies exist in the salivary analysis of adiponectin in relation to type 2 diabetes risk in both youth and adults. Goodson et al47 studied metabolic disease risk in children by salivary biomarker analysis (Table 1). The proposal of this study was to test a non-invasive approach to study inflammatory parameters in at-risk children. A total of 744 eleven-year-old children were selected from underweight, normal-weight, overweight, and obese categories (n=186 for each sub-group).47 Results showed that salivary adiponectin concentrations were 30% lower in obese children compared with their healthy counterparts (P<0.0001). Median and interquartile ranges (pg/mL) for salivary adiponectin were obtained, respectively: Underweight: 4421, 6424 (M) and 5060, 4573 (F); Normal weight: 4220, 5303 (M), and 3994, 5052 (F). Overweight: 2402, 3785 (M), and 3222, 3693(F). Obese: 2548, 2779 (M) and 3062, 3752 (F).47 Given their findings for salivary adiponectin, along with additional promising correlations with several other biomarkers tested, there is
a potential to use salivary biomarker analysis as a non-invasive tool for diabetes risk. However, to date, findings have not been consistent. Thanakun et al\(^{27}\) compared both salivary and plasma adiponectin and leptin concentrations in 46 healthy participants and 82 patients with newly diagnosed metabolic syndrome (Table 1). Total plasma adiponectin (μg/mL) was 2.3 times higher in the control group than the metabolic syndrome patients (P<0.001).\(^{27}\) In contrary to the results from the blood, total salivary adiponectin was similar between the two groups (Control: 2.92 μg/mL; Metabolic Syndrome: 2.78 μg/mL), with additional results showing no correlation between salivary concentrations of adiponectin and metabolic syndrome.\(^{27}\) However, despite the observations that adiponectin in plasma was 11.0 times higher than in saliva in the control group and 5.0 times higher in the metabolic syndrome group (P<0.001), a positive correlation between salivary and plasma adiponectin was observed (r=0.211, P=0.018).\(^{27}\) Collectively, this study concluded that only plasma adiponectin may be useful for the prediction of metabolic syndrome. Similarly, Mamali et al\(^ {28}\) studied the measurement of salivary resistin, visfatin and adiponectin levels with the purpose of evaluating their associations with serum levels (Table 1). Resistin, visfatin, and adiponectin were measured in both serum and saliva of 50 healthy adult volunteers (17M, 33F).\(^ {28}\) Mean serum and salivary adiponectin levels were 12.27 μg/mL (interquartile range 8.15–14.70) and 10.92 ng/mL (interquartile range 3.22–28.71), respectively.\(^ {28}\) Levels of adiponectin in saliva and serum showed a significant correlation (r=0.347, P<0.05), confirming their association to each other.\(^ {28}\) Data from this study suggest that salivary adiponectin may be combined with serum adiponectin in the field of research for clinical conditions such as obesity, insulin resistance, energy imbalance, and more, given more research is conducted with advanced technical procedures with standardized analytical tools. From the previous literatures, there is a common agreement on the significant correlation between salivary and plasma adiponectin concentrations.\(^ {28,47}\)

**Leptin**

Leptin is a pro-inflammatory, adipose-derived hormone that is mainly secreted by adipocytes as well as in the stomach, placenta, vascular smooth muscle cells, and cardiomyocytes.\(^ {48,49}\) Leptin receptors are located in the hypothalamus, pancreas, kidneys, liver, and heart.\(^ {50,51}\) Leptin functions as a satiety signal in the hypothalamus to regulate body weight and metabolism, with low post-prandial levels, and high levels during hunger.\(^ {48,49}\) It mediates the suppression of thyroid hormone, growth hormone, and reproductive hormones.\(^ {49}\) Hyperleptinemia is seen exclusively in obese humans, with obese individuals showing resistance to the effects of leptin on appetite.\(^ {49}\) A number of studies have reported elevated leptin concentrations in obese individuals, type 2 diabetics, and patients with metabolic syndrome (Figure 1).\(^ {52-54}\)

Metabolic syndrome participants are shown to have dramatically higher serum leptin concentration levels, with studies reporting a 56–237% increase in serum leptin concentrations in participants with metabolic syndrome vs controls.\(^ {44,54,55}\) Obese diabetic patients report a 31–54% increase in serum leptin levels than non-obese diabetic patients.\(^ {52,56}\) Additionally, drastic differences are seen in serum leptin concentrations between obese and normal-weight participants, with studies showing an average 318% increase in serum leptin concentrations in obese participants.\(^ {53,57}\)

In contrast to the aforementioned studies focusing on the leptin concentrations in blood, conclusive findings on salivary leptin are limited. Tvarijonaviciute et al\(^ {29}\) conducted a pilot study on leptin in the saliva of adults (n=65; 34 control and 31 type 2 diabetics) (Table 1). The aims of this study were to compare salivary concentrations of leptin in patients with type 2 diabetes and healthy participants, examine the changes that occur in the saliva of type 2 diabetic patients, and determine the potential of finding additional biomarkers to identify patients at greater risk of developing type 2 diabetes.\(^ {29}\) Results showed that salivary leptin was 2.1 times higher in the adults with type 2 diabetes compared with non-diabetic adults (P<0.001).\(^ {29}\) Additionally, leptin was positively correlated with insulin and pro-inflammatory cytokines (r=0.393), suggesting salivary leptin be further explored as a non-invasive biomarker of type 2 diabetes.\(^ {29}\) Similar to the aforementioned study, Goodson et al\(^ {47}\) showed promising results in 744 children with salivary leptin demonstrating a threefold increase in obese children compared to their healthy counterparts (Table 1). Furthermore, Thanakun et al\(^ {27}\) compared salivary and plasma adiponectin and leptin concentrations, as noted previously. Leptin concentration was higher in plasma (total plasma=6.04 ng/mL in healthy, 13.23 ng/mL in T2D) than in saliva (total saliva=33.80 pg/mL in healthy, and 34.88 pg/mL in T2D), (P<0.001).\(^ {27}\) Salivary and plasma leptin had a tendency toward significant bivariate relationship (r=−0.161, P=0.069).\(^ {27}\) Plasma...
leptin increased twofold in patients with metabolic syndrome (P<0.001); however, salivary leptin showed no differences between healthy participants and metabolic syndrome patients.27 Body mass index and high-density lipoprotein cholesterol were associated with a high level of plasma leptin (OR = 1.621; 95% CI 1.212–2.168 and OR = 0.966; 95% CI 0.938–0.996).27 A significant correlation was found between the salivary adiponectin/salivary leptin ratio and plasma adiponectin (r=0.371, P<0.001); however, no association with metabolic syndrome was found.27 It should be noted that such inconsistent results could be potentially caused by the heterogeneity of study design (i.e. participant characteristics and methodological protocols).27,29,47 Two studies included adult participants,27,29 whereas Goodson et al examined children only.47 Furthermore, Goodson et al conducted their study based on increasing weight status, not diabetic status.47 Taken together, it is not conclusive that salivary leptin is useful in the prediction of metabolic syndrome.

1,5-Anhydroglucitol (1,5-AG)

1,5-AG is a monosaccharide similar in molecular structure to glucose that is maintained at high concentrations in circulation under normal physiological conditions via filtration by the kidneys and reabsorption into the blood.58–62 1,5-AG is commonly used to measure short-term glucose status and hyperglycemia.58,60,63 Studies report a 35–58% decrease in concentrations of 1,5-AG in type 2 diabetics versus healthy controls.64,65 Decreased serum concentrations of 1,5-AG result from its ability to be reabsorbed by renal tubules, a process inhibited by glucosuria.58,60,64 The greater the degree of glucosuria, the lower the amount of 1,5-AG present in circulation, and the greater the amount of glucose in blood.58,60,64,66 Moreover, increased serum glucose concentrations are associated with limited glucose reabsorption by the kidneys in diabetic patients.58,60,64

To date, limited data exist on salivary 1,5-AG and its associate to type 2 diabetes. 1,5-AG is a reliable marker for glycemic control in serum, with detectable concentrations in saliva,58,60,64,67,68 suggesting its potential use as a noninvasive marker for screening at-risk diabetic patients. Mook-Kanamori et al64 conducted a cross-sectional study on salivary metabolic biomarkers of type 2 diabetes with the objective of identifying a non-invasive metabolic marker of type 2 diabetes (Table 1). A total of 188 type 2 diabetic patients and 181 controls of Arab and Asian ethnicities was included (n=369), testing a total of 2178 metabolites in their saliva, plasma, and urine.64 After the standardization and adjusting for covariates, a mere total of 94 metabolites were significantly associated with type 2 diabetes (after Bonferroni correction, P<0.000023).64 Notably, only 3 of the 94 metabolites were detectable in saliva, with 1,5-anhydroglucitol (P=3.6×10−13) being one of them.64 Salivary 1,5-AG’s association to type 2 diabetes was observed equally across all three ethnicity groups (P<0.005, for each group) and 1,5-AG in saliva correlates with 1,5-AG in serum (r=0.74), while it anti-correlates with serum glucose (r=−0.51) and HbA1c (r=−0.59).64 It was concluded that salivary 1,5-AG can be used in national screening programs for undiagnosed diabetes. Moreover, Asha et al evaluated 1,5-AG as a salivary biomarker in type 2 diabetes patients (60 participants were recruited and divided into two groups) (Table 1).69 The first group consists of 30 adult participants that have HbA1c-confirmed type 2 diabetes and the second group consists of 30 healthy adult controls confirmed by HbA1c.69 The relationship between salivary and serum 1,5-AG levels in both groups was found using the Spearman correlation test.69 A moderate, positive correlation between 1,5-AG levels in serum and saliva was found among both groups (P<0.03 and 0.009, respectively).69 Statistical analysis of the obtained data found that the group with type 2 diabetes demonstrated decreased salivary and serum concentrations of 1,5-AG compared to the healthy group.69 As such, the study concludes that serum 1,5-AG levels can be predicted when using salivary 1,5-AG levels. Despite the limited number of studies on salivary 1,5-AG as a diabetes risk, consistent findings from the studies reviewed here would potentially suggest that salivary 1,5-AG levels can be used as a biomarker for increased risk of type 2 diabetes.64,69

Interleukin-6 (IL-6)

IL-6 is a pro-inflammatory cytokine involved in the formation and progression of diabetic dyslipidemia and hyperinsulinemia.9,70,71 Studies report a 47–76% increase in serum IL-6 concentrations in type 2 diabetic patients compared to healthy controls.3,71 Likewise, altered IL-6 profiles are reported in conditions associated with the pathophysiology of type 2 diabetes (Figure 1), such as inflammation,9,30,71 insulin resistance,30,72–74 and β-cell dysfunction.74,75 Moreover, IL-6 utilizes classic and trans-signaling pathways, where classical signaling utilizes anti-inflammatory and regenerative processes, and trans-signaling utilizes pro-inflammatory responses.74,76 Elevated circulating levels of IL-6 lead to an inflammatory
response, which poses concern, as a chronic, low-grade inflammatory state is associated with the pathogenesis of type 2 diabetes.\textsuperscript{9,30,71,73,74}

Research on salivary IL-6 as a biomarker is emerging. Monea et al\textsuperscript{33} examined IL-6 as a salivary biomarker in three subgroups of patients (Table 1). Twenty adult participants with type 2 diabetes and periodontal disease, 21 systemically healthy adult participants with periodontal disease, and 21 healthy adult participants were enrolled (n=62). Salivary IL-6 levels were highest (91.1 ± 36.9 pg/mL) in patients with type 2 diabetes and periodontal disease, and the healthy controls showed the lowest (7.4 ± 2.5 pg/mL) concentration of salivary IL-6 (P<0.0001).\textsuperscript{31} A significant correlation was found between salivary and serum IL-6 concentrations (r=0.500, P=0.057) after Spearman’s nonparametric test was applied.\textsuperscript{31} It was hypothesized that cells, in diabetic patients, exert a stronger inflammatory response compared to controls. The data obtained confirmed their hypothesis. However, these findings have not been consistent amongst studies. Costa et al\textsuperscript{30} analyzed salivary IL-6 in patients with periodontal disease and type 2 diabetes (Table 1). Ninety adult participants were enrolled and categorized into four groups; healthy (n=22), untreated periodontitis (n=24), type 2 diabetes (n=20, glycated hemoglobin [HbA1c] test >6%), and untreated periodontitis and type 2 diabetes (n=24).\textsuperscript{30} All subjects in the untreated periodontitis or type 2 diabetes groups were receiving insulin and/or oral hypoglycemic agents. The untreated periodontitis, and untreated periodontitis and type 2 diabetes groups exhibited higher salivary IL-6 concentrations than the control and diabetic groups (P<0.01).\textsuperscript{30} Thus, elevated salivary IL-6 concentrations are reported in patients with periodontitis independent of diabetic status. However, given that Costa et al\textsuperscript{30} examined patients with untreated periodontitis with or without type 2 diabetes, there might be a possibility that salivary IL-6 concentrations were overestimated. Such possibility could explain inconsistent results among the studies described above. Additional studies on salivary IL-6 in relation to type 2 diabetes risk are required to draw a conclusion on its clinical utility.

**C-Reactive Protein (CRP)**

CRP is a liver-derived, sensitive and systematic biomarker of low-grade inflammation that is commonly used to predict type 2 diabetes (Figure 1).\textsuperscript{77–79} Elevated CRP can contribute to inflammatory responses by activating complement proteins, which can lead to an increase in the production of thrombogenic components, which are bound to the membranes of injured vascular cells, thus contributing to the development of insulin resistance.\textsuperscript{80,81} Furthermore, CRP is regulated by pro-inflammatory cytokines, like IL-6 and TNF-α, that have correspondingly been reported as elevated in diabetic patients.\textsuperscript{80,81} Altered profiles of these regulatory cytokines can directly impact CRP concentrations as well.\textsuperscript{81–83} Elevated CRP concentrations can result from insufficient insulin-induced suppression of CRP synthesis.\textsuperscript{81–83} CRP has shown to be effective in the risk identification of type 2 diabetes in all races, genders, and ages.\textsuperscript{77,78,80–82,84} Moreover, studies report type 2 diabetic participants having a 41–125% increase in serum CRP concentrations compared to non-diabetic patients.\textsuperscript{77,79,84} Inconsistencies exist regarding the relationship between CRP concentrations and BMI; however, a number of studies have found a significant correlation between the two.\textsuperscript{77,79,84} Strikingly, non-diabetic participants who later developed type 2 diabetes had elevated baseline median CRP levels, suggesting CRP as a potential biomarker of type 2 diabetes.\textsuperscript{77–79,84}

Contrary to the well-established data of serum CRP, there is an increasing interest in measuring salivary CRP as a less invasive diagnostic tool.\textsuperscript{47,85,86} Naidoo et al\textsuperscript{37} conducted a cross-sectional study, examining the association between salivary CRP, low cardiorespiratory fitness, and body composition in South African children (Table 1). A total of 170 South African children, ages 9.4 ± 1.6 years (grades 3 to 7), were enrolled and categorized into groups based on their weight status as either normal-weight, overweight, or obese, using the CDC guidelines.\textsuperscript{87} Obese participants showed elevated CRP concentrations (7.3 ± 0.9 pg/mL) compared to normal-weight children (6.7 ± 0.9 pg/mL, P<0.05).\textsuperscript{87} Serum samples were analyzed and CRP concentrations in saliva and serum displayed a significant correlation (r=0.70).\textsuperscript{87} The data reported poor cardiorespiratory fitness (OR 2.7, 95% CI: 1.2–6.1, P=0.02) and overweight and obesity weight status (BMI ≥ 85th percentile; OR 2.5, 95% CI: 1.1–5.9, P=0.03) as independent predictors of elevated salivary CRP secretion rate.\textsuperscript{87} These findings show a strong correlation between weight status, cardiorespiratory fitness, and inflammatory status in children. In the aforementioned study, Goodson et al\textsuperscript{17} examined metabolic disease risk in children by salivary biomarker analysis, categorizing 744 children into four categories based on their weight status (Table 1). Salivary concentrations of CRP in the obese children were sixfold higher than in normal-weight children, with
the obese children demonstrating a median CRP concentration of 435 pg/mL and normal-weight children demonstrating a median CRP concentration of 76 pg/mL. It was concluded that increased levels of CRP are associated with type 2 diabetes, with salivary and serum CRP concentrations displaying a correlation ($r=0.70$). Given the consistencies among the studies described above, it is positive to suggest that salivary CRP can be used as a biomarker of type 2 diabetes risk.\(^\text{47,87}\)

**Tumor Necrosis Factor Alpha (TNF-$\alpha$)**

TNF-$\alpha$ is a pro-inflammatory, cell signaling protein that plays critical roles in inflammatory and autoimmune responses, cellular events such as apoptosis and cell differentiation, and oxidative stress.\(^\text{88-96}\) Elevated concentrations of TNF-$\alpha$ are associated with insulin resistance, making it a contributory biomarker of type 2 diabetes.\(^\text{88-91}\) TNF-$\alpha$ works to reduce the expression of glucose transporter type 4 (GLUT-4) and serine phosphorylation of insulin receptor substrate 1 (IRS-1), both of which are critical for the synthesis of insulin.\(^\text{92-94}\) TNF-$\alpha$ activates the endothelial production of adhesion molecules such as intracellular adhesion molecule-1.\(^\text{92-94}\) Intracellular adhesion molecule-1 promotes the development of insulin resistance.\(^\text{92-94}\) If an increase in TNF-$\alpha$ concentration occurs, additional GLUT-4 and serine phosphorylation of IRS-1 are suppressed, thereby activating additional intracellular adhesion molecule-1, leading to an increased risk of developing type 2 diabetes.\(^\text{88-90}\) Furthermore, studies report type 2 diabetics demonstrating a 17–39% increase in TNF-$\alpha$ concentrations versus healthy controls.\(^\text{90,95}\) Multiple studies have tested different races, ages, and genders and have continued to show elevated TNF-$\alpha$ concentrations in patients with type 2 diabetes, suggesting TNF-$\alpha$ as a risk factor of type 2 diabetes (Figure 1).\(^\text{88,89,91}\)

Despite the established method of collecting TNF-$\alpha$ through serum samples, recent interest in measuring salivary samples has emerged.\(^\text{29,47,86,91,96}\) Mentioned prior, Tvarijonavičius et al.\(^\text{29}\) examined various salivary cytokines in type 2 diabetics with the objective of comparing their data to healthy controls (Table 1). Salivary TNF-$\alpha$ was 1.9-fold greater in type 2 diabetics versus nondiabetics ($P<0.049$).\(^\text{29}\) However, these promising findings are not consistent amongst additional studies. Srinivasan et al.\(^\text{96}\) conducted a study comparing salivary concentrations of pro-inflammatory adipokines and anti-inflammatory adipokines in self-reported type 2 diabetic participants versus healthy controls (Table 1). Forty participants were enrolled, with 20 healthy participants and 20 participants with type 2 diabetes.\(^\text{96}\) Minimal differences in TNF-$\alpha$ concentration (50 pg/mL increase in diabetic participants) were observed between the diabetic participants and healthy controls. The difference in outcomes from both studies could be possibly attributed to the fact that participants in the Srinivasan et al.\(^\text{96}\) were self-reported type 2 diabetics whereas participants in the Tvarijonavičius et al.\(^\text{29}\) were diagnosed type 2 diabetics based on the American Diabetes Association criteria. Given these inconsistencies, more research needs to be conducted on TNF-$\alpha$ as a salivary biomarker of type 2 diabetes in order to determine its clinical utility.\(^\text{96}\)

**Additional Consideration: Obesity, Oxidative Stress, Inflammation, and Type 2 Diabetes**

Oxidative stress, defined by an elevation in reactive oxygen species (ROS) and a decreased antioxidant capacity,\(^\text{97,98}\) plays an important role in the pathogenesis of insulin resistance and type 2 diabetes (Figure 1). Increased adipose tissue, a characteristic of obesity, disrupts metabolic homeostasis via overproduction of pro-inflammatory adipokines including leptin, IL-6, and TNF-$\alpha$, followed by an accelerated stimulation of ROS production which induces a prolonged state of oxidative stress.\(^\text{97,98}\) Based on this potential mechanism linking obesity and type 2 diabetes, it would be germane to investigate any salivary markers for their association with oxidative stress that could further support the consistency and reliability of saliva-based inflammatory markers discussed here.

**Conclusion**

With rapid advances in nanotechnology and molecular diagnostics, the use of saliva as a diagnostic is evolving and poses increasing clinical relevance. Saliva contains biomarkers that are ideal for early identification for oral and systemic diseases, disease diagnostics, and clinical monitoring.\(^\text{99}\) Serum sampling has been the traditional form of measuring biomarkers, primarily due to the high concentration of molecular components found in blood. Blood analysis looks at protein-bound, serum-circulating compounds, while saliva analysis looks at biologically active molecular compounds at the cellular level, suggesting salivary analysis has a greater clinical relevance.\(^\text{26}\) If saliva sampling can be used as a diagnostic, it would pose many advantages, as it would be less-intrusive, non-
invasive, cost-effective, rapid, portable, and more accessible.21,25

Nonetheless, there are a number of challenges regarding the clinical success of using saliva as a diagnostic. First, concentrations of molecular components in saliva are found in exceptionally low concentrations.23,25 Second, research on established, disease-specific salivary biomarkers is limited.25 However, contributions from genomic technologies for saliva sampling are evolving,26,100 with increasing research conducted on disease-specific biomarkers.23,25

This review focused on biomarker-specific research, with varying results on the clinical utility of saliva for each biomarker reviewed. Research on salivary 1,5-AG and CRP as a clinical biomarker of type 2 diabetes produced desirable, consistent results, while the remaining salivary biomarkers reviewed, adiponectin, leptin, IL-6, and TNF-α produced inconsistent results with some studies suggesting a clinical relevance and use as a diagnostic, while other studies suggesting further research to be conducted. These findings suggest a potential for salivary analysis to be used as a diagnostic for early identification and risk prediction for inflammatory diseases such as type 2 diabetes. Standardized methods for salivary analysis, processing, and collection must be created to reduce inconsistencies and errors amongst published data. Additional research on saliva’s diagnostic utility, using longitudinal data and standardized methods, should be further explored in order for salivary analysis to be incorporated into diagnostic settings.

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