INTRODUCTION

Celiac disease (CD) is a complex metabolic and autoimmune disorder associated with the malabsorption of gluten. Gluten protein is mainly found in wheat, rye, barley and is absorbed in the small intestine. Glutenin and gliadin proteins form a complex that leads to immune activation in patients. These immune reactions primarily affect the small intestine (mainly the proximal part), leading to the malabsorption and deficiency of minerals and vitamins. The clinical presentations of CD are widely heterogeneous and are misdiagnosed as overlapping bowel disease and other diseases. The most specific, sensitive, and preferred test is the human tissue transglutaminase (HTTG) IgA antibody, which may establish the CD diagnosis alone.
celiac disease is not completely known, although it might be due to the combination of genetic, environmental, and immunological factors.\textsuperscript{8,9} It is estimated that 1% of CD cases occur in America, and the cases have been reported globally. The CD prevalence has been reported to increase gradually with improved diagnostic practices.\textsuperscript{6} Hence, the prevalence of CD is variable from 0.33% to 2.5% globally. In Western Sahara, the highest prevalence was found, up to 5.6\%.\textsuperscript{10} The occurrence of CD in Saudi Arabia ranges from 0.5% and 1%.\textsuperscript{11,12}

The proximal small intestine has a role in mineral absorption like iron, but chronic inflammation in celiac disease affects the absorptive capacity of the proximal small intestine. The mucosal damage leads to the development of iron deficiency due to loss of iron absorption site.\textsuperscript{3,13,14} The entry of iron in duodenal mucosa requires a divalent metal transporter (DMT1) which helps iron absorption parallel to the DMT1 expression.\textsuperscript{15} Iron has an essential role in the heme group of hemoglobin for the transport of oxygen. Iron is also crucial for cytochromes and other proteins like myoglobin.\textsuperscript{3}

Moreover, CD may affect the liver due to the hallmark injury of the small intestine. The presentations of liver dysfunction are nonspecific types of hepatitis, such as discon tent and exhaustion.\textsuperscript{16} CD-associated liver diseases is probably developed by variable factors, comprising intestinal bacterial overgrowth, chronic infection and inflammation, gut permeability, genetic, molecular mimicry, and predisposition of genes. Liver diseases may develop as primary biliary cirrhosis, autoimmune hepatitis, and non-alcoholic steatohepatitis. These liver defects can be improved by the intervention of a gluten-free diet (GFD), particularly when morbid obesity is not present.\textsuperscript{17}

There are various studies in which the role of lipids has been reported. Mostly, hypocholesterolemia is associated with celiac disease, which is more severe in men than in women.\textsuperscript{18} The possible mechanism of this understanding is due to the loss of mucosa of the small intestine that cannot absorb the cholesterol in blood circulation. Our study aimed to evaluate the role of biochemical profiles (serum iron level, lipids level, and liver enzymes) and also to determine blood constituents (hematocrit, mean cell volume (MCV) in patients with celiac disease (CD) in the Saudi population.

METHODS

Prior to starting the study, ethical permission was obtained from the Local Ethical Board of Jof University (HAP-13-s-001), and Helsinki Declaration (modified 2013) was followed to include the subjects. In this cross-sectional study, a total of 70 subjects comprised of 40 celiac disease patients and 30 healthy controls (sample size calculation was carried out by applying a statistical formula keeping power of study 5% and frequency of disease in the population 0.5%) were investigated. A random selection technique was applied to collect samples. The experiments were carried out at the Department of Clinical Laboratory Sciences from November 2020 to October 2021. The celiac disease patients were included with features; a) clinical features like diarrhea and weight loss; b) endoscopic and pathological features like duodenal atrophy and goblet cell hyperplasia on small intestine biopsy; and c) biochemical testing of antibodies (positive human tissue transglutaminase (HTTG) IgA antibody. The patients were screened based on the positive biopsy and high antibody levels. The patients with malabsorption syndrome, dermatitis herpetiformis, Wilson’s disease, and Crohn’s disease were excluded from the study.

After written informed consent, about five milliliters of blood was collected aseptically into serum separating gel and EDTA vacutainers for biochemical (serum iron, lipid profile and liver enzymes) and hematological analyses, respectively. Sample analyses for hematological and biochemical were carried out according to the standard protocols.

The concentration of hematocrit (Hct) and mean cell volume (MCV) was recorded in an auto-analyzer for cell counting (Sysmex 21). Serum biochemical levels of iron (Fe), serum liver enzymes (AST, ALT, and ALP), and lipids profile were analyzed by an automated chemistry analyzer (Dimension Vista 1500 Analyzer, Siemens). Determination of human tissue transglutaminase (HTTG) IgA antibodies was done by ELISA technique that detects semi-quantitatively IgA autoantibodies to tissue transglutaminase (TTG) protein in human serum (ETI-MAX3000, Diasurin).

Statistical analysis of the data was done using SPSS v.26.0. The normally distributed quantitative variables were given as mean and standard deviation (SD) and non-normally distributed as the median and interquartile range (IQR). The comparison of groups for normally distributed data was analyzed by the Student “t” test and Mann-Whitney U test. Association was carried out by Spearman’s rho correlation (rho) and Pearson correlation (r). The categorical data were compared
between studied groups by Chi-square test. P-value <0.05 was considered statistically significant.

RESULTS

This study was comprised of 70 individuals, including 40 cases of celiac disease and 30 age and sex-matched healthy controls. The gender frequency of patients was 55% male (n = 22) and 45% females (n = 18). On the other hand, the gender frequency of controls was 46.7% (n = 13) males and 53.3% females (n = 17).

The mean ages of celiac disease patients and healthy individuals were 30.9 ± 6.8 (mean ± SD) and 31.5 ± 9.0 years, respectively, without any statistically significant difference (p = 0.774). The comparison of serum iron (Fe), serum high-density lipoprotein (HDL) and blood hematocrit and means cell volume (MCV), which were significantly reduced in CD patients in comparison to healthy controls (p <0.05) is shown in Table-I. The serum liver aminotransferases (ALT and AST) and serum anti-HTTG IgA antibodies were significantly high in CD patients as compared to the controls (p <0.05).

The correlation between anti-HTTG IgA, serum Fe, MCV, and hematocrit was analyzed in groups. A strong negative correlation was found between anti-HTTG-IgA and serum Fe levels with r-value -0.991 (p = 0.000) in the patients’ group. On the other hand, serum anti-HTTG-IgA was negatively associated with hematocrit and MCV in the patients’ group (Table-II). A weak negative association between anti-HTTG-IgA and serum Fe levels with

Table-I: Comparison of biochemical and hematological parameters between CD patients and controls (n=70).

| Parameters                     | Patients (mean ± SD) n = 40 | Controls (mean ± SD) n = 30 | Mann-Whitney U Test* / t-test† | p-value |
|--------------------------------|----------------------------|----------------------------|--------------------------------|---------|
| Age (years)                    | 30.9 ± 6.8                 | 31.5 ± 9.0                 | 24.7*                          | 0.774*  |
| Iron level (11 - 31 µmol/L)    | 8.5 ± 4.2                  | 18.9 ± 5.9                 | 13.0†                          | 0.000†  |
| AST (10-35 U/L)                | 76.9 ± 9.2                 | 36.0 ± 13.8                | 4.39*                          | 0.000*  |
| ALT (5-40 U/L)                 | 66.8 ± 8.8                 | 30.7 ± 10.1                | 31.4†                          | 0.000†  |
| ALP (40-125 U/L)               | 82.8 ± 11.4                | 77.7 ± 15.1                | 1.176*                         | 0.135*  |
| TG (<150 mg/dL)                | 100.2 ± 13.4               | 106.9 ± 21.3               | 1.184*                         | 0.136*  |
| T. cholesterol (<200 mg/dL)    | 154.2 ± 17.9               | 152.9 ± 11.6               | 1.70*                          | 0.722*  |
| HDL (>45 mg/dL)                | 49.8 ± 5.6                 | 53.6 ± 6.4                 | 3.87*                          | 0.011*  |
| LDL (<130 mg/dL)               | 72.3 ± 17.1                | 72.3 ± 11.5                | 2.47*                          | 0.963*  |
| Hct (38-48%)                   | 28.0 ± 4.1                 | 38.8 ± 6.0                 | 4.248*                         | 0.000*  |
| MCV (80-97fl)                  | 72.8 ± 6.0                 | 84.5 ± 6.5                 | 11.7†                          | 0.000†  |
| HTTG-IGA (U/mL)                | 217.4 ± 58.4               | 18.5 ± 9.5                 | 4.248†                         | 0.000†  |

*p-value generated by Mann-Whitney U Test; † p-value generated by Independent Sample “t”-Test; p-value ≤ 0.05 is considered statistically significant.

Table-II: Correlation matrix of biochemical parameters in celiac disease patients.

| Correlations (Patients) | Iron   | HTTG-IgA | Hct    | MCV    |
|-------------------------|--------|----------|--------|--------|
| Iron                    | Pearson Correlation | 1       | -0.991* | 0.979* | 0.952* |
|                         | Sig. (2-tailed)     | 0.000*  | 0.000* | 0.000* |
| HTTG-IgA                | Pearson Correlation | 1       | -0.967* | -0.946*|
|                         | Sig. (2-tailed)     | 0.000*  | 0.000* |        |
| Hct                     | Pearson Correlation | 1       |        | 0.958* |
|                         | Sig. (2-tailed)     | 0.000*  |        |        |
| MCV                     | Pearson Correlation | 1       |        |        |
|                         | Sig. (2-tailed)     |        |        |        |

* Two-tailed Pearson correlation, p-value ≤ 0.05 is considered statistically significant.
an r value -0.401 (p = 0.028) was also observed in apparently healthy controls, while the serum anti-HTTG-IgA did not link with other parameters like hematocrit and MCV (Table-III).

Association of serum iron levels, MCV, hematocrit, and anti-HTTG-IgA, were observed between patients and controls by Chi-square test, as shown in Table-IV. The categories of serum anti-HTTG IgA were divided into four groups (negative, weakly positive, positive, and strong positive), and serum iron was distributed into three categories (normal, low, and high). The CD patients were positive (100%) for anti-HTTG-IgA, which was >30 U/mL, while it was positive only in 5 (16.6%) controls. The association of higher anti-HTTG-IgA was statistically significant ($X^2$ value = 55.27, p = 0.000) and lower serum Fe was statistically significant ($X^2$ value 32.9 with p-value = 0.000) in CD patients than controls. Similarly, hematocrit and MCV were low in 98% and 82.5% of CD patients, while it was normal in most of the individuals in the control group with a statistically strong association (p <0.05).

**DISCUSSION**

In the present study, we determined the role of biochemical profiles in celiac disease patients. The impact of serum iron, lipid profile, and liver enzymes was compared. Also, the blood concentration of MCV and hematocrit were determined to assess

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**Table-III: Correlation matrix of Biochemical parameters in apparently healthy individuals.**

| Correlations (Controls) | Iron | HTTG-IgA | Hct | MCV |
|-------------------------|------|----------|-----|-----|
| Iron                    | Pearson Correlation | 1    |      |     |
|                         | Sig. (2-tailed)      | 0.028*|      |     |
| HTTG-IgA                | Pearson Correlation | -0.401*| 1    |
|                         | Sig. (2-tailed)      | 0.028*|      |     |
| Hct                     | Pearson Correlation | 0.333 | 0.265| 1   |
|                         | Sig. (2-tailed)      | 0.072 | 0.158|     |
| MCV                     | Pearson Correlation | 0.091 | 0.066| 0.029| 1   |
|                         | Sig. (2-tailed)      | 0.633 | 0.730| 0.877|     |

* Two-tailed Pearson correlation, *p-value ≤ 0.05 is considered statistically significant

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**Table-IV: Associations of qualitative parameters between CD patients and controls.**

| Variables   | Categories | Patients (n = 40) | Controls (n = 30) | Statistics |
|-------------|------------|------------------|------------------|------------|
|             |            | Frequency (%)     | Frequency (%)     | $X^2$      | *p-value* |
| Sex         | Male       | 22 (55)          | 14 (46.7)        | 0.47       | 0.48      |
|             | Female     | 18 (45)          | 16 (53.3)        |            |           |
|             | Normal     | 10 (25)          | 29 (96.7)        |            |           |
| Serum Iron  | Low        | 27 (67.5)        | 0                | 32.9       | 0.000*    |
|             | High       | 03 (7.5)         | 01(3.3)          |            |           |
| Hematocrit  | Low        | 39 (97.5)        | 11 (36.7)        | 31.0       | 0.0001*   |
|             | Normal     | 01 (2.5)         | 19 (50)          |            |           |
| MCV         | Low        | 33 (82.5)        | 09 (30)          | 19.68      | 0.000*    |
|             | Normal     | 07 (17.5)        | 21 (70)          |            |           |
|             | 1-20       | ----             | 17 (56.7)        |            |           |
|             | 21-30      | ----             | 08 (26.7)        | 55.27      | 0.000*    |
|             | 31-100     | 01 (2.5)         | 04 (13.3)        |            |           |
|             | >100       | 39 (97.5)        | 1 (3.3)          |            |           |

*p-value ≤ 0.05 is considered statistically significant.
iron deficiency anemia diagnosis. Serum iron and hematological parameters were significantly lower in CD patients than in healthy controls. The correlation study found a positive relationship between low serum with MCV and hematocrit while negatively associated with anti-HTTG-IgA antibodies.

Iron deficiency postulates the mechanism that develops malabsorption in CD patients, and iron cannot absorb into the blood. Our results are consistent with a previous study in which iron deficiency is associated with celiac disease. In another study, anemia was detected in 11% of patients and 81.5% CD patients, while low ferritin was found in 70% of cases. There are very few studies reported in the Saudi population that only targeted epidemiological studies. Micronutrient deficiencies are due to abnormal regulatory proteins having a key role in iron absorption at the stage of the enterocyte. The mechanism for iron deficiency development is not well understood.

We also determined that the serum liver enzymes, including transaminases (AST, ALT) and ALP, were significantly raised, explaining the development of liver dysfunction in celiac disease. Our results are comparable to a study conducted on the Saudi population. Although various studies documented altered or elevated serum transaminases, however, no association was found between CD patients and controls in our study. In CD cases, the development of autoimmune and cryptogenic liver defects (having a positive response with GFD) ensues. Usually, the aminotransferase elevations are mild to moderate, mainly with a low AST/ALT ratio, and show huge manifestation variability. Some other studies documented ameliorated liver functions after the adoption of GFD. The possible mechanism is likely multifactorial, and gluten-induced damage to the gut lining could lead to elevated liver aminotransferases in celiac disease.

The lipid profile was also estimated, but there was no significant association except serum HDL, which was significantly low in celiac patients. The association studies were also carried out, but there was no link with CD. A previous study documented the low serum HDL-cholesterol at baseline, which was improved with GFD intervention. The mechanism of reduced HDL is likely due to intestinal inflammation or conformational changes in apolipoprotein. In other reports, hypcholesterolemia occurs with reduced total cholesterol, LDL, and HDL in celiac disease. The role of GFD in altered lipid profile is unclear in CD. Although some reports described the improvement in lipoprotein with saturated fat-rich food, other data reported the worsening of cardiovascular disease. In another study, raised HTTG IgA antibodies and reduced serum cholesterol were reported in CD patients compared to non-celiacs. The status of biochemical parameters is very significant in celiac disease; these biomarkers might be helpful in the diagnosis and prognosis of the disease for future prospects.

Limitations: The study’s population was small and limited biomarkers were used in the diagnosis.

CONCLUSION

The serum iron was remarkably reduced in CD patients compared to the controls establishing the iron deficiency. In patients, serum levels of liver transaminases (ALT and AST) and serum anti-HTTG IgA were significantly higher. A negative correlation was found between anti-HTTG IgA and serum iron in CD patients. A positive correlation between serum iron was observed with hematocrit and MCV in CD patients. In contrast, the correlation between altered liver enzymes and lipid profiles did not observe in CD patients and controls. Moreover, establishing and exploring the roles of biochemical parameters would be helpful in disease diagnosis and tailored medicine.

Recommendations: 1. It is recommended to conduct the studies on a large sample scale to devise the mechanism for celiac disease development. 2. It is proposed to utilize the results of this study to establish the differential diagnosis of celiac disease and tailored management.

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Author’s Contribution:
MIU: Conceived the idea, Analysis, Original draft and is responsible and accountable for the accuracy or integrity of the work.
AA: Methodology and analysis.
MA: Literature review, investigation and analysis.
IS: Specimen analysis and results interpretation.
SH: Specimen collection and literature review.
HE: Literature review, analysis and editing.