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Prevalence and Morbidity of Undiagnosed Celiac Disease From a Community-based Study

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Abstract

Background & Aims—Little is known about the prevalence and burden of undiagnosed celiac disease in individuals younger than 50 years old. We determined the prevalence and morbidity of undiagnosed celiac disease in individuals younger than 50 years in a community.

Methods—We tested sera from 31,255 residents of Olmsted County, Minnesota (younger than 50 years old) without a prior diagnosis of celiac disease assay using an assay for immunoglobulin A (IgA) against tissue transglutaminase (tTG); in subjects with positive test results, celiac disease was confirmed using an assay for endomysial IgA. We performed a nested case—control study to compare the proportion of comorbidities between undiagnosed cases of celiac disease and age-and sex-matched seronegative controls (1:2). Medical records were abstracted to identify potential comorbidities.

Results—We identified 338 of 30,425 adults with positive results from both serologic tests. Based on this finding, we estimated the prevalence of celiac disease to be 1.1% (95% CI, 1.0%–

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1.2%); 8 of 830 children tested positive for IgA against tTG (1.0%, 95% CI, 0.4%–1.9%). No typical symptoms or classic consequences of diagnosed celiac disease (diarrhea, anemia, or fracture) were associated with undiagnosed celiac disease. Undiagnosed celiac disease was associated with increased rates of hypothyroidism (odds ratio, 2.2; P<.01) and lower than average level of cholesterol (P=.03) and ferritin (P=.01). During a median follow-up period of 6.3 years, the cumulative incidence of subsequent diagnosis with celiac disease at 5 years after testing was 10.8% in persons with undiagnosed celiac disease vs 0.1% in seronegative persons (P<.01). Celiac disease status was not associated with overall survival.

Conclusions—Based on serologic tests of a community population for celiac disease, we estimated the prevalence of undiagnosed celiac disease to be 1%. Undiagnosed celiac disease appeared to be clinically silent and remained undetected, but long-term outcomes have not been determined.

Keywords

mortality; epidemiology; small intestine; autoimmune disease

INTRODUCTION

Celiac disease is an immune-mediated multisystem disorder triggered by the ingestion of gluten, possibly affecting approximately 1% of the US white population.^{1, 2} Symptomatic celiac disease is associated with substantial morbidity and a modest increased mortality risk.^{3–10} A study from Sweden demonstrated increased mortality risk in patients with villous atrophy, inflammation (Marsh I/II), and latent celiac disease (defined by positive serology with normal histologic findings) in both children and adults.¹¹ However, a recent large database study of England did not show an increased mortality rate for patients with diagnosed celiac disease.¹² Strict adherence to a gluten-free diet (GFD) improves gastrointestinal symptoms within weeks and may decrease the risks of both severe complications and death, especially when mucosal recovery has been achieved.^{13–16}

Despite the availability of effective therapy, most individuals with celiac disease remain undetected and subject to the development of consequences of hidden disease.^{17, 18} Previously, we demonstrated that individuals with undiagnosed celiac disease had increased risk of comorbid conditions in people aged 50 years or older.¹⁹ In addition, based on a military based cohort, we demonstrated that undiagnosed celiac disease in adults has been shown a nearly 4-fold increased risk of death over 45 years of follow-up²; however other studies have not shown an increased risk of death^{12, 20}, and information regarding morbidity in adults with undiagnosed celiac disease is scarce. Further, the natural history of undiagnosed celiac disease is incompletely understood:^{21–24} whether any of them will stay clinically silent or progress to clinically obvious celiac disease. The aims of this study were to determine the prevalence, determine the identifying features, and evaluate morbidity of undiagnosed celiac disease in adults aged 18 to 50 years in a community. We also evaluated celiac disease outcomes, including subsequent diagnosis and survival in a community-based cohort of adults identified with undiagnosed celiac disease.

MATERIALS AND METHODS

This study is a community- based nested case-control study of individuals aged less than 50 years old, whose waste blood samples were tested for antibodies associated with celiac disease in 2006 through 2011. Our study was approved by the institutional review boards of Mayo Clinic and Olmsted Medical Center, Rochester, Minnesota.

Setting

Subjects are residents of Olmsted County, Minnesota, which has about 120,000 residents. The population has been repeatedly studied because its medical care is virtually selfcontained within the community; it is one of a few US communities where population-based studies can be conducted.^{25–28} The 2010 US census data showed that about 89% of residents in this community are White, and the county has been determined to be representative of the United States.^{28, 29} Mayo Clinic and Olmsted Medical Center are the major providers of medical care to residents in the county, and over any 4-year span, almost all residents see a provider at one of these health centers.^{25, 26, 28, 29} The Rochester Epidemiology Project is a study funded by the National Institutes of Health that links all providers in Olmsted County and provides a unique records linkage system so that accurate medical information and representative samples can be obtained.

Subjects

Approximately 65,000 adults (age, 18–50 years) lived in Olmsted County per year from 2006 through 2011, based on the 2010 US census data. Over this time, we collected serum samples of patients in this age-group who were seen at Mayo Clinic. We identified patients who had residual blood samples available for general research. Patients with known celiac disease diagnosed before or 2 weeks after serum draw or patients without research authorization were excluded from the study. Samples were retrieved from 30,425 adults whose samples were screened for celiac disease serologic findings. In addition, we tested samples of 250 children who had residual blood samples available for general research and samples of 580 children who were from age-stratified random cohorts of healthy children in Rochester, MN.^{30, 31} The Pediatric samples were only tested for tissue transglutaminase antibody-immunoglobulin A (tTGA) due to restriction of sample availability.

Case Definition

Samples were screened for celiac disease with sequential testing, with initial screening through tTGA enzyme-linked immunosorbent assay (Inova Diagnostics), as previously described.¹⁹ Positive test results were confirmed with endomysial antibody (EMA) immunofluorescence assay (Inova Diagnostics). Undiagnosed celiac disease was defined as tTGA 2.0 U/mL and a positive EMA test. A tTGA level <2.0 U/mL was considered negative, and therefore no EMA test was done. In addition, samples were considered negative when tTGA was 2.0 to 4.0 U/mL and EMA test was negative. Samples were considered equivocal when tTGA exceeded 4.0 U/mL and EMA test was negative. As previously reported, a positive fluorescence at dilutions equal to or greater than 1:5 was considered positive for EMA immunofluorescence assay.³² This sequential testing paradigm, using subsequent EMA testing for positive tTGA result, showed a sensitivity of 97% and a

specificity of 100% for celiac disease in a prior investigation done with parallel study of this exact serologic strategy and histologic evaluation.^{33, 34}

Each case was matched by age, gender, the time of serum collection, and Mayo Clinic registration year, two controls with negative celiac disease serology that were selected from the same population. All samples were removed of patient identifiers and testing was performed with no knowledge of the patients' clinical data. Ultimately 281 individuals could be used in the nested study of undiagnosed celiac disease based on serology testing and authorization of medical records for research.

Data Collection

Serum status was masked to the persons who reviewed the medical records for information before the serum draw date. The records contained inpatient, outpatient, and emergency department data. To obtain information of known comorbid conditions related to celiac disease, the diagnosis lists, outpatient notes, hospital notes, and laboratory results were evaluated, as well as follow-up data on mortality. In laboratory testing review, the values used were those obtained nearest the serum draw date.

For identifying the subsequent diagnosis of celiac disease, several data sources including celiac disease registry database were utilized, which were used in our previous study estimating the incidence of celiac disease in residents of Olmsted County. Briefly, the diagnosis codes for celiac disease through the Rochester Epidemiology Project that were used in our previous studies were sensitive to find potential celiac disease patients, and then the diagnoses of these patients were confirmed by the electronic medical record that contains information on serology and histopathology data.

Statistical Analysis

Descriptive statistics used to summarize baseline data included number (percentage) for categorical data and mean (standard deviation [SD]) or median (interquartile range [IQR]) for continuous data. Discrete numeric variables that represent counts were reported as mean (IQR) which is a more sensitive statistic than the median when there are excessive ties in the data (e.g., counts of 0). The extent of undiagnosed celiac disease among adults 50 years old in the community was estimated as the observed proportion with positive serology, along with a 95% "exact" confidence interval (CI, using the Clopper-Pearson method) to reflect this estimate's variability. Analyzed for the entire cohort, the Kaplan-Meier product limit method was used to estimate survival (from all-cause mortality) and cumulative event rate of subsequent celiac disease diagnosis, with group comparison of these rates evaluated with log-rank test. Median follow-up was estimated using the reverse Kaplan-Meier method, in which the standard calculation is modified by reversing the event indicator as if censoring were the outcome of interest.

On the matched subset of individuals with and without positive serology, we collected information on candidate risk factors, including comorbidities that occurred before the serum draw and laboratory measures collected within 1 year before or after serum draw. Each candidate risk factor was analyzed separately for an association with seropositive case status through conditional logistic regression or through regular logistic regression when

extensive amounts of data (eg, laboratory parameters) were missing. In all models, the matching variables age and sex were included as adjusting covariates to control for any residual confounding not prevented by the matching itself. Odds ratios (ORs) (95% CIs) are reported to measure the strength of association of the candidate predictor with serology status. For laboratory and hospital utilization measures with skewed distributions, data was analyzed in conditional logistic regression based on log-transformed values (ORs were expressed per 1 SD increase on log-scale). We also performed a sensitivity analysis retesting these associations on an augmented matched subset in which tTGA result, not serology result, was the basis of the matching. For this we retained all original matched sets in which the seropositive case had positive tTGA (4.0 U/mL) and both controls had negative tTGA (<4.0 U/mL), and additionally included those with equivocal serology who had positive tTGA along with corresponding matched controls.

All analyses were uncorrected for multiple testing and adhered to a liberal significance level of .05. Given that numerous factors were analyzed for an association (in all, nearly 80 comorbid conditions, diseases, and laboratory variables), these findings should be considered hypothesis-generating and interpreted cautiously. All analyses were performed using statistical software (version 9.3; SAS Institute Inc).

RESULTS

Prevalence of Undiagnosed celiac disease

From residents (age, 18-50 years) of Olmsted County who sought health care at Mayo Clinic from 2006 through 2011, we obtained and tested waste serum samples of 30,425 individuals. In total, 372 persons (1.2%) tested positive for tTGA (4.0 U/mL), 241 (0.8%) had borderline tTGA levels (2.0 tTG < 4.0 U/mL), and 29,812 (98.0%) tested negative. Those with a positive or borderline tTGA result underwent confirmatory EMA testing. On the basis of a combined serology status of tTGA and EMA results, 57 persons were considered of equivocal status (tTGA 4.0 U/mL, with negative EMA test) and were removed from the denominator for estimating the rate with positive serology. Based on the 338 persons with a combined seropositive result for celiac disease from the final denominator of 30,368 persons, the estimated prevalence of undiagnosed celiac disease in the Olmsted County residents age 18–50 was 1.1% (95% CI, 1.0%–1.2%). The estimate appeared to be slightly higher in women (1.2%; 95% CI,1.0%-1.4%) than men (1.0%, 95% CI=0.8%-1.2%), but this difference was not statistically significant (P=.07). In addition to adults, we collected and tested sera of 830 children aged less than 18 years who resided at Olmsted County, in whom 8 children were found to have positive tTGA. Figure 1 showed the overall prevalence of undiagnosed celiac disease based on tTG positivity by age categories.

Subsequent Clinical Detection

During a median follow-up of 6.3 years, 74 individuals from the cohort had received a subsequent clinical diagnosis of celiac disease at time of analysis, which corresponded to a 5-year cumulative event rate of 0.2% (95% CI, 0.2%–0.3%). Women accounted for 52 of these celiac disease diagnoses and showed slightly higher rates compared to men (5–year

event rate, 0.3% vs. 0.2%, respectively), although formal comparison showed only a borderline difference that was not statistically significant (P=.05).

Figure 2 shows the cumulative incidence of clinical diagnosis among persons in the cohort, grouped in the first panel according to serology result (after those with equivocal serology were excluded) and in the second panel according to tTGA result (all cohort members were included). Compared to rates in seronegative persons (5-year event rate, 0.1%; 95% CI, 0.1%–0.2%), the frequency of clinical diagnosis after blood draw was significantly higher in persons with undiagnosed celiac disease (5–year event rate, 10.8%; 95% CI, 7.2%–14.4%; P<.01). Of the 38 seropositive patients who received a celiac disease diagnosis, abdominal cramping (53%) and abdominal distention (39%) were the most common presenting symptoms. Interestingly, only 8% of these patients had presented with classic symptoms of diarrhea and weight loss at the time of celiac disease diagnosis.

Figure 3 shows the cumulative event rate of clinical diagnosis estimated as a function of age rather than time after blood draw. Among individuals with positive serology who had available follow-up beyond age 25, the rate of diagnosis appeared to be fairly linear across age, with approximately half of sero-positives diagnosed with celiac disease by age 53. Likewise, approximately 50% of individuals with tTGA 10 U/mL received a celiac disease diagnosis by age 50, compared to 30% of those with tTGA 4–10 U/mL and <1% of those with tTGA <4 U/mL.

Survival

In this adult cohort, rates of all-cause mortality were generally low across study follow-up (overall 5-year survival rate, 99%). Of the 301 persons who died within 5 years of serum draw, 298 had a negative serologic result at baseline and 2 had positive serology. Survival was not significantly worse in persons with undiagnosed celiac disease than those with negative serology (P=.18) (Figure 4).

Comorbid Conditions

Table 1 shows the demographic characteristics on the matched sample of cases with positive celiac disease serology and age- and sex-matched controls with negative serology. In general, patient characteristics were similar between the 2 groups, with the exception of race, which showed non-Caucasians were at lower risk of undiagnosed celiac disease than Caucasians (P=.02). Interestingly, celiac disease serology status (undiagnosed celiac disease vs controls) was not found to be associated with typical celiac disease symptoms including diarrhea, weight loss, abdominal pain, anemia, and IBS (supplemental table 1). In particular, no difference was found in the proportion reported as having irritable bowel syndrome (3% vs 4%), weight loss (3% vs 2%), diarrhea (3% vs 3%), and anemia (9% vs 10%). Table 2 summarizes the distribution of comorbid conditions and laboratory findings in persons with undiagnosed celiac disease and their matched controls. Odds of having undiagnosed celiac disease were found to be increased in persons with hypothyroidism (OR [95% CI], 2.2 [1.2–4.0]; P<.01) and those with a history of autoimmune disease (OR [95% CI], 1.6 [1.1–2.3]; P=.02). However, no significant differences were identified for hyperthyroidism (1% in persons with undiagnosed celiac disease vs <1% in matched controls), type 1 diabetes

mellitus (2% vs 1%), glucose intolerance (3% vs 3%), or neuropathy (2% vs 1%). On the basis of unmatched comparisons on laboratory findings (Table 2), undiagnosed celiac disease was associated with reduced levels of ferritin (median, 13 ug/L vs 32 ug/L; n=98) and cholesterol (mean, 180 mg/dL vs 188 mg/dL; n=526) in the respective subgroups with available measurements. None of these differences remained significant after adjustment for multiple comparisons.

Sensitivity analysis was performed on an augmented matched subset that was identified and compared on the basis of tTGA result, thereby allowing the inclusion of tTGA positive (4.0 U/mL) individuals with negative EMA (i.e., equivocal serology) in addition to the original serology-based matched sets (Supplemental Table 2). These matched comparisons on 319 individuals with positive tTGA (regardless of EMA result) and 638 matched tTGA-negative controls demonstrated similar findings, although chronic fatigue was found to be significantly associated with positive tTGA (OR [95% CI], 1.9 [1.2–3.0]; *P*<.01). Since the current study is complementary to that previously performed among older adults (aged 50 years old) in the same community, the comparison with the previous study¹⁹ is shown in supplemental table 3.

Health Care Utilization

Health care utilization, as measured by numbers of inpatient, outpatient, and emergency visits, was additionally compared between groups during the period of five years prior and one year after serum collection. Table 3 summarizes the hospitalization visits according to celiac disease serology status. Undiagnosed celiac disease was associated with fewer ER visits in the six years around the time of serum collection, while frequencies of inpatient and outpatient visits were comparable between matched groups.

DISCUSSION

In this large, community-based cohort of health care–attending adults aged 18 to 50 years, the prevalence of undiagnosed celiac disease was 1.1% (95% CI, 1.0%–1.2%), which is similar to other epidemiologic data in the United States.^{19, 35} To our knowledge, this study is the largest and most comprehensively analyzed on comorbid conditions in adults aged 18 to 50 years with undiagnosed celiac disease. We found that undiagnosed celiac disease was associated with subtle differences in nutrition, including decreased levels of cholesterol and ferritin, but was not associated with typical symptoms of celiac disease. Further, the most comorbid conditions commonly related to clinically diagnosed celiac disease were not related to undiagnosed celiac disease, except hypothyroidism and a history of autoimmune disorders, compared with seronegative matched controls. Interestingly, we found about 11% of undiagnosed celiac disease persons were subsequently diagnosed (clinically) within 5 years of their positive test result.

Although the majority of patients with celiac disease (more than 80%) have not received the diagnosis, serologic screening for celiac disease is still debated, mainly because evidence is lacking on the clinically deleterious outcome and benefits after a GFD has been followed.^{36–38} Moreover, recent birth cohort studies have shown that in the patients with celiac disease susceptibility genes, celiac disease autoimmunity develops in early life,

especially at an age <5 years. ^{39, 40} Thus, patients with undiagnosed celiac disease conceivably may have atypical or mild symptoms, but their celiac disease continues to be unrecognized over a long duration and perhaps even for life.

In contrast to most of other studies in children,^{39–41} which reported a female preponderance in celiac disease, our study showed that the seroprevalence of celiac disease in adults was not significantly different between men and women. Another study based on a representative sample of the US population, which included children and adults, we also showed that seroprevalence of undiagnosed celiac disease was similar in men and women.⁴² This discrepancy data could be from the different study population such as high risk population vs. general population. In addition, biological difference could be much more influential in the early occurrence of disease. Further studies are warranted to explore the gender differences in celiac disease.

Our study showed from analyses uncorrected for multiple comparisons that hypothyroidism was associated with undiagnosed celiac disease. Moreover, we found some laboratory measures, including cholesterol and ferritin, were lower in undiagnosed celiac disease than in matched controls. These findings are partly consistent with results of a previous study¹⁹ showing that undiagnosed celiac disease is associated with increased risk of comorbid conditions in older adults, as well as findings of other studies of patients with clinically detected celiac disease.^{43, 44} Furthermore, we found the difference of laboratory parameters between those with and without positive serology. In particular, we found lower ferritin levels in undiagnosed celiac disease than controls, and although proportions with anemia did not differ between groups, the association between deficient ferritin and undiagnosed celiac disease shows some parallels to the association between anemia and diagnosed celiac disease.⁴⁵ Excess comorbid conditions and reduced nutritional markers such as lower cholesterol may result from a long duration of disease progression.^{19, 46} Recently, Kurppa et al¹⁶ reported that screen found asymptomatic adults with celiac disease benefited from a GFD, compared with untreated screen found celiac disease patients. Although current practice for the detection of celiac disease largely relies on the development of sufficient symptoms allied with clinical suspicion to enable the diagnosis of celiac disease, we found only 11% of undiagnosed celiac disease was subsequently diagnosed over a median of 6.3 years follow-up. This result is suggestive that the most persons with undiagnosed celiac disease do not have clinical symptoms or problems that sufficiently incite clinical attention to celiac disease in Olmsted County, a community with excellent access to medical care and a sustained increased rate of celiac disease diagnosis.⁴⁷ In a study of pediatric patients, van Koppen et al²² conducted a prospective 10-year follow-up of 32 children who had celiac disease identified through mass screening. Interestingly, they found that 59% of the children with undiagnosed celiac disease, who had positive serology for celiac disease and villous atrophy, did not have symptoms at screening. Among 8 of the asymptomatic children who continued to consume a gluten-containing diet, 6 children stayed symptom-free over the 10year follow-up, including 5 with persistently high levels of EMA and tTGA in serum. Moreover, following a GFD is not easy for many people and may have a negative effect on a person's life, related to psychological, sociological, and economic burden.^{48, 49} Whitaker et al⁴⁹ conducted a study about patient perceptions on the burden of GFD and showed that 27% of study participants, with a diagnosis of celiac disease but without classic celiac disease

symptoms, regretted being diagnosed with their condition. Our study also showed that the presence of comorbid conditions in undiagnosed celiac disease was similar to the presence in controls. Therefore, further large longitudinal studies are warranted to identify which persons with undetected celiac disease are at high risk for the further development of symptomatic celiac disease.

Several population-based studies have shown an association of increased mortality risk in patients with celiac disease,^{3, 5, 10, 11} but other studies have not.^{12, 19, 20} Moreover, studies of mortality in undetected celiac disease report contradictory findings, suggestive that it is either 4-fold that of the general population or not increased at all.^{2, 19, 20, 50} Our study did not find an increased mortality rate in undiagnosed celiac disease compared with seronegative controls. However, based on a small number of mortality cases and relatively short-term follow-up, our results are not conclusive and much longer follow up is needed.

A potential limitation of this study is that we did not confirm the presence of celiac disease by small intestinal biopsy in our patients with celiac disease and controls. However, the sequential serologic approach vs duodenal biopsy showed high sensitivity and high specificity for celiac disease diagnosis in previous studies.^{34, 51} Thus, misclassification bias should have been minimal. The symptoms and morbid conditions were not captured uniformly like by a survey or interview, rather these data were abstracted through medical chart review. Thus, systematic error including recall bias could be minimized since the most of medical conditions can be accurately identified through the review of medical chart. Even though we did not test IgA deficiency in the controls, the possibility of having celiac disease cases in this group is low because about 1 in 400 persons of the general population have IgA deficiency. Our data was derived from a study cohort that is predominately white (92% within the matched subset), mirroring the ethnic composition of the Olmsted County population, which limits applicability of our findings to more diverse populations.²⁹. Given that the prevalence of celiac disease has been shown to be more common in whites than in other ethnicities, we would expect our study estimate to more closely reflect the prevalence of undiagnosed celiac disease in a white US population.¹⁸ Furthermore, we tested only the waste samples collected from individuals who sought health care. Since this sample represents a potentially less healthy selection of adults in Olmsted County, our data may not be representative of the general population. Due to the relatively young age and short followup in our study population, it is difficult to assess the relative hazard of mortality in individuals with undiagnosed celiac disease, and a substantially longer period under observation is needed.

In conclusion, the estimated prevalence of undiagnosed celiac disease was 1.1% among adults aged 18–50 years old from this North American community. The group with undiagnosed celiac disease had few clues to the diagnosis and little excess morbidity rate, except hypothyroidism, compared with age- and sex-matched seronegative controls in a community sample of Olmsted County residents age 18 to 50. Undiagnosed celiac disease is common in younger adults and children, although it appears to be clinically silent, and so far no excess mortality has been demonstrated.

Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

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Abbreviations

EMA	endomysial antibody
GFD	gluten-free diet
IgA	immunoglobulin A
OR	odds ratio
tTGA	tissue transglutaminase antibody

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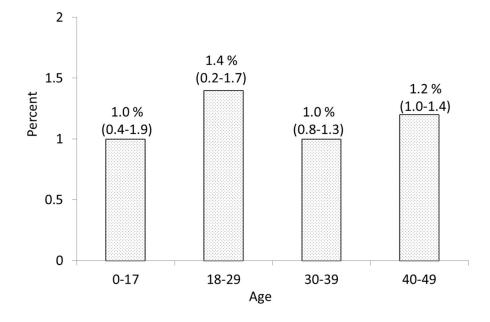


Figure 1.

The prevalence of undiagnosed celiac disease based on tissue transglutaminase IgA (tTGA) according to age categories. Positive was defined by tTGA 4 U/mL

А Serology 20 Positive Cumulative incidence of diagnosed CD (percent) Negative 15 10 5 Log-Rank test, p<.001 0 2 0 4 6 8 Years of Follow-up Number at risk 238 20007 Positive 338 289 163 46 25820 16641 Negative 30029 5285 В 25 tTGA result Cumulative incidence of diagnosed CD (percent) < 4 4-10 20 >10 15 10 5 Log-Rank test, p<.001 0 0 2 4 6 8 Years of Follow-up tTGA Number at risk 25840 <4 30053 23025 16653 5288 4-10 110 66 20 127 94 > 10 204 29 245 167 114

Figure 2.

Cumulative incidence of clinical diagnosis based on celiac disease serology status. A. Individuals are grouped according to serology result after those with equivocal serology were excluded. The cumulative incidence of clinical diagnosis was significantly higher in individuals with undiagnosed celiac disease compared to controls. B. According to categories of tTG titers (<4, 4–10, or 10 U/mL), event rates were significantly higher in individuals with strong tTG positive.

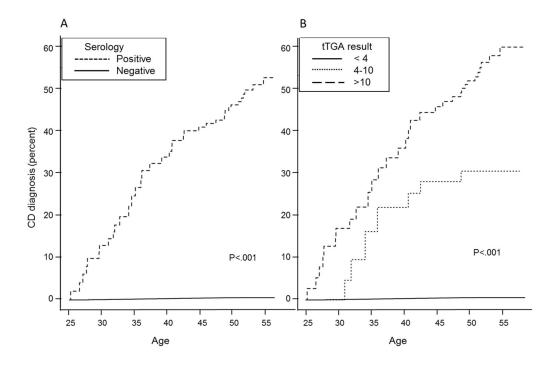
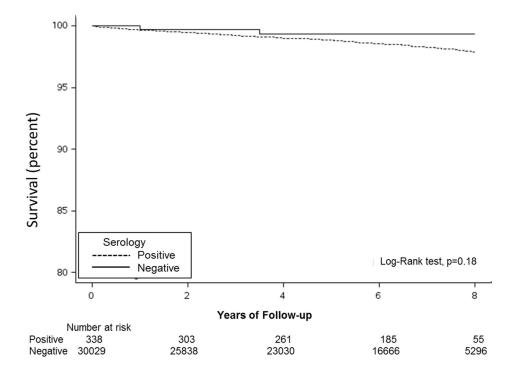


Figure 3.

Cumulative event rate of clinical diagnosis for cohort members according to their age, estimated as a function of their age rather than time after blood draw. A. Individuals are grouped according to serology result after those with equivocal serology were excluded. The cumulative rate of celiac disease diagnosis across age was markedly higher in seropositive persons (p<.001), in whom median age at celiac disease diagnosis was 53.1 years. B. All individuals in the cohort are included and grouped according to tTGA <4 4–10, or 10 U/mL, event rates across age were higher for increasing tTGA levels (p<.001), with median age at celiac disease diagnosis of 49.7 years for individuals with tTGA 10 U/mL. tTGA=Tissue transglutaminase immunoglobulin A





Kaplan-Meier survival curve for persons with positive and negative serology.

Table 1

Demographic Characteristics of Participants in the Nested Case Control Cohort ^a

Characteristic (Total No.) ^a	Missing, No. (%)	Serology Negative (n=562)	Serology Positive (n=281)	Matched Result ^b OR (95% CI) (P Value)
Age at serum draw, y (n=843),	0 (0)	35.1 (9.5)	35.1 (9.5)	NA
Female sex (n=843)	0 (0)	358 (64)	179 (64)	NA
Race (n=755)	88 (10)			Overall P=.02
Caucasian		453 (90%)	243 (96%)	1.0 (reference)
African American		20 (4%)	2 (1%)	0.13 (0.03-0.66)
Asian		24 (5%)	3 (1%)	0.23 (0.07-0.82)
Other		6 (1%)	4 (2%)	1.13 (0.29–4.36)
Weight, kg (n=763)	80 (9)	81.1 (23.6)	80.7 (19.6)	$0.99 (0.92 - 1.08) (.85)^{\mathcal{C}}$
Height, cm (n=773)	70 (8)	169.5 (9.8)	169.3 (9.6)	$0.98 (0.79 - 1.22) (.88)^{\mathcal{C}}$
BMI, kg/m ² (n=737)	106 (13)	28.1 (7.3)	28.1 (6.1)	$1.00 (0.79 - 1.27) (.99)^{\mathcal{C}}$
BMI categories (n=737) ^a	106 (13)			Overall P=.09
<25		203 (42)	90 (36)	1.0 (reference)
25-30		127 (26)	83 (33)	1.66 (1.10–2.50)
30–35		76 (16)	40 (16)	1.34 (0.82–2.19)
>35		81 (17)	37 (15)	1.10 (0.66–1.82)

Abbreviations: BMI, body mass index; NA, not applicable; OR, odds ratio; CI, confidence interval.

 a Categorical variables are presented as number (percentage) of available responses and continuous data as mean (±SD).

 b Each variable was assessed for an association with positive serology using matched analysis (conditional logistic regression analyzed on matched sets, with matching variables included as adjusting covariates in each model).

 c Data were rescaled in regression model to express the OR per 10-unit increase (ie, a 10-kg increase in weight).

Table 2

Comorbid Conditions and Laboratory Evaluations of Individuals With Undiagnosed Celiac Disease Compared With Serology-Negative Controls

	Serology Negative, No. (%) (n=562)	Serology Positive,No. (%) (n=281)	Matched Result, ^b OR (95% CI) (P Value)
Comorbid conditions ^a			
Diabetes mellitus type 1 (n=834)	3 (1)	5 (2)	3.05 (0.72–12.86) (.13)
Glucose intolerance (n=820)	14 (3)	9 (3)	1.40 (0.60–3.27) (.44)
Diabetes mellitus type 2 (n=819)	12 (2)	10 (4)	1.81 (0.74–4.43) (.19)
Hypothyroid (n=834)	26 (5)	25 (9)	2.20 (1.21–4.01) (<.01) ^C
Hyperthyroid (n=817)	1 (<1)	4 (1)	7.68 (0.84–70.13) (.07)
Any autoimmune disorder (n=835)	98 (18)	67 (24)	1.58 (1.08, 2.30) (.02) ^C
Gastroenteritis (n=826)	37 (7)	17 (6)	0.92 (0.50–1.66) (.77)
Neuropathy (n=831)	6 (1)	5 (2)	1.79 (0.51–6.35) (.36)
Depression (n=828)	128 (23)	58 (21)	0.91 (0.62–1.33) (.62)
Fibromyalgia (n=827)	11 (2)	1 (<1)	0.15 (0.02–1.24) (.08)
Chronic fatigue (n=836)	43 (8)	30 (11)	1.52 (0.93–2.49) (.10)
Ever had a fracture (n=843)	58 (10)	27 (10)	0.91 (0.55–1.49) (.70)
Ever had osteopenia (n=843)	8 (1)	2 (1)	0.48 (0.10–2.29) (.36)
Ever had osteoporosis (n=843)	2 (<1)	1 (<1)	0.99 (0.09–10.95) (>.99)
Laboratory findings			
Hemoglobin, g/dL ^C (n=564)	13.7 (1.5)	13.7 (1.5)	1.05 (0.91–1.21) (.52)
Cholesterol, mg/dL ^C (n=526)	187.8 (36.6)	180.4 (40.3)	0.95 (0.90–1.00) (. 03) ^d
Vitamin B ₁₂ , ng/Le (n=48)	422.0 (287.0–552.0)	423.0 (266.0–437.0)	0.54 (0.27–1.11) (.10)
Serum folate, mcg/L ^{e} (n=34)	13.2 (10.0–15.7)	11.1 (7.5–15.7)	0.63 (0.27–1.50) (.30)
Iron, mcg/dL ^{e} (n=52)	78.5 (55.0–106.0)	77.5 (34.0–114.0)	0.85 (0.47–1.54) (.59)
Ferritin, mcg/L ^e (n=98)	32.0 (15.0–74.0)	13.0 (5.0–39.0)	0.45 (0.24–0.86) (.02)
Total vitamin D, ng/mL ^e (n=30)	25.5 (20.5–34.5)	32.5 (20.0-40.0)	1.02 (0.48–2.17) (.97)
Albumin, g/dL^e (n=76)	4.1 (3.9–4.5)	4.3 (3.9–4.5)	1.18 (0.68–2.04) (.56)

Abbreviation: OR, odds ratio; CI confidence interval; SD, standard deviation

^aComorbid conditions are presented as count (percentage) of available responses.

 b_{Each} comorbid condition was assessed for an association with positive serology using matched analysis (conditional logistic regression performed on matched sets, with matching variables included as adjusting covariates in each model). Bold indicates statistical significance.

 C Laboratory variables are presented as mean (SD) and as unit OR to describe the effect on odds of positive serology from increasing the variable by 1 unit.

 d Reported OR describes the effect on odds of positive serology to affect cholesterol by 10 mg/dL.

 e Due to skewness, laboratory variables are described with median (interquartile range) and modeled in logistic regression based on log-transformed data, with reported OR expressed per 1 SD increase on log-scale.

Table 3

Health Care Utilization over 6 years around serum collection according to celiac disease serology status

Variable	Serology Negative (n=562)	Serology Positive (n=281)	OR (95% CI) [P-value]
Number of Encounters (n=843)	31.6 (13.0, 43.0)	29.9 (15.0, 37.0)	0.99 (0.84–1.16) [.87]
Number of Inpatient Encounters (n=843)	0.7 (0.0, 1.0)	0.6 (0.0, 1.0)	0.92 (0.80–1.07) [.30]
Number of Outpatient Encounters (n=843)	30.9 (13.0, 43.0)	29.2 (15.0, 36.0)	0.99 (0.84–1.16) [.87]
Number of ER Visits (n=843)	1.9 (0.0, 2.0)	1.5 (0.0, 2.0)	0.80 (0.68–0.94) [<.01]

Each count variable is reported as mean (IQR), with odds ratio expressed per 1 SD increase in log-transformed values