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Detection of Celiac Disease and Lymphocytic Enteropathy by Parallel Serology and Histopathology in a Population-Based

Study

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Abstract

Background & Aims—Although serological analysis is used in diagnosis of celiac disease, histopathology is considered most reliable. We performed a prospective study to determine the clinical, pathological and serological spectrum of celiac disease in a general population (Kalixanda study).

Methods—A random sample of an adult general population (n=1000) was analyzed by upper endoscopy, duodenal biopsy, and serological analysis of tissue transglutaminase (tTg) levels; endomysial antibody (EMA) levels were analyzed in samples that were tTg+. The cutoff values for diagnosis of celiac disease were villous atrophy with 40 intraepithelial lymphocytes (IELs)/100 enterocytes (ECs).

Results—Samples from 33 subjects were tTg+ and 16 were EMA+. Histological analysis identified 7/1000 subjects (0.7%) with celiac disease; all were tTg+ and 6/7 were EMA+. Another 26 subjects were tTg+ (7/26 EMA+). This was addressed by a second quantitative pathology study, (nested case-control design) using a threshold of 25 IELS/100 ECs. In this analysis, all 13 samples that were tTg + and EMA+ had \geq 25 IELs/100ECs. In total, 16 subjects (1.6%) had serological and histological evidence of gluten-sensitive enteropathy. IELs were quantified in duodenal biopsy samples from seronegative individuals (n=500); 19 (3.8%) had >25 IELs and lymphocytic duodenosis (LD).

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All authors were responsible for study concept and design; acquisition of data; analysis and interpretation of data; drafting of the manuscript; critical revision of the manuscript for important intellectual content; obtained funding, technical, or material support; Lars Agréus, Nicholas Talley were responsible for study supervision; Brian Lahr undertook statistical analysis, Mauro D'Amato HLA typing.

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Conclusions—Measurement of \geq 25 IELs/100 ECs correlated with serological indicators of celiac disease; a higher IEL threshold could miss 50% of cases. Quantification of tTg is a sensitive test for celiac disease; diagnosis can be confirmed by observation of \geq 25 IELs/100ECs in duodenal biopsies. Lymphocytic enteropathy (celiac disease and LD) is common in the population (5.4%).

Keywords

Celiac disease; Lymphocytic enteropathy; Serology; Histology; Epidemiology

Background

Celiac disease, previously thought to be rare, may occur in up to 1% of the adult Caucasian population. ¹ In rigorous studies estimates of the prevalence of celiac disease are based on the rate of clinically diagnosed cases using either a geographically restricted or birth cohort denominator. ^{2–6,7,8} However, even these studies are still likely to underestimate the prevalence of undiagnosed celiac disease. ⁹

There are two main strategies for detecting celiac disease, firstly, testing members of the population with increased risk,10 and secondly, screening general population samples by serology.¹¹ The case detection and confirmation in these studies is limited by the accuracy of serologic methods. Over the last two decades serology for celiac disease has evolved to more specific and sensitive autoantibody detection, using endomysial (EMA) or tissue transglutaminase antibodies (tTg), however, the performance of these is still variable. The sensitivity may not as good as originally reported, and may be less for individuals with partial villous atrophy. ¹² 13 14 15 ¹⁶ Conversely, serologic tests including the most specific endomysial antibody may be positive in the absence of the architectural changes that defines celiac disease. In some individuals, biopsies reveal evidence of an inflammatory response such as increased intraepithelial lymphocytosis without architectural change or increased immunoglobulin A (IgA) deposition. ¹⁷,18

The histological features of established celiac disease are easily recognized using established criteria, 19^{, 20} but subtle and early changes are difficult to diagnose with assurance. In particular, the "normal" intraepithelial lymphocyte (IEL) count has raised considerable debate, with the normal range of IELs/ 100 enterocytes (ECs) quoted from 10–40 in varying studies in different centers. ²¹, 22

Celiac disease is defined by the enteropathy that occurs in response to ingestion of gluten and resolves when it is removed from the diet. 23 An important characteristic is that of increased IELs without which celiac disease is considered unlikely even with substantial villous changes. 24 However, the converse, increased IELs without architectural changes, has been included in the spectrum of so-called gluten sensitivity. ¹⁹ Increased IELs in the absence of other histological features of celiac disease, hereafter referred to as lymphocytic duodenosis (LD) is only associated with celiac disease in a minority and may be a response to other inflammatory process in the gut. ²⁵ The clinical implications of LD are largely unknown, apart from some disease associations, but are a frequent issue given the widespread use of upper endoscopy with duodenal biopsies for the investigation of iron deficiency anemia and chronic diarrhea. 25, 26 Possible etiologies of LD are legion including infection, drugs and autoimmune disease. 25, ^{27–31} The prevalence of LD, whether associated with gluten sensitivity or otherwise in the normal population is unknown.

There are no prospective studies where celiac serology and duodenal biopsies have been performed in parallel in a sample of a normal general population to determine the spectrum of lymphocytic enteropathy that includes celiac disease and lymphocytic duodenosis. The

The aims of this study are to define the prevalence of celiac disease in the general population correlating histology and serology, and to explore the frequency of lymphocytic enteropathy, to define the true nature of the celiac iceberg. Additionally, we correlated symptoms with serological and pathological findings.

Materials and Methods

Study population

The population is from two neighboring communities in Northern Sweden, Kalix and Haparanda, with 18,408 and 10,580 inhabitants, respectively (December 1998). The demographics were similar to the national average in Sweden in both communities. The study was approved by the ethics committees of Umeå University, Sweden, and the Mayo Clinic and conducted in accordance with the revised Declaration of Helsinki.

Random sampling

Using the computerized national population register, a representative sample was generated. Every seventh adult (n = 3000) from the target population (20–80 years of age, n = 21 610) was identified, equivalent to random sampling.

Study design and response rate

A total of 2122 individuals (response rate 74.2%) completed the validated postal questionnaire, the Abdominal Symptom Questionnaire (ASQ) ³³ Of these subjects, 1001 underwent endoscopy. Of the 1001 subjects with complete EGD and extended ASQ, one refused biopsies and another two not evaluable due to missing data. Age and gender distribution in the 1001 subjects who responded to the questionnaire at both assessments (488 males [48.8%], mean age 54 years) reflected the local and Swedish population. 32

Esophagogastroduodenoscopy

Upper gastrointestinal endoscopy was performed by three experienced endoscopists using a predefined endoscopy protocol. ³² At endoscopy, biopsies were taken from the stomach (Sydney protocol) and 2 biopsies each from the bulb (D1) and second part (D2) of the duodenum. ³⁴

Celiac Serology

All subjects in the study (n=1001) had serum saved at the time of endoscopy. The basic screening test used was based on the recombinant human tissue transglutaminase (tTg) ELISA IgA antibodies assay (The Binding Site, UK). In all samples that were either weak positive (20–30 U/ml), or positive (>30 U/ml), endomysial antibody (EMA) testing was performed by incubating diluted (1:5, 1:10 and 1:20) serum samples on 5 μ m cryostat sections of monkey esophagus. The immunofluorescent antibody test was interpreted while unaware of the identity or dilution of the sera.

Histopathology

Sections were stained with haematoxylin and eosin. *H. pylori* was detected in gastric biopsies by Warthin-Starry staining. Gastric pathology was recorded as per the Sydney system. ³⁴

Histopathology I (all subjects)—There were 2 separate pathological examinations of the duodenal biopsies from D1 and D2. In the first stage, all specimens were assessed for

architectural change, (total, partial or no villous atrophy) alongside an estimation of IEL counts using the then contemporary criteria for diagnosis of celiac disease. 20

Histopathology II (nested case-control based on serology)—After serology, with changing criteria for celiac disease, a systematic re-evaluation of duodenal biopsies (blinded to serological results) a nested case control study was undertaken, each positive serological case matched to 2 seronegative controls, matched for age and gender. 21, 35 IEL counts were noted for all cases in D1 and D2 biopsies. The histological criteria used were: Non atrophic (grade A), and atrophic (grade B), Grade B subdivided into B1 –villus: crypt ratio less than 3:1 with detectable villi, and B2 with flat mucosa i.e. partial and total villous atrophy. Non atrophic (grade A) lesions were characterised by an increase in intraepithelial lymphocytes (>25) with normal villous architecture. 36. Villus height: crypt depth ratio and crypt hyperplasia were also recorded.

Validation of IEL counting method

A validated method was developed to ensure time efficient and reproducible method of determining IEL counts in paired duodenal biopsies from both D1 and D2 Five samples were subjected to detailed counts. Observers (2) each performed IEL counts, in groups of 10 enterocytes and selected 4 villi with epithelial nuclei aligned to the basement membrane, marked 1 - 4. At villus 1, IELs/10 enterocytes were counted and recorded, starting at the base of the crypt (lowest point between two adjacent villi) and continuing till the next base, (Figure 1) IEL counts of villi marked 1 - 4 were recorded, to establish intravillus differences in IEL counts. The IEL count/100 ECs was extrapolated for cumulative groups of 10 enterocytes and plotted on a graph to find at which point the counts became stable. (Figure 2) We also studied additional benefit in immunostaining lymphocytes with CD3, but found no additional sensitivity which gave largely similar findings (data not shown)

Quantitative analysis of IELs in 500 subjects

To determine the prevalence of LD in a larger sample of seronegative subjects in the cohort (500/1000) subjects were examined for D1 and D2 pathologies. IEL counts in D1 and D2 were compared. Architecture was assessed and IELs/ 100 ECs counted. A threshold of 25 IELs per 100 ECs was used as the threshold for intraepithelial lymphocytosis as quoted in European and other studies and hence defined LD in those without villous atrophy. ²¹, 26, 37

Symptom correlation

Symptoms as reported in the validated questionnaires were analyzed individually or in groups that represented dyspepsia, or irritable bowel syndrome by Rome II. ³⁸ The ASQ has been applied in a previous study of celiac disease. ³⁹

H. pylori infection

Samples from the antrum and corpus were cultured for *H. pylori* and analysed as previously reported. ⁴⁰ Current *H. pylori* infection was defined as a positive culture or histological finding.

HLA typing

HLA-DQ risk types were predicted using the method described,41 and validated in several populations of European origin.⁴² Briefly, this method uses six HLA-tagging single nucleotide polymorphisms (SNPs), to identify DQ2.2, DQ2.5, DQ7 and DQ8 risk variants based on strong linkage disequilibrium (LD) at HLA-loci. SNPs rs2187668, rs2395182, rs4713586, rs7775228, rs4639334 and rs7454108 were genotyped with TaqMan assays (Applied Biosystems, Foster City, California, USA) on a 7500 Fast Real-Time PCR system (Applied Biosystems). All markers had a genotyping success rate > 99% and did not deviate (p > 0.05) from Hardy-

Weinberg Equilibrium (HWE), with the exception of SNP rs4713586 which failed to produce clear allele clusters in the TaqMan assay, and was therefore excluded from further analyses. Genotype at the rs4713586 locus allows discrimination between DQ2.2 and the rare DQ4 type, and is therefore needed to predict DQ2 risk carriage in DQ7/DQ2.2 individuals. Only two individuals carried DQ2.2 (or DQ4) in combination with DQ7 in our sample, and these were excluded from further analyses. DQ2 and DQ8 risk variants were predicted for all other subjects as previously described. ⁴¹

Statistical analysis

Descriptive statistics were used to summarize the data, including counts and percentages for categorical data and medians and ranges for continuous data. The chi-square or Fisher's exact test was used to test for associations between two categorical variables, while the Wilcoxon Rank Sum test was used to test for continuous data differences between two groups. For two continuous parameters, the strength of their linear relationship was measured using Spearman's correlation coefficients (ρ). To assess the agreement between two measurements or the reliability between two raters, Lin's concordance correlation coefficients were used for continuous data and Cohen's kappa (κ) statistics for categorical data. The level of significance for statistical testing was defined as *P*<.05 (2-sided). All analyses were carried out using SAS version 8 software (SAS Institute Inc, Cary, NC).

Results

Serology

There were 33 subjects seropositive for tTg-IgA, widely dispersed across age (median=60, range 25–75) and evenly distributed by gender (16 males). Two known celiac subjects who had been on a gluten free diet were negative. EMA were tested on these 33 samples; 16 were positive for EMA, 3 of whom also had smooth muscle staining. All but one of the EMA positive samples were moderately or strongly positive on tTg-IGA testing. The EMA positives had higher tTg-IgA levels than the EMA negative subjects (median [range]: 89.3 [22.3 – 178.0] vs. 26.7 [20.8 – 87.1], respectively; P=0.0002)

Histopathology I

Initial review of histology using the presence of at least partial villous atrophy and 40 IELs/ 100 enterocytes as the cut off, found 7 /1000 subjects with celiac disease based on histology: 4 had total villous atrophy and 3, partial villous atrophy. All 7 were positive for tTg-IgA and 6/7 were EMA positive. However, by serology, a further 26 were positive for tTg-IgA and of these, 7/26 were also EMA positive.

Histopathology II, Nested case control study

The nested case control study comprised a blinded review of D1 and D2 biopsies of 99 subjects. One patient's slides previously found to have total villous atrophy and IELS > 40 could not be retrieved nor could the D1 slides for 2 seronegative controls. A total of 16 subjects had an IEL count of >25/ 100 enterocytes and all of these were tTg-IgA positive (14 were EMA positive, and 2 negative). Of these, 6 had total villous atrophy on at least one sample, 7 had partial villous atrophy and 3 had normal architecture (LD). Two of 19 tTg-IgA positive but EMA negative had partial villous atrophy and IELs> 25. One subject with positive tTg and strong smooth muscle staining on EMA had normal D1 and D2 biopsies. There was high agreement between serology and indication of histological abnormality associated with celiac disease (κ =0.956) though the density of IELs correlated weakly with tTg-IgA levels (ρ = 0.36, P=0.0003)

Two patients in the control group had partial villous atrophy but neither of these or the rest of seronegative controls (n=66) had > 25 IELs/100 enterocytes. The mean counts for IELs for all subjects were 14.0 (+/- 6.6) in D1 and 14.2 (+/- 6.4) in D2 (Figure 2). The mean IEL count/ 100 enterocytes in the tTg-IgA and EMA positive subjects was 33 (7–70) in D1 and 33 (7–70) in D2. The IEL counts in the biopsies from the first and second part of the duodenum were highly concordant (Lin's concordance correlation coefficient=0.854, 95% CI, 0.788– 0.901). Results are shown in Table 1

Celiac Disease, Composite Results

Thus, in total, 16/1000 subjects had serological and histological evidence of gluten sensitive enteropathy (> 25 IELs/100enterocytes with both tTg-IgA and EMA-IgA double positivity. (Figure 3) In this population the prevalence of undiagnosed celiac disease (based on tTg-IgA positive, EMA positive and IELs> 25 was 1.6% (95% CI, 0.92–2.58). If we include the 2 subjects that were previously diagnosed and treated the prevalence is 1.8%.

HLA genotyping

From the case control study, (n=97 available) with measured genotype data, 16 were seropositive. All 16 had the presence of DQ2 compared to only 17% of 81 seronegatives (p<. 001, Fisher's Exact test). There was no association between seropositivity and DQ8 (12.5% vs. 21.0%, p=0.43 Chi-Square test). The median (IQR) IEL counts of those with DQ2 vs. without are 25.5 (11.0, 34.0) vs. 10.8 (8.5, 14.0), respectively with a strong correlation between the DQ2 and the IEL count and the serology status (Rank sum test p<0.0001)

Accurate assessment of IELs by histology

Extrapolation of IEL counts/group of 10 enterocytes showed that when 50 enterocytes have been counted there is a consistent count/100 ECs with minor variation per villus. (Figure 2) Analysis of interobserver concordance in IEL counts showed substantial agreement with an unweighted kappa value of 0.74 (95% CI 0.69–0.78).

Prevalence of Lymphocytic Duodenosis (raised IELs with normal villous architecture and negative celiac serology)

In seronegative subjects, 19/ 500 subjects had LD in either D1 or D2. In this group, 72% with LD had *H. pylori* infection compared to only 30% without LD (p value = 0.0001). The rate of *H. pylori* infection did not differ according to tTg result, with 12 of 33 (36.4%) tTg-positive subjects with infection compared to 33.9% subjects in the tTg-negative group (p =0.77). One subject had taken NSAIDs and one had associated lymphocytic gastritis. No concurrent autoimmune disease, Giardia or inflammatory bowel disease was reported in these subjects. The prevalence of LD in seronegative subjects is 3.8%. (95% CI: 2.3%-6.0%)

Symptom correlation with Serology and Histology

Symptoms did not influence participation in the endoscopy study, and there was no evidence of selection bias based on a careful review of socio-demographic and symptom data as previously documented. ³²

In general, there was a lack of association between seropositivity and bowel disease symptoms, adjusted for age and gender. Among the symptoms that did show a trend, constipation with hard stool (OR=6.61, 95% CI, 1.06–41.24, p=0.043) and poor appetite (OR=4.60, 95% CI, 0.95–22.20, p=0.058) were each associated with EMA-confirmed seropositivity. Interestingly, a BMI over 25 was found to be protective from positive serology (OR=0.34, 95% CI, 0.12–0.96, p=0.042). Factors showing a trend toward association with a positive tTg result included poor appetite (OR=3.41, 95% CI, 0.96–12.11, p=0.058) and weight loss (OR=3.73, 95% CI, 0.96–12.11, p=0.058) and weight loss (OR=3.74, 95% CI, 0.96–12.11, p=0.058) and weight loss (OR=3.74, 95% CI, 0.96–12.11) and p=0.058) and weight loss (OR=3.74, 95% CI, 0.96–12.11) and p=0.058) and weight loss (OR=3.74, 95% CI, 0.96–12.11) and p=0.058) and p=0.058)

0.81-17.12, p=0.090). Abdominal pain any time in the last 3 months (OR=0.49, 95% CI, 0.23– 1.04, p=0.062) and symptoms of irritable bowel syndrome any time in the last 3 months (OR=0.44, 95% CI, 0.17–1.15, p=0.092) were negatively associated with tTg positivity, as was an elevated BMI (OR=0.46, 95% CI, 0.22–0.98, p=0.045). In addition to serology, most gastrointestinal symptoms including diarrhea were not positively associated with having an increased IEL count (>25). The one exception were any minor symptoms, not amounting to criteria for functional disorders, which corresponded to a greater than 3-fold higher odds of an increased IEL count (OR=3.63, 95% CI, 1.74–7.58, p=0.001). Abdominal pain any time in the last 3 months, (OR=0.27, 95% CI, 0.12–0.62, p=0.002), dyspepsia any time in the last 3 months (OR=0.43, 95% CI, 0.19–0.98, p=0.045) and having retrosternal pain (OR=0.30, 95% CI, 0.09– 0.99, p=0.049) were negatively associated with having an increased IEL count. Correlation for height, weight, and BMI with IEL counts showed no results that were close to significance (height had nearly no correlation with IELs, r=0.01, p-value=0.87)

Discussion

This study is the first endoscopic population-based study to examine the prevalence and spectrum of celiac disease by parallel histology and serology with confirmation by HLA genotyping. Whilst prior studies have examined prevalence of celiac disease by counting clinically detected cases, serological testing of groups at especially high risk or serum surveys of the general population, biopsies have largely been reserved for confirmation of seropositive individuals. These previous studies have failed to detect potential gluten sensitive enteropathy in the absence of serological markers as may be seen in clinically detected disease. ^{15, 43}.

This study demonstrates one of the highest prevalence rates of celiac disease found in any population except for Mexico and similar to that in Finland. ^{44–46} Including patients with serology and increased IELs without atrophy and 2 previously treated patients, the prevalence approaches 2%. The high prevalence of celiac disease in this population likely is the result of the carriage of the at risk genotype that is common in Finnish and Swedish populations. ⁴⁷ If one also assumes that some of the seronegative subjects with increased IELs and normal architecture may also belong to the celiac spectrum then the prevalence of intestinal gluten sensitivity or latent celiac disease may be even higher. ⁴⁸ Such a remarkably high prevalence may represent longstanding celiac disease though it has recently been suggested that the historic rate of celiac disease based on serology may be increasing especially in adults. ^{45, 49, 50}

It is remarkable that so few subjects had symptoms of celiac disease, so called "silent" celiac disease. This makes it more difficult to identify those in a population who may have celiac disease beyond those already identified in high risk groups but also suggests that undiagnosed celiac disease may have little clinical impact, at least as detected by our questionnaire, which has been used in celiac disease, but not validated in this condition. ⁵¹

This observation is supported by work from the Cambridge heart health study, which found a lower body mass index, cholesterol and higher quality of life in seropositive individuals despite anemia and low bone density ⁵². Given these subtle manifestations of celiac disease and the increasing age at diagnosis, there may be a long presymptomatic phase prior to the development of symptoms. This hypothesis is supported by a high seroprevalence that developed by age 7 in a US birth cohort. ⁵³

In this study there was a strong correlation between quantitative morphological analysis of the duodenal biopsy (with a threshold of 25 IELs/100 ECs) and human tTg-IgA serology. EMA provided important confirmatory secondary testing in our population. It may be reasonable to use this sequential testing strategy when screening a general population sample where the pretest prevalence of celiac disease is less than 2%.

The cut off for IEL counts in duodenal biopsies is controversial and normal ranges are cited as 10-30/100 ECs in the UK and up to 40/100 ECs in Europe and the United States. ⁶, ²⁵, ²⁶, ³⁸, ⁵⁴, ⁵⁵ We found using counts of >25 IELs/100 ECs is the likely cut-off for abnormality. Higher thresholds of IELs would have missed 50% of cases in the population sample evaluated.

A raised IEL count with normal villous architecture is not uncommon. One study found 2.2% (14/ 626) of duodenal biopsies revealed LD in routine practice. ²² If the IEL count is raised then celiac disease needs consideration. 56^-58

Lymphocytic enteropathy (celiac disease and LD) affected over 5% of this population. The estimated prevalence of LD in subjects seronegative for celiac disease in this setting is 3.8%. Here, there was no concurrent autoimmune disease, or inflammatory bowel disease reported, and one case with Giardia had normal duodenal pathology. There are many disease associations of a raised IEL count, including cow's milk protein sensitivity, IgA deficiency, tropical sprue and post infective malabsorption ¹¹, 12, 15, 16, ²⁷, 30, 31, ^{59–61} dermatitis herpetiformis, bacterial overgrowth, NSAIDs, systemic immune disease, microscopic colitis, lymphoma and neoplastic lymphoid disease are also possible associations. ²⁵, ⁶², ⁶³

Our observation of a strong association between lymphocytic duodenosis and *H. pylori* raises one such explanation. This suggests the small intestine can produce a distinct chronic inflammatory response to a variety of immune stimuli. The understanding of how this occurs would provide insight into basic immune homeostasis of the proximal small intestine that differs from those of IBD, which typically affects more distal parts of the intestine where the major luminal antigens are bacterial. Some experimental evidence also suggests that gluten may induce functional changes in the setting of mild lymphocytic duodenosis. ⁶⁴

In summary, lymphocytic enteropathy affecting the proximal small intestine is common, and often but not exclusively associated with celiac disease. Celiac disease appears to be more common than previously suspected, generally clinically silent, and most cases can be detected by modern serologic techniques. Specificity can be maximized by the use of the endomysial antibody test as a second line. The impact of undiagnosed celiac disease and its natural history is crucial to determining the need for detection. We also need to understand which perturbation of the intestinal milieu lead to lymphocytic enteropathy and its consequences. Clinicians should be aware of the diversity of these associations when contemplating the management of patients in whom lymphocytic enteropathy is discovered.

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| IELs | intraepithelial lymphocytes | | |
|------|-----------------------------|--|--|
| ECs | enterocytes | | |
| tTg | tissue transglutaminase | | |
| EMA | endomysial antibody | | |
| LD | Lymphocytic duodenosis | | |
| | | | |

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Walker et al.

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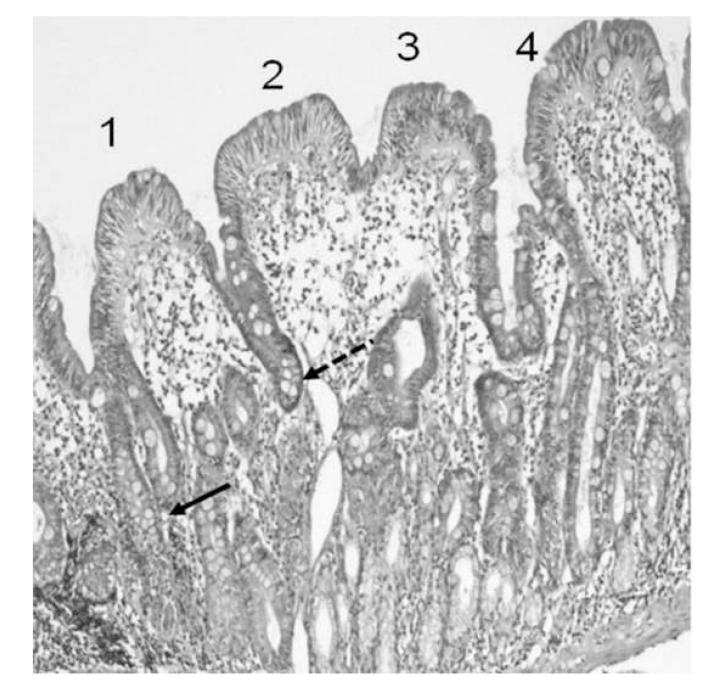


Figure 1.

Select 4 villi with epithelial nuclei aligned to basement membrane (marked 1-4). To count IELs, from villus 1: count and record IELs/10 enterocytes, starting at base of crypt (arrow, lowest point between two adjacent villi) continue till next base (dashed arrow), continue counts in of villi marked 2-4

Walker et al.

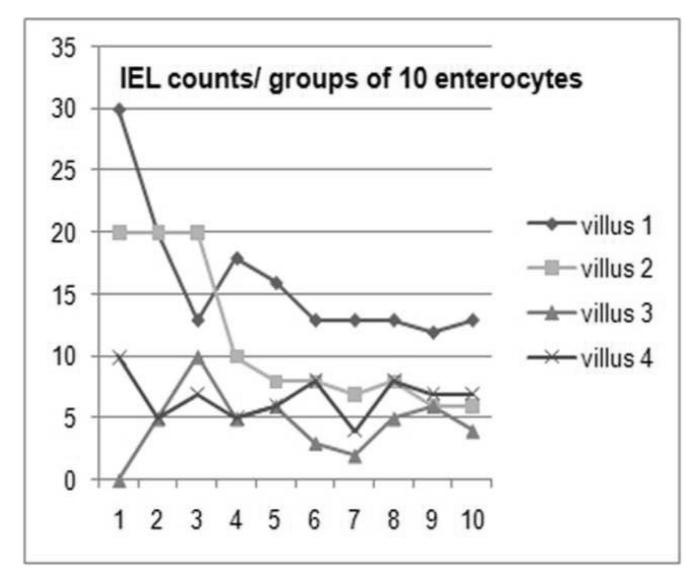


Figure 2.

Extrapolation of IEL counts/ 10 enterocytes in 4 villi, the graph levels out at 50 enterocytes.

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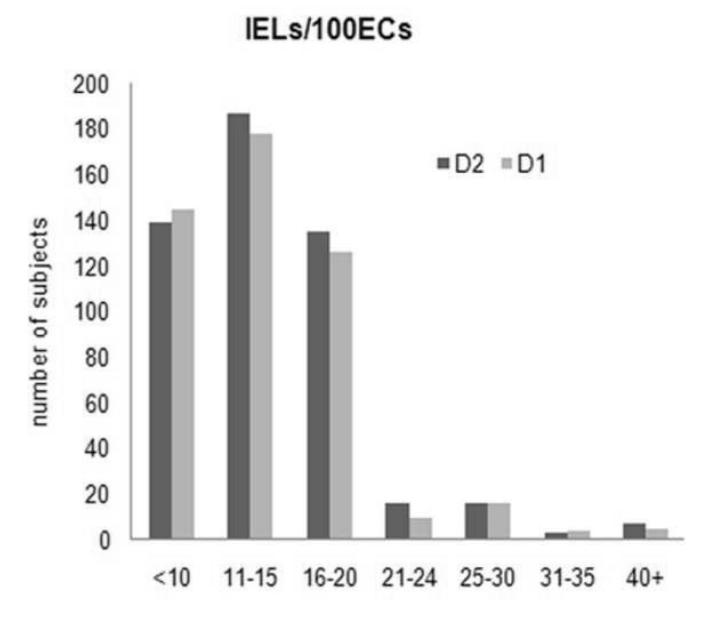


Figure 3. Comparison of D1 and D2 IEL counts in all subjects

Walker et al.

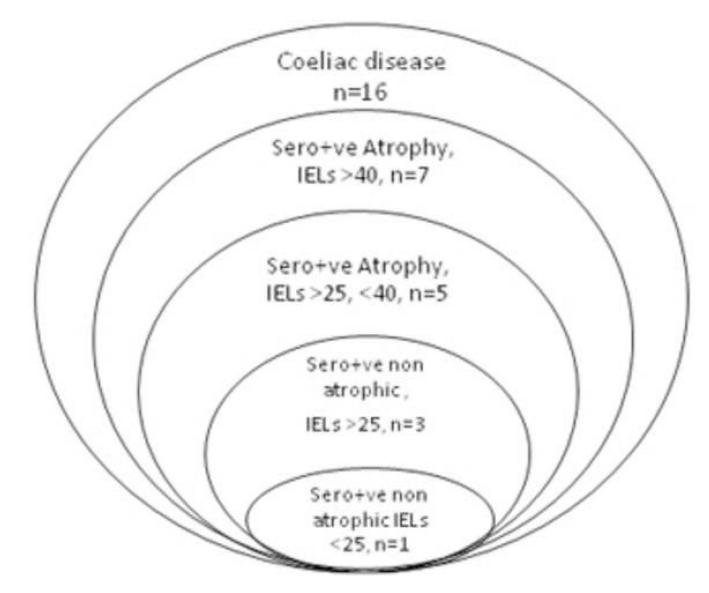


Figure 4.

Comparison of serology, villous architecture and intraepithelial lymphocyte (IELs) counts/100 enterocytes of >40 versus <25/100 enterocytes in 16 subjects with diagnosed celiac disease. Sero+ve = positive serology for tTg and EMA, Atrophy = atrophy of villi, (total, B1 or partial, B2)

Table 1

Celiac Disease Serology and Histology

| Celiac Disease by histology/ serology n=16 | | | | |
|--|---------|---|-------------------|---|
| Serology +ve | IELs>25 | D1 or/ and 2 Atrophic Grade B1 = 6, Atrophic Grade B2 = 6 | Crypt hyperplasia | Villus height: crypt depth ratio 1:1 =6 (B1) 2:1 =6 (B2) |
| 11 | 12 | 12 | 12 | 12 |
| | IELs>25 | D1 or 2 Non-Atrophic Grade A | Crypt hyperplasia | Villus height: crypt depth ratio 3:1 |
| 4 | 3 | 3 | 2 | 3 |
| | IELs<25 | D1 or 2 Non- Atrophic Grade A | Crypt hyperplasia | Villus height: crypt depth ratio 3:1 |
| 1 | 1 | 1 | 0 | 1 |