

**Type 1 diabetes**

<b>Bibliographic reference</b>	<b>Adlercreutz., EH. (2014).</b>
<b>Study type</b>	Cohort study
<b>Study quality</b>	<p>The Joanna Briggs Institute Prevalence Critical Appraisal Tool (<a href="http://ijhpm.com/article_2870_607.html">http://ijhpm.com/article_2870_607.html</a>)</p> <ol style="list-style-type: none"> <li>1. Was the sample representative of the target population? YES</li> <li>2. Were study participants recruited in an appropriate way? NO – Unclear is consecutive sample recruited</li> <li>3. Was the sample size adequate? YES</li> <li>4. Were the study subjects and the setting described in detail? YES</li> <li>5. Was the data analysis conducted with sufficient coverage of the identified sample? YES</li> <li>6. Were objective, standard criteria used for the measurement of the condition? YES</li> <li>7. Was the condition measured reliably? YES</li> <li>8. Was there appropriate statistical analysis? YES</li> </ol>

	<p>9. Are all important confounding factors/subgroups/differences identified and accounted for? YES</p> <p>10. Were subpopulations identified using objective criteria? NA</p> <p>Overall risk of bias = MODERATE</p>
<b>Patient characteristics</b>	<p>N = 662 Swedish children with T1D; 1080 Danish children with T1D; 309 healthy children from Sweden; 283 healthy Danish children.</p> <p>All children were diagnosed with T1D between 1995 and 2006. Samples were collected at the time of diagnosis. Healthy controls were recruited from local schools in both Denmark and Sweden.</p> <p>Swedish T1D:</p> <ul style="list-style-type: none"> <li>• N=662</li> <li>• 305 = female</li> <li>• Median age = 10.2 (1 - 17.9)</li> </ul> <p>Danish T1D</p> <ul style="list-style-type: none"> <li>• N=1080</li> <li>• Female = 518</li> <li>• Median age = 10.3 (0.6 - 17.8)</li> </ul>
<b>Comorbid condition</b>	Type 1 diabetes (T1D)
<b>Investigations</b>	<p>Serological testing</p> <ul style="list-style-type: none"> <li>• Conjugated IgA/IgG DGP tTG</li> <li>• IgG tTG</li> </ul> <p>Celiac disease autoimmunity was defined as being positive for both IgA/G DGP tTG and IgG tTG</p> <p>HLA genotyping</p> <ul style="list-style-type: none"> <li>• HLA DQ genotyping</li> </ul>
<b>Results</b>	<p>Swedish T1D</p> <ul style="list-style-type: none"> <li>• Prevalence of CD = 17.2% (114/662)</li> </ul> <p>Danish T1D</p> <ul style="list-style-type: none"> <li>• Prevalence of CD = 11.7% (126/1080)</li> </ul>
<b>Funding</b>	None listed
<b>Other comments</b>	NO Biopsy confirmed diagnosis of CD

<b>Bibliographic reference</b>	<b>Barbato et al. (1998)</b>
<b>Study type</b>	Cross-sectional survey
<b>Study quality</b>	<p>The Joanna Briggs Institute Prevalence Critical Appraisal Tool (<a href="http://ijhpm.com/article_2870_607.html">http://ijhpm.com/article_2870_607.html</a>)</p> <ol style="list-style-type: none"> <li>1. Was the sample representative of the target population? YES</li> <li>2. Were study participants recruited in an appropriate way? YES (consecutive sample recruited)</li> <li>3. Was the sample size adequate? YES</li> <li>4. Were the study subjects and the setting described in detail? YES</li> <li>5. Was the data analysis conducted with sufficient coverage of the identified sample? YES</li> <li>6. Were objective, standard criteria used for the measurement of the condition? YES</li> <li>7. Was the condition measured reliably? YES</li> <li>8. Was there appropriate statistical analysis? YES</li> <li>9. Are all important confounding factors/subgroups/differences identified and accounted for? YES</li> <li>10. Were subpopulations identified using objective criteria? NA</li> </ol> <p>Overall risk of bias = LOW</p>
<b>Country</b>	Italy
<b>Number of patients</b>	N=175 patients with insulin dependent diabetes
<b>Study population</b>	<p>Inclusion: Patients with insulin dependent diabetes mellitus (no further inclusion criteria provided)</p> <p>51.4% male</p> <p>Age from 1 to 30 years (102 were paediatric [between 6 and 14 years] and 73 were adults)</p>
<b>Control</b>	none
<b>Length of follow-up</b>	n/a
<b>Details of coeliac testing</b>	<p>IgA and IgG AGA (using fluorescent immunoenzymatic test, Eurospital)</p> <p>Anti-endomysium antibodies (AEA) (using indirect immuno-fluorescence with those who fluoresced only in reticular tissues as positive, Medic)</p> <p>anti-reticulin antibodies (ARA) (using indirect immuno-fluorescence with those who fluoresced only in reticular tissues as positive, Eurospital)</p> <p>If tests positive for AEA (with or without positivity for ARA and AGA), intestinal biopsy was performed</p>
<b>Results</b>	<p>Overall seroprevalence: 25.6% (45/175)</p> <p>Anti-endomysium antibodies (AEA) – 21 had pathological values</p>

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	<p>23 had biopsy (21 with pathological values for AEA and 2 with pathological values for only ARA) – all 21 with pathological values for AEA had villous atrophy</p> <p>Prevalence of CD in children with diabetes: 8.8% (9/102; 95% CI 3.3 to 14.3) Prevalence of CD in adults with diabetes: 16.4% (12/73; 95% CI 7.9 to 24.9)</p> <p>Presenting symptoms at diagnosis included diarrhoea and weight loss in 2 (16 and 17 years old) and others had one or more of growth failure in height and/or weight, recurrent abdominal pain, abdominal distension, lack of appetite, mood changes, headache, sideropenic anaemia. (all symptoms disappeared after GFD was introduced)</p>
<b>Source of funding</b>	Not reported
<b>Conflicts of interest</b>	Not reported
<b>Comments</b>	

Definitions of abbreviations are given at the end of this document.

<b>Bibliographic reference</b>	<b>Cev et al. (2010)</b>
<b>Study type</b>	Cross-sectional data from case series
<b>Study quality</b>	<p>The Joanna Briggs Institute Prevalence Critical Appraisal Tool (<a href="http://ijhpm.com/article_2870_607.html">http://ijhpm.com/article_2870_607.html</a>)</p> <ol style="list-style-type: none"> <li>1. Was the sample representative of the target population? YES</li> <li>2. Were study participants recruited in an appropriate way? NO – Unclear is consecutive sample recruited</li> <li>3. Was the sample size adequate? YES</li> <li>4. Were the study subjects and the setting described in detail? YES</li> <li>5. Was the data analysis conducted with sufficient coverage of the identified sample? YES</li> <li>6. Were objective, standard criteria used for the measurement of the condition? YES</li> <li>7. Was the condition measured reliably? YES</li> <li>8. Was there appropriate statistical analysis? YES</li> <li>9. Are all important confounding factors/subgroups/differences identified and accounted for? YES</li> <li>10. Were subpopulations identified using objective criteria? NA</li> </ol> <p>Overall risk of bias = MODERATE</p>
<b>Country</b>	Romania
<b>Number of patients</b>	N=307 patients with T1D

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<b>Study population</b>	Inclusion: patients with T1D prospectively enrolled from January 2004 to December 2008 who had presented at a centre for evaluation and rehabilitation for children and adolescents 158 females, 149 males Median age 27 years (range 14-38)
<b>Control</b>	None
<b>Length of follow-up</b>	n/a
<b>Details of coeliac testing</b>	tTGA (IgA and IgG) (ELISA with human recombinant tTG as antigen with Test ESKULISA, CeliCheck, Germany; values greater than 24 U/ml were considered positive) If positive, IgA EMA (indirect immunofluorescence using unfixed cryosections of monkey oesophagus) If positive on tTGA, duodenal biopsy assessed with Marsh system
<b>Results</b>	5.5% (17) with positive tTGA 16 has biopsy 3.9% (12) with biopsy-confirmed CD - four Marsh 0 (not considered CD) - 2 Marsh 1 - 1 Marsh 2 - 9 Marsh 3
<b>Source of funding</b>	Not reported
<b>Conflicts of interest</b>	Not reported
<b>Comments</b>	Study also reports results after treatment on GFD but this was not extracted here

Definitions of abbreviations are given at the end of this document.

<b>Bibliographic reference</b>	<b>Djurić et al. (2010)</b>
<b>Study type</b>	Cross-sectional survey
<b>Study quality</b>	The Joanna Briggs Institute Prevalence Critical Appraisal Tool ( <a href="http://ijhpm.com/article_2870_607.html">http://ijhpm.com/article_2870_607.html</a> ) <ol style="list-style-type: none"> <li>1. Was the sample representative of the target population? YES</li> <li>2. Were study participants recruited in an appropriate way? NO – Unclear is consecutive sample recruited</li> <li>3. Was the sample size adequate? YES</li> <li>4. Were the study subjects and the setting described in detail? YES</li> <li>5. Was the data analysis conducted with sufficient coverage of the identified sample? YES</li> <li>6. Were objective, standard criteria used for the measurement of the condition? YES</li> <li>7. Was the condition measured reliably? YES</li> </ol>

Appendix D: Evidence Tables

	<p>8. Was there appropriate statistical analysis? YES</p> <p>9. Are all important confounding factors/subgroups/differences identified and accounted for? YES</p> <p>10. Were subpopulations identified using objective criteria? NA</p> <p>Overall risk of bias = MODERATE</p>
<b>Country</b>	Serbia
<b>Number of patients</b>	N=121 children and adolescents with T1D N=125 healthy children and adolescents as control
<b>Study population</b>	Inclusion: children and adolescent with T1D who were admitted to a university hospital or observed on an outpatient basis from October 2004 to December 2007
	70 girls, 51 boys Mean age 10.4 years
<b>Control</b>	Healthy children and adolescents identified as healthy from their medical records and routine physical examinations from south east Serbia
<b>Length of follow-up</b>	n/a
<b>Details of coeliac testing</b>	Serum IgA Anti-tTG IgA (ELISA, Euroimmun; 20 RU/ml was cut-off) / anti-tTG IgG (ELISA) if IgA deficient Biopsy if serologically positive (ESPGHAN criteria)
<b>Results</b>	9 (7.4%) were serologically positive on tTG IgA
	Of 4 with selective IgA deficiency, all had negative IgG tTG
	Biopsy-proven CD: 5.79%(7) vs 0.8% (1) (p < 0.05) (T1D group: 2 had Marsh IIIa, 3 had Marsh IIIb, 2 had Marsh IIIc; the positive control participant had Marsh IIIa)
<b>Source of funding</b>	Not reported
<b>Conflicts of interest</b>	Not reported
<b>Comments</b>	

Definitions of abbreviations are given at the end of this document.

<b>Bibliographic reference</b>	<b>Galván et al. (2008)</b>
<b>Study type</b>	Cross-sectional survey
<b>Study quality</b>	The Joanna Briggs Institute Prevalence Critical Appraisal Tool ( <a href="http://ijhpm.com/article_2870_607.html">http://ijhpm.com/article_2870_607.html</a> )

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	<ol style="list-style-type: none"> <li>1. Was the sample representative of the target population? YES</li> <li>2. Were study participants recruited in an appropriate way? NO – Unclear is consecutive sample recruited</li> <li>3. Was the sample size adequate? YES</li> <li>4. Were the study subjects and the setting described in detail? YES</li> <li>5. Was the data analysis conducted with sufficient coverage of the identified sample? YES</li> <li>6. Were objective, standard criteria used for the measurement of the condition? YES</li> <li>7. Was the condition measured reliably? YES</li> <li>8. Was there appropriate statistical analysis? YES</li> <li>9. Are all important confounding factors/subgroups/differences identified and accounted for? YES</li> <li>10. Were subpopulations identified using objective criteria? NA</li> </ol> <p>Overall risk of bias = MODERATE</p>
<b>Country</b>	Cuba
<b>Number of patients</b>	N=208 patients with T1D
<b>Study population</b>	<p>Inclusion: patients with T1D who were diagnosed as positive for antibodies against islet cells and/or glutamic acid decarboxylase isoform 65 (antibodies against GAD65) requiring insulin treatment at diagnosis</p> <p>Mean 19 years old (range 2-58) 116 male, 92 female</p>
<b>Control</b>	none
<b>Length of follow-up</b>	n/a
<b>Details of coeliac testing</b>	tTGA IgA (immunochromatographic test, HeberFast Line® anti-transglutaminase and also ELISA) Biopsy if positive tTGA
<b>Results</b>	<p>14 patients were positive on both arrays (2 had symptoms)</p> <p>6 agreed to biopsy (including the 2 with symptoms) and had features consistent with CD with 2.88% (6/208) biopsy-confirmed prevalence:</p> <ul style="list-style-type: none"> <li>- 5 had partial villous atrophy with elevated IEL counts</li> <li>- 1 had subtotal villous atrophy</li> <li>- (mean age at diagnosis: 11.00 ±4.56 years)</li> </ul>
<b>Source of funding</b>	Not reported
<b>Conflicts of interest</b>	Not reported

Comments	
Definitions of abbreviations are given at the end of this document.	
<b>Bibliographic reference</b>	<b>Kakleas et al. (2010)</b>
<b>Study type</b>	Comparative cross-sectional survey
<b>Study quality</b>	<p>The Joanna Briggs Institute Prevalence Critical Appraisal Tool (<a href="http://ijhpm.com/article_2870_607.html">http://ijhpm.com/article_2870_607.html</a>)</p> <ol style="list-style-type: none"> <li>1. Was the sample representative of the target population? YES</li> <li>2. Were study participants recruited in an appropriate way? NO – Unclear is consecutive sample recruited</li> <li>3. Was the sample size adequate? YES</li> <li>4. Were the study subjects and the setting described in detail? YES</li> <li>5. Was the data analysis conducted with sufficient coverage of the identified sample? YES</li> <li>6. Were objective, standard criteria used for the measurement of the condition? YES</li> <li>7. Was the condition measured reliably? YES</li> <li>8. Was there appropriate statistical analysis? YES</li> <li>9. Are all important confounding factors/subgroups/differences identified and accounted for? YES</li> <li>10. Were subpopulations identified using objective criteria? NA</li> </ol> <p>Overall risk of bias = MODERATE</p>
<b>Country</b>	Greece
<b>Number of patients</b>	N=105 children and adolescents with type 1 diabetes mellitus
<b>Study population</b>	<p>Inclusion: children and adolescents with T1DM regularly followed at the Diabetic Clinic of the Second University Department of Paediatrics between 2005 and 2007</p> <p>Mean <math>\pm</math> SD:            Age: 12.44 <math>\pm</math> 4.76 years            Duration of diabetes: 4.41 <math>\pm</math> 3.70            Age at diabetes diagnosis: 8.01 <math>\pm</math> 3.17 years            50.4% male            HbA1c levels: 8.13 <math>\pm</math> 1.70%</p>
<b>Control</b>	Study compared those with and without tTG IgA seropositivity for CD
<b>Length of follow-up</b>	n/a
<b>Details of coeliac</b>	Anti-tTG IgA class antibodies were detected by ELISA (using DYNEX DSX ELISA analyser; human native tissues transglutaminase



<b>testing</b>	<p>from red blood cells was used; 20-30 units was considered to be weakly positive [Inova Diagnostics, USA]                  If high values of tTG IgA was discovered on 2 consecutive measurements (60 units or more), jejunal biopsy was performed                  Conclusive diagnosis on typical mucosal findings including lymphocytic infiltration, hypertrophy of the crypts and villous atrophy( Marsh II)                  Serum total IgA levels were determined to detect IgA deficiency</p>
<b>Results</b>	<p>Serological results:  <b>Anti-tTG IgA positivity: 8.6% (9/105)</b>                  (only 5 had mild intestinal symptoms, iron deficiency anaemia and growth retardation)</p> <p>No differences between males/females, BMIHbA1c levels, but patients with positive anti-tTG IgA were significant younger (p=0.038), had shorter T1DM duration (p=0.056) and shorter height (p=0.055)</p> <p>Univariate regression analyses showed that the likelihood of anti-tTG IgA positivity was:                  - approximately 18% greater [95% CI 0.68-0.99] in younger patients with T1DM                  - 30% greater in those with short T1DM duration [95% CI: 0.48, 1.04]</p> <p>Multivariate logistic regression indicated that the patients' present age was the only determinant associated with anti-tTG IgA positivity: younger children with T1DM had 22% more odds of presenting with anti-tTG IgA positivity (OR:1.22, 95% CI 1.01-1.45)</p> <p>Biopsy results:  <b>5 patients (4.8%) had biopsy-proven CD</b> (the same 5 were those who had symptoms and anti-tTG IgA positivity with high titres 60 or more units)</p>
<b>Source of funding</b>	Not reported
<b>Conflicts of interest</b>	Study reports that there are none
<b>Comments</b>	

Definitions of abbreviations are given at the end of this document.

<b>Bibliographic reference</b>	<b>Leeds et al. (2010)</b>
<b>Study type</b>	Cross-sectional data (for prevalence) and case-control
<b>Study quality</b>	<p>The Joanna Briggs Institute Prevalence Critical Appraisal Tool (<a href="http://ijhpm.com/article_2870_607.html">http://ijhpm.com/article_2870_607.html</a>)</p> <ol style="list-style-type: none"> <li>1. Was the sample representative of the target population? YES</li> <li>2. Were study participants recruited in an appropriate way? NO – Unclear is consecutive sample recruited</li> <li>3. Was the sample size adequate? YES</li> <li>4. Were the study subjects and the setting described in detail? YES</li> </ol>

	<p>5. Was the data analysis conducted with sufficient coverage of the identified sample? YES</p> <p>6. Were objective, standard criteria used for the measurement of the condition? YES</p> <p>7. Was the condition measured reliably? YES</p> <p>8. Was there appropriate statistical analysis? YES</p> <p>9. Are all important confounding factors/subgroups/differences identified and accounted for? YES</p> <p>10. Were subpopulations identified using objective criteria? NA</p> <p>Overall risk of bias = MODERATE</p>										
<b>Country</b>	UK										
<b>Number of patients</b>	For cross-sectional data: N=1000 with T1D; N=1200 healthy controls For case-control: N=12 with newly diagnosed CD and T1D, N=24 matched controls with T1D but not CD										
<b>Study population</b>	<p>Inclusion: patients with T1D aged &gt;16 years</p> <p>Exclusion: patients &lt; 16 years, inability to consent , diabetes other than type 1</p> <p>43 patients refused to participate, resulting in 1000 included overall</p> <p>Mean age 43.2 years</p> <p>439 females</p> <p>21 patients already had established CD and T1D and were included in the analyses</p>										
<b>Control</b>	For cross-sectional data: screening of 1200 healthy volunteers from 5 separate general practices in Sheffield For case-control: 2 control subjects with T1D, matched for every case by age, sex, weight, and diabetes duration										
<b>Length of follow-up</b>	Not reported here										
<b>Details of coeliac testing</b>	IgA EMA, IgA anti-tTG and total IgA All with either positive antibody or low IgA level were offered a duodenal biopsy; histological features consistent with CD were classified according to Marsh staging with grade 3 changes(villous atrophy) considered diagnostic for CD										
<b>Results</b>	<p>Prevalence of CD:</p> <table border="1"> <thead> <tr> <th></th> <th>Newly diagnosed</th> <th>Including the 21 who already had established CD</th> <th>Control group</th> <th>Comparison of all CD patients with control group</th> </tr> </thead> <tbody> <tr> <td>Prevalence of CD</td> <td>12% (12/1000)*</td> <td>3.3% (33/1000; 95%CI 2.3-4.6)</td> <td>1% (12/1200; 95%CI 0.5-1.7)</td> <td>OR 3.3 (95%CI 1.7-6.6, p&lt; 0.0001)</td> </tr> </tbody> </table> <p>*6 had GI symptoms, 1 was anaemic, 2 were negative for EMA) – 12% undetected CD</p> <p>4 patients with positive antibodies refused to be tested. Authors calculated that if all had biopsy-proven CD, the prevalence would be 3.7% (37/1000; 95%CI 2.6-5.1)</p> <p>21 patients tested positive for EMA but did not have biopsy considered CD so were considered to have potential CD – 18 of these had completely normal biopsies but 3 had increased IELs; these patients were not included in the overall rate of CD and were excluded from</p>		Newly diagnosed	Including the 21 who already had established CD	Control group	Comparison of all CD patients with control group	Prevalence of CD	12% (12/1000)*	3.3% (33/1000; 95%CI 2.3-4.6)	1% (12/1200; 95%CI 0.5-1.7)	OR 3.3 (95%CI 1.7-6.6, p< 0.0001)
	Newly diagnosed	Including the 21 who already had established CD	Control group	Comparison of all CD patients with control group							
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	<p>investigations in this study</p> <p>A comparison between those with T1D and newly diagnosed CD and matched controls showed that patients were well matched but that those with CD and T1D had significantly higher HbA1C (median 8.2% vs 7.5%, p=0.05), significantly lower cholesterol (median 4.1 vs 4.9 mmol/L, p=0.014), and significantly lower HDL (median 1.1 vs 1.56 mmol/L, p=0.017). These patients also had a significantly higher proportion with nephrology stage &gt; 3 (41.6% vs 4.2%) and advanced retinopathy (58.3% vs 25%). However, there was no difference in quality of life, cholesterol-toHDL ratio, triglycerides, eGFR, or proportion with peripheral neuropathy.</p> <p>Of those with newly identified CD, 3/12 had abnormal bone density (on DEXA scan) and 16.7% (2/12) were considered as having osteoporosis and 8.3% (1/12) considered as having osteopenia.</p>
<b>Source of funding</b>	Bardhan Research and Education Trust of Rotherham and Solvay
<b>Conflicts of interest</b>	Paper reports no potential conflicts relevant to the article
<b>Comments</b>	This is data from a larger study considering the prevalence of microvascular complications in adults with T1D and newly diagnosed CD; data was available after 1 year but as this included patients on a GFD, this data was not extracted here.

Definitions of abbreviations are given at the end of this document.

<b>Bibliographic reference</b>	<b>Pham-Short et al. (2010)</b>
<b>Study type</b>	Case series
<b>Study quality</b>	<p>The Joanna Briggs Institute Prevalence Critical Appraisal Tool (<a href="http://ijhpm.com/article_2870_607.html">http://ijhpm.com/article_2870_607.html</a>)</p> <ol style="list-style-type: none"> <li>1. Was the sample representative of the target population? YES</li> <li>2. Were study participants recruited in an appropriate way? NO – Unclear is consecutive sample recruited</li> <li>3. Was the sample size adequate? YES</li> <li>4. Were the study subjects and the setting described in detail? YES</li> <li>5. Was the data analysis conducted with sufficient coverage of the identified sample? YES</li> <li>6. Were objective, standard criteria used for the measurement of the condition? YES</li> <li>7. Was the condition measured reliably? YES</li> <li>8. Was there appropriate statistical analysis? YES</li> <li>9. Are all important confounding factors/subgroups/differences identified and accounted for? YES</li> <li>10. Were subpopulations identified using objective criteria? NA</li> </ol> <p>Overall risk of bias = MODERATE</p>
<b>Country</b>	Australia
<b>Number of</b>	N=4379 young people with T1D

<b>patients</b>																																					
<b>Study population</b>	<p>Inclusion: people aged 18 years or younger with T1D attending a tertiary diabetes centre in New South Wales between January 1990 and December 2009</p> <p>49% (2147) male Mean age at diabetes diagnosis was <math>6.6 \pm 4.0</math> compared with <math>8.4 \pm 4.1</math> in those without CD (<math>P &lt; 0.001</math>)</p>																																				
<b>Control</b>	None																																				
<b>Length of follow-up</b>	Study conducted over a 20-year period																																				
<b>Details of coeliac testing</b>	<p>Screening for coeliac disease at diagnosis and 1-2 yearly using anti-EMA IgA and/or anti-tTG IgA antibodies (EMA used until June 2004 with indirect immunofluorescence and anti-tTG IgA after June 2004 with enzyme-linked immunosorbent assay)</p> <p>CD diagnosed with small bowel biopsy based on Marsh scores III or greater</p>																																				
<b>Results</b>	<p>4.2% (185/4379) were diagnosed with coeliac disease (45% within 2 years, 78% within 5 years, and 94% within 10 years of diabetes diagnosis)</p> <p>Of these 33% (61) were EMA or anti tTG IgA positive at diagnosis of diabetes</p> <p>Incidence of coeliac disease:</p> <table border="1"> <thead> <tr> <th>Time period</th> <th>Incidence of CD (95% CI)</th> </tr> </thead> <tbody> <tr> <td>Over entire 20 year period</td> <td>7.7 per 1000 person years (6.6-8.9)</td> </tr> <tr> <td>1990-1999</td> <td>7.5 per 1000 person years (5.8-9.5)</td> </tr> <tr> <td>2000-2009</td> <td>7.7 per 1000 person years (6.4-9.3)</td> </tr> </tbody> </table> <p>(difference between the 2 decades was not significant)</p> <p>In 2009, the prevalence of CD was 7.1% (95% CI 5.6-8.8) (75 were biopsy-proven over 1051 clinic population)</p> <p>Comparison of age at diagnosis of diabetes:</p> <table border="1"> <thead> <tr> <th rowspan="2"></th> <th colspan="3">Age at diabetes diagnosis</th> <th rowspan="2">p value*</th> </tr> <tr> <th>&lt; 5 years (n=80)</th> <th>5-10 years (n=61)</th> <th><math>\geq 10</math> years (n=44)</th> </tr> </thead> <tbody> <tr> <td>Mean age at CD diagnosis (SD)</td> <td>7.1 (3.4)</td> <td>10.5 (2.6)</td> <td>13.3 (1.6)</td> <td>Not reported</td> </tr> <tr> <td>Male gender</td> <td>50%</td> <td>46%</td> <td>51%</td> <td>NS</td> </tr> <tr> <td>Median time in years to diagnosis of CD after diabetes diagnosis (range)</td> <td>3.0 (0.1-14.3)</td> <td>2.1 (0.1-10)</td> <td>0.7 (0.2-3.8)</td> <td>&lt; 0.001</td> </tr> <tr> <td>Diagnosed with CD within</td> <td>33%</td> <td>48%</td> <td>75%</td> <td>&lt;0.01</td> </tr> </tbody> </table>	Time period	Incidence of CD (95% CI)	Over entire 20 year period	7.7 per 1000 person years (6.6-8.9)	1990-1999	7.5 per 1000 person years (5.8-9.5)	2000-2009	7.7 per 1000 person years (6.4-9.3)		Age at diabetes diagnosis			p value*	< 5 years (n=80)	5-10 years (n=61)	$\geq 10$ years (n=44)	Mean age at CD diagnosis (SD)	7.1 (3.4)	10.5 (2.6)	13.3 (1.6)	Not reported	Male gender	50%	46%	51%	NS	Median time in years to diagnosis of CD after diabetes diagnosis (range)	3.0 (0.1-14.3)	2.1 (0.1-10)	0.7 (0.2-3.8)	< 0.001	Diagnosed with CD within	33%	48%	75%	<0.01
Time period	Incidence of CD (95% CI)																																				
Over entire 20 year period	7.7 per 1000 person years (6.6-8.9)																																				
1990-1999	7.5 per 1000 person years (5.8-9.5)																																				
2000-2009	7.7 per 1000 person years (6.4-9.3)																																				
	Age at diabetes diagnosis			p value*																																	
	< 5 years (n=80)	5-10 years (n=61)	$\geq 10$ years (n=44)																																		
Mean age at CD diagnosis (SD)	7.1 (3.4)	10.5 (2.6)	13.3 (1.6)	Not reported																																	
Male gender	50%	46%	51%	NS																																	
Median time in years to diagnosis of CD after diabetes diagnosis (range)	3.0 (0.1-14.3)	2.1 (0.1-10)	0.7 (0.2-3.8)	< 0.001																																	
Diagnosed with CD within	33%	48%	75%	<0.01																																	

Appendix D: Evidence Tables

	2 years of diabetes				
	Incidence of CD per 100 person years (95%CI)	10.4 (8.2-13.0)	6.5 (4.7-8.8)	6.4 (4.9-8.2)	<0.01
	* <5 years compared to ≥10 years				
<b>Source of funding</b>	Not reported				
<b>Conflicts of interest</b>	Authors state that there is nothing to declare				
<b>Comments</b>					

Definitions of abbreviations are given at the end of this document.

<b>Bibliographic reference</b>	<b>Picarelli et al. (2005)</b>
<b>Study type</b>	Case control
<b>Study quality</b>	<p>The Joanna Briggs Institute Prevalence Critical Appraisal Tool (<a href="http://ijhpm.com/article_2870_607.html">http://ijhpm.com/article_2870_607.html</a>)</p> <ol style="list-style-type: none"> <li>1. Was the sample representative of the target population? YES</li> <li>2. Were study participants recruited in an appropriate way? Yes (consecutive sample recruited)</li> <li>3. Was the sample size adequate? YES</li> <li>4. Were the study subjects and the setting described in detail? YES</li> <li>5. Was the data analysis conducted with sufficient coverage of the identified sample? YES</li> <li>6. Were objective, standard criteria used for the measurement of the condition? YES</li> <li>7. Was the condition measured reliably? YES</li> <li>8. Was there appropriate statistical analysis? YES</li> <li>9. Are all important confounding factors/subgroups/differences identified and accounted for? YES</li> <li>10. Were subpopulations identified using objective criteria? NA</li> </ol> <p>Overall risk of bias = LOW</p>
<b>Country</b>	Italy
<b>Number of patients</b>	N=94 adults with insulin-dependent diabetes mellitus 1 N=83 control
<b>Study population</b>	Inclusion: consecutive adult patients with IDDM1 regularly attending a centre for the study of diabetes, N=43 male, N=51 female, mean age 46.9yrs (range 18 to 70yrs), none had any symptoms attributable to enteropathy, any evidence of malabsorption or been previously diagnosed with coeliac disease, all on gluten containing diet
<b>Control</b>	blood donors without IDDM1, CD, other auto-immune conditions, or first-degree relative with any autoimmune condition

Appendix D: Evidence Tables

<b>Details of coeliac testing</b>	
<b>Results</b>	All had IDDM1 for >15yrs and satisfactory metabolic control N=13 (6.4%) with coeliac disease EMA =ve vs. EMA -ve
<b>Source of funding</b>	Ministry of University and Research (MIUR), the non-governmental association for research on coeliac disease and diabetes mellitus
<b>Conflicts of interest</b>	
<b>Comments</b>	

Definitions of abbreviations are given at the end of this document.

<b>Bibliographic reference</b>	<b>Salardi et al. (2008)</b>
<b>Study type</b>	Case series (retrospective and prospective)
<b>Study quality</b>	The Joanna Briggs Institute Prevalence Critical Appraisal Tool ( <a href="http://ijhpm.com/article_2870_607.html">http://ijhpm.com/article_2870_607.html</a> ) <ol style="list-style-type: none"> <li>1. Was the sample representative of the target population? YES</li> <li>2. Were study participants recruited in an appropriate way? YES (consecutive sample recruited)</li> <li>3. Was the sample size adequate? YES</li> <li>4. Were the study subjects and the setting described in detail? YES</li> <li>5. Was the data analysis conducted with sufficient coverage of the identified sample? YES</li> <li>6. Were objective, standard criteria used for the measurement of the condition? YES</li> <li>7. Was the condition measured reliably? YES</li> <li>8. Was there appropriate statistical analysis? YES</li> <li>9. Are all important confounding factors/subgroups/differences identified and accounted for? YES</li> <li>10. Were subpopulations identified using objective criteria? NA</li> </ol> Overall risk of bias = LOW
<b>Country</b>	Italy
<b>Number of patients</b>	N=331 children with type I diabetes
<b>Study population</b>	Consecutive children newly diagnosed with type I diabetes mellitus in a paediatric clinic between 1987 and 2004 (sera was stored between 1987 and 1993 and this was retrospectively tested for CD-related antibodies)  Mean age: 8.1 ± 4.3 years (range 0.08-14.9)

Appendix D: Evidence Tables

<b>Control</b>	None
<b>Length of follow-up</b>	Immunological evaluation at diagnosis of diabetes, every 6 to 12mths after (duration 1 to 18yrs, mean 9yrs)
<b>Details of coeliac testing</b>	IgA EMA (indirect immunofluorescence using monkey oesophagus commercial kits, Eurospital, Trieste, Italy) and human umbilical cord cryostat sections (were tested to the dilution of 1:5 and were titrated to the end point if positive) Diagnosis was confirmed by intestinal biopsy with gastroduodenoscopy and multiple biopsies with specimens graded according to Marsh classification
<b>Results</b>	Apart from 2/331 patients who were diagnosed with CD before they were diagnosed with diabetes, 29 additional patients had positive EMA assay – 6 did not have biopsy as they had borderline EMA positivity (n=2) or because EMA became negative without a GFD (n=4).  23 patients had biopsy – 18 had typical CD lesions and 5 had normal mucosa; however, 2 of these 5 had a second biopsy at 1 and 4.5 years after the onset of symptoms showing typical CD lesions  6.0% (20/331) had biopsy-proven CD (an additional 2 patients had been diagnosed with CD before being diagnosed with diabetes and were on a GFD)  (After 1994, the prevalence was 10.6% [16/151] and before 1994 it was 3.3% [6/180] [p=0.015])
<b>Source of funding</b>	Not reported
<b>Conflicts of interest</b>	Not reported
<b>Comments</b>	(author's comment: same screening methods (EMA), all tests carried out in the same reference lab, consistent assay performance, population referring to the clinic did not change over time, suggest that the risk of CD increased in diabetic children after 1994)

Definitions of abbreviations are given at the end of this document.

<b>Bibliographic reference</b>	<b>Smith et al. (2000)</b>
<b>Study type</b>	Cross-sectional data (for prevalence) from case series
<b>Study quality</b>	The Joanna Briggs Institute Prevalence Critical Appraisal Tool ( <a href="http://ijhpm.com/article_2870_607.html">http://ijhpm.com/article_2870_607.html</a> ) <ol style="list-style-type: none"> <li>1. Was the sample representative of the target population? YES</li> <li>2. Were study participants recruited in an appropriate way? YES – (unselected population)</li> <li>3. Was the sample size adequate? YES</li> <li>4. Were the study subjects and the setting described in detail? YES</li> <li>5. Was the data analysis conducted with sufficient coverage of the identified sample? YES</li> <li>6. Were objective, standard criteria used for the measurement of the condition? YES</li> <li>7. Was the condition measured reliably? YES</li> </ol>

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	<p>8. Was there appropriate statistical analysis? YES</p> <p>9. Are all important confounding factors/subgroups/differences identified and accounted for? YES</p> <p>10. Were subpopulations identified using objective criteria? NA</p> <p>Overall risk of bias = LOW</p>
<b>Country</b>	Australia
<b>Number of patients</b>	N=281 children and adolescents with T1D
<b>Study population</b>	<p>Inclusion: children and adolescents with diabetes mellitus attending a paediatric diabetes clinic between January 1993 and December 1998</p> <p>Mean and SD: Age 9.9 ± 3.8 years (range 1.3 to 18 years) 133 females/136 males One patient had prior diagnosis of CD before onset of diabetes</p>
<b>Control</b>	None
<b>Length of follow-up</b>	Only cross-sectional data extracted
<b>Details of coeliac testing</b>	<p>AGA-IgG and AGA-IgA If positive AGA-IgG and undetectable AGA-IgA, total serum was measured to exclude IgA deficiency Those with double positive AGAs had gastro-duodenoscopy and multiple biopsy to confirm CD according to ESPGAN criteria</p> <p>CD diagnosis was based on increased IELs, crypt hyperplasia and/or increase in inflammatory cells in the lamina propria in addition to either total or partial villous atrophy</p>
<b>Results</b>	<p>Double positive AGAs: 12.5% (35/280) None had IgA deficiency <b>Overall CD prevalence: 5.7% (16/281)</b> (with initial biopsies confirming CD diagnosis; this rate includes the one patient with previously diagnosed CD)</p> <p>Of those diagnosed on biopsy, 7 had gluten challenge and third a biopsy under ESPGHAN criteria to confirm the diagnosis, and 4 have completed 2 biopsies; one declined a gluten challenge after the initial biopsy due to extreme gluten sensitivity and four had yet to complete a confirmatory biopsy on a GFD and/or gluten challenge at the writing of the paper) 5 with double positive antibodies did not have biopsy: one because of loss to follow-up and 4 declined because they were asymptomatic</p>
<b>Source of funding</b>	Not reported
<b>Conflicts of interest</b>	Not reported
<b>Comments</b>	The purpose of the study was to look at the prevalence of CD in diabetes mellitus and also consider the longitudinal changes in AGA status – only the cross-sectional data on prevalence was included here.



Definitions of abbreviations are given at the end of this document.

<b>Bibliographic reference</b>	<b>Uibo et al. (2010)</b>
<b>Study type</b>	Cross-sectional survey and prospective case series of some patients
<b>Study quality</b>	<p>The Joanna Briggs Institute Prevalence Critical Appraisal Tool (<a href="http://ijhpm.com/article_2870_607.html">http://ijhpm.com/article_2870_607.html</a>)</p> <ol style="list-style-type: none"> <li>1. Was the sample representative of the target population? YES</li> <li>2. Were study participants recruited in an appropriate way? NO – Unclear is consecutive sample recruited</li> <li>3. Was the sample size adequate? YES</li> <li>4. Were the study subjects and the setting described in detail? YES</li> <li>5. Was the data analysis conducted with sufficient coverage of the identified sample? YES</li> <li>6. Were objective, standard criteria used for the measurement of the condition? YES</li> <li>7. Was the condition measured reliably? YES</li> <li>8. Was there appropriate statistical analysis? YES</li> <li>9. Are all important confounding factors/subgroups/differences identified and accounted for? YES</li> <li>10. Were subpopulations identified using objective criteria? NA</li> </ol> <p>Overall risk of bias = MODERATE</p>
<b>Country</b>	Estonia
<b>Number of patients</b>	N=271 children with type 1 diabetes
<b>Study population</b>	<p>Inclusion: T1D patients from 2 main children's hospitals in Estonia who were investigated between 1995 and 2006 (T1D definition made according to the WHO and International Society for Paediatric Adolescent Diabetes criteria)</p> <p>For cross-sectional data/initial screening study (n=271):  57% male  Mean age: 10.6 years (range 1.7-18.0)  Mean age at diagnosis of T1D: 8.3 years (range 1.6-17.7)  N=122 at diagnosis of T1D  N=149 after diagnosis of T1D (0.1 to 14.8 years after diagnosis)</p> <p>For prospective case series:  N=73 of the 271 patients included in the initial screening study (56.2% male, age range: 1.7-16.2)</p>
<b>Control</b>	none
<b>Length of follow-</b>	n/a for cross-sectional data Not reported for case series

<b>up</b>						
<b>Details of coeliac testing</b>	IgA EMA and IgA tTGA (until 2000, only EMA; in 2003 all who had been tested so far were re-tested with tTGA) (IgA levels were tested to rule out IgA deficiency with DPS Immulite assay) Those with antibodies and/or with coeliac-disease related symptoms were invited for small intestinal biopsy Diagnosis of CD according to criteria recommended by ESPGHAN					
<b>Results</b>	Results of testing:					
	Initial screening/cross-sectional survey (n=271)			Prospective follow-up (n=73)		
	Rate with/without symptoms (95% CI) <sup>1</sup>	tTGA/EMA results	Biopsy results <sup>4</sup>	Those who continued in prospective study	tTGA/EMA results	Biopsy results
With symptoms	2.2% (6/271; 95% CI 0.90-4.99)	5/6 negative	Marsh 0-5 <sup>2</sup>	0	n/a	n/a
		1/6 positive	n/a (refused)	1	Negative	n/a
Without symptoms	265/271	254 negative	NA	73	71 negative 2 positive	n/a MIIIa&IIIb <sup>3</sup>
		11 positive	1 M0 1 MIIIa <sup>3</sup> 1 MIIIb <sup>3</sup> 1 n/a (refused)	0	n/a	n/a
<sup>1</sup> not a statistically significant difference, <sup>2</sup> authors considered this to be normal mucosa, <sup>3</sup> considered to be coeliac disease, <sup>4</sup> rate of those with biopsy-proven CD was considered statically significant than the EMA/tTG negative group (p<0.01)						
(none had IgA deficiency)						
Rate of CD:						
	Rate (95% CI)	Patient characteristics of those diagnosed		Presence of symptoms in those diagnosed		
Primary screening (n=271)	3.3% (9/271; 95% CI 1.63-6.42)*	Mean age 9.9 years (3.1-16.2)		None		
Prospective case series (n=73)	2.7% (2/73; 95% CI 0-0.072)	Both 10 years with duration of T1D 3.2 and 3.3 years		None		
Overall in 1995-2006	4.1% (11/271)	7 girls, 4 boys		None		
* CD was diagnosed simultaneously with T1D in 2 patients but mean 3.4 years (range 0.9-6.9) after the T1D diagnosis in the other 7.						
<b>Source of funding</b>	Estonian Science Foundation and Estonian Ministry of Education and research					
<b>Conflicts of</b>	Not reported					

interest	
Comments	

Definitions of abbreviations are given at the end of this document.