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Low childhood high density lipoprotein cholesterol levels and subsequent risk for chronic inflammatory bowel disease

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LOW CHILDHOOD HIGH DENSITY LIPOPROTEIN CHOLESTEROL LEVELS AND SUBSEQUENT RISK FOR CHRONIC INFLAMMATORY BOWEL DISEASE

Short title: Childhood HDL-cholesterol and subsequent IBD

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**ABSTRACT**

**Background and aims.** Several genetic and environmental risk factors have been linked to chronic inflammatory bowel disease (IBD). The incidence of IBD has significantly increased in developed countries during last decades. The aim of the present study was to examine childhood risk factors for subsequent IBD diagnosis in a longitudinal cohort study of children and adolescents.

**Methods:** A Finnish study population consisting of 3551 children and adolescents originally evaluated as part of the Cardiovascular Risk in Young Finns study in 1980. At baseline, participant BMI, insulin, lipid, C-reactive protein and blood pressure levels, socioeconomic position, dietary habits, and physical activity, were evaluated. In addition, information was gathered on rural residency, severe infections, breast feeding, parental smoking and birth weight. Subsequent IBD diagnosis status was evaluated based on nationwide registries on hospitalisations and drug reimbursement decisions.

**Results.** Altogether, 49 participants (1.4%) had IBD diagnosed during the 34 years of register follow-up, of which 31 had ulcerative colitis, 12 Crohn’s disease and 6 undetermined colitis. In univariate analyses, significant correlations were observed between childhood HDL-cholesterol (risk ratio (95% CI) for 1-SD change (0.58(0.42-0.79)) and CRP concentrations (1.20(1.01-1.43)) with IBD. The inverse association between HDL-cholesterol and IBD remained significant (0.57(0.39-0.82)) in a multivariable model including data on age, sex and CRP. In addition, a weighted genetic z-score of 71 single nucleotide polymorphisms associated with elevated HDL-cholesterol levels was significantly lower in IBD patients ,=0.01).
**Conclusion.** Low childhood HDL-cholesterol levels are associated with subsequent IBD diagnosis. In addition, a genetic risk score associated with low HDL-cholesterol levels predict later IBD suggesting that HDL-cholesterol metabolism might have a role in the pathogenesis of IBD.

**Keywords.** Inflammatory bowel disease, ulcerative colitis, Crohn’s disease, high density lipoprotein cholesterol, genetics

**INTRODUCTION**

In developed countries, the incidence of chronic inflammatory bowel disease (IBD), i.e. ulcerative colitis and Crohn’s disease, has increased more than tenfold during last decades (1). IBD is incurable and 80% of patients with Crohn’s disease need surgical therapy for the complications of the disease. Among children, the incidence of IBD has increased more than threefold in Finland during the last three decades and the increase has been most pronounced among adolescents (2). In children aged less than 10 years, the increase has been lower suggesting possible environmental triggers for IBD (2). The incidence of pediatric IBD in Finland is comparable to other developed countries (3). Within Finland, a geographical variation in the prevalence of ulcerative colitis, but not in Crohn’s disease, has been reported with a higher prevalence in the northern parts of the country (4), and the incidence of paediatric IBD has been shown to be higher in areas with a low density of child population (5). In other countries, a low environmental hygiene level has been associated with lower IBD prevalence(6).

In addition to environmental risk factors, genetic factors contribute to the risk of IBD. Family members of patients with Crohn’s disease have an increased risk. Compared to dizygotic twins, monozygotic twins have a significantly higher risk of Crohn’s disease (7). Childhood-onset Crohn’s disease has more often aggressive phenotype (8). Although more than 160 genetic loci are associated with the risk of developing IBD, the known risk alleles may explain only a quarter of the heritability of Crohn’s disease (9).
At present, there is a lack of prospective data concerning childhood risk factors for subsequent IBD. Using data from the longitudinal Cardiovascular Risk in Young Finns Study cohort, our aim was to examine the association between possible childhood (age 3-18 years) risk factors and subsequent IBD occurrence during 34 years of follow-up among 3551 individuals. These data could provide novel information on childhood exposures predicting later IBD.
METHODS

Participants
The Cardiovascular Risk in Young Finns Study is an ongoing prospective study focusing cardiovascular risk factors from childhood to adulthood. The baseline examination was in 1980, when participants were aged 3 – 18 years. The latest clinical examination was performed in 2011. In addition, these individuals have been followed up through national registries including data on clinical diagnoses. The current sample included 3,551 individuals with baseline risk exposure data. The study complies with the Declaration of Helsinki and has institutional ethics approval. Written informed consent was obtained from all participants.

IBD diagnoses data
Diagnose codes for Crohn’s disease (ICD-9 diagnoses 555.X and ICD-10 diagnoses K50.X) and ulcerative colitis (ICD-9 diagnoses 556.X and ICD-10 diagnoses K51.X) were extracted from the Finnish national hospitalization database (which commenced in 1969). In addition, similar data extraction was performed from the Finnish Social Insurance Institution’s drug reimbursement register. Since 1986, the Social Insurance Institution has recorded all decisions on the entitlements for the special refunds of IBD patients in a nationwide register. Between 1986 and 1993, CD and UC had separate reimbursement codes in the register. Between 1994 and 1999, CD and UC were recorded with the same IBD-code, and diagnosis codes of subtypes were not registered. Therefore the prevalence of CD and UC cannot be calculated separately between 1994 and 1999. Since 2000, the ICD-10-codes K50 and K51 have been available to separately identify patients with CD and patients with UC.

Physical variables
Height and weight, rounded to the nearest 0·5 cm and 0·1 kg respectively, were measured at all time-points using standardized protocols, and BMI was calculated as weight (kg) divided by height (m) squared. Baseline blood pressure at three years of age was measured by an ultrasound device and at other childhood ages by a mercury sphygmomanometer. A random zero sphygmomanometer was used in adults.
The first and fifth Korotkoff sounds were used to define systolic and diastolic blood pressures, which were averaged from three measurements.

Laboratory variables

Blood samples were obtained following a 12-hour fast. Standard enzymatic methods were used for serum total cholesterol, triglycerides, high-density lipoprotein (HDL) cholesterol, and plasma glucose. HDL cholesterol was measured after dextran sulfate precipitation and low-density lipoprotein (LDL) cholesterol was calculated using the Friedewald formula. High sensitivity C-reactive protein (hsCRP) was measured by an automated analyzer using a latex turbidimetric immunoassay. For hsCRP analyses, childhood serum samples were taken in 1980 and stored in -20°C. These samples were analyzed in 2005. During the storage, the samples were not thawed or refrozen.

Anamnestic variables

In childhood, questionnaires completed by the participant or their parent were used to obtain data on physical activity, birth weight, family income, rural/urban residence, breast feeding in infancy, fruit and vegetable consumption, and parental smoking. Physical activity was assessed with questions concerning the frequency and intensity of physical activity and a physical activity index was calculated based on the variables as previously described (10). There were two different kinds of physical activity questionnaires for the younger (three to six year olds, a parent completed questionnaire) and older children (nine to 18 year olds, self-completed questionnaire). The calculated physical activity indices were age-standardized to allow comparison across age groups. Annual family income strata at the time of enrolment was determined as follows: [category 1] <12500 Finnish marks (FIM) (~5850 EUR); [2] 15001 - 25000 FIM; [3] 25001 - 35000 FIM; [4] 35001 - 45000 FIM; [5] 45001 - 55000 FIM; [6] 55001 - 75000 FIM; [7] 75001 - 100000 FIM; [8] > 100000 FIM).

Hospital discharge data on infections
Infection-related hospitalization was defined as a hospital discharge diagnosis that included at least one International Classification of Disease (ICD) infection-related code as either a primary or secondary code. Hospitalization was defined as an admission that included at least one overnight stay. We used both primary and secondary codes to ensure the capture of all infections, an approach we and others have used previously (11).

Genotyping and construction of the genetic risk score

Genotyping was performed using the Illumina Human 670K Bead Chip, and imputation was performed using SHAPEIT v1 (12) and IMPUTE2 (13) software and the 1000G Phase I Integrated Release Version 3 as a reference panel (14).

Statistical Analyses

Baseline (in 1980) risk factor levels between individuals with and without IBD diagnoses during the follow-up were compared by t-test for continuous and chi-square test for categorical variables. Thereafter, those factors significantly associated with IBD were included in a multivariable logistic regression model. This model also included age and sex. In addition, among those factors with significant association for IBD, sensitivity analyses were performed separately for males and females, as well as for both IBD outcomes, i.e. ulcerative colitis and Crohn’s disease.

A weighted genetic risk score comprising 71 SNPs associated with HDL-C in a recent large GWAS (15) was calculated as a sum of the genotyped risk alleles or imputed allele dosages carried by an individual each multiplied by the externally reported effect size. The weighted genetic risk score was standardized according to the mean and SD before the analyses. Comparison of genetic risk score values between individuals with and without IBD was performed with t-test.
Statistical analyses were performed using SAS® 9.3 or in the case of the genetic analyses with R software version 3.3.2.
RESULTS

Characteristics of the study cohort

Baseline characteristics are shown in Table 1. At the beginning of the study in 1980, 3551 persons were included, of which 1805 (50.8%) were female. At the end of the follow-up, 49 persons (1.4%) had IBD diagnosed, of those 26 were female (Table 2). Of the IBD patients, 31 had ulcerative colitis, 12 Crohn’s disease and 6 had undetermined colitis. Sex ratios (M:F) were 1:1.1, 1:2 and 2:1 for ulcerative colitis, Crohn’s disease and undetermined colitis, respectively.

Childhood risk factors predicting subsequent IBD

Univariate age and sex adjusted comparisons between IBD diagnosis, and demographic, clinical and laboratory variables revealed statistically significant inverse correlation between HDL-cholesterol and CRP concentrations with IBD (Table 3). In a multivariable analysis including HDL-cholesterol concentration, C-reactive protein concentration, age, sex an inverse association childhood HDL-cholesterol and subsequent IBD diagnosis remained statistically significant (OR for 1-SD change in HDL-cholesterol 0.57; 95% confidence interval 0.39 -0.82, P=0.006) (Table 4).

HDL-cholesterol levels in adulthood (in 2011) were lower among IBD patients compared to the non-IBD individuals (1.22±0.32 vs. 1.32±0.32 mmol/L, P=0.02). When the results shown in Table 4 were additionally adjusted with adulthood HDL-cholesterol level (in 2011), the association between childhood HDL-cholesterol and subsequent IBD remained significant (P=0.003).

As shown in Table 5, in sex-specific analyses, the association between childhood HDL-cholesterol and IBD was significant in both genders. In outcome-specific analyses, the association was significant for ulcerative colitis.

Genetic analyses between IBD and HDL-cholesterol associated SNPs
Compared with persons without IBD, weighted genetic z-score of 71 single nucleotide polymorphism (SNP) associated with HDL cholesterol was significantly lower in IBD patients (Figure 1).

**DISCUSSION**

In the present prospective cohort study, we observed a significant inverse correlation between childhood HDL-cholesterol levels and subsequent IBD diagnosis. This finding was further confirmed by the observation that a genetically derived SNPs related to higher HDL-cholesterol levels were detected less frequently among individuals with IBD diagnosis. To our knowledge, this is the first observation linking low genetically determined childhood HDL-cholesterol level to increased IBD risk.

There is a paucity of data concerning the association between early life HDL-cholesterol levels and later risk of IBD. In contrast to our findings based on a longitudinal population-based cohort study of over 3500 children, a smaller, cross-sectional Polish study reported similar lipid profiles in 30 children with first IBD flare and 20 healthy controls (16). Among young adults (mean age 31 years), Cappello et al. observed that total and LDL-cholesterol levels were statistically significantly lower in IBD patients compared to controls whereas the difference was non-significant in HDL-cholesterol levels (17). Low HDL-cholesterol level is an established cardiovascular risk factor and previous studies have found an increased risk for coronary artery disease (CAD) in adult IBD patients. In Finnish studies, it has been observed that cardiovascular mortality risk is higher among IBD patients (18, 19). These results are concordant with Canadian and Danish studies that detected an increased coronary artery disease risk in IBD patient cohorts of over 8000 and 28 000 patients, respectively (20, 21). A study from Denmark revealed that active IBD worsened patient’s prognosis after myocardial infarction (22). However, the above-mentioned findings were not confirmed in UK and US studies (23, 24). The pathogenesis of atherosclerotic disease and the role of conventional cardiovascular risk factors, such as low HDL-cholesterol levels, are still not completely understood among IBD patients. It is noteworthy, that in addition to traditional CAD risk factors, IBD patients have also
nonconventional risk factors such as inflammatory activity of IBD and the use of disease-modifying therapy that could modify CAD morbidity and mortality (25).

The treatment of adverse lipid levels may have impact on the risk and course of IBD. The use of statins significantly decreases the risk on both new onset ulcerative colitis and Crohn’s disease (26). Statin use is also associated with reduced use of steroids in ulcerative colitis (27). The impact of statins on IBD may be due to their anti-inflammatory effect, because they also attenuate disease activity in rheumatoid arthritis (28) and systemic lupus erythematosus (29). Present results suggest that increased emphasis on HDL levels already in childhood might be beneficial.

The pathogenesis of IBD is complex. An immunological dysfunction with primary and secondary inflammatory reactions is the hallmark of IBD, and genetic and environmental factors as well as gut microbiota are involved (30). Results of a large international study suggest that genetic factors determine mainly disease location and type (ulcerative colitis, Crohn’s colitis and ileal Crohn’s disease), but environmental factors such as diet, smoking and microbiota may be even stronger contributors to IBD phenotypes (31). Lipids and lipoproteins may have impact on the risk of IBD and its course. HDL-cholesterol and its main apolipoprotein apoA-I reduce cholesterol in lipid rafts of several immune cell types leading to the modulation of key receptors involved in both innate and adaptive immune responses (32). HDL has anti-inflammatory effects as a part of innate immune system and low levels of HDL cholesterol are associated with inflammatory and immune diseases (33). Furthermore, inflammation also reportedly decreases HDL-cholesterol levels and induces changes in HDL-associated proteins possibly leading to an increased risk of atherosclerosis (34).

The strength of the present study is its study population representing a randomly selected ethnically homogeneous young age cohort with large data on socioeconomic, clinical and laboratory factors. Furthermore, the follow-up time was over thirty years and well into middle adulthood of our participants.
As IBD is most commonly diagnosed in adolescence and early adulthood (35), we are likely to have covered most cases in our sample.

A major limitation in the present study is the low number of diagnosed IBD cases with different IBD types possibly resulting in selection bias. IBD diagnoses were not acquired by clinical follow-up, but were derived from national registries. Therefore, possible misclassification of non-IBD cases might have occurred. However, earlier Finnish reports have shown that the completeness and accuracy in Finnish registers are very good, e.g. positive predictive value for gastrointestinal diseases is over 90% in the hospital discharge register(36). The accuracy of the drug reimbursement data is also high. A study group evaluated 50 randomly selected IBD reimbursement reports to assess consistency in diagnostic criteria. Only one of the 50 diagnoses was questionable (37). According to a Finnish survey, IBD patients have been diagnosed by specialist (in Finland usually a gastroenterologist or a specialist in internal medicine) in over 96% of cases (38). Moreover, the combined IBD incidence rates (CD + UC) in our cohort are in line with prior reports in Finland (4).

Another limitation is that the age of patients at the time of IBD diagnosis was not available. As the analyses were derived from a cardiovascular cohort study initiated in 1980, we were not able to include all possible risk factors for subsequent IBD in our analyses. Furthermore, the present results are generalizable only to Caucasian populations. Finally, we were not able to perform replication analyses in another cohort to confirm our findings. Further studies are required to investigate the association between HDL-cholesterol and IBD.

In conclusion, the present study revealed that low childhood HDL-cholesterol levels associate with the subsequent risk of clinical IBD. Moreover, a genetic risk score related with elevated HDL-cholesterol levels was lower among IBD patients. These data suggest that low HDL-cholesterol level is a possible risk factor for IBD and therefore warrant further research to explore the mechanisms explaining this association to
possibly provide novel therapeutic options. From a clinical point of view, it is important to evaluate whether HDL-cholesterol data can improve IBD risk prediction.

**Author contributions:**

Markku Voutilainen: study concept and design; drafting of the manuscript; analysis and interpretation of data; study supervision

Nina Hutri-Kähönen: acquisition of data; critical revision of the manuscript for important intellectual content; obtained funding

Päivi Tossavainen: acquisition of data; critical revision of the manuscript for important intellectual content; technical support

Taina Sipponen: study concept and design; interpretation of data; critical revision of the manuscript for important intellectual content

Niina Pitkänen: analysis and interpretation of data; drafting of the manuscript; critical revision of the manuscript for important intellectual content; statistical analysis

Tomi Laitinen: acquisition of data; analysis and interpretation of data; obtained funding

Eero Jokinen: acquisition of data; analysis and interpretation of data; administrative and material support

Tapani Rönnemaa: study concept and design; acquisition of data; critical revision of the manuscript for important intellectual content; obtained funding

Jorma S.A. Viikari: study concept and design; acquisition of data; critical revision of the manuscript for important intellectual content; obtained funding; study supervision

Olli T. Raitakari: study concept and design; acquisition of data; analysis and interpretation of data; critical revision of the manuscript for important intellectual content; obtained funding; study supervision

Markus Juonala: study concept and design; acquisition of data; analysis and interpretation of data; critical revision of the manuscript for important intellectual content; statistical analysis; obtained funding
Financial Disclosure: The authors have no financial relationships relevant to this article to disclose.

Conflict of Interest: The authors have no conflicts of interest to disclose.
References


Figure legend

Figure 1. Standardized weighted genetic risk score constructed using 71 variants associated with HDL-C in individuals with and without IBD. P-value for group comparison 0.01.
<table>
<thead>
<tr>
<th>Variable</th>
<th>No IBD during follow-up</th>
<th>IBD during follow-up</th>
<th>P-value</th>
</tr>
</thead>
<tbody>
<tr>
<td>N</td>
<td>3502</td>
<td>49</td>
<td></td>
</tr>
<tr>
<td>Age (years)</td>
<td>10.5±5.0</td>
<td>10.7±5.2</td>
<td>0.84</td>
</tr>
<tr>
<td>Males (%)</td>
<td>49.1</td>
<td>52.0</td>
<td>0.75</td>
</tr>
<tr>
<td>BMI (kg/m²)</td>
<td>17.9±3.1</td>
<td>17.8±2.8</td>
<td>0.91</td>
</tr>
<tr>
<td>Systolic BP (mmHg)</td>
<td>113±12</td>
<td>113±12</td>
<td>0.96</td>
</tr>
<tr>
<td>LDL-cholesterol (mmol/l)</td>
<td>3.43±0.83</td>
<td>3.30±0.82</td>
<td>0.28</td>
</tr>
<tr>
<td>HDL-cholesterol (mmol/l)</td>
<td>1.56±0.31</td>
<td>1.41±0.26</td>
<td>0.0006</td>
</tr>
<tr>
<td>Triglycerides (mmol/l)</td>
<td>0.66±0.31</td>
<td>0.69±0.27</td>
<td>0.58</td>
</tr>
<tr>
<td>Insulin (IU/l)</td>
<td>9.6±6.0</td>
<td>9.1±4.9</td>
<td>0.61</td>
</tr>
<tr>
<td>CRP (mg/l)</td>
<td>1.0±3.0</td>
<td>2.0±4.4</td>
<td>0.04</td>
</tr>
<tr>
<td>Fruit consumption (servings/week)</td>
<td>6.9±2.8</td>
<td>7.3±2.6</td>
<td>0.34</td>
</tr>
<tr>
<td>Vegetable consumption (servings/week)</td>
<td>6.3±2.9</td>
<td>6.4±2.9</td>
<td>0.78</td>
</tr>
<tr>
<td>Physical activity index</td>
<td>10.9±3.7</td>
<td>10.6±3.3</td>
<td>0.61</td>
</tr>
<tr>
<td>Birth weight (kg)</td>
<td>3.51±0.55</td>
<td>3.41±0.59</td>
<td>0.25</td>
</tr>
<tr>
<td>Breast feeding (%)</td>
<td>85.3</td>
<td>85.0</td>
<td>0.97</td>
</tr>
<tr>
<td>Rural resident (%)</td>
<td>47.2</td>
<td>52.0</td>
<td>0.42</td>
</tr>
<tr>
<td>Family income (categorical variable 1-8)</td>
<td>4.8±1.9</td>
<td>4.7±2.2</td>
<td>0.65</td>
</tr>
<tr>
<td>Parental smoking (%)</td>
<td>46.5</td>
<td>55.6</td>
<td>0.17</td>
</tr>
<tr>
<td>Childhood infection related hospitalisations(%)</td>
<td>7.6</td>
<td>10.0</td>
<td>0.47</td>
</tr>
</tbody>
</table>

Plus-minus values are mean±SD. P-values from t-test for continuous and chi-square test for categorical variables.
Table 2. Prevalence of inflammatory bowel disease

<table>
<thead>
<tr>
<th>Disease</th>
<th>Females</th>
<th>Males</th>
</tr>
</thead>
<tbody>
<tr>
<td>Inflammatory bowel disease</td>
<td>26 (1.4%)</td>
<td>23 (1.3%)</td>
</tr>
<tr>
<td>Ulcerative colitis</td>
<td>16 (0.9%)</td>
<td>15 (0.9%)</td>
</tr>
<tr>
<td>Crohn disease</td>
<td>8 (0.5%)</td>
<td>4 (0.2%)</td>
</tr>
<tr>
<td>Unspecific colitis</td>
<td>2 (0.1%)</td>
<td>4 (0.2%)</td>
</tr>
</tbody>
</table>

Table 3. Univariate associations between childhood factors (in 1980) and later IBD.

<table>
<thead>
<tr>
<th>Variable</th>
<th>OR (95% CI)</th>
</tr>
</thead>
<tbody>
<tr>
<td>BMI</td>
<td>0.94 (0.64-1.38)</td>
</tr>
<tr>
<td>Systolic BP</td>
<td>1.00 (0.72-1.39)</td>
</tr>
<tr>
<td>LDL-cholesterol</td>
<td>0.84 (0.62-1.14)</td>
</tr>
<tr>
<td>HDL-cholesterol</td>
<td>0.58 (0.42-0.79)</td>
</tr>
<tr>
<td>Triglycerides</td>
<td>1.07 (0.82-1.40)</td>
</tr>
<tr>
<td>Insulin</td>
<td>0.87 (0.60-1.25)</td>
</tr>
<tr>
<td>CRP</td>
<td>1.20 (1.01-1.43)</td>
</tr>
<tr>
<td>Fruit consumption</td>
<td>1.19 (0.87-1.61)</td>
</tr>
<tr>
<td>Vegetable consumption</td>
<td>1.02 (0.76-1.36)</td>
</tr>
<tr>
<td>Physical activity index</td>
<td>1.03 (0.85-1.25)</td>
</tr>
<tr>
<td>Birth weight</td>
<td>0.84 (0.62-1.13)</td>
</tr>
<tr>
<td>Breast feeding</td>
<td>0.96 (0.40-2.33)</td>
</tr>
<tr>
<td>Rural resident</td>
<td>1.21 (0.69-2.12)</td>
</tr>
<tr>
<td>Family income</td>
<td>0.95 (0.72-1.26)</td>
</tr>
<tr>
<td>Parental smoking</td>
<td>1.43 (0.79-2.60)</td>
</tr>
<tr>
<td>Childhood infection related hospitalisations</td>
<td>1.39 (0.53-3.65)</td>
</tr>
</tbody>
</table>

OR-values are for 1-SD change in continuous variables. All analyses are adjusted with age and sex.
Table 4. Multivariate model of childhood predictors of IBD.

<table>
<thead>
<tr>
<th>Variable</th>
<th>OR (95% CI)</th>
</tr>
</thead>
<tbody>
<tr>
<td>HDL-cholesterol</td>
<td>0.57 (0.39-0.82)</td>
</tr>
<tr>
<td>CRP</td>
<td>1.16 (0.96-1.41)</td>
</tr>
<tr>
<td>Age</td>
<td>1.02 (0.96-1.09)</td>
</tr>
<tr>
<td>Male sex</td>
<td>1.00 (0.52-1.93)</td>
</tr>
</tbody>
</table>

OR-values are from logistic regression model for 1-SD change in continuous variables.

Table 5. Sensitivity analyses for the association between childhood HDL-cholesterol and later IBD in different subgroups.

<table>
<thead>
<tr>
<th>Analysis group</th>
<th>OR (95% CI)</th>
</tr>
</thead>
<tbody>
<tr>
<td>Females</td>
<td>0.56 (0.36-0.86)</td>
</tr>
<tr>
<td>Males</td>
<td>0.59 (0.38-0.93)</td>
</tr>
<tr>
<td>Outcome: ulcerative colitis</td>
<td>0.63 (0.43-0.92)</td>
</tr>
<tr>
<td>Outcome: Crohn disease</td>
<td>0.66 (0.36-1.22)</td>
</tr>
</tbody>
</table>

OR-values are from logistic regression model for 1-SD change in continuous variables. All analyses are adjusted with age, outcome-specific analyses are also adjusted with sex.