



OPEN ACCESS

EDITED BY

Dasheng Lin,
Fujian Medical University Union Hospital,
China

REVIEWED BY

Shangmin Chen,
Shantou University, China
Sadiq Umar,
University of Illinois Chicago, United States

*CORRESPONDENCE

Xiaoqiang Yang
✉ ll006000@163.com
Lan Luo
✉ 394032462@qq.com

[†]These authors have contributed equally to this work

RECEIVED 23 June 2024

ACCEPTED 23 September 2024

PUBLISHED 09 October 2024

CITATION

Xiang J, Zheng X, Luo L and Yang X (2024)
Role of interleukin-18 in mediating the
impacts of celiac disease on osteoporosis: a
Mendelian randomization study.
Front. Immunol. 15:1453657.
doi: 10.3389/fimmu.2024.1453657

COPYRIGHT

© 2024 Xiang, Zheng, Luo and Yang. This is an open-access article distributed under the terms of the [Creative Commons Attribution License \(CC BY\)](https://creativecommons.org/licenses/by/4.0/). The use, distribution or reproduction in other forums is permitted, provided the original author(s) and the copyright owner(s) are credited and that the original publication in this journal is cited, in accordance with accepted academic practice. No use, distribution or reproduction is permitted which does not comply with these terms.

Role of interleukin-18 in mediating the impacts of celiac disease on osteoporosis: a Mendelian randomization study

Jie Xiang^{1†}, Xiaoyu Zheng^{1†}, Lan Luo^{2*} and Xiaoqiang Yang^{1*}

¹Department of Gastroenterology, The Central Hospital of Enshi Tujia and Miao Autonomous Prefecture, Enshi, China, ²Department of Anesthesiology, The Central Hospital of Enshi Tujia and Miao Autonomous Prefecture, Enshi, China

Background: Extensive observational data suggest a link between celiac disease (CeD) and osteoporosis, but the causality and mediating mechanism remain undetermined. Herein, we performed a Mendelian randomization (MR) study to address these concerns.

Methods: We obtained the summary-level statistics for CeD from a large genome-wide association study (GWAS) comprising 4,533 cases and 10,750 controls of European ancestry. The GWAS data for osteoporosis-related traits and inflammatory cytokines were derived from the UK Biobank, FinnGen, IEU OpenGWAS database, or GWAS catalog. Two-sample MR with the inverse variance-weighted methods were employed to evaluate the genetic association between CeD and osteoporosis-related traits. The potential inflammatory mediators from CeD to osteoporosis were explored using two-step mediation analyses.

Results: The primary MR analyses demonstrated causal associations between genetically predicted CeD and osteoporosis (odds ratio [OR]: 1.110, 95% confidence interval [CI]: 1.043–1.182, $p=0.001$), total body bone mineral density (β : -0.025, $p=0.039$), and osteoporotic fracture (OR: 1.124, 95% CI: 1.009–1.253, $p=0.034$). Extensive sensitivity analyses consolidated these findings. Among the candidate inflammatory cytokines, only interleukin-18 was observed to mediate the effects of CeD on osteoporosis, with an indirect OR of 1.020 (95% CI: 1.000–1.040, $p=0.048$) and a mediation proportion of 18.9%. The mediation effects of interleukin-18 could be validated in other datasets (OR: 1.015, 95% CI: 1.001–1.029, $p=0.041$). Bayesian colocalization analysis supported the role of interleukin-18 in osteoporosis.

Conclusion: The present MR study reveals that CeD is associated with an increased risk of developing osteoporosis, which may be partly mediated by upregulation of interleukin-18.

KEYWORDS

celiac disease, osteoporosis, inflammation, Mendelian randomization, mediation

Introduction

Celiac disease (CeD) is a chronic immune-mediated enteropathy triggered by intolerance to gluten proteins in genetically predisposed subjects (1). Approximately 0.7–1.4% of the general population worldwide are affected by this illness, and the prevalence appears to be raising over time (2, 3). Within the small bowel of individuals with CeD, gluten digestion products (e.g., omega-5-gliadin) penetrating into the lamina propria can directly, or after deamination by tissue transglutaminases in the submucosa, activate the innate immune system (4). This mechanism induces an abnormal inflammatory cascade followed by damage to the structure and function of intestinal tissue (5). Although CeD primarily attacks the small intestine, it is increasingly recognized as a systemic autoimmune disorder that may present with a diverse of extraintestinal comorbidities, such as type 1 diabetes, autoimmune liver disease, and psoriasis (6).

Osteoporosis is a skeletal disorder featured by low bone mineral density (BMD) and deterioration of bone microarchitecture, with a consequent increase in susceptibility to bone fragility or fractures (7). As a global prevalent disease in the elderly, osteoporosis causes more than 8.9 million pathological fractures each year, casting a heavy economical burden to many regions (8). Nearly 14.4% of patients with CeD are suffering from osteoporosis (9). Accumulating data from population-based studies have also demonstrated that CeD was associated with reduced BMD and might represent an independent risk factor for osteoporotic fracture (10). However, whether these relationships are causal or driven by shared environmental factors remains undetermined, mainly owing to the inherent drawbacks of observational study designs. Traditional observational studies are susceptible to reverse causality as they are always problematic to determine which of two associated variables is the cause; and, confounding bias is more difficult to control for because it is mainly due to social, behavioral, or physiological factors that are difficult to measure and deal with (11). Therefore, previous observational data are insufficient to establish causal insights between CeD and osteoporosis.

Additionally, the etiopathology of osteoporosis in CeD remains largely under-investigated. The osteoporosis may be caused by malabsorption of calcium or vitamin D, but also other factors such as chronic inflammation in CeD can exert an crucial role (12). CeD is characterized by an intestinal Th1 response to dietary gluten, presenting with hypersecretion of proinflammatory proteins in the damaged mucosa or serum, particularly interferon (IFN)- γ , tumor necrosis factor (TNF), interleukin (IL)-1, and IL-6 (13). These cytokines are implicated in bone metabolism as they regulate the differentiation and activation of osteoblasts or osteoclasts (14). IL-18, also known as an IFN- γ inducing factor, are produced linked to gluten intake and associated with Th1 activity in CeD (15). Recent reports have shown that IL-18 maintain a long-standing inflammation status in CeD patients (16) and can up-regulate the expression of key osteoclastogenic regulators (17), pointing towards a mediation role in the CeD-induced osteoporosis.

Mendelian randomization (MR) is an epidemiological strategy widely used to strengthen the causal inference by employing single

nucleotide polymorphisms (SNPs) as unbiased instrumental variables (IVs) for exposures (18). Because genetic variants are allocated randomly during gametogenesis and would not be modified by acquired factors, MR procedure furnishes several advantages over observational designs: 1) it ensures the temporality of exposure and outcome, preventing reverse causation; 2) it minimizes the impact of residual or unmeasured confounding factors; 3) it reflects the long-term risk estimates of exposure, as the IVs remain valid throughout a lifetime (19). MR can thus represent an analogue to randomized controlled trials that utilizes genetic variation as the method for randomization ultimately providing causal inferences. In this study, we applied the MR approach to explore the causal association between CeD and osteoporosis-related traits, and simultaneously examined the mediating relationship of inflammatory cytokines between CeD and osteoporosis.

Methods

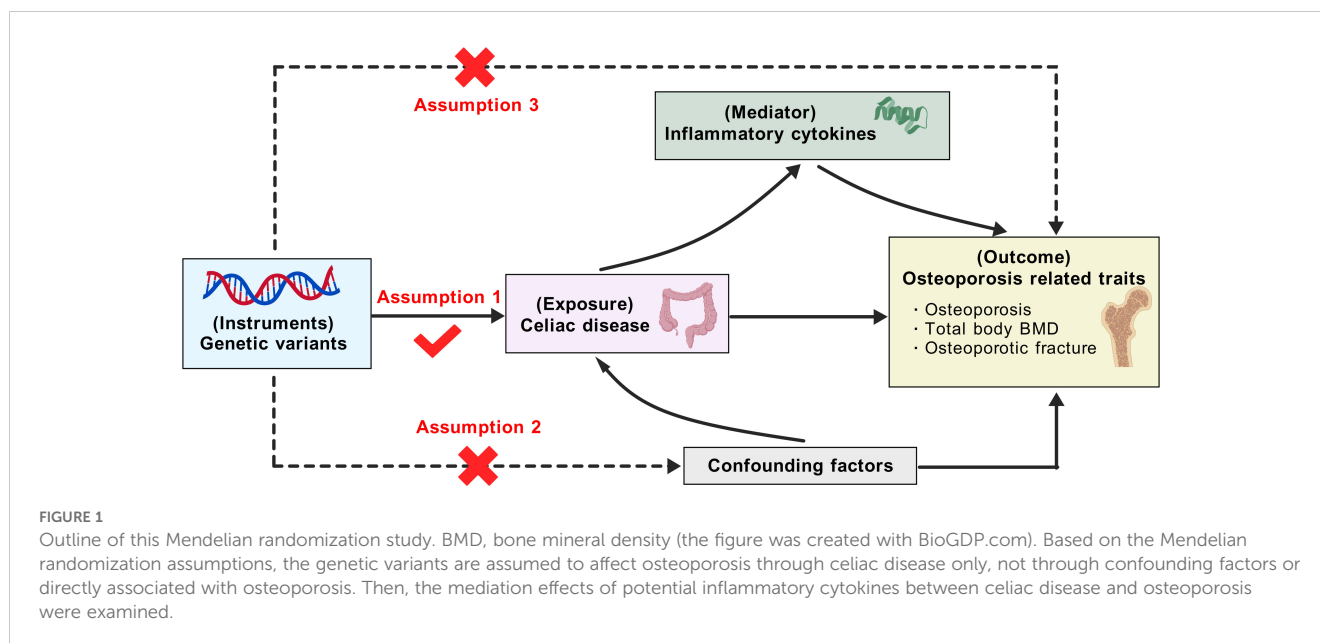
Study design

The overall design of this work was shown in [Figure 1](#). Briefly, we first employed two-sample MR methods to assess the associations of CeD with osteoporosis-related traits, including osteoporosis, total body BMD, and osteoporotic fracture. Then, two-step MR strategies were leveraged to explore the mediating effects of inflammatory candidates in the associations. MR analyses can provide unbiased causal inference if the selected IVs satisfy the following three assumptions: 1) the IVs must be strongly associated with exposures; 2) the IVs should be free of confounding factors; 3) the IVs affect the outcomes solely through the exposures. This study complies with the Strengthening the Reporting of Observational Studies in Epidemiology Using Mendelian Randomization (STROBE-MR) statement (seeing the checklist in [Supplementary Table S1](#)).

Data source

Our MR study utilized publicly accessible data from genome-wide association studies (GWAS) or databases ([Supplementary Table S2](#)). The original articles have provided the ethical clearance and consent to participates, thus there was no need for additional approvals. The summary-level genetic statistics for CeD were derived from a GWAS study involving 4,533 cases and 10,750 controls of European ancestry (IEU OpenGWAS ID: ieu-a-276) (20). The summary-level GWAS data pertaining to osteoporosis and osteoporotic fracture were acquired from the UK Biobank (21) and FinnGen (release 10) (22) consortium, respectively. The former consisted of 6,484 osteoporosis individuals and 401,279 controls, and the latter comprised 1,822 cases and 311,210 controls. The genetic statistics for total body BMD were sourced from a GWAS meta-analysis of 30 cohorts totaling 56,284 European participates (23).

Based on recent literature reviews (13, 24), we examined the mediating effects of nine inflammatory proteins in the CeD-



osteoporosis association, including IL-1 α , IL-1 β , IL-6, IL-10, IL-12, IL-18, TNF- α , TNF- β , and IFN- γ . The GWAS summary statistics correlated with these cytokines were retrieved from three meta-analyses (25–27) of pQTL, with the number of participants ranging from 3,309 to 21758 as described in [Supplementary Table S2](#). If a cytokine was measured in at least two studies, we used the pQTL data with the largest sample size.

Instrumental variable selection

For CeD, we extracted SNPs that were highly associated with this illness at a genome-wide significance level ($p < 5 \times 10^{-8}$). For each inflammatory protein, cis-pQTLs strongly correlated with the protein level ($p < 5 \times 10^{-8}$) were collected. Cis-pQTLs were defined as the pQTLs locating at ± 1 Mb from the encoding gene. To identify independent IVs, the selected genetic variants were pruned through linkage disequilibrium (LD) $R^2 < 0.001$ within 10-Mb windows, using the 1000 Genomes Project of European ancestry as reference panel (28). Confounding biases were minimized by removing any pleiotropic SNPs using the LDTrait tool (29), and SNPs that directly influenced the outcomes ($p < 5 \times 10^{-8}$) were excluded. During MR harmonization process, SNPs not present in the outcome GWAS data or those being palindromic were further discarded. The remaining genetic variants were screened out as IVs, and their strengths were assessed using the F statistics, calculated as dividing the square of β coefficient by the square of standard error.

Statistical analysis

In the primary MR analyses, we used the inverse-variance weighted (IVW) method or the Wald ratio to estimate the causal effects of exposures on outcomes. Heterogeneity was detected using

Cochrane's Q-test, with $p > 0.05$ indicating no substantial heterogeneity. We applied the MR-Egger intercept test to evaluate potential horizontal pleiotropy, where the difference between the intercept of the MR-Egger regression and zero were tested (30). The IVW approach has the highest statistical power but may be biased when pleiotropy exists; therefore, we further introduced the MR-Egger with bootstrapping (30) and the weighted median (31) methods for reanalysis. The MR-Egger approach can provide genetic estimates corrected for pleiotropy because it allows nearly all of SNPs to be horizontally pleiotropic; and, the weighted median can generate unbiased causal estimates when more than half of the IVs are valid. MR-pleiotropy residual sum and outlier (MR-PRESSO) test was also employed to detect the outliers involved in horizontal pleiotropy and produce the corrected results (32). Leave-one-out analyses were performed to assess the influence of individual SNPs on the MR results.

Two-step MR was conducted to identify the potential inflammatory proteins that mediate the effects of CeD on osteoporosis. In the first step, we performed MR analyses with the IVW method to examine the causal effects of CeD on the nine inflammatory proteins. The proteins that passed the significance threshold ($p < 0.05$) were entered into the second step, in which their causal effects on the risk of osteoporosis were investigated. The MR estimates of CeD on osteoporosis, CeD on each protein, and each protein on osteoporosis were recorded as β_0 , β_1 , and β_2 , respectively. For an inflammatory cytokine with both β_1 and β_2 being significant, we conducted mediation analysis to further elucidate whether it could mediate the effect of CeD on osteoporosis risk. Indirect effect, which refers to the effect of CeD on osteoporosis through the inflammatory cytokine, was computed using the "Product of coefficients" method (33) ($\beta_1 \times \beta_2$). We also estimated the corresponding proportion of mediation as the indirect effect divided by the total effect ($\beta_1 \times \beta_2 / \beta_0$). The 95% confidence intervals (CIs) were obtained from the delta method.

The MR-Steiger test was performed to validate the directionality of causal relationships between CeD, inflammatory mediator, and osteoporosis.

For a cytokine with significant mediating effect, Bayesian colocalization analysis (34) was conducted to reinforce the MR assumption in osteoporosis. This approach calculates the posterior probabilities (PP) for five hypothesis testing: H0 (no causal variants), H1 (causal variant for the cytokine only), H2 (causal variant for osteoporosis only), H3 (separate causal variants for the cytokine and osteoporosis), and H4 (shared causal variant for the cytokine and osteoporosis). We selected all SNPs located within ± 100 kb around the lead cis-pQTL of the cytokine encoding gene for colocalization. $PP.H4 > 0.5$ was considered evidence of colocalization, implying that colocalization is more likely than any other situations combined.

All statistical analyses were implemented using R version 4.1.0 (The R Foundation for Statistical Computing, Vienna, Austria) software. The “TwoSampleMR” (version 0.5.6), “RMeditation” (version 1.2.2), and “coloc” (version 5.2.3) R packages were used for the main analyses. A two-sided p value of < 0.05 was deemed as of significance.

Results

Instrument variables for CeD

The correlation analyses and LD clumping generated 12 independent genetic variants that were strongly associated with CeD. **Supplementary Table S3** summarized the details of these IVs for CeD. Of them, rs653178 was interfered with other traits such as smoking and diabetes; therefore, we removed this pleiotropic SNP from the subsequent MR analyses of the CeD-osteoporosis association. The F statistics ranged from 29 to 101 for the selected

instrumental SNPs, signifying that our MR analyses were robust against weak instrument bias.

CeD and osteoporosis-related traits

The MR associations between CeD and osteoporosis-related phenotypes were exhibited in **Figure 2** and **Supplementary Table S4**. The IVW-MR analyses indicated that genetically instrumented CeD was related to higher risks of developing osteoporosis (odds ratio [OR]: 1.110, 95% CI: 1.043–1.182, $p=0.001$; **Figure 2A**) and osteoporotic fracture (OR: 1.124, 95% CI: 1.009–1.253, $p=0.034$). Meanwhile, we found that genetic liability to CeD could decrease total body BMD (β : -0.025, $p=0.039$; **Figure 2B**). The results from the MR-Egger and the weighted median methods, which provide adjustments for potential pleiotropic effects, were directionally consistent with the IVW estimates albeit with noticeably wider CIs (**Supplementary Table S4**). The Cochran Q-test documented no significant heterogeneity for the above findings (**Supplementary Table S5**). There were no evidence of horizontal pleiotropy from the MR-Egger regression intercept test and the MR-PRESSO Global test (**Supplementary Table S5**), implying that the IVW estimates were credible. Leave-one-out sensitivity analyses indicated that our MR results were not driven by a single instrumental SNP (**Supplementary Table S6**), further reinforcing the robustness of the MR conclusions.

Mediation effects of inflammatory cytokines

Among the nine candidate inflammatory cytokines, we observed that only IL-6 and IL-18 could be affected by genetically

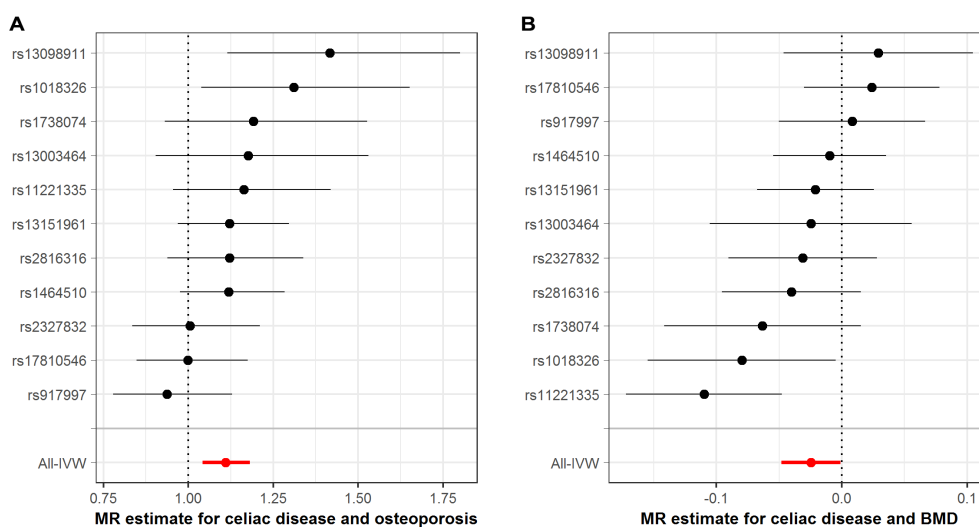


FIGURE 2

Association between genetically proxied celiac disease with osteoporosis-related traits. (A), estimates for celiac disease and osteoporosis risk; (B), estimates for celiac disease and BMD. BMD, bone mineral density; IVW, inverse-variance weighted; MR, Mendelian randomization. The dots represent the odds ratios of estimates, and the bars represent the corresponding 95% confidence intervals.

surrogated CeD. The results showed that CeD was linked to an increased expression of plasma IL-6 (β : 0.038 95% CI: 0.002–0.073, $p=0.037$) and IL-18 (β : 0.066, 95% CI: 0.018–0.114, $p=0.007$; **Supplementary Table S7**). However, there was no significant cis-pQTLs identified for IL-6 in the pQTL datasets; therefore, we only retained IL-18 for the following analyses. By using one cis-pQTL (rs5744249, located in the intron of IL18 gene) as the instrument, we found that plasma level of IL-18 was positively associated with osteoporosis risk (OR: 1.347, 95% CI: 1.115–1.626, $p=0.002$). These findings indicated that IL-18 might serve as a mediator from CeD to osteoporosis. As expected, the mediation analysis suggested that the indirect effect of IL-18 was 1.020 (95% CI: 1.000–1.040, $p=0.048$), with a mediation proportion of 18.9% in the CeD-osteoporosis association (**Figure 3**). The MR-Steiger test confirmed the causal directions from CeD to IL-18 and from IL-18 to osteoporosis (**Supplementary Table S8**). To validate the mediation effects, we analyzed another two pQTL datasets (25, 26) for IL-18 (seeing **Supplementary Table S2**). We first combined the MR estimates (β_1 or β_2) of IL-18 from each pQTL dataset using random-effect meta-

analysis and then repeated the aforementioned mediation analyses. The results also demonstrated that IL-18 could mediate the effect of CeD on osteoporosis (indirect effect, OR: 1.015, 95% CI: 1.001–1.029, $p=0.041$; **Supplementary Table S9**), with a mediation proportion of 14.0%.

Colocalization between IL-18 and osteoporosis

Considering that the above analysis identified IL-18 as a mediator between CeD and osteoporosis, we further performed colocalization analyses to investigate whether the cis-pQTLs of IL-18 shared the same casual variants with osteoporosis. The results provided evidence for colocalization between the protein expression of IL-18 and osteoporosis (PP.H4 = 0.56, **Figure 4** and **Supplementary Table S10**). This finding reinforces the aforementioned MR results and underlines that IL18 may represent a potential therapeutic target in osteoporosis.

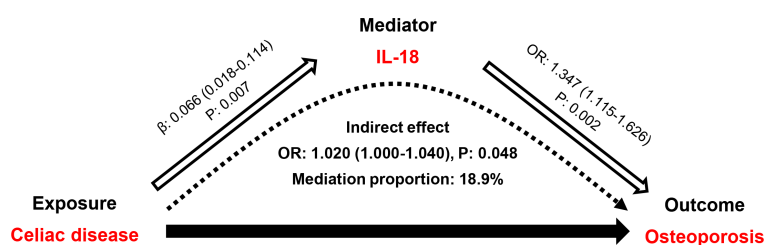


FIGURE 3 Mediation effect of IL-18 in the association between celiac disease and osteoporosis. IL-18, interleukin-18; OR, odds ratio. The indirect effect refers to the effect of celiac disease on osteoporosis through IL-18.

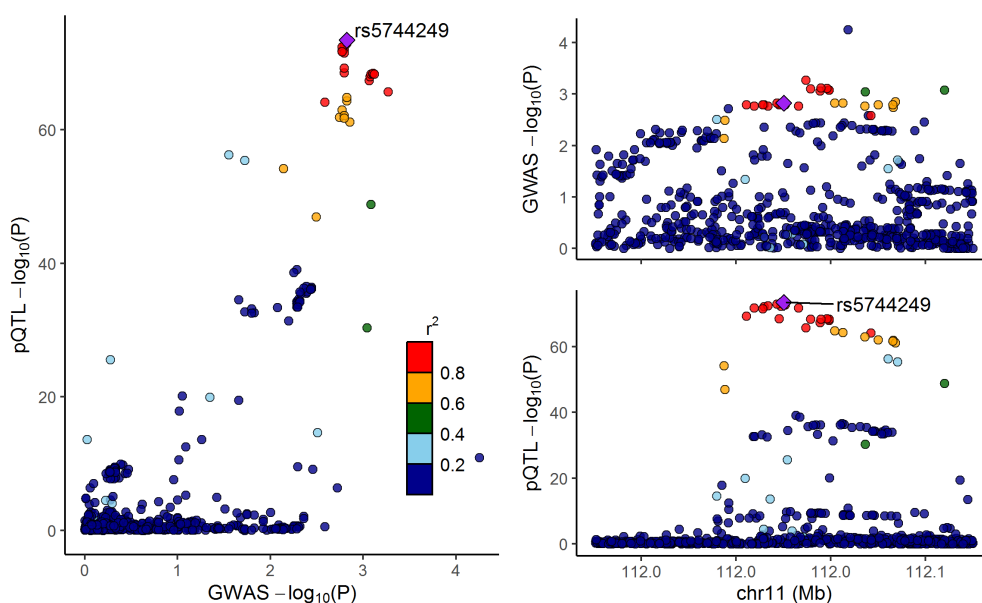


FIGURE 4 LocusCompare plot for the colocalization between interleukin-18 pQTLs and osteoporosis. The dots represent the genetic variants at chromosome 11, with color indicating the magnitude of the r^2 value. The purple rhombus stands for the shared causal variant, rs5744249.

Discussion

CeD is a chronic immuno-inflammatory disease with a broad spectrum of extraintestinal comorbidities. In the present MR investigation, we found that genetic predisposition to CeD could increase the risks of osteoporosis and osteoporotic fracture and reduce total body BMD. Additionally, plasma IL-18 levels appeared to play an important role in mediating the relationship between CeD and osteoporosis. The colocalization analysis supported the connection between IL-18 and osteoporosis.

CeD has been recognized for decades as a secondary cause of osteoporosis, but most of the relevant data were sourced from observational studies. For example, a nationwide cohort study of 103,361 individuals demonstrated that CeD was an independent risk factor for developing osteoporosis and osteoporotic fractures, both before or after the diagnosis (35). A meta-analysis of prospective cohort studies indicated that CeD at baseline conferred a 30% increase in the risk of any fractures and a 69% increase in the risk of hip fracture (36). As mentioned above, MR is effective in avoiding the bias from observational nature including residual confounding and reverse causation. The present MR analyses revealed causal links of CeD with osteoporosis-related traits, reinforcing the conclusion from conventional epidemiological studies. Our results further highlight the importance of BMD monitoring in the clinical management of patients with CeD, as it proposed in recent guidelines (37, 38).

The mechanisms underlying osteoporosis in CeD remain incompletely understood. One of the theories assumes an impaired intestinal absorption of vitamin D and calcium in CeD, leading to secondary hyperparathyroidism and subsequent osteoclast-mediated bone turnover (13). In symptomatic CeD, the bone loss appears directly associated with intestinal malabsorption of vitamin D, calcium, and other nutrients that are essential to bone health. However, low BMD can be observed even in patients with atypical or asymptomatic CeD at the time of diagnosis (39), raising the possibility of other determinants for the origin of osteoporosis in CeD. In recent years, accelerating evidence have emphasized the role of both local and systemic inflammation in the pathological process of CeD-related bone loss, characterized by a chronic increase in both mucosal and circulating pro-inflammatory cytokines (40). Pro-inflammatory cytokines may imbalance the receptor activator of nuclear kappa-B ligand (RANKL)/osteoprotegerin (OPN) pathway by lowering the OPN to RANKL ratio thus favoring osteoclastogenesis (41). Previous studies have also shown a decreased OPG/RANKL ratio under the condition of CeD, which was positively correlated with BMD at the spine (42). Among the inflammatory cytokines, we observed that IL-18 might mediate the effect of CeD on osteoporosis. IL-18 was reported to be activated at a post-translational level in patients with CeD (15), which in turn sustained a long-standing intestinal inflammation (16). As a powerful inflammatory cytokine, IL-18 can facilitate osteoclast differentiation by boosting inflammatory response via inducing the secretion of critical inflammatory factors (e.g., IFN- γ and TNF- α) as well as acting on T lymphocytes (43). IL-18 also suppress the secretion of osteogenic related proteins or transcription factors, such as Wnt-10b, Runx-2, and BMP-2 (44). Overall, IL-18 seems to play a pivotal role in creating and

maintaining a chronic immuno-inflammatory microenvironment that provokes osteoclasts and inhibits osteogenesis in CeD (45). Accordingly, administration of IL-18 antagonists can improve osteoporosis and reduce pro-inflammatory cytokines in ovariectomized mice (46). In this study, we found that the protein expression of IL-18 was colocalized with osteoporosis, further supporting its role in the bone mass deterioration in CeD patients.

Our findings undoubtedly supported the assessment of BMD for CeD at the time of diagnosis, but whether it should be routinely implemented in all patients, particularly in those being atypical or asymptomatic, remains debated in recent guidelines (40). This has risen the need for clinical or biochemical marker to select high-risk patients of developing bone loss. In the present study, we identified IL-18 as mediator between CeD and osteoporosis, reflecting that IL-18 measurement may help the risk classification of osteoporosis in atypical or asymptomatic CeD patients. Currently, a lifelong gluten-free diet (GFD) is still considered to be the mainstay treatment for patients with CeD and bone diseases (47). Nevertheless, some studies suggested that despite strict adherence to GFD, more than half of patients displayed low BMD and continued to experience a higher rate of osteoporosis, denoting that the persistent activation of inflammation should be considered in such residual risk (48, 49). As mentioned above, IL-18 is a dominate cytokine to maintain the long-standing inflammatory nature in CeD. Targeting IL-18 was found to be effective in experimental models of autoimmune diseases including inflammatory bowel disease (50) and rheumatoid arthritis (51), as well as to be safe in clinical patients (52). Alongside with these reports, our results indicated that IL-18 may also serve as a potential target for the prevention and treatment of osteoporosis in patients with CeD.

Several limitations should be acknowledged. First of all, although we have performed extensive analyses to reinforce the MR inferences, the horizontal pleiotropy cannot be completely eliminated. Secondly, due to the lack of relevant data, it is challenging to determine whether our MR findings are affected by potential confounding factors, such as age, gender, reproductive status, smoke, steroid exposure, and endocrine disorders. Thirdly, despite being documented as a potential mediator, the exact biological pathway through which IL-18 promotes osteoporosis in CeD remains unclear. This is a research topic and should be taken into consideration in future works. Fourthly, the original GWAS datasets only include European individuals, limiting the generalization of our results to other ethnics.

In summary, our MR research indicates that patients with CeD are at higher risks of developing osteoporosis, which may be partly mediated by the increased level of IL-18. These findings support the assessment of BMD in CeD management, and put forward that IL-18 may represent a potential target in the management of CeD-induced osteoporosis, which requires further exploration in future researches.

Data availability statement

The original contributions presented in the study are included in the article/Supplementary Material. Further inquiries can be directed to the corresponding authors.

Ethics statement

Ethical approval was not required for the study involving humans in accordance with the local legislation and institutional requirements. Written informed consent to participate in this study was not required from the participants or the participants' legal guardians/next of kin in accordance with the national legislation and the institutional requirements.

Author contributions

JX: Data curation, Formal analysis, Investigation, Methodology, Validation, Writing – original draft. XZ: Data curation, Formal analysis, Investigation, Methodology, Resources, Software, Validation, Writing – review & editing. LL: Conceptualization, Project administration, Resources, Supervision, Validation, Visualization, Writing – review & editing. XY: Conceptualization, Project administration, Resources, Software, Supervision, Validation, Visualization, Writing – review & editing.

Funding

The author(s) declare that no financial support was received for the research, authorship, and/or publication of this article.

References

- Ludvigsson JF, Leffler DA, Bai JC, Biagi F, Fasano A, Green PH, et al. The Oslo definitions for coeliac disease and related terms. *Gut*. (2013) 62:43–52. doi: 10.1136/gutjnl-2011-301346
- Makharia GK, Chauhan A, Singh P, Ahuja V. Review article: Epidemiology of coeliac disease. *Aliment Pharmacol Ther*. (2022) 56 Suppl 1:S3–S17. doi: 10.1111/apt.v56.s1
- King JA, Jeong J, Underwood FE, Quan J, Panaccione N, Windsor JW, et al. Incidence of celiac disease is increasing over time: A systematic review and meta-analysis. *Am J Gastroenterol*. (2020) 115:507–25. doi: 10.14309/ajg.0000000000000523
- Lebwohl B, Sanders DS, Green PHR. Coeliac disease. *Lancet*. (2018) 391:70–81. doi: 10.1016/S0140-6736(17)31796-8
- Iversen R, Sollid LM. The immunobiology and pathogenesis of celiac disease. *Annu Rev Pathol*. (2023) 18:47–70. doi: 10.1146/annurev-pathmechdis-031521-032634
- Zingone F, Bai JC, Cellier C, Ludvigsson JF. Celiac disease-related conditions: who to test? *Gastroenterology*. (2024) 167:64–78. doi: 10.1053/j.gastro.2024.02.044
- Dimai HP, Fahrleitner-Pammer A. Osteoporosis and Fragility Fractures: currently available pharmacological options and future directions. *Best Pract Res Clin Rheumatol*. (2022) 36:101780. doi: 10.1016/j.berh.2022.101780
- Wong RMY, Wong PY, Liu C, Wong HW, Chung YL, Chow SKH, et al. The imminent risk of a fracture-existing worldwide data: a systematic review and meta-analysis. *Osteoporos Int*. (2022) 33:2453–66. doi: 10.1007/s00198-022-06473-0
- Ganji R, Moghbeli M, Sadeghi R, Bayat G, Ganji A. Prevalence of osteoporosis and osteopenia in men and premenopausal women with celiac disease: a systematic review. *Nutr J*. (2019) 18:9. doi: 10.1186/s12937-019-0434-6
- Kamycheva E, Goto T, Camargo CA Jr. Celiac disease is associated with reduced bone mineral density and increased FRAX scores in the US National Health and Nutrition Examination Survey. *Osteoporos Int*. (2017) 28:781–90. doi: 10.1007/s00198-016-3791-4
- Sheehan NA, Didelez V, Burton PR, Tobin MD. Mendelian randomisation and causal inference in observational epidemiology. *PLoS Med*. (2008) 5:e177. doi: 10.1371/journal.pmed.0050177
- Lungaro L, Manza F, Costanzini A, Barbalinardo M, Gentili D, Caputo F, et al. Osteoporosis and celiac disease: updates and hidden pitfalls. *Nutrients*. (2023) 15:1089. doi: 10.3390/nu15051089

Acknowledgments

We thank the participants and investigators for providing publicly available summary statistics.

Conflict of interest

The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

Publisher's note

All claims expressed in this article are solely those of the authors and do not necessarily represent those of their affiliated organizations, or those of the publisher, the editors and the reviewers. Any product that may be evaluated in this article, or claim that may be made by its manufacturer, is not guaranteed or endorsed by the publisher.

Supplementary material

The Supplementary Material for this article can be found online at: <https://www.frontiersin.org/articles/10.3389/fimmu.2024.1453657/full#supplementary-material>

- Di Stefano M, Mengoli C, Bergonzi M, Corazza GR. Bone mass and mineral metabolism alterations in adult celiac disease: pathophysiology and clinical approach. *Nutrients*. (2013) 5:4786–99. doi: 10.3390/nu5114786
- Krupa-Kozak U. Pathologic bone alterations in celiac disease: etiology, epidemiology, and treatment. *Nutrition*. (2014) 30:16–24. doi: 10.1016/j.nut.2013.05.027
- Salvati VM, MacDonald TT, Bajaj-Elliott M, Borrelli M, Staiano A, Auricchio S, et al. Interleukin 18 and associated markers of T helper cell type 1 activity in coeliac disease. *Gut*. (2002) 50:186–90. doi: 10.1136/gut.50.2.186
- Leon AJ, Garrote JA, Blanco-Quiros A, Calvo C, Fernandez-Salazar L, Del Villar A, et al. Interleukin 18 maintains a long-standing inflammation in coeliac disease patients. *Clin Exp Immunol*. (2006) 146:479–85. doi: 10.1111/j.1365-2249.2006.03239.x
- Zhang W, Cong XL, Qin YH, He ZW, He DY, Dai SM. IL-18 upregulates the production of key regulators of osteoclastogenesis from fibroblast-like synoviocytes in rheumatoid arthritis. *Inflammation*. (2013) 36:103–9. doi: 10.1007/s10753-012-9524-8
- Sanderson E, Glymour MM, Holmes MV, Kang H, Morrison J, Munafa MR, et al. Mendelian randomization. *Nat Rev Methods Primers*. (2022) 2:6. doi: 10.1038/s43586-021-00092-5
- Lawlor DA, Harbord RM, Sterne JA, Timpson N, Davey Smith G. Mendelian randomization: using genes as instruments for making causal inferences in epidemiology. *Stat Med*. (2008) 27:1133–63. doi: 10.1002/sim.v27:8
- Dubois PC, Trynka G, Franke L, Hunt KA, Romanos J, Curtotti A, et al. Multiple common variants for celiac disease influencing immune gene expression. *Nat Genet*. (2010) 42:295–302. doi: 10.1038/ng.543
- Gagliano Taliun SA, VandeHaar P, Boughton AP, Welch RP, Taliun D, Schmidt EM, et al. Exploring and visualizing large-scale genetic associations by using PheWeb. *Nat Genet*. (2020) 52:550–2. doi: 10.1038/s41588-020-0622-5
- Kurki MI, Karjalainen J, Palta P, Sipila TP, Kristiansson K, Donner KM, et al. FinnGen provides genetic insights from a well-phenotyped isolated population. *Nature*. (2023) 613:508–18. doi: 10.1038/s41586-022-05473-8
- Medina-Gomez C, Kemp JP, Trajanoska K, Luan J, Chesi A, Ahluwalia TS, et al. Life-course genome-wide association study meta-analysis of total body BMD and

- assessment of age-specific effects. *Am J Hum Genet.* (2018) 102:88–102. doi: 10.1016/j.ajhg.2017.12.005
24. Skoracka K, Hryhorowicz S, Tovoli F, Raiteri A, Rychter AM, Slomski R, et al. Genetic, immunological, dietary, gut microbiota, and environmental determinants of osteoporosis in the course of celiac disease: which factor plays the first violin in this orchestra? *Calcif Tissue Int.* (2024) 114:98–109. doi: 10.1007/s00223-023-01155-3
25. Ahola-Olli AV, Wurtz P, Havulinna AS, Aalto K, Pitkanen N, Lehtimäki T, et al. Genome-wide association study identifies 27 loci influencing concentrations of circulating cytokines and growth factors. *Am J Hum Genet.* (2017) 100:40–50. doi: 10.1016/j.ajhg.2016.11.007
26. Zhao JH, Stacey D, Eriksson N, Macdonald-Dunlop E, Hedman AK, Kalnapenkis A, et al. Genetics of circulating inflammatory proteins identifies drivers of immune-mediated disease risk and therapeutic targets. *Nat Immunol.* (2023) 24:1540–51. doi: 10.1038/s41590-023-01588-w
27. Folkersen L, Gustafsson S, Wang Q, Hansen DH, Hedman AK, Schork A, et al. Genomic and drug target evaluation of 90 cardiovascular proteins in 30,931 individuals. *Nat Metab.* (2020) 2:1135–48. doi: 10.1038/s42255-020-00287-2
28. 1000 Genomes Project Consortium, Abecasis GR, Altshuler D, Auton A, Brooks LD, Durbin RM, et al. A map of human genome variation from population-scale sequencing. *Nature.* (2010) 467:1061–73. doi: 10.1038/nature09534
29. Machiela MJ, Chanock SJ. LDlink: a web-based application for exploring population-specific haplotype structure and linking correlated alleles of possible functional variants. *Bioinformatics.* (2015) 31:3555–7. doi: 10.1093/bioinformatics/btv402
30. Bowden J, Davey Smith G, Burgess S. Mendelian randomization with invalid instruments: effect estimation and bias detection through Egger regression. *Int J Epidemiol.* (2015) 44:512–25. doi: 10.1093/ije/dyv080
31. Bowden J, Davey Smith G, Haycock PC, Burgess S. Consistent estimation in mendelian randomization with some invalid instruments using a weighted median estimator. *Genet Epidemiol.* (2016) 40:304–14. doi: 10.1002/gepi.2016.40.issue-4
32. Verbanck M, Chen CY, Neale B, Do R. Detection of widespread horizontal pleiotropy in causal relationships inferred from Mendelian randomization between complex traits and diseases. *Nat Genet.* (2018) 50:693–98. doi: 10.1038/s41588-018-0099-7
33. MacKinnon DP, Fairchild AJ, Fritz MS. Mediation analysis. *Annu Rev Psychol.* (2007) 58:593–614. doi: 10.1146/annurev.psych.58.110405.085542
34. Giambartolomei C, Vukcevic D, Schadt EE, Franke L, Hingorani AD, Wallace C, et al. Bayesian test for colocalisation between pairs of genetic association studies using summary statistics. *PLoS Genet.* (2014) 10:e1004383. doi: 10.1371/journal.pgen.1004383
35. Hansen S, Schwarz P, Rumessen J, Linneberg A, Karhus LL. Osteoporosis and bone fractures in patients with celiac disease: A nationwide cohort study. *Bone.* (2023) 177:116913. doi: 10.1016/j.bone.2023.116913
36. Heikkilä K, Pearce J, Maki M, Kaukinen K. Celiac disease and bone fractures: a systematic review and meta-analysis. *J Clin Endocrinol Metab.* (2015) 100:25–34. doi: 10.1210/jc.2014-1858
37. Downey L, Houten R, Murch S, Longson D. Guideline Development Group. Recognition, assessment, and management of coeliac disease: summary of updated NICE guidance. *BMJ.* (2015) 351:h4513. doi: 10.1136/bmj.h4513
38. Al-Toma A, Volta U, Auricchio R, Castillejo G, Sanders DS, Cellier C, et al. European Society for the Study of Coeliac Disease (ESsCD) guideline for coeliac disease and other gluten-related disorders. *United Eur Gastroenterol J.* (2019) 7:583–613. doi: 10.1177/2050640619844125
39. Mazure R, Vazquez H, Gonzalez D, Mautalen C, Pedreira S, Boerr L, et al. Bone mineral affection in asymptomatic adult patients with celiac disease. *Am J Gastroenterol.* (1994) 89:2130–4.
40. Kondapalli AV, Walker MD. Celiac disease and bone. *Arch Endocrinol Metab.* (2022) 66:756–64. doi: 10.20945/2359-399700000561
41. Khosla S. Minireview: the OPG/RANKL/RANK system. *Endocrinology.* (2001) 142:5050–5. doi: 10.1210/endo.142.12.8536
42. Fiore CE, Pennisi P, Ferro G, Ximenes B, Privitelli L, Mangiafico RA, et al. Altered osteoprotegerin/RANKL ratio and low bone mineral density in celiac patients on long-term treatment with gluten-free diet. *Horm Metab Res.* (2006) 38:417–22. doi: 10.1055/s-2006-944548
43. Tao Z, Wang J, Wen K, Yao R, Da W, Zhou S, et al. Pyroptosis in osteoblasts: A novel hypothesis underlying the pathogenesis of osteoporosis. *Front Endocrinol (Lausanne).* (2021) 11:548812. doi: 10.3389/fendo.2020.548812
44. Jiang N, An J, Yang K, Liu J, Guan C, Ma C, et al. NLRP3 inflammasome: A new target for prevention and control of osteoporosis? *Front Endocrinol (Lausanne).* (2021) 12:752546. doi: 10.3389/fendo.2021.752546
45. Chen T, Jin L, Li J, Liu Y. Pyroptosis mediates osteoporosis via the inflammation immune microenvironment. *Front Immunol.* (2024) 15:1371463. doi: 10.3389/fimmu.2024.1371463
46. Mansoori MN, Shukla P, Kakaji M, Tyagi AM, Srivastava K, Shukla M, et al. IL-18BP is decreased in osteoporotic women: Prevents inflammasome mediated IL-18 activation and reduces Th17 differentiation. *Sci Rep.* (2016) 6:33680. doi: 10.1038/srep33680
47. Al-Toma A, Herman A, Lems WF, Mulder CJJ. The dietary and non-dietary management of osteoporosis in adult-onset celiac disease: current status and practical guidance. *Nutrients.* (2022) 14:4554. doi: 10.3390/nu14214554
48. Sayar S, Aykut H, Kaya O, Kuruz K, Ak C, Gokcen P, et al. Bone mineral density screening and the frequency of osteopenia/osteoporosis in turkish adult patients with celiac disease. *Turk J Gastroenterol.* (2021) 32:600–7. doi: 10.5152/tjg.2021.20313
49. Di Stefano M, Bergonzi M, Benedetti I, De Amici M, Torre C, Brondino N, et al. Alterations of inflammatory and matrix production indices in celiac disease with low bone mass on long-term gluten-free diet. *J Clin Gastroenterol.* (2019) 53:e221–6. doi: 10.1097/MCG.0000000000001032
50. Ikegami S, Maeda K, Urano T, Mu J, Nakamura M, Yamamura T, et al. Monoclonal antibody against mature interleukin-18 ameliorates colitis in mice and improves epithelial barrier function. *Inflammation Bowel Dis.* (2024) 30:1353–66. doi: 10.1093/ibd/izad292
51. Plater-Zyberk C, Joosten LA, Helsen MM, Sattoune-Roche P, Siegfried C, Alouani S, et al. Therapeutic effect of neutralizing endogenous IL-18 activity in the collagen-induced model of arthritis. *J Clin Invest.* (2001) 108:1825–32. doi: 10.1172/JCI200112097
52. McKie EA, Reid JL, Mistry PC, DeWall SL, Abberley L, Ambery PD, et al. A study to investigate the efficacy and safety of an anti-interleukin-18 monoclonal antibody in the treatment of type 2 diabetes mellitus. *PLoS One.* (2016) 11:e0150018. doi: 10.1371/journal.pone.0150018