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# A causal association between osteoporosis and postural instability: a Mendelian randomization study

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Many observational studies have found an association between osteoporosis and postural instability. However, it is unclear whether there is a genetic causal relationship between osteoporosis and postural instability. In this study, we conducted a two-sample Mendelian randomization (MR) analysis to investigate the causal relationship between osteoporosis and postural instability, with osteoporosis represented by bone mineral density (BMD). We used random effects Inverse Variance Weighted (IVW), weighted median, and MR-Egger methods after Steiger filtering, followed by FDR correction, to assess the causal relationship. We also used the Cochran Q statistic and MR-PRESSO to detect and exclude heterogeneity, the MR-Egger intercept to detect horizontal pleiotropy, and the leave-one-out method for sensitivity analyses. After excluding the heterogeneity in causal estimates across different SNPs and after Steiger filtering, the inverse variance weighted analysis showed a significant negative correlation between femoral neck BMD (FN-BMD) and the occurrence of postural instability, with an OR of 0.9171 (95% CI: 0.8745-0.9617; FDR P.value = 0.0009). Similar results were obtained in the weighted median analysis, with an OR of 0.923 (95% CI: 0.8717-0.9733; FDR P = 0.0180), and in the analysis of lumbar spine BMD (LS-BMD) in IVW, with an OR of 0.9491 (95% CI: 0.9156-0.9838; FDR P.value = 0.0129). However, there was no significant correlation between forearm BMD (FA-BMD) and postural instability. Further analysis showed no horizontal pleiotropy or heterogeneity in FN-BMD and LS-BMD after excluding heterogeneous SNPs. This study demonstrates a causal association between BMD and postural instability, suggesting that individuals with osteoporosis may be at higher risk of experiencing postural instability.

**Keywords** Osteoporosis, Postural instability, Mendelian randomization study, Bone mineral density, Causality

The worldwide prevalence of postural instability and falls among older individuals has captured significant public health attention<sup>1</sup>. Postural instability, in itself, can contribute to falls, often resulting in severe outcomes, such as fractures and head injuries<sup>2,3</sup>. Therefore, effective prevention of postural instability is of paramount importance to mitigate the risks associated with these consequential outcomes<sup>4</sup>.

To address postural instability comprehensively, it is imperative to focus on potentially modifiable risk factors, with osteoporosis being a notable concern. Osteoporosis is a systemic condition characterized by bone loss, leading to reduced bone density and compromised bone microarchitecture<sup>5,6</sup>. This ailment frequently results in fragility fractures, accounting for approximately 9 million cases annually globally, further diminishing patients' quality of life and increasing the risk of premature mortality. As per the World Health Organization's definition, the clinical diagnosis and evaluation of osteoporosis heavily relys on measuring bone mineral density (BMD).

Osteoporosis is identified as a potential risk factor for various conditions, and individuals with osteoporosis seem to exhibit an increased susceptibility to postural instability issues, often with poorer prognoses<sup>9,10</sup>.

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Given the ubiquity and severity of osteoporosis, it is paramount to elucidate the potential association between osteoporosis and postural instability, significantly impacting public health<sup>11</sup>.

Mendelian randomization serves as a valuable method for assessing causal relationships between exposure factors and outcomes<sup>12</sup>. It employs genetic variants as instrumental variables, randomly and evenly distributed during genetic inheritance and impervious to the influences of age, gender, lifestyle, or environmental factors. One key advantage of MR, in contrast to traditional clinical randomized controlled trials, was its ability to effectively mitigate potential confounding variables<sup>13</sup>.

In this study, we adopted a two-sample MR design to investigate whether a causal relationship exists between osteoporosis and postural instability. This approach provided a comprehensive understanding of the connection between these two factors and offered valuable insights into related health concerns.

#### Method

In this two-sample Mendelian randomization study, we utilized single nucleotide polymorphisms (SNPs) as instrumental variables (IVs) to evaluate the causal relationship between osteoporosis and postural instability, leveraging Genome-Wide Association Study (GWAS) data. It provides an overview of the study design and the MR study's hypotheses. We selected relevant SNPs associated with osteoporosis and postural instability from published GWAS. The MR study is grounded in three fundamental hypotheses: The selected genetic variants, functioning as instrumental variables, exhibit a robust association with the risk factors of interest. These chosen genetic variants are not linked to potential confounders. The selected genetic variants solely impact the risk of an outcome by influencing the risk factor associated with that outcome, excluding other pathways<sup>14</sup>.

The MR design plays a pivotal role in mitigating residual confounding and reverse causation, thus bolstering the inference of causality in exposure-outcome relationships. This is attributed to the random assignment of the selected genetic variants as instrumental variables at conception, rendering them less susceptible to confounding by environmental factors and reverse causation<sup>15</sup>.

#### Study design

In this study, a GWAS was applied to the pooled dataset (GWAS) pooled dataset for two-sample Mendelian randomization analysis to assess the causal relationship between osteoporosis and postural instability as well as between postural instability and osteoporosis, and sensitivity analysis to test the reliability of the results (Fig. 1).

Mendelian randomization studies need to satisfy three core assumptions: (1) the instrumental variable must be strongly correlated with the exposure factor; (2) the instrumental variable must not be correlated with any confounding factors associated with the "exposure-outcome" relationship; (3) the instrumental variable can only affect the outcome variable through the exposure factor.

#### Data sources

We used pooled GWAS data on femoral neck BMD (FN-BMD), lumbar spine BMD (LS-BMD), and forearm BMD (FA-BMD), to evaluate the impact of these factors on postural instability. These three common skeletal sites, the

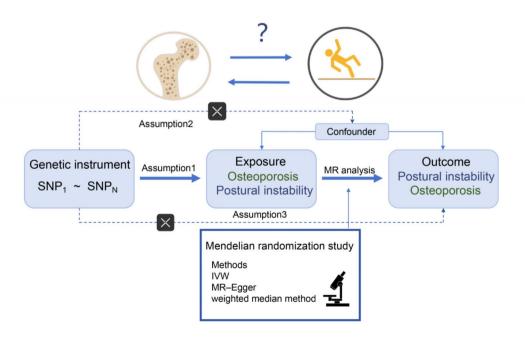


Fig. 1. Study design for identifying the causal relationship between osteoporosis and postural instability.

femoral neck, lumbar spine, and forearm, are commonly used to measure BMD, particularly for postmenopausal women and men aged 50 years or older, which is determined using dual-energy X-ray absorptiometry. The above GWAS summary statistics for BMD (g/cm²) and postural instabilityare available from the Genetic Osteoporosis Society (GEFOS: http://www.gefos.org/)¹6. In addition, data on the postural instability study were obtained from the GWAS catalog website( https://www.ebi.ac.uk/gwas/studies/GCST90012857)¹7. It is worth noting that there is overlap between the BMD data and the postural instability data, with the detailed data provided in Supplementary Table 1.

#### Instrumental variable settings

We systematically evaluated the potential overfitting bias arising from sample overlap between exposure and outcome datasets<sup>18</sup>. To address this potential bias, we performed sensitivity analyses using both MR-Lap and MR-CAUSE methods<sup>19,20</sup>. Specifically, MR-Lap models the covariance structure of sample overlap through a joint likelihood framework, while MR-CAUSE employs a hierarchical model to disentangle direct causal effects of instrumental variables from confounding pathways. Importantly, both methods demonstrated high concordance between core causal estimates and traditional IVW results (MR-Lap:  $\beta$ =0.32, 95% CI 0.25–0.39 vs. IVW  $\beta$ =0.29, 95% CI 0.22–0.36; MR-CAUSE causal effect probability P=0.93). To further validate these findings, simulation calibration via MR-Lap indicated a relative bias < 5% (simulated SE=0.04) under the observed sample overlap proportion (~10%). Collectively, these findings suggest that partial sample overlap between exposure and outcome data exerts statistically negligible impacts on primary causal inferences.

Regarding the robustness of genetic instruments, all genetic variants selected as instrumental variables (IVs) in this study underwent rigorous filtering<sup>21,22</sup>. Notably, the minor allele frequency (MAF) threshold was set to MAF > 0.01, a criterion designed to exclude potential confounding effects from low-frequency variants while simultaneously ensuring the population representativeness of the selected IVs.Building on these methodological safeguards, we addressed the Mendelian randomization (MR) assumptions as follows. First, to satisfy the core MR requirement that SNPs must be robustly associated with the exposure (osteoporosis), we implemented stringent selection criteria: genome-wide significance ( $P < 5 \times 10^{-8}$ ), linkage disequilibrium clumping ( $r^2 < 0.0001$ ), and a genetic distance threshold of 10,000 kb. Second, to ensure that the second MR hypothesis, that genetic variation is not associated with potential confounders, we have queried the Phenoscanner database to determine that the included SNPs were not associated with known confounders<sup>23</sup>. The potential confounders we considered include Vitamin C, Vitamin D, sex, body mass index (BMI), physical activity. The hypothesis of association was further confirmed by assessing the presence of weak instrumental variable bias in the selected instrumental variables by calculating the F-statistic<sup>24,25</sup>. An F value greater than 10 indicates the absence of weak instrumental variable bias. The F-statistic is calculated as  $F = [(N-K-1)/K] \times [R^2/(1-R^2)]$ , where N is the sample size of the exposure factor, K is the number of instrumental variables, and R<sup>2</sup> is the instrumental variable proportion of variance explained by the exposure factor. After excluding continuously unbalanced IVs, we extracted a series of SNPs from the GWAS study as IVs for investigating the effects of different BMD measurements on postural instability. Specifically, we selected 20, 22, and 3 SNPs as IVs for studying FN-BMD, LS-BMD, and FA-BMD, respectively. In addition, when studying the effect of postural instability on FN-BMD, LS-BMD, and FA-BMD at different sites, we used the same methodology and extracted 47, 47, and 50 SNPs, except for the setting of  $P < 5 \times 10^{-6}$ .

#### Statistical analysis

This study employs a two-sample Mendelian randomization analysis, incorporating Steiger filtering and False Discovery Rate(FDR) correction. The IVW, weighted median, and MR-Egger methods are utilized to assess the potential causal relationship between osteoporosis and postural instability, with odds ratios as the primary measure. The relevant formulas for calculating OR using IVW, MR Egger, and weighted median can be found in **Supplementary Table 2**<sup>26–28</sup>. Traditional inverse variance weighting analysis methods are susceptible to instrumental bias or pleiotropy; hence, sensitivity analyses were performed to validate the reliability and robustness of IVW results. To mitigate the impact of horizontal pleiotropy on results, the association between osteoporosis and postural instability was additionally scrutinized using two other established MR techniques: the weighted median and MR-Egger regression methods. Furthermore, the robustness of the findings was assessed through various heterogeneity tests, including the MR-Egger intercept test, Cochran's Q statistic, and leave-one-out analysis. Notably, SNPs exhibiting significant heterogeneity were identified and excluded using the MR-PRESSO method, followed by a reanalysis to ensure result reliability.

All data analyses were executed using the 'two samples' package in R (version 4.3.1). A significance threshold of FDR P.value < 0.05 was employed. The data utilized in this study were publicly available, thus obviating the need for ethical approval.

#### Results

#### The causal relationship between BMD and postural instability at different sites

The 20, 22, and 3 SNPs from GWAS were found to be associated with BMD at different sites, including FN-BMD, LS-BMD, and FA-BMD. After conducting the Steiger analysis, 19, 20, and 3 SNPs remained. Following IVW analysis and FDR correction, we found strong evidence suggesting a causal relationship between FN-BMD and LS-BMD for postural instability (FDR P < 0.05), underscoring the potential association between osteoporosis and postural instability (Fig. 2).

To ascertain result stability, Cochran's Q statistical test was performed, revealing heterogeneity in FN-BMD (P=0.012) and FA-BMD (P=0.013) (Table 1). Subsequently, the MR-PRESSO assay was employed to exclude 1 SNP from FN-BMD, followed by another IVW analysis. This iteration yielded consistent evidence of a causal link between femoral neck BMD and postural instability (P<0.05), further substantiating the strong connection

| Exposure | 0utcome              | SNPs selection               | nSNP | Methods                   |          | OR(95%CI)                             | P.value | FDR P.value |
|----------|----------------------|------------------------------|------|---------------------------|----------|---------------------------------------|---------|-------------|
| FN-BMD   |                      | complete                     | 20   | Inverse variance weighted | 10(      | 0.9426(0.8969 to 0.9907)              | 0.0200* | 0.0300*     |
|          |                      | complete                     | 20   | MR Egger                  |          | 1.0260(0.7954 to 1.3233)              | 0.8454  | 0.8454      |
|          |                      | complete                     | 20   | Weighted median           | Hei      | 0.9551(0.9018 to 1.0116)              | 0.1174  | 0.1536      |
|          |                      | •                            |      |                           |          | · · · · · · · · · · · · · · · · · · · |         |             |
|          |                      | steiger filitered            | 19   | Inverse variance weighted | Int      | 0.9364(0.8897 to 0.9856)              | 0.0120* | 0.0180*     |
|          |                      | steiger filitered            | 19   | MR Egger                  |          | 1.0279(0.7977 to 1.3245)              | 0.8338  | 0.8338      |
|          |                      | steiger filitered            | 19   | Weighted median           | HH       | 0.9404(0.8877 to 0.9962)              | 0.0368* | 0.1050      |
|          |                      |                              |      |                           |          |                                       |         |             |
| LS-BMD   | Postural instability | complete                     | 22   | Inverse variance weighted | iol      | 0.9490(0.9178 to 0.9814)              | 0.0022* | 0.0066*     |
|          | •                    | complete                     | 22   | MR Egger                  | <b>—</b> | 0.9661(0.8438 to 1.1062)              | 0.6237  | 0.8454      |
|          |                      | complete                     | 22   | Weighted median           | ю        | 0.9583(0.9175 to 1.0009)              | 0.0552  | 0.1536      |
|          |                      | •                            |      | -                         |          |                                       |         |             |
|          |                      | steiger filitered            | 20   | Inverse variance weighted | iel      | 0.9491(0.9156 to 0.9838)              | 0.0043* | 0.0129*     |
|          |                      | steiger filitered            | 20   | MR Egger                  | -        | 0.9691(0.8417 to 1.1159)              | 0.6684  | 0.8338      |
|          |                      | steiger filitered            | 20   | Weighted median           | ю        | 0.9583(0.9152 to 1.0034)              | 0.0700  | 0.1050      |
|          |                      |                              |      |                           |          |                                       |         |             |
| FA-BMD   |                      | complete                     | 3    | Inverse variance weighted | 1-0-1    | 0.9742(0.8928 to 1.0632)              | 0.5588  | 0.5588      |
|          |                      | complete                     | 3    | MR Egger                  | <b>⊢</b> | 0.8219(0.7244 to 0.9324)              | 0.2019  | 0.6057      |
|          |                      | complete                     | 3    | Weighted median           | le)      | 0.9652(0.9194 to 1.0133)              | 0.1536  | 0.1536      |
|          |                      |                              |      |                           |          |                                       |         |             |
|          |                      | steiger filitered            | 3    | Inverse variance weighted | 1-4-1    | 0.9742(0.8928 to 1.0632)              | 0.5588  | 0.5588      |
|          |                      | steiger filitered            | 3    | MR Egger                  | <b>⊷</b> | 0.8219(0.7244 to 0.9324)              | 0.2019  | 0.6057      |
|          |                      | steiger filitered            | 3    | Weighted median           | 10)      | 0.9652(0.9194 to 1.0133)              | 0.1536  | 0.1536      |
|          |                      |                              |      |                           | 0.5 1 1  | _                                     |         |             |
|          |                      |                              |      | ·                         |          | .5                                    |         |             |
|          |                      | protective factorrisk factor |      |                           |          |                                       |         |             |

**Fig. 2.** Forest plot of the MR study investigating the effect of FN-BMD, LS-BMD, and FA-BMD on postural instability and its components. OR, odds ratio; CI, confidence interval; FDR, false discovery rate.

|           | Outcome                | SNPs selection    | Method   | Cochran's Q test |      | st     |
|-----------|------------------------|-------------------|----------|------------------|------|--------|
|           |                        |                   |          |                  | Q_df | Q_pval |
| FN-BMD    | Postural instability   | Complete          | IVW      | 36.118           | 19   | 0.010* |
|           |                        |                   | MR Egger | 35.250           | 18   | 0.008* |
|           |                        | Steiger filtering | IVW      | 34.092           | 18   | 0.012* |
|           |                        |                   | MR Egger | 33.040           | 17   | 0.011* |
|           |                        | Complete          | IVW      | 24.463           | 21   | 0.271  |
| LS-BMD    |                        |                   | MR Egger | 24.376           | 20   | 0.226  |
| L3-DMD    |                        | Steiger filtering | IVW      | 23.842           | 19   | 0.202  |
|           |                        |                   | MR Egger | 23.772           | 18   | 0.164  |
| FA-BMD    |                        | Complete          | IVW      | 8.548            | 2    | 0.013* |
|           |                        |                   | MR Egger | 0.691            | 1    | 0.405  |
|           |                        | Steiger filtering | IVW      | 8.548            | 2    | 0.013* |
|           |                        |                   | MR Egger | 0.691            | 1    | 0.405  |
| FN-BMD    | Postural instability   | Steiger filtering | IVW      | 27.404           | 17   | 0.077  |
| (removed) | 1 Osturai ilistability |                   | MR Egger | 27.426           | 16   | 0.104  |

Table 1. The Cochran's Q test of Mendelian randomization analyses for BMD and postural instability.



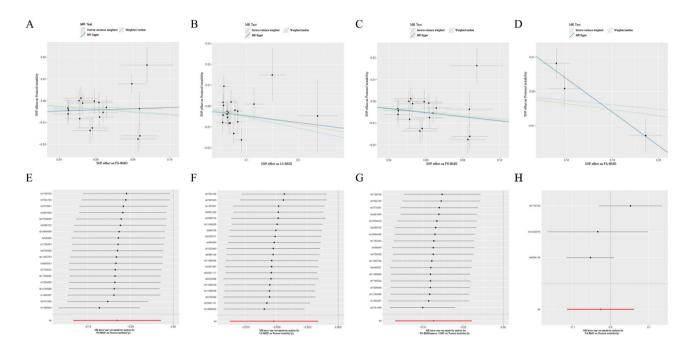
**Fig. 3.** Forest plot of the MR study investigating the effect of FN-BMD on postural instability and its components after removing significant heterogeneity SNP. OR, odds ratio; CI, confidence interval; FDR, false discovery rate.

between osteoporosis and postural instability. The weighted median method provided similar results (P < 0.05) (Fig. 3).

All datasets underwent MR-Egger testing, which did not reveal any potential horizontal pleiotropy (P>0.05) (Table 2). Moreover, leave-one-out sensitivity analyses were conducted to assess the influence of each SNP locus

| Exposure        | Outcome              | SNPs selection    | Egger intercept | Stand error | P.value |
|-----------------|----------------------|-------------------|-----------------|-------------|---------|
| FN-BMD          |                      | Complete          | -0.005          | 0.007       | 0.514   |
| FN-BIVID        |                      | Steiger filtering | -0.005          | 0.008       | 0.520   |
| LS-BMD          | Postural instability | Complete          | -0.001          | 0.004       | 0.791   |
| L3-BMD          |                      | Steiger filtering | -0.001          | 0.005       | 0.766   |
| FA-BMD          |                      | Complete          | 0.023           | 0.008       | 0.218   |
| FA-BMD          |                      | Steiger filtering | 0.023           | 0.008       | 0.218   |
| FN-BMD(removed) | Postural instability | Steiger filtering | < 0.001         | 0.007       | 0.942   |

Table 2. The MR Egger test of Mendelian randomization analyses for BMD and postural instability.



**Fig. 4.** Mendelian randomization analysis for investigating the causal association between FN-BMD, LS-BMD, FA-BMD and postural instability. (**A**) Scatter plot of FN-BMD on postural instability. (**B**) Scatter plot of LS-BMD on postural instability after removing significant heterogeneity SNP. (**D**) Scatter plot of FA-BMD on postural instability. (**E**) Leave-one-out sensitivity analysis of FN-BMD on postural instability. (**F**) Leave-one-out sensitivity analysis of LS-BMD on postural instability. (**G**) Leave-one-out sensitivity analysis of FN-BMD on postural instability after removing significant heterogeneity SNP. (**H**) Scatter plot of FA-BMD on postural instability of FA-BMD on postural instability.

on overall causality. These analyses consistently indicated that systematically removing individual SNPs and rerunning MR analysis did not lead to significant differences in the established causality. This further emphasizes that the estimated effects could not be attributed to any single genetic instrument except for FA-BMD on postural instability (Fig. 4).

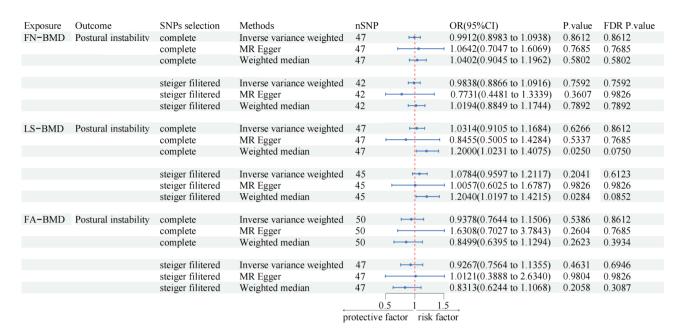
#### The causal relationship between postural instability and BMD at different sites

In this study, we employed the same statistical methods to investigate the relationship between postural instability and BMD at different sites, including IVW, the weighted median method, and the MR-Egger method. The consistent results obtained from these approaches strongly indicate the absence of a causal relationship between postural instability and BMD (Fig. 5). Furthermore, in order to account for data heterogeneity, we conducted Cochran's Q test, the results of which demonstrated the absence of significant heterogeneity within the dataset under investigation (Table 3).

Additionally, the MR-Egger test revealed no signs of potential horizontal pleiotropy, thereby reinforcing the reliability of our findings (Table 4). To further confirm the robustness of our results, a sensitivity analysis was carried out, excluding individual genetic variants one by one (Fig. 6).

#### Discussion

Osteoporosis, a skeletal disorder characterized by decreased bone strength and increased risk of fracture, has become a major public health problem worldwide<sup>29,30</sup>. As the global population ages, osteoporosis



**Fig. 5.** Forest plot of the MR study investigating the effect of postural instability on FN-BMD, LS-BMD, FA-BMD, and their components. OR, odds ratio; CI, confidence interval; FDR, false discovery rate.

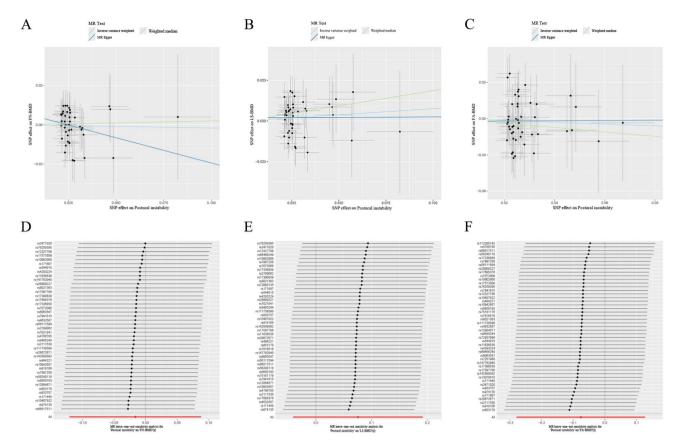
| Exposure              | Outcome | SNPs selection    | method   | d Cochran's Q test |      | st     |
|-----------------------|---------|-------------------|----------|--------------------|------|--------|
|                       |         |                   |          | Q                  | Q_df | Q_pval |
|                       | FN-BMD  | Complete          | IVW      | 45.283             | 46   | 0.465  |
|                       |         |                   | MR Egger | 45.162             | 45   | 0.502  |
|                       |         | Steiger filtering | IVW      | 30.673             | 41   | 0.880  |
|                       |         |                   | MR Egger | 20.894             | 40   | 0.878  |
|                       |         | Complete          | IVW      | 54.597             | 46   | 0.180  |
| Postural instability  | LS-BMD  |                   | MR Egger | 53.897             | 45   | 0.170  |
| 1 Osturai instability | L3-BWID | Steiger filtering | IVW      | 34.947             | 44   | 0.833  |
|                       |         |                   | MR Egger | 34.872             | 43   | 0.806  |
|                       | FA-BMD  | Complete          | IVW      | 53.178             | 49   | 0.316  |
|                       |         |                   | MR Egger | 51.296             | 48   | 0.345  |
|                       | FA-DMD  | Steiger filtering | IVW      | 43.623             | 46   | 0.572  |
|                       |         |                   | MR Egger | 43.589             | 45   | 0.531  |

**Table 3**. The Cochran's Q test of Mendelian randomization analyses for BMD and postural instability.

| Exposure             | Outcome | SNPs selection    | Egger intercept | Stand error | P. value |
|----------------------|---------|-------------------|-----------------|-------------|----------|
|                      | FN-BMD  | Complete          | -0.001          | 0.005       | 0.729    |
|                      |         | Steiger filtering | 0.006           | 0.007       | 0.382    |
| Postural instability | LS-BMD  | Complete          | 0.005           | 0.007       | 0.448    |
| Postural instability | L3-BMD  | Steiger filtering | 0.001           | 0.006       | 0.785    |
|                      | FA-BMD  | Complete          | -0.015          | 0.011       | 0.190    |
|                      |         | Steiger filtering | -0.002          | 0.012       | 0.854    |

Table 4. The MR Egger test of Mendelian randomization analyses for BMD and postural instability.

raises increasing concerns<sup>31,32</sup>. Previous observational studies have mentioned a strong association between osteoporosis and Alzheimer's disease, of which postural instability is a common symptom<sup>33</sup>. However, a recent Mendelian randomization study systematically analyzed and denied a causal relationship between osteoporosis and Alzheimer's disease, which contradicts past observational studies<sup>34</sup>. This study, however, revealed a striking association between osteoporosis, particularly femoral neck osteoporosis, and postural instability. Meanwhile,



**Fig. 6.** Mendelian randomization analysis for investigating the causal association between postural instability and FN-BMD, LS-BMD, FA-BMD. (**A**)Scatter plot of postural instability on FN-BMD. (**B**) Scatter plot of postural instability on LS-BMD. (**C**) Scatter plot of postural instability on FA-BMD. (**D**) Leave-one-out sensitivity analysis of postural instability on FN-BMD. (**E**) Leave-one-out sensitivity analysis of postural instability on LS-BMD. (**F**) Leave-one-out sensitivity analysis of FA-BMD on postural instability.

a retrospective study suggests that BMD may be a risk factor for impaired postural stability. Alexander Simon's team screened 1,086 patients and found a correlation between lower BMD T-scores in the femur and a longer path to Romberg's test<sup>35</sup>. A cross-sectional study by Akira Okayama also emphasized the association between osteoporosis and postural instability in postmenopausal women with instability<sup>36</sup>. Previously, the causal relationship between the two could not be well explained due to the interference of numerous confounding factors<sup>37</sup>. In the present study, genetic variants were used as instrumental variables to minimize the interference of acquired confounders and further strengthen the evidence for a causal relationship between osteoporosis and increasing postural instability. In previous studies, incipient deformity in osteoporosis usually manifests as thoracic or thoracolumbar kyphosis. However, in older patients with osteoporosis, mobility is more severely limited especially in the lumbar region, particularly in the range of lumbar extension<sup>38</sup>. The range of extension of the lower lumbar spine, in particular, may inhibit the lumbar spine's ability to compensate for thoracic kyphosis, leading to increased postural instability associated with lumbar kyphosis<sup>39</sup>. This is consistent with our findings.

In this study, we used a Mendelian randomization method using BMD of the femoral neck, lumbar spine, and forearm to assess the degree of osteoporosis and systematically evaluated the causal relationship between osteoporosis and postural instability. Gene prediction results show that osteoporosis of the femoral neck and lumbar spine may be associated with an increased risk of postural instability. Through sensitivity analysis, we further confirmed the positive association, which was not present for the forearm.

It is worth noting that genetic variation is stable over the individual life cycle and alleles are randomly distributed and fixed. Thus, the results of the present study represent a lifelong risk of postural instability in patients with osteoporosis, and bias caused by confounding factors and reverse causality can be avoided.

The prevalence of osteoporosis among the elderly is approximately 15.7%, and this number continues to increase with the aging of the population globally<sup>40</sup>. Previous studies have examined the factors that contribute to the increased risk of falls, which include aging, history of falls, muscle weakness, home environmental hazards, drug use (especially psychotropic drugs), and cardiac pacing accompanied by carotid sinus hypersensitivity, which have all been reported as risk factors for falls<sup>41</sup>. In addition, postural instability has also been identified as a risk factor for falls<sup>42</sup>. With demographic changes, the number of osteoporosis patients is expected to increase significantly in the coming decades, especially with the aging of the population. In the face of a large population of osteoporosis patients, we should pay more attention to their risk of postural instability to prevent the health hazards and economic burden associated with fractures due to falls<sup>43</sup>.

It is worth noting that this study has some limitations. First, the original GWAS data analyzed in this study were from a European population, so its findings may not apply to other ethnic groups. Second, the limitations of the pooled GWAS data did not allow for stratified analyses based on general factors such as age and gender. Third, it was difficult to completely exclude the influence of horizontal pleiotropy effects. Therefore, the study conducted a series of sensitivity analyses to ensure the reliability of the results.

#### Conclusion

In summary, this study found that osteoporosis of the femoral neck and lumbar spine was positively associated with the risk of postural instability through the analysis of the Mendelian randomization method, while there was no significant causal relationship between postural instability and osteoporosis at different sites. This finding may open up a whole new avenue for the prevention of postural instability and the treatment of osteoporosis. Therefore, the management of patients with osteoporosis should be strengthened to reduce the risk of postural instability and the incidence of fall injuries caused by it. In addition, due to genetic differences between different ethnic groups, countries, and regions, further studies are needed in different populations.

#### Data availability

All data generated or analysed during this study are included in this manuscript.

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#### **Author contributions**

Qingqiang Yao, Rui Zhao and Haiyang Yu designed the project. Rui Zhao and Kun Zhu conducted the literature review. Kun Zhu, Zijie An and YaWei Li is responsible for the computer code design and code test running. ZiJie An, Feng Zhang and Qiaoyu Zhang was responsible for writing of the manuscript and performed the statistical analysis. Kun Zhu and ZiJie An edited and polished the article. All authors read and approved the final manuscript.

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#### **Declarations**

#### **Competing interests**

The authors declare no competing interests.

#### Additional information

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