

Supplementary Table 3. Multivariable Mendelian randomization estimates for the association of genetically proxied appendicular lean mass (adjusted for fat mass) with AD after adjustment for genetic predictors of height. Results from fixed-effects meta-analysis are displayed in the last row. CI: confidence interval; IVW: inverse-variance weighted.

Cohort	Beta [95% CI]	P value
UKB (682 total SNPs)	0.91 [0.80-1.05]	0.19
FinnGen (679 total SNPs)	0.86 [0.72-1.03]	0.11
Meta-analysis	0.90 [0.80-0.997]	0.04