

Supplementary Table 2. Mendelian randomization estimates for the association of genetically proxied liability to AD with appendicular lean mass adjusted for fat mass. 16 SNPs were used in MR analysis. CI: confidence interval; IVW: inverse-variance weighted.

Method	Beta [95% CI]	P value
IVW	0.01 [0.001-0.02]	0.03
Weighted Median	0.01 [0.004-0.02]	0.002
MR Egger	0.01 [-0.005-0.02]	0.28
Penalised weighted median	0.01 [0.005-0.02]	0.001