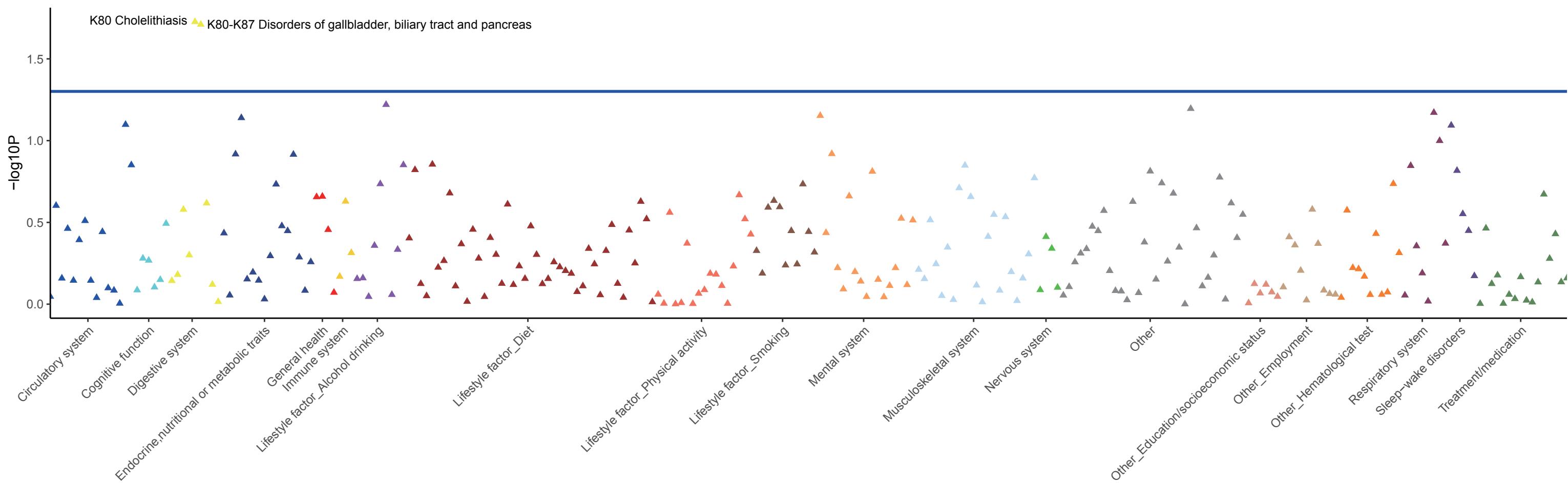


Fig. S1 The data filtering process.

A



B

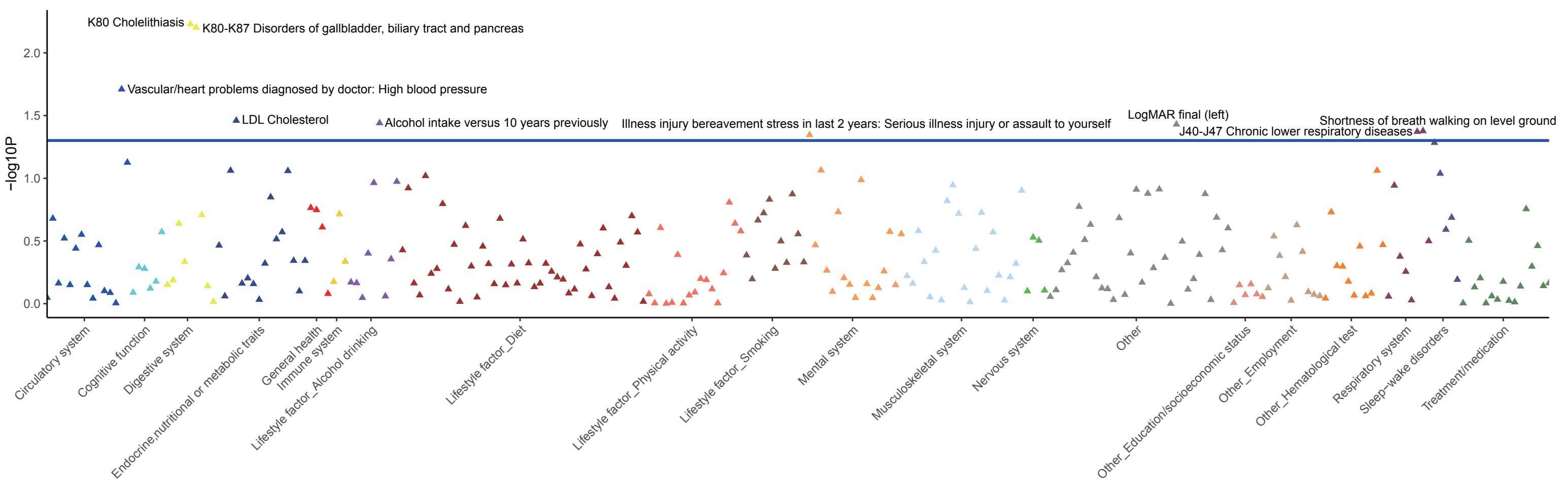


Fig. S2 Manhattan plot showing the pleiotropy assessment results for MR-Egger intercept test (A) and Q-Q' test (B). For each plot, the blue horizontal line represents the threshold for significance ($P < 0.05$).

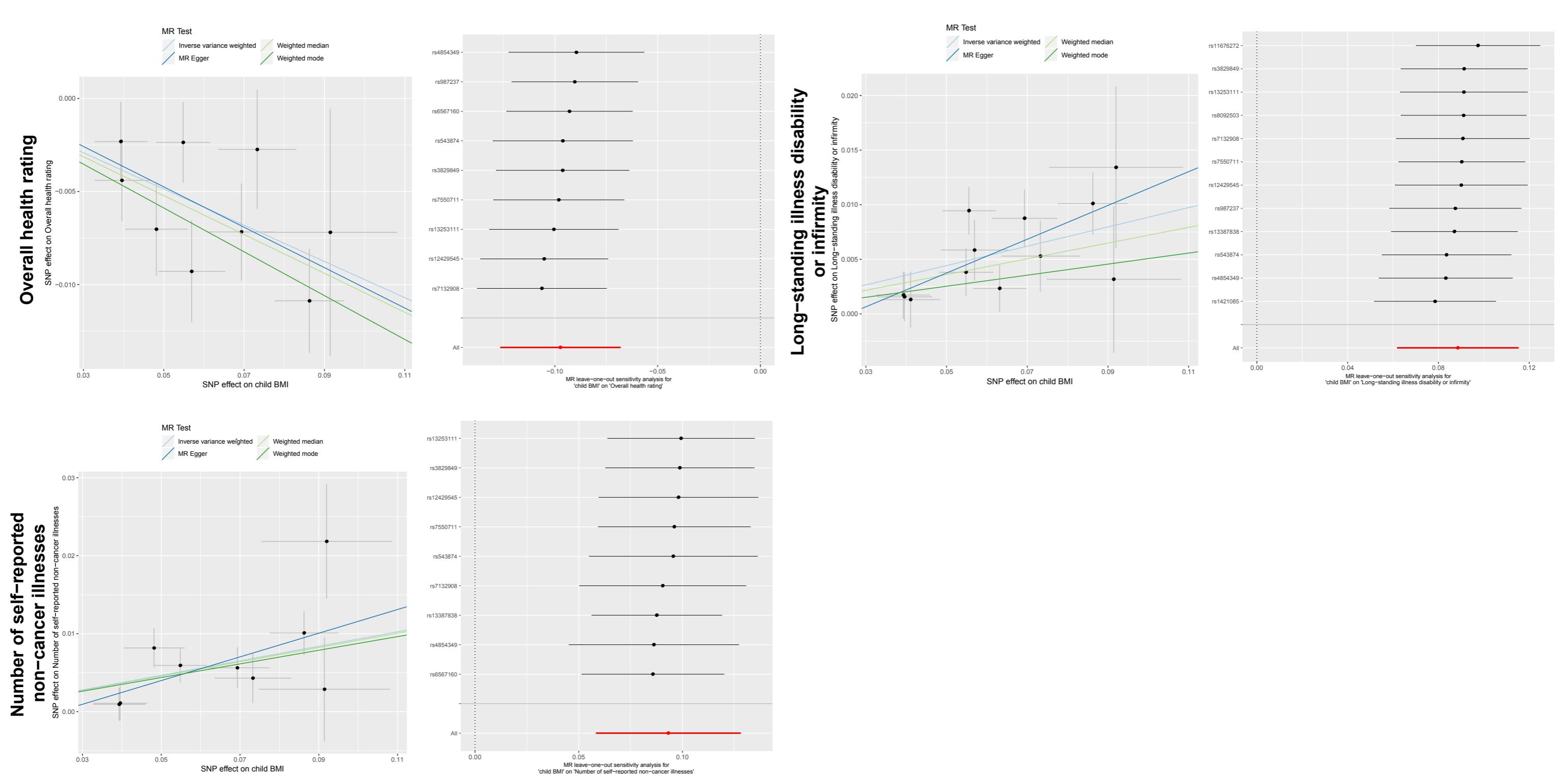
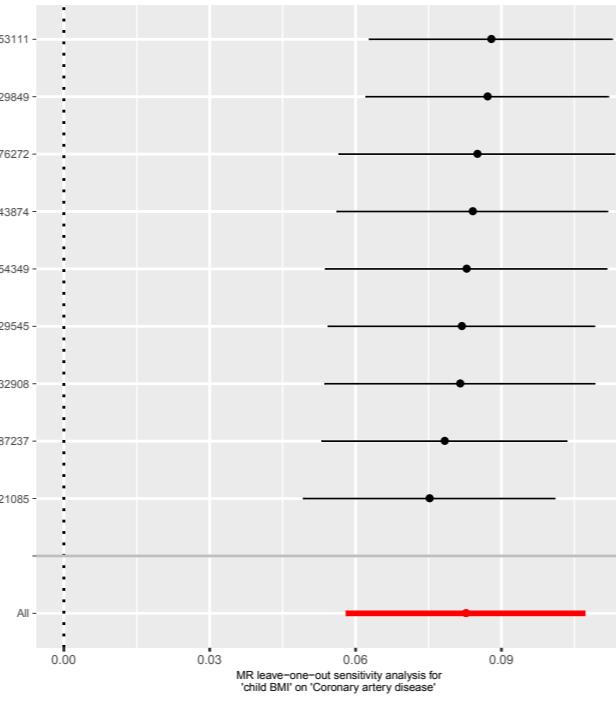
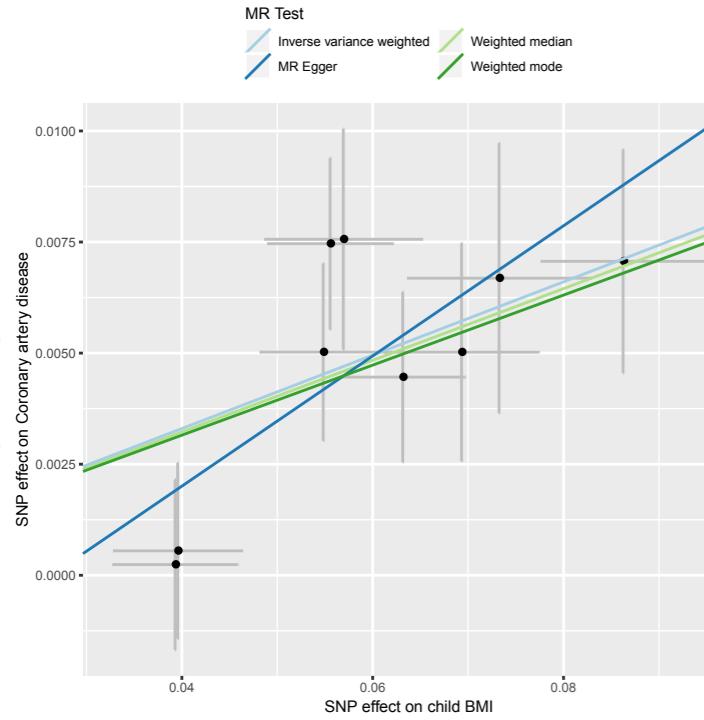
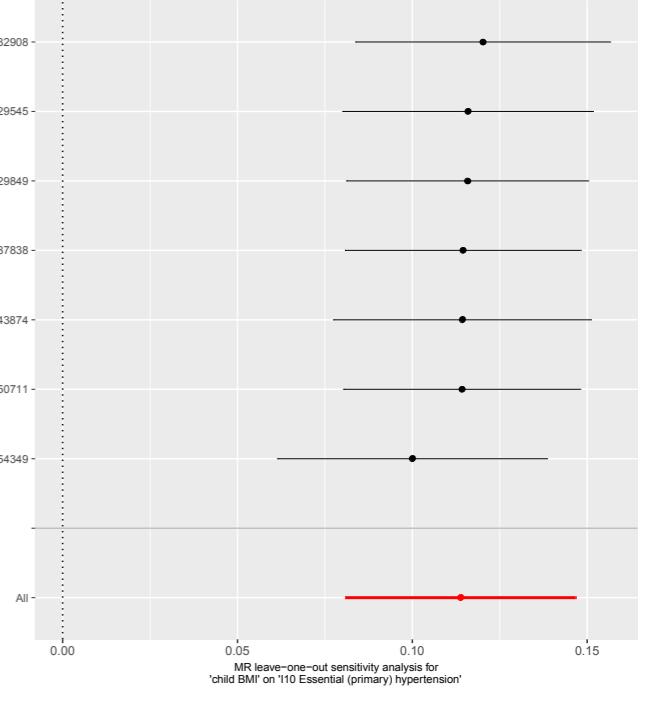
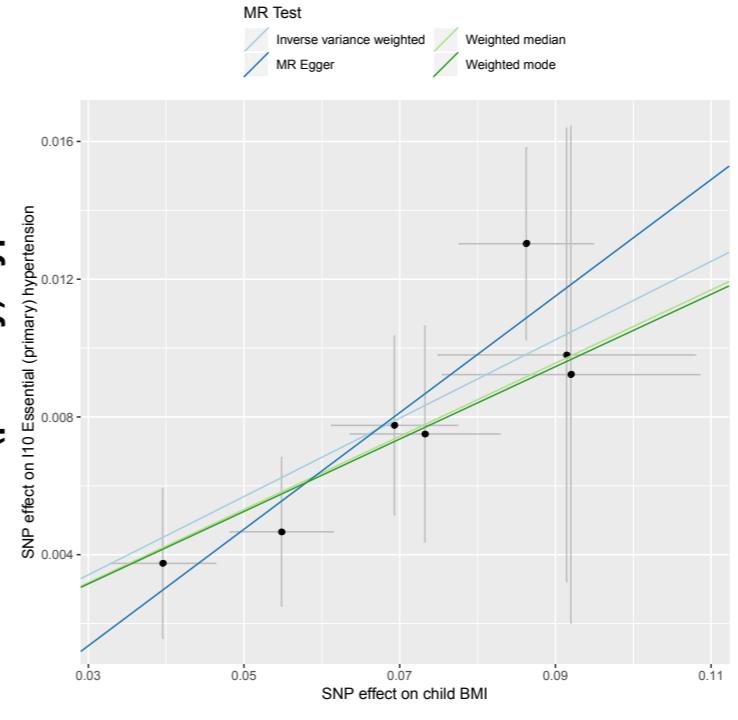


Fig. S3 Scatter plot and leave one out analysis plot for overall health rating, long-standing illness disability or infirmity, and number of self-reported non-cancer illnesses.

Coronary artery disease



I10 Essential (primary) hypertension



Vascular/heart problems diagnosed by doctor: High blood pressure

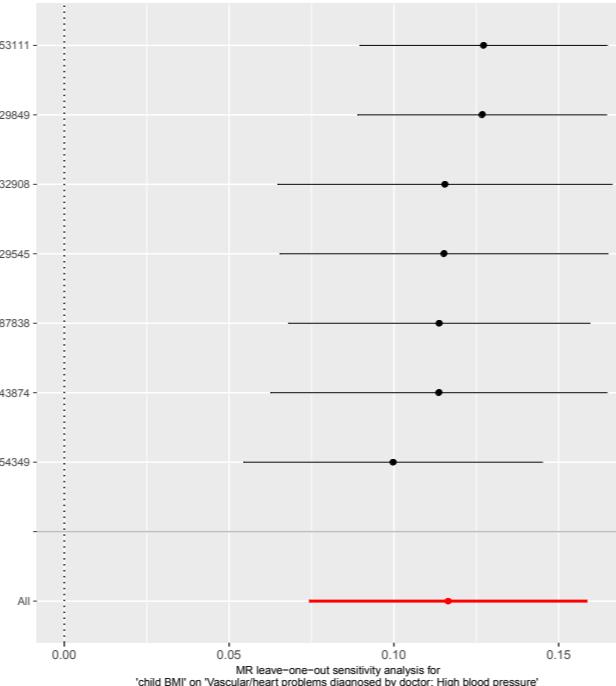
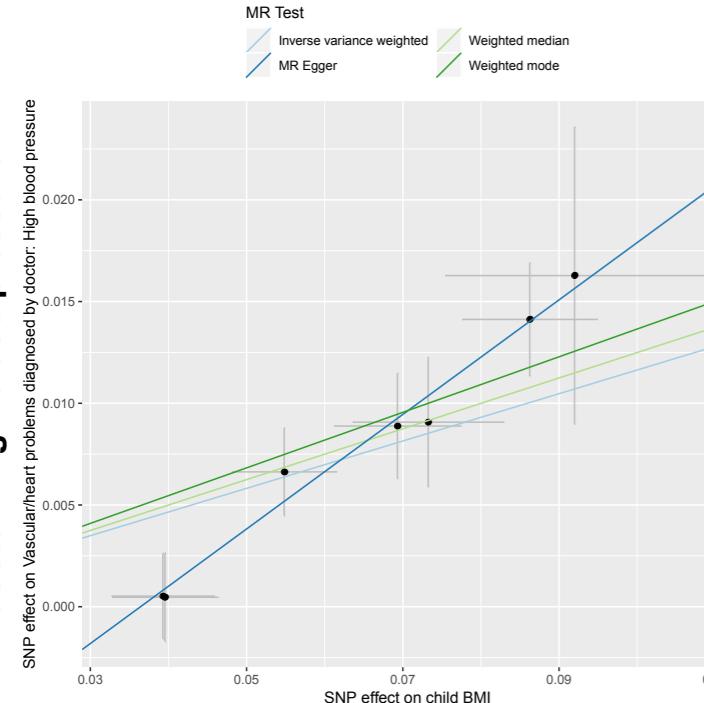
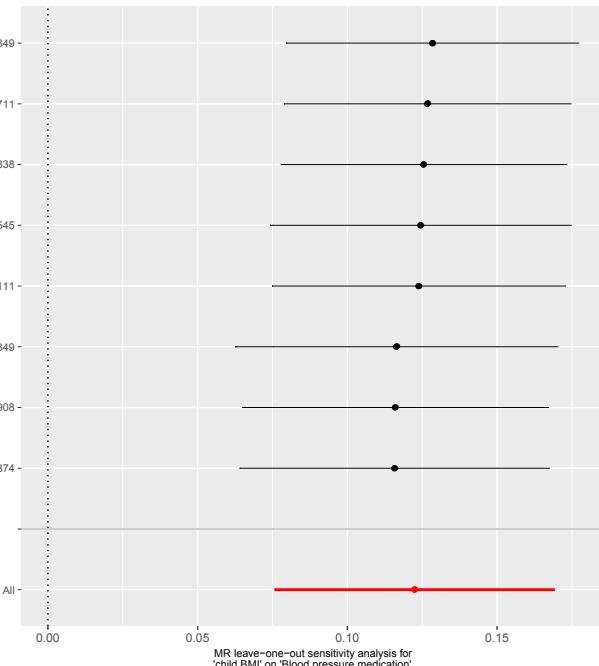
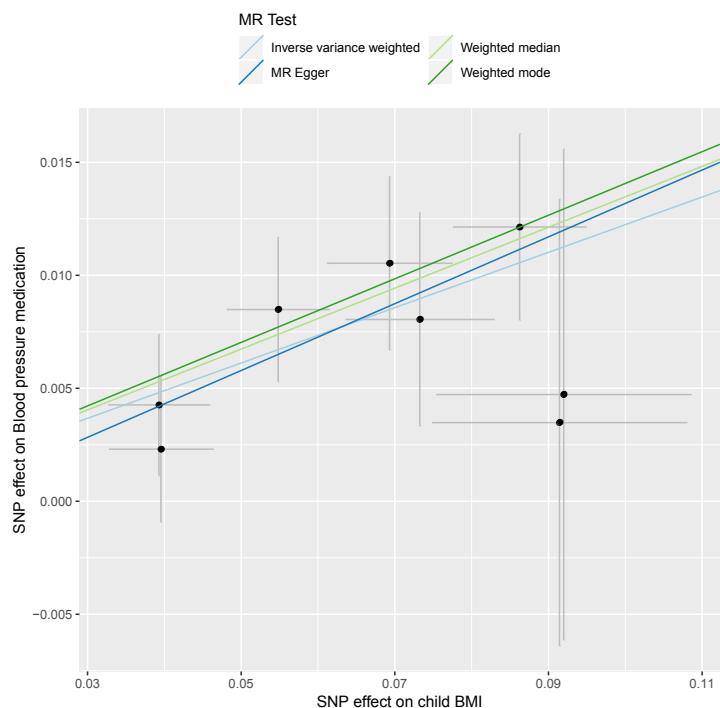


Fig. S4 Scatter plot and leave one out analysis plot for circulatory system traits.

Blood pressure medication



Metformin

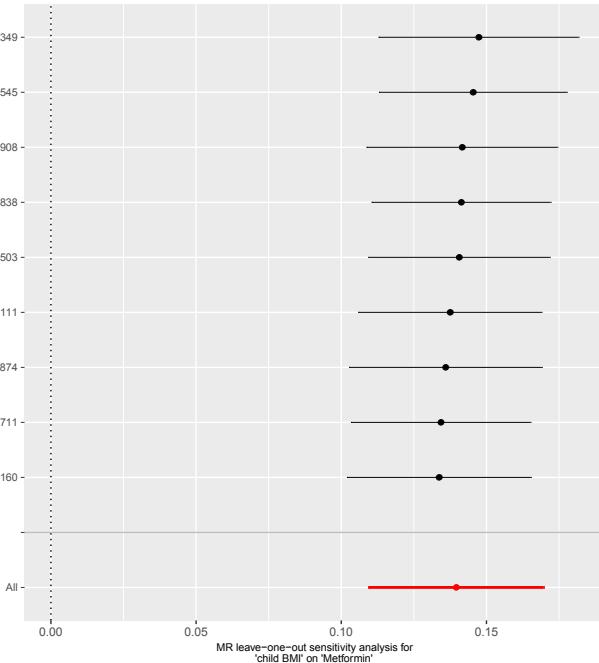
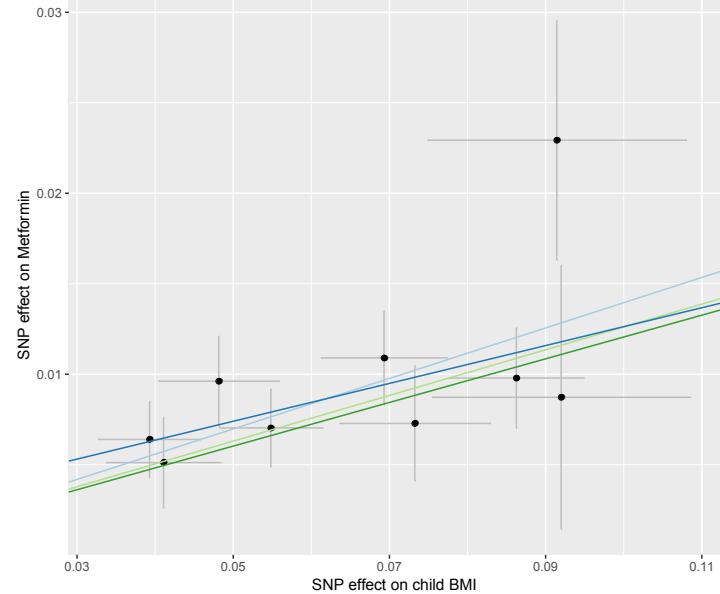


Fig. S5 Scatter plot and leave one out analysis plot for traits of received treatment/medication.

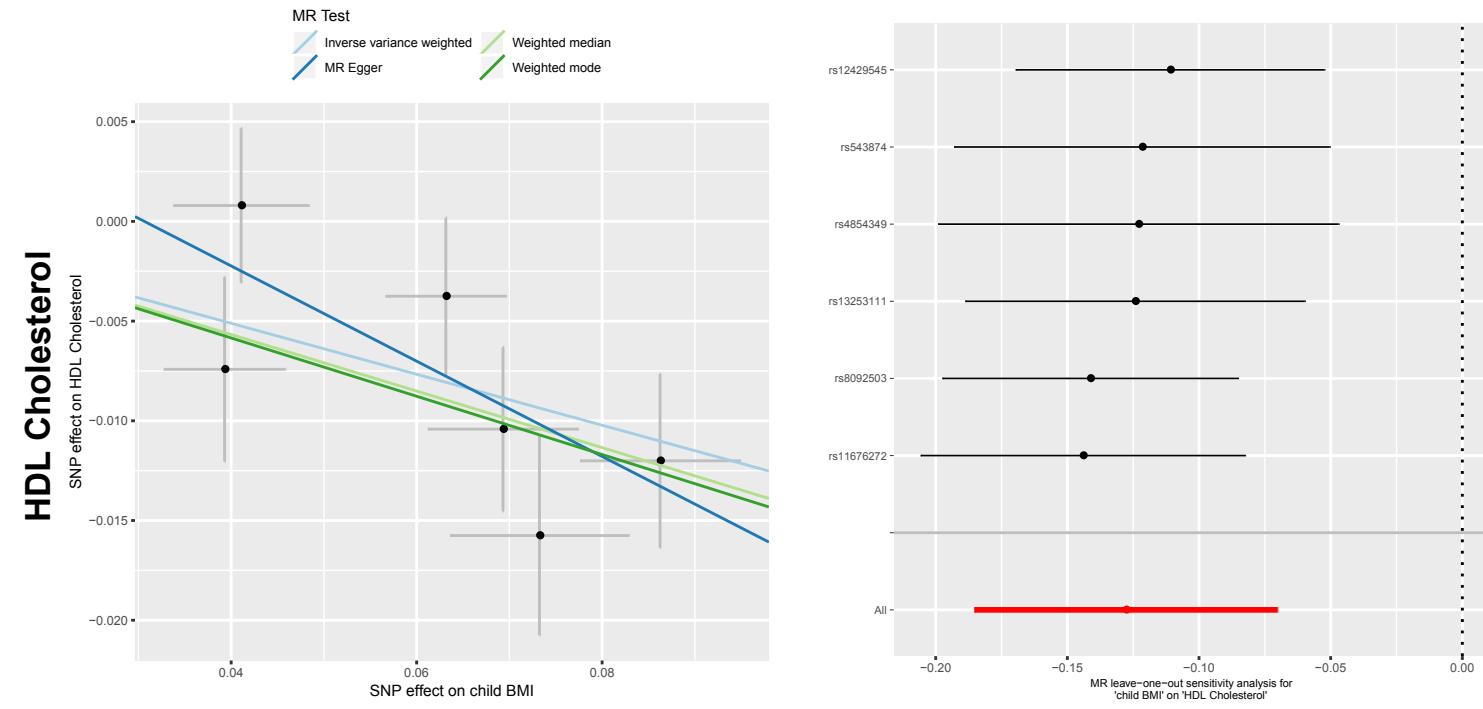
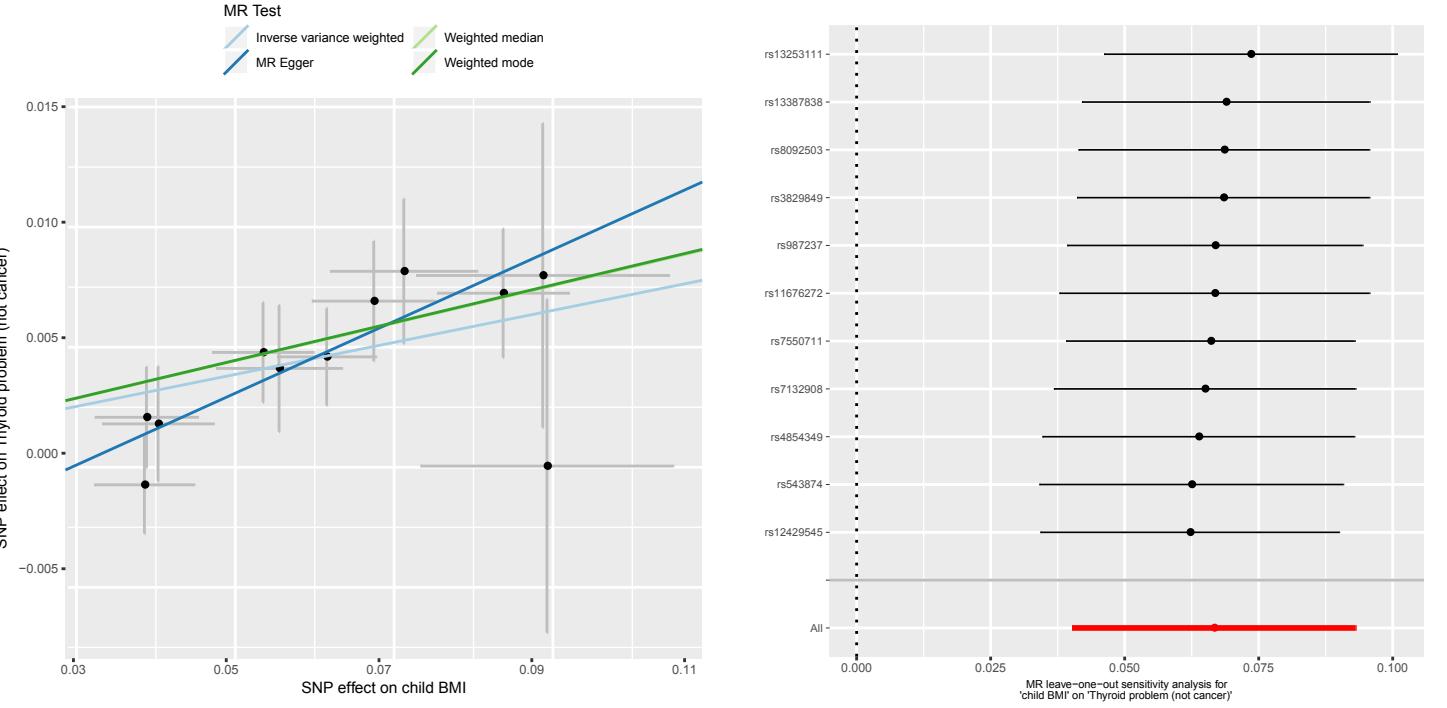
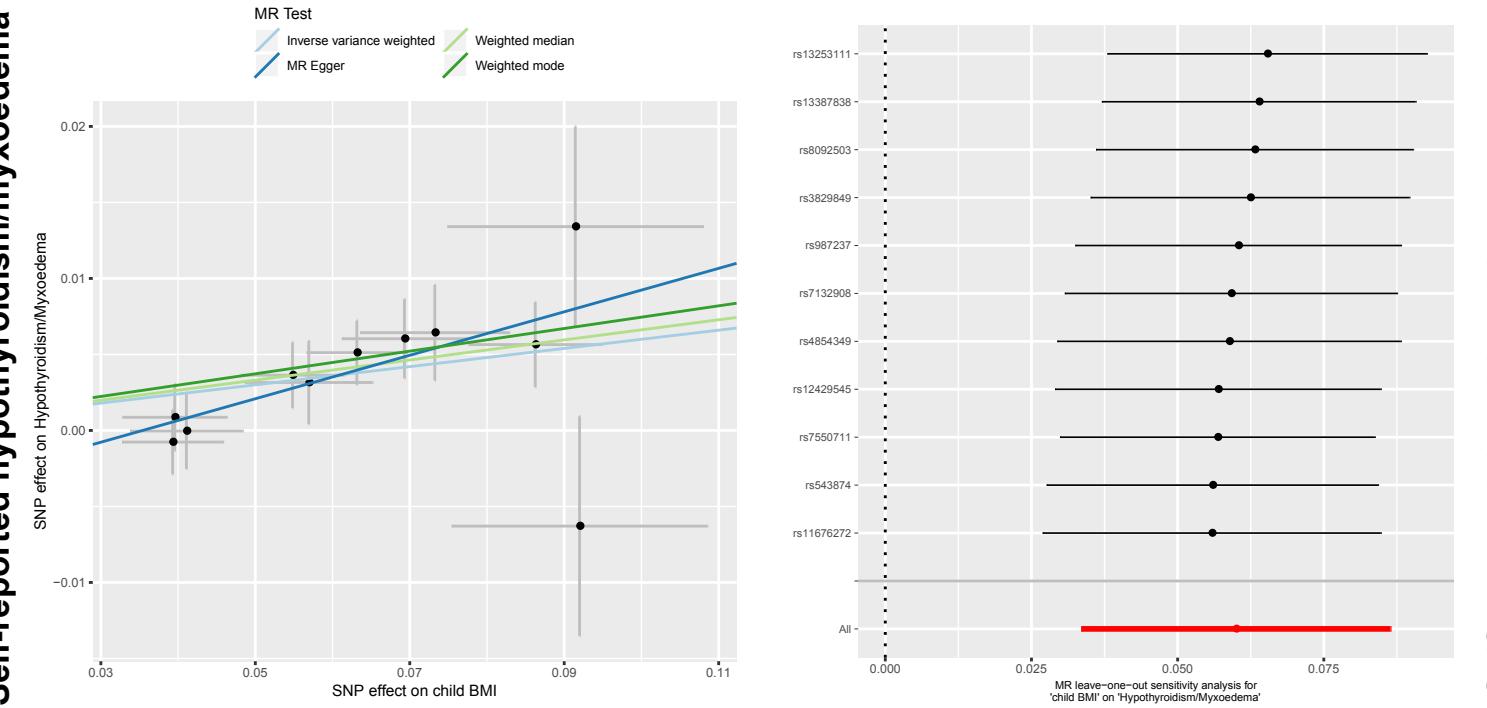
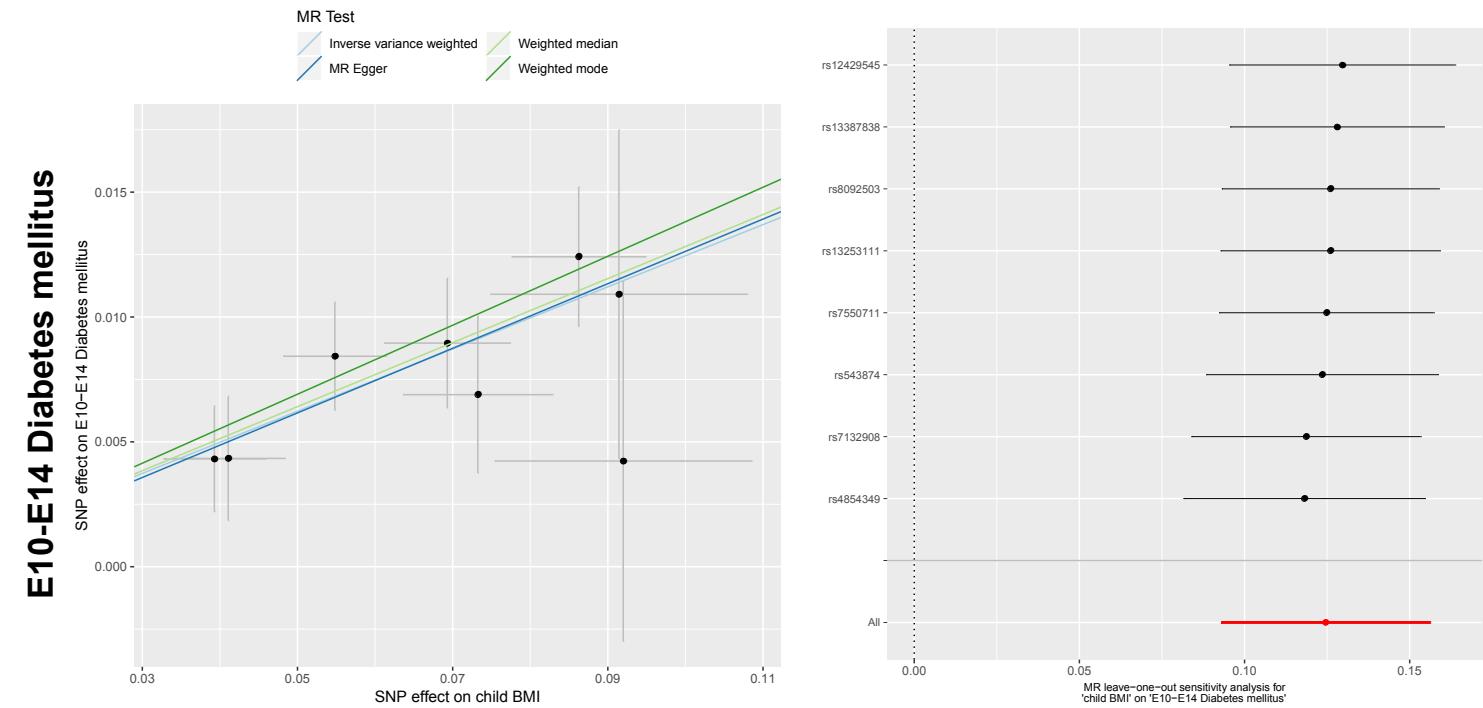
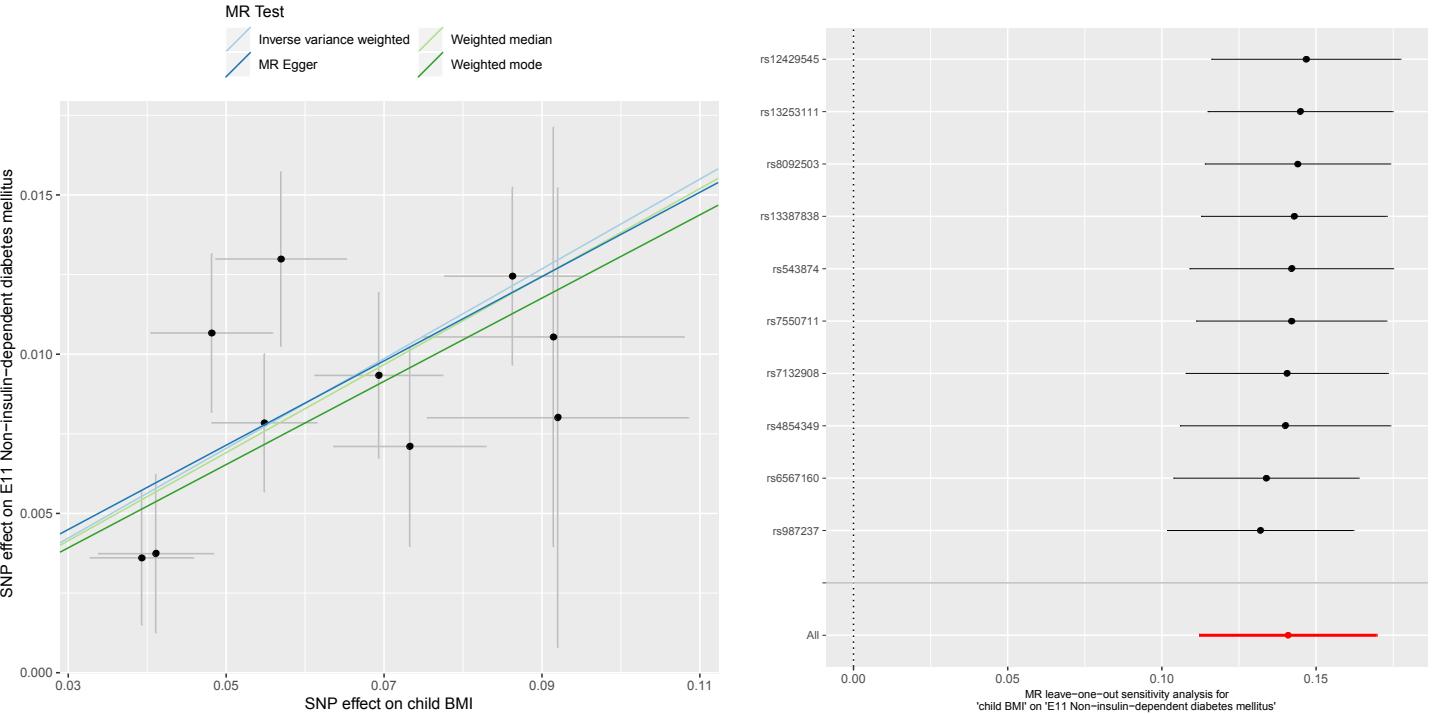
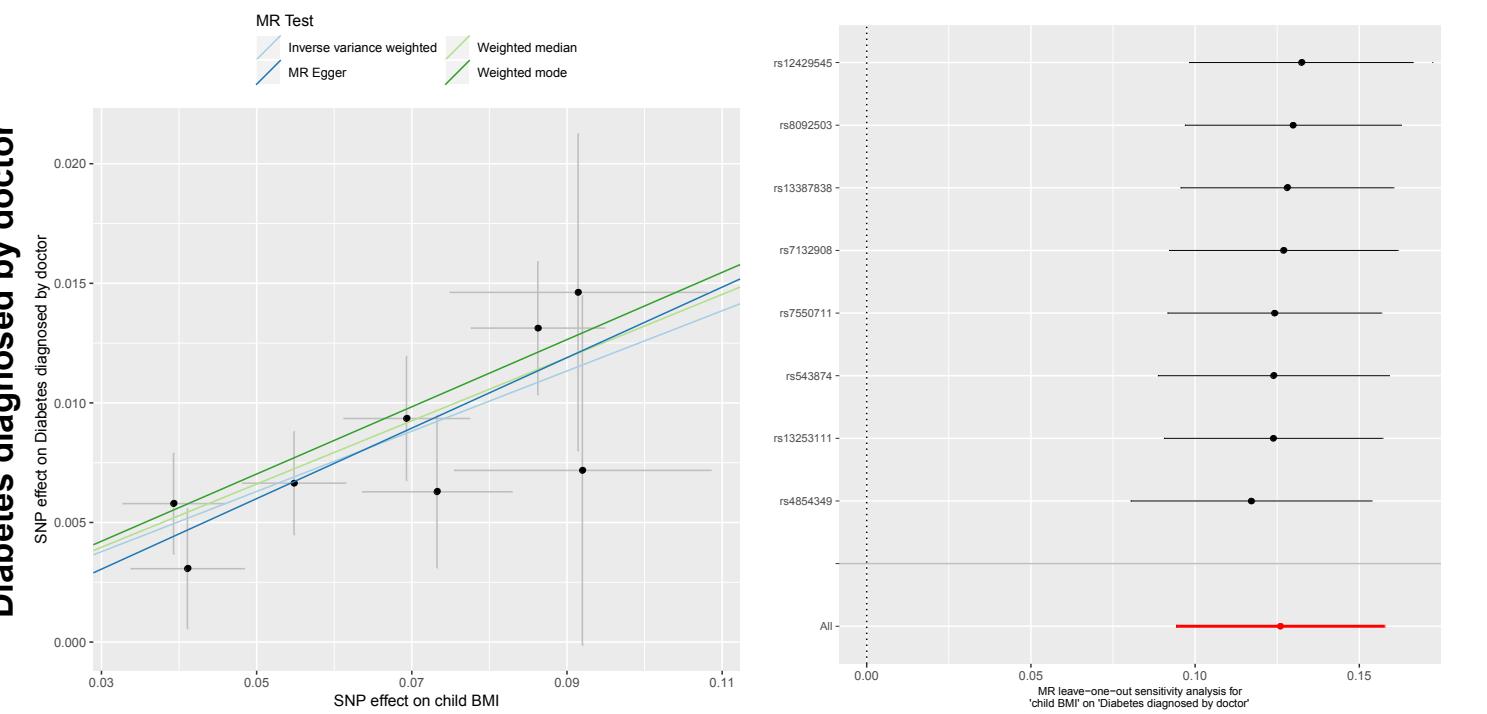
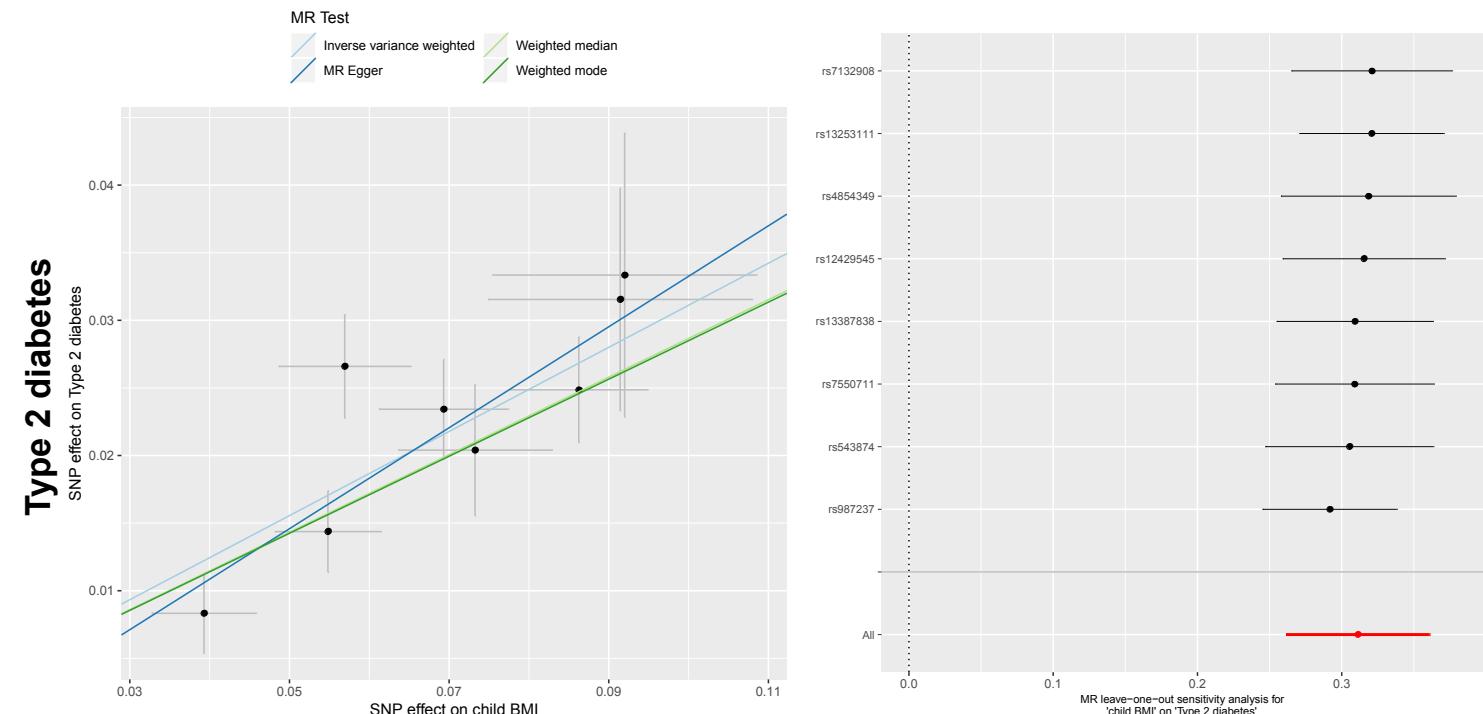


Fig. S6 Scatter plot and leave one out analysis plot for endocrine, nutritional or metabolic traits.

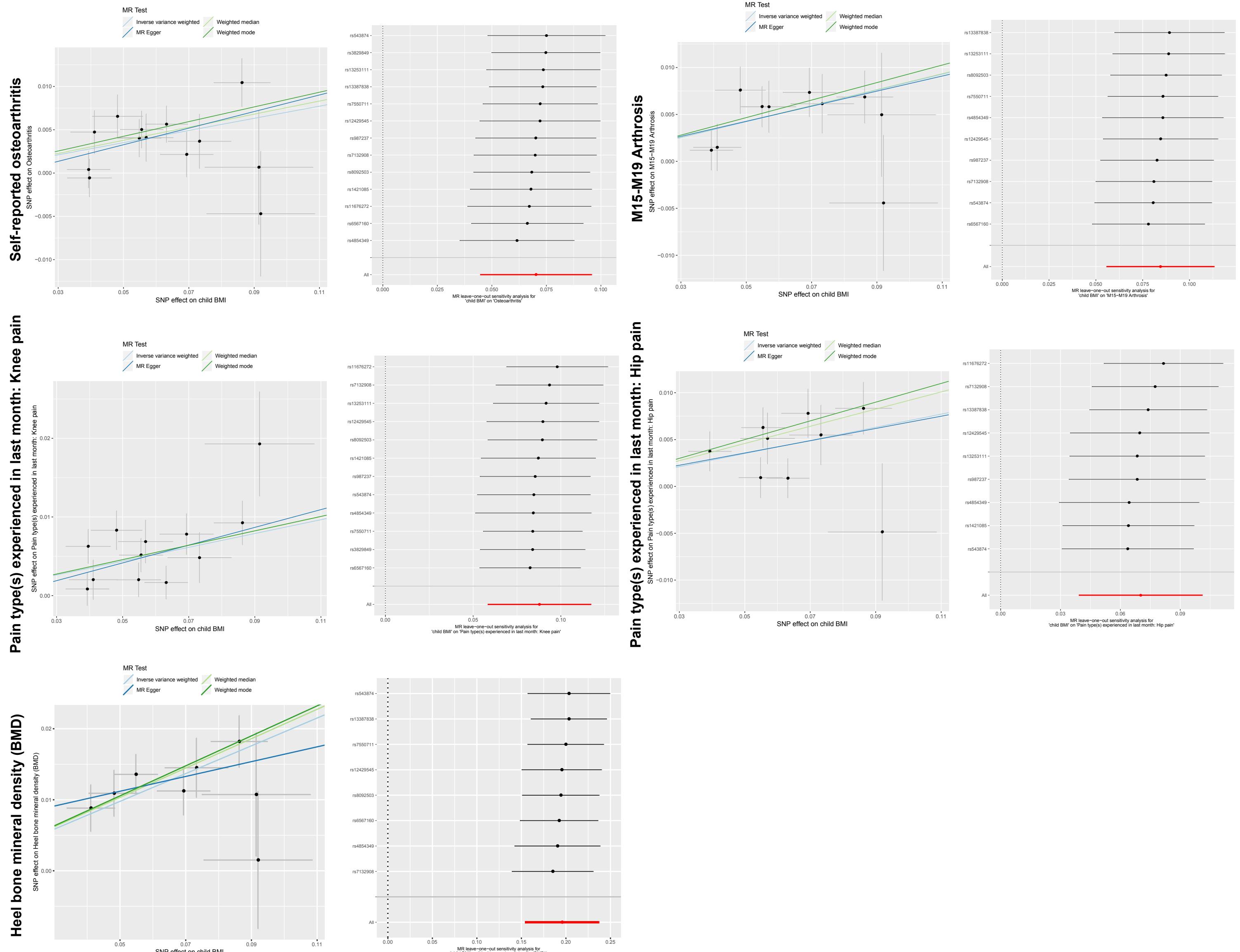
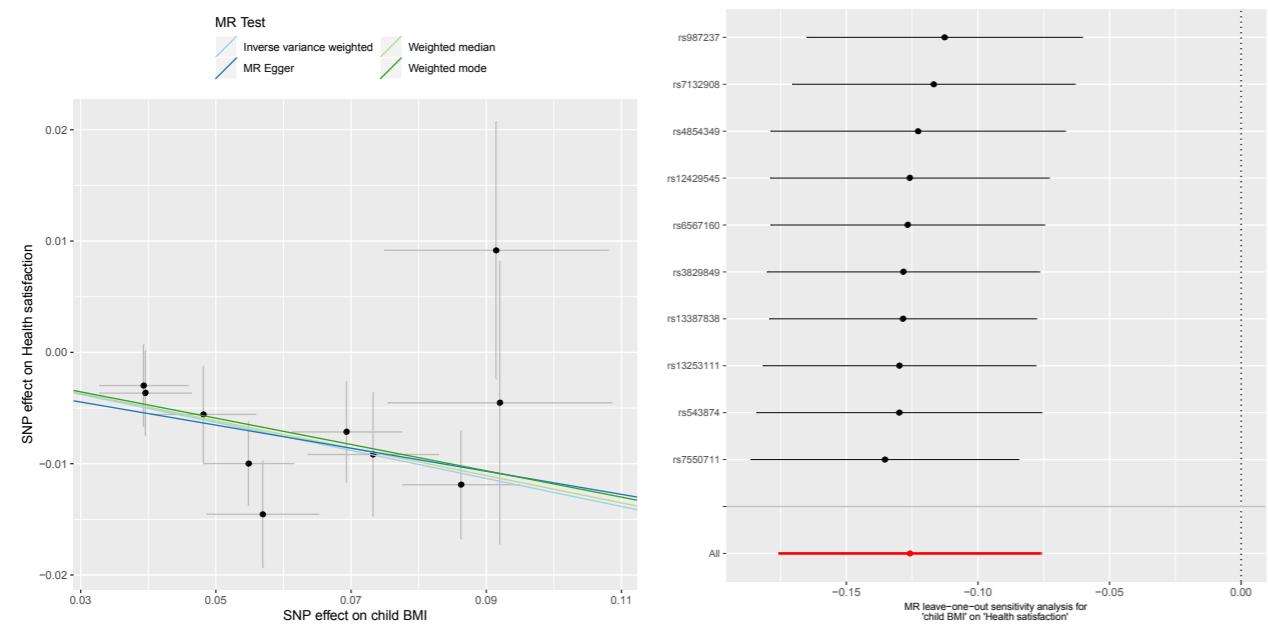
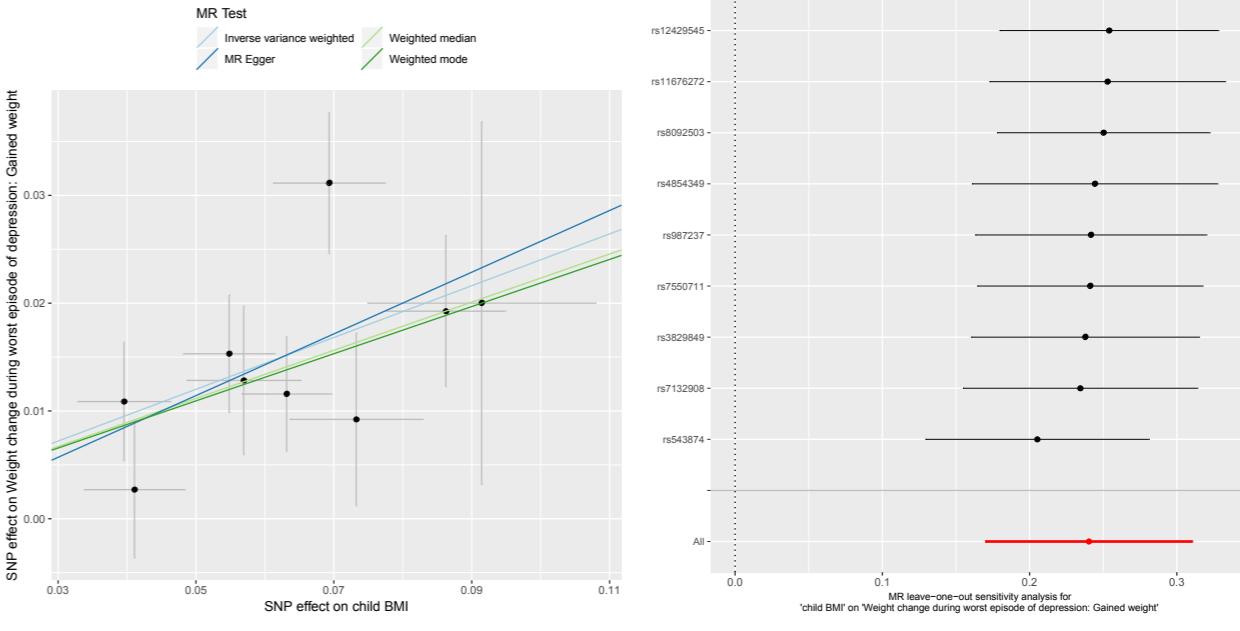


Fig. S7 Scatter plot and leave one out analysis plot for musculoskeletal system traits.

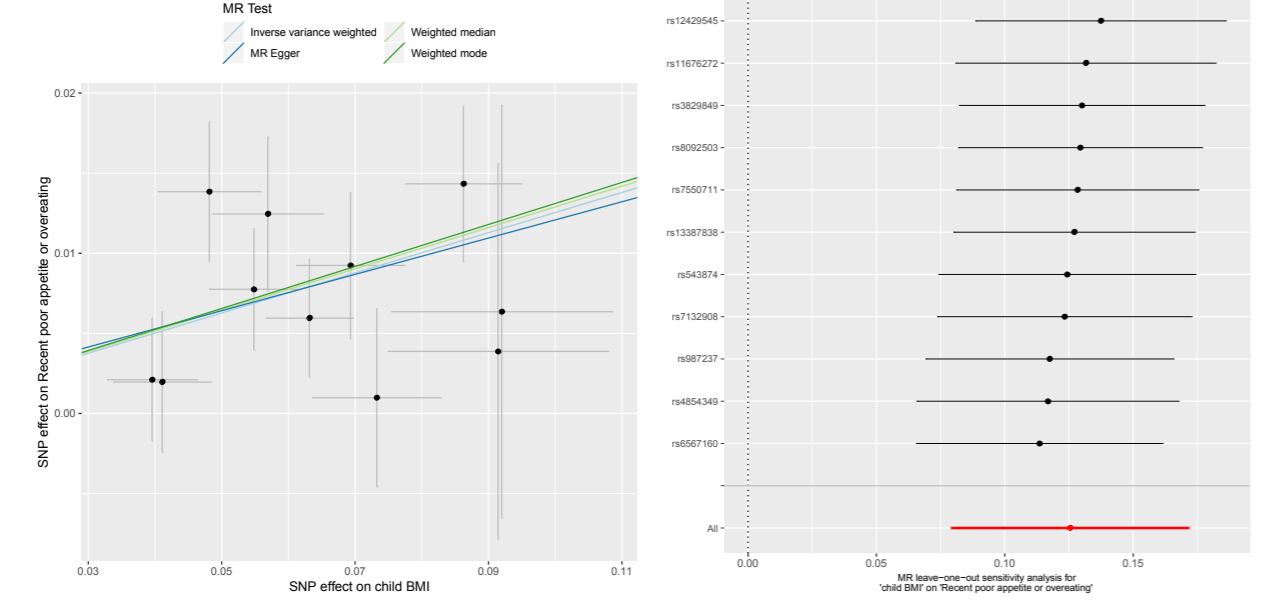
Health satisfaction



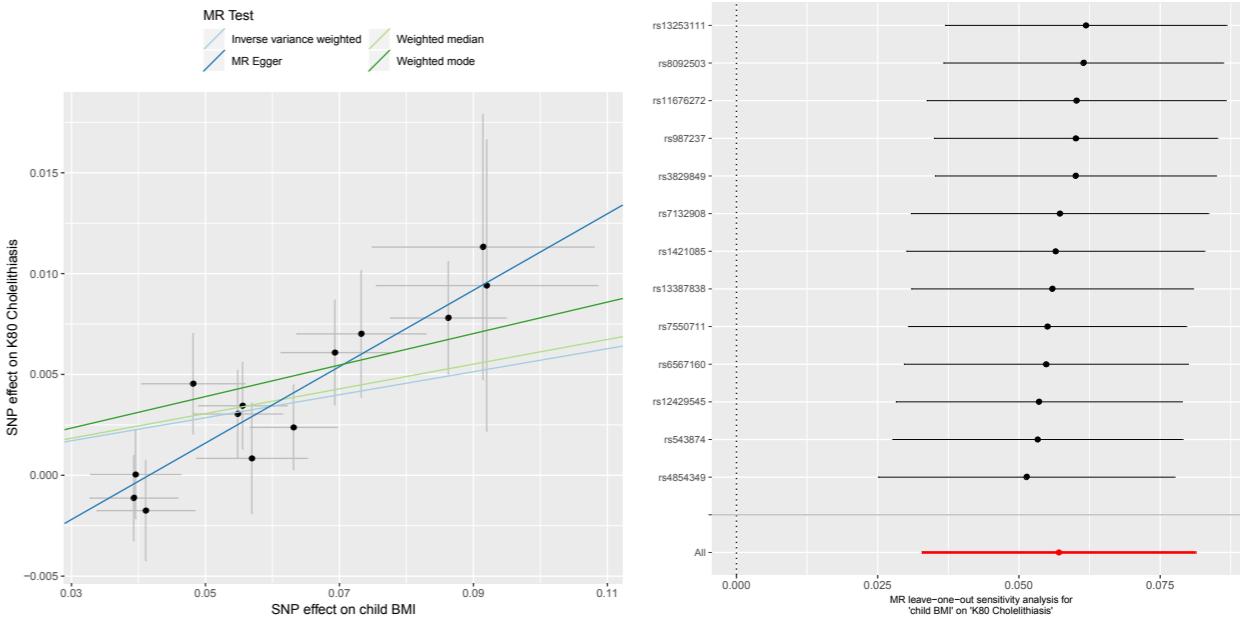
Weight change during worst episode of depression: Gained weight



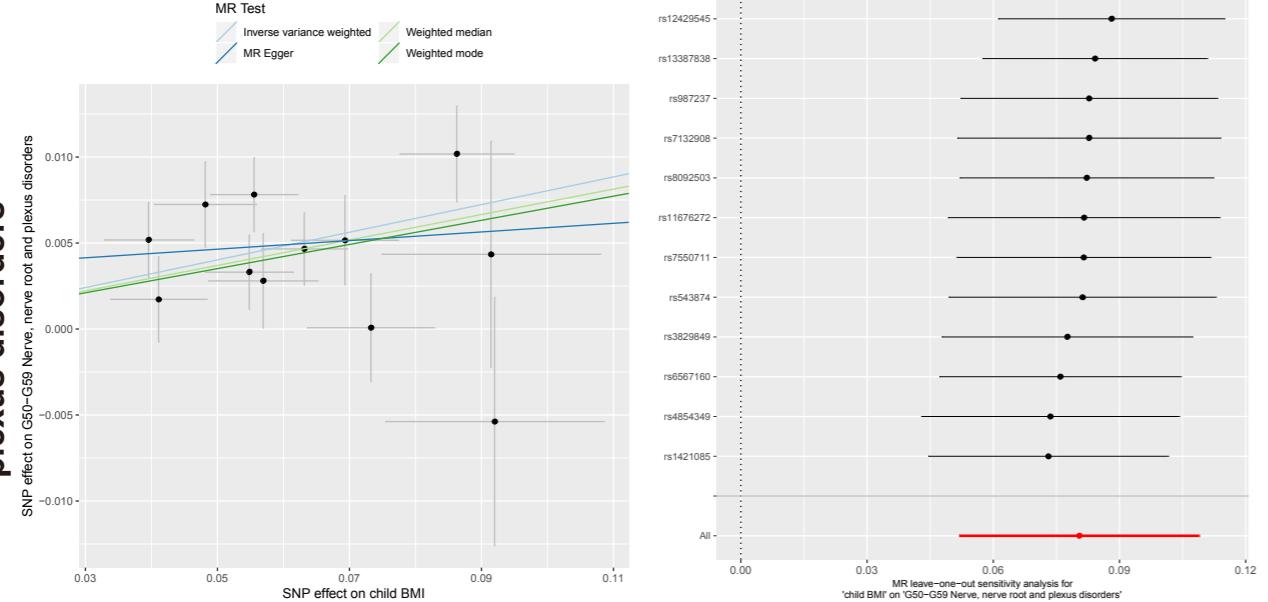
Recent poor appetite or overeating



K80 Cholelithiasis



G50-G59 Nerve, nerve root and plexus disorders



Forced vital capacity (FVC) Best measure

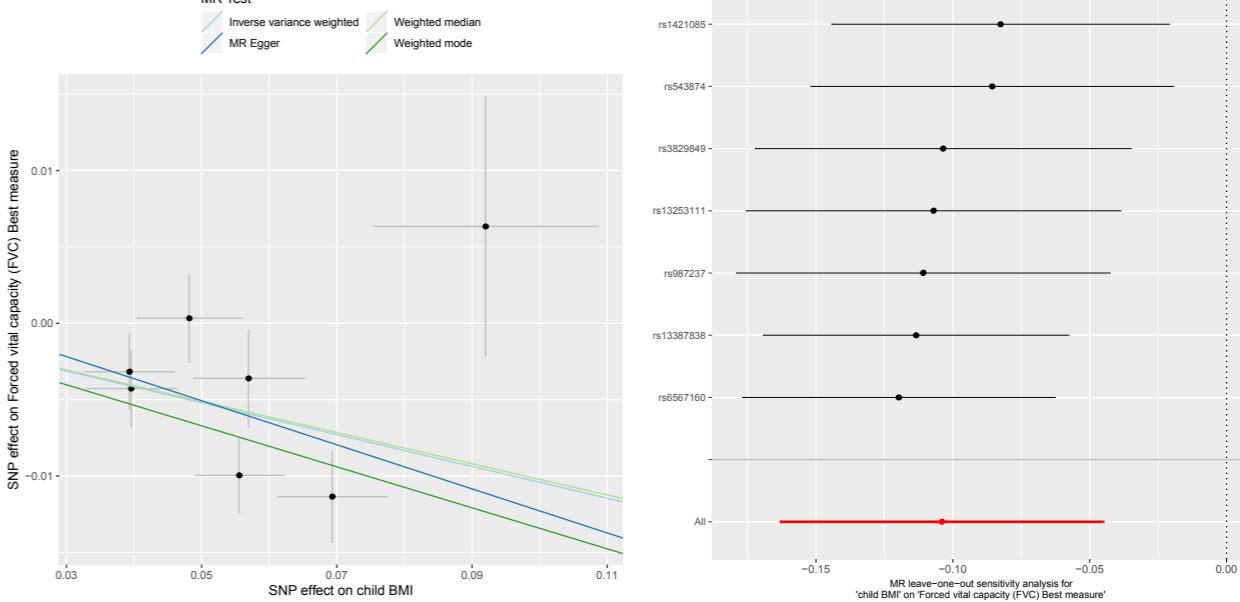
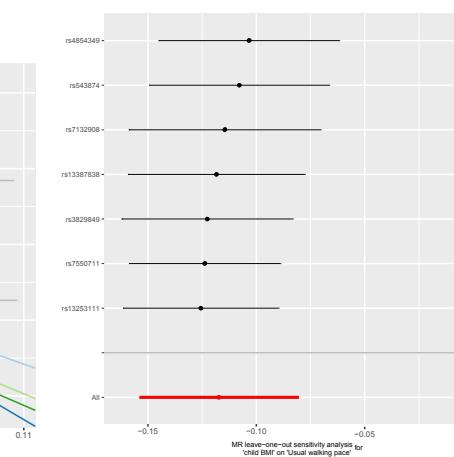
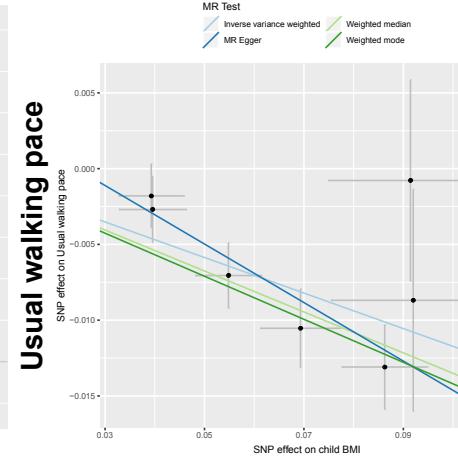
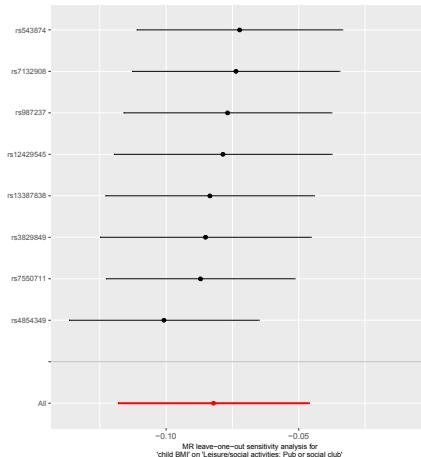
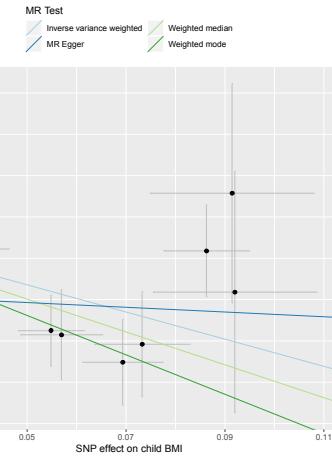
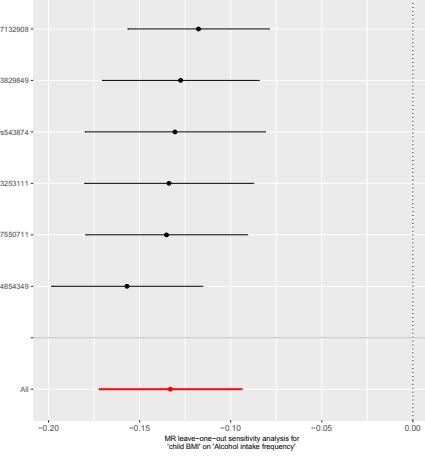
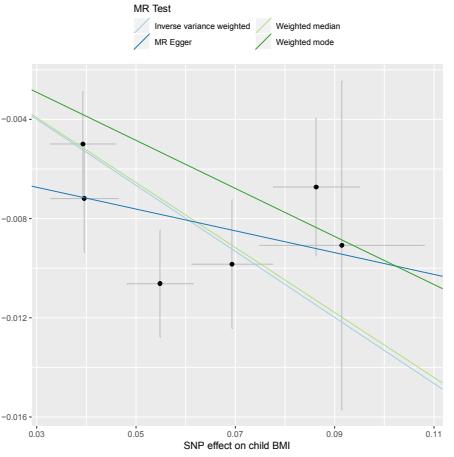


Fig. S8 Scatter plot and leave one out analysis plot for the rest disease-related traits.

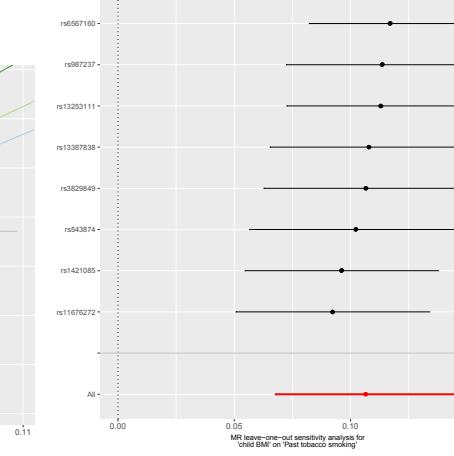
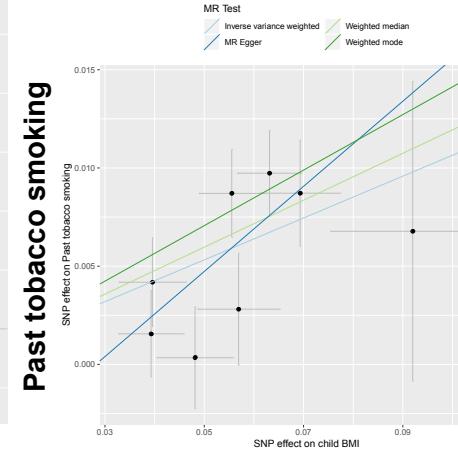
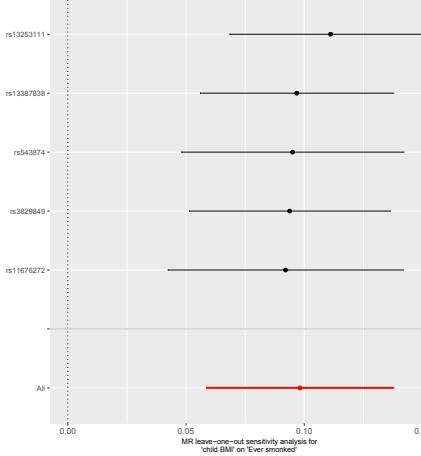
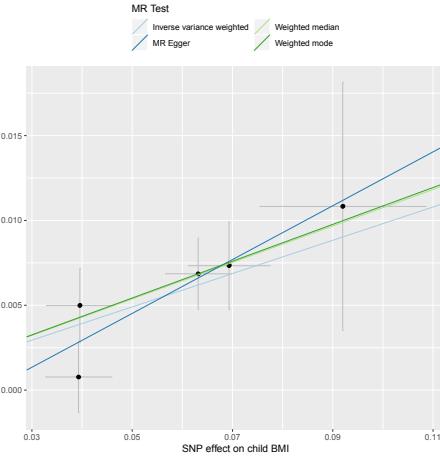
Leisure/social activities: Pub or social club



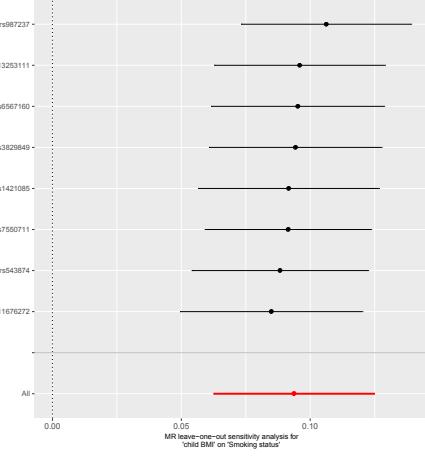
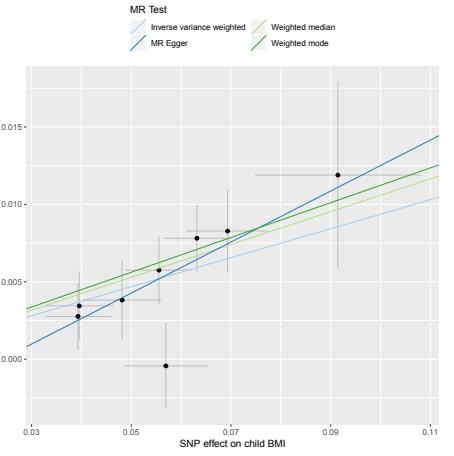
Alcohol intake frequency



Ever smoked



Smoking status



Smoking status: Never

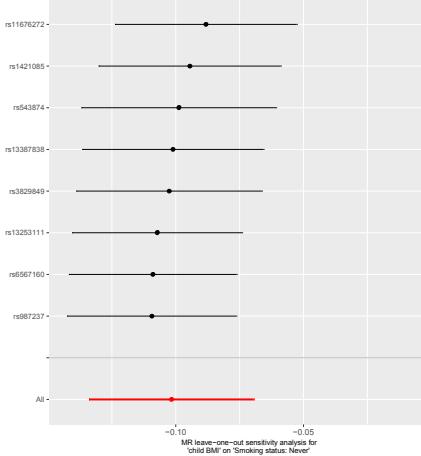
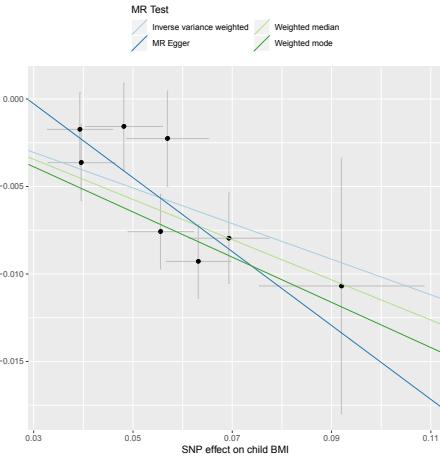


Fig. S9 Scatter plot and leave one out analysis plot for physical activity, smoking behaviors, and alcohol intake frequency.

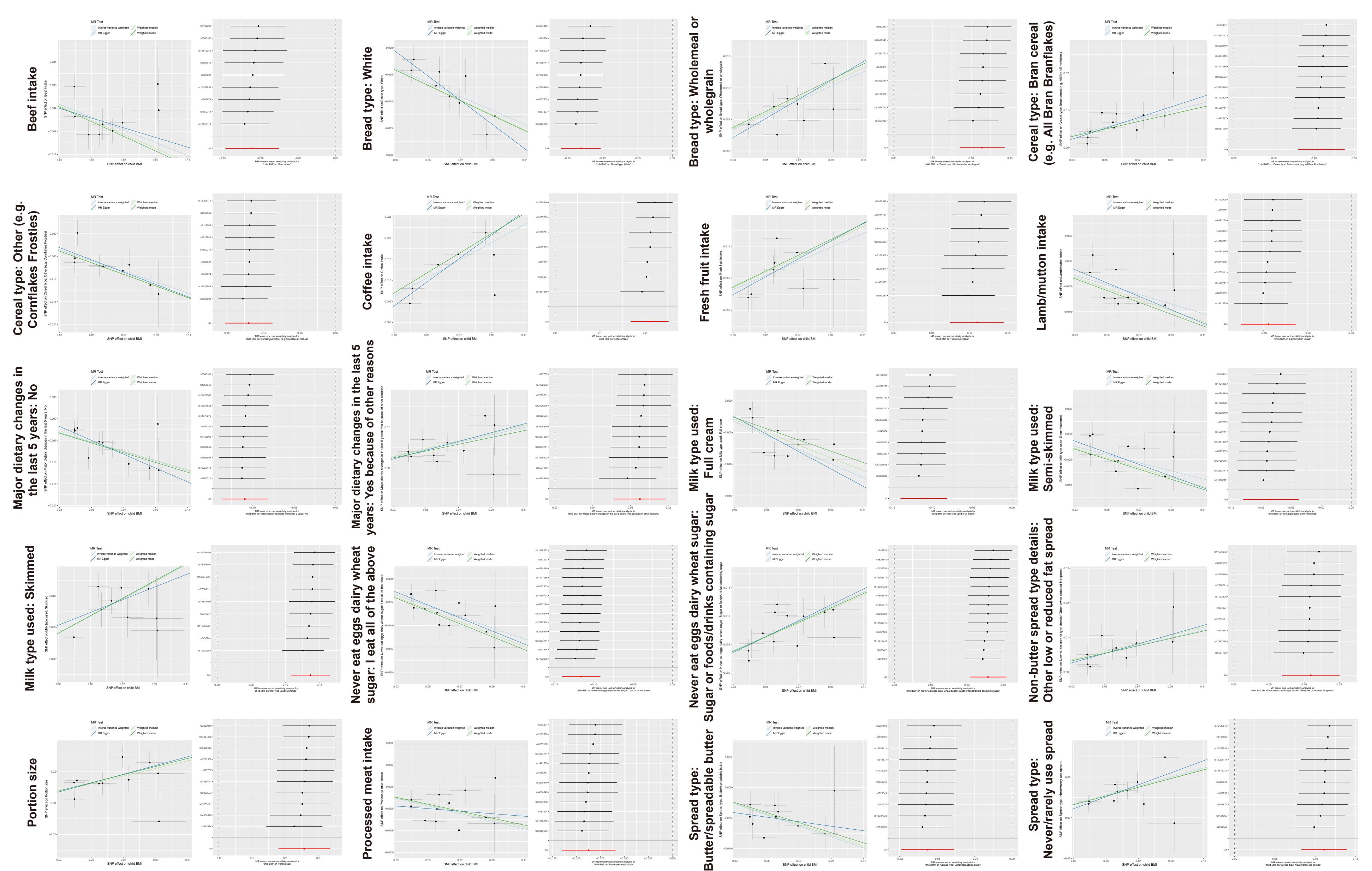
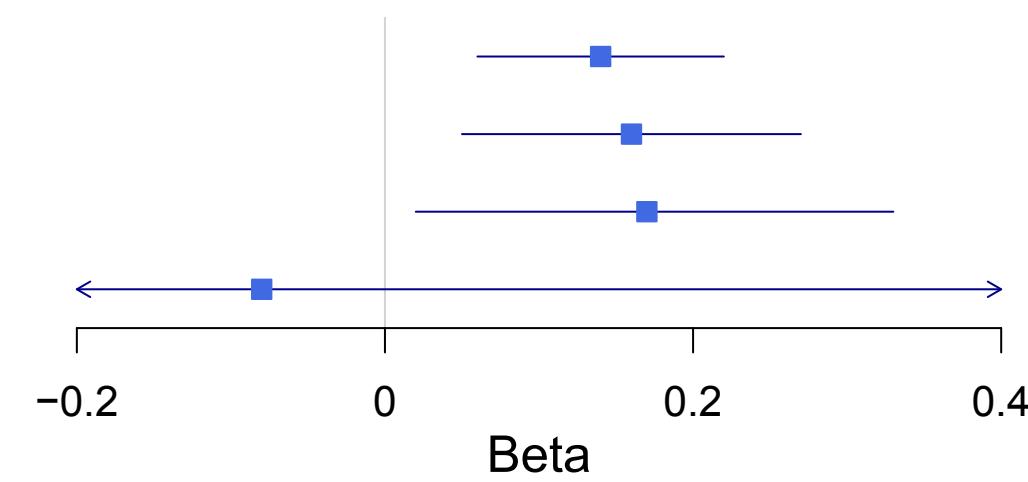


Fig. S10 Scatter plot and leave one out analysis plot for dietary habits.

Disease count

Method

Inverse variance weighted
Weighted median
Weighted mode
MR-Egger (SIMEX)

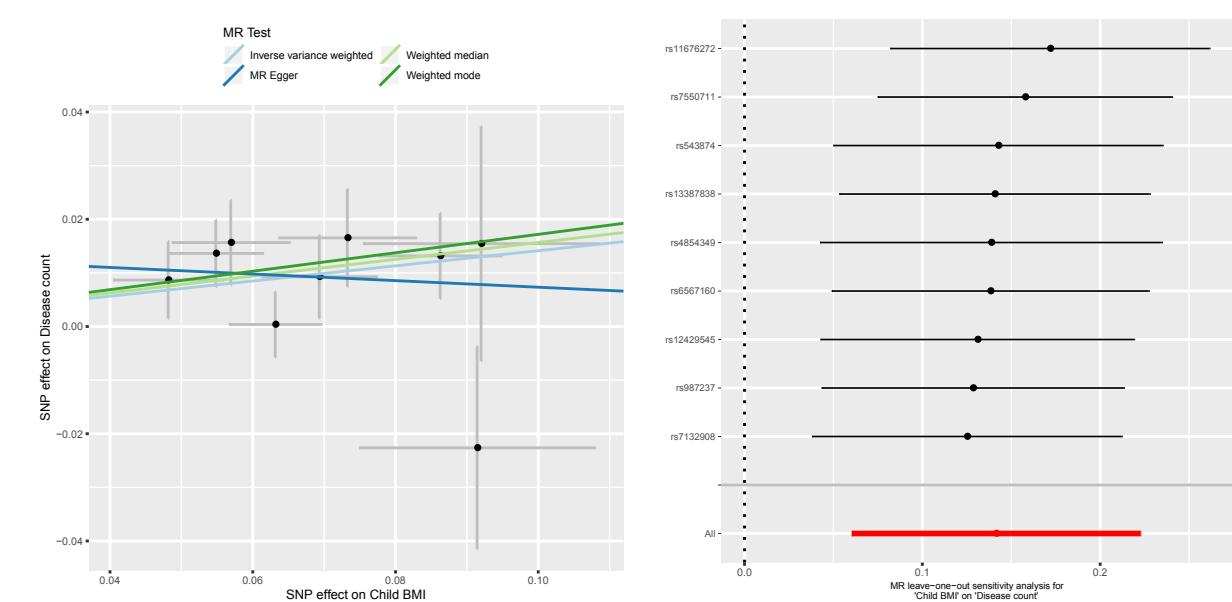


Beta (95% CI)

0.14 (0.06 to 0.22)
0.16 (0.05 to 0.27)
0.17 (0.02 to 0.33)
-0.08 (-0.63 to 0.46)

P value

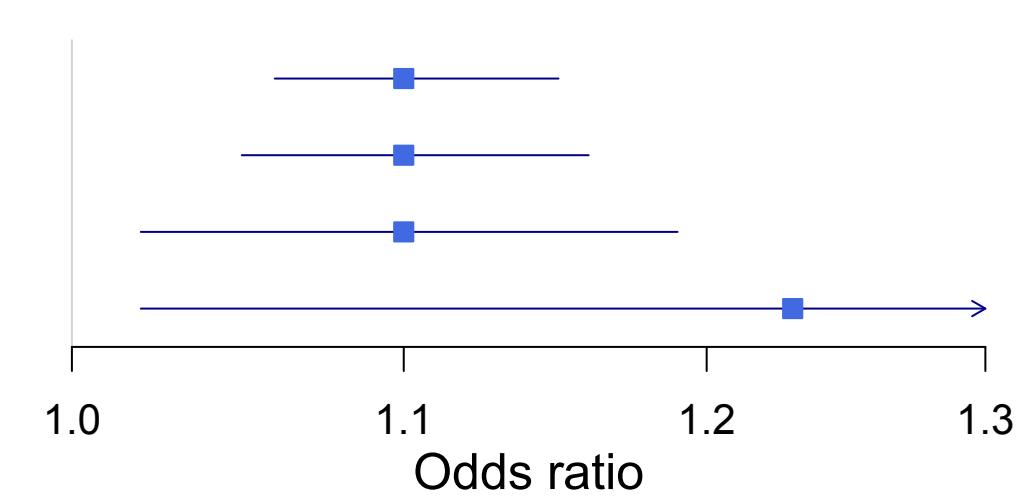
6.32×10^{-4}
 5.72×10^{-3}
 6.17×10^{-2}
 7.76×10^{-1}



Coronary artery disease

Method

Inverse variance weighted
Weighted median
Weighted mode
MR-Egger (SIMEX)

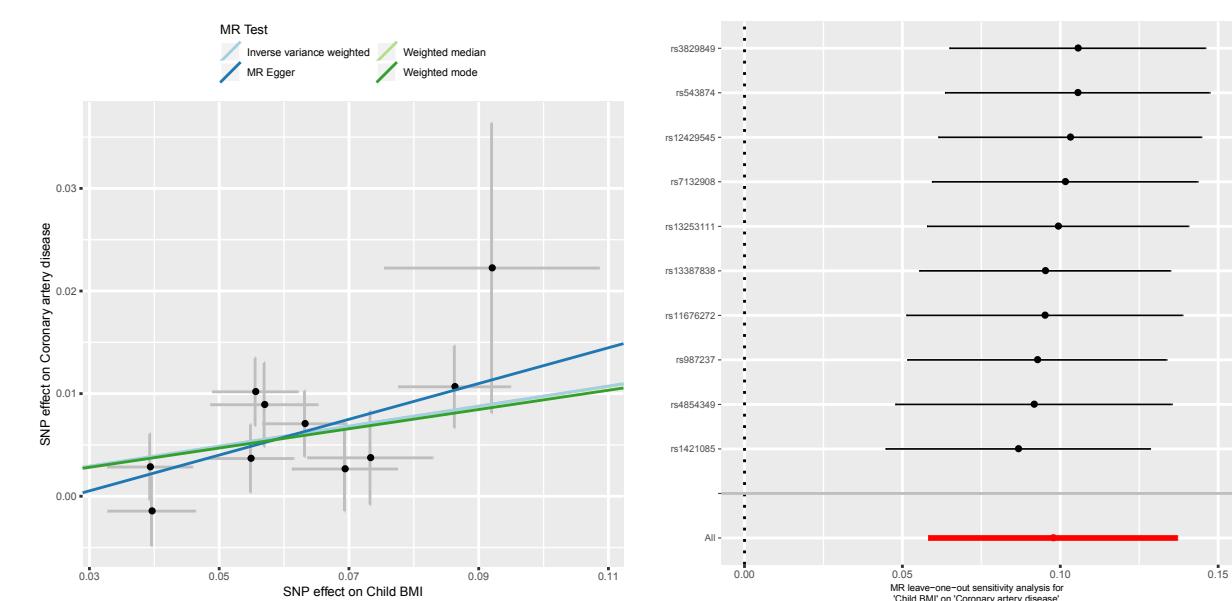


Odds ratio (95% CI)

1.10 (1.06 to 1.15)
1.10 (1.05 to 1.16)
1.10 (1.02 to 1.19)
1.23 (1.02 to 1.49)

P value

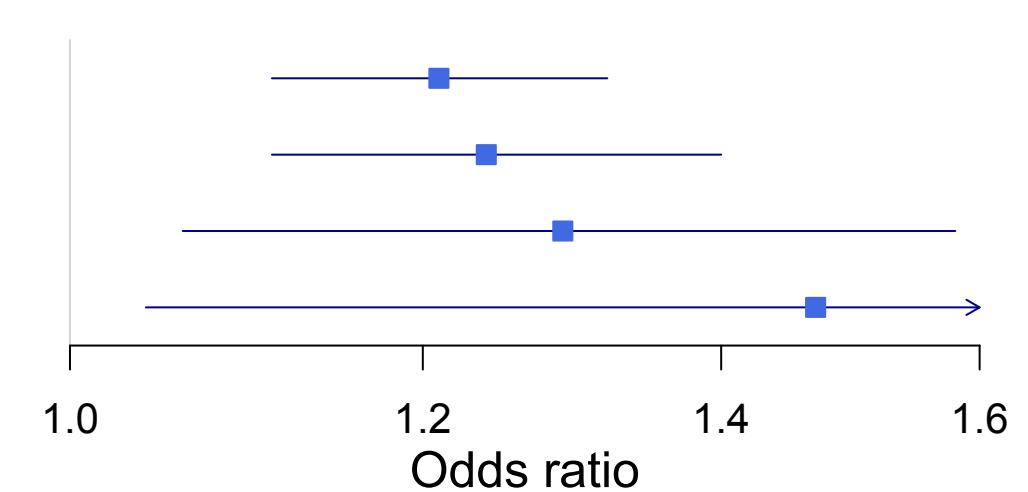
1.20×10^{-6}
 2.65×10^{-4}
 4.16×10^{-2}
 5.99×10^{-2}



Hypertensive Disease

Method

Inverse variance weighted
Weighted median
Weighted mode
MR-Egger (SIMEX)

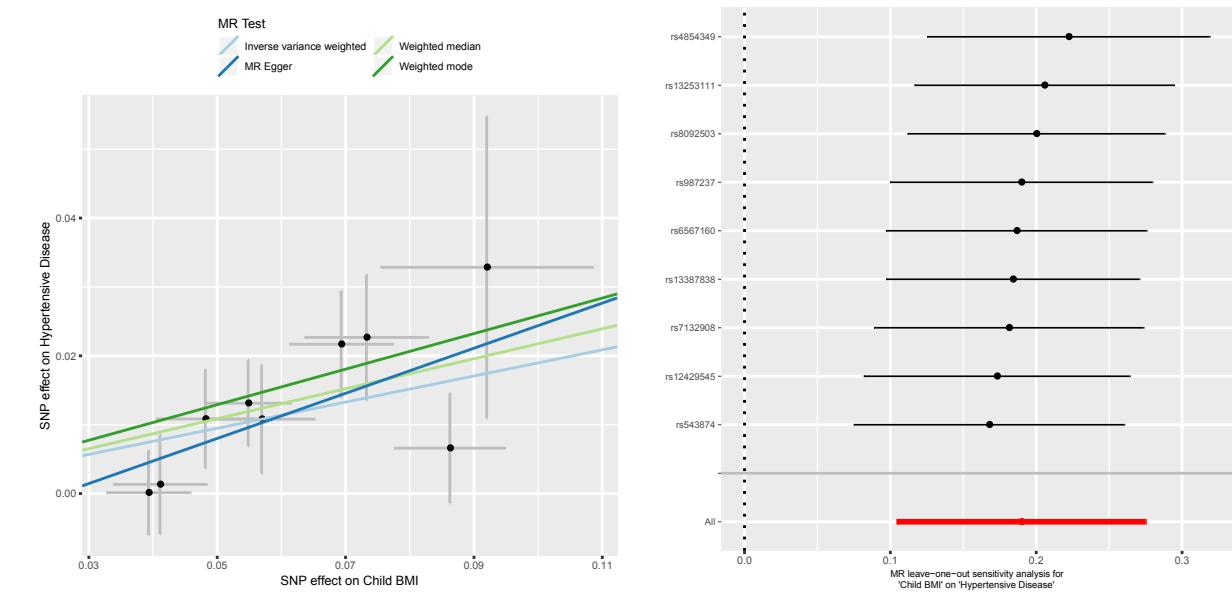


Odds ratio (95% CI)

1.21 (1.11 to 1.32)
1.24 (1.11 to 1.40)
1.29 (1.06 to 1.58)
1.47 (1.04 to 2.09)

P value

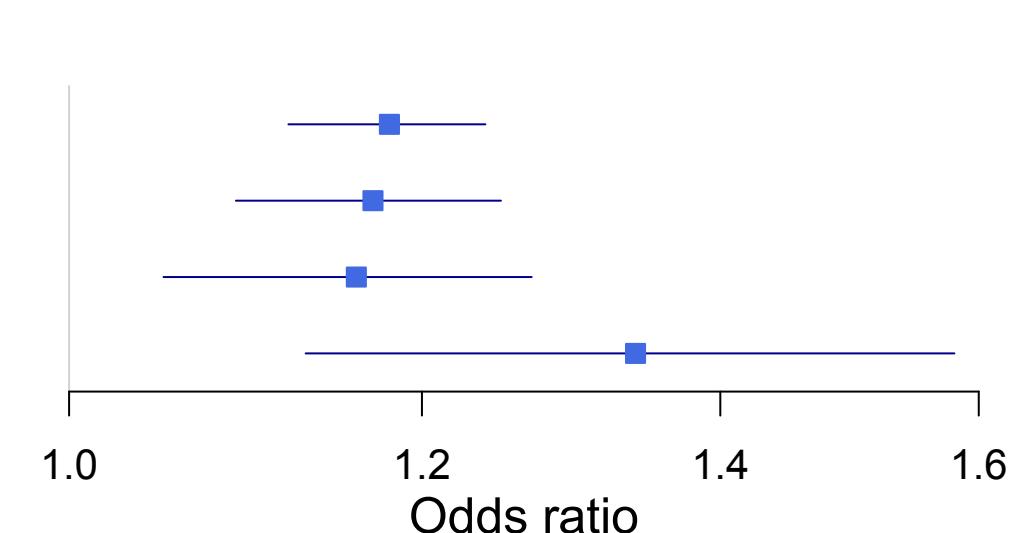
1.33×10^{-5}
 2.78×10^{-4}
 3.30×10^{-2}
 6.84×10^{-2}



Type 2 diabetes

Method

Inverse variance weighted
Weighted median
Weighted mode
MR-Egger (SIMEX)

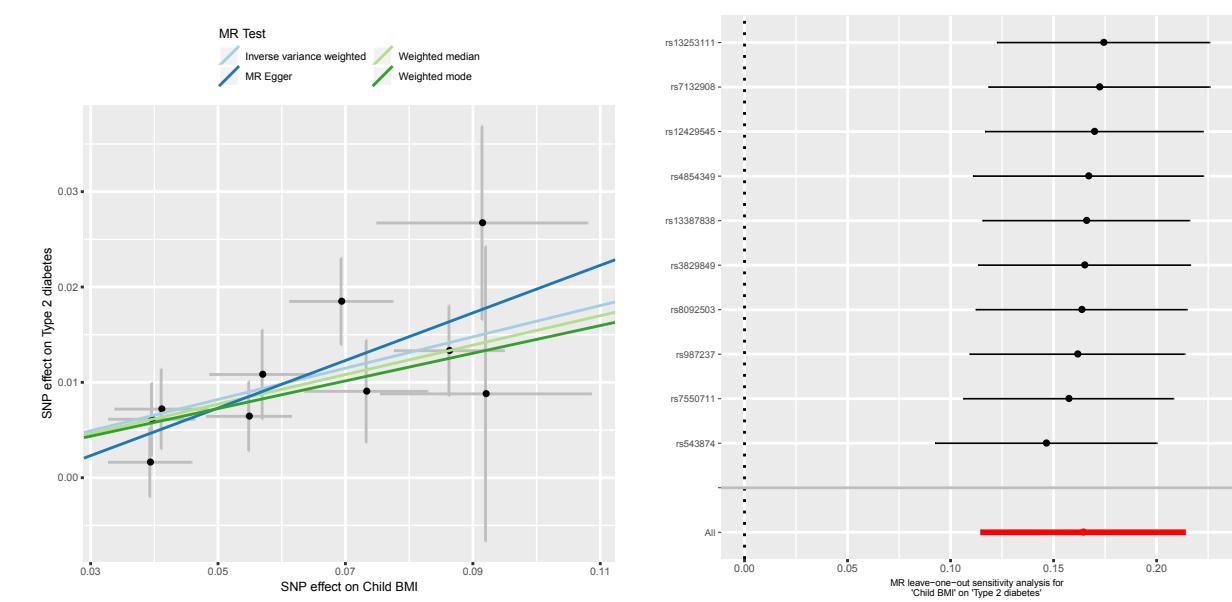


Odds ratio (95% CI)

1.18 (1.12 to 1.24)
1.17 (1.09 to 1.25)
1.16 (1.05 to 1.27)
1.34 (1.13 to 1.58)

P value

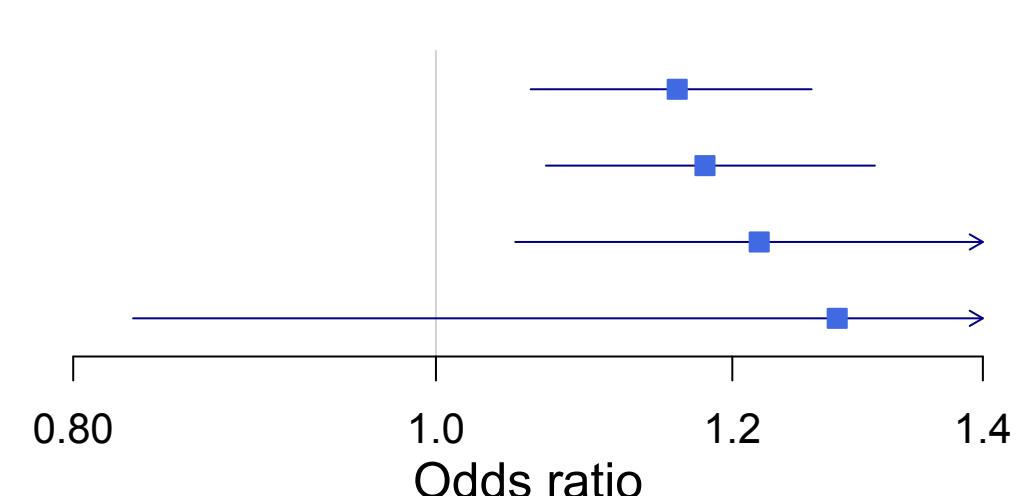
8.85×10^{-11}
 2.33×10^{-5}
 1.29×10^{-2}
 9.24×10^{-3}



Osteoarthritis

Method

Inverse variance weighted
Weighted median
Weighted mode
MR-Egger (SIMEX)



Odds ratio (95% CI)

1.16 (1.06 to 1.26)
1.18 (1.07 to 1.31)
1.22 (1.05 to 1.41)
1.28 (0.83 to 1.99)

P value

1.04×10^{-3}
 1.40×10^{-3}
 2.53×10^{-2}
 2.93×10^{-1}

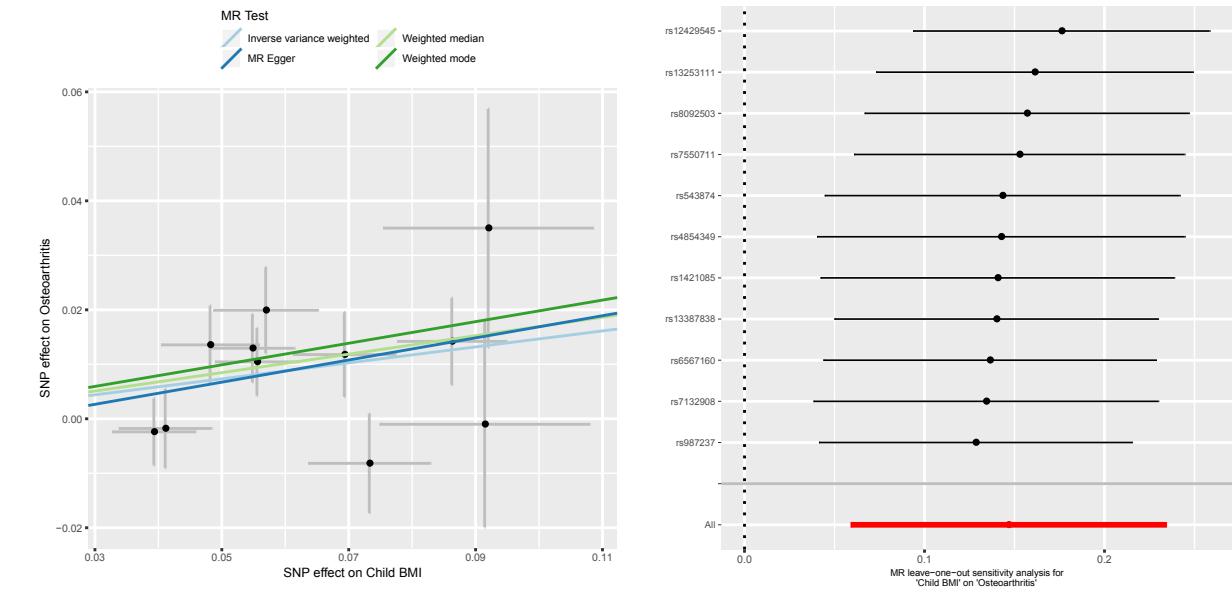
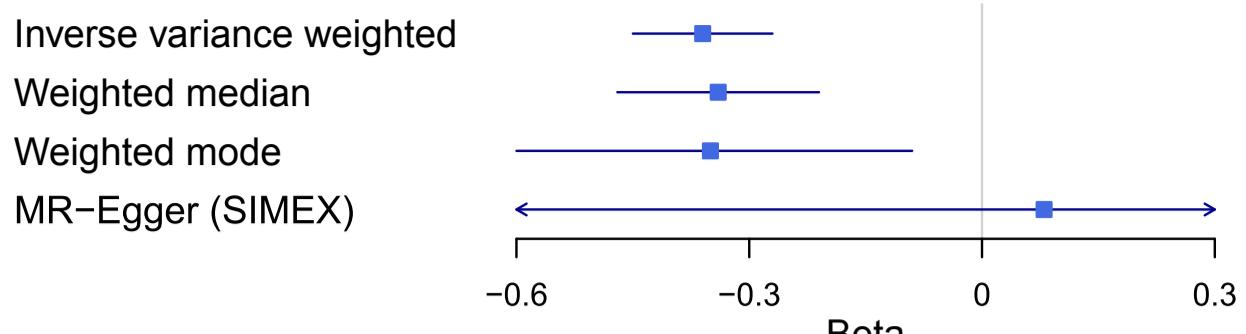


Fig. S11 Summary Mendelian randomization (MR) estimates derived from the inverse-variance weighted, MR-Egger, weighted median and weighted mode-based methods for additional datasets. Childhood BMI was used as exposure and significant associations were detected for these traits. Scatter plot and leave one out analysis plot for each trait are also shown.

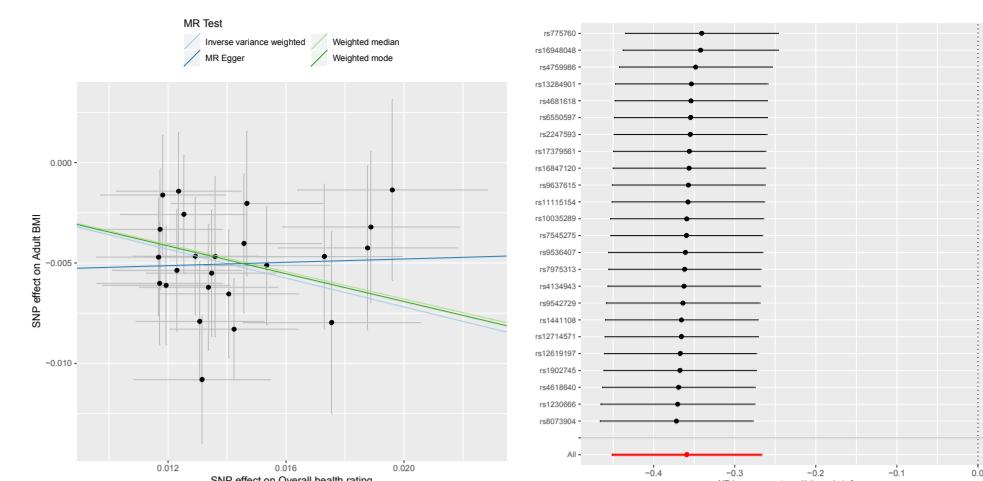
Overall health rating

Method



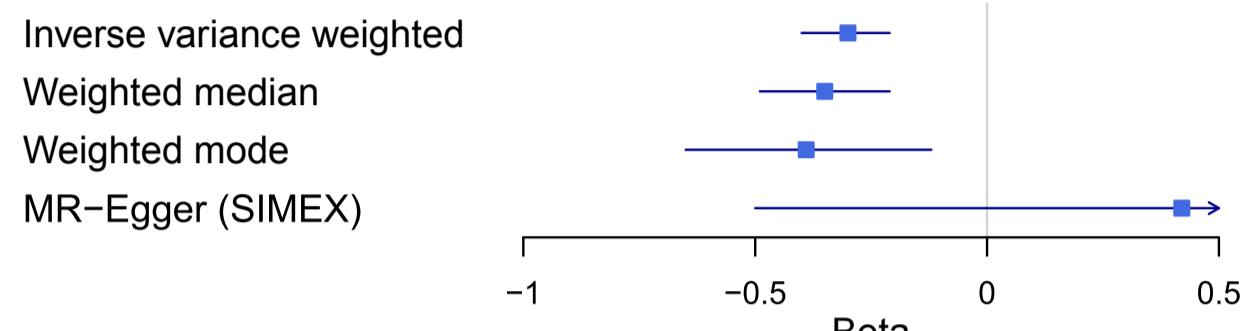
Beta (95% CI)

	Beta (95% CI)	P value
Inverse variance weighted	-0.36 (-0.45 to -0.27)	2.41×10^{-14}
Weighted median	-0.34 (-0.47 to -0.21)	1.91×10^{-7}
Weighted mode	-0.35 (-0.60 to -0.09)	1.31×10^{-2}
MR-Egger (SIMEX)	0.08 (-0.65 to 0.81)	8.39×10^{-1}



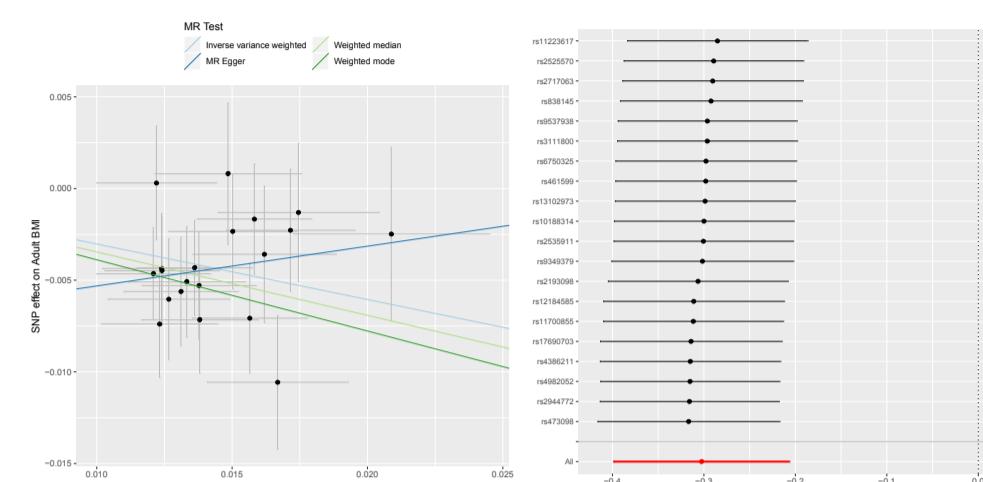
Alcohol intake frequency

Method



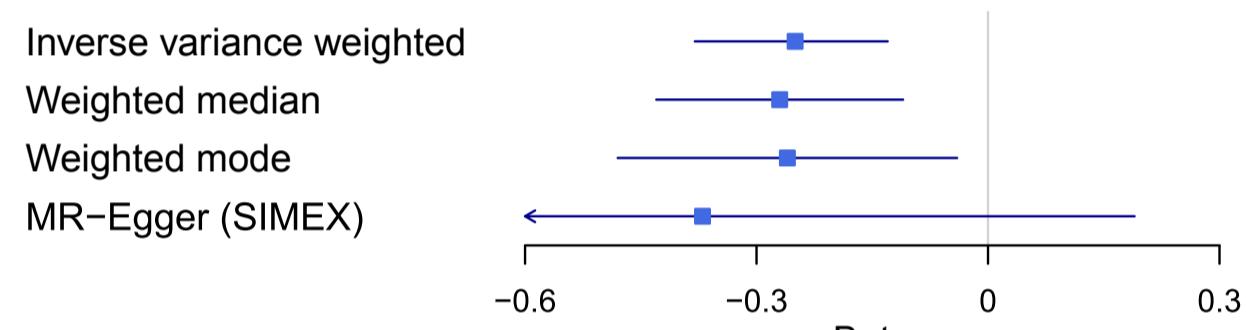
Beta (95% CI)

	Beta (95% CI)	P value
Inverse variance weighted	-0.30 (-0.40 to -0.21)	6.86×10^{-10}
Weighted median	-0.35 (-0.49 to -0.21)	1.16×10^{-6}
Weighted mode	-0.39 (-0.65 to -0.12)	9.97×10^{-3}
MR-Egger (SIMEX)	0.42 (-0.50 to 1.35)	3.83×10^{-1}



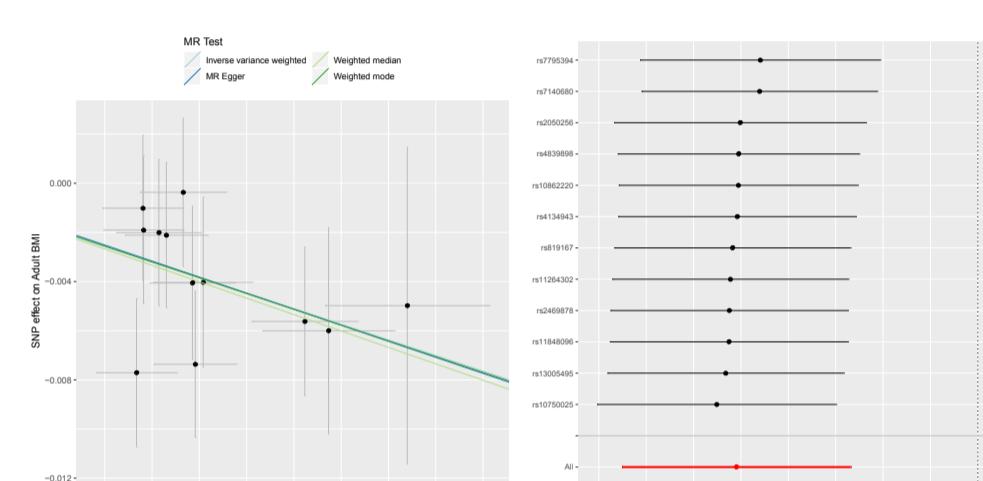
Usual walking pace

Method



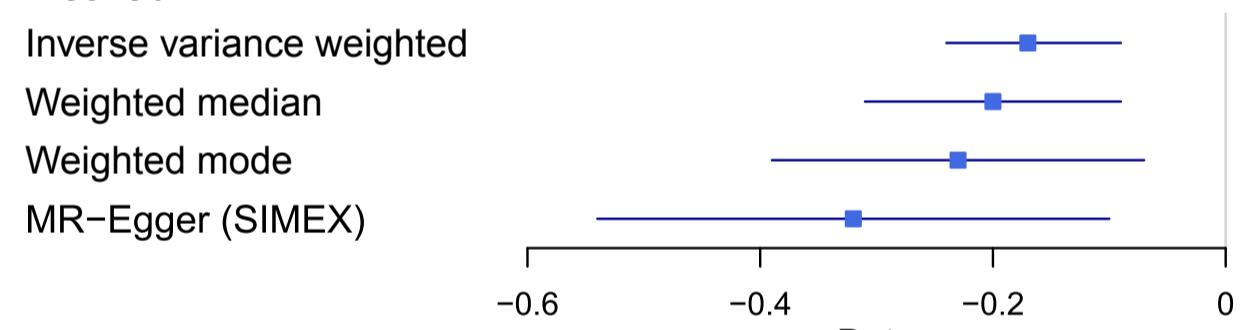
Beta (95% CI)

	Beta (95% CI)	P value
Inverse variance weighted	-0.25 (-0.38 to -0.13)	3.53×10^{-5}
Weighted median	-0.27 (-0.43 to -0.11)	1.23×10^{-3}
Weighted mode	-0.26 (-0.48 to -0.04)	4.29×10^{-2}
MR-Egger (SIMEX)	-0.37 (-0.94 to 0.19)	2.26×10^{-1}



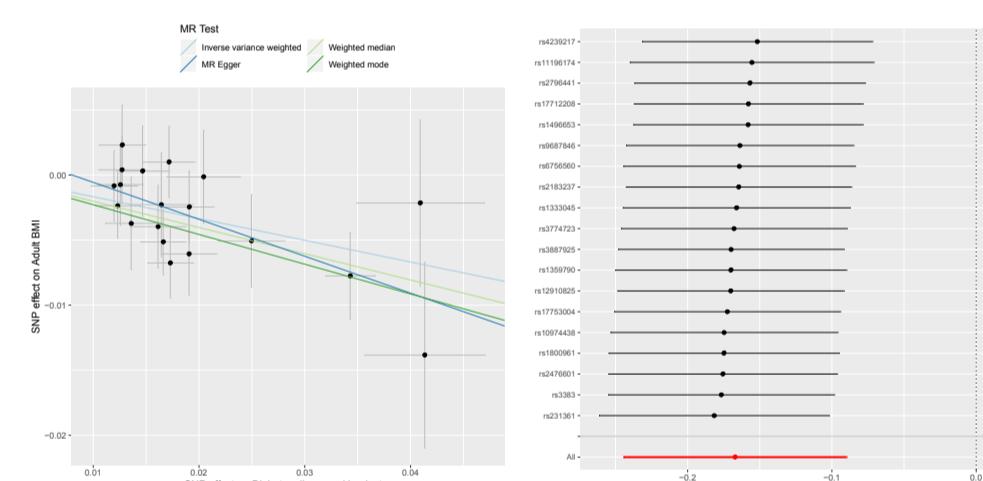
Diabetes diagnosed by doctor

Method



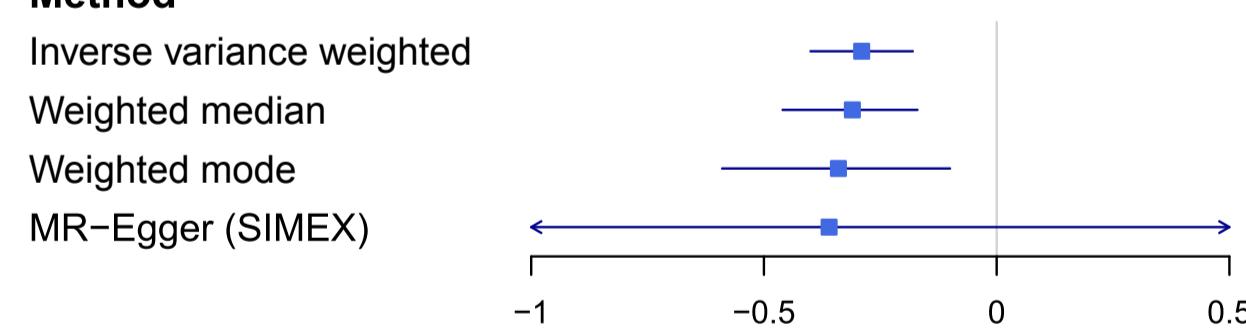
Beta (95% CI)

	Beta (95% CI)	P value
Inverse variance weighted	-0.17 (-0.24 to -0.09)	2.22×10^{-5}
Weighted median	-0.20 (-0.31 to -0.09)	3.24×10^{-4}
Weighted mode	-0.23 (-0.39 to -0.07)	1.09×10^{-2}
MR-Egger (SIMEX)	-0.32 (-0.54 to -0.10)	9.98×10^{-3}



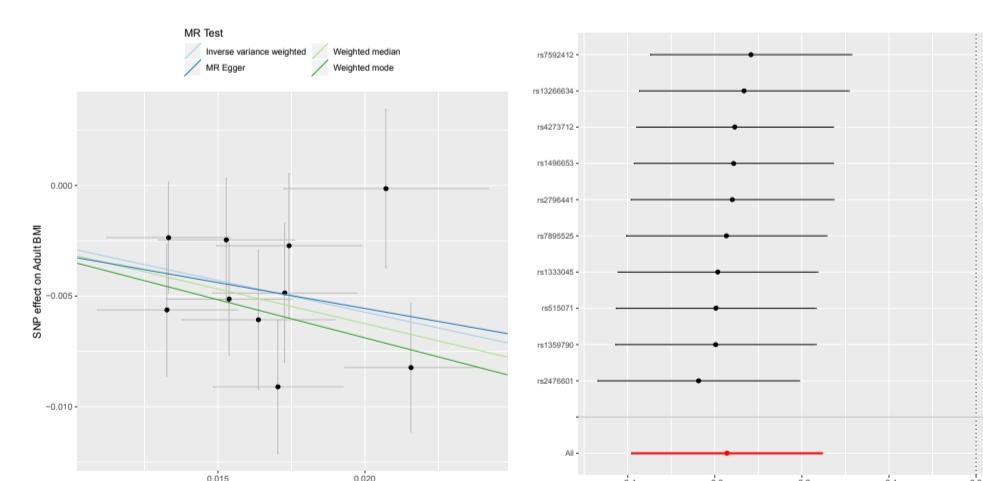
E10-E14 Diabetes mellitus

Method



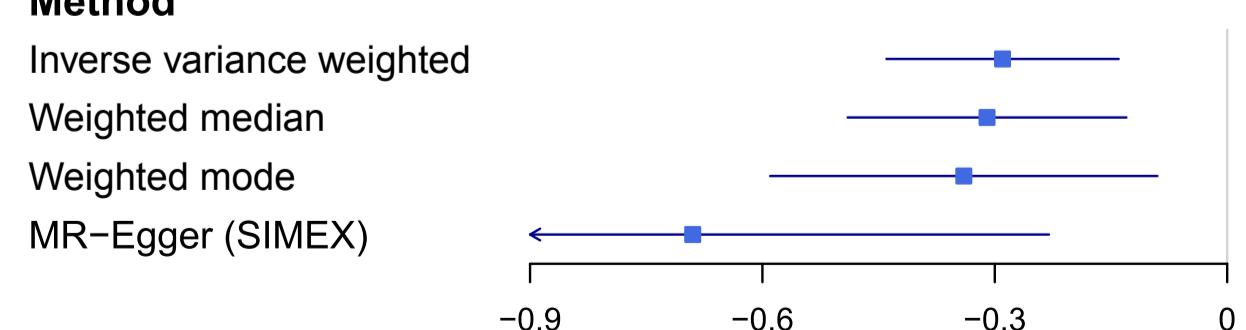
Beta (95% CI)

	Beta (95% CI)	P value
Inverse variance weighted	-0.29 (-0.40 to -0.18)	3.05×10^{-7}
Weighted median	-0.31 (-0.46 to -0.17)	2.53×10^{-5}
Weighted mode	-0.34 (-0.59 to -0.10)	2.24×10^{-2}
MR-Egger (SIMEX)	-0.36 (-1.33 to 0.60)	4.84×10^{-1}



E11 Non-insulin-dependent diabetes mellitus

Method



Beta (95% CI)

	Beta (95% CI)	P value
Inverse variance weighted	-0.29 (-0.44 to -0.14)	1.26×10^{-4}
Weighted median	-0.31 (-0.49 to -0.13)	7.79×10^{-4}
Weighted mode	-0.34 (-0.59 to -0.09)	5.61×10^{-2}
MR-Egger (SIMEX)	-0.69 (-1.16 to -0.23)	6.25×10^{-2}

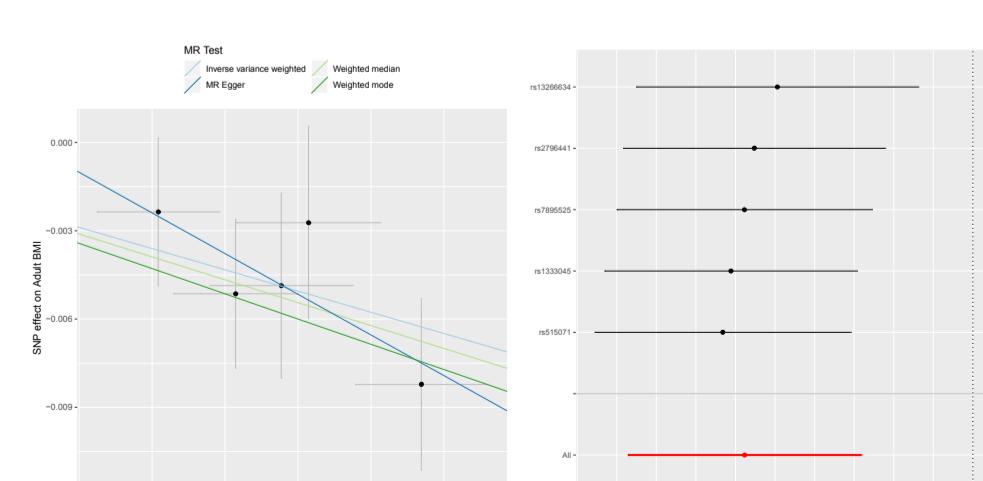


Fig. S12 Summary Mendelian randomization (MR) estimates derived from the main inverse-variance weighted, MR-Egger, weighted median and weighted mode-based methods for reverse analyses using adulthood BMI as outcome. Scatter plot and leave one out analysis plot for each trait are also shown.