EMIF Deliverable 6.6: The use of Mendelian Randomisation methods to investigate the causal significance of associated biomarkers

Executive summary

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Combined genetic and biomarker data on (a) candidate biomarkers identified in WP5 or other WP6 tasks and (b) targeted and untargeted metabolomics profiles collected in medium-sized well-phenotyped cohorts were used to investigate the causal association of biomarkers with the metabolic consequences using a Mendelian randomisation framework.

Contacts

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