# Annual review: year2

Matthew Lee 2020-01-09

#### Title

What lies behind the causal impact of body mass index (BMI) level and change on human health? Added value from complementary study design and deep metabolomic phenotyping

**Start date**: 01/10/2017

End date: 01/04/2021

#### Aim

Identify metabolites that sit on the causal pathway from increased adiposity to disease

#### **Objectives**

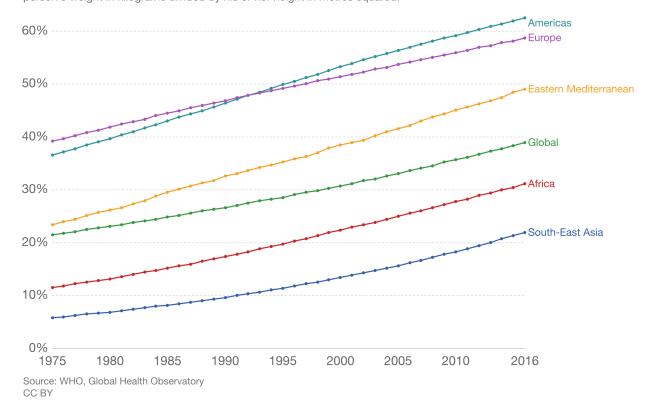
- 1. Identify diseases causally associated with increased adiposity
- 2. Identify and describe appropriate instrumentation of increased adiposity
- 3. Identify metabolites causally associated with increased adiposity
- 4. Design and implement methods and rules to cluster metabolites
- 5. Identify diseases causally associated with metabolites

### Overview

#### Share of adults that are overweight



Share of adults aged 18 years and older who have a body-mass index (BMI) greater than or equal to 25. BMI is a person's weight in kilograms divided by his or her height in metres squared.

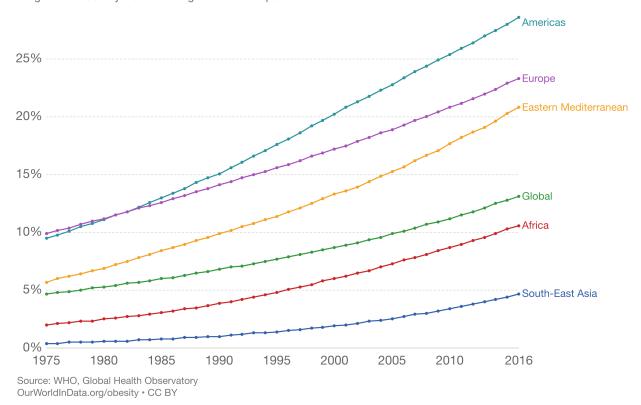


Proportion of overweight individuals

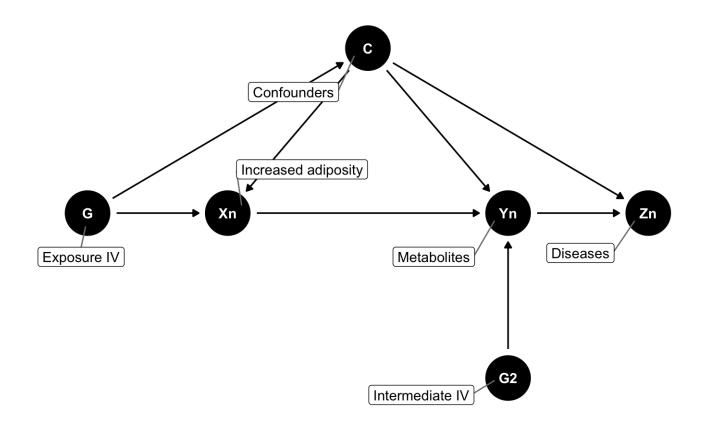
#### Share of adults that are obese



Obesity is defined as having a body-mass index (BMI) equal to or greater than 30. BMI is a person's weight in kilograms divided by his or her height in metres squared.



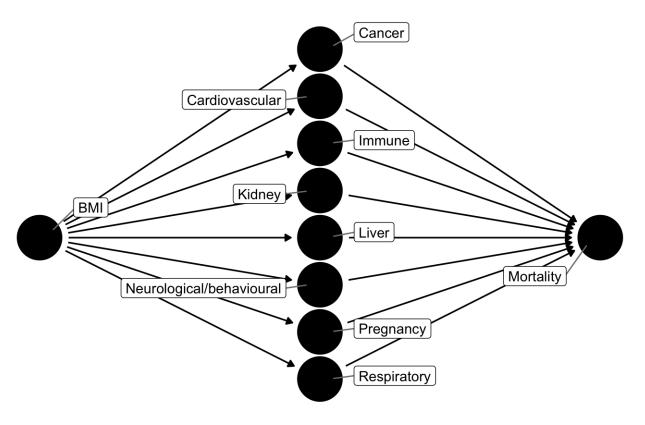
Proportion of obese individuals



Can we build a flowchart of decisions to achieve this?

#### Intro/overview

- What is the problem (100%)
- What is adiposity (100%)
  - inc. adipose cells idea of genetics and metabolites introduced
  - inc. different measures
- Broad overview of diseases (90%)
- Application of metabolites and MR (0%)



MELODI analysis: 975,402 articles; 10,828 enriched terms, filtering left 77 terms, 8 categories plus 'Other'

#### Intermediates identified between BMI and all targets using MELODI

Cancer	Cardiovascular	Immune	Kidney	Liver	Neuro_behav	Pregnancy	Respiratory	Other
Primary carcinoma of the liver cells	Heart failure	Pancreatitis	End stage renal failure	Liver diseases	Depressive disorder	Pre-Eclampsia	Tuberculosis	Metabolic syndrome
Malignant neoplasm of stomach	Anemia	Inflammatory disorder	Kidney Failure, Chronic	Non- alcoholic fatty liver	Dementia	Pregnancy	Sleep Apnea, Obstructive	Cessation of life
Malignant neoplasm of prostate	Dyslipidemias	Immunocompromised Host	Glomerular Filtration Rate	Liver and Intrahepatic Biliary Tract Carcinoma		Hypertension induced by pregnancy	Pneumonia	Malnutrition
Malignant neoplasm of lung	Cerebrovascular accident	Bacteremia	Kidney Diseases	Chronic liver disease			Chronic Obstructive Airway Disease	Diabetic
Common Neoplasm	Cardiovascular Diseases	Septicemia	Kidney Failure				Respiration Disorders	Multiple Organ Failure
Liver neoplasms	Atherosclerosis	Lupus Erythematosus, Systemic	Renal function				Respiratory Distress Syndrome, Adult	Fibrosis
Malignant disease Carcinoma	Myocardial Infarction	Sepsis Syndrome					Respiratory Tract Infections	Deglutition Disorders
of the Large Intestine	Ischemic stroke						Respiratory Failure	Vitamin D Deficiency
Pancreatic carcinoma	Acute coronary syndrome Atrial Fibrillation Coronary heart						Acute respiratory failure	
	disease Systemic arterial pressure							
	Thrombosis Cerebrovascular Disorders							
	Acute myocardial infarction							
	Sinus rhythm Cardiomyopathies Myocardial Ischemia							
	Peripheral Vascular Diseases							9/24
	Vascular calcification							

### Systematic review

All MR studies using a measure of increased adiposity as the exposure; ~150 articles included

- · Literature search (100%)
- · Screening (100%)
- Data extraction (30%)
- Writing (40%)

#### Instrumentation

- Current instrumentation practices (will get from SR)
- GWAS descriptions (80%)
- Selecting instruments (0%)
- Writing (30%)

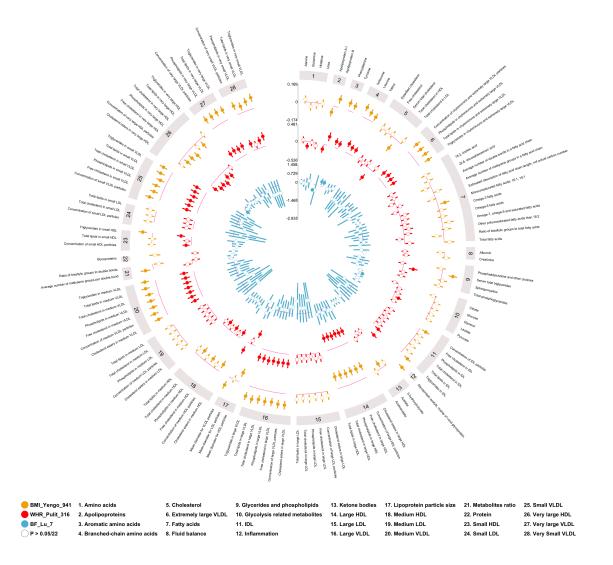
### Observational analysis

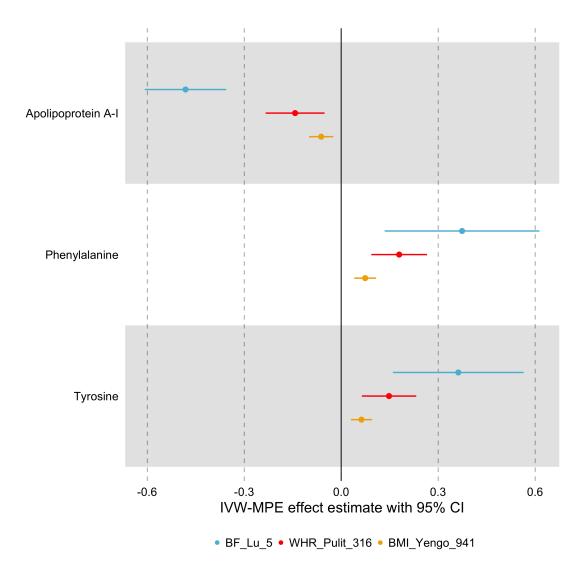
- BMI/WHR/BF% -> metabolites in ALSPAC/FGFP/Biobank (20%)
- · Writing (0%)

Waiting on Biobank data...

### MR step 1

- BMI/WHR/BF% -> metabolites analysis (100%)
- Wiriting/paper (60%)

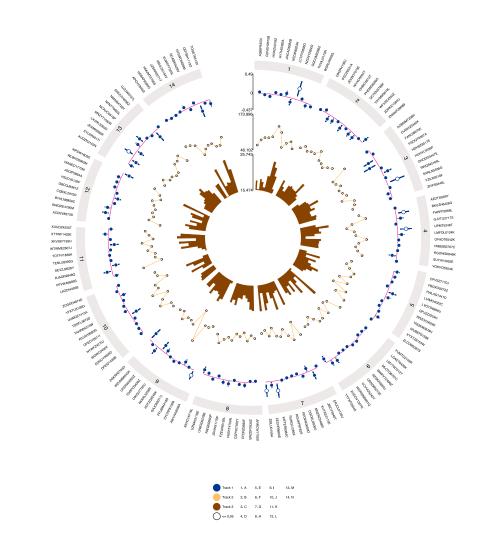




#### MR Viz

- web app (90%)
- · R package (90%)
- Writing/paper (60%)
- · App / R package

```
library(EpiCircos)
circos plot(track number = 3,
            track1 data = EpiCircos::EpiCircos data,
            track2 data = EpiCircos::EpiCircos data,
            track3 data = EpiCircos::EpiCircos data,
            track1 type = "points", track2_type = "lines", track3_type = "bar",
            label column = 1, section column = 2,
            estimate column = 4, pvalue column = 5,
            pvalue adjustment = 1,
            lower ci = 7, upper ci = 8,
            lines column = 10, lines type = "o",
            bar column = 9,
            legend = TRUE,
            track1 label = "Track 1",
            track2 label = "Track 2",
            track3 label = "Track 3",
            pvalue label = "<= 0.05",
            circle size = 25)
```



### Clustering metabolites

- not started
- Writing (0%)
- Priors
  - class
  - subclass
  - biological pathway
  - size
  - shared genetic variants

- No priors
  - PCA
  - factor analysis
  - Hierarchical clustering
  - density clustering
  - self organising map
  - LDSR
  - ontology
    - have discussed with Ben Elsworth (pipeline can be adpated)

### Rules for instrumenting clusters

- have ideas
- not started in parallel with Chapter 7
- · Writing (0%)

### MR step 2

- metabolites -> diseases analysis (20%) will be scaled quickly
- Final figure for global overview (1%)
  - combining MR step 1 and MR step 2 results together
  - possible to incorporate multiple clustering methods
- · Writing (0%)

#### Discussion/limitations/conclusion

- Final flowchart figure of pipeline
- not started
- · Writing (0%)