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Unpacking the Exposome's Influence on Chronic Kidney Disease: The Mediating Role of Biomarkers

Computational Epidemiology Group 3

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Background

Chronic Kidney Disease and the Exposome

Status quo in the UK:

- 22% of adults over 35 and 50% of adults over 75 years of age have CKD (~7.2m individuals)
- A substantial proportion of CKD cases remain undiagnosed in the general population
- CKD is an established risk factor for cardiovascular disease, cognitive impairment, and stroke, which collectively cost the UK economy >£100bn p.a.

A plethora of exposures are associated with CKD risk:

- Smoking
- Alcohol Consumption
- Obesity and diet
- PM2.5
- Deprivation

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Our research question:

To what extent does the exposome contribute to the development of chronic kidney disease, and how are these effects mediated via biomarkers?

Study Motivations

Inform targeted prevention and personalised approaches

Aid early intervention strategies

Comprehending how biomarkers mediate exposure-related CKD risk will...

Reduce disease burden and associated costs

Help identify new therapeutic targets

Data Processing

Preprocessing steps

Exclusion Criteria:

- Individuals not in UK Biobank
- Prevalent cases
- Limitations

Scaling/One-Hot Encoding:

 Reference dummy column removed to aid interpretation and reduce collinearity in analysis

Imputation (MICE):

 Test data imputed based off training data, preventing data leakage

Case Definition:

 Cases - persons diagnosed with chronic kidney disease (CKD) as reported by HES and/or registry data.

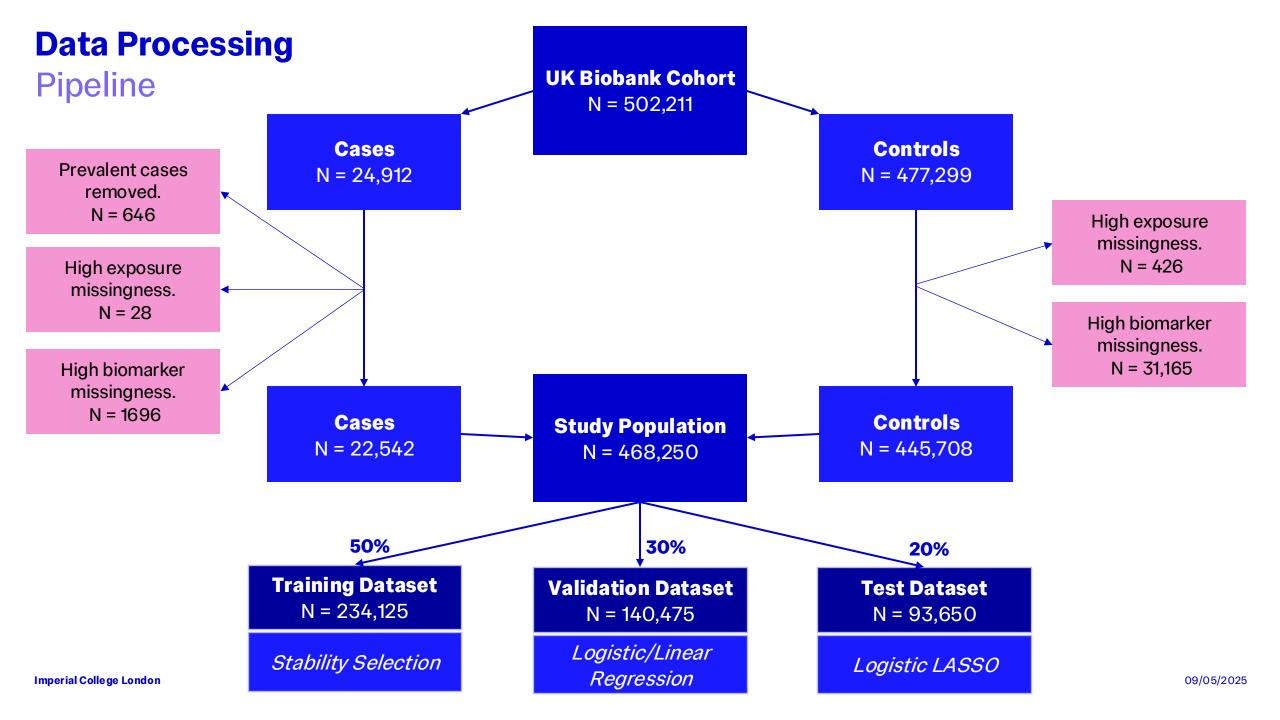
Missingness:

- Individuals with >50% missing biomarker and/or exposure data removed
- Biomarkers with >50% missingness removed (rheumatoid factor, oestradiol)

Data Splits:

50:30:20 train-validation-test split

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Methods

Overview

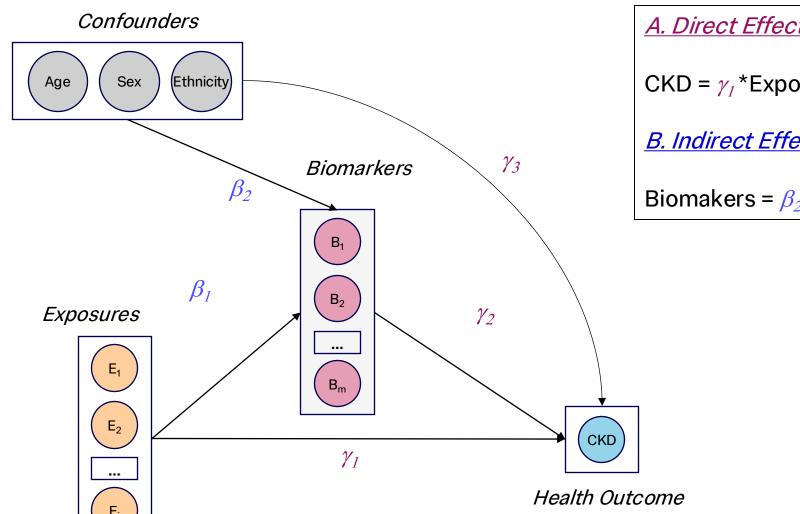
Stability Selection

Logistic and Linear Regression

Prediction with LASSO and elastic net

Modelling approach

Selected DAG of informative relationships



A. Direct Effect Model:

CKD = γ_1 *Exposures + γ_2 * Biomarkers + γ_3 * Confounders

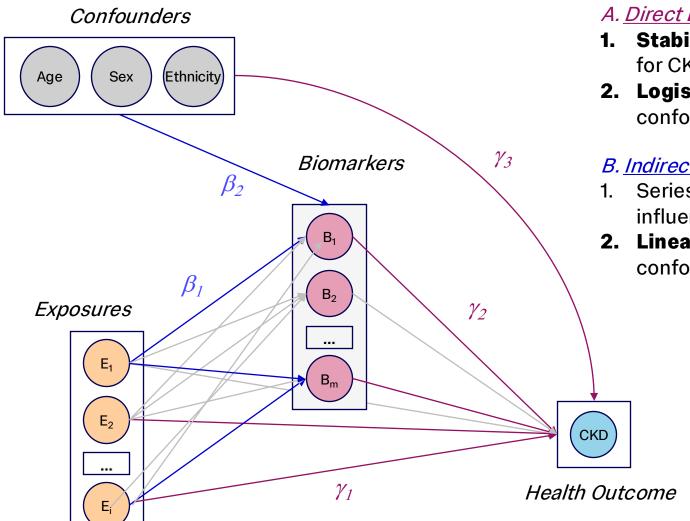
B. Indirect Effect Model:

Biomakers = β_2 * Confounders + β_1 * Exposures

Exposures: N = 34 Biomarkers: N = 41

Modelling approach

Selected DAG of informative relationships



A. Direct Effect Model:

- Stability LASSO to select essential exposures and biomarkers for CKD.
- **2. Logistic regression** with only the selected variables and confounders to retrieve <u>unbiased</u> estimates.

B. Indirect Effect Model:

- 1. Series of **Stability LASSOs** for the selection of exposures that influence previously selected biomarkers.
- 2. Linear regression with only the selected exposures and confounders to retrieve unbiased estimates on each biomarkers.
 - **Direct Effect** of exposure $i(\gamma_{1i})$
 - Indirect Effect of exposure i

$$\sum_{\substack{j \text{ selected} \\ \text{biomarkers}}} \gamma_{2_j} \, \beta_{1_ij}$$

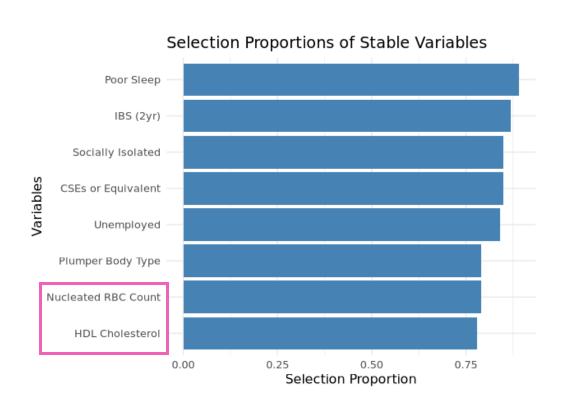
 γ_{1i}, γ_{2i} : log-odds $\beta_{1_{ij}}$: linear coefficient

Descriptive Analysis

	Total (n = 468,250)			
	Case	Control	p-value	Missing(%)
Sample Characteristics	22,542	445,708	-	
Demographics				
Sex = Male (%)	11659 (51.7)	202665 (45.5)	<0.001	0
Age (median [IQR])	64.00 [60.00, 67.00]	57.00 [50.00, 63.00]	<0.001	0
Ethnicity (%)	-	<u> </u>	<0.001	0.1
White (reference)	21137 (93.9)	420369 (94.4)		
Asian	561 (2.5)	9933 (2.2)		
Black	443 (2.0)	6808 (1.5)		
Other	362 (1.6)	8045 (1.8)		
Clinical Variables				
BMI (median [IQR])	28.85 [25.93, 32.56]	26.64 [24.07, 29.75]	< 0.001	0.4
Systolic Blood Pressure (median [IQR])	142.00 [130.00, 155.50]	136.00 [124.50, 149.00]	< 0.001	8.6
Socioeconomic Variables				
Employment Status (%)			< 0.001	0.5
Employed (reference)	6399 (28.5)	252158 (56.9)		
Looking after home and/or family	309 (1.4)	12604 (2.8)		
Other	219 (1.0)	5794 (1.3)		
Retired	13808 (61.6)	151814 (34.2)		
Unable to work due to sickness	1404 (6.3)	13838 (3.1)		
Unemployed	286 (1.3)	7334 (1.7)		
Own or Rent (%)			< 0.001	0
Own (reference)	18796 (83.4)	395908 (88.8)		
Rent	3091 (13.7)	39689 (8.9)		
Other	655 (2.9)	10111 (2.3)		
Lifestyle Variables				
Alcohol Intake (%)			< 0.001	0.6
Never/Abstinent (Excluding previous drinkers)	1580 (7.1)	18993 (4.3)		
Previous Drinker (who currently never drink)	1407 (6.3)	15362 (3.5)		
Light (reference)	12810 (57.3)	242718 (54.8)		
Moderate	3378 (15.1)	84242 (19.0)		
Heavy*	1922 (8.6)	50260 (11.3)		
Abusive*	1258 (5.6)	31482 (7.1)		
Smoking Status (%)			<0.001	0.5
Never (reference)	10161 (45.4)	244906 (55.2)		
Previous	9651 (43.1)	152099 (34.3)		
Current	2557 (11.4)	46600 (10.5)		

Stably Selected Variables - Direct Model

Selection Proportions of Stable Variables Predicting CKD (Direct Model)



Stability selection threshold (π)	0.78
Optimal lambda (λ)	0.000492

- 8 variables were selected:
 - 6 are exposures: poor sleep, having IBS in last 2
 years, being socially isolated, having education up to
 CSEs or equivalent, being unemployed, having a
 plumper body composition
 - 2 are biomarkers: Nucleated RBC count, HDL cholesterol
- These variables were included in the logistic regression model to estimate direct effects on CKD risk.

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Alcohol: Moderate

Alcohol: Heavy Alcohol: Abusive

MET Score

0.00

0.25

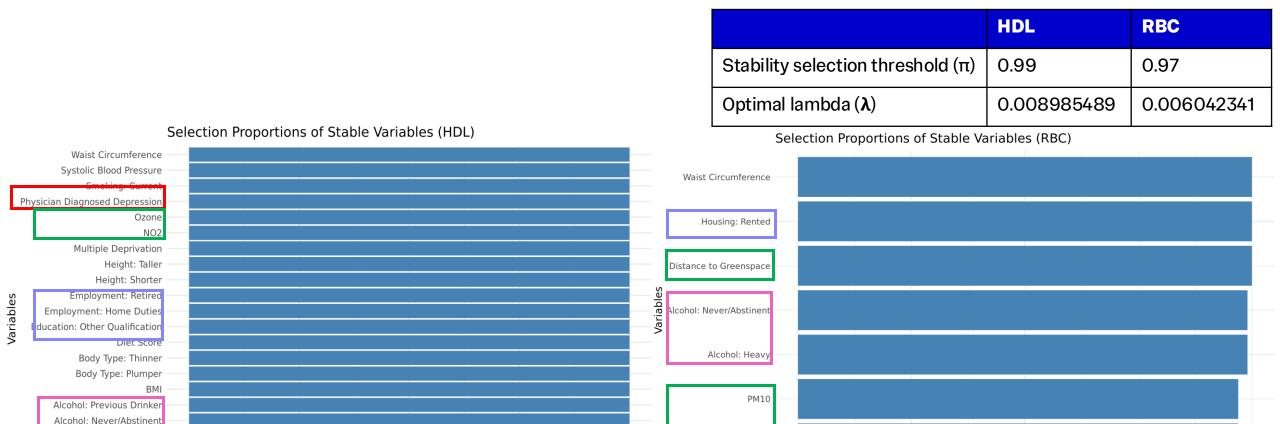
0.50

Selection Proportion

0.75

Stably Selected Variables - Indirect Model

Selection Proportions of Stable Variables Predicting CKD (Indirect Model)



1.00

Ozone

0.00

0.25

0.50

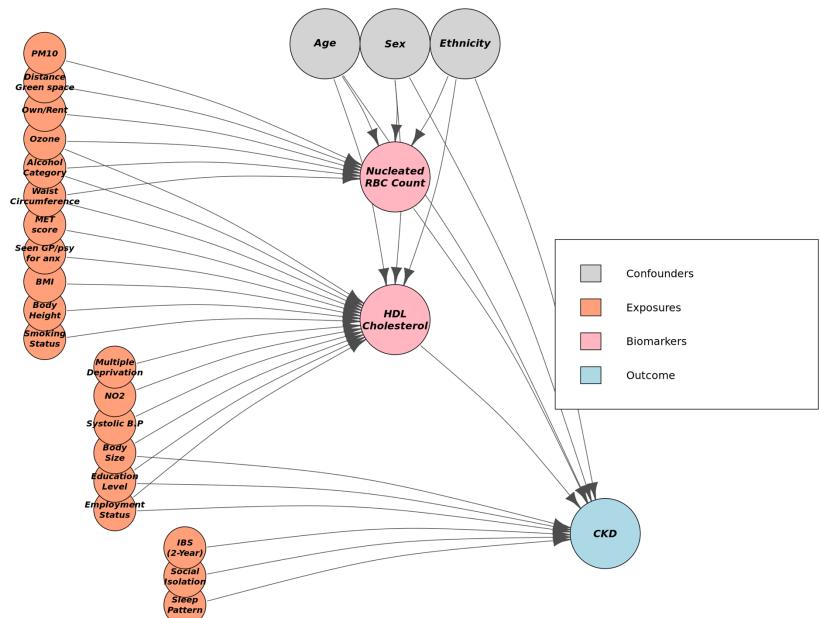
Selection Proportion

0.75

1.00

ResultsSparse DAG

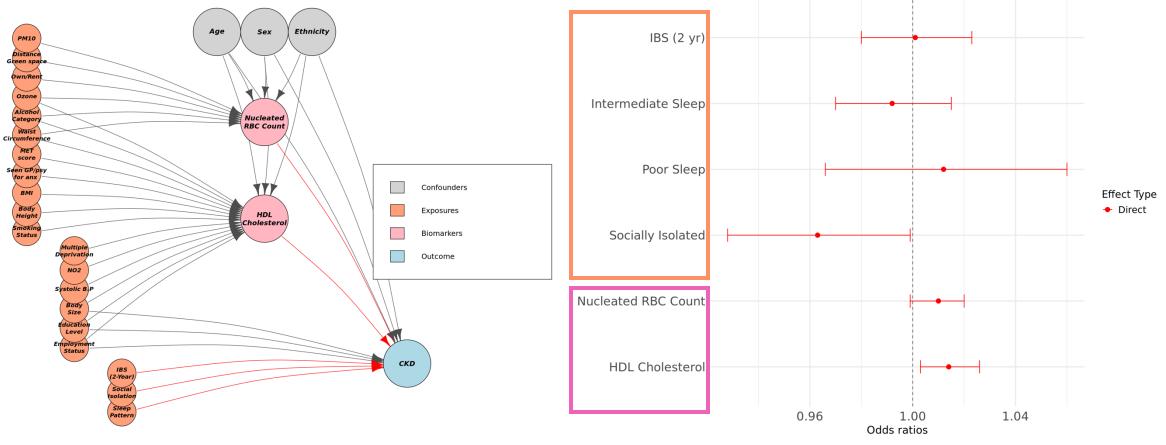
Sparse DAG - Mediation Analysis



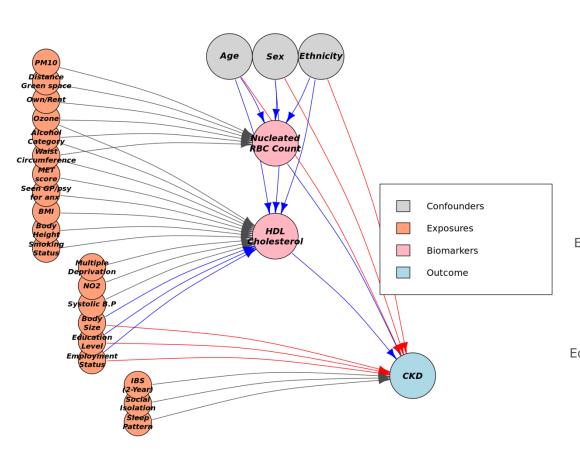
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Direct effect

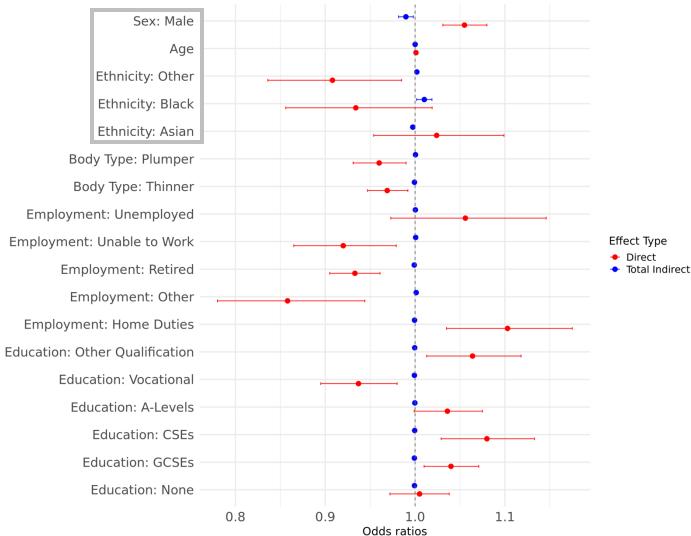
Direct Effects:



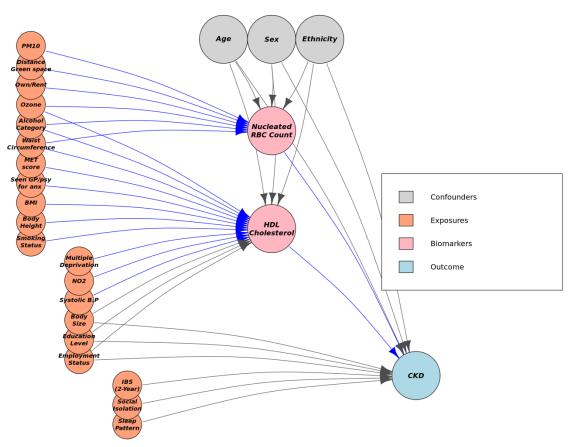
Direct and indirect effect



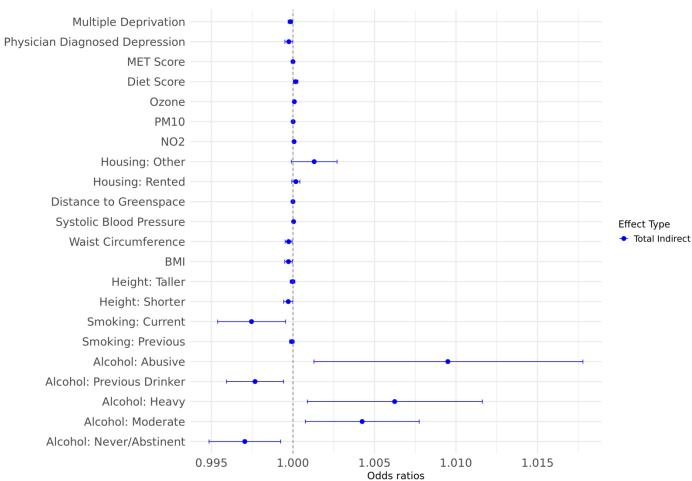
Direct and Indirect Effects:



Indirect/Mediated effect



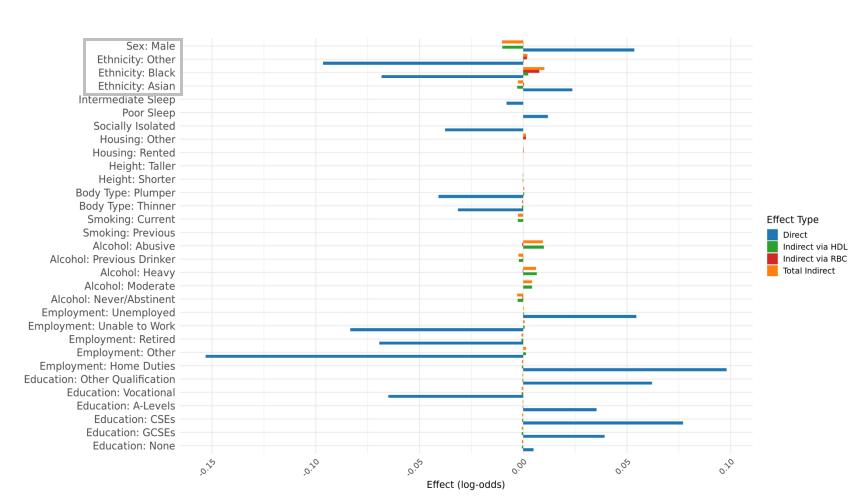
Mediated Effects:



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Modelling approach

Summary of direct and indirect (mediated) effects – Categorical variables

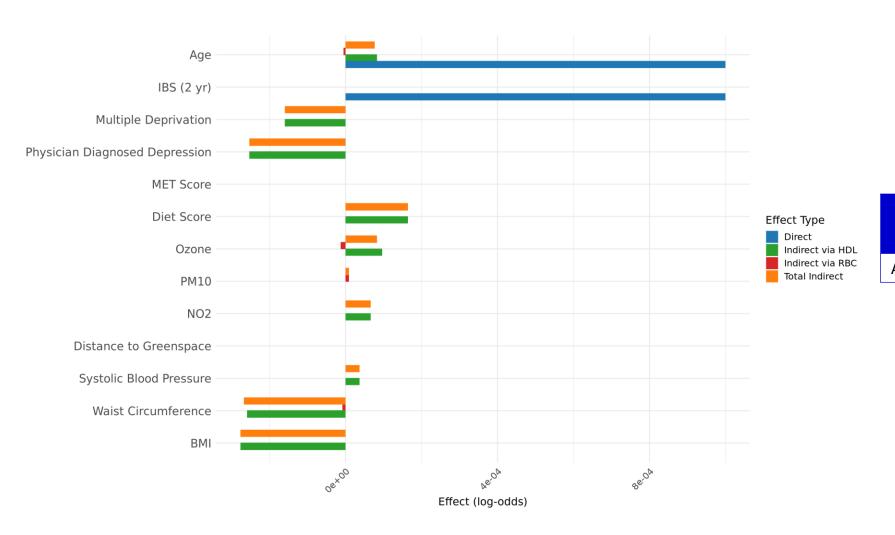


Variable	Proportion mediated effect (%)
Sex(Male)	16%
Ethnicity (Other)	2%
Ethnicity (Black)	13%
Ethnicity (Asian)	10%
Body Type : Plumper	1%
Body Type : Thinner	2%
Employment (Unemployed)	1%
Employment (Unable to work)	1%
Employment (Retired)	1%
Employment (Other)	1%
Employment (Home duties)	1%
Employment (Other qualification)	12%
Education (Vocational)	1%
Education (A-level)	1%
Education (CSEs)	0.5%
Education (GCSEs)	2%
Education (None)	12%

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Modelling approach

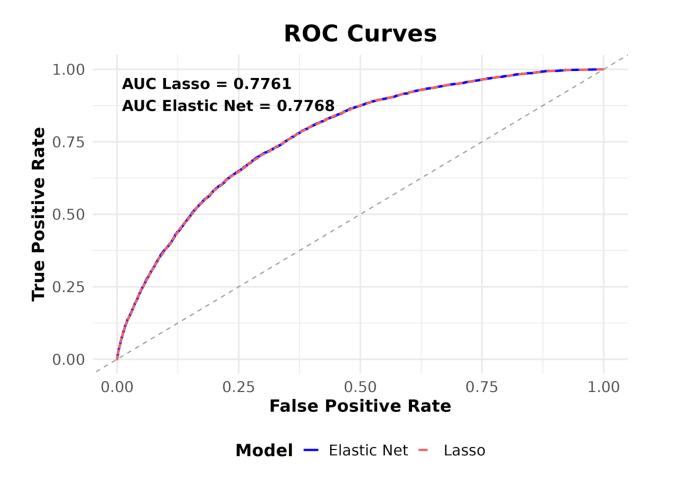
Summary of direct and indirect (mediated) effects – Continuous variables



	Variable	Proportion mediated effect (%)
Age		7%

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Prediction Results



- Out of all 25 variables, both models selected 22
- Lasso excluded "Distance to greenspace" and "Smoking status"
- Elastic Net excluded "Social isolation" and "IBS in 2 years"
- Both models excluded "Nucleated RBC Count"

Model	Accuracy	Sensitivity	Specificity
Lasso	0.68	0.10	0.98
Elastic net	0.68	0.10	0.98

Discussion

Conclusions

Indirect and Direct Effect

- Direct effects of exposome on CKD were found to be stronger than effects mediated HDL and nucleated RBC count
 - Effects might be mediated through other, potentially unmeasured, biological pathways.

HDL Cholesterol

- HDL cholesterol was found to be positively associated with CKD:
 - May reflect underlying metabolic dysfunction or HDL particle impairment⁶
 - Its effect on CKD could be mediated/confounded unmeasured factors affecting HDL
- BMI, waist circumference, and alcohol consumption show significant effect via HDL
 - Affect lipid metabolism → influence kidney health

nRBC and Prediction

- nRBC emerged to be a weak mediator → also weak predictor (not selected by LASSO & EINet)
- Predictive models are not good at identifying true cases (low sensitivity) → limits clinical use

Study Limitation

Exposure and biomarkers measurement

- Behavioural factors were measured by self-reported questionnaires
- Exposures and biomarkers were measured once at baseline
- Potential temporal mismatch between exposures and mediators

Outcome measurement

- HES captured hospitalised and death registery cases only
- CKD is often silent until late stage
- Risk of reverse causality

UK Biobank cohort

 Elder and pre-dominantly white population of UK Biobank limits the generalisability of our findings to diverse populations

Future Considerations

Enhance exposure/biomarkers measurement

- Collect exposures and biomarkers at multiple time point during the follow-up
- Expand biomarker panels (e.g KIM-1)

Subgroup analysis

- Subgroup analysis stratified by sex, ethnicity, etc
- Inform public health policy making

External validation

Validate signatures in diverse populations

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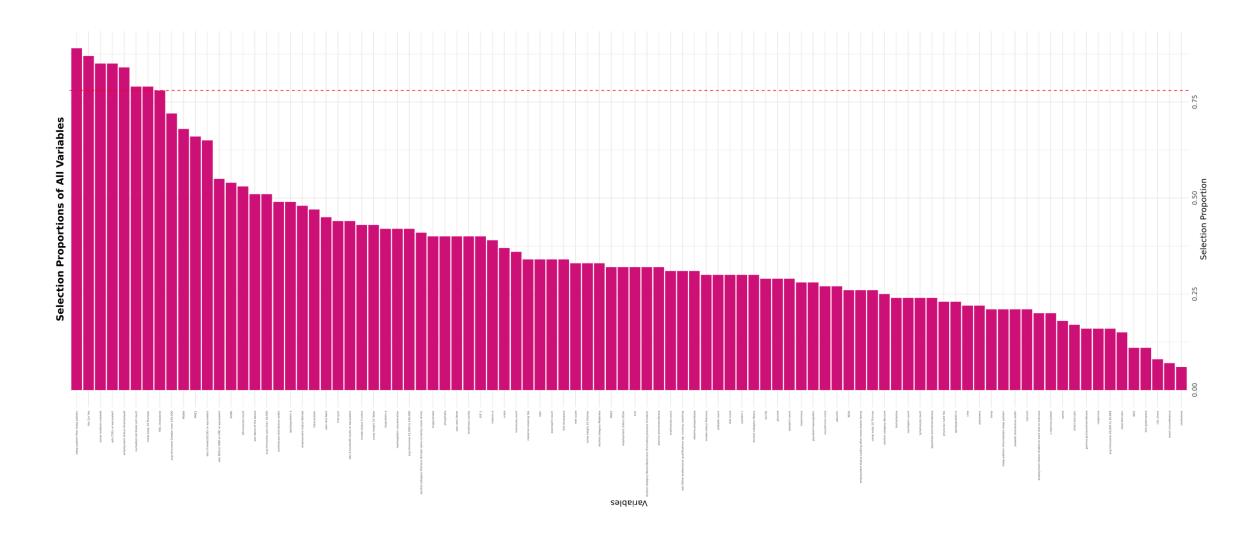
Thank you Questions welcomed

References

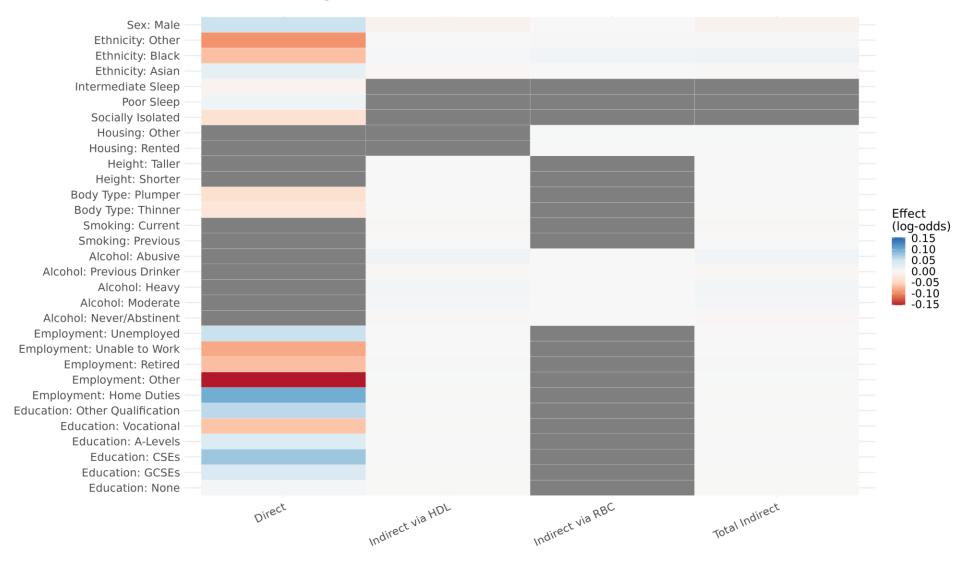
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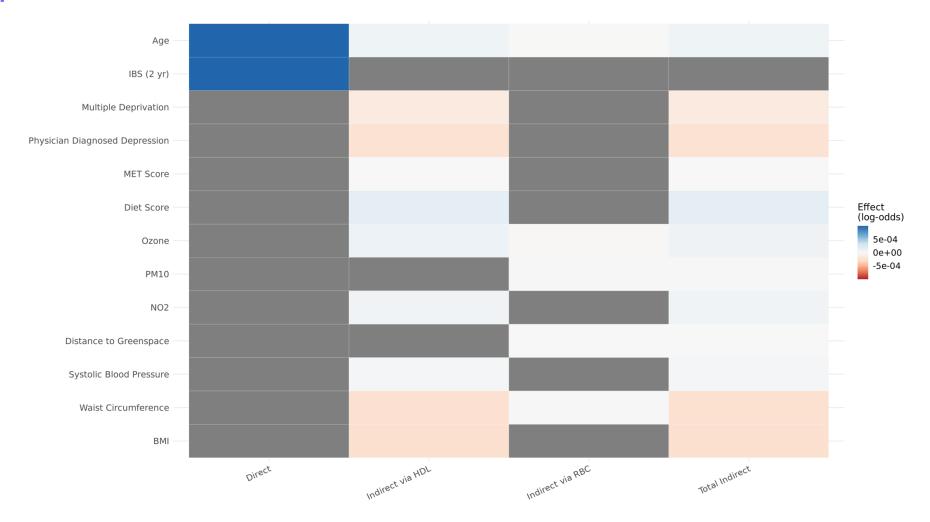
Selection Proportion of All Variables (from Direct Model)



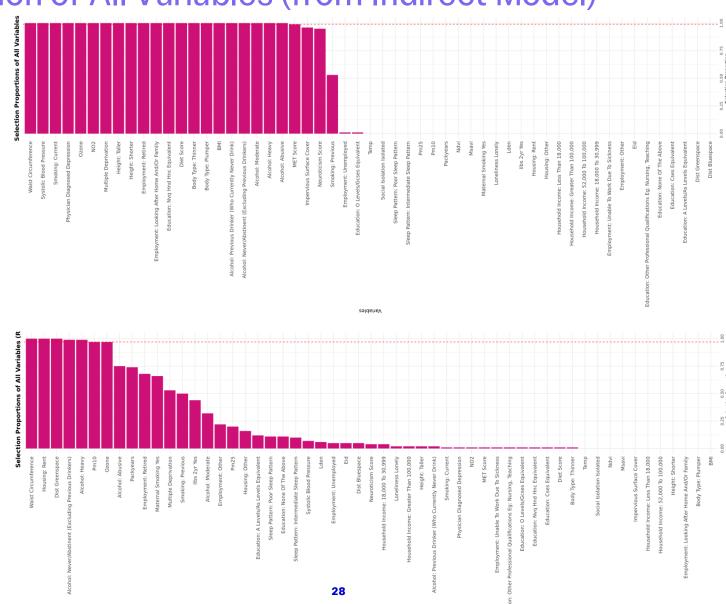
Heatmap of the effects - categorical



Heatmap of the effects - continous



Selection Proportion of All Variables (from Indirect Model)



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Confusion Matrix (lasso & elastic net)

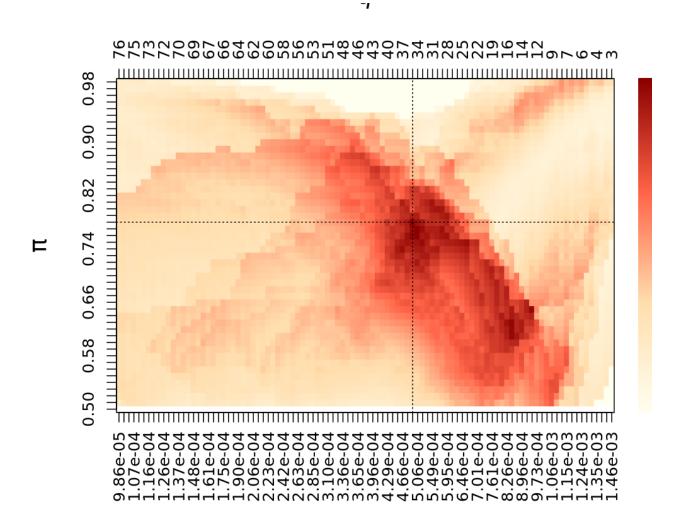
Lasso:

	Actual 0	Actual 1
Prediction 0	60610	1235
Prediction 1	28621	3184

Elastic net:

	Actual 0	Actual 1
Prediction 0	60453	1216
Prediction 1	28678	3303

Stability Selection plot - direct effect



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Stability Selection plot – indirect effect

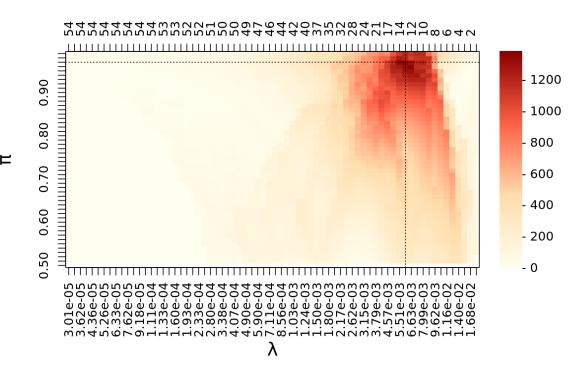


q

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nRBC

q



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