

# REPLICATED BLOOD-BASED BIOMARKERS FOR MYALGIC ENCEPHALOMYELITIS NOT EXPLICABLE BY INACTIVITY



Sjoerd Viktor Beentjes<sup>1,2</sup>, Artur Miralles Méharon<sup>1</sup>, Julia Kaczmarczyk<sup>1</sup>, Amanda Cassar<sup>1</sup>, Gemma Louise Samms<sup>1</sup>, Nima S Hejazi<sup>3</sup>, Ava Khamseh<sup>1,2</sup>, Chris P Ponting<sup>1</sup>

<sup>1</sup>University of Edinburgh, <sup>2</sup>University of California, Berkeley, <sup>3</sup>Harvard T.H. Chan School of Public Health

AI4Bi  
UKRI AI Centre for Doctoral Training  
in Biomedical Innovation  
THE UNIVERSITY  
of EDINBURGH

UK  
RI  
UK Research  
and Innovation

## Background & Motivation

### What is Myalgic Encephalomyelitis / Chronic Fatigue Syndrome (ME/CFS)?

- Debilitating multi-system disease affecting  $\approx 0.6\%$  of the UK population (predominantly women)
- Key symptom: post-exertional malaise (worsening after activity)
- No diagnostic biomarker**; biology poorly understood
- No cure and no widely effective therapy

### The Problem: Confounding by Activity

- ME/CFS  $\rightarrow$  reduced physical activity
- Many biomarkers are activity-dependent
- Key question:** Are biomarker differences disease-driven, or explained indirectly through reduced activity?

## Data

### UK Biobank (UKB), UK

- 1,455 ME/CFS cases and 131,303 controls
- Blood traits (63), NMR metabolites (251), Proteins (2923)
- Activity measures: walking duration, days of moderate activity, minutes of moderate activity

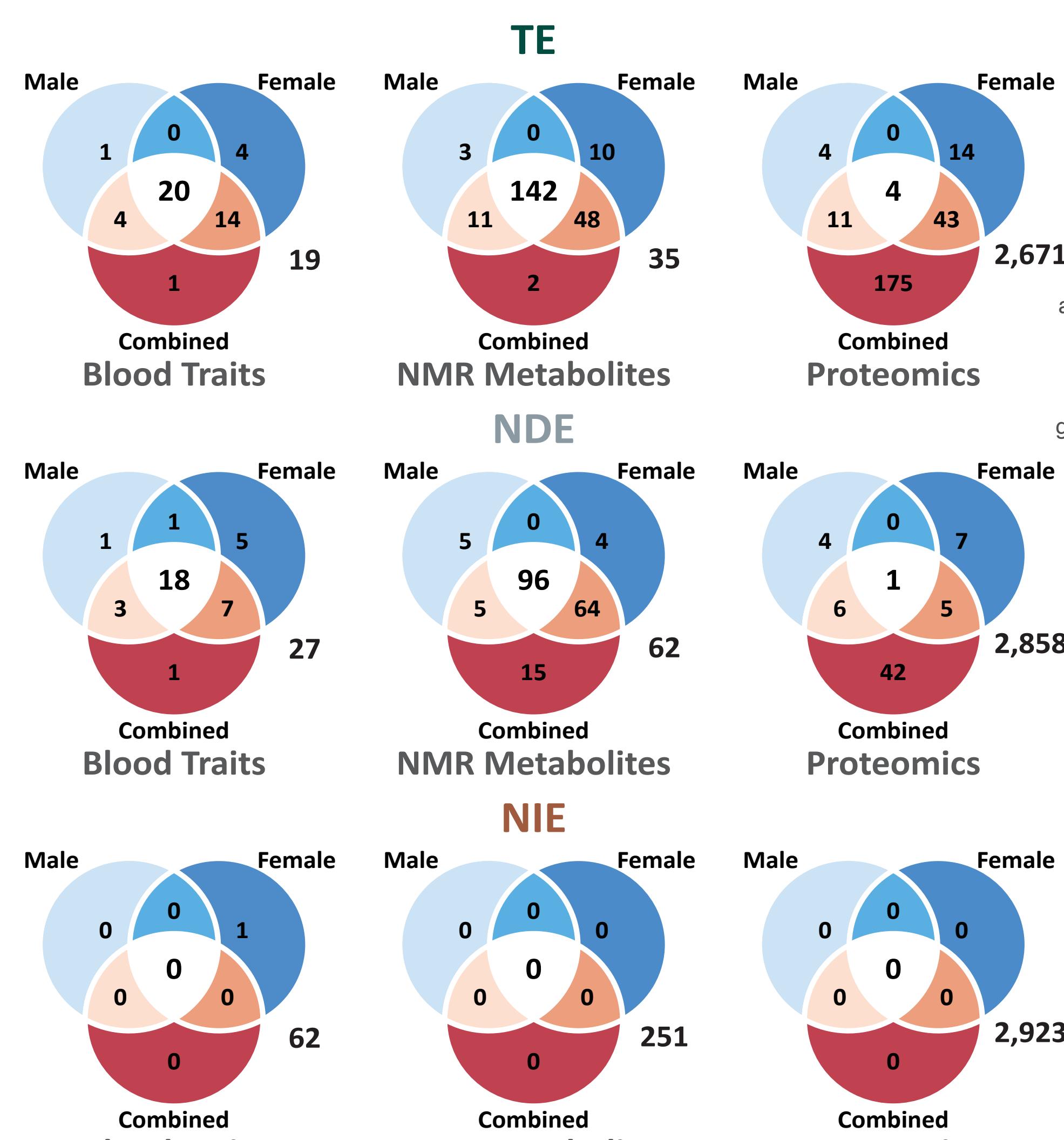
### PEM subgroup (UK Biobank)

- Defined using the UKB Pain Questionnaire (fatigue 6 months, post-exertional worsening, not relieved by rest).
- 297 identified; 239 used after matching to non-PEM ME/CFS cases.

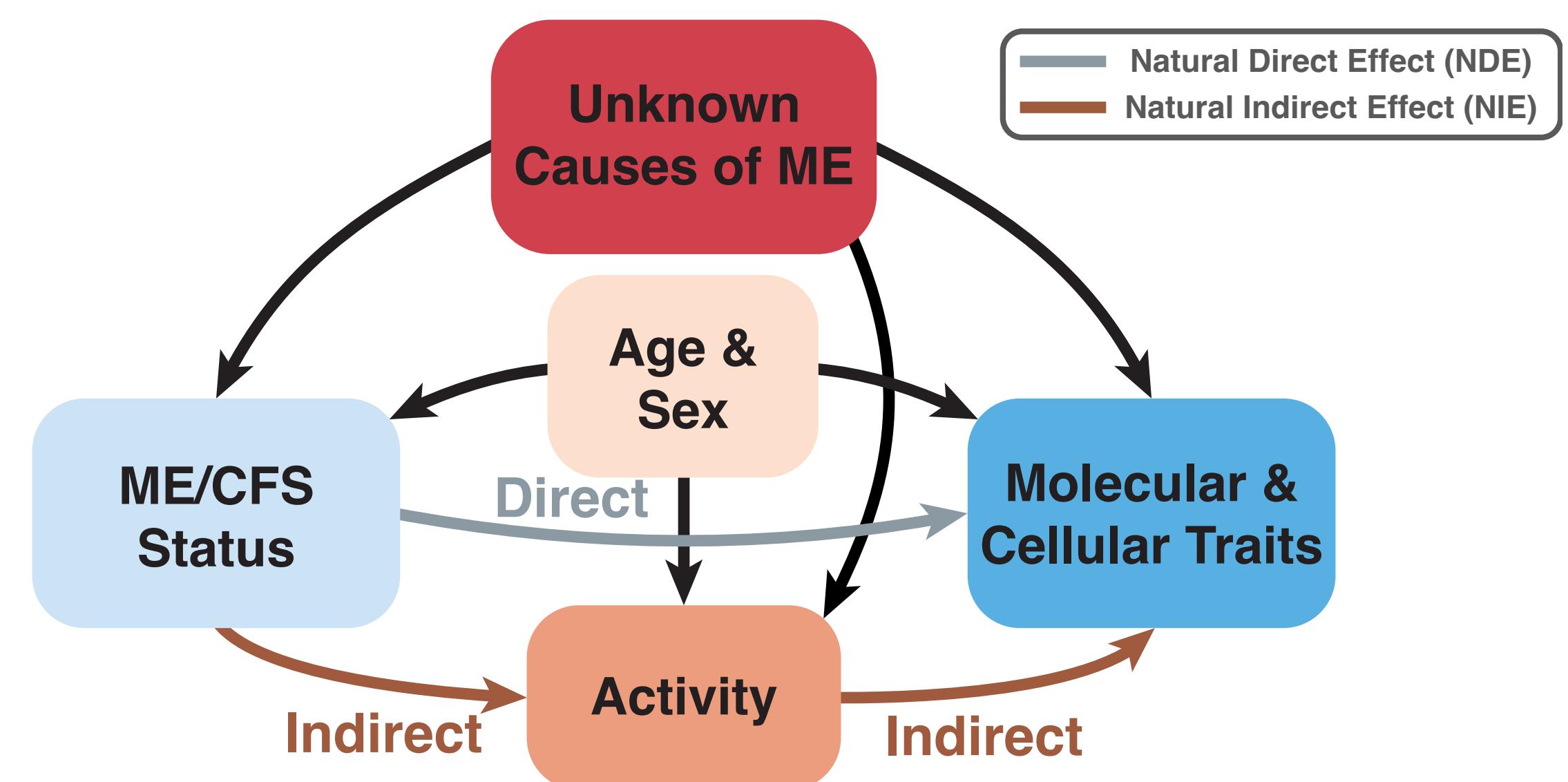
### All of Us (AoU), USA (Replication)

- 903 cases and 75,943 controls
- 14 overlapping blood traits

## Results



## Methods



Note: Full causal identifiability is not guaranteed; mediation effects are interpreted as statistical quantities adjusted for age and sex.

### Semi-parametric mediation analysis

We used ensemble machine learning (Super Learner) to flexibly estimate all nuisance functions required for the one-step estimation of:

- Total Effect (TE)
- Natural Direct Effect (NDE)
- Natural Indirect Effect (NIE)

$$\underbrace{\mathbb{E}[Y(1)] - \mathbb{E}[Y(0)]}_{\text{TE}} = \underbrace{\mathbb{E}[Y(1, M(0))] - \mathbb{E}[Y(0, M(0))]}_{\text{NDE}} + \underbrace{\mathbb{E}[Y(1, M(1))] - \mathbb{E}[Y(1, M(0))]}_{\text{NIE}}$$

### Super Learner Library

- Earth (MARS), GLMnet (LASSO)
- GLM with interactions
- XGBoost/LightGBM
- Highly Adaptive Lasso (HAL)

### Software (R packages)

- TE: npcausal (Kennedy, 2021) + SuperLearner (Polley et al, 2011)
- NDE & NIE: medoutcon (Hejazi et al, 2022) + s13 (Coyle et al, 2021).

## Key findings

- 511 biomarkers** differ between ME/CFS cases and controls across blood traits, NMR metabolites, and proteomics.
- Effects are **consistent across males and females**: 166 biomarkers are significant in both sexes.
- Mediation via physical activity is negligible**: almost no NIEs are significant despite strong TEs.
- External replication** in All of Us: 9 of 14 blood traits tested show consistent effects (only urea differs).
- Sensitivity analysis**: individuals with PEM-like symptoms show stronger biomarker shifts.

### References

I Diaz, N S Hejazi, K E Rudolph, and M J van Der Laan. Nonparametric efficient causal mediation with intermediate confounders. *Biometrika*, 108(3):627–641, August 2021.  
Nima Hejazi, Kara Rudolph, and Iván Díaz. medoutcon: Nonparametric efficient causal mediation analysis with machine learning in R. *J. Open Source Softw.*, 7(69):3979, January 2022.  
Edward H Kennedy. Nonparametric causal effects based on incremental propensity score interventions. *J. Am. Stat. Assoc.*, 114(526):645–656, April 2019.  
Mark J. Van Der Laan and Sherri Rose. *Targeted Learning: Causal Inference for Observational and Experimental Data*. Springer Series in Statistics. Springer, New York, NY, 2011. ISBN 978-1-4419-9781-4. doi: 10.1007/978-1-4419-9782-1.  
Mark J van der Laan, Eric C Polley, and Alan E Hubbard. Super learner. *Stat. Appl. Genet. Mol. Biol.*, 6(1), January 2007.