

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



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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

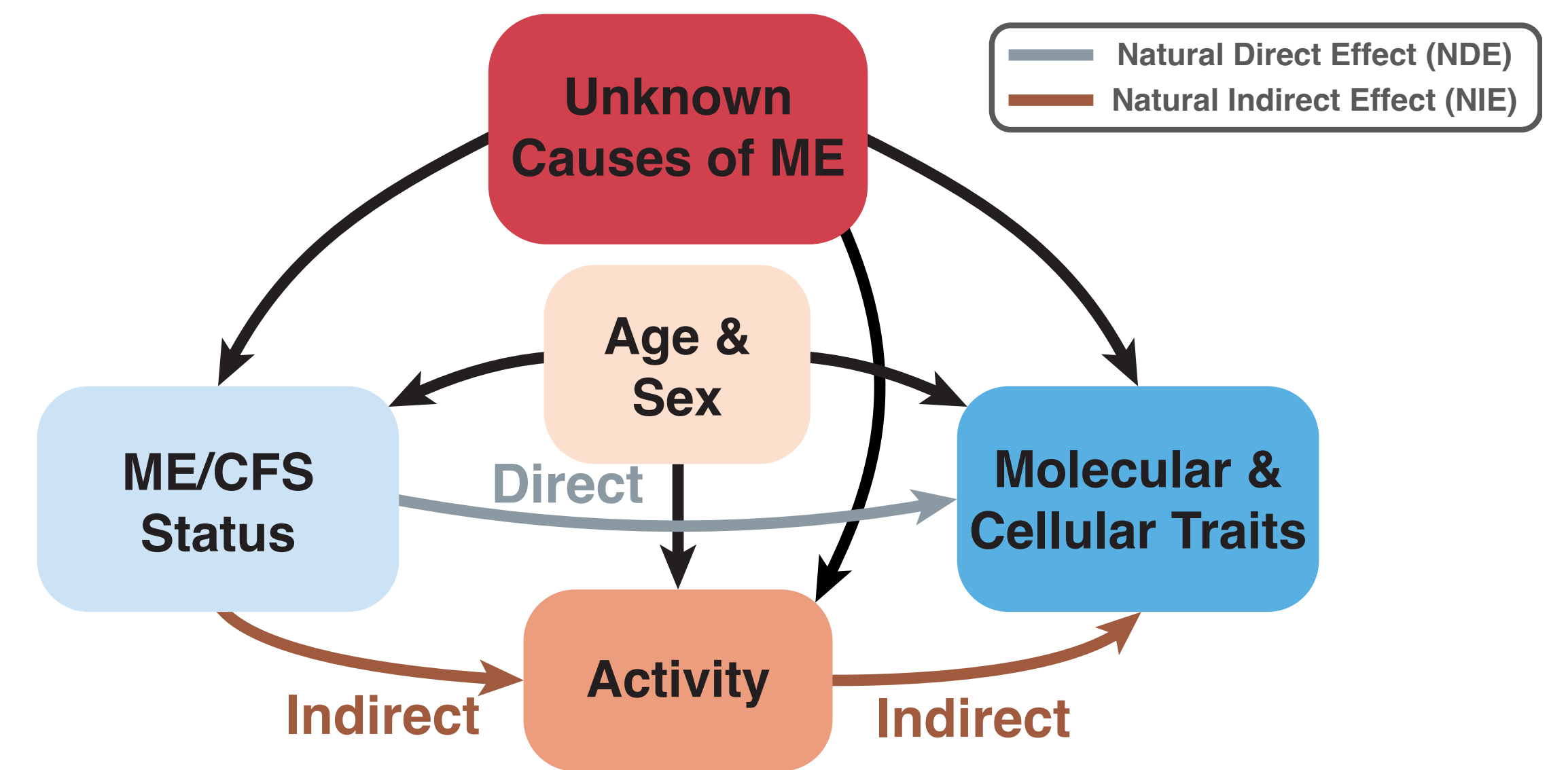
PEM subgroup (UK Biobank)

- 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

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)

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

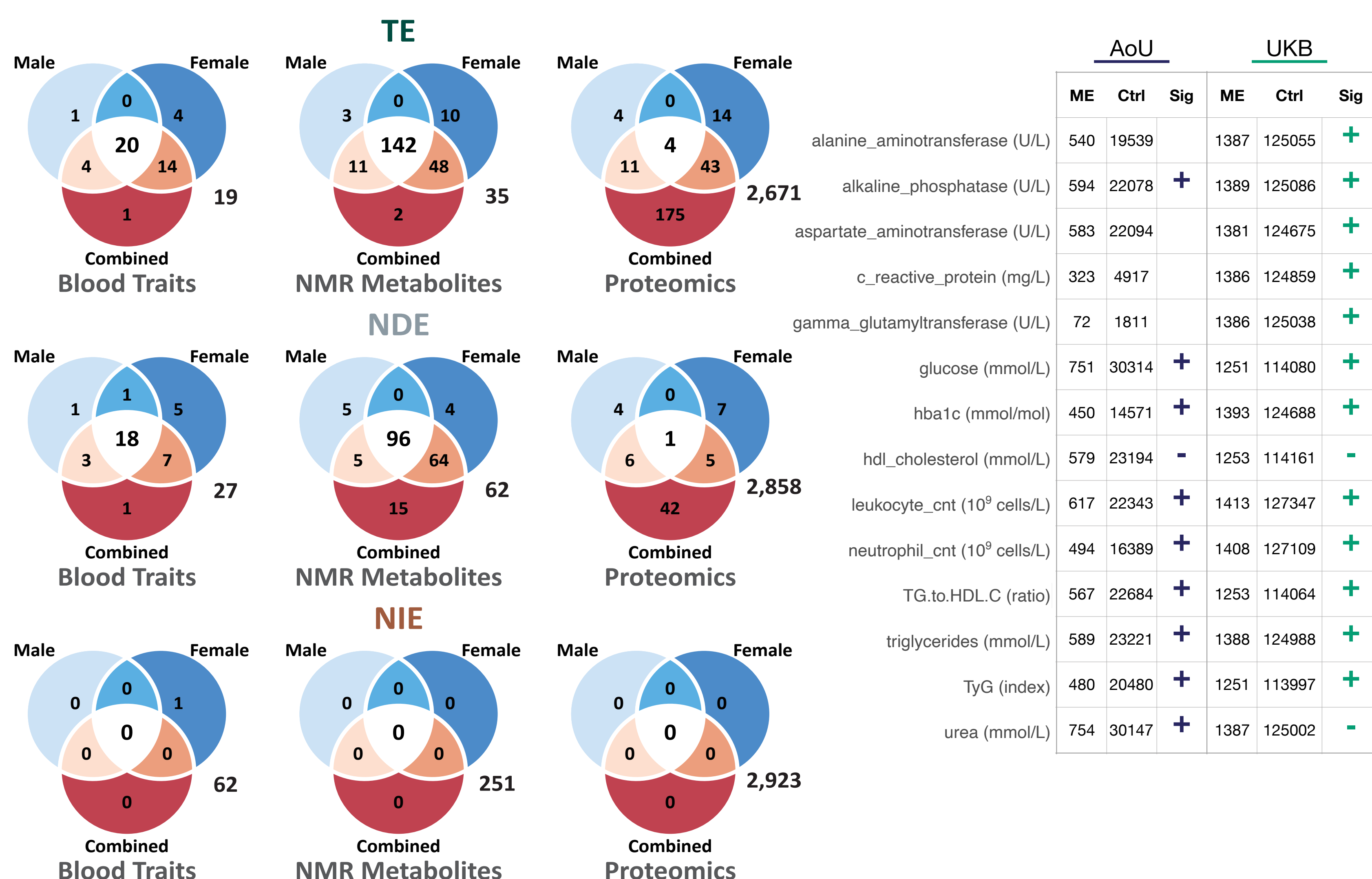
Super Learner Library

- 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).

Results



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

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