

**Early diagnosis of Dementia prediction
using accessible exposome-based features:
A comparison of Statistical and Machine
Learning models on the UK Biobank**

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What are we doing?

- We are using UK-Biobank data (~500,000 individuals).
- To create a predictive test for Dementia affordable and exposome-based.
- We used different algorithms.

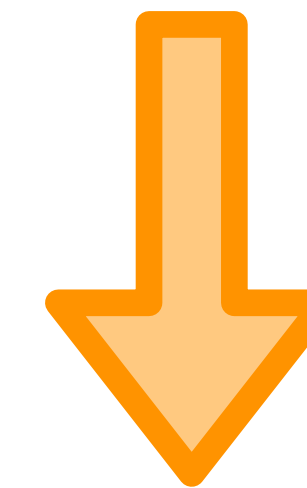




Why a Dementia predictor?



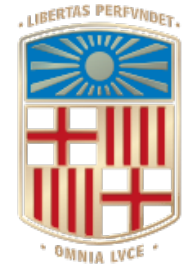
Dementia is a syndrome without cure, and with a challenging diagnosis.



That's why currently there is more focus on:

Risk reduction, early intervention and timely diagnosis.

1. Geldmacher, David S., and Peter J. Whitehouse. "Evaluation of dementia." *New England Journal of Medicine* 335.5 (1996): 330-336.
2. (February 28, 2022) Dementia statistics. Alzheimer's Disease International. <https://www.alzint.org/about/dementia-facts-figures/dementia-statistics/>
3. Breitner, John CS. "Dementia—epidemiological considerations, nomenclature, and a tacit consensus definition." *Journal of geriatric psychiatry and neurology* 19.3 (2006): 129-136.
4. (February 28, 2022) Treatment. NHS. <https://www.nhs.uk/conditions/alzheimers-disease/treatment/>
5. (February 28, 2022) Treatments for dementia. Alzheimer's Society. <https://www.alzheimers.org.uk/about-dementia/treatments>



Why an Early predictor?



In clinical practice, Dementia is diagnosed at late stages when symptoms become highly pronounced.

~18 years before a diagnosis



Additionally...



FOR ECONOMIC REASONS

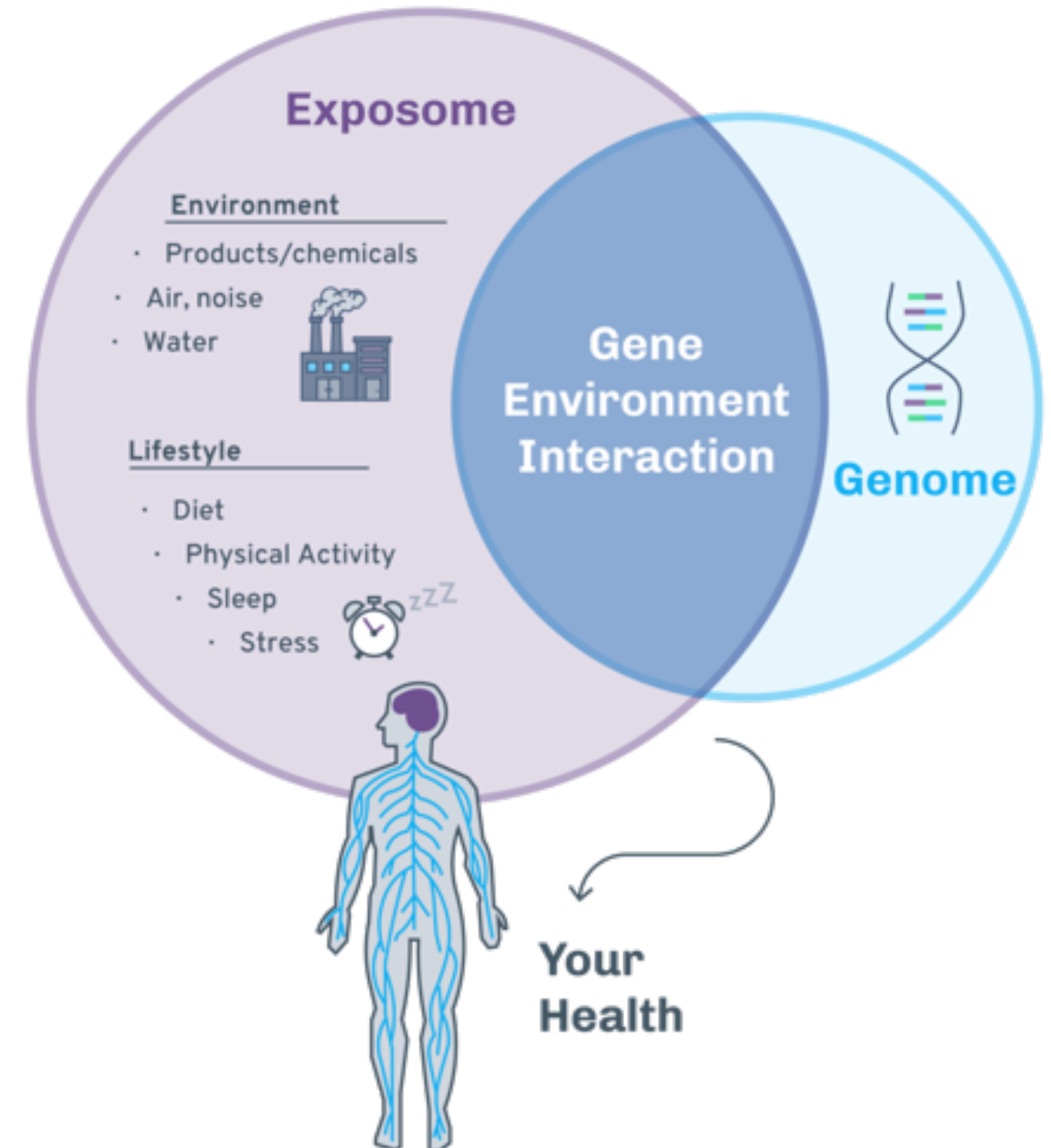
- Dementia costs in the UK:
- ~31 billion euros per year
 - Diagnosis: Several test and brain images over a period of time



** Additionally, our method could be applied in other cohorts and for other disorders, which will be extremely interesting for low-income countries or those having a private health care system.*

Why exposome-based?

<< Unlike genetic factors which are stable and unmodifiable, the exposome has large spatiotemporal variability and can be modified at different levels. >>





Papers about: UK Biobank + Exposome + Dementia

- Sleep, major depressive disorder, and Alzheimer disease
- Meat consumption and risk of incident dementia: cohort study of 493,888 UK Biobank participants
- Is neuroticism differentially associated with risk of Alzheimer's disease, vascular dementia, and frontotemporal
- Diet and Dementia: A Prospective Study
- Identifying dementia outcomes in UK Biobank: a validation study of primary care, hospital admissions and mortality data
- Associations between vascular risk factors and brain MRI indices in UK Biobank
- High coffee consumption, brain volume and risk of dementia and stroke
- Sex differences in the association between major cardiovascular risk factors in midlife and dementia: a cohort study using data from the UK Biobank

1. *Development and validation of a predictive algorithm for risk of dementia in the community setting (CANADA)*
2. *Machine learning prediction of incidence of Alzheimer's disease using large-scale administrative health data (KOREA)*

Previous study

External validation of four dementia prediction models for use in the general community-dwelling population: a comparative analysis from the Rotterdam Study

- All models showed similar discriminative ability when compared to prediction based on age alone. These findings highlight the urgent need for updated or new models to predict dementia risk in the general population.

Prediction model	C-statistics at various follow-up horizons (95% CI)			
	2 years n/N = 63/6667	5 years n/N = 233/6667	10 years n/N = 515/6667	15 years n/N = 847/6667
CAIDE	0.49 (0.42–0.56)	0.54 (0.50–0.58)	0.55 (0.53–0.58)	0.55 (0.53–0.57)
Age only	NA	NA	NA	NA
Without age	0.49 (0.42–0.56)	0.54 (0.50–0.58)	0.55 (0.53–0.58)	0.55 (0.53–0.57)
BDSI	0.83 (0.75–0.90)	0.80 (0.76–0.84)	0.78 (0.76–0.81)	0.76 (0.74–0.78)
Age only	0.83 (0.76–0.90)	0.81 (0.78–0.85)	0.81 (0.78–0.83)	0.79 (0.77–0.81)
Without age	0.64 (0.57–0.71)	0.63 (0.59–0.66)	0.60 (0.58–0.63)	0.59 (0.57–0.61)
ANU-ADRI	0.81 (0.77–0.86)	0.78 (0.76–0.81)	0.75 (0.74–0.77)	0.70 (0.69–0.72)
Age only	0.83 (0.79–0.87)	0.80 (0.77–0.82)	0.77 (0.75–0.79)	0.72 (0.71–0.74)
Without age	0.56 (0.49–0.64)	0.51 (0.47–0.55)	0.52 (0.49–0.54)	0.51 (0.49–0.53)
DRS	0.84 (0.77–0.92)	0.82 (0.78–0.86)	0.81 (0.78–0.83)	0.79 (0.77–0.81)
Age only	0.83 (0.76–0.90)	0.81 (0.78–0.85)	0.81 (0.78–0.83)	0.79 (0.77–0.81)
Without age	0.63 (0.56–0.70)	0.58 (0.54–0.62)	0.57 (0.55–0.60)	0.55 (0.53–0.57)

CI confidence interval, *n* number of cases, *N* number of people at risk, *CAIDE* cardiovascular risk factors, aging, and dementia study, *NA* not applicable, *BDSI* brief dementia screening indicator, *ANU-ADRI* Australian National University Alzheimer’s Disease Risk Index and, *DRS* dementia risk score



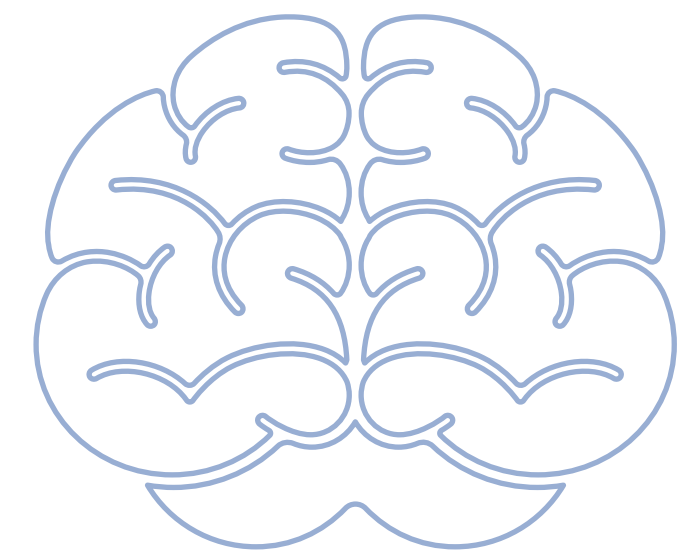
Our study



Early diagnosis of Dementia prediction using accessible exposome-based features: A comparison of Statistical and Machine Learning models on the UK Biobank

Objectives:

1. Demonstrate that accessible exposome variables are powerful enough to predict Dementia.
2. Demonstrate that ML outperforms SL in our classification problem.
3. Demonstrate that our models work well in external validation.
4. Demonstrate that we are not predicting just age.



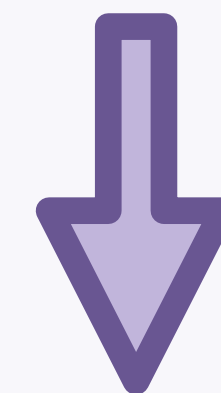


Dementia ICD-10



1. Alzheimer (AD)
2. Vascular Dementia (VD)
3. Frontotemporal dementia (FTD)
4. Other causes of dementia (OD)

Rejected cases with a Dementia diagnosis before 2011, hence before or during Baseline assessment.



1,523 individuals



Overview



Sample to Build the models and Perform Internal Validation **90%**

2.740

Dementia Group after 2011 **1370**

Healthy Control Group **1370**

Assessment Center Locations:

19/22 (All except: Edinburgh, Oxford, Barts)

Mean Birth: 1944.025

Treatment —> **1943.918**

Control —> **1944.132**

For Fairness Purposes:

Treatment —> Female = **623** Male = **747**

Control —> Female = **747** Male = **623**



Sample to perform External Validation **10%**

306

Dementia Group after 2011 **153**

Healthy Control Group **153**

Assessment Center Locations:

Edinburgh = **108**

Oxford = **107**

Barts = **91**

Mean Birth: 1943.973

Treatment —> **1943.300**

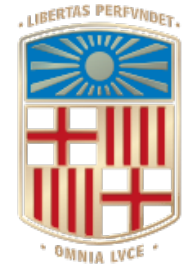
Control —> **1944.647**

For Fairness Purposes:

Treatment —> Female = **66** Male = **87**

Control —> Female = **87** Male = **66**





Exposome features



1. Physical measurements —> Weight
2. Sociodemographics —> Qualifications
3. Lifestyle —> Sleeplessness
4. Environmental factors —> Average evening noise
5. Early life factors —> Adopted as a child
6. Traumatic events —> Victim of sexual assault
7. Mental health —> Happiness

Exposome



Methods

1. Exposome data

502,664 patients from UK Biobank and 128 exposome features

2. Data pre-processing

Selection of Dementia patients, and 1,370 Healthy patients

80% missing value threshold for features leading to 78 exposome features

MissForest for Imputation

3. Predictive disease modeling

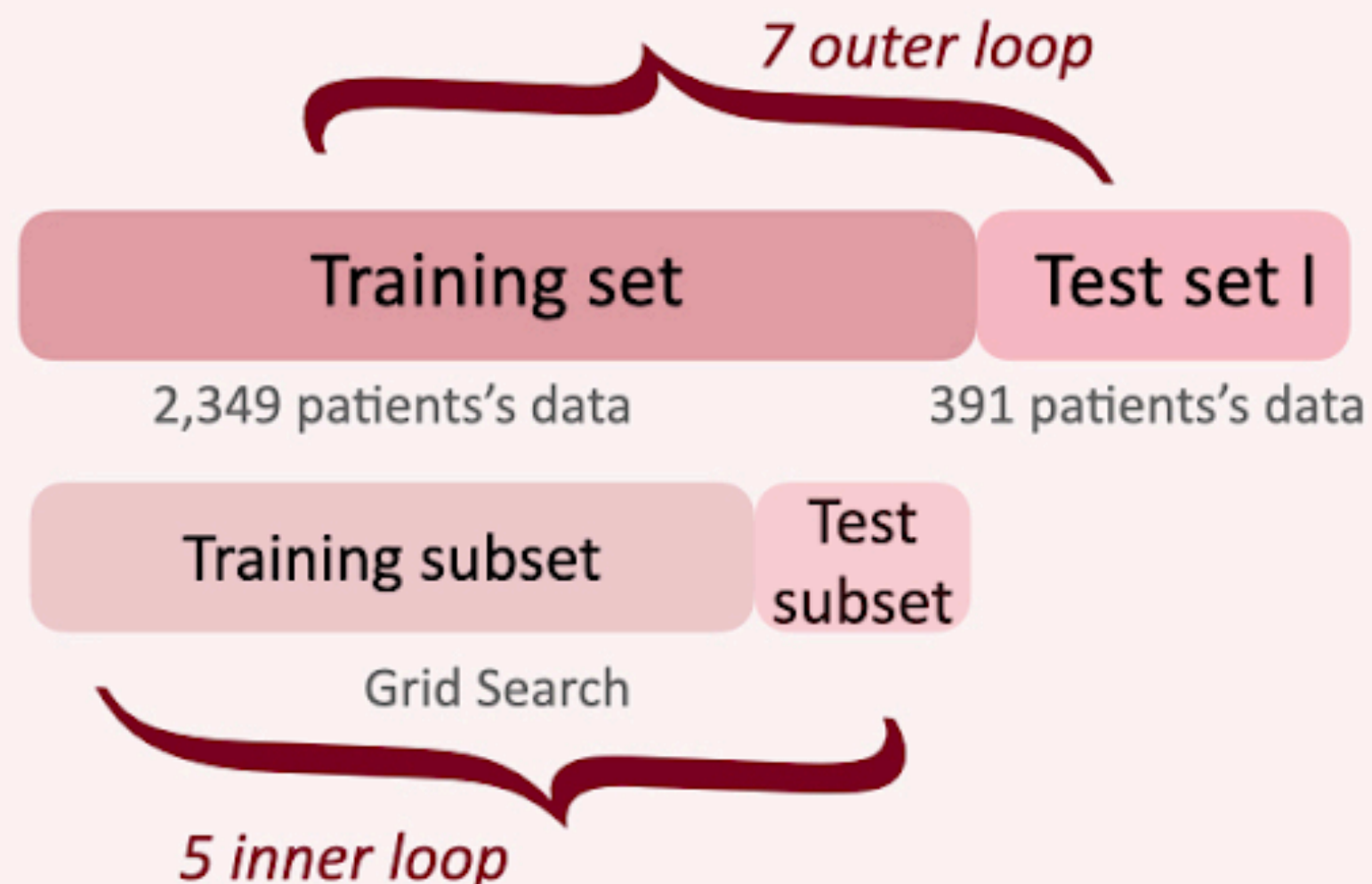
Logistic Regression (SL): linear based model

XGBoost (ML): decision-tree-based ensemble model

3.1. Internal Validation

2,740 patients's data obtained from 19 different assessment centers

Nested cross-validation



3.2. External Validation

306 patients's data obtained from 3 different assessment centers: Oxford, Edinburgh and Barts.

Test set II

4. Model Interpretability

SL → Coefficients

ML → Gini Importance

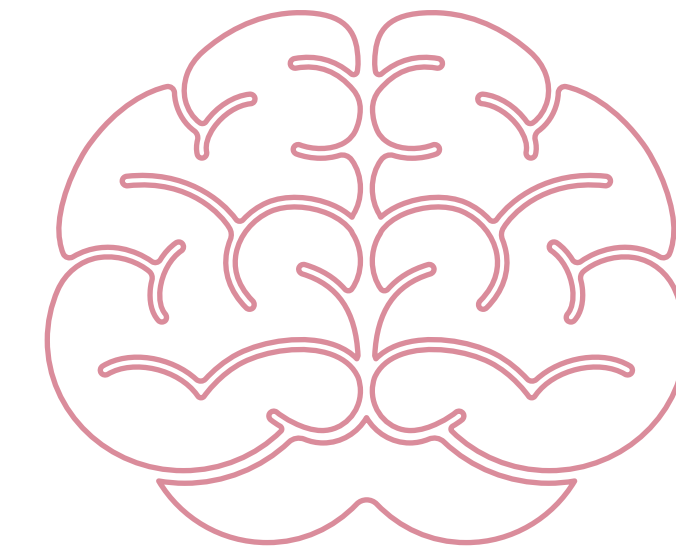
Flowchart for early prediction of Dementia using exposures data in UK Biobank



Experiments



- **Experiment 1:** Exposome with Age
- **Experiment 2:** Exposome without Age
- **Experiment 3:** Age
- **Experiment 4:** Accessible with Age (30)
- **Experiment 5:** Accessible without Age (30)



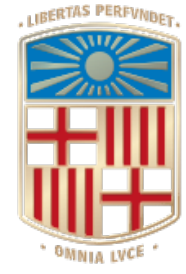


AUC can be interpreted as:

- 1) No discrimination [$AUC = 0.5$]
- 2) Poor discrimination [$0.6 \geq AUC > 0.5$]
- 3) Acceptable discrimination [$0.7 \geq AUC > 0.6$]
- 4) Excellent discrimination [$0.8 \geq AUC > 0.7$]
- 5) Outstanding discrimination [$AUC > 0.9$]

Our Goal:

Obtain an accessible model able to perform an Excellent or Outstanding discrimination.



Logistic Regression: Linear Model

	Exposome with Age	Exposome without Age	Age	Accessible with Age	Accessible without Age
AUC	0.75±0.02 0.77±0.01	0.75±0.01 0.76±0.01	0.58 ± 0.02 0.68±0.03	0.74±0.02 0.75±0.01	0.75±0.02 0.77±0.01
F1	0.74±0.02 0.73±0.01	0.74±0.01 0.73±0.01	0.58 ± 0.03 0.68±0.01	0.73±0.01 0.72±0.01	0.74±0.01 0.75±0.01
precision	0.77±0.02 0.86±0.01	0.77±0.02 0.85±0.01	0.59 ± 0.03 0.68±0.04	0.76±0.03 0.84±0.01	0.78±0.03 0.84±0.01
sensitivity	0.75±0.02 0.64±0.01	0.72±0.02 0.64±0.01	0.59 ± 0.07 0.69±0.02	0.71±0.02 0.63±0.01	0.71±0.02 0.67±0.02



Machine Learning Results

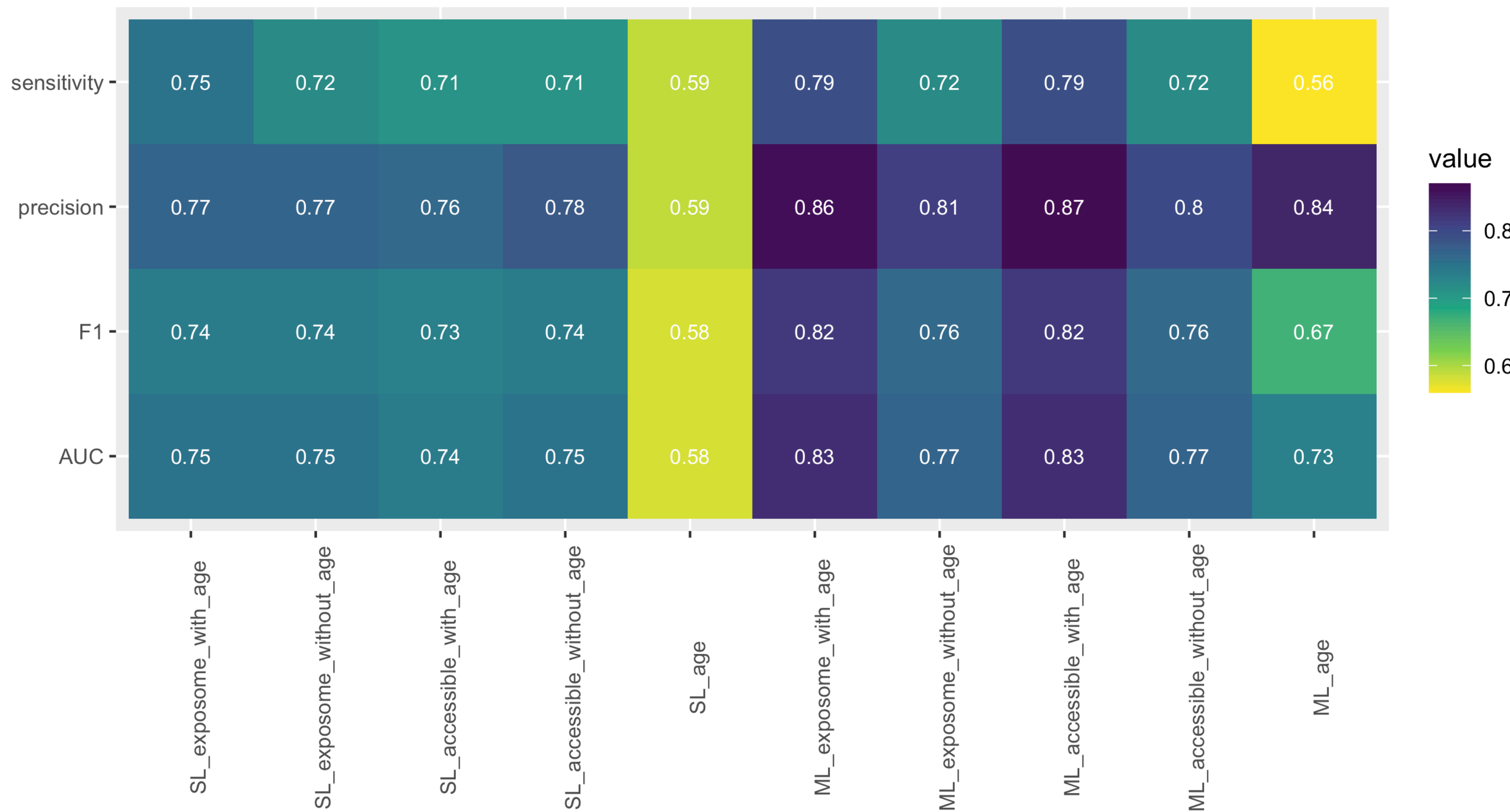


XGBoost: Non-Linear Model

	Exposome with Age	Exposome without Age	Age	Accessible with Age	Accessible without Age
AUC	0.83±0.02 0.89±0.01	0.77±0.01 0.79±0.01	0.73±0.02 0.84±0.00	0.83±0.02 0.88±0.01	0.77±0.01 0.78±0.02
F1	0.82±0.01 0.88±0.01	0.76±0.01 0.76±0.01	0.67±0.02 0.82±0.00	0.82±0.02 0.87±0.01	0.76±0.01 0.75±0.02
precision	0.86±0.02 0.93±0.01	0.81±0.03 0.87±0.02	0.84±0.01 0.96±0.00	0.87±0.04 0.94±0.01	0.80±0.02 0.86±0.03
sensitivity	0.79±0.01 0.83±0.01	0.72±0.02 0.68±0.03	0.56±0.03 0.71±0.00	0.79±0.02 0.81±0.02	0.72±0.02 0.67±0.02



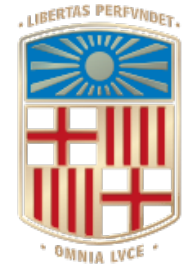
Internal Validation





External Validation



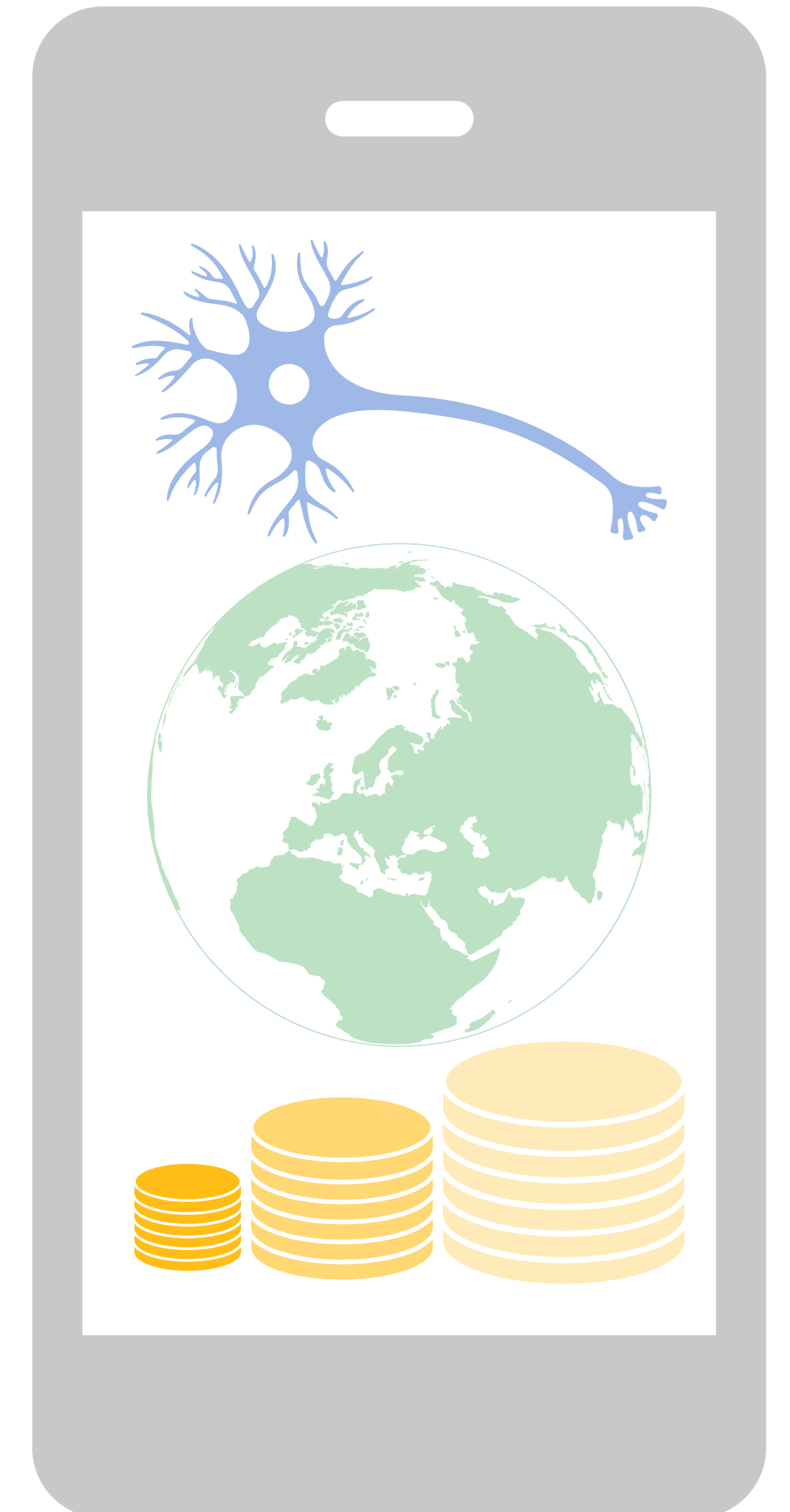


Accessible Models



Create new models with just 30 accessible variables.

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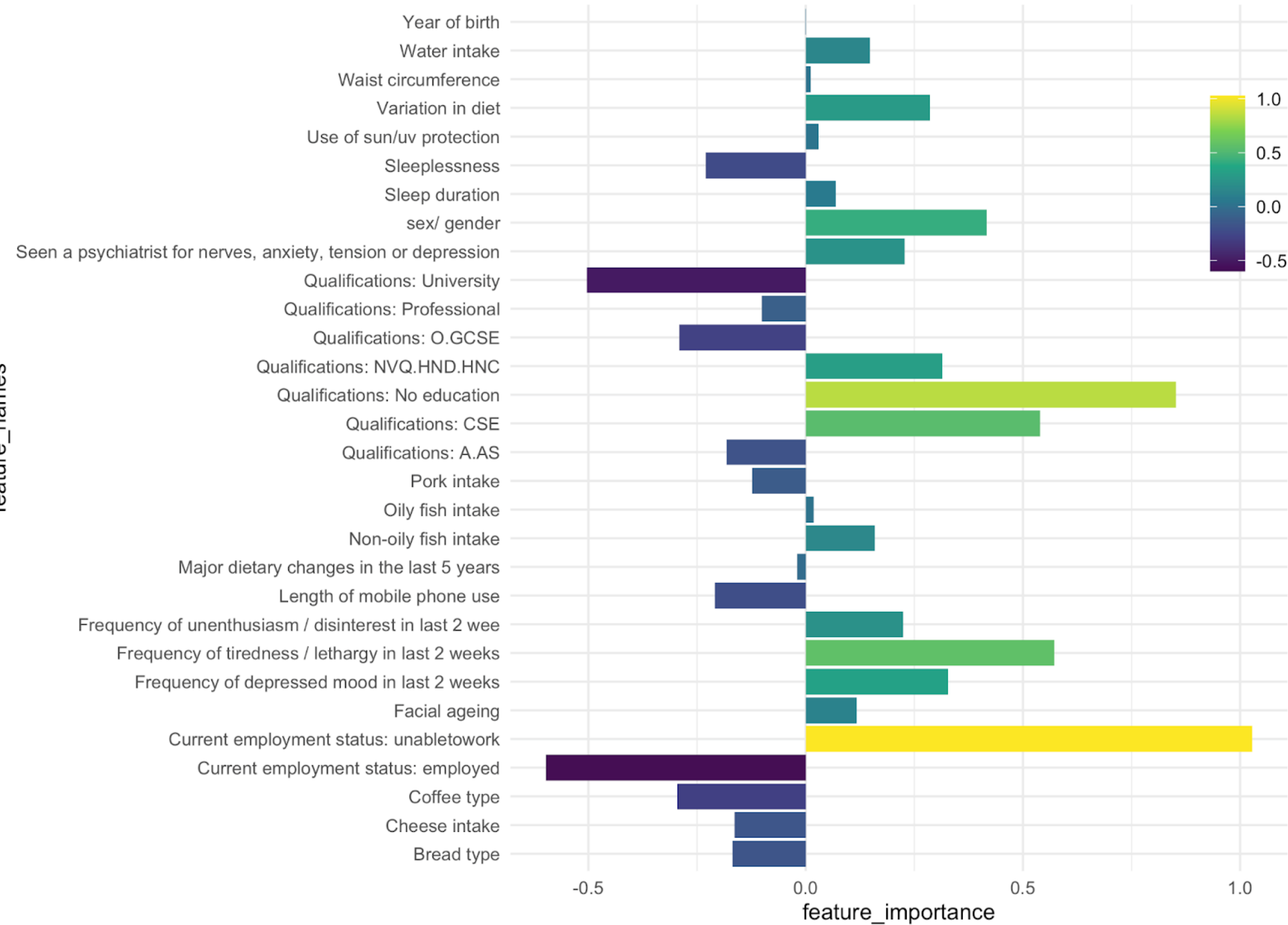




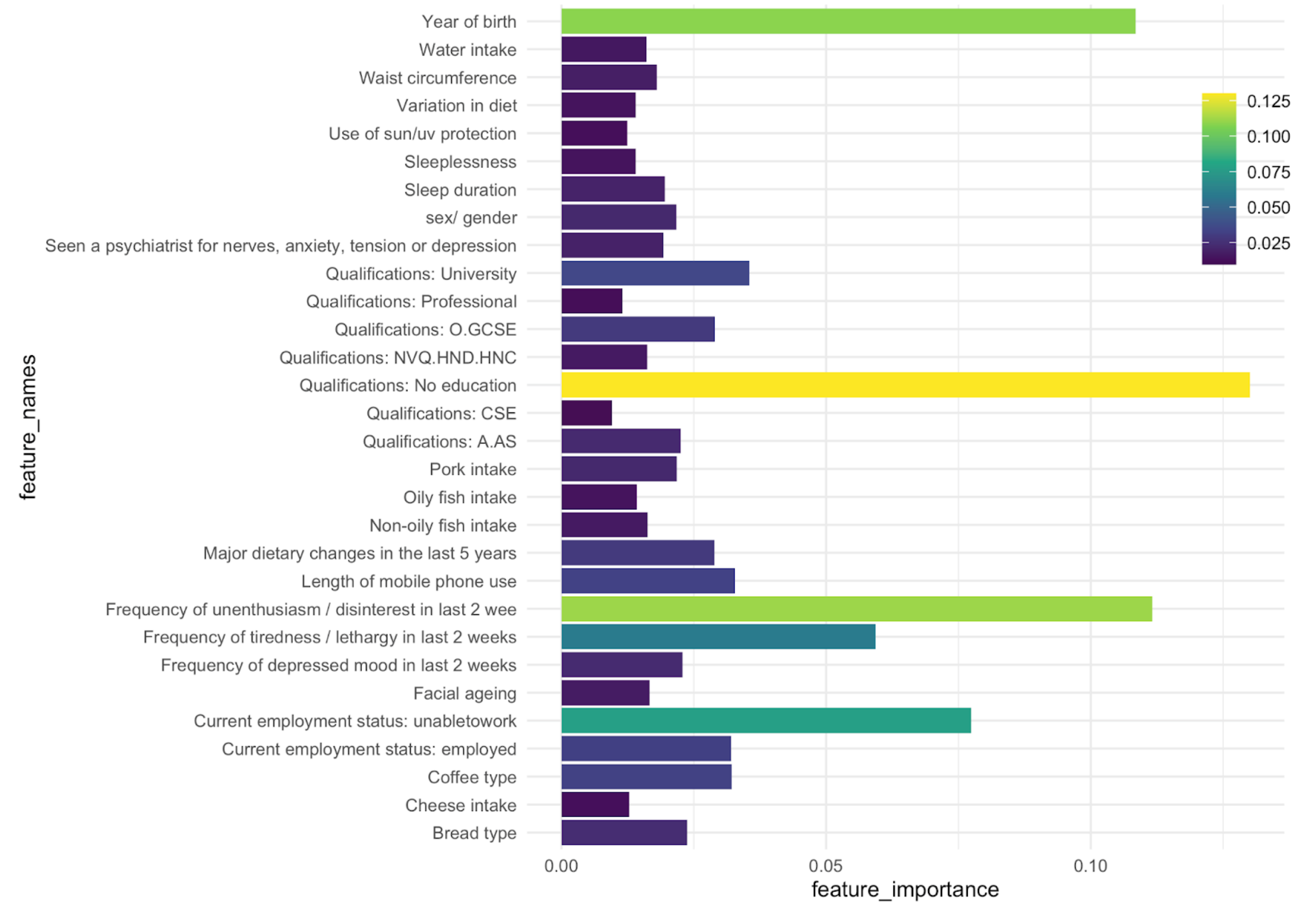
Accessible with Age



Accessible with age experiment: Statistical Learning



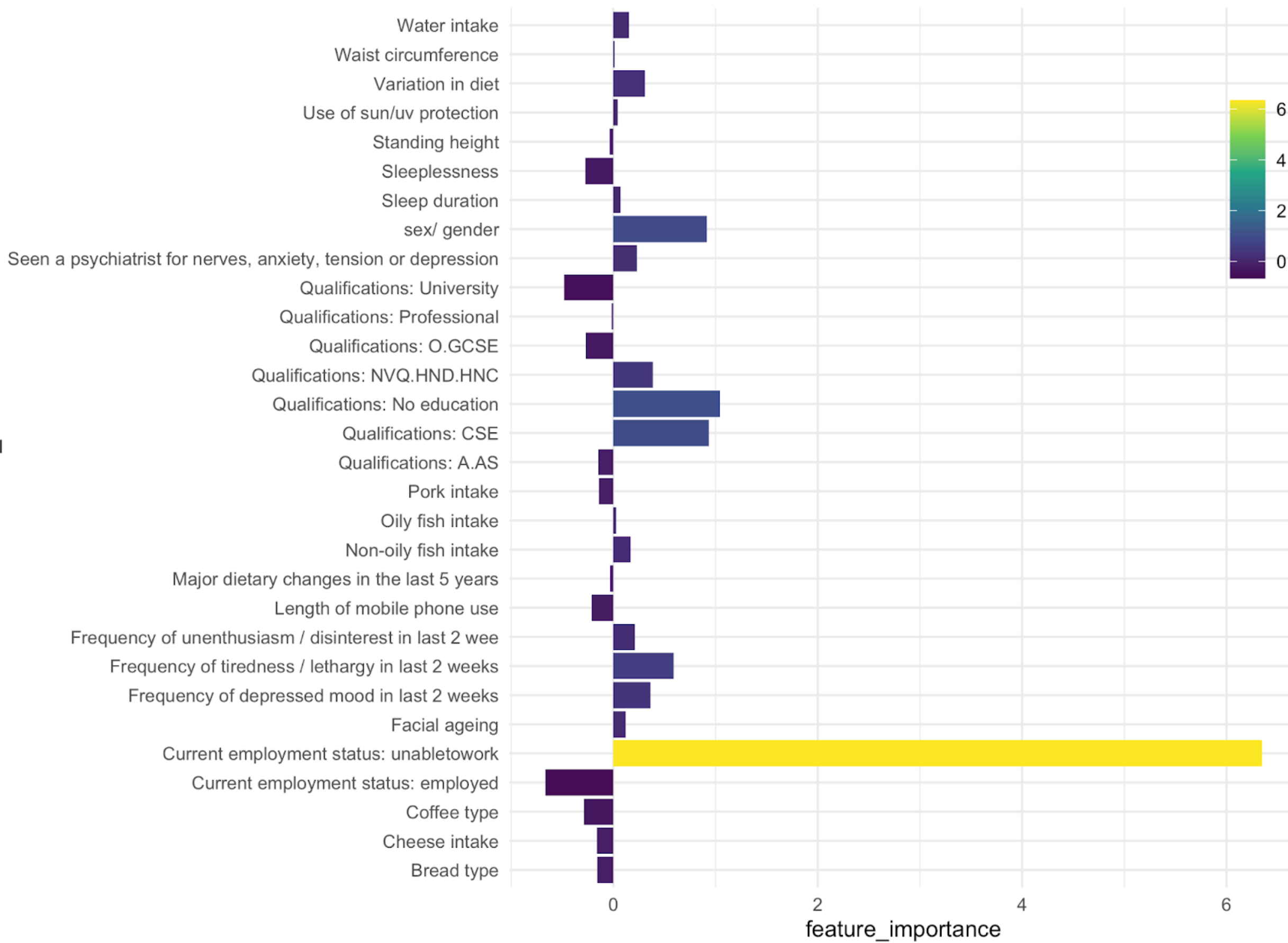
Accessible with age experiment: Machine Learning



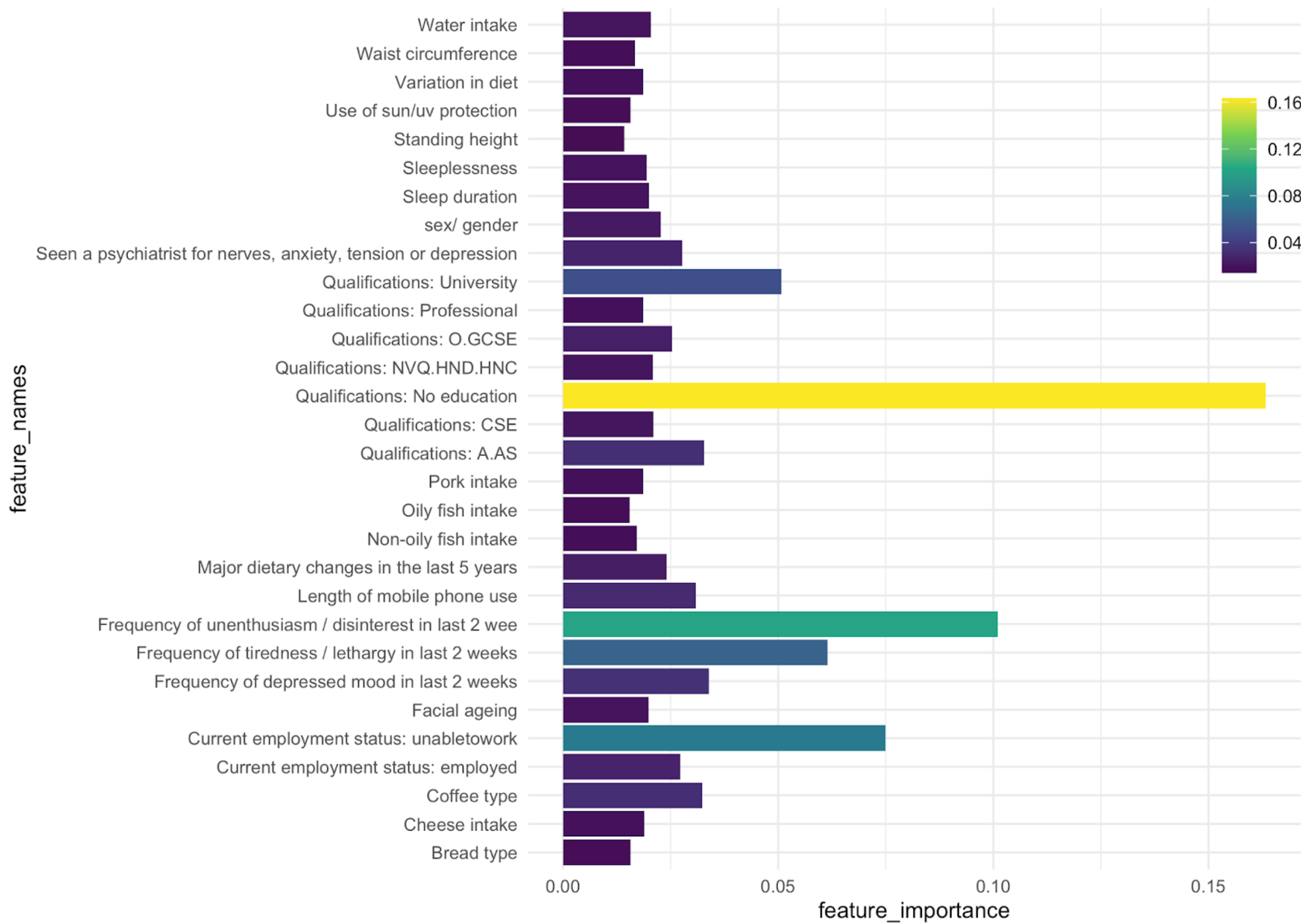
30 accessible variables with age best models and their importance in experiment 4. At right results displayed for Statistical Learning using Logistic Regression, and at left results for machine learning using XGBoost.

Accessible without Age

Accessible without age experiment: Statistical Learning



Accessible without age experiment: Machine Learning



30 accessible variables without age best models and their importance in experiment 5. At right results displayed for Statistical Learning using Logistic Regression, and at left results for machine learning using XGBoost.

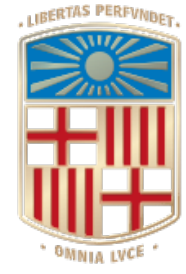


Conclusions



1. If we move from SL to ML for dementia prediction we we gain accuracy.
2. Exposome data is good enough for an accurate prediction of the syndrome.
3. Our models are predicting not just age.
4. We got a high accuracy in external validation.
5. Most relevant exposures found in this study could be used in the future to select risk patients for drug studies.



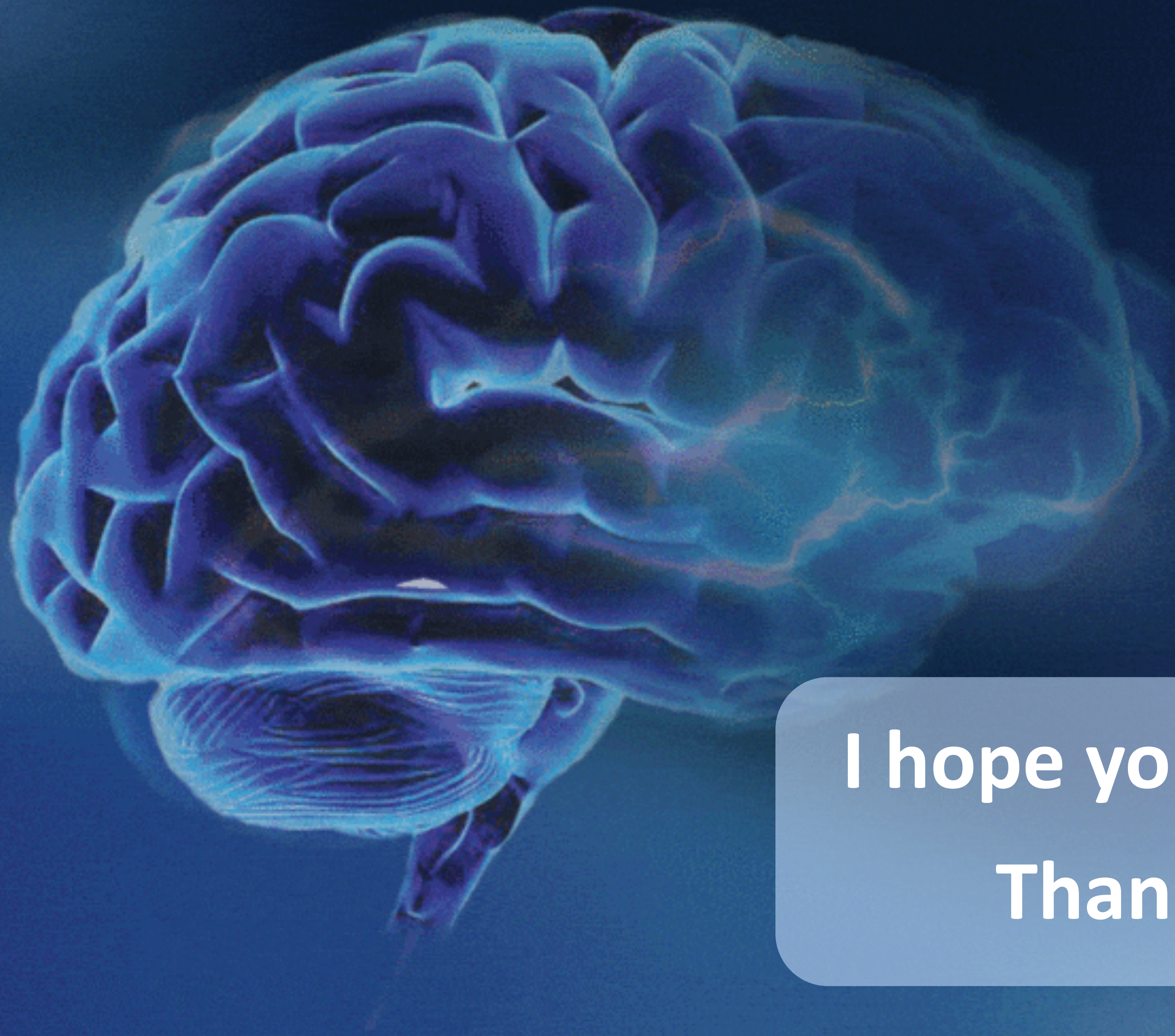


Mapping the exposome effect on the brain

An Artificial Intelligence approach using MRI images from the UK Biobank cohort



- DIET
- Coffe intake
- Tea intake
- ...
- BEHAVIOUR
- Length of mobile phone use
- Nap during the day
- ...
- EARLY LIFE FACTORS
- Maternal smoking around birth
- ...
- ENVIROMENTAL FACTORS
-



**I hope you find it interesting.
Thanks for listening!**