



Per person profiling

Gut microbiome

16S rRNA
Metagenomics



Blood tests



Questionnaires

Food frequency
Lifestyle
Medical



Anthropometrics



Diary (food, sleep, physical activity)

Using smartphone-adjusted website

5,435 days, 46,898 meals, 9.8M Calories, 2,532 exercises

Continuous glucose monitoring

Using a subcutaneous sensor (iPro2)

130K hours, 1.56M glucose measurements

Standardized meals (50g available carbohydrates)

Day 1 Day 2 Day 3 Day 4 Day 5 Day 6 Day 7



Bread

Bread

Bread &
butter

Bread &
butter

Glucose

Glucose

Fructose

Computational analysis

Main cohort



800 Participants

PPGR prediction



Validation cohort



100 Participants

Dietary intervention

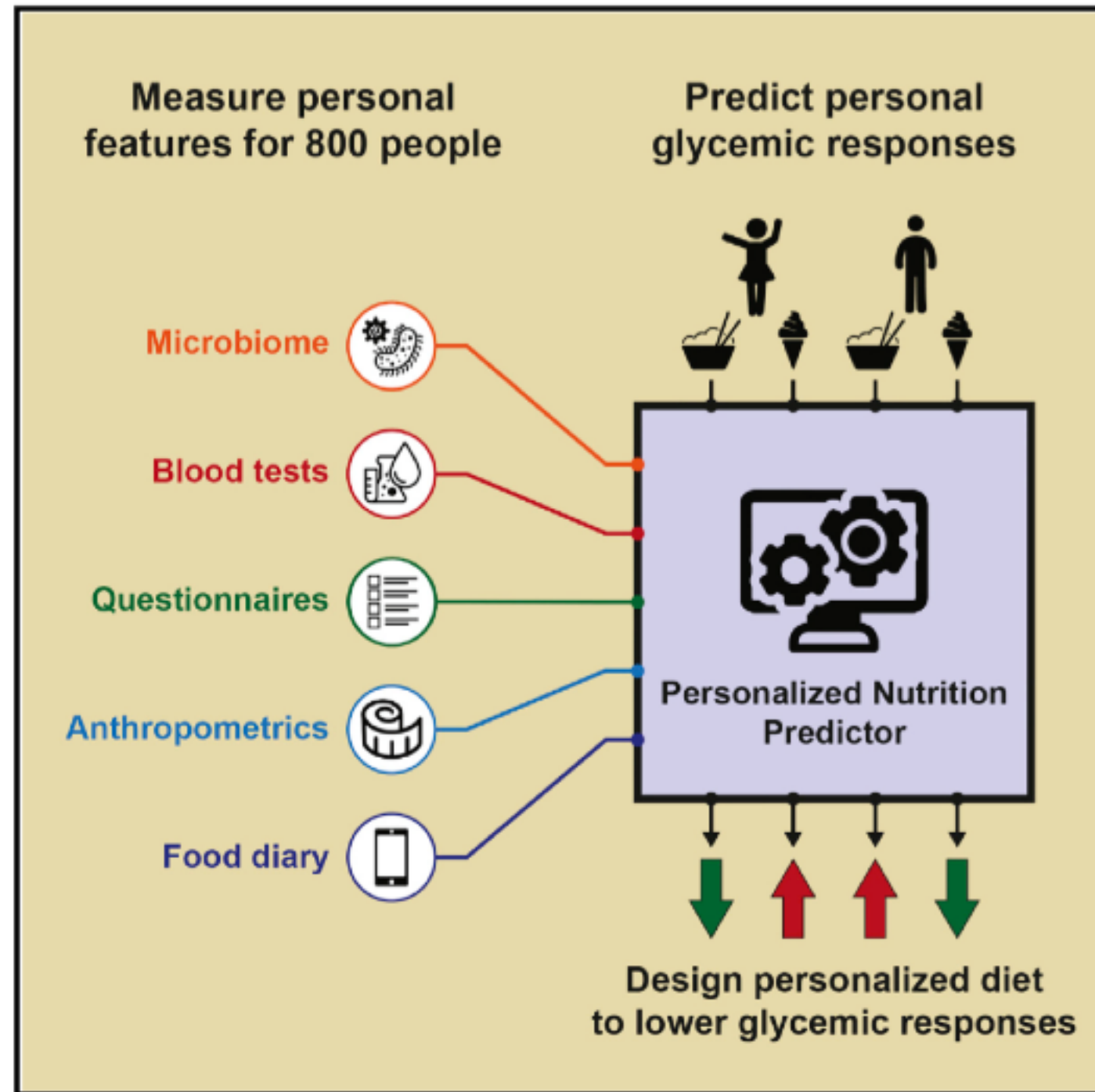
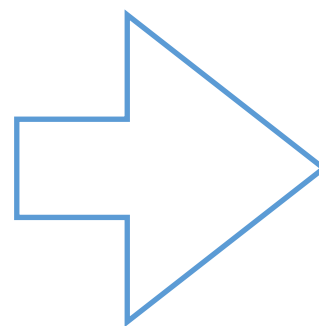
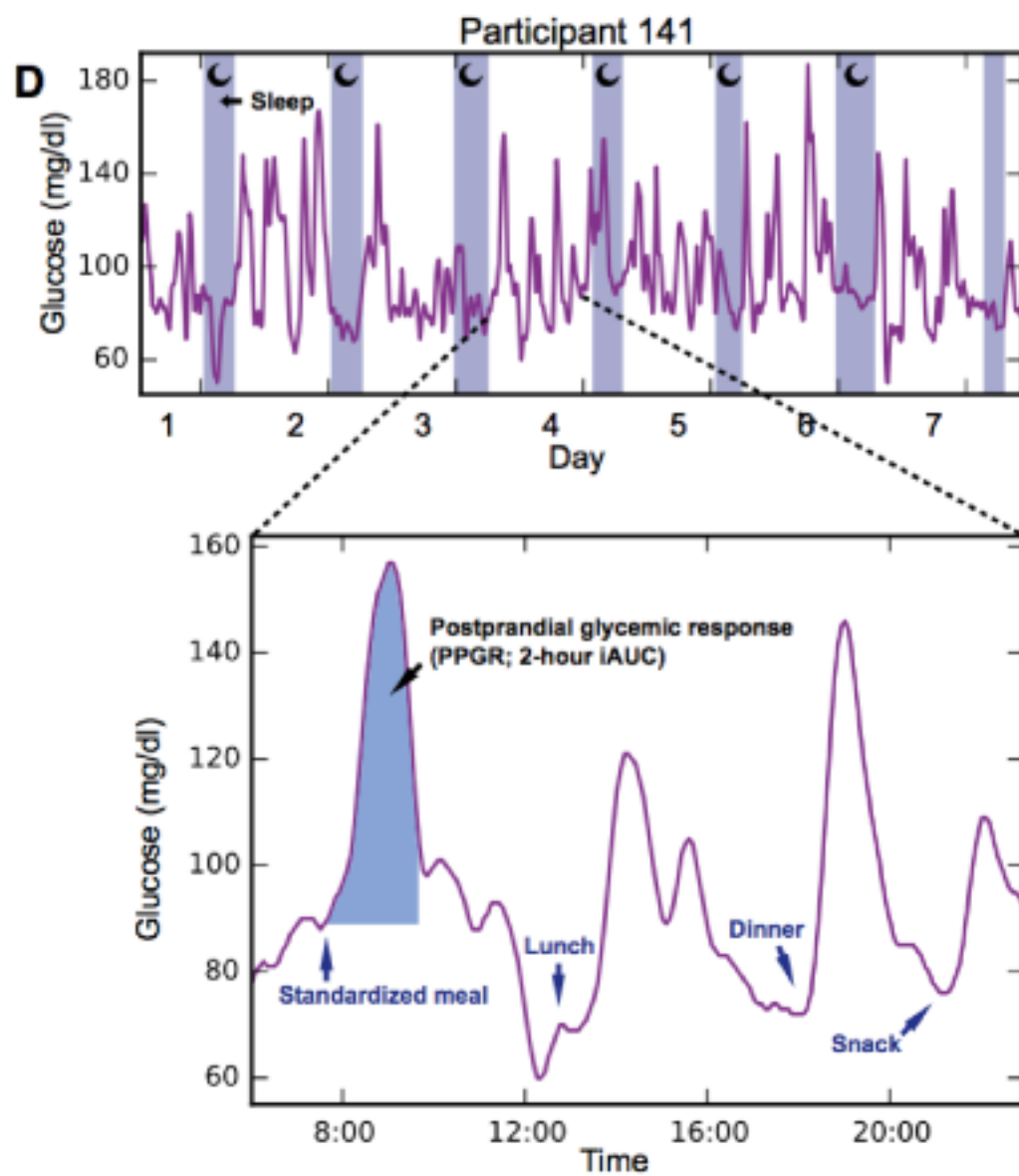


26 Participants



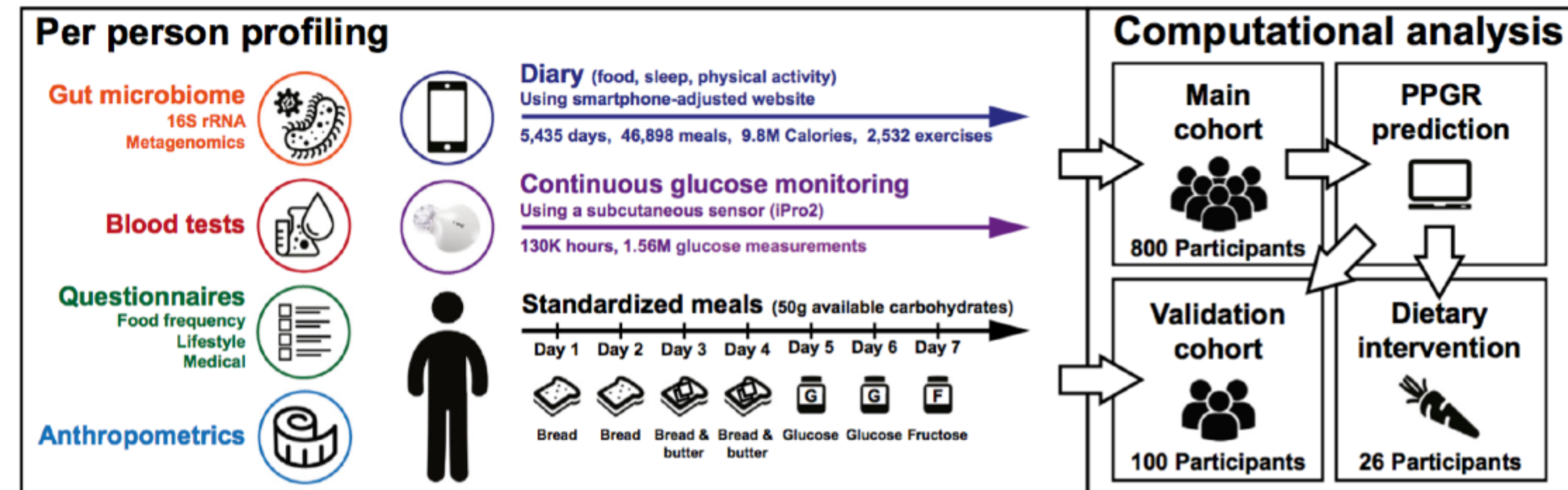
Personalized Nutrition by Prediction of Glycemic Responses

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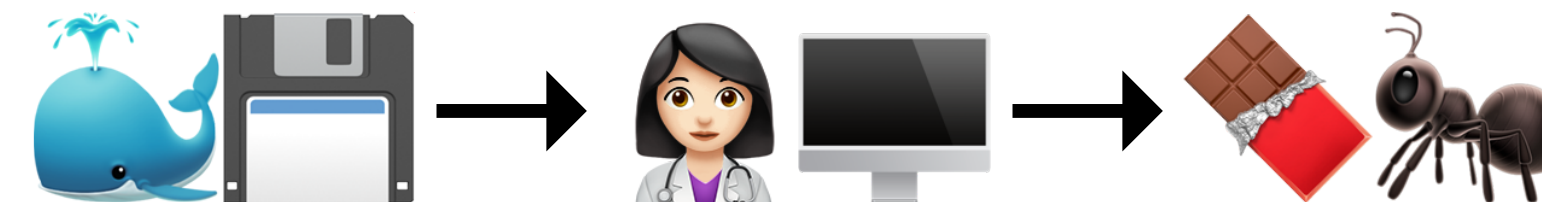
Personalized nutrition

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LETTERS

<https://doi.org/10.1038/s41591-019-0485-4>

nature
medicine

Meta-omics analysis of elite athletes identifies a performance-enhancing microbe that functions via lactate metabolism

Model_1 < -lme(*Veillonella*~time + sex + weight + BMI + age + race + menstruation + vegetables + fruits + grains + protein + dairy + dietary_protein_supp, random = ~1|subjectID, data = marathon16S)

