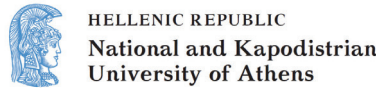


Next generation protein biomarker signature in blood for early detection of colorectal cancer

Christos Fotis<sup>1,2</sup>, Nikolaos Meimetis<sup>3</sup>, Nikolaos Tsolakos<sup>1</sup>, Athanasios Stamogiannos<sup>1</sup>, Vaia Pliaka<sup>1</sup>, Ioannis Temponeras<sup>1</sup>, Eleni Koukouloti<sup>4</sup>, Elli Karatza<sup>4</sup>, Pinelopi Antonopoulou<sup>4</sup>, Krystalina Kapantai<sup>4</sup>, Georgia Anagnostou<sup>4</sup>, Douglas A. Lauffenburger<sup>3</sup>, Ioannis S. Papanikolaou<sup>4</sup>, Leonidas Alexopoulos<sup>1,2</sup>

- 1)Protavio Ltd, Demokritos Science Park, Athens, Greece
- 2)Biomedical Systems Laboratory, National Technical University of Athens, Athens, Greece
- 3)Department of Biological Engineering,Massachusetts Institute of Technology, Cambridge, MA, USA,
- 4)Hepatogastroenterology Unit, Second Department of Internal Medicine- Propaedeutic, Medical School, National and Kapodistrian University of Athens, Attikon University General Hospital, Athens, Greece



NATIONAL  
TECHNICAL  
UNIVERSITY  
OF ATHENS

PROTAVIO  
Proteomics for Better Health

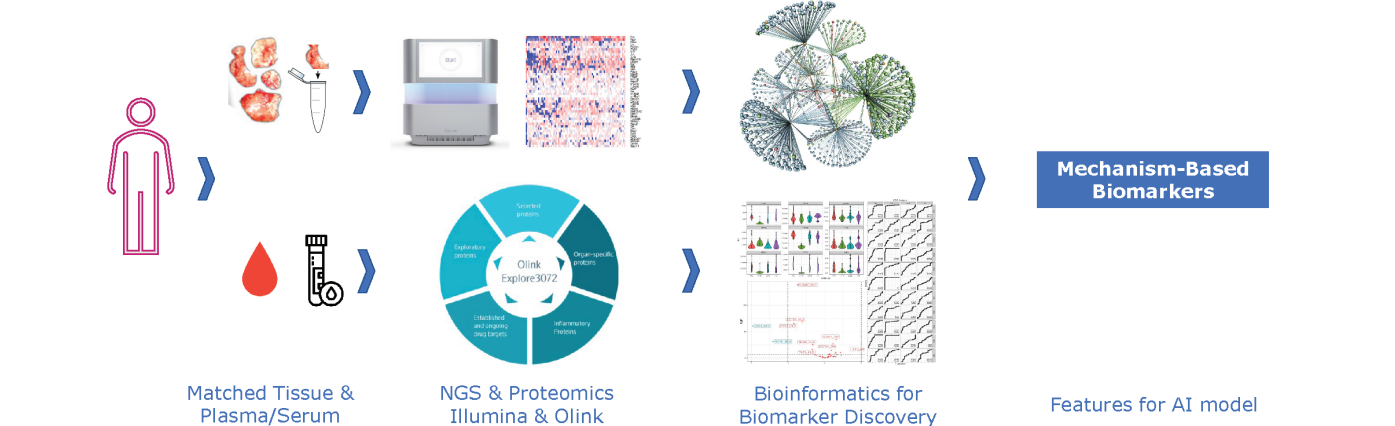
MIT BE  
BIOLOGICAL ENGINEERING

DALLAB

Abstract

**Importance:** Colorectal cancer is the second leading cause of cancer-related deaths in Europe and the United States.

**Aim:** Discover novel blood-based diagnostic protein biomarkers that are directly related to the CRC mechanism.

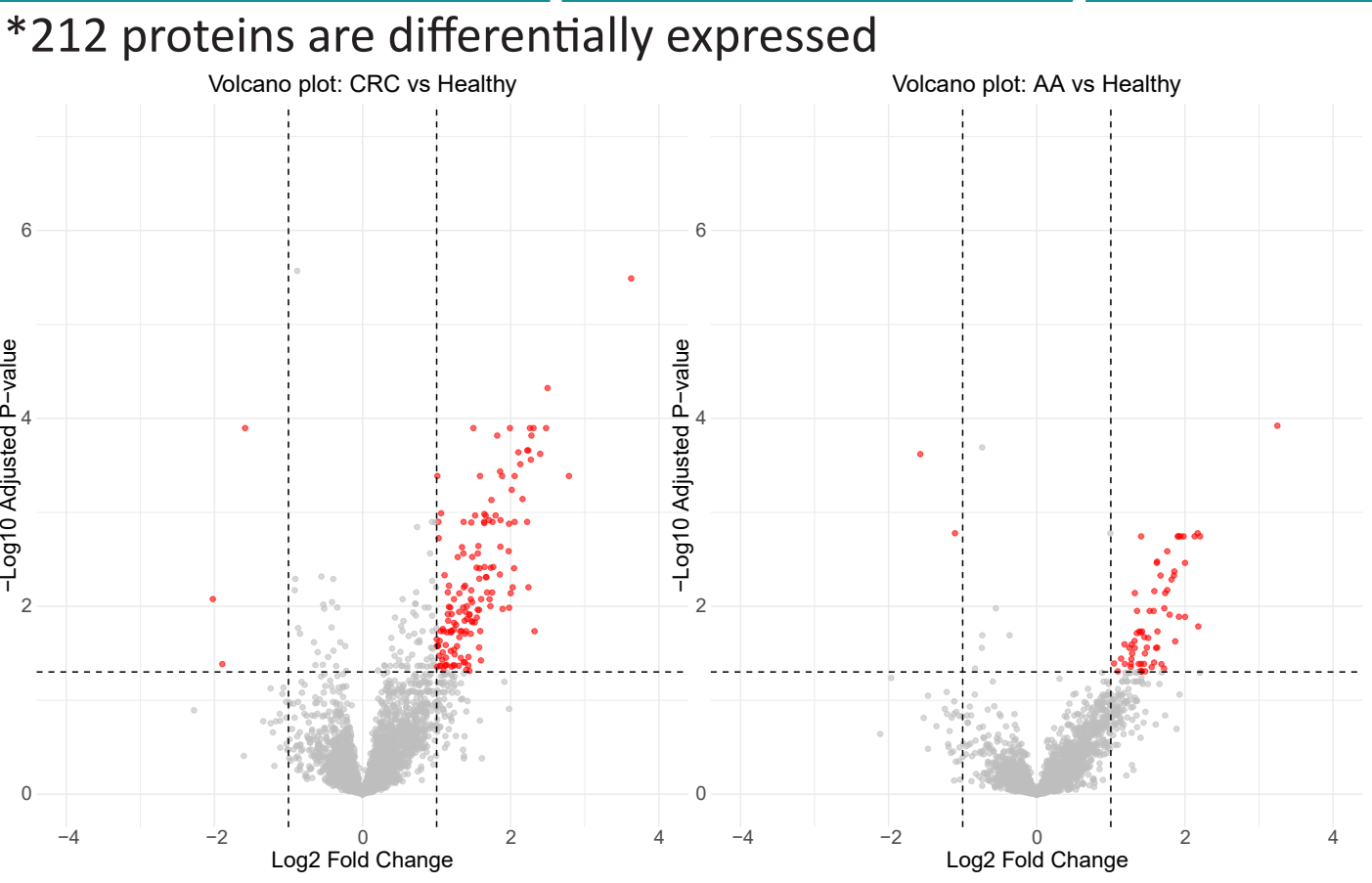


Discovery Clinical Cohort

Group	Participants	Age (median)	Age (IQR)	Sex (%)
CRC	65	65.0	15.0	Female: 45.4 Male: 54.6
AA	106	65.0	13.0	Female: 40.9 Male: 59.1
NAA	50	63.5	10.8	Female: 47.6 Male: 52.4
Healthy	75	56.5	17.3	Female: 54.1 Male: 45.9

- CRC = Colorectal cancer patients
- AA = Advanced Adenomas
- NAA = Patients with non neoplastic findings
- Healthy = Patients with no findings

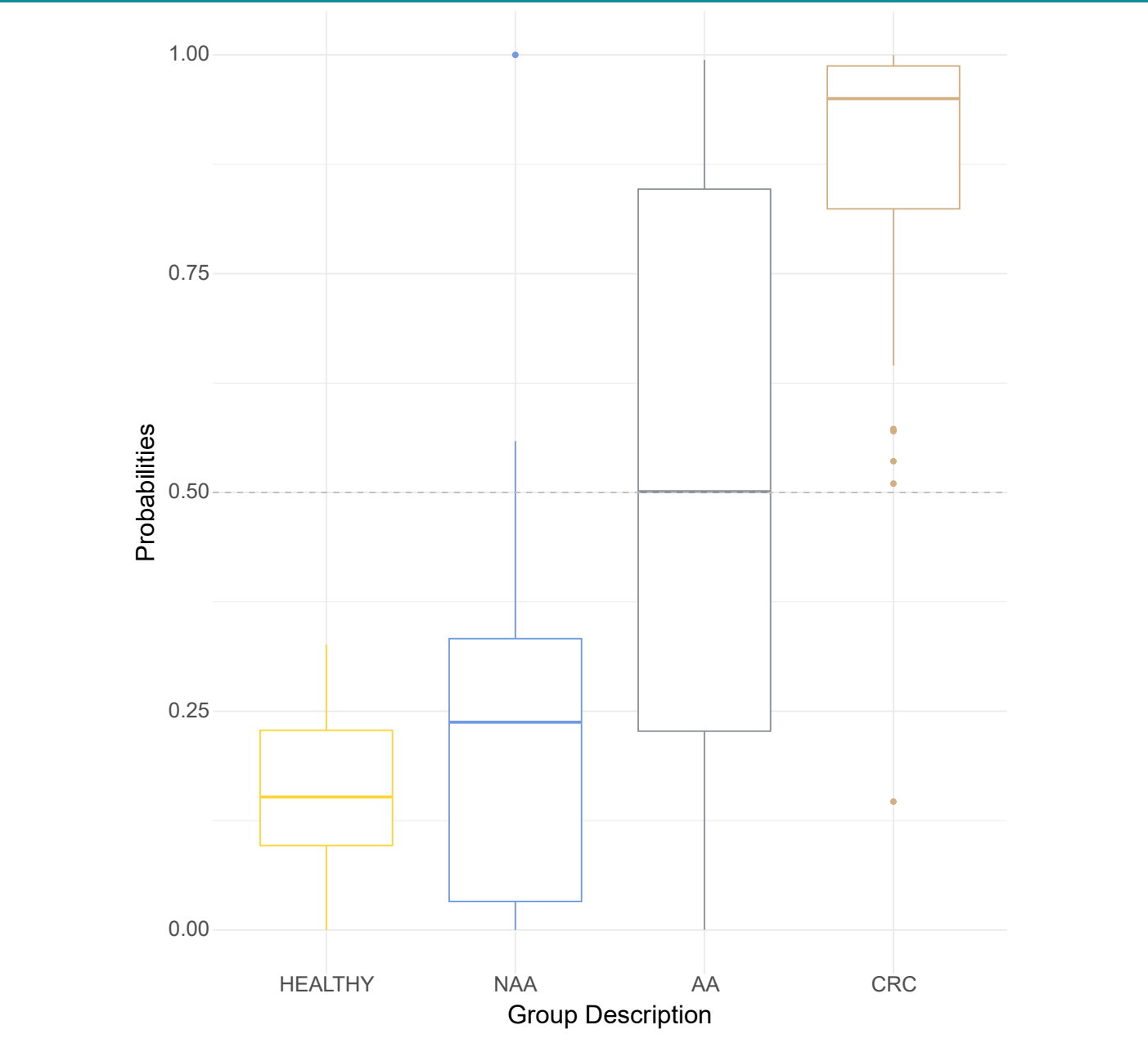
Differential expression analysis



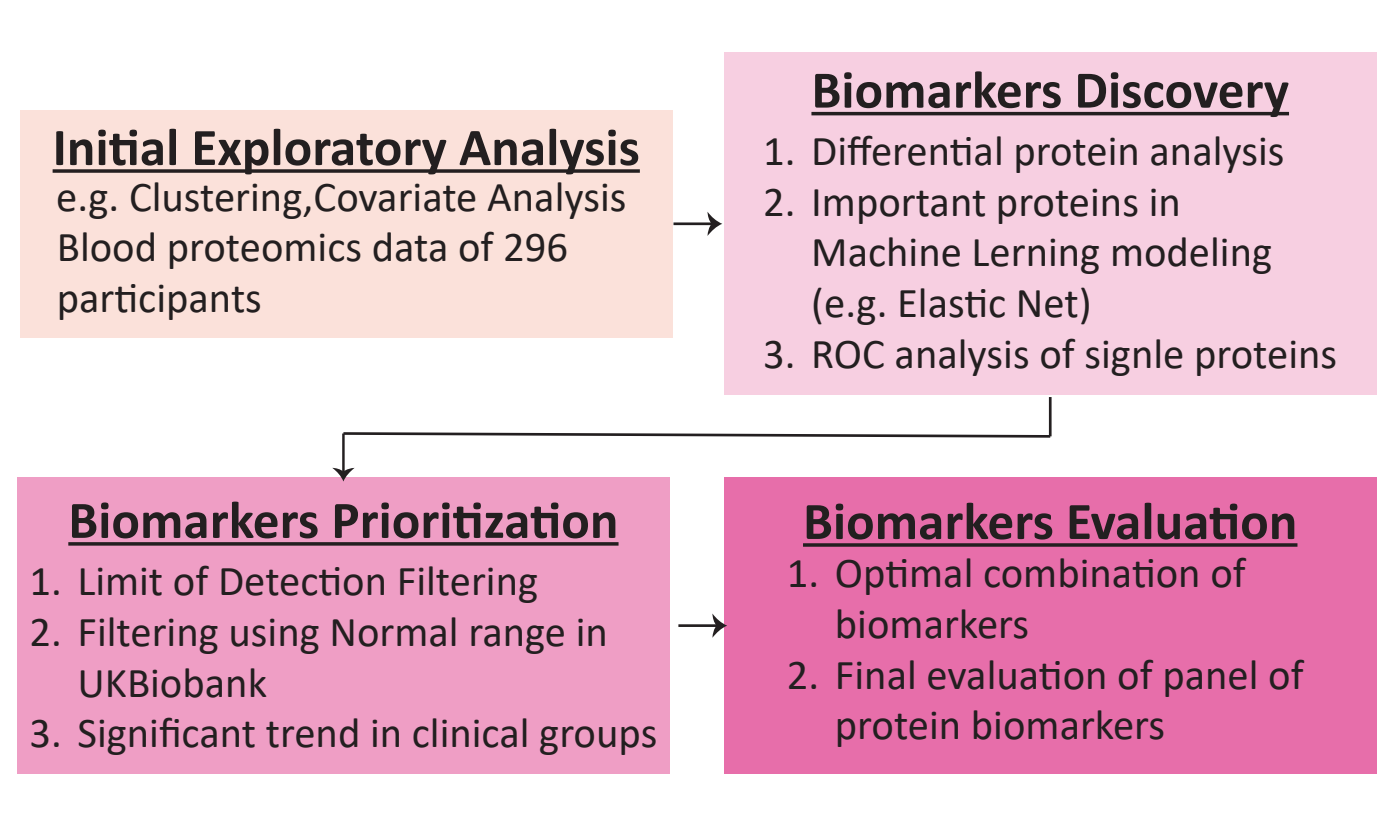
The final model has high performance across all groups

Comparison	AUC (%)	Sensitivity (%)	Sensitivity 95% CI	Specificity (%)	Specificity 95% CI
CRC vs Normal	95.9	98.1	90.1-100.0	93.1	77.2-99.2
SI vs Normal	96.9	100.0	71.5-100.0	93.1	77.2-99.2
SII vs Normal	97.6	100.0	75.3-100.0	93.1	77.2-99.2
SI/SII vs Normal	97.3	100.0	85.8-100.0	93.1	77.2-99.2
AA vs Normal	78.0	50.0	37.0-63.0	93.1	77.2-99.2
HGD vs Normal	74.7	43.5	23.2-65.5	93.1	77.2-99.2
CIS vs Normal	66.4	50.0	6.8-93.2	93.1	77.2-99.2
AA10 vs Normal	85.8	61.1	35.7-82.7	93.1	77.2-99.2
V vs Normal	74.1	50.0	21.1-78.9	93.1	77.2-99.2

Predicted probabilities separate different groups

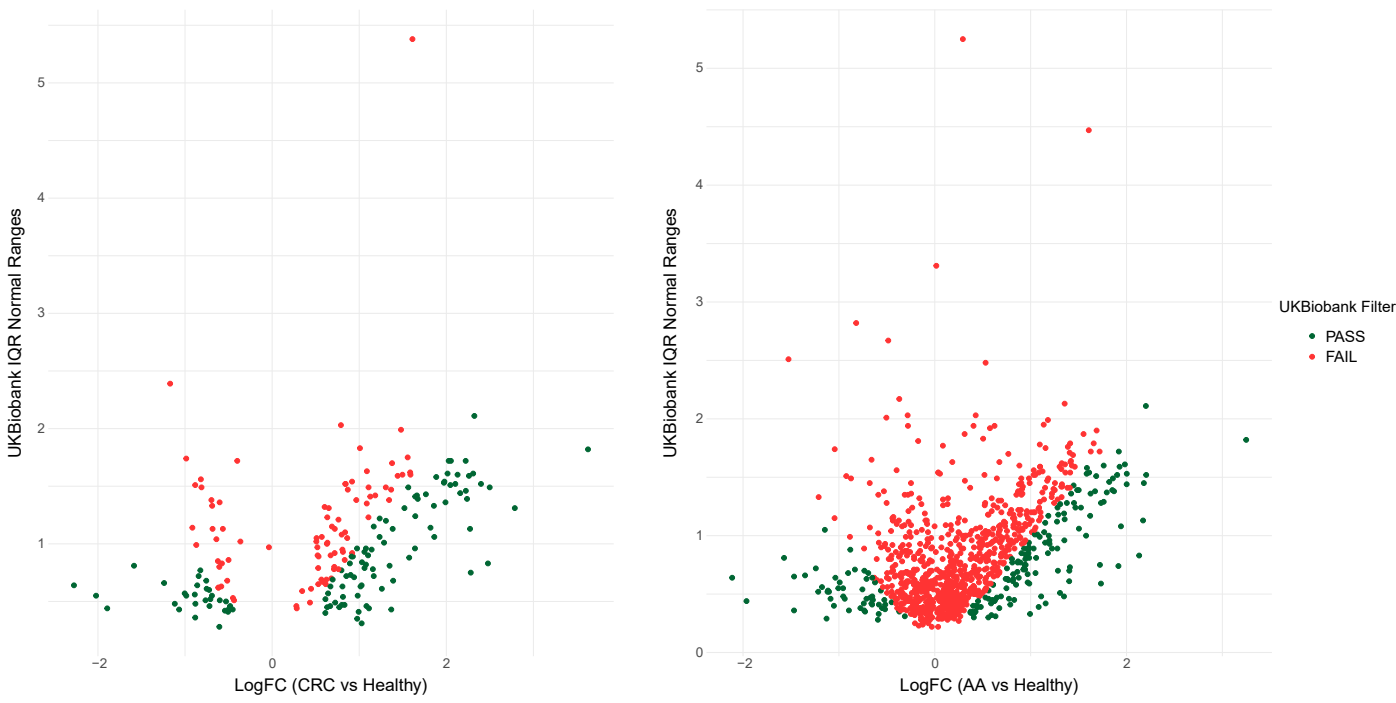


Analysis Pipeline

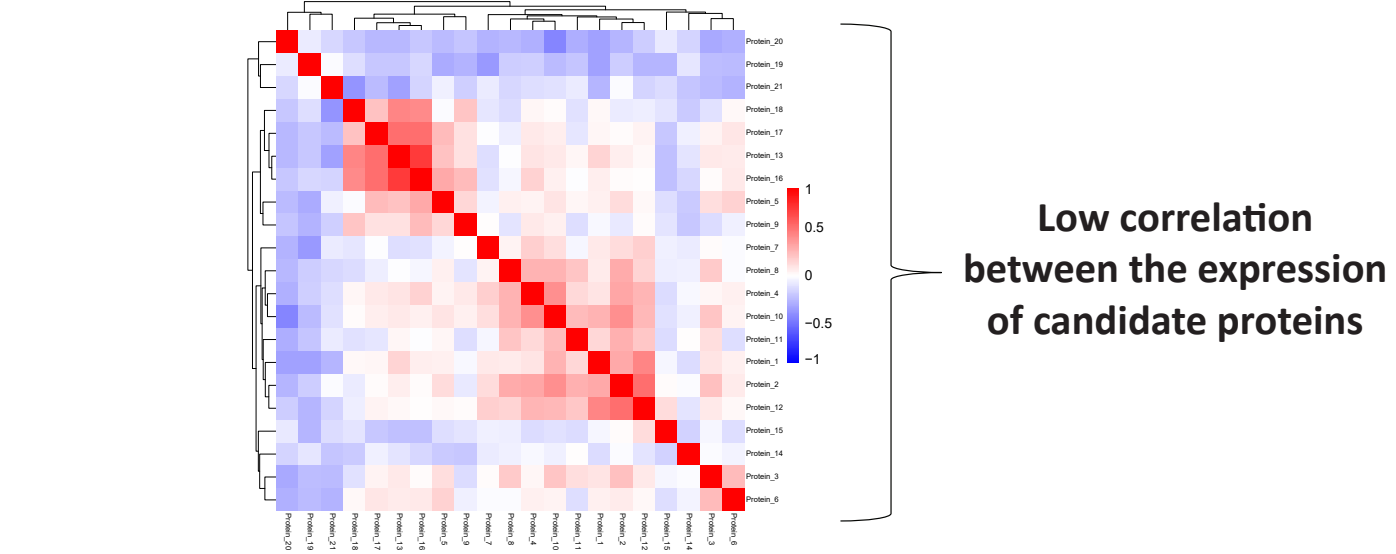


Biomarker prioritization criteria

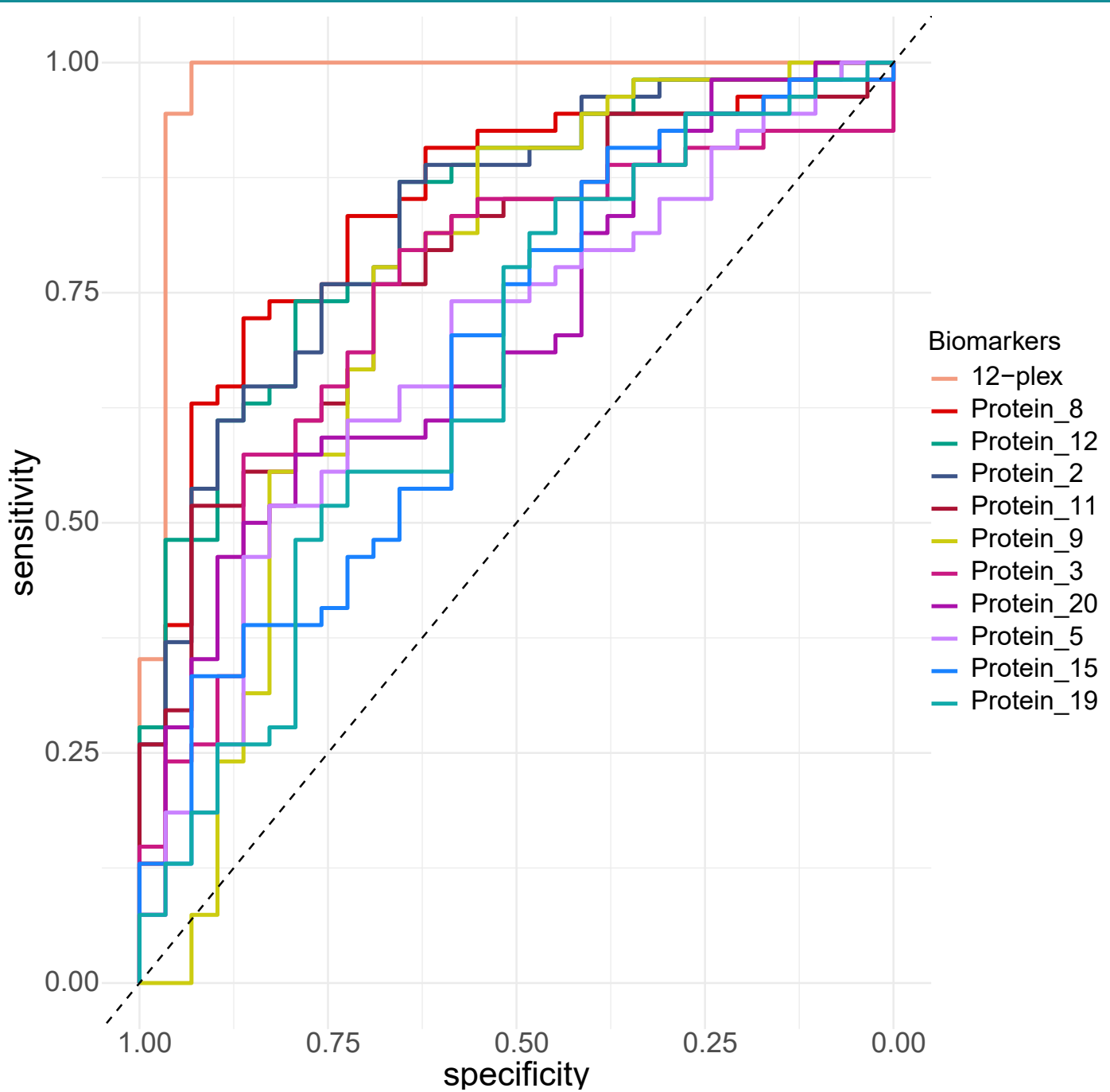
One prioritization criterion is based on the comparison of biological effect size with the Normal ranges from the UK Biobank. We applied this criterion in order to ensure that the final multiplex assay can potentially achieve greater specificity. We select for proteins with a log(FC) greater than the interquartile range in normal cohorts in the UKBiobank (shown in green).



Candidates capture unique information



Performance of 12-plex assay and single biomarkers for CRC detection



Funding & Aknowledgements

This work was funded by the European Union’s Mission Cancer initiative (GA 101096649). Partners from the Biobank of MU Graz and Agios Savvas cancer hospital contributed to the discovery clinical cohort.

dioptra-project.eu

@dioptra\_project

Funded by the European Union

UK Research and Innovation

Project funded by

Schweizerische Eidgenossenschaft  
Confédération suisse  
Confederazione Svizzera  
Confederaziun svizra  
Swiss Confederation  
Federal Department of Economic Affairs,  
Education and Research EAEF  
State Secretariat for Education,  
Research and Innovation SERI