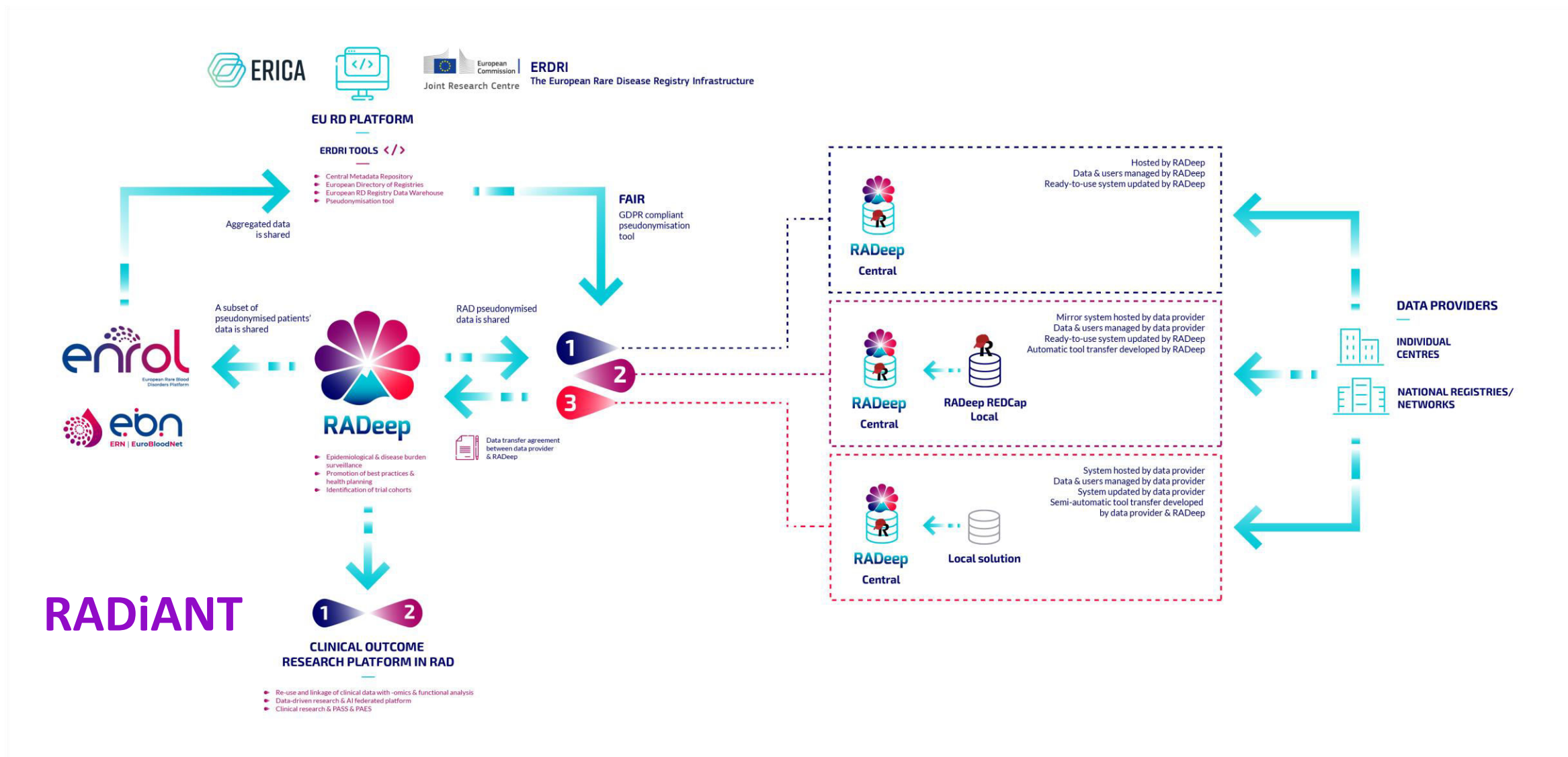


RADeep

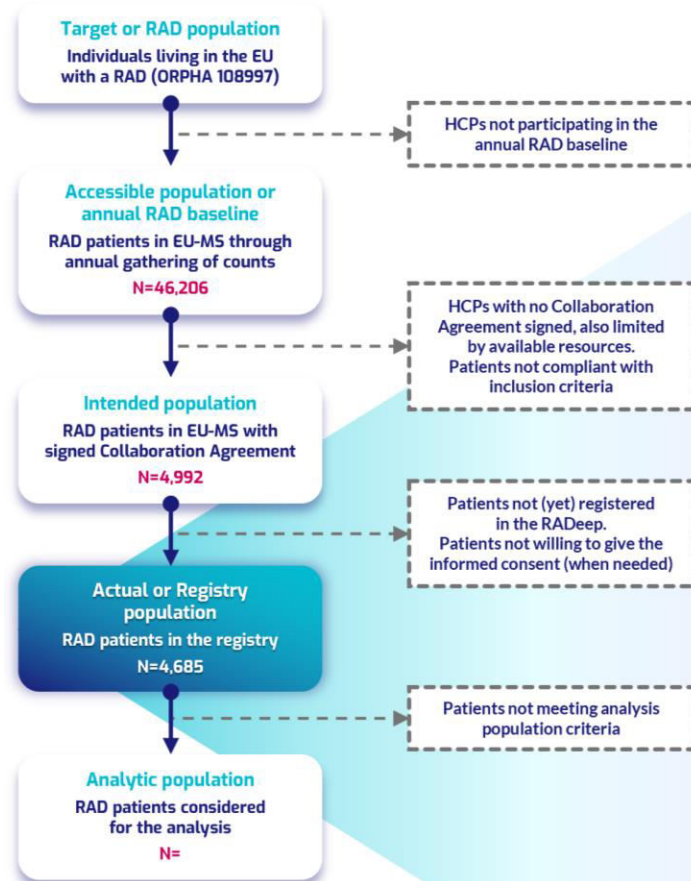
AI Federated platform in anemia

June 2025

RADeep: Clinical data standardization in RADs



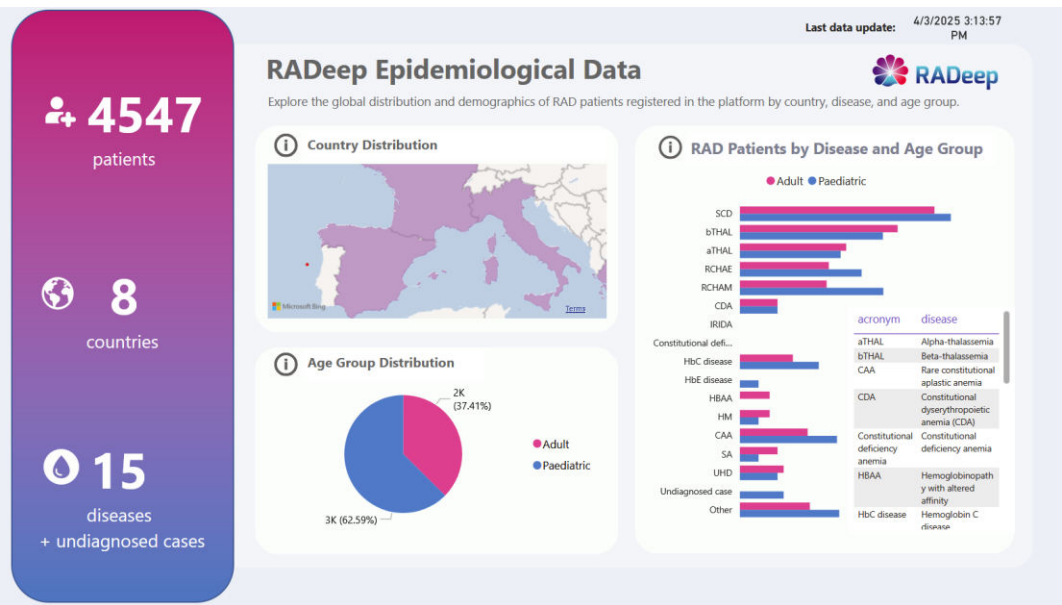
RADeep study populations and CRF summary



Group		Subgroup	Parameters	Mandatory	Longitudinal
1	Patient pseudonym, permissions and biobanking information	1.1. Pseudonym	1	1	0
		1.2. Permissions	5	5	0
		1.3. Biobanking	3	3	0
2	Demographics	2.1. Population distribution	12	8	4
		2.2. Mortality and comorbidities	7	7	6
3	Diagnosis	3.1. Diagnosis	6	6	0
		3.2. Genotype	5	5	0
		3.3. Undiagnosis	2	2	0
		3.4. Disease Onset	4	2	0
4	Physical examination	3.5. Neonatal Manifestations	5	0	0
		4.1. Development	0	0	0
5	Organ Damage	5.1. Organ damage assessment	14	3	14
		5.2. Chronic complications of bones and extremities	8	4	8
		5.3. Chronic cardiac and pulmonary disease	6	5	6
		5.4. Chronic neurological disease	7	6	7
		5.5. Chronic endocrinologic disease	8	7	8
		5.6. Chronic liver and renal disease	8	7	8
		5.7. Visual and hearing disease	5	4	5
6	Acute Complications	6.1. Acute complications in RADs (Except SCD) requiring hospitalization or emergency admission for more than 24 hours	4	3	4
		6.2. Acute complications in SCD requiring hospitalization or emergency admission for more than 24 hours	4	3	4
		6.3. Intensive Care Unit Admission in the last 12 months	1	1	1
7	Clinical manifestations and surgery	7.1. Spleen	4	4	4
		7.2. Gallbladder	3	0	3
8	Treatments	8.1. Blood transfusion	14	7	13
		8.2. Chelation	5	5	5
		8.3. Hydroxyurea	4	1	4
		8.4. Specific treatment(s)	2	2	2
		8.5. Haematopoietic stem cell transplantation (HSCT) / gene therapy	5	5	5
		8.6. Inclusion in clinical trial protocol	2	2	2
9	Fertility	9.1. Fertility and Offspring	2	2	2
10	Disability	10.1. Disability	3	1	3
11	Laboratory tests	11.1. Complete blood count	17	17	17
		11.2. Biochemical tests	14	14	14
		11.3. Hemoglobin tests	7	4	1
		11.4. Enzyme tests	17	17	0
		11.5. Membrane tests	3	3	0
		11.6. Other Laboratory tests	4	4	0
Total			221	170	150

Gliklich RE, Dreyer NA, Leavy MB, editors. Registries for Evaluating Patient Outcomes: A User's Guide [Internet]. 3rd ed. Rockville (MD): Agency for Healthcare Research and Quality (US); 2014 Apr. Report No.: 13(14)-EHC111. PMID: 24945055.

Public dashboard: <https://www.radeepnetwork.eu/epidemiological-data/>



Private dashboards: <https://radeep.vhir.org/>

VHIR Shiny Applications

Interactive data visualization tools

RADeep Descriptive Tables
RADeep Descriptive tables with granular information per country, disease, sub-disease, age-group and gender.

Launch Application

Last updated: March 2025

RADeep Data Completeness Report
RADeep Completeness reports by variable with filters for: mandatory and optional parameters, active patients, country, hcpid and diagnosis.

Launch Application

Last updated: March 2025

RADeep Data Coverage Report
RADeep coverage reports by data parameter with filters for: country, RADs and mapping status

Launch Application

Last updated: May 2025

- Real time data quality rules
 - Alerts when data is being entered

Name of the Healthcare Provider (HCP)

Name of the physician in charge of the patient in the current care unit

Select the date of the patient's first contact with a specialized healthcare center, or choose "Never" or "Not available" if applicable

* must provide value

Enter the date of the patient's first contact with the specialized healthcare center in the format dd-mm-yyyy Today D-M-Y

* must provide value

If dd and/or mm is not available, complete with 07 for mm and 15 for dd

Form Status

Alert

The value you provided is outside the suggested range (05-04-2000 - 03-04-2025). This value is admissible, but you may wish to double check it.

Close

- Asynchronous data quality rules:
 - Data quality module accessible for data entrists

1	[genetic_label_hbb] The genetic variations do not match with the subdiagnosis for the HBB gene.	(([hbb_label] = '1' AND [sub_orpha_id_scd] <> '1') OR ([hbb_label] = '2' AND [sub_orpha_id_scd] <> '4') OR ([hbb_label] = '3' AND [sub_orpha_id_scd] <> '8') AND	✓	0					
2	[globaluniqueid] (Patient's pseudonym) should have a value but is missing.	(([globaluniqueid] = "") AND [patient_registration_complete] = 2	✓	1	export view	0	0	0	0
3	[consent_registry] (I confirm that the legal basis allowing the processing of this pseudonymized clinical information within the registry is secured, warranting participants' rights according to GDPR) should have a value but is missing.	(([consent_registry(1)] = '0') AND [patient_registration_complete] = 2	✓	2	export view	0	0	0	0
4	[consent_reuse] (I confirm that patient consent has been obtained for pseudonymized data in RADeep to be re-used by third parties in order to contribute to projects whose objectives are directly connected to improve healthcare provision for rare anemia disorders) should have a value but is missing.	(([consent_reuse] = "") AND [patient_registration_complete] = 2	✓	2	export view	0	0	0	0
5	[consent_no_eu] (I confirm that patient consent has been obtained for transferring pseudonymized data in RADeep to non-EU countries in order to contribute to projects whose objectives are directly connected to improving healthcare provision for rare anemia disorders) should have a value but is missing.	(([consent_no_eu] = "") AND [patient_registration_complete] = 2	✓	3	export view	0	0	0	0

Quality management

Data Quality flow



- Central Solution

1. **RADeep Data Monitor** runs the data quality rules.
2. **RADeep Data Monitor** open and assigns the query.
3. **Data Entries** resolves the query assigned to them.
4. **RADeep Data Monitor** closes the query.
5. **RADeep Data Monitor** generates periodically a report for each HCP with open queries.

- Local Solution

1. **Data Entries** generates their own internal data quality flow.
2. **Data Entries** resolves the query before sending the data for batch transfer.
3. **RADeep Data Monitor** runs the data quality rules.
4. **RADeep Data Monitor** generates a report for each HCP.

The query in RAdDeep REDCap central is closed when a new data batch transfer is performed.



RADeep



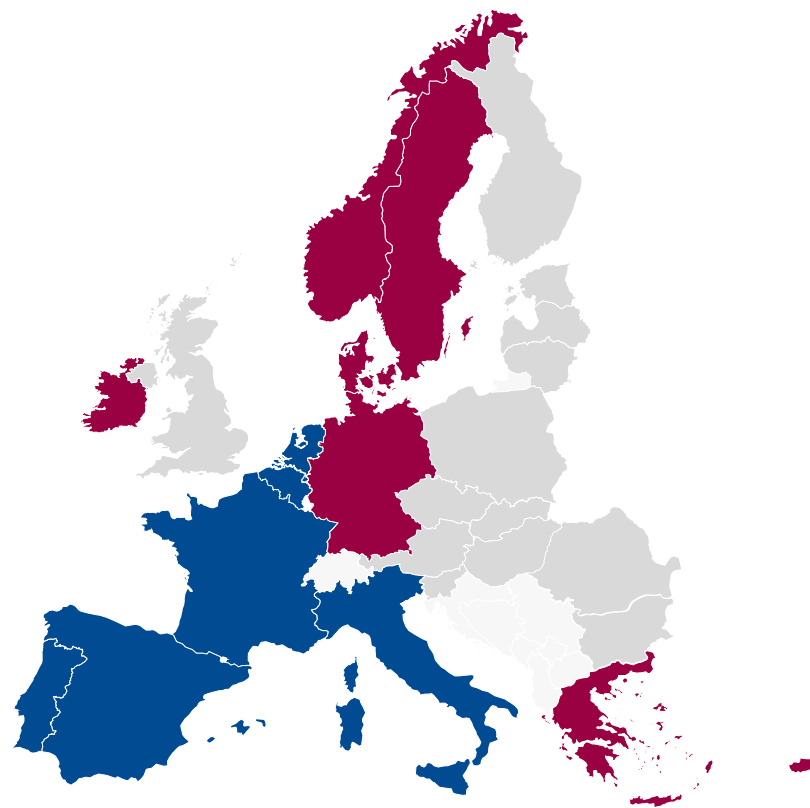
RADiANT

**14 Ongoing collaboration
agreements involving 202 HCPs
in 13 EU countries:
12 Member States**

- Belgium
- Cyprus
- Denmark
- France (2)
- Germany
- Greece
- Ireland
- Italy
- Portugal
- Spain
- Sweden
- The Netherlands
- + Norway

Countries onboard in
the research platform
in RADs: 4 Members
States

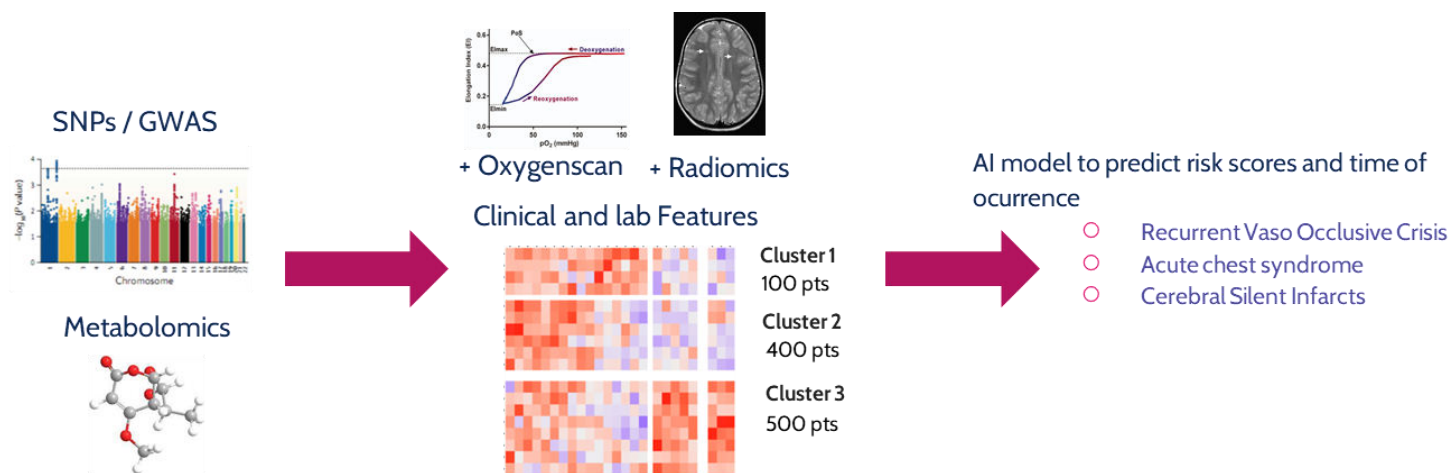
- France (2/2)
- Italy (1/1)
- Spain (1/10)
- The Netherlands (1/4)
- Belgium (2/2)
- Portugal (3/3)



AI aims in SCD

Development of AI Solutions to Improve SCD Clinical Management through a Personalized Precision Medicine Approach:

- ❑ Omics-based Classification and Prognosis of SCD and Omics-based Clinical Decision Making in SCD
 - Aim 1/2: Cluster SCD patients according to their genomic /metabolomic profile
 - Aim 3: To develop a probability score using AI-based brain MRI image analysis (radiomics) to detect existing lesions for early diagnosis of cerebral silent infarcts
 - Aim 4: Based on results from objectives 1 to 3, to develop predictive risk scores for recurrent vaso-occlusive crisis, acute chest syndrome, cerebral silent infarct, kidney disease
- ❑ Synthetic data generation in Kidney disease



SCD Use Case

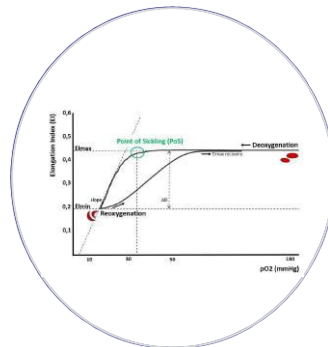
Target 1,000 SCD Patients



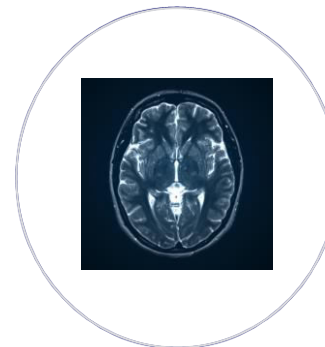
Genetic
modifiers
1000+ GWAS



Metabolomics
1900+
metabolites



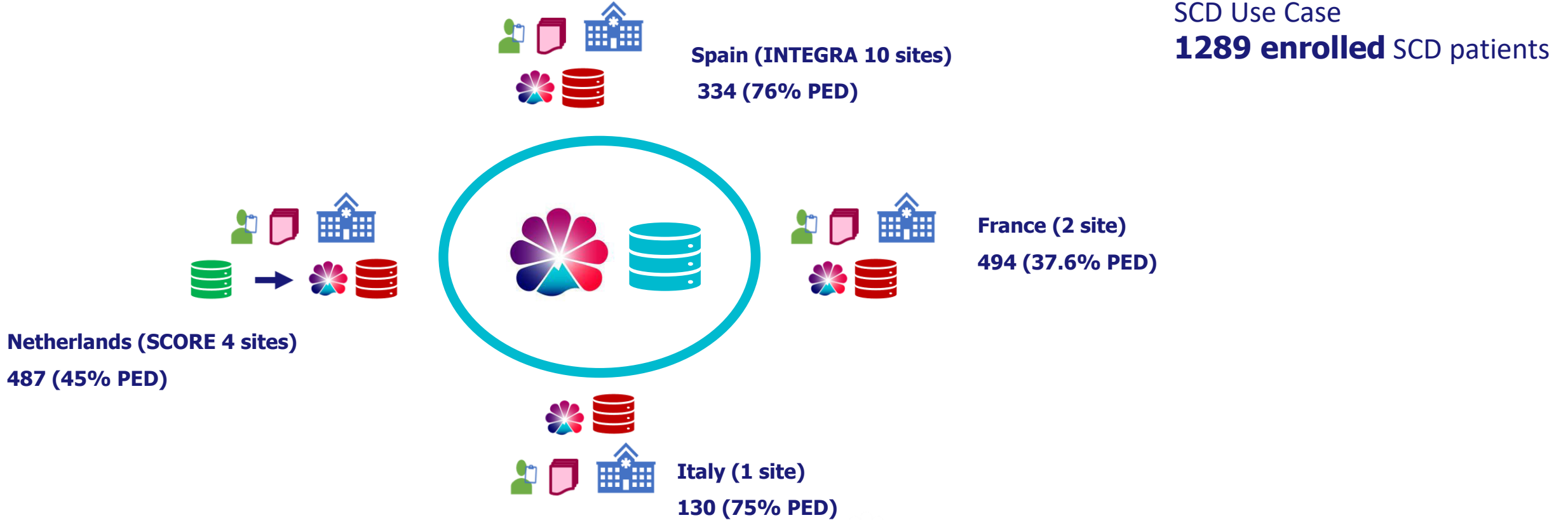
Functional test
Rheology
LoRRca -
Oxygenscan



MRIs



Clinical and lab
data

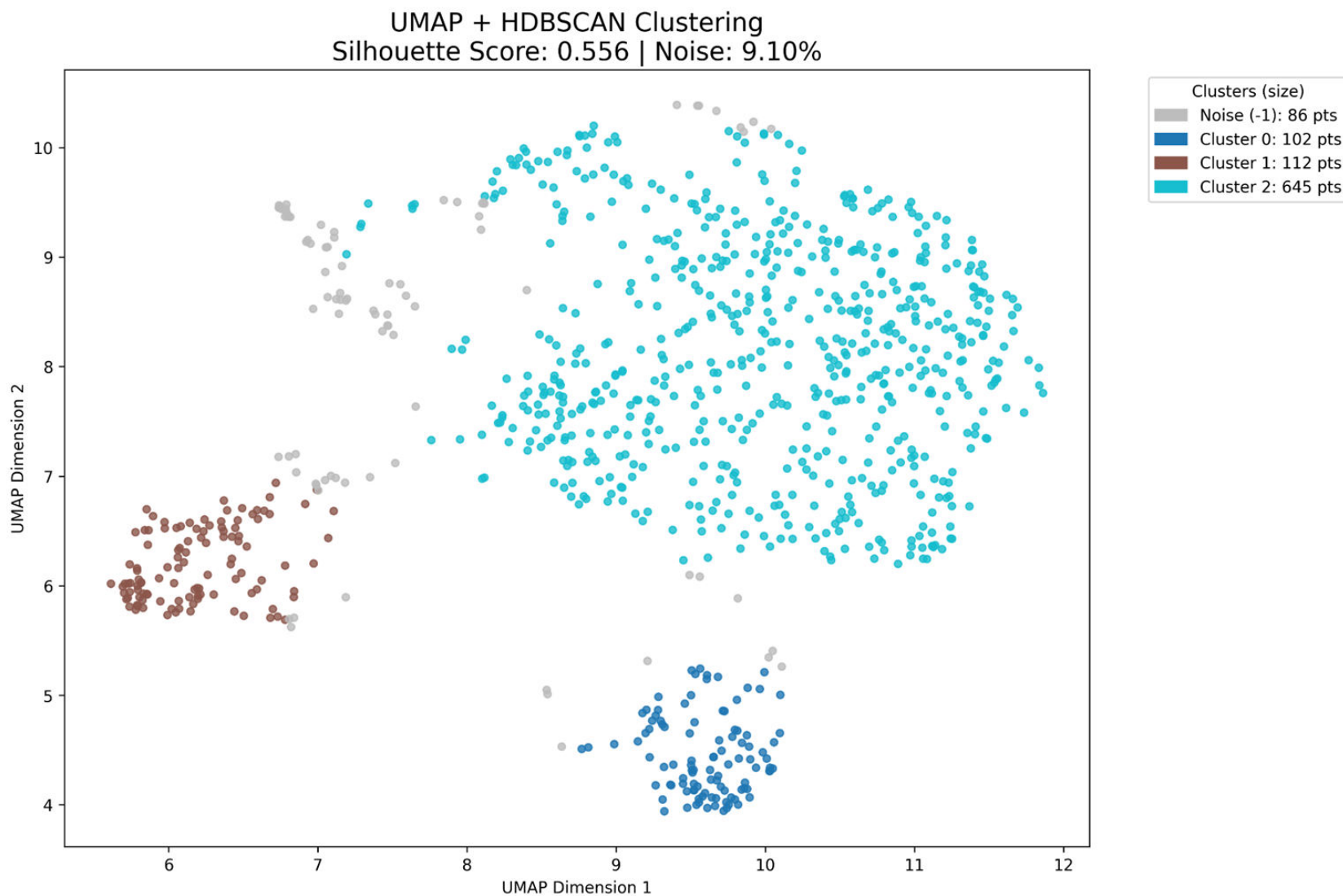


AI Methods

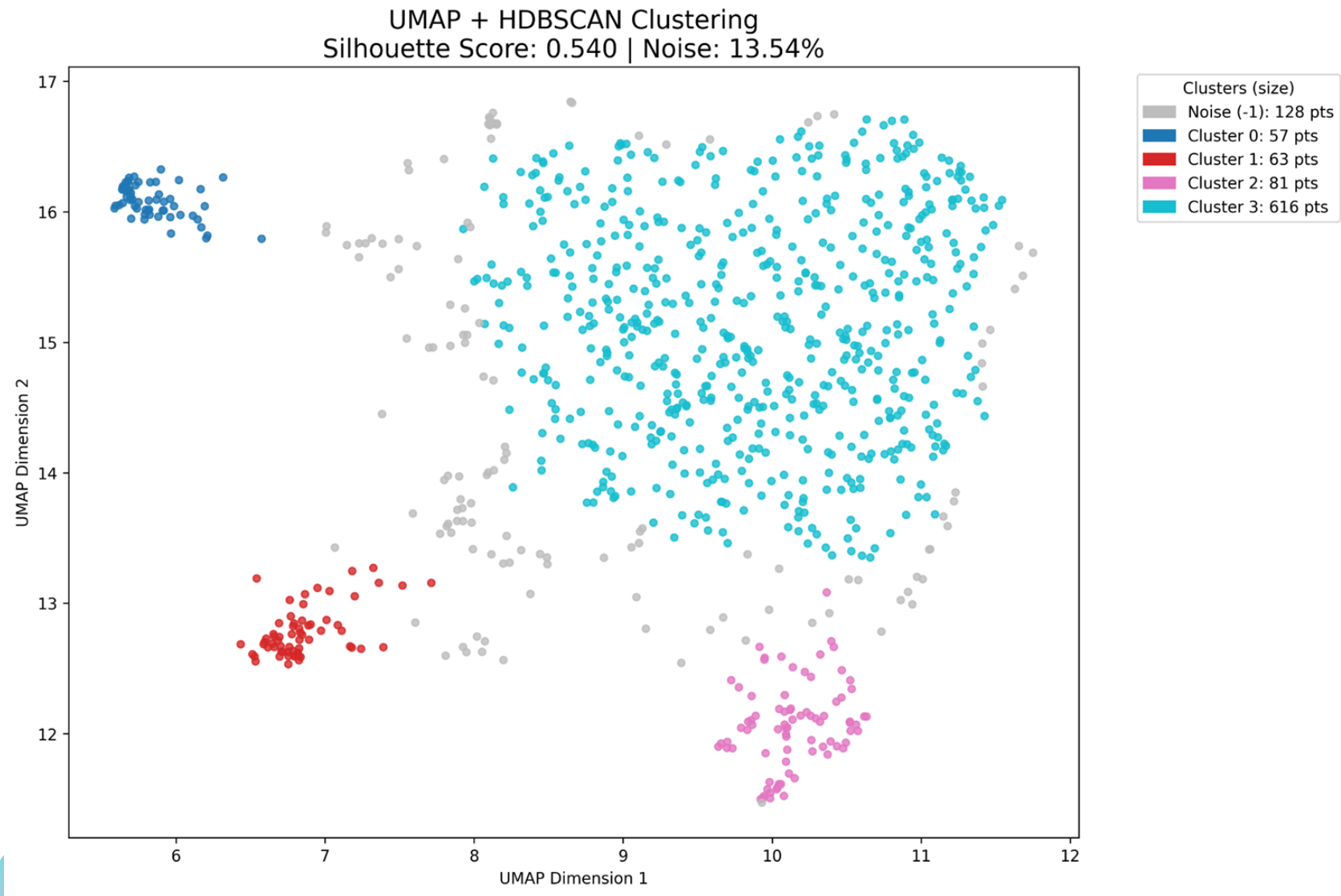
- Clinical dataset: 1274 patients
 - GWAS dataset: 991 patients
-
- Overlap: 959 patients
-
- Clinical dataset: 1274 patients
 - Metabolomics dataset: 979 patients
-
- Overlap: 924 patients



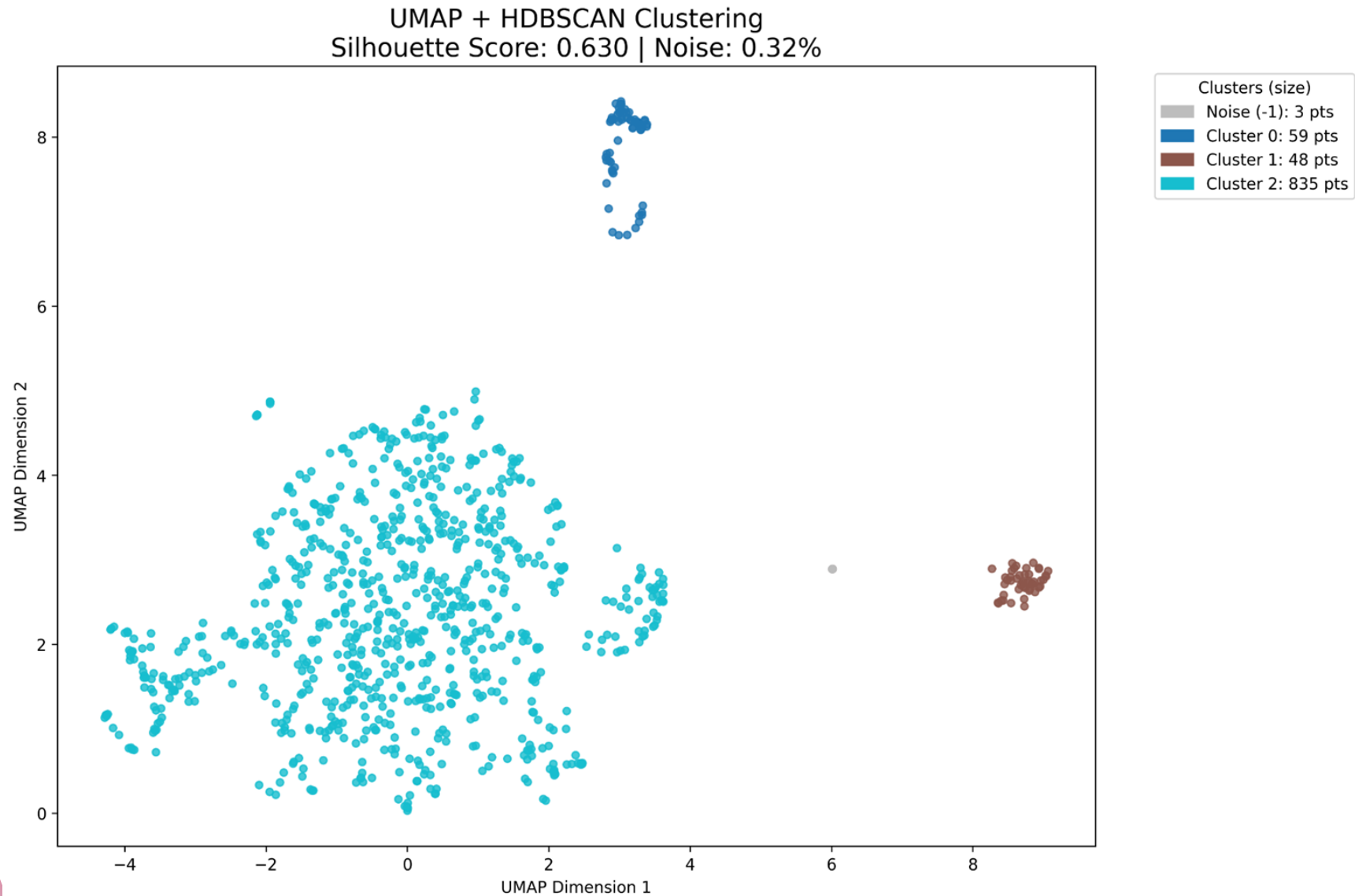
GWAS Clustering (945 samples): Binarized data



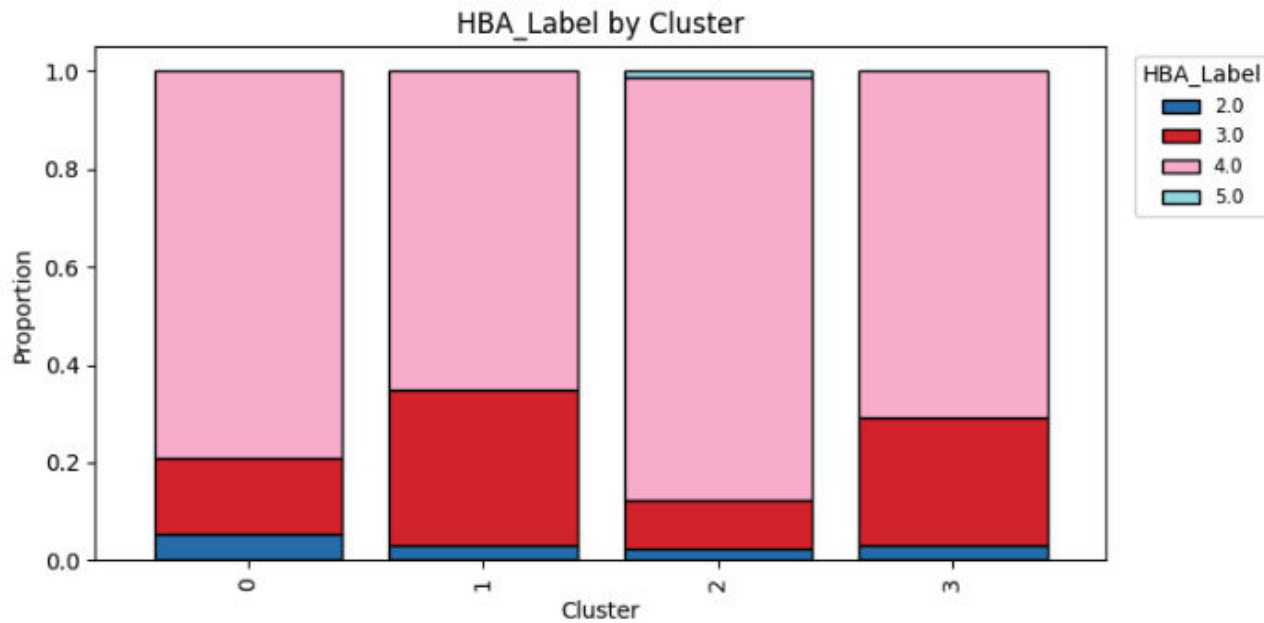
GWAS Clustering (945 samples): {0,0.5,1} mapping



GWAS Clustering (945 samples): One-Hot Encoding



GWAS Clustering (945 samples): {0,0.5,1} mapping



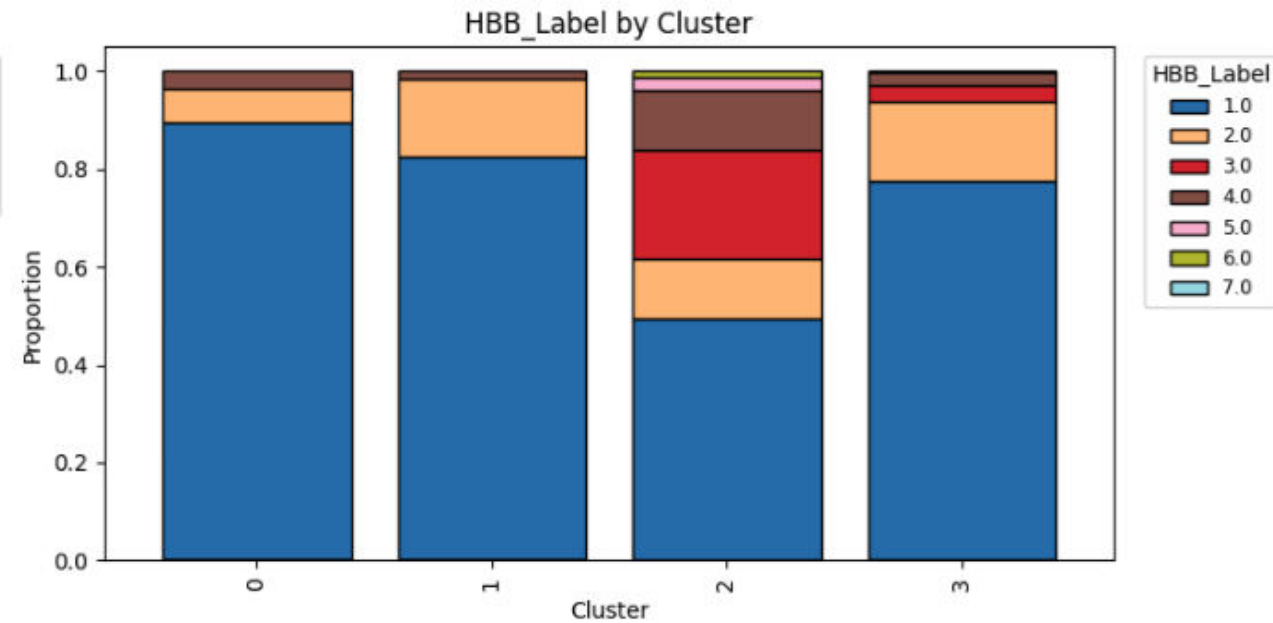
HBA:

5, $\alpha\alpha/\alpha\alpha$ |
4, $\alpha\alpha/\alpha\alpha$ |
3, $\alpha\alpha/\alpha-$ |
2, $\alpha-/ \alpha-$ |
1, $\alpha-/-$ |
8, Inconclusive

Signif. pvalues:

(2, 3) **

(2, 1) *



HBB:

1, HbSS |
2, HbSC |
3, HbSbeta0 |
4, HbSbeta+ |
5, HbSD |
6, Other |
7, HbS HPHF | 8,
Inconclusive

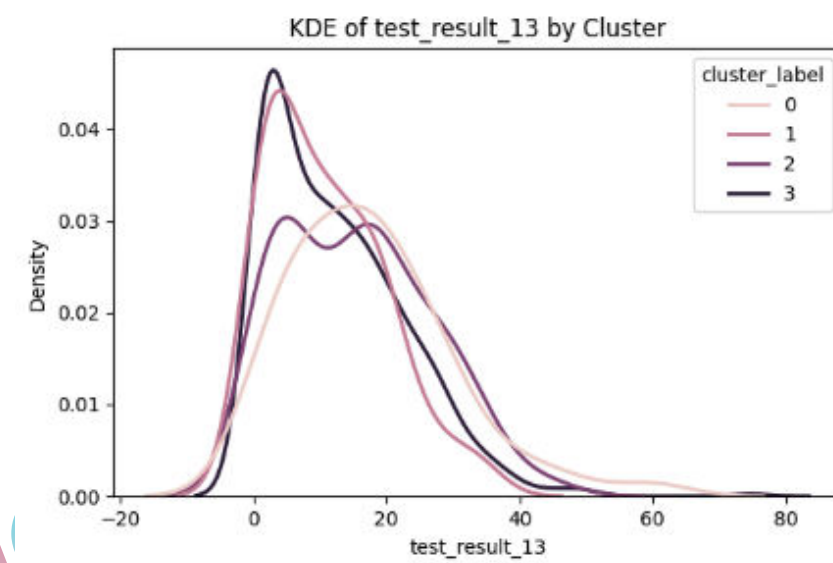
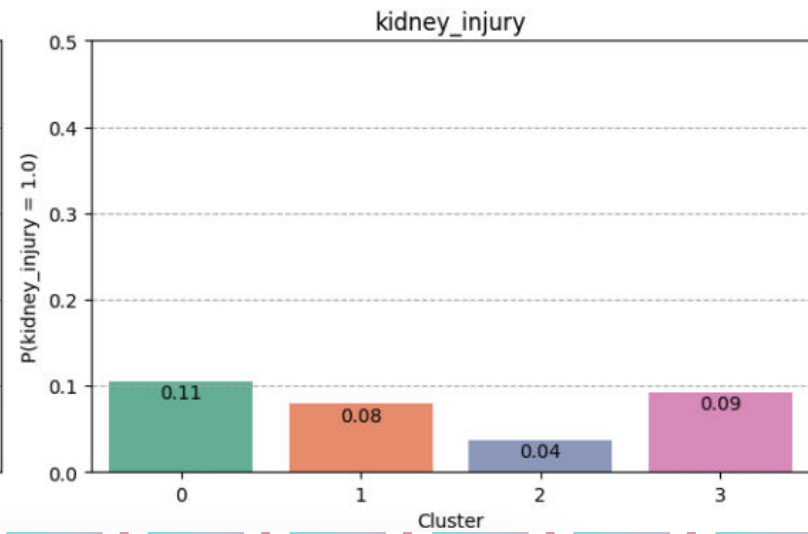
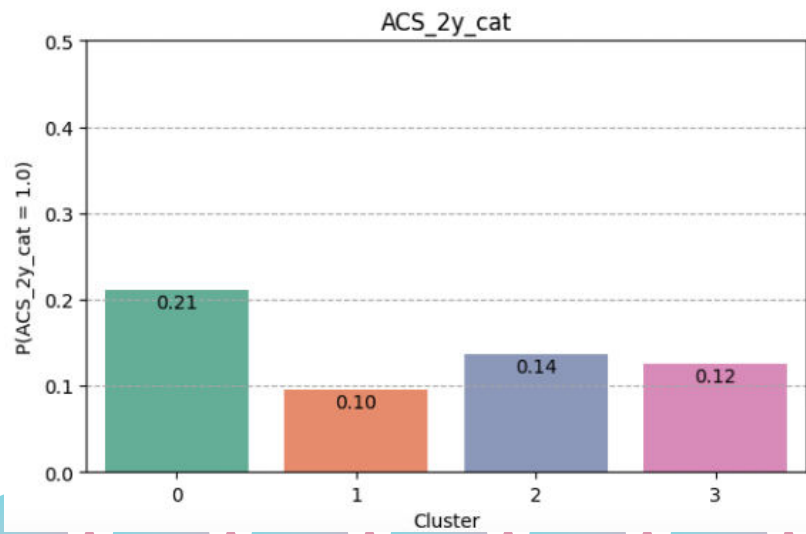
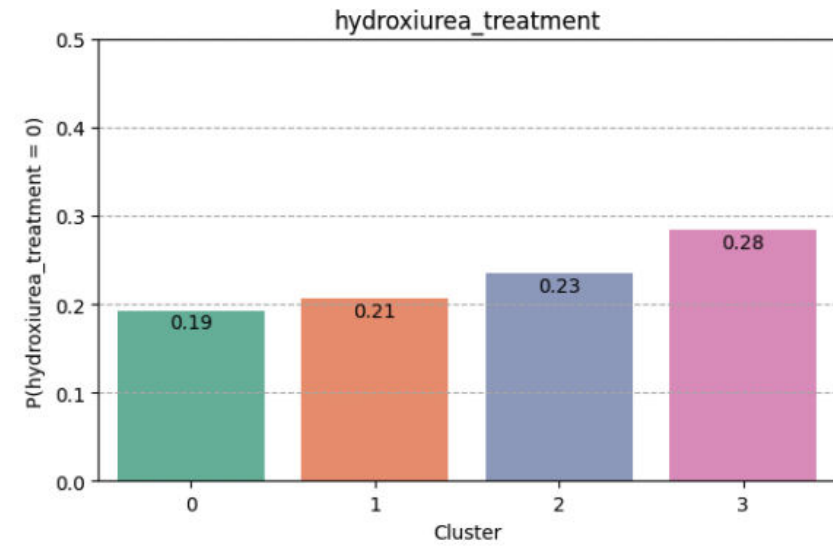
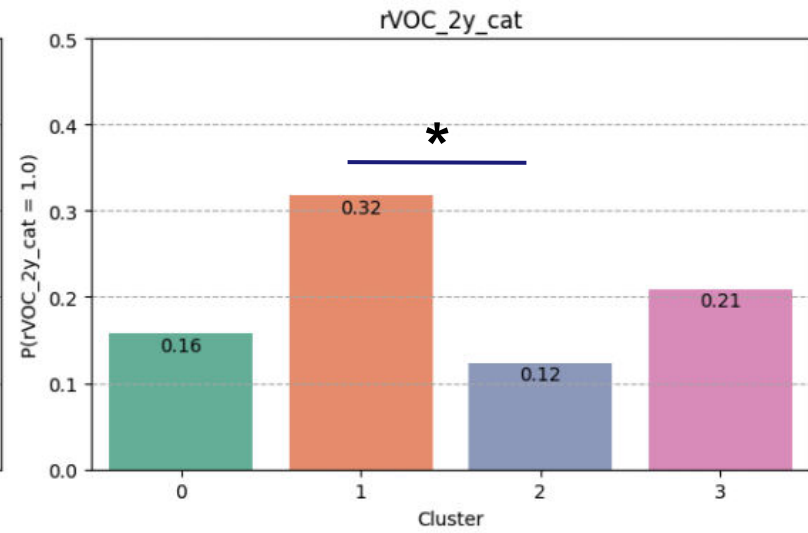
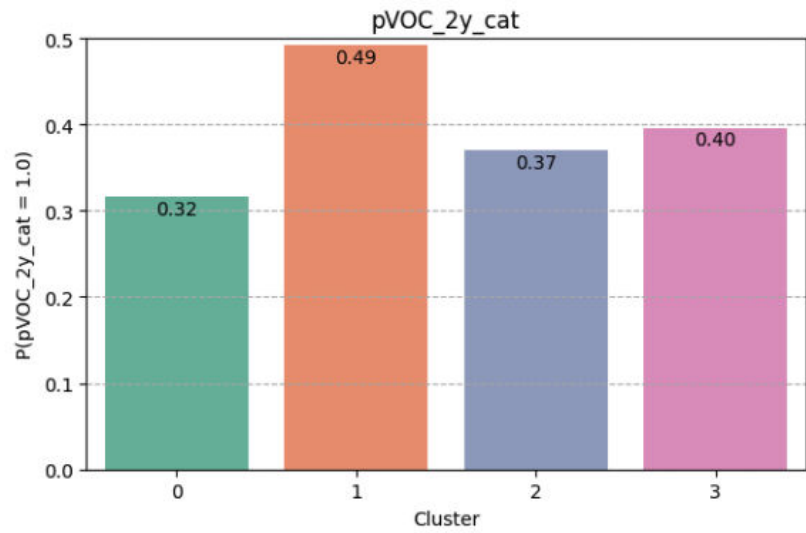
Signif. pvalues:

(2, 3) ***

(2, 1) ***

(2, 0) ***

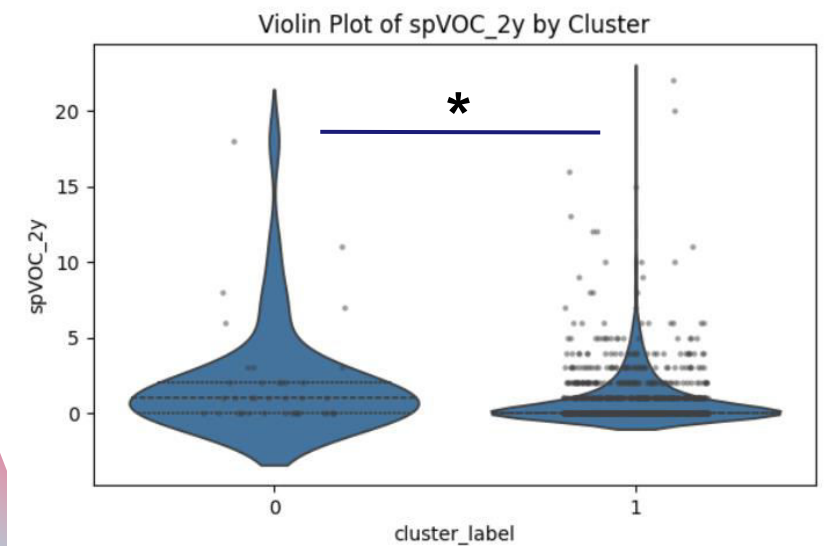
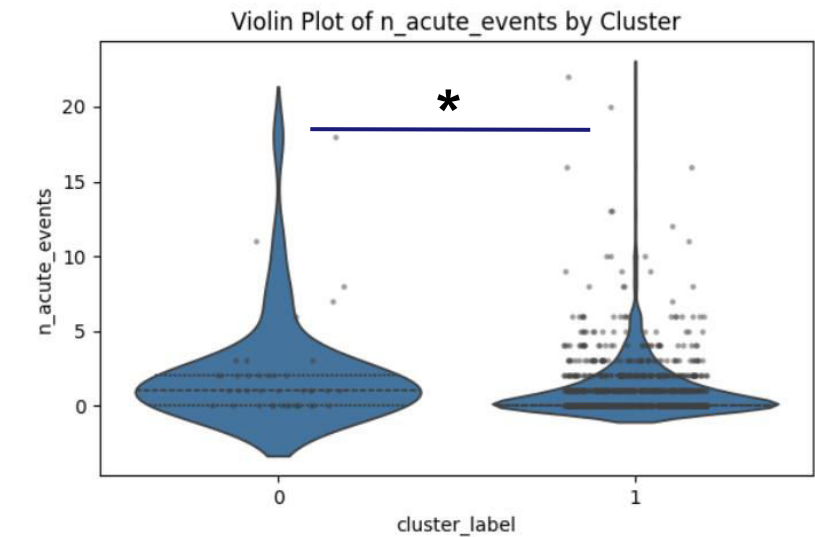
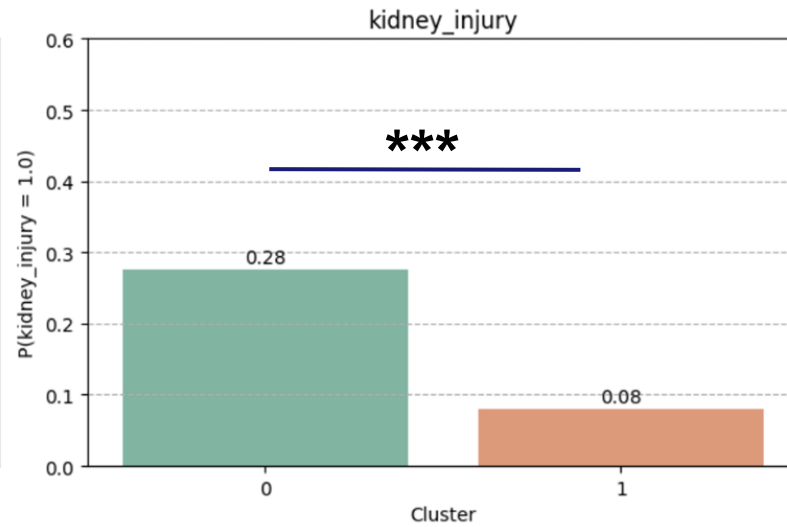
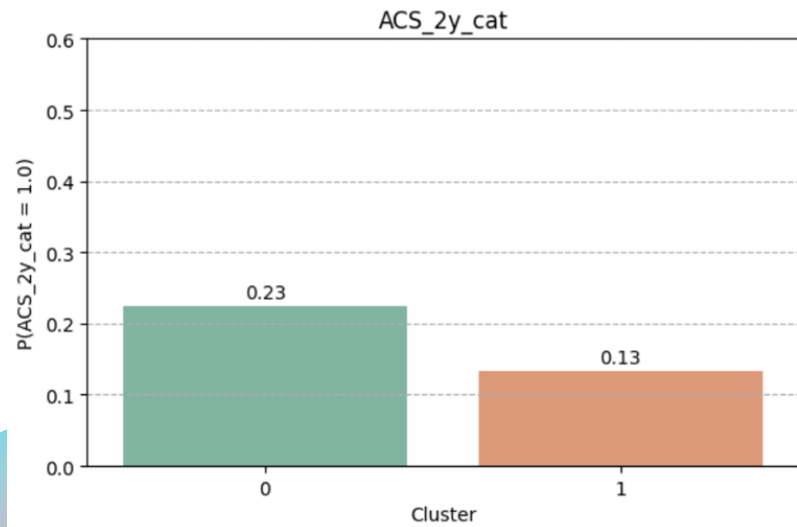
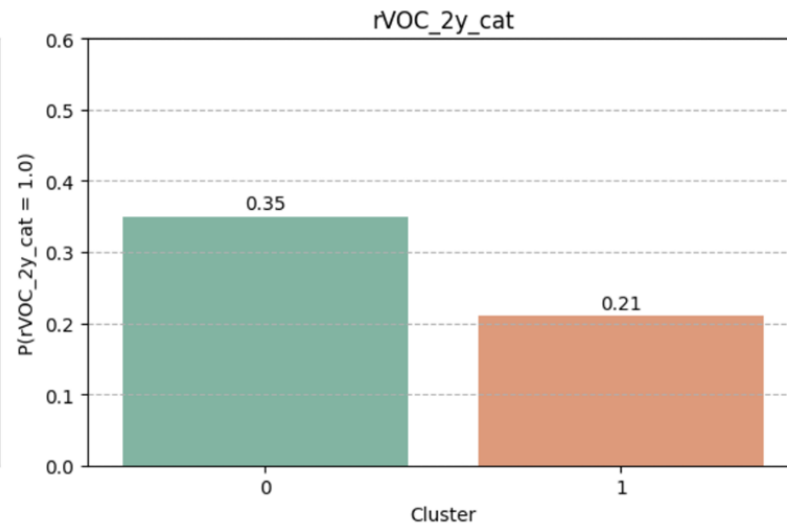
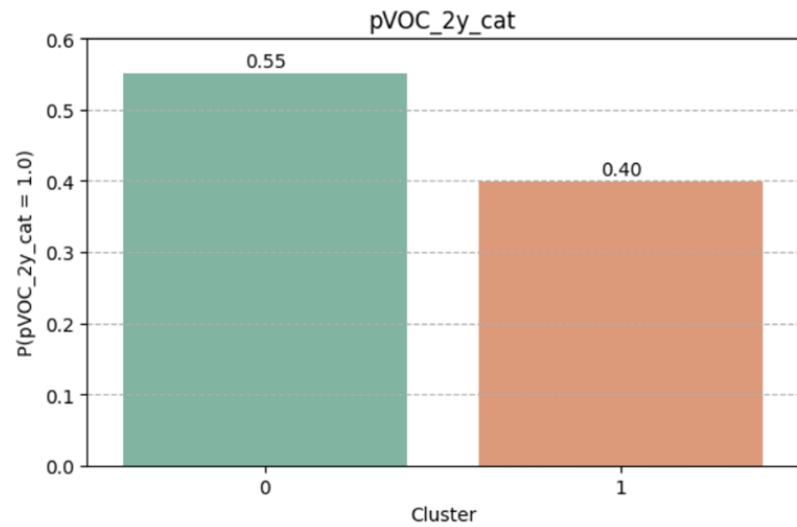
GWAS Clustering (945 samples): {0,0.5,1} mapping



Clustering on Metabolomics (on 961 samples)



Clustering on Metabolomics (on 961 samples)



Supervised model

Outcomes to predict:

- ❖ ACS (0/1)
- ❖ pVOC (0/1)
- ❖ spVOC (cont.)
- ❖ ACS (0/1)
- ❖ Kidney Injury (0/1)

How:

Train a unique Neural Network model that exploits at the same time:

- ❖ Clinical and laboratory test data
- ❖ Metabolomics (1800 metabolites)
- ❖ GWAS (800 SNPs selected)
- ❖ OxygenScan data

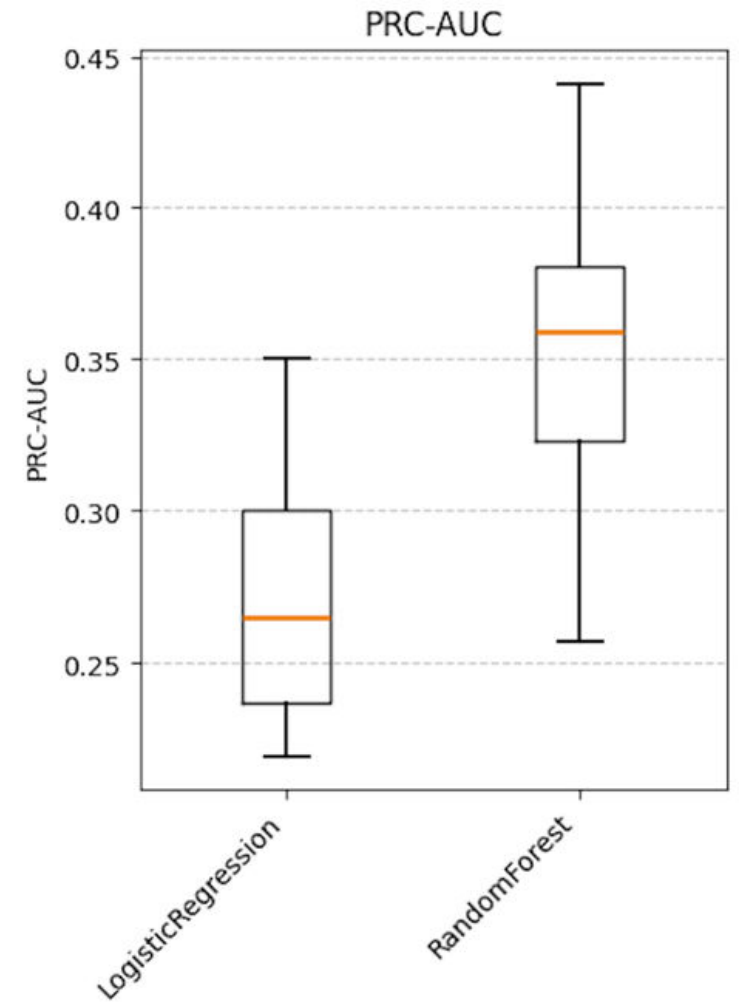
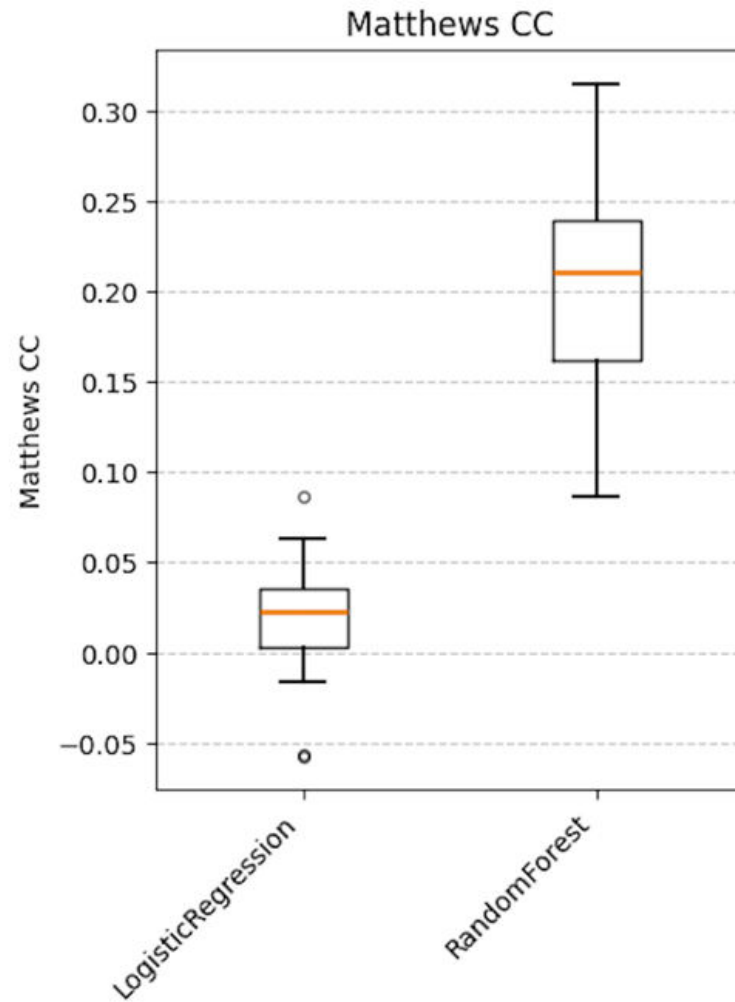
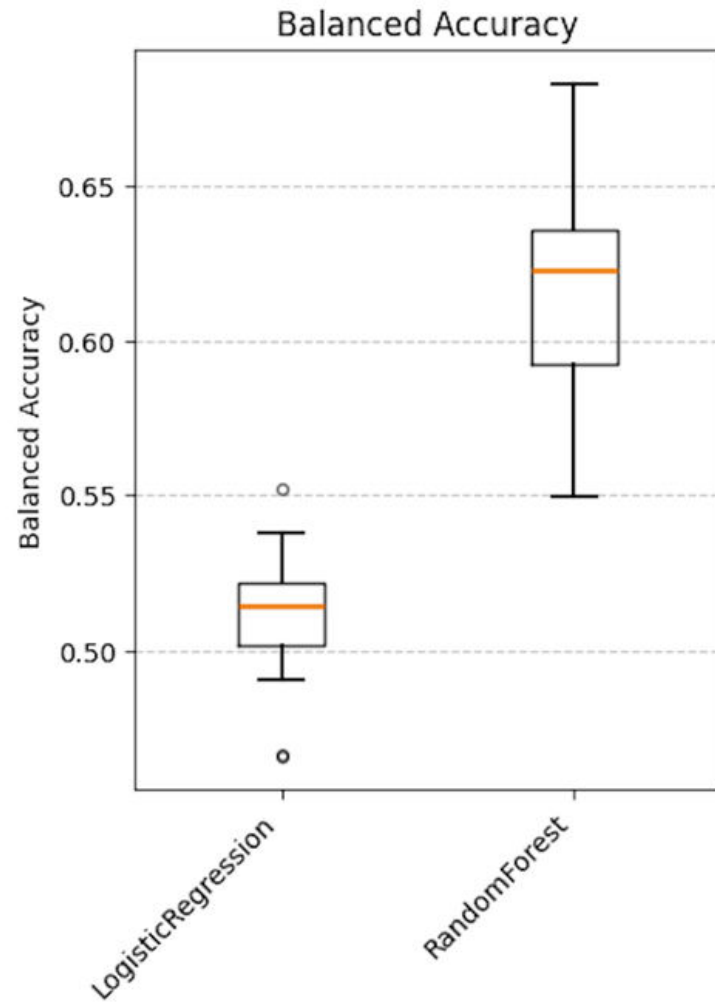


Supervised model: Clinical variables

gender	test_result_1	test_result_81	neuro_answer_3
first_hcp_contact_cum	test_result_2	test_result_82	neuro_answer_4
clin_diag_cum	test_result_3	test_result_23	priapism_2y_cat
age	test_result_4	test_result_83	splenic_seq_2y_cat
bmi	test_result_5	test_result_10	bones_complication_answer_1
oxygen_saturation_rate	test_result_6	test_result_62	bones_complication_answer_4
heart_rate	test_result_7	test_result_63	bones_complication_answer_5
diastolic_blood_pressue	test_result_8	test_result_64	bones_complication_answer_3
systolic_blood_pressure	test_result_9	test_result_65	endo_answer_2
hydroxiurea_treatment	test_result_11	test_result_66	lkd_answer_2
cumulative_years	test_result_13	test_result_84	lkd_answer_4
cumulative_years_age	test_result_69	test_result_86	lkd_answer_1
hydroxiurea_dose	test_result_70	test_result_29	lkd_answer_5
chelation_treat	test_result_71	test_result_88	lkd_answer_6
splenomegaly	test_result_72	test_result_87	vhd_answer_3
cholelithiasis	test_result_73	test_result_91	cpd_value_5
cholecystectomy	test_result_74	test_result_92_recalc	splenectomy
ab_test_id___1	test_result_24	test_result_89	Stroke
treatid___4	test_result_25	test_result_90_recalc	Cerebral0th
treatid___5	test_result_26	comorbiditiesid___1	
treatid___11	test_result_27	comorbiditiesid___2	sub_orpha_id
treatid___20	test_result_18	comorbiditiesid___4	HBA_Label
treatid___12	test_result_19	comorbiditiesid___14	HBB_Label
treatid___3	test_result_20	comorbiditiesid___22	
treatid___19	test_result_21	comorbiditiesid___19	
treatid___21	test_result_75	comorbiditiesid___29	
treatid___18			

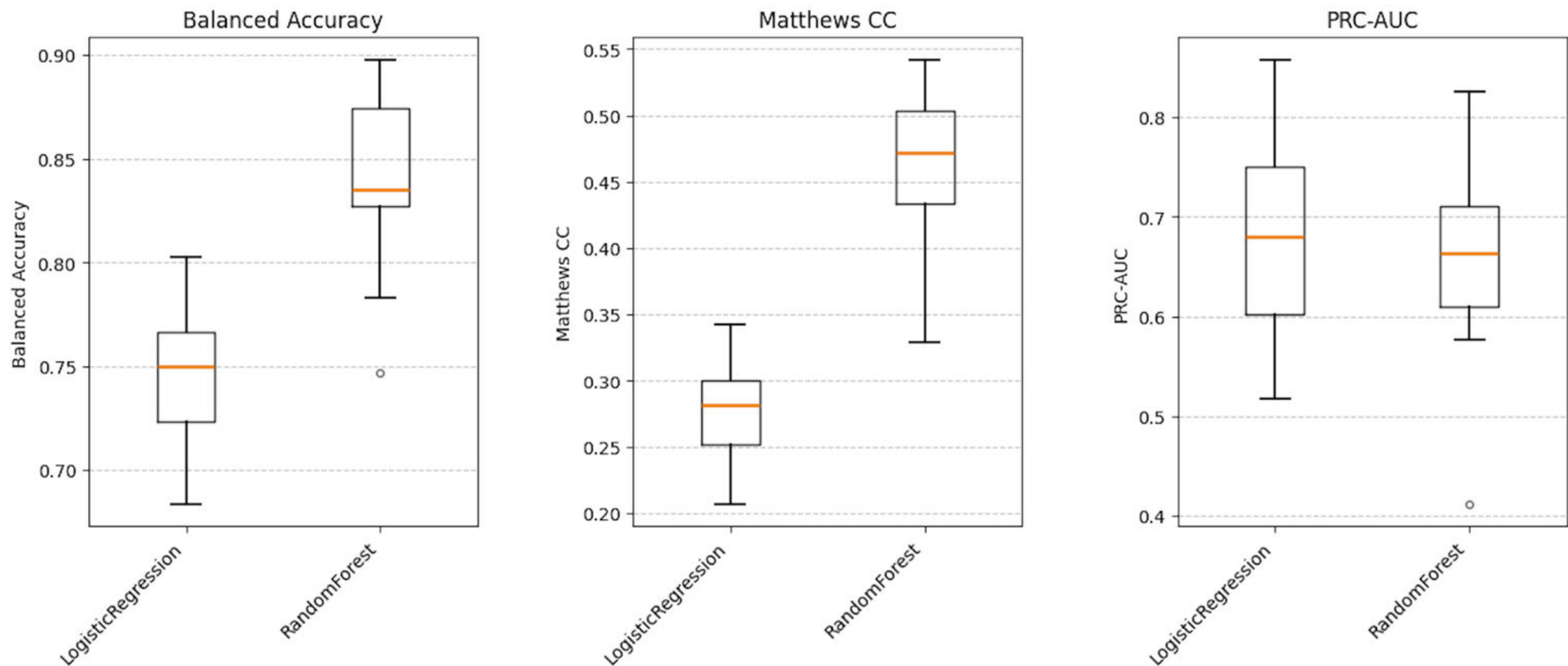
rVOC outcome - Baseline (only clinical) models

rVOC_2y_cat — Test-Set Distributions

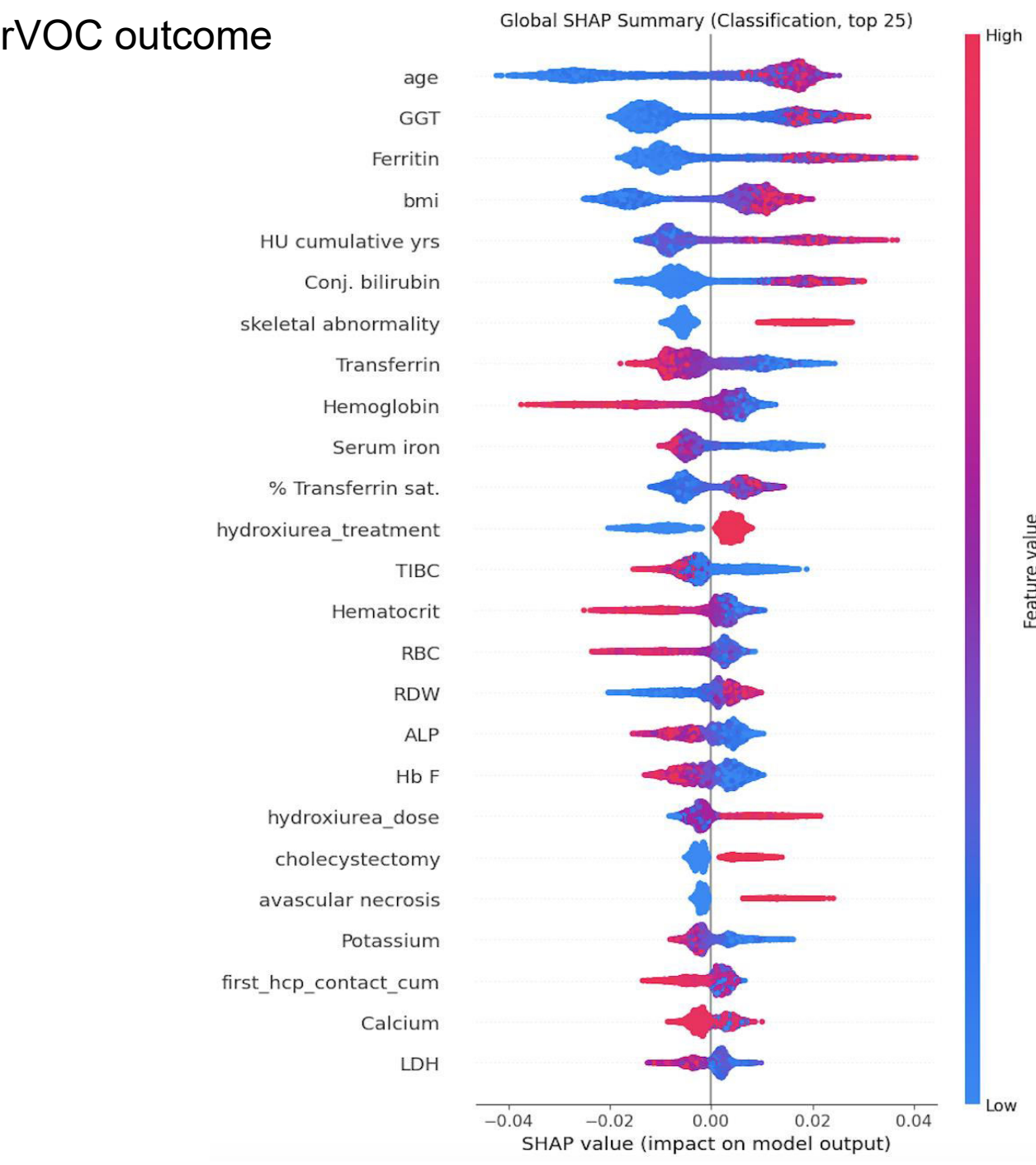


Kidney injury outcome - Baseline (only clinical) models

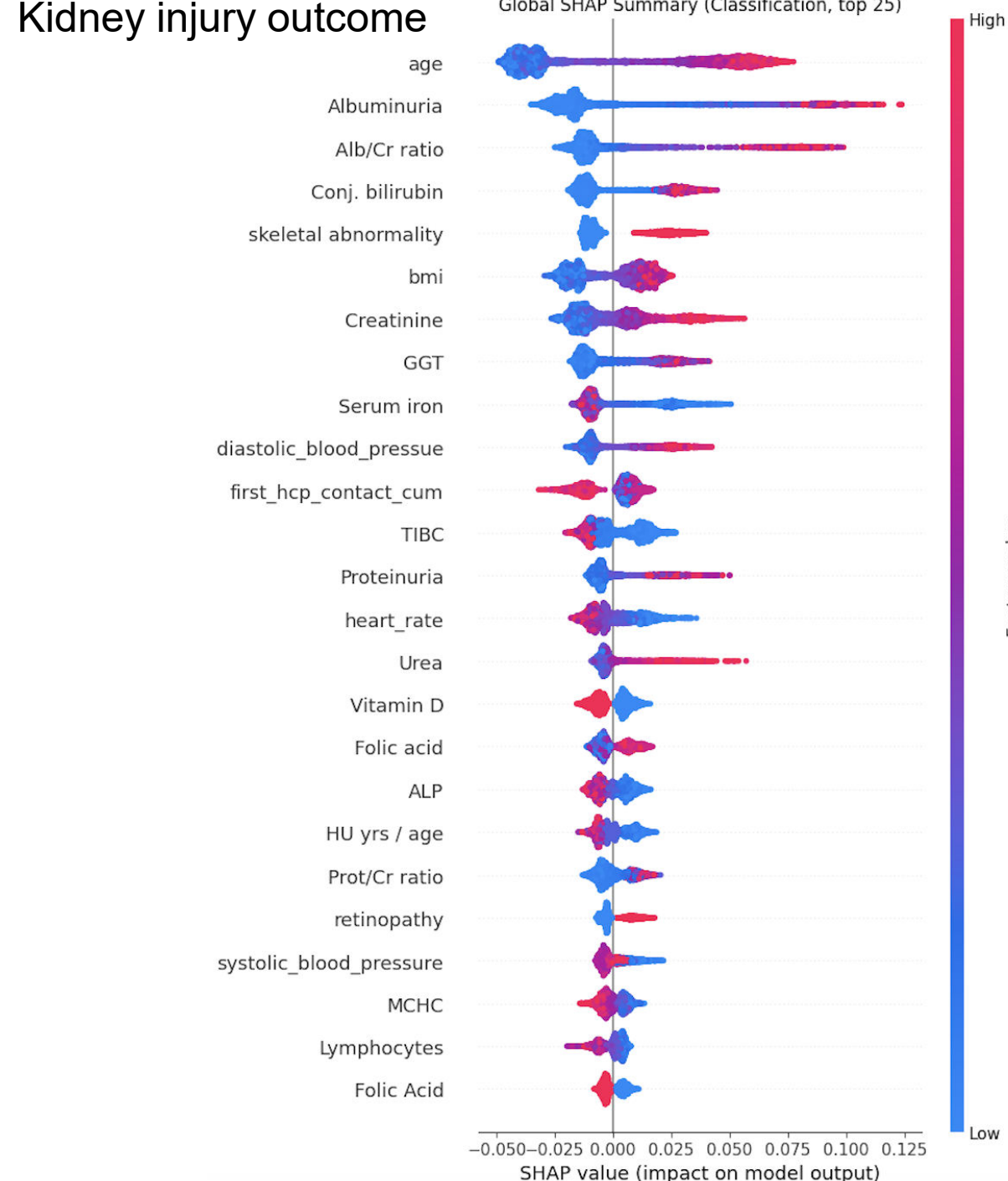
kidney_injury — Test-Set Distributions



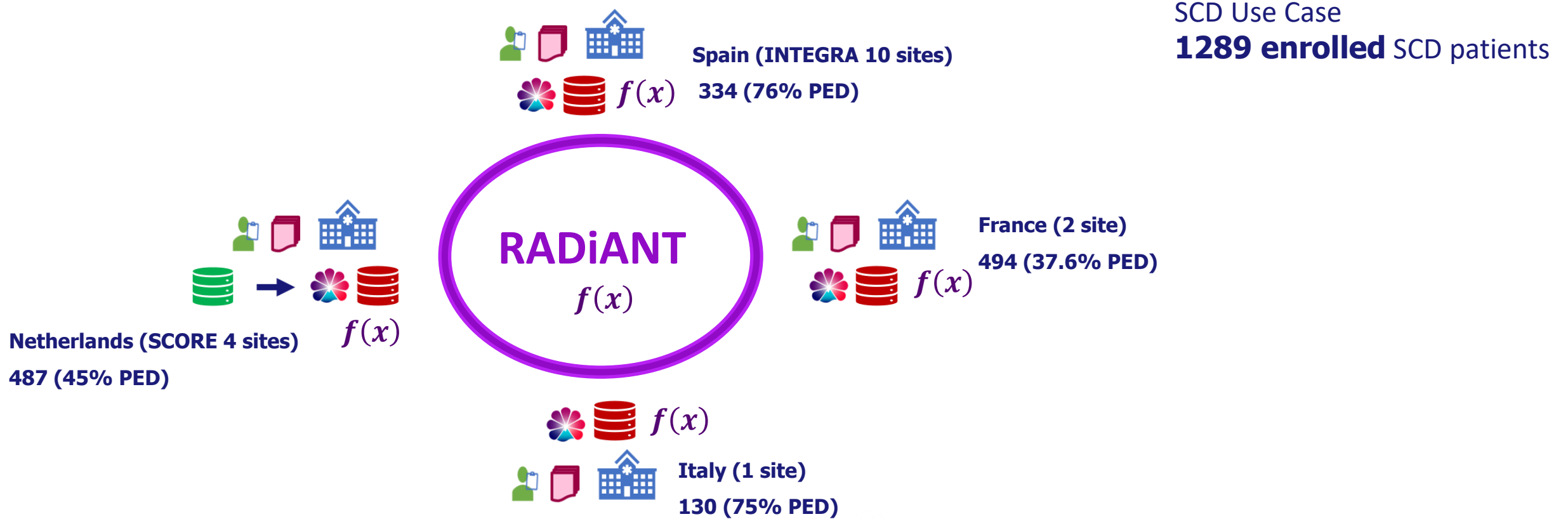
rVOC outcome



Kidney injury outcome

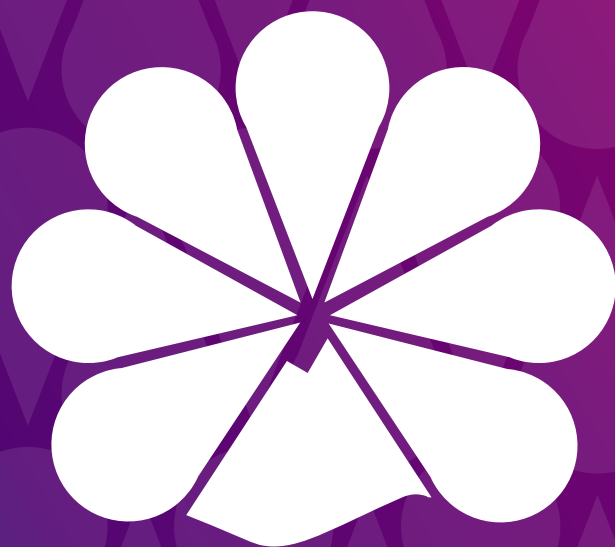


RADiANT



SCD Use Case

1289 enrolled SCD patients



RADeep
RADiANT

www.radeepnetwork.eu