

Linking and sharing multi-modal data with Knowledge Graphs

Application to intracranial aneurysms

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Comprendre les pathologies

cardio-vasculaires, métaboliques et respiratoires pour mieux les **prévenir**

Médecine translationnelle :

Patients ← → Recherche

Lien entre **fonction** biologique et séquences **génétiques** ?

Bio-informatique

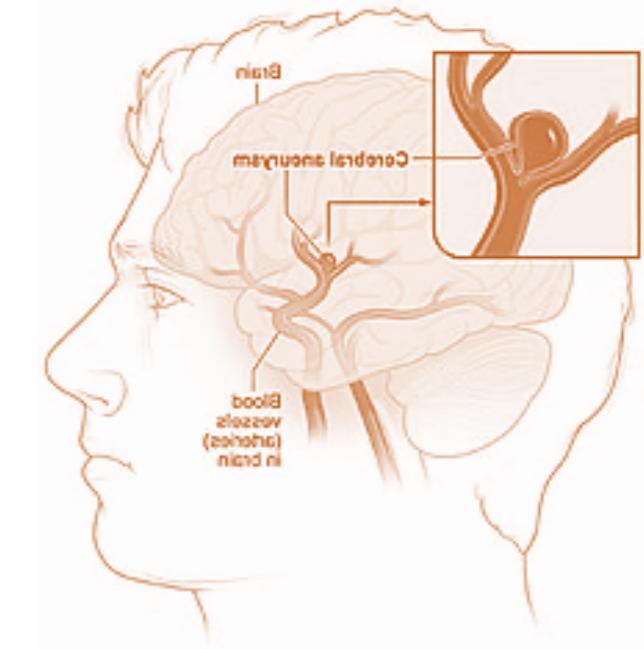
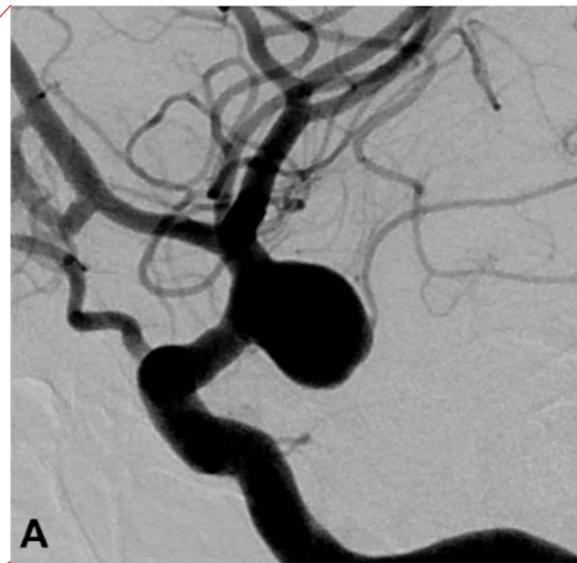
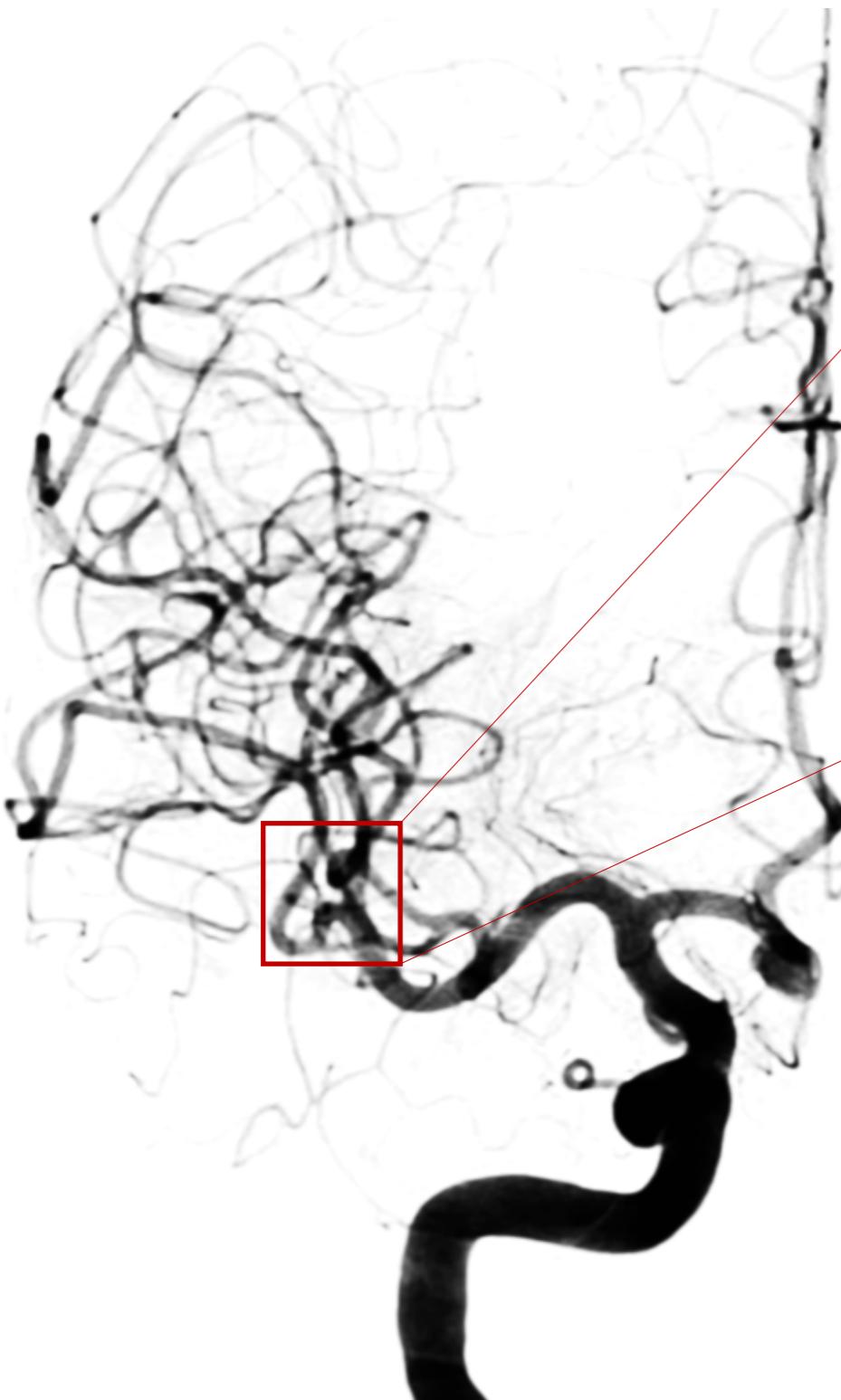
Données de santé en grand nombre

- ▶ **calcul haut-débit** : infrastructure distribuées
- ▶ **intégration de données** hétérogènes
- ▶ **modèles explicatifs / prédictifs (IA)**

données massives → nouvelles connaissances

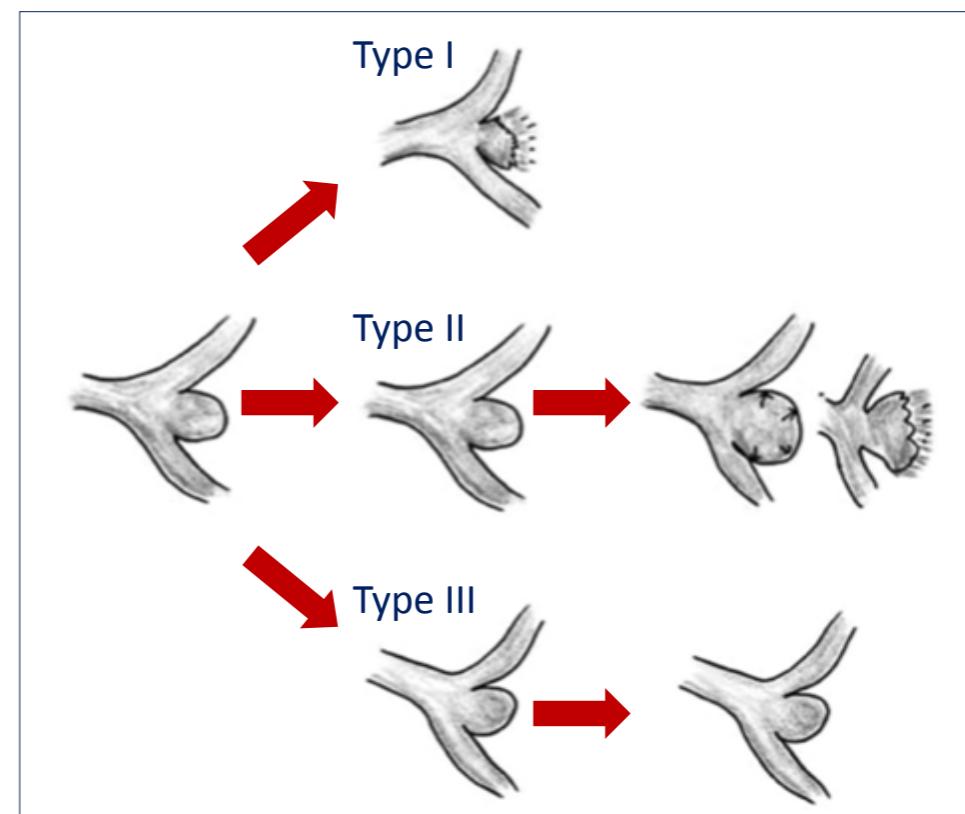
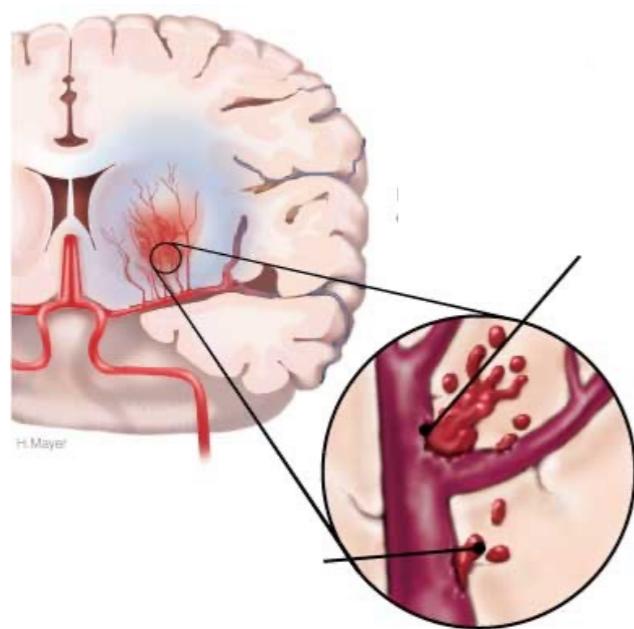
Intracranial aneurysms

Intracranial aneurysm (ICA)



- ▶ 3% of the general population
- ▶ unpredictable rupture
- ▶ 50% of death in case of rupture

ICA rupture / stroke



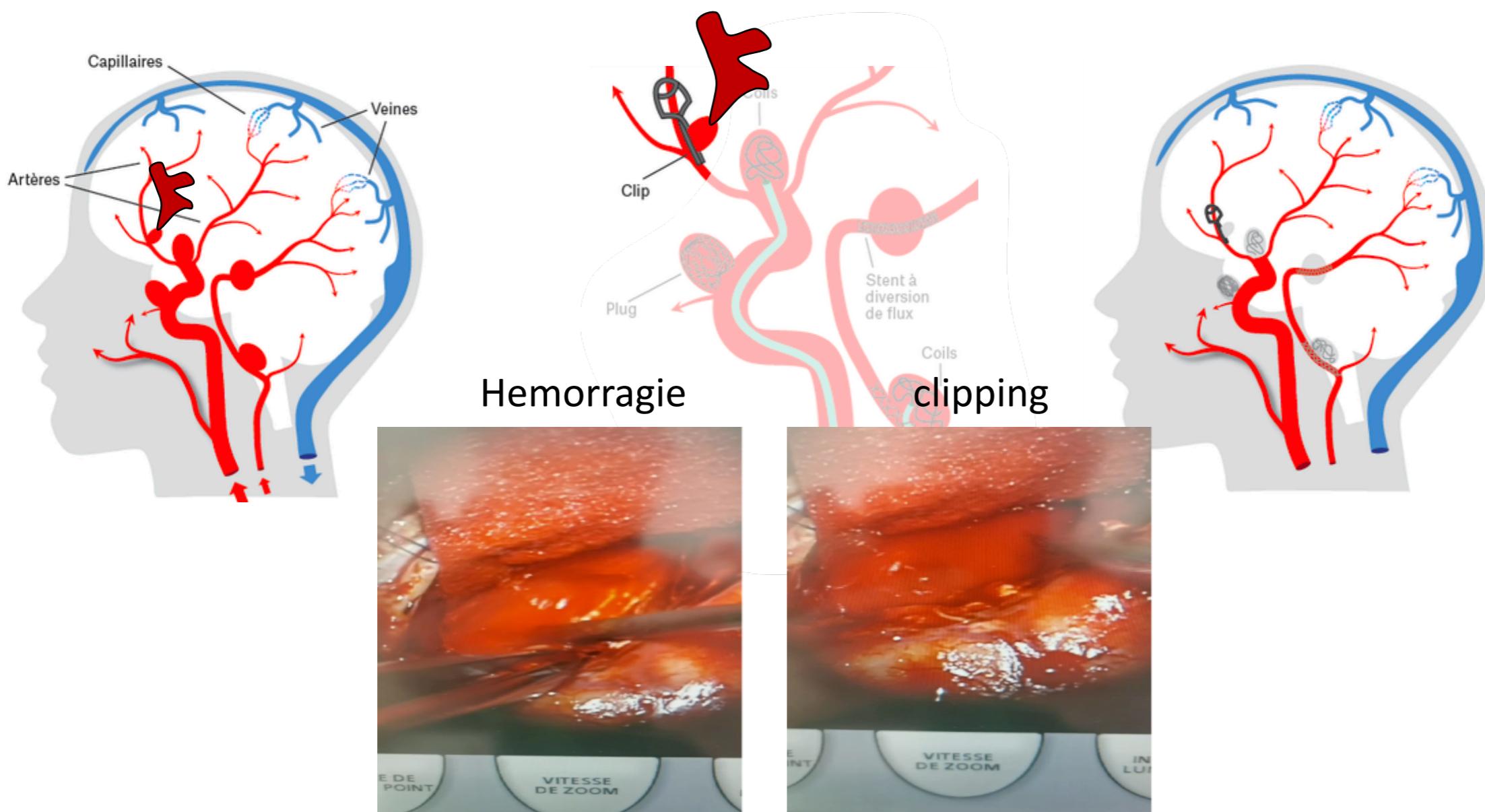
Risk factors

- ▶ smoking, alcohol consumption, hypertension, female sex, age
- ▶ familial history (RRx4)

In 10% of ICA cases = familial form of AIC with at least 2 first-degree relatives affected

How are they treated ?

Surgical clipping

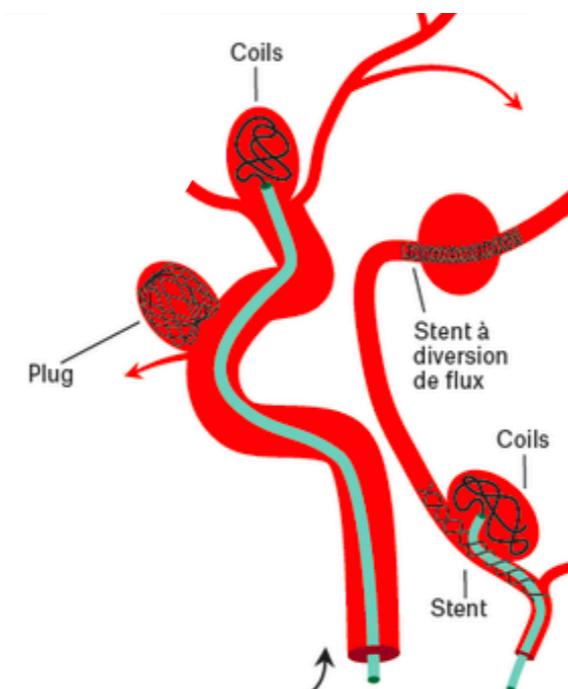
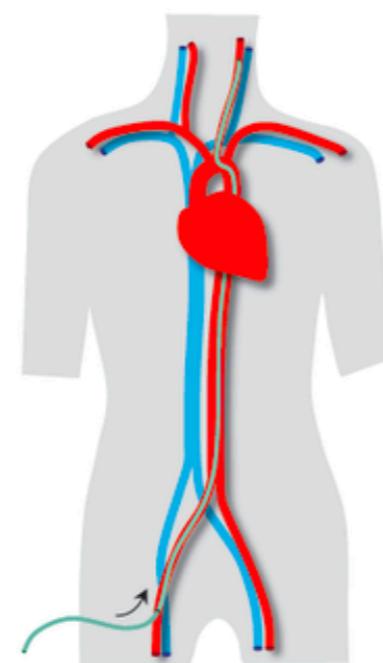
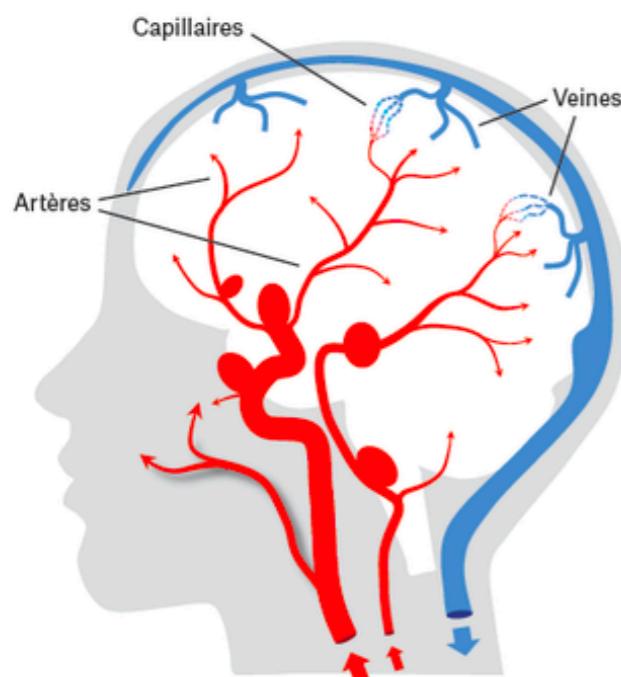


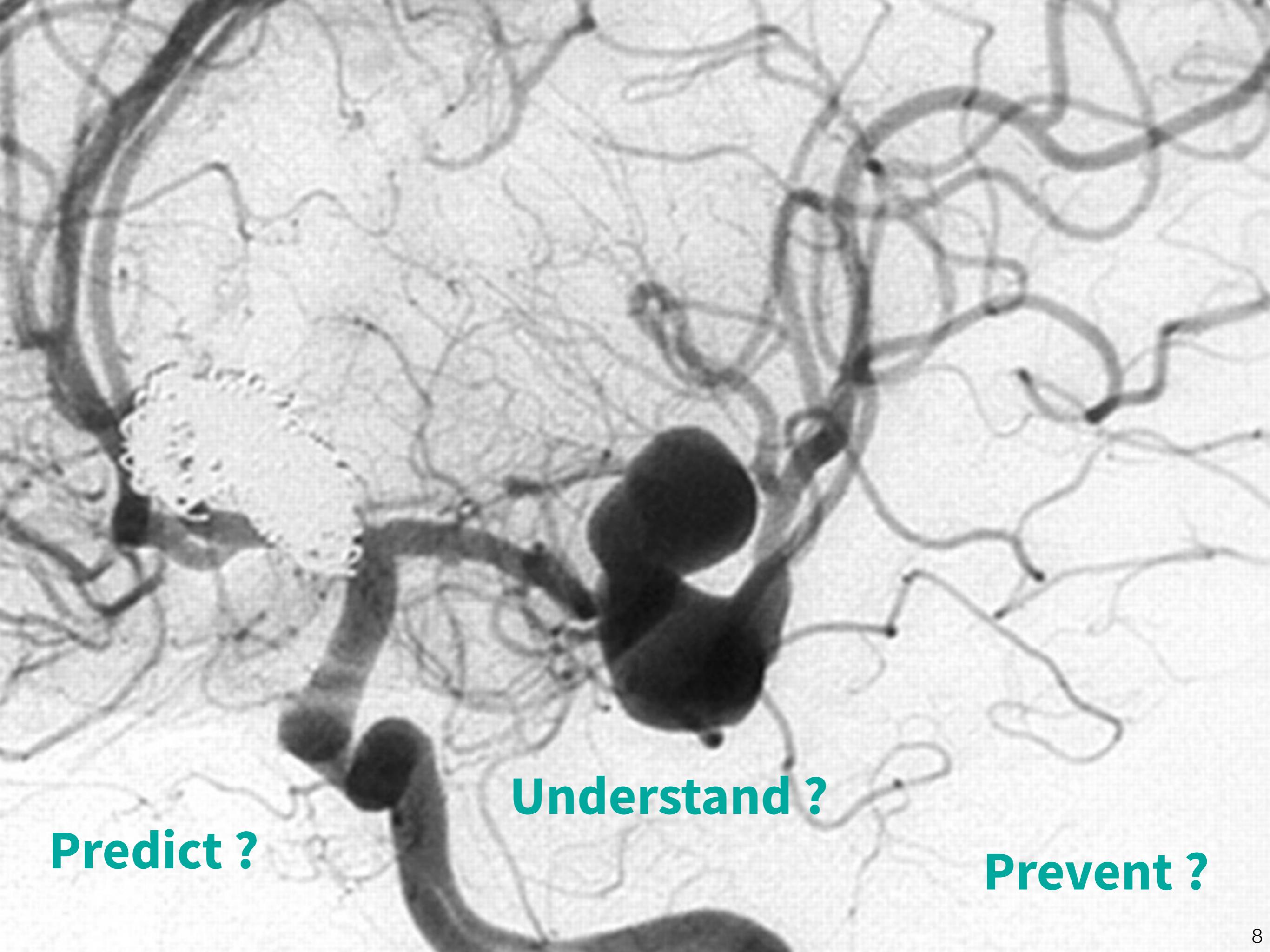
How are they treated ?

Regular MRI



+ endovascular surgery





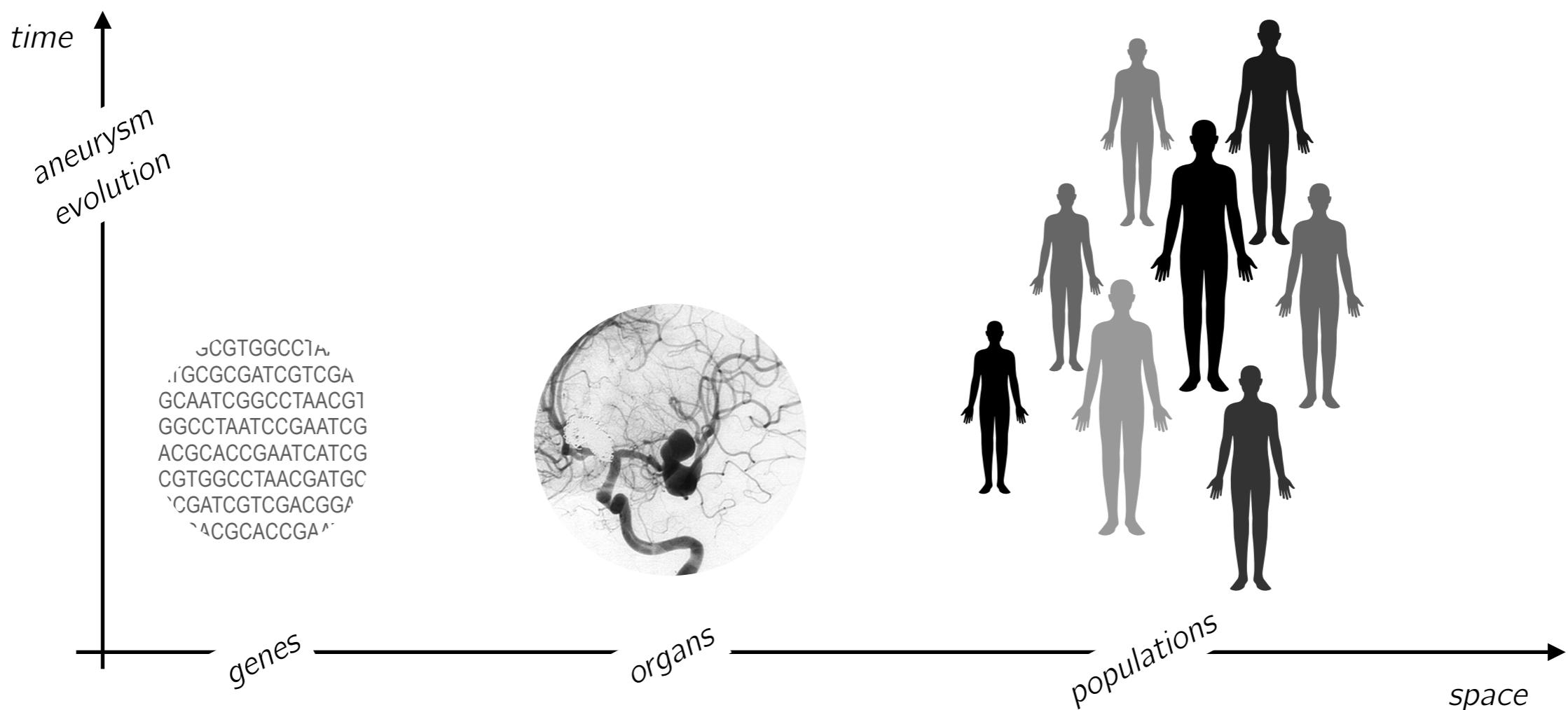
Predict ?

Understand ?

Prevent ?

Multi-factorial disease → multi-scale data

- ▶ Intracranial aneurysms: a complex & multifactorial disease
- ▶ Inter-disciplinary efforts needed for a better understanding of the pathology
- ▶ Specific data produced at very specific scales



Multi-centric study: 29 contributing centers



ICAN, UCAN, weCAN projects

+ recently funded eCAN RHU project



2023

4 000 ICA cases
TOF MRIs
800+ whole genomes



2024 +

> 3 000 patients per year



Multi-modal data analysis

ARTICLE

Using deep learning for an automatic detection and classification of the vascular bifurcations along the Circle of Willis

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Keywords:

Vascular bifurcations
Circle of Willis
Deep learning

ABSTRACT

Most of the intracranial aneurysms (ICAs) occur on a specific portion of the cerebral vascular tree named the Circle of Willis (CoW). More particularly, they mainly arise onto fifteen of the major arterial bifurcations constituting this circular structure. Hence, for an efficient and timely diagnosis it is critical to develop some methods being able to accurately recognize each Bifurcation of Interest (BoI). Indeed, an automatic extraction of the bifurcations presenting the higher risk of developing an ICA would offer the neuroradiologists a quick glance at the most alarming areas. Due to the recent efforts on Artificial Intelligence, Deep Learning turned out to be the best performing technology for many pattern recognition tasks. Moreover, various methods have been particularly designed for medical image analysis purposes. This study intends to assist the neuroradiologists to promptly locate any bifurcation presenting a high risk of ICA occurrence. It can be seen as a Computer Aided Diagnosis scheme, where the Artificial Intelligence facilitates the access to the regions of interest within the MRI. In this work, we propose a method for a fully automatic detection and recognition of the bifurcations of interest forming the Circle of Willis. Several neural networks architectures have been tested, and we thoroughly evaluate the bifurcation recognition rate.

Segmented vascular tree

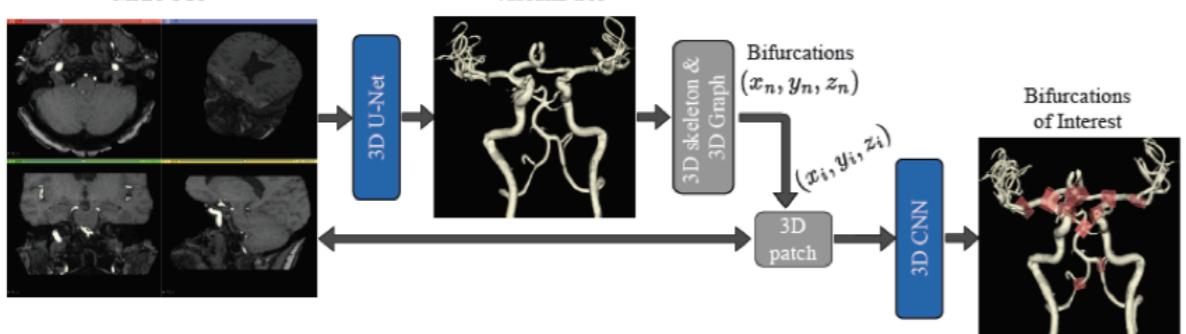


Fig. 1: General flowchart of the bifurcation recognition process.

Observational Study > J Neurol Neurosurg Psychiatry. 2021 Feb;92(2):122-128.

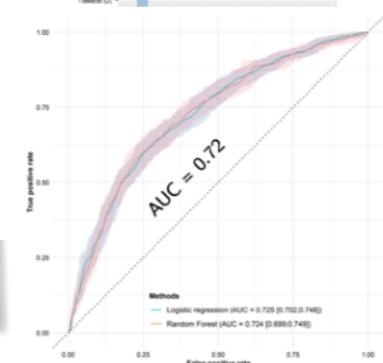
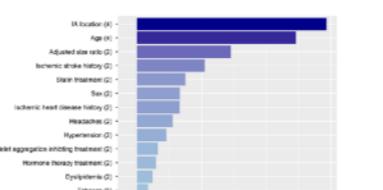
doi: 10.1136/jnnp-2020-324371. Epub 2020 Oct 23.

Location of intracranial aneurysms is the main factor associated with rupture in the ICAN population

Olivia Rousseau¹, Matilde Karakachoff¹, Alban Gaignard², Lise Bellanger³, Philippe Bijlenga⁴, Pacôme Constant Dit Beaufils¹, Vincent L'Allinec⁵, Olivier Levrier⁶, Pierre Aguetzaz⁷, Jean-Philippe Desilles⁸, Caterina Michelozzi⁹, Gaultier Marnat¹⁰, Anne-Clémence Vion², Gervaise Loirand², Hubert Desal¹¹, Richard Redon², Pierre-Antoine Gourraud¹, Romain Bourcier¹²; ICAN Investigators

Collaborators, Affiliations + expand

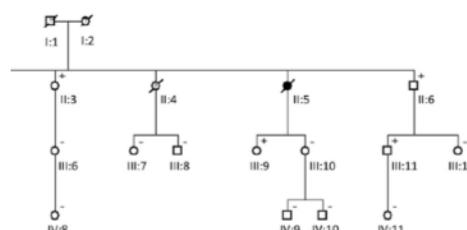
PMID: 33097563 DOI: 10.1136/jnnp-2020-324371



Rare Coding Variants in ANGPTL6 Are Associated with Familial Forms of Intracranial Aneurysm

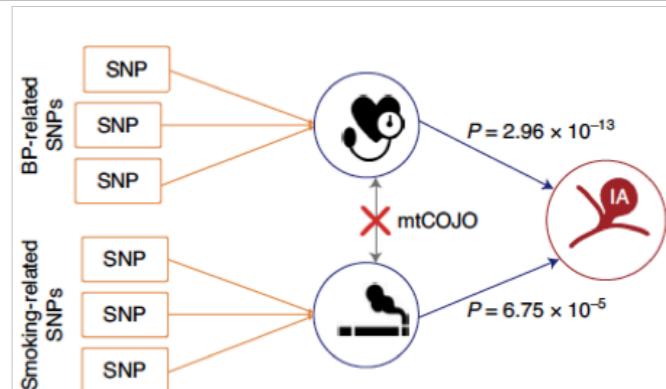
Romain Bourcier,^{1,2} Solena Le Scouarnec,¹ Stéphanie Bonnaud,^{1,3} Matilde Karakachoff,^{1,3} Emmanuelle Bourcereau,³ Sandrine Heurtelise-Chrétien,¹ Céline Menguy,¹ Christian Dina,^{1,3} Floriane Simonet,^{1,3} Alexis Moles,⁴ Cédric Lenoble,² Pierre Lindenbaum,¹ Stéphanie Chatel,^{1,3} Bertrand Isidor,⁵ Emmanuelle Génin,⁶ Jean-François Deleuze,⁷ Jean-Jacques Schott,^{1,3} Hervé Le Marec,^{1,3} ICAN Study Group, Gervaise Loirand,^{1,3,8} Hubert Desal,^{1,2,8,*} and Richard Redon^{1,3,8,*}

Intracranial aneurysms (ICAs) are acquired cerebrovascular abnormalities characterized by localized dilation and wall thinning in intracranial arteries, possibly leading to subarachnoid hemorrhage and severe outcome in case of rupture. Here, we identified one rare nonsense variant (*c.1378A>T*) in the last exon of *ANGPTL6* (Angiopoietin-Like 6)—which encodes a circulating pro-angiogenic factor mainly secreted from the liver—shared by the four tested affected members of a large pedigree with multiple IA-affected case subjects. We showed a 50% reduction of *ANGPTL6* serum concentration in individuals heterozygous for the *c.1378A>T* allele (*p.Lys460Ter*) compared to relatives homozygous for the normal allele, probably due to the non-secretion of the truncated protein produced by the *c.1378A>T* transcript. Considering *ANGPTL6* in a series of 61 additional index case subjects with familial IA identified three other



Filtering steps	Remaining variants	
	Individuals : III-1	III-5
Functional variants	25,674	23,456
MAF < 0,1% in ExAC (NFE)	456	436
Shared by III.1 & III.5	29	
In IBD haplotypes	10	
Shared by all affected	8	
Predicted LOF	1 (<i>ANGPTL6</i>)	

Genome-wide association study of intracranial aneurysms identifies 17 risk loci and genetic overlap with clinical risk factors



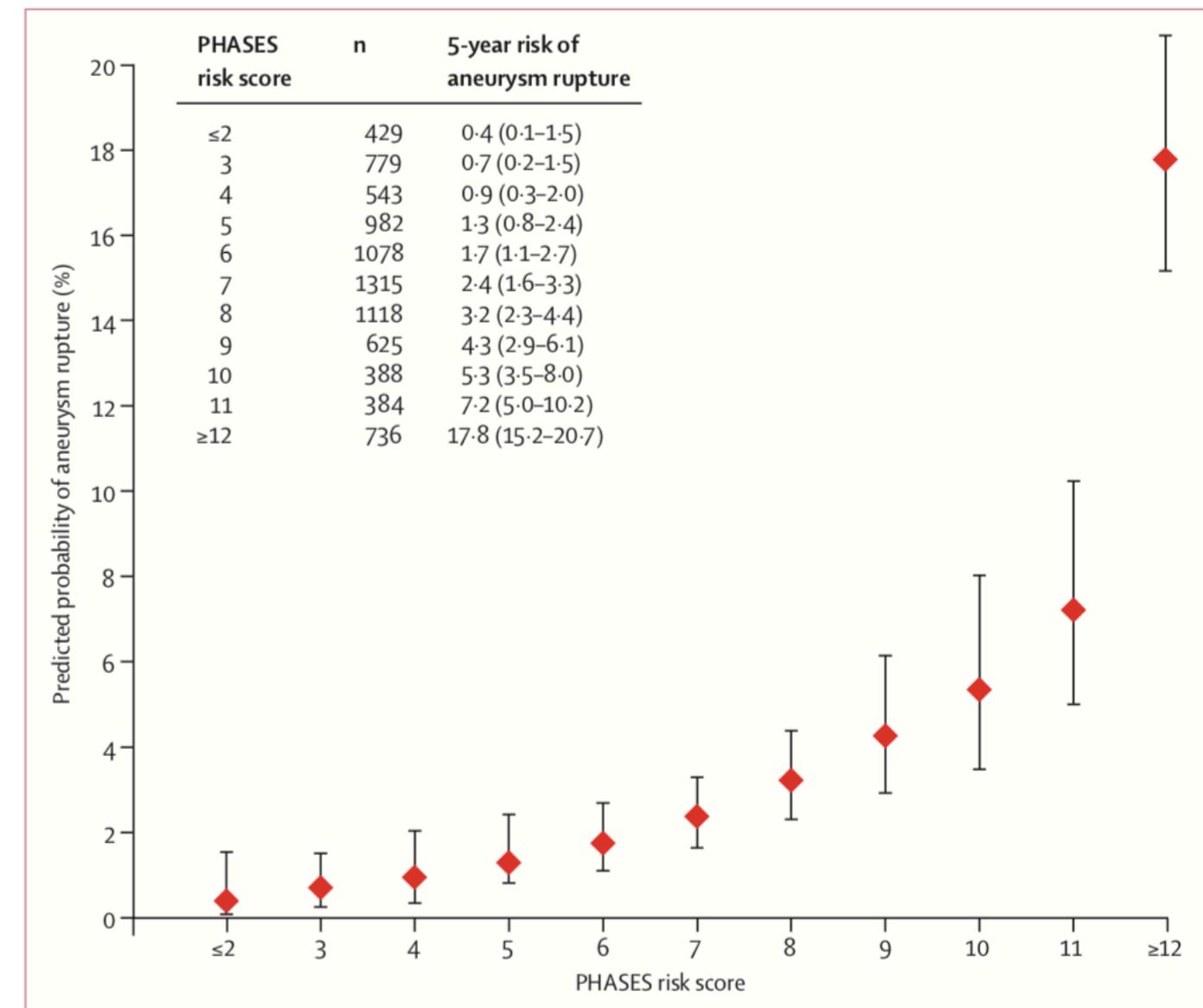


Olivia Rousseau

Is it possible to **predict IA
rupture** from **clinical data** ?

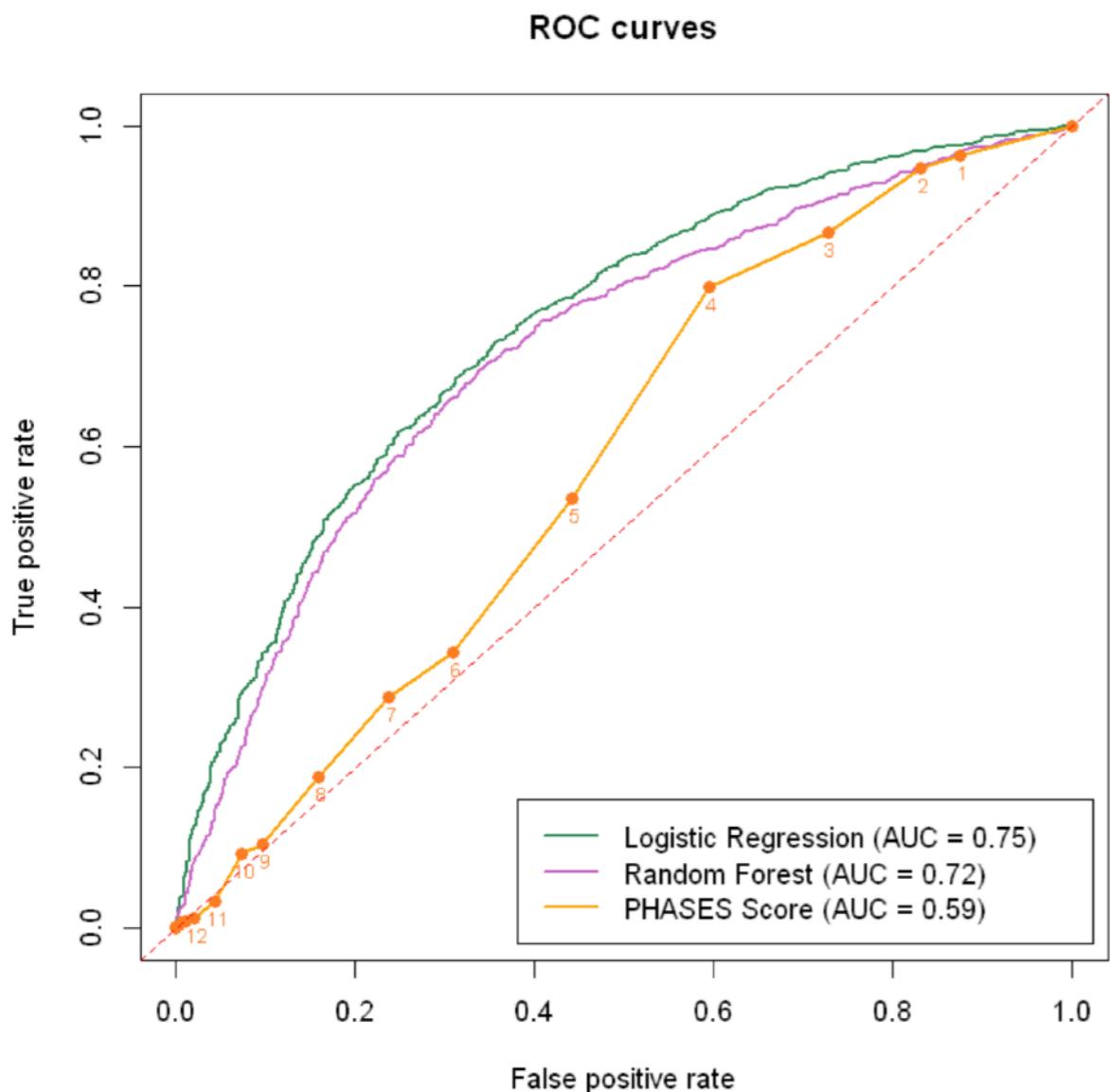
PHASES score

PHASES aneurysm risk score	Points
(P) Population	
North American, European (other than Finnish)	0
Japanese	3
Finnish	5
(H) Hypertension	
No	0
Yes	1
(A) Age	
<70 years	0
≥70 years	1
(S) Size of aneurysm	
<7.0 mm	0
7.0–9.9 mm	3
10.0–19.9 mm	6
≥20 mm	10
(E) Earlier SAH from another aneurysm	
No	0
Yes	1
(S) Site of aneurysm	
ICA	0
MCA	2
ACA/Pcom/posterior	4



Greving et. al. 2014 : Development of the PHASES score for prediction of risk of rupture of intracranial aneurysms: a pooled analysis of six prospective cohort studies.

Applying PHASES score to ICAN clinical data



- Unfortunately, poor prediction accuracy for PHASES (0.59)
- Better prediction accuracy with state-of-the-art Machine Learning / Statistical methods + other variables
- **Predictions are still failing in around 25% of cases ...**

integrating imaging or genomic markers ?

Intracranial aneurysms, current challenges

Complex and **multi-scale** biological phenomena involved :

- Impact of life habits ? Image-based evaluation ?
- Any genetic variant specific to image-based phenotypes ?
- Mirror aneurysms patient sub-grouping ? (Vincent Lallinec)

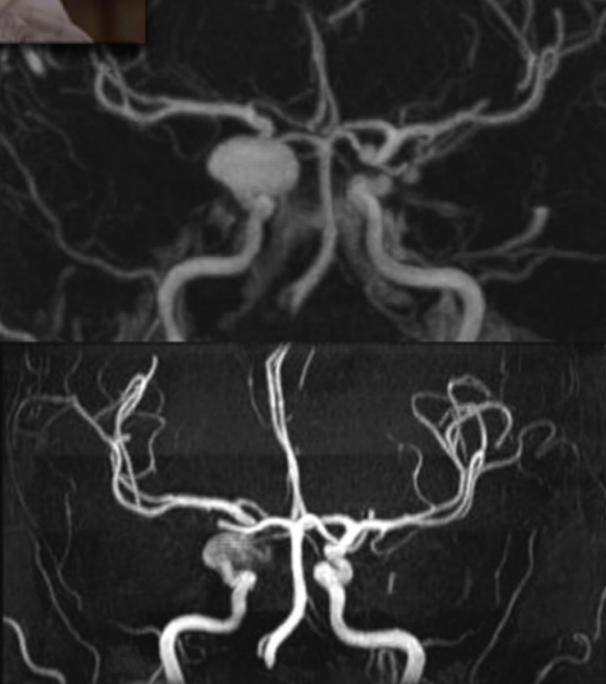
Needs for **jointly** exploiting genomic data, MR imaging and clinical observations ...

... to better **understand** and **predict** evolution schemes.

#CHROM	POS	ID	REF	ALT	QUAL	FILTER	INFO	FORMAT	39458_2001	39458_2003	39458_2005
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39458_CD12099		39458_CD12100	39458_CD12252	39458_CD12350	39458_CD12380	39458	CD12406	39458_CD12466	39458_CD12480	39458_CD12480	39458_CD12480
39458_CD12501		39458_CD12586	39458_CD12587	39458_CD12588	39458_CD12704	39458	CD13016	39458_CD13018	39458_CD13178	39458_CD13178	39458_CD13178
39458_CD13363											
	1	752566	rs3094315	A	G	.	.	PR	GT	./.	./.
	./.	./.	./.	./.	0/1	0/0	0/1	0/1	0/1	0/0	0/1
	0/1	0/0	0/0	0/1	0/1	1/1					
	1	754105	rs12184325	C	.	.	.	PR	GT	./.	./.
	./.	./.	./.	./.	0/0	0/0	0/0	0/0	0/0	0/0	0/0
	0/0	0/0	0/0	0/0	0/0	0/0	0/0	0/0	0/0	0/0	0/0
	1	754182	rs3131969	G	A	.	.	PR	GT	./.	./.
	./.	./.	./.	./.	0/1	0/0	0/1	0/1	0/1	0/1	0/1
	0/1	0/0	0/0	0/1	0/1	1/1					
	1	758626	rs3131954	T	.	.	.	PR	GT	./.	./.
	./.	./.	./.	./.	0/0	0/0	0/0	0/0	0/0	0/0	0/0
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	./.	./.	./.	./.	0/0	0/0	0/0	0/0	0/0	0/0	0/0
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	1	798801	rs12132517	G	A	.	.	PR	GT	./.	./.
	./.	./.	./.	./.	0/1	0/0	0/0	0/1	0/0	0/1	0/1
	0/1	0/0	0/0	0/0	0/0	0/0	0/0	0/0	0/0	0/0	0/1
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	./.	./.	./.	./.	0/1	0/0	0/0	0/1	0/1	0/1	0/1
	0/1	0/0	0/0	0/0	0/1	0/0	0/0	0/1	0/1	0/1	0/1



Ante attein	circonstances d'	consommation c'	quantité alcool p	status phenotyp	nb d'anévrismes	date du 1er diag	AIC 1 Grand axe	AIC 1 Localstatik	AIC 2 Grand axe	AIC 2 Localstatik
Porteur	Sangage > 3ans	96.6	certain	1.0	1/2/16	7.0	5			
Débutage familial	Toboggane usif	277.2	certain	1.0	3/2/16	4.0	10			
Porteur	Sangage > 3 ans	109.2	certain	3.0	12/28/15	6.0	4	14.0	4	
Ruturale AIC	Jenais fami	0.0	certain	2.0	4/0/16	7.0	M.D.A.Ghe	5.0	M.C.A.P	
Ruturale AIC	Sangage > 3ans	certain	4.0	1/1/16			18			
Porteur	Jenais fami	35.2	certain	1.0	1/2/16	5.0	21			
Porteur	Sangage < 3ans	50.4	certain	3.0	9/2/15	5.0	18	6.0	118	
Ruturale AIC	Jenais fami	certain	1.0	8/1/15			118			
Ruturale AIC	Jenais fami	certain	1.0	1/2/16	9.0	107				
Ruturale AIC	Toboggane usif	certain	1.0				105			
Porteur	Sangage > 3ans	58.0	certain	1.0	1/1/10	2.0	10			
Ruturale AIC	Jenais fami	certain	1.0				10			
Porteur	Jenais fami	46.2	certain	1.0			8.0	10		
Porteur	Sangage > 3ans	certain	2.0				8.0	18	5.0	10
Porteur	Jenais fami	certain	1.0	8/1/16	2.0		104			
Débutage familial	Jenais fami	certain	1.0	1/1/16			118			
Porteur	Toboggane usif	certain	1.0				20			
Ruturale AIC	Toboggane usif	25.2	certain	4.0	9/2/18	7.0	17	4.0	104	105
Ruturale AIC	Toboggane usif	398.0	certain	1.0			132			
Ruturale AIC	Jenais fami	certain	3.0	1/1/10	3.0	10	4.0	10	3.0	10.0
Ruturale AIC	Toboggane usif	certain	2.0	6/23/05	12.0	10	3.0	100		
Débutage familial	Sangage > 3 ans	378.0	certain	1.0	1/1/16	2.0	18			
Ruturale AIC	Jenais fami	certain	1.0	2/1/12	2.0	6				
Ruturale AIC	Jenais fami	certain	1.0	1/1/13	6.0	6				
Ruturale AIC	Toboggane usif	84.0	certain	3.0	1/1/16	4.0	10	2.0	118	100
Ruturale AIC	Jenais fami	certain	2.0	3/1/16	5.0	118	1.5	18		
Débutage familial	Sangage > 3ans	certain	1.0				18			
Ruturale AIC	Toboggane usif	0.0	certain	3.0	6/14/16	2.0	18	5.0	21	121.0
Ruturale AIC	Sangage > 3ans	12.6	certain	3.0	11/1/14	6.0	10	2.0	16	119.0
Ruturale AIC	Sangage > 3ans	25.2	certain	1.0	6/2/16	2.0	10			
Porteur	Jenais fami	certain	1.0				125/16			
Ruturale AIC	Sangage > 3 ans	470.4	certain	2.0	3/2/15	2.0	10	3.0	12	
Porteur	Toboggane usif	12.6	certain	1.0	1/1/18	8.0	10			
Porteur	Toboggane usif	0.0	certain	2.0	7/1/15	11.0	118	5.5	18	
Porteur	Jenais fami	certain	1.0	1/2/16	7.0	10				
Ruturale AIC	Jenais fami	0.0	certain	4.0			18	4.0	108	4.0
Ruturale AIC	Jenais fami	0.0	certain	1.0			12	2.5	18	
Porteur	Jenais fami	0.0	certain	1.0			723/12	5.0	132	
Ruturale AIC	Jenais fami	0.0	certain	2.0	4/24/15	0.5	105	13.0	5	
Porteur	Toboggane usif	42.0	certain	1.0	1/1/15	7.0	105			
Débutage familial	Sangage > 3 ans	21.0	certain	1.0	6/23/16	8.0	109			
Porteur	Toboggane usif	11.76	certain	2.0	6/3/16	2.0	5	3.0	185	
Ruturale AIC	Sangage > 3 ans	35.28	certain	1.0	8/4/18	5.0	111			
Porteur	Sangage > 3ans	0.0	certain	1.0	4/1/18	8.5	121			
Ruturale AIC	Jenais fami	11.78	certain	1.0	2/1/14	8.5	10			
Ruturale AIC	Sangage > 3ans	35.28	certain	1.0	6/9/15	6.7	10			
Porteur	Jenais fami	11.78	certain	1.0	2/22/16	6.0	111			
Débutage familial	Toboggane usif	11.76	certain	1.0	4/1/16	9.0	107			
Porteur	Sangage > 3ans	200.29	certain	2.0	1/1/14	4.4	17	5.2	10	



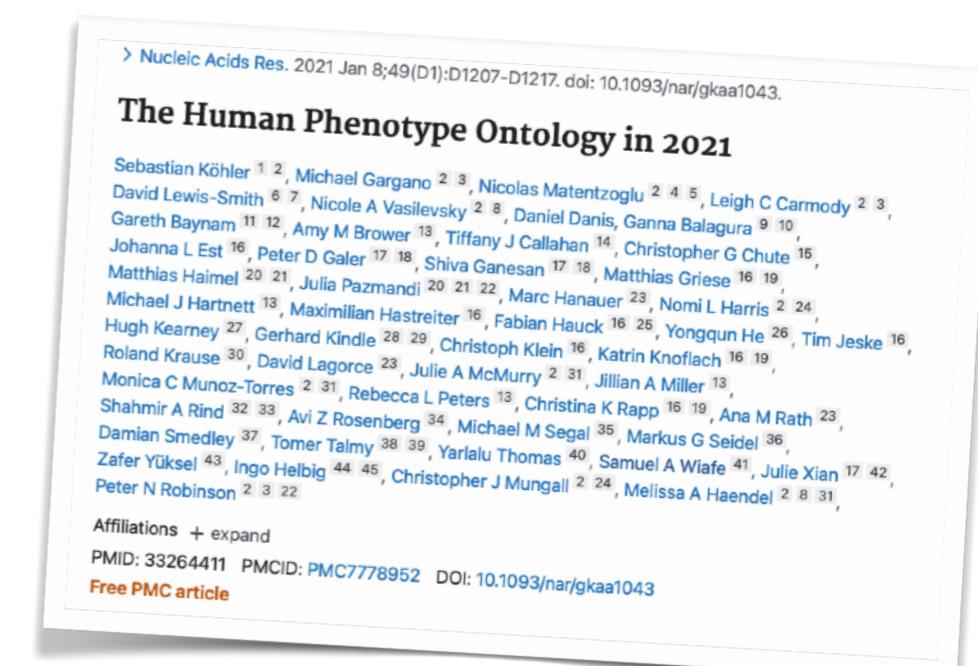
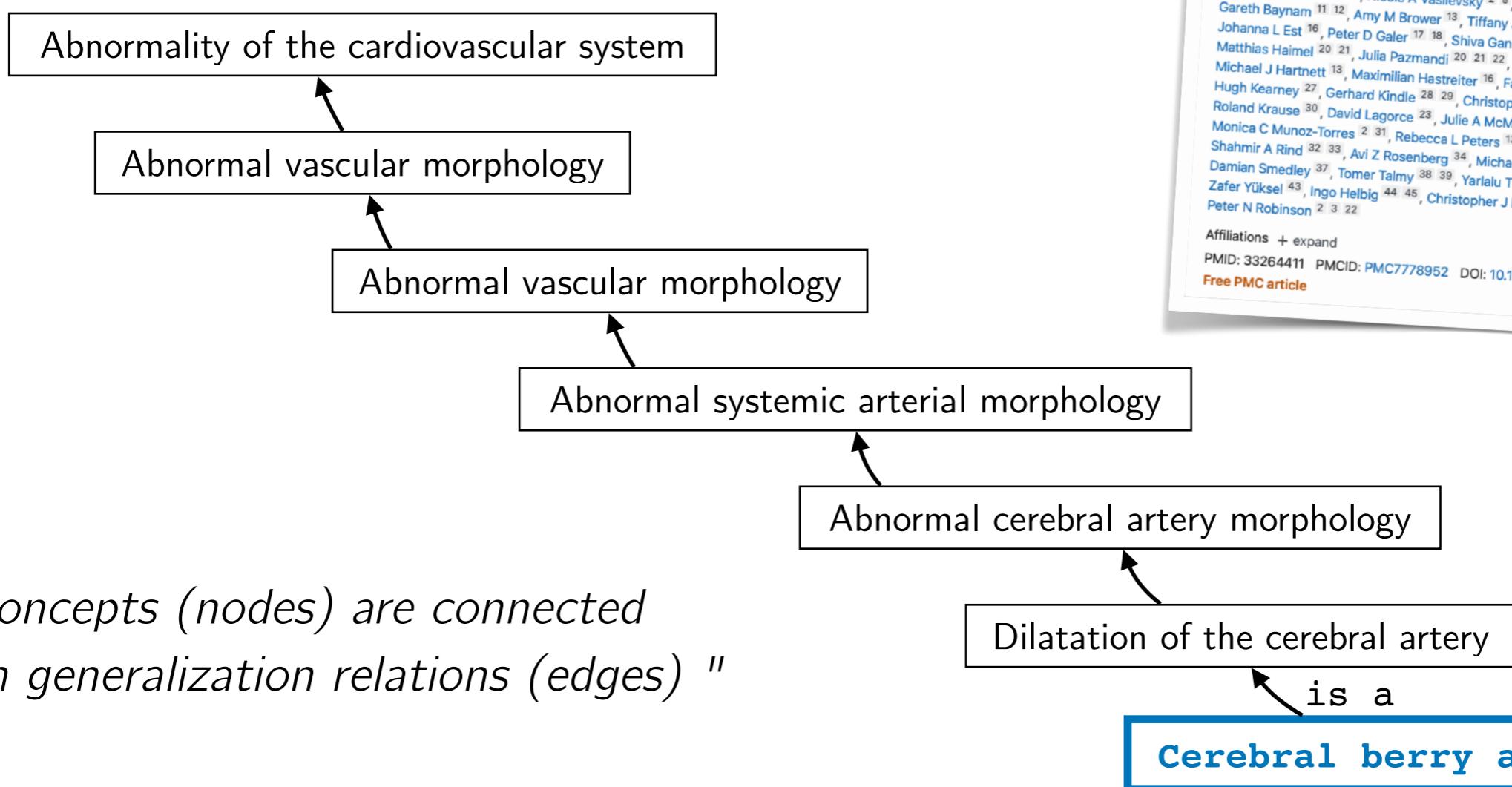
How to jointly analyse
these diverse datasets ?

① what is observed with data ?

Computational ontology

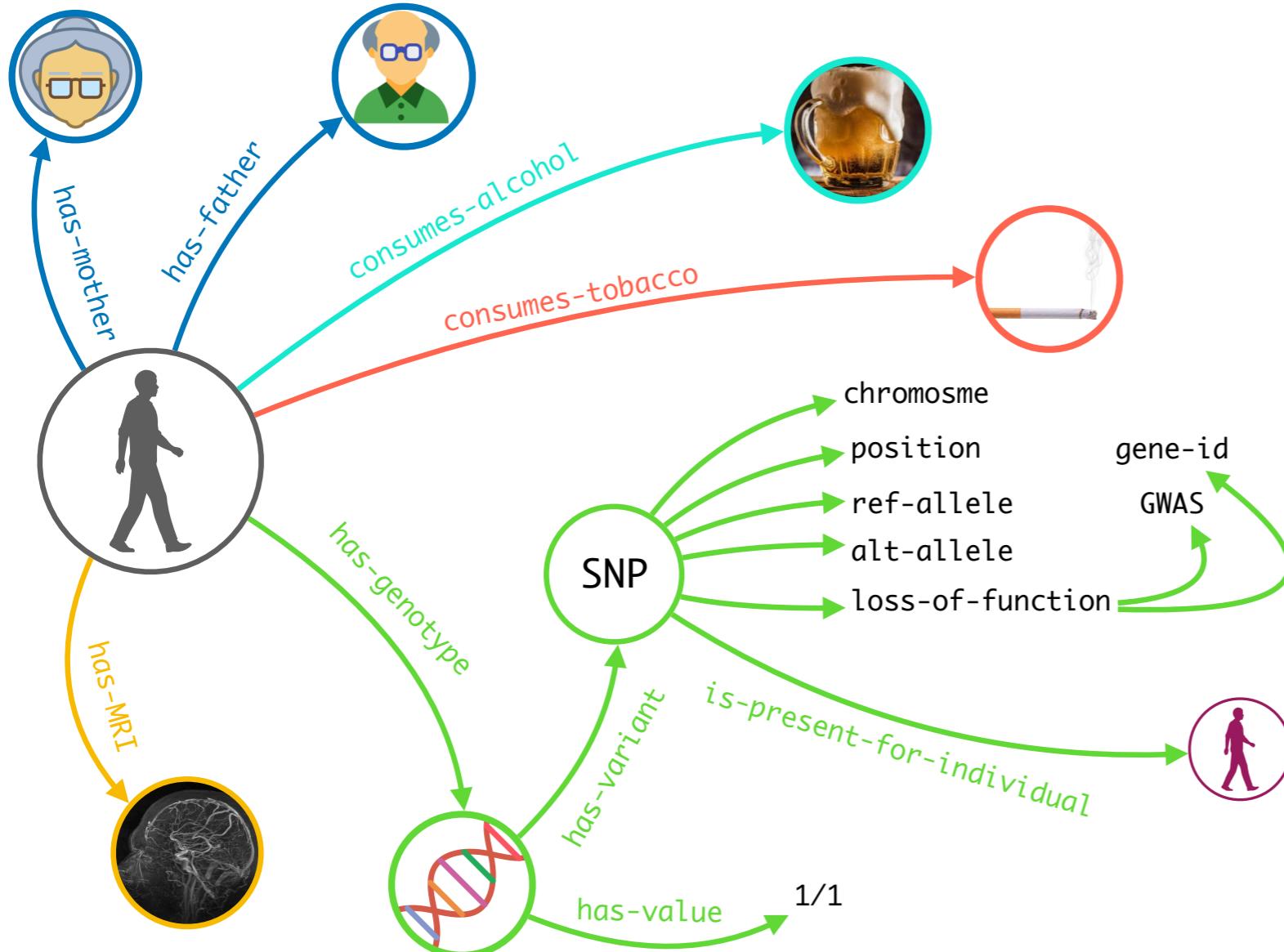
« a **formal specification** of a **shared conceptualization** » (Borst, 1997)
→ 1,049 life science ontologies registered in BioPortal (2023)

Human Phenotype Ontology



② how to link data from multiple scales ?

Knowledge graphs: « a collection of interlinked descriptions of things (real-word objects, abstract concepts, events, etc.) »



= **database:** information storage / extraction

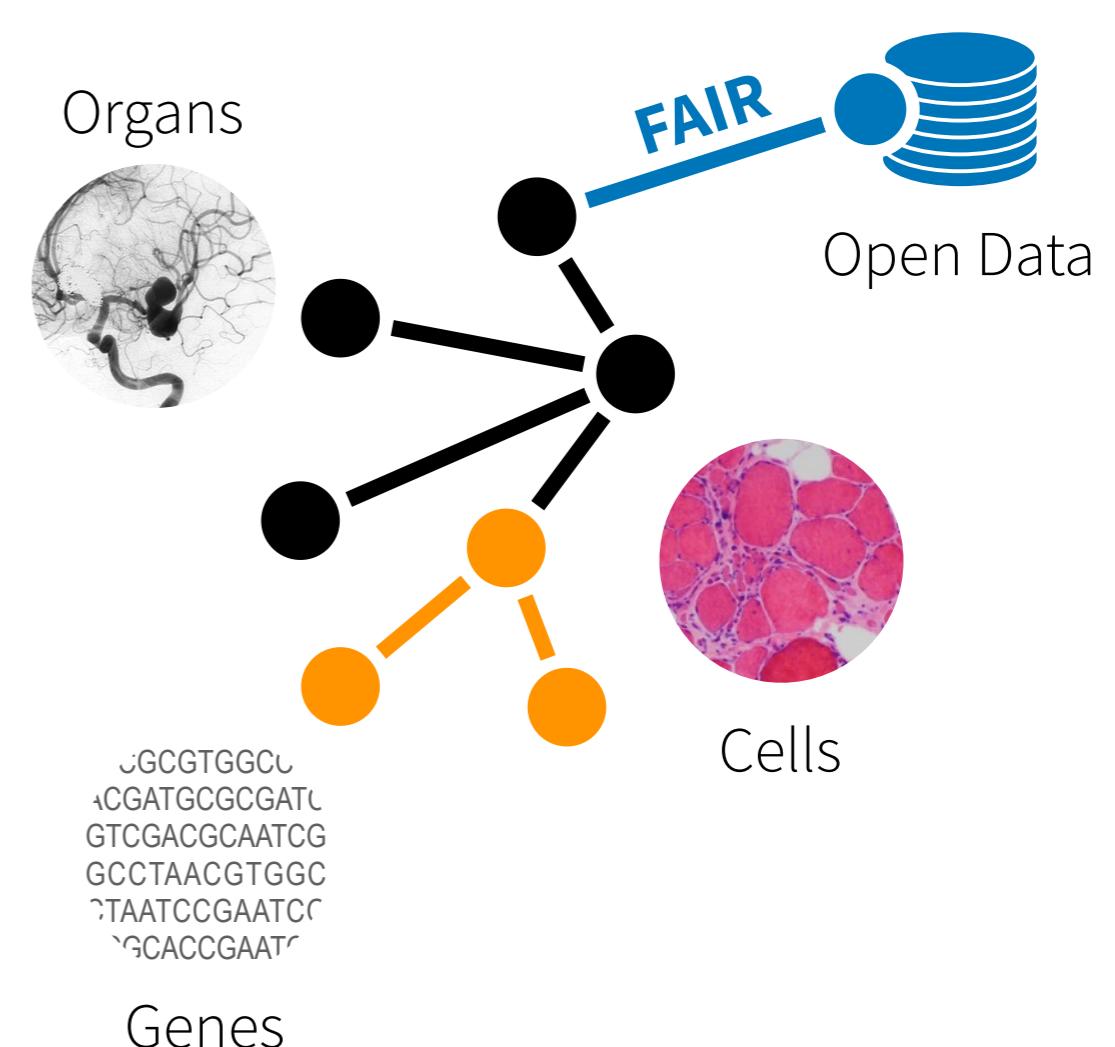
= **graph:** network analysis

= **knowledge base** with formal semantics :

- + logical facts
- + logical inferences

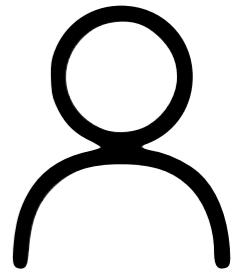
Bridging imaging-omics- clinical data

INEX-MED

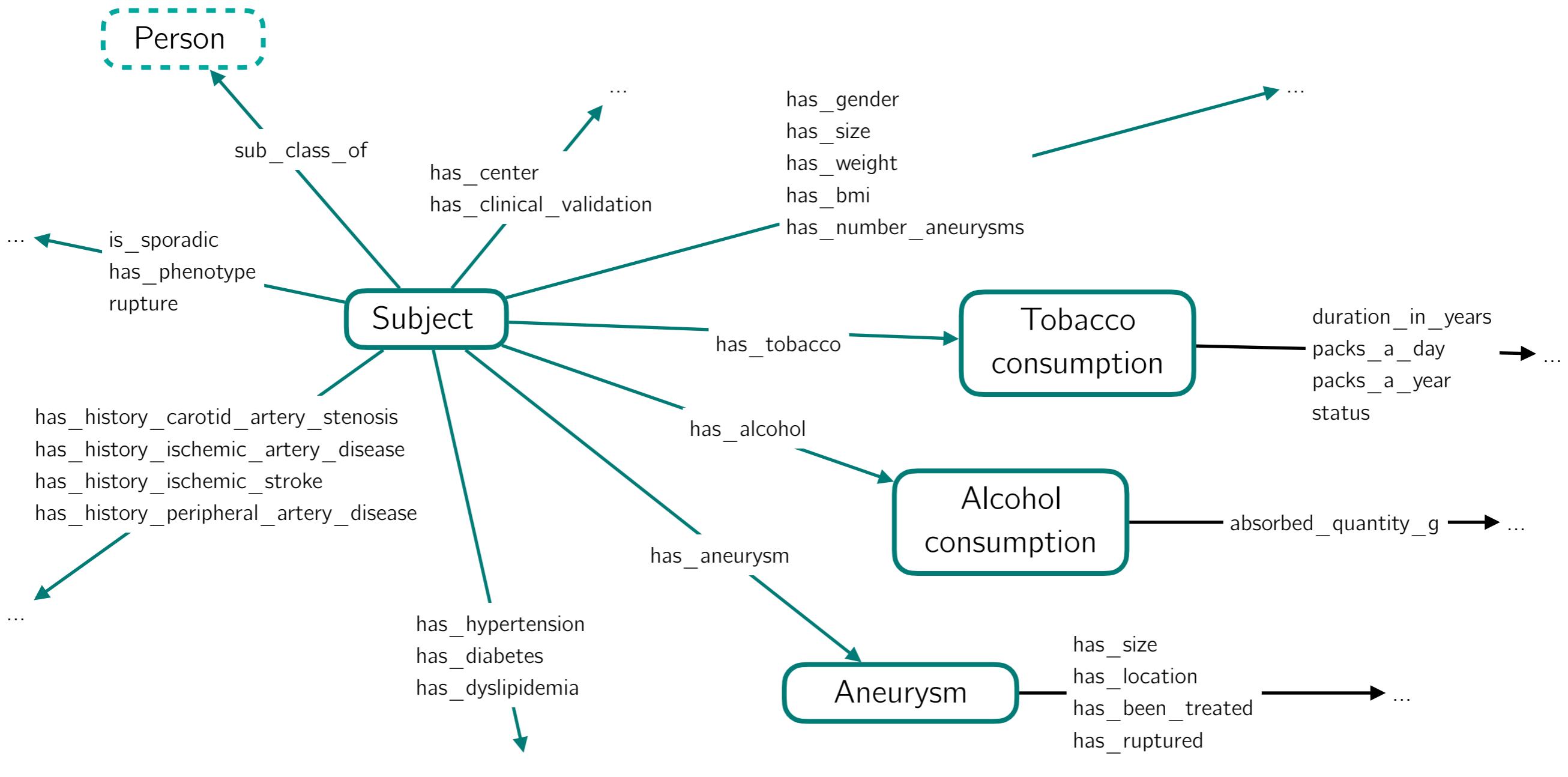


Multi-scale « Knowledge Graph »
145 Million facts

Clinical observations



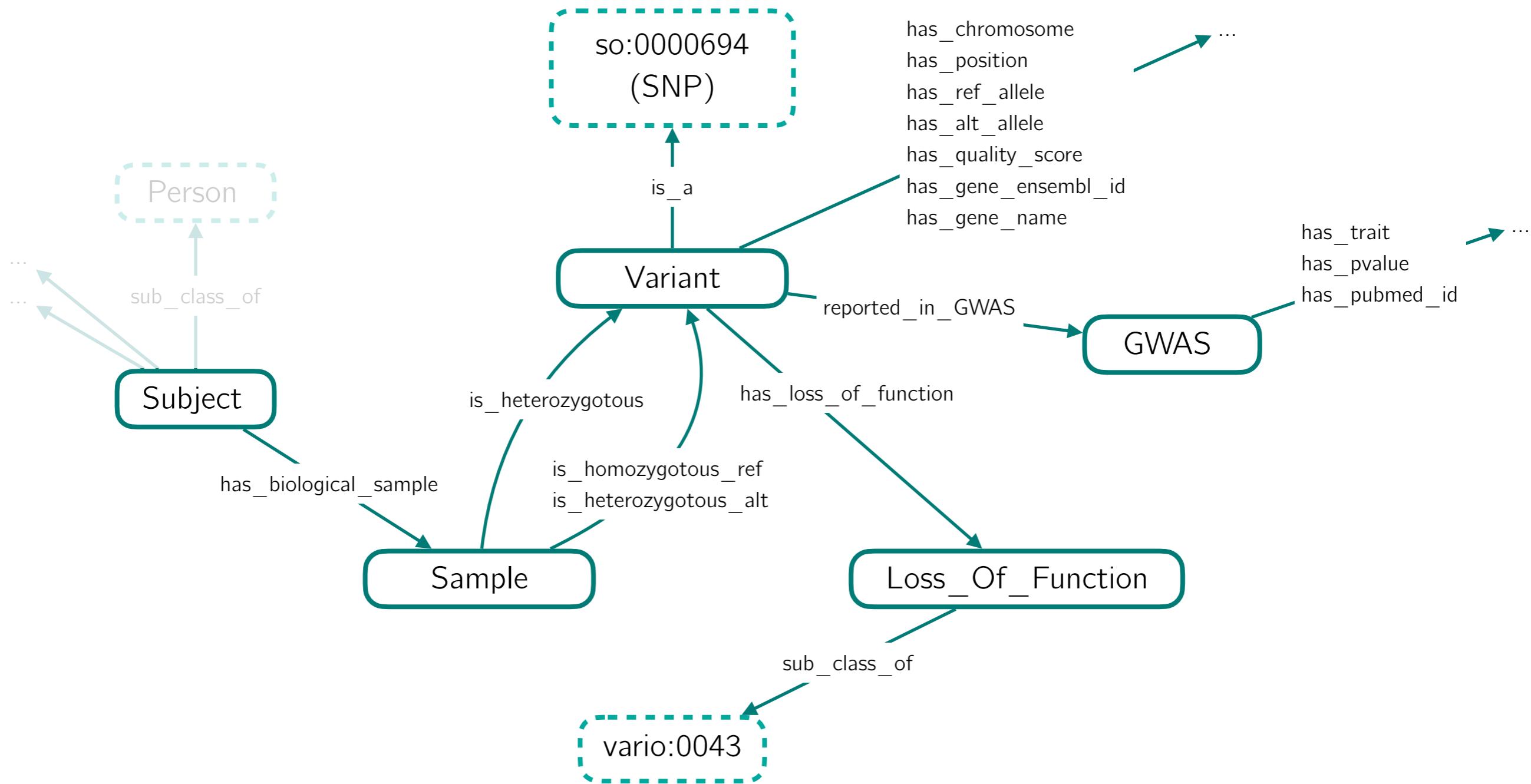
- ▶ Aim : representing key characteristics of ICA patients



Genomic markers



- ▶ Aim : representing associated genetic variations by **extending reference ontologies** (Sequence Ontology, VARIO)

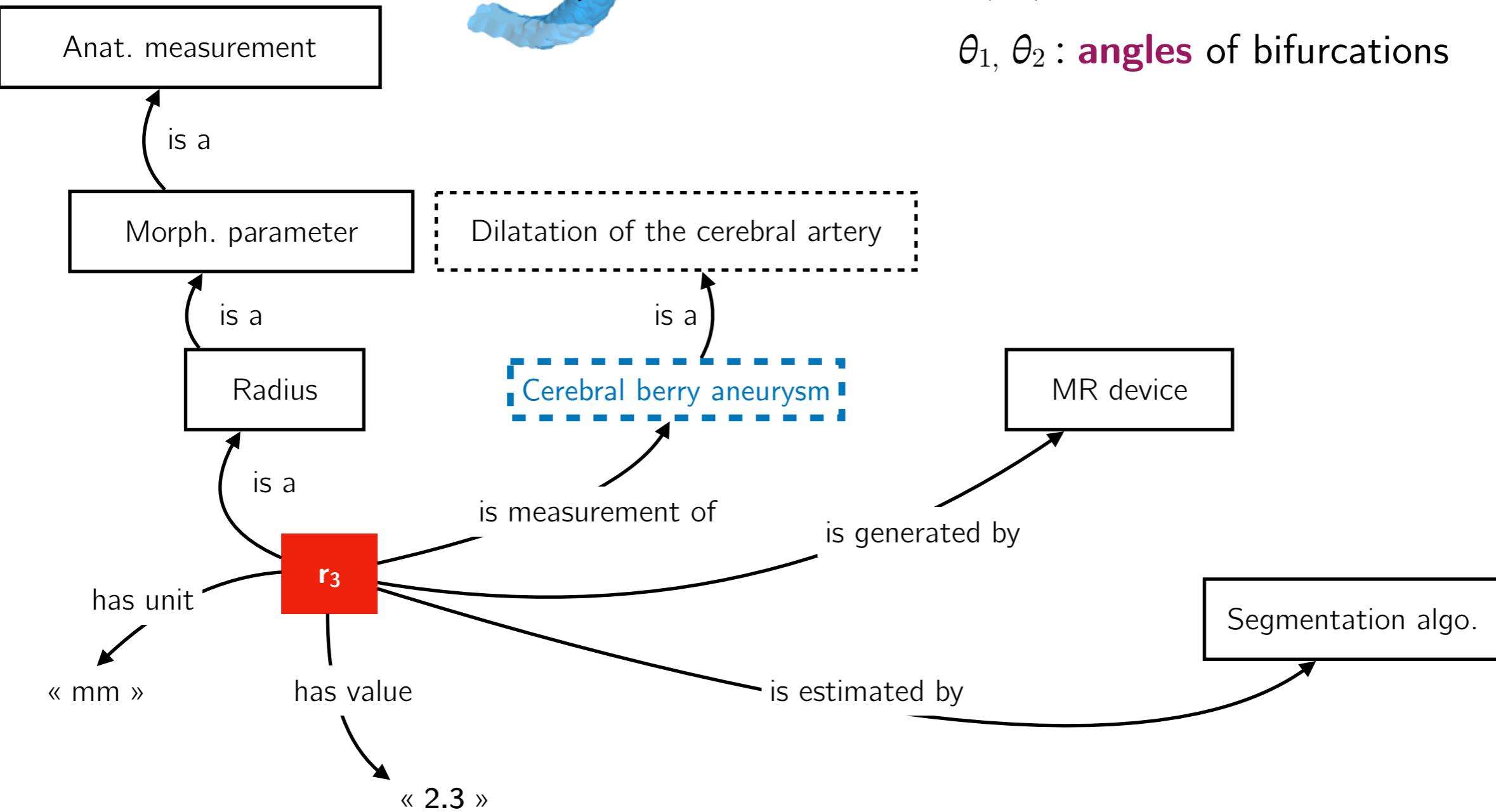
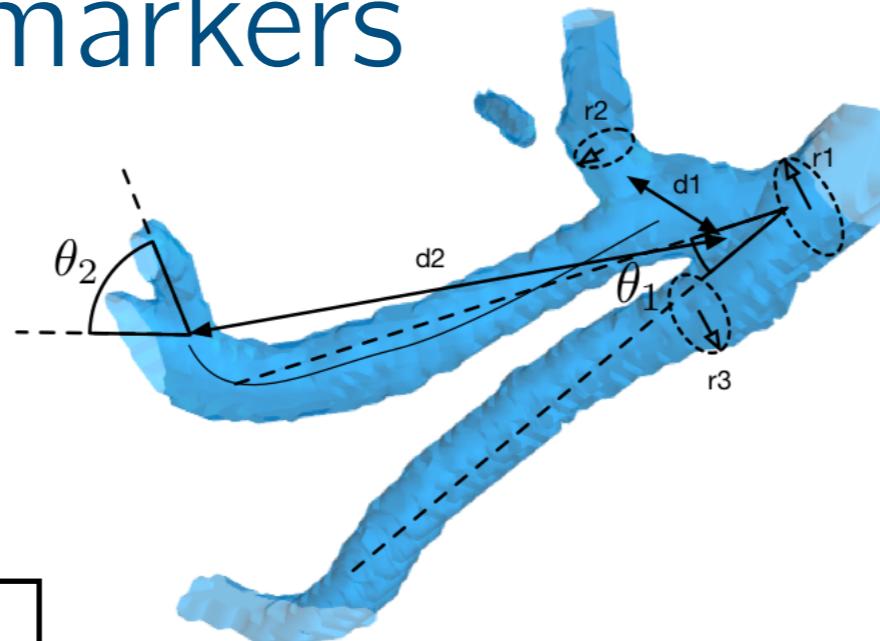


Imaging markers



Anass Nouri

Florent Autrusseau



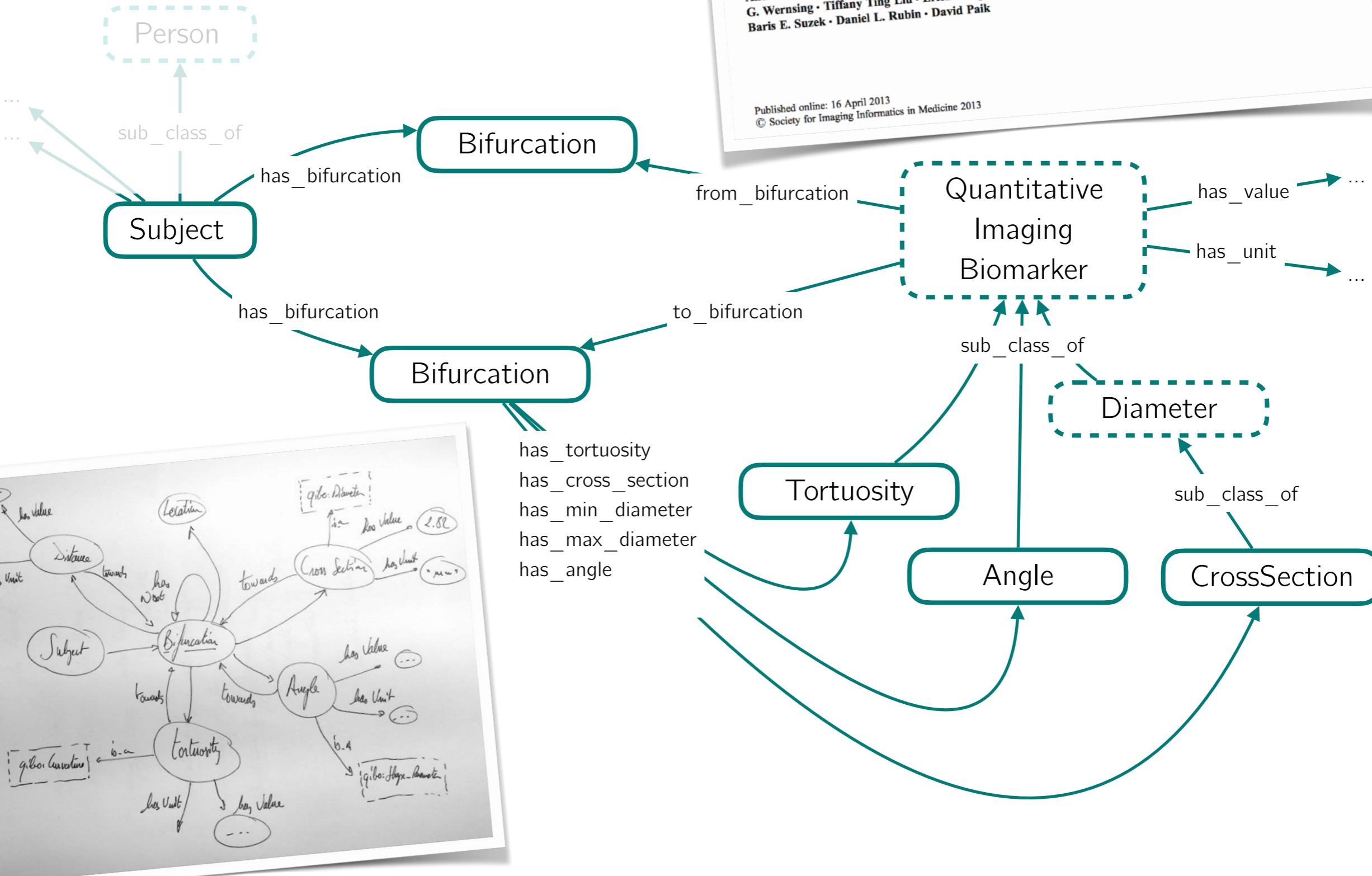
Imaging markers

J Digit Imaging (2013) 26:630–641
DOI 10.1007/s10278-013-9599-2

Quantitative Imaging Biomarker Ontology (QIBO) for Knowledge Representation of Biomedical Imaging Biomarkers

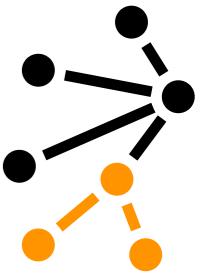
Andrew J. Buckler · M. Ouellette · J. Danagoulian ·
G. Wernsing · Tiffany Ting Liu · Erica Savig ·
Baris E. Suzek · Daniel L. Rubin · David Paik

Published online: 16 April 2013
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Searching
multi-scale data ?

Queries with graph patterns



? "Give all patients with a mutation in ANGPTL6, in the case of a rupture, give me also the age and the location of the ruptured aneurysm."

```
SELECT * WHERE {
  # Search variants
  ?subject ican:has_dna_sample ?sample .
  ?sample ican:is_heterozygous ?variant .
  ?variant ican:has_chromosome "19" ;
            ican:has_position ?position ;
            ican:has_alt_allele ?alt_allele ;
            ican:has_ref_allele ?ref_allele .
  FILTER (?position > 10092337 && ?position < 10106407)
  # dbSNP identifier
  OPTIONAL { ?variant ican:has_dbsnp_id ?dbSNPid . }
  # First aneurysm rupture: age and localisation
  OPTIONAL { ?subject ican:age_first_rupture ?ageRupt . }
  OPTIONAL { ?subject ican:loc_first_rupture ?locRupt . }
}
```

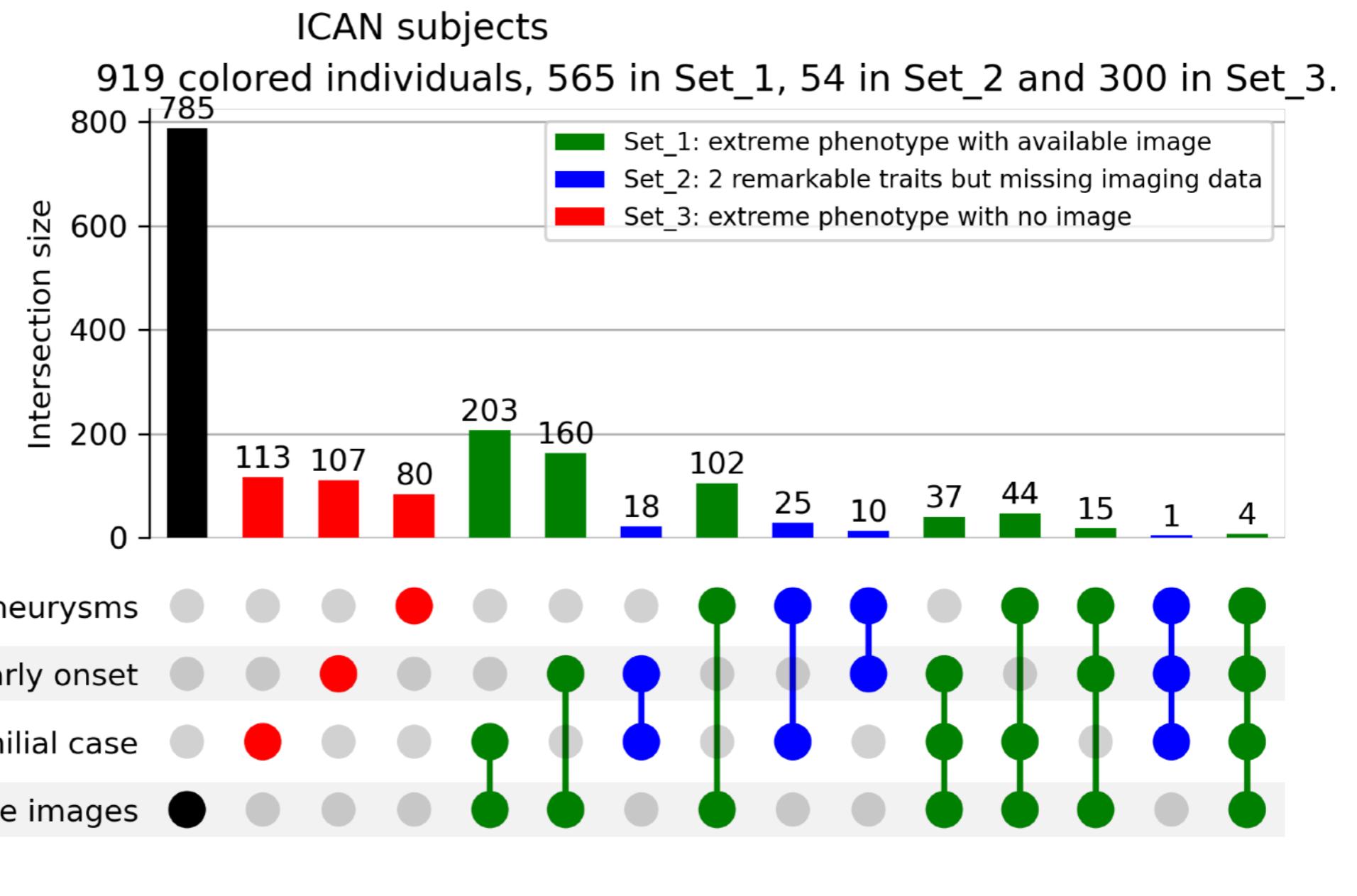
Select "extreme phenotypes" for genomic studies

usable images ?

< 30 years old ?

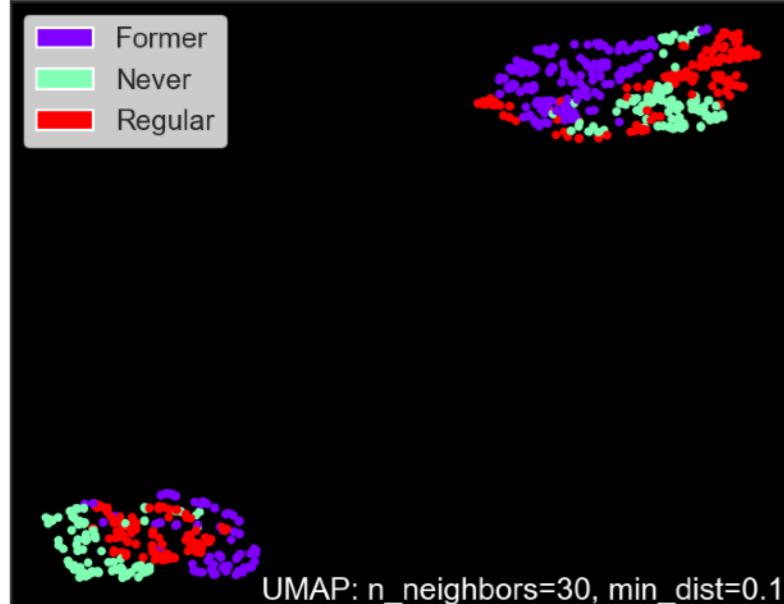
2+ aneurysms ?

familial forms ?

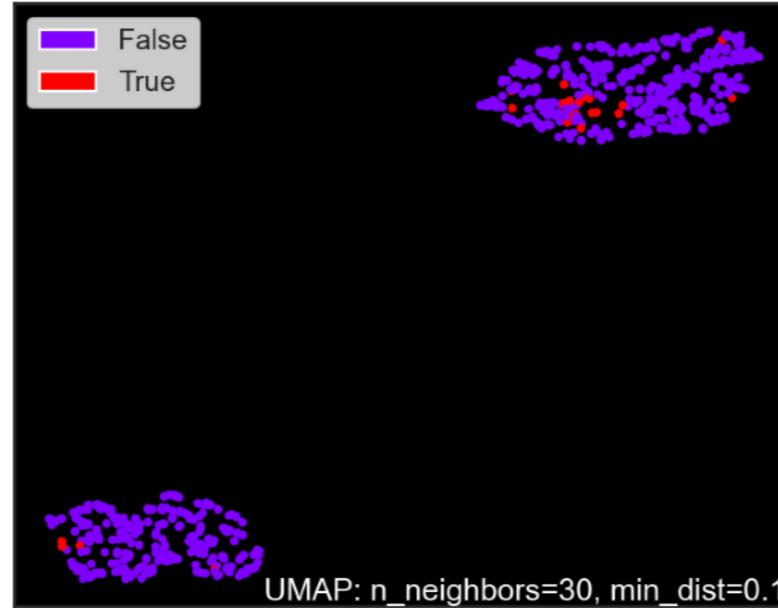


Train machine-learning models ...

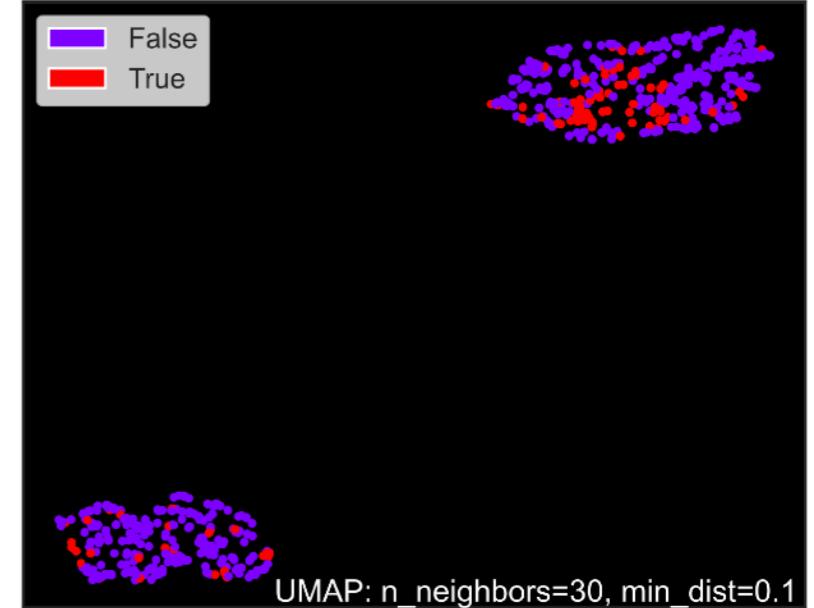
Tobacco consumption



Ischemic heart disease history



Platelet aggregation inhibiting treatment



Observational Study > J Neurol Neurosurg Psychiatry. 2021 Feb;92(2):122-128.

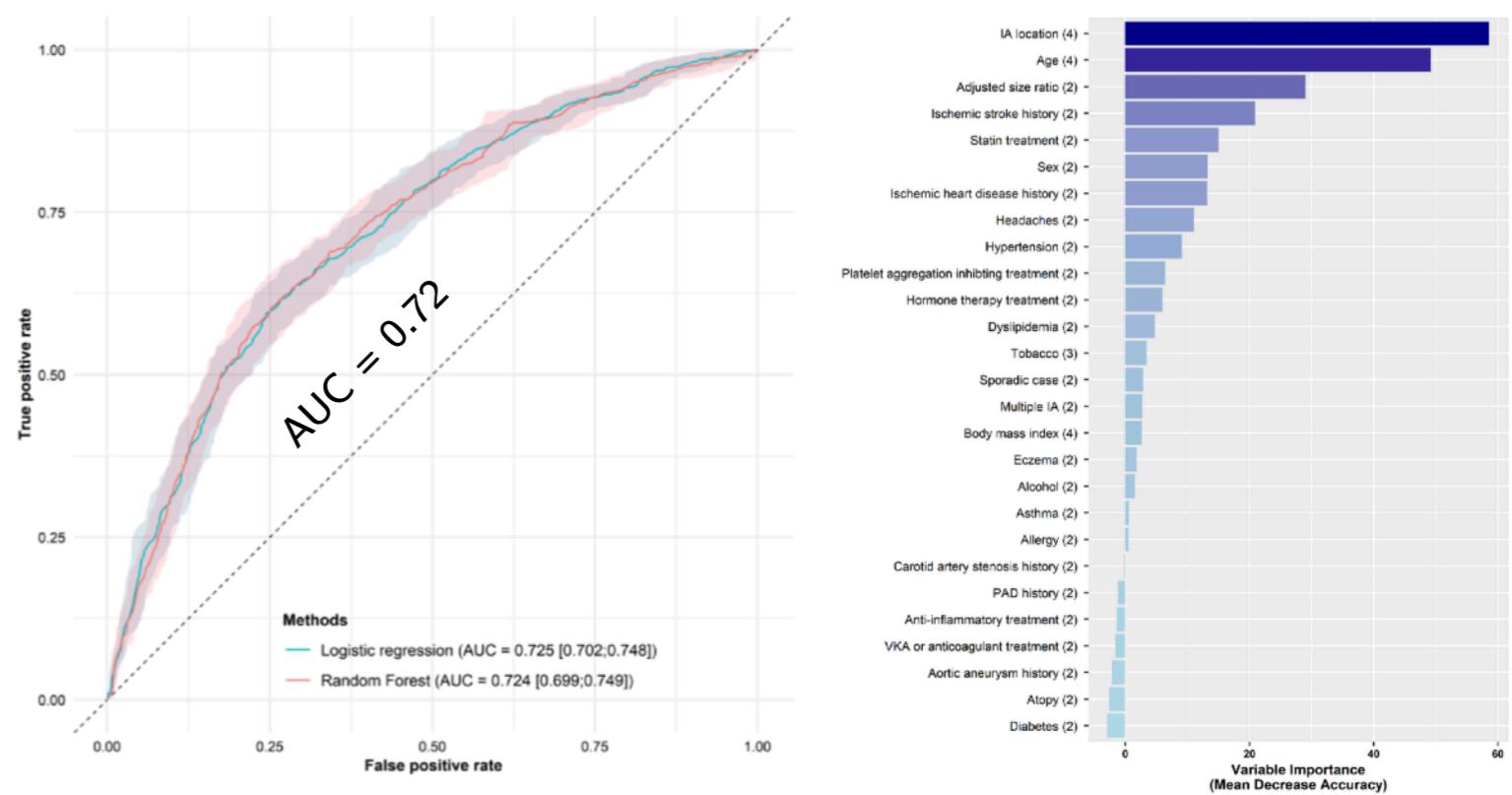
doi: 10.1136/jnnp-2020-324371. Epub 2020 Oct 23.

Location of intracranial aneurysms is the main factor associated with rupture in the ICAN population

Olivia Rousseau ¹, Matilde Karakachoff ¹, Alban Gaignard ², Lise Bellanger ³, Philippe Bijlenga ⁴, Pacôme Constant Dit Beaufils ¹, Vincent L'Allinec ⁵, Olivier Levrier ⁶, Pierre Aguettaz ⁷, Jean-Philippe Desilles ⁸, Caterina Michelozzi ⁹, Gaultier Marnat ¹⁰, Anne-Clémence Vion ², Gervaise Loirand ², Hubert Desal ¹¹, Richard Redon ², Pierre-Antoine Gourraud ¹, Romain Bourcier ¹²; ICAN Investigators

Collaborators, Affiliations + expand

PMID: 33097563 DOI: 10.1136/jnnp-2020-324371



(controlled) access and sharing

INEX-MED data hub ≡

GENERAL

- Dashboard
- Admin

SEARCH

- Centers
- Gender Distribution

ANEURYSMS

- Aneurysms Position
- Genomic Variants

AUTHENTICATION

- Sign-in
- Sign-out

INEX-MED data hub

Exploring clinical, genomics, and imaging data to better understand congenital myopathies and intracranial aneurysms

See the [official INEX-MED wiki page](#) for more information.

SUBJECTS	CLINICAL OBSERVATIONS	BIFURCATIONS	GENETIC VARIANTS
2,974	2,613	9,490	1,549,005

FACTS

143,745,265

Web demonstrator → <https://inexmed-api.univ-nantes.fr>

Related works

Unruptured IA: 51 "core" common data elements / 1254

Age value, Birth date, Birth sex assigned type, Body system category, Race USA category, Sex genotype type, Ethnicity USA category, Gender identity type, Gender type,

Aneurysm diameter maximum measurement, Aneurysm height measurement, Aneurysm morphology type, Aneurysm width measurement,

Hypertension indicator,

Tobacco current use indicator, Tobacco prior use indicator, Tobacco use started age value, Tobacco use stopped age value,

World Federation of Neurological Surgeons (WFNS) - grading system subarachnoid hemorrhage scale

Behavioral history assessment date and time,

Biospecimen collection other text, Biospecimen collection type,

Cerebrospinal fluid collection anatomic site, Cerebrospinal fluid collection anatomic site other text, Cerebrospinal fluid collection method other text, Cerebrospinal fluid collection method type,

Data collected date and time, Data source,

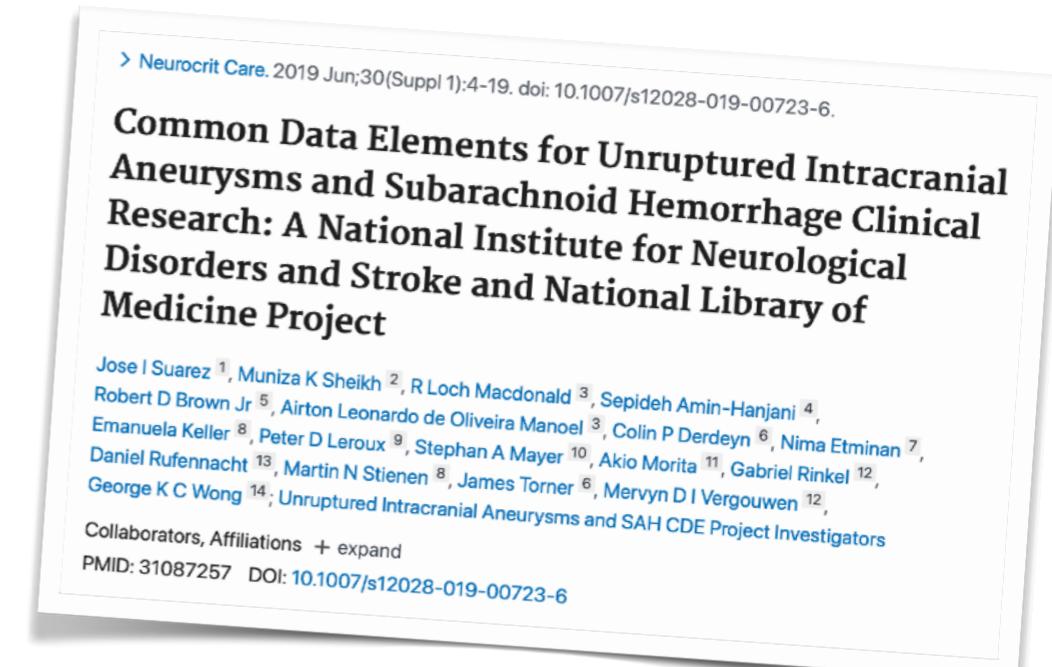
History data not obtained reason,

Imaging modality other text, Imaging modality type, Imaging modality vessel imaging angiography type, Imaging vessel angiography arterial anatomic site, Imaging vessel angiography arterial result,

Intervention endovascular status, Intervention surgical status, Lab specimen collection date and time,

Medical history condition SNOMED CT code, Medical history condition text, Medical history taken date and time,

Specimen acquisition anatomical site, Specimen baseline collection indicator, Specimen biologic source type, Specimen centrifugation duration, Specimen centrifugation revolutions per minute value, Specimen centrifugation temperature value, Specimen collection tube other text, Specimen collection tube type, Specimen not collected reason, Specimen storage temperature value, Specimen time collection process interval,



✓ Large community effort

✓ Consensual definitions

⚠ Flat list of terms

⚠ No recommendation data structures

Sample knowledge graphs : SPHN network

nature > scientific data > articles > article

Article | [Open Access](#) | Published: 10 March 2023

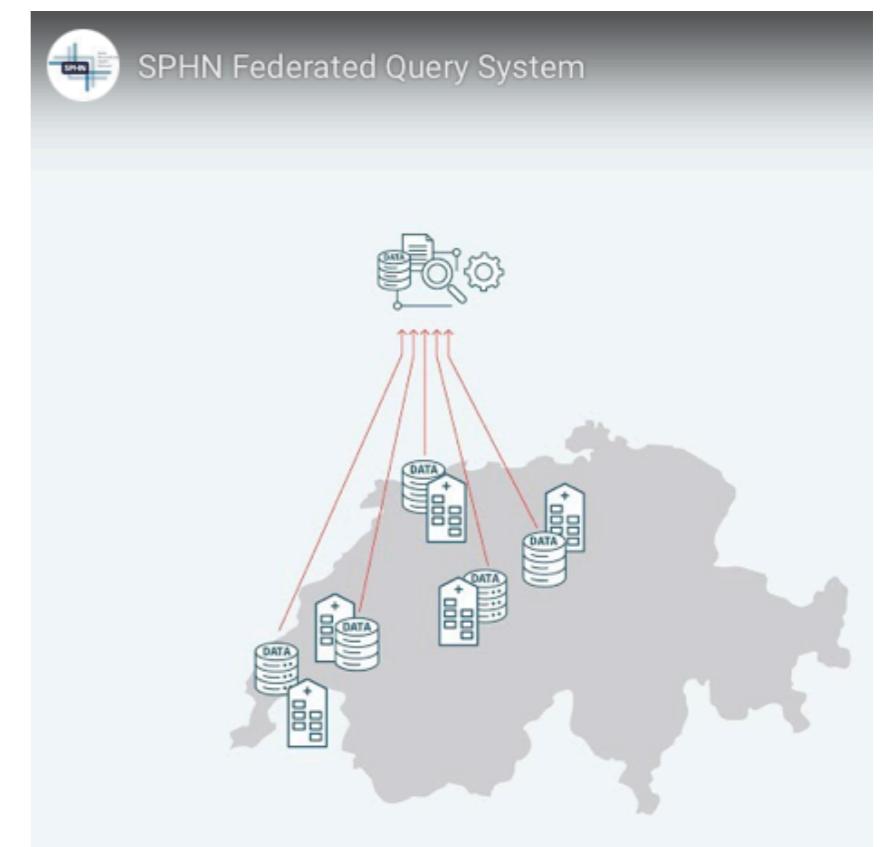
FAIRification of health-related data using semantic web technologies in the Swiss Personalized Health Network

Vasundra Touré, Philip Krauss, Kristin Gnodtke, Jascha Buchhorn, Deepak Unni, Petar Horki, Jean Louis Raisaro, Katie Kalt, Daniel Teixeira, Katrin Cramer & Sabine Österle 

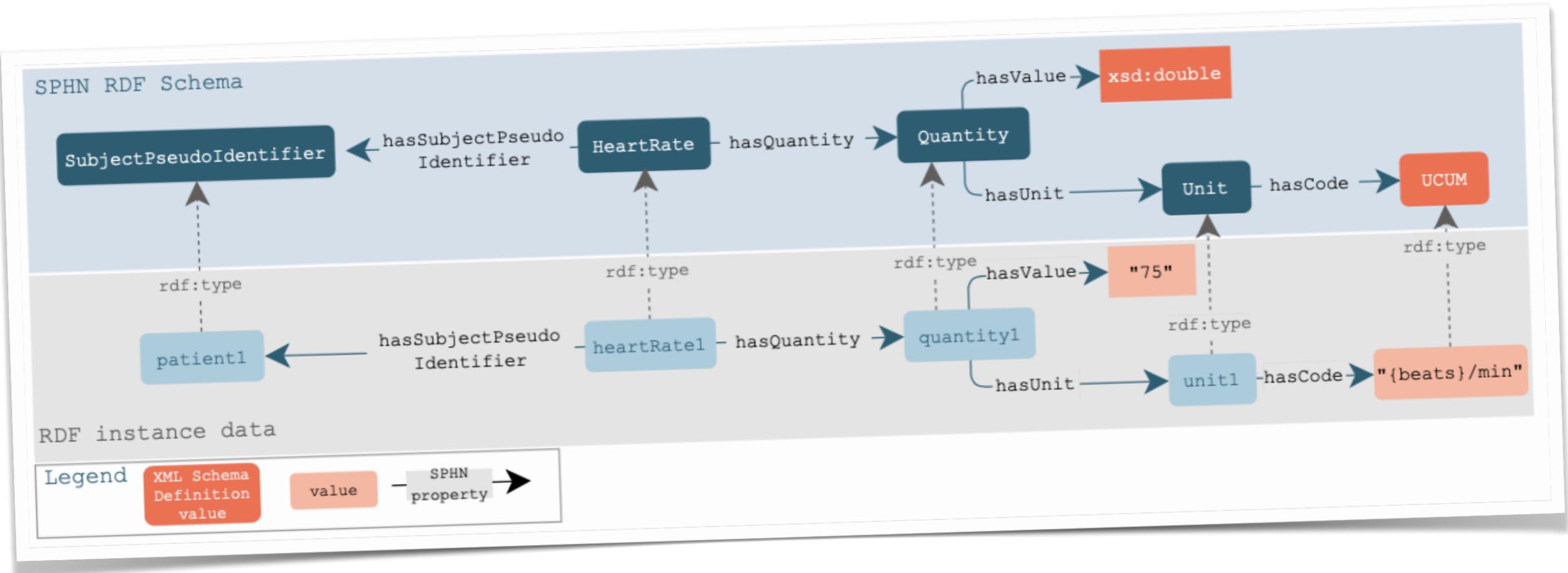
[Scientific Data](#) 10, Article number: 127 (2023) | [Cite this article](#)

Abstract

The Swiss Personalized Health Network (SPHN) is a government-funded initiative developing federated infrastructures for a responsible and efficient secondary use of health data for research purposes in compliance with the FAIR principles (Findable, Accessible, Interoperable and Reusable). We built a common standard infrastructure with

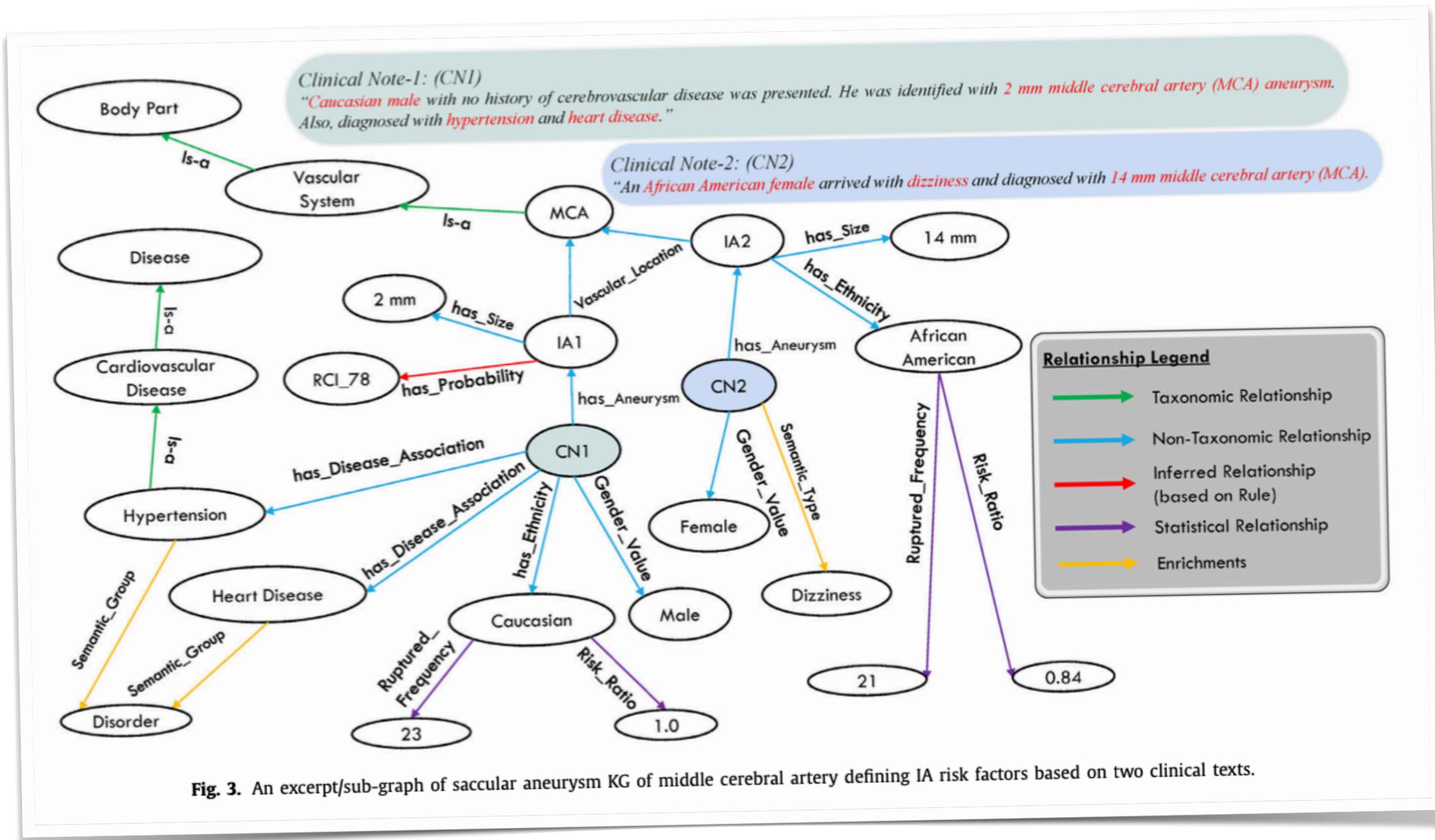


Sample knowledge graphs : SPHN network



- + Allergy
- + Biobank
- + Consent
- + Radiology diagnosis
- + Genomic variants position
- + ECG

Sample knowledge graphs: Malik et. al. 2020

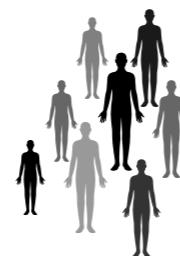
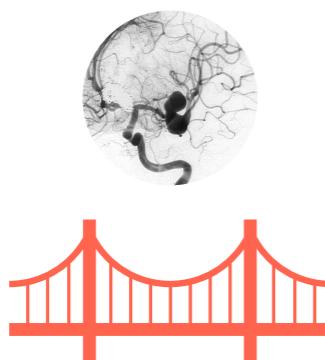


Future works

Wrap up & next steps

- ▶ Gather **consensual expert knowledge** with computational **Ontologies**
- ▶ Knowledge Graphs are "**machine-processable**"
- ▶ Link **multi-scale data** with Knowledge Graphs
- ▶ Query **multi-source** data

CGCGTC
CCTAACGAT
GCGCGATCC
TGACGCA



PEPR Santé numérique

- ▶ Include familial relations to query IA **familial forms**



- ▶ Represent **care pathways and guidelines**
- ▶ Develop a national **data hub** with FAIR intracranial aneurysms data
- ▶ **Share** and enrich data models with **international** initiatives, *but which ones ?*



- ▶ Further develop **predictive approaches** (rupture risk prediction)

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BiRD facility,
NeurlInfo facility

