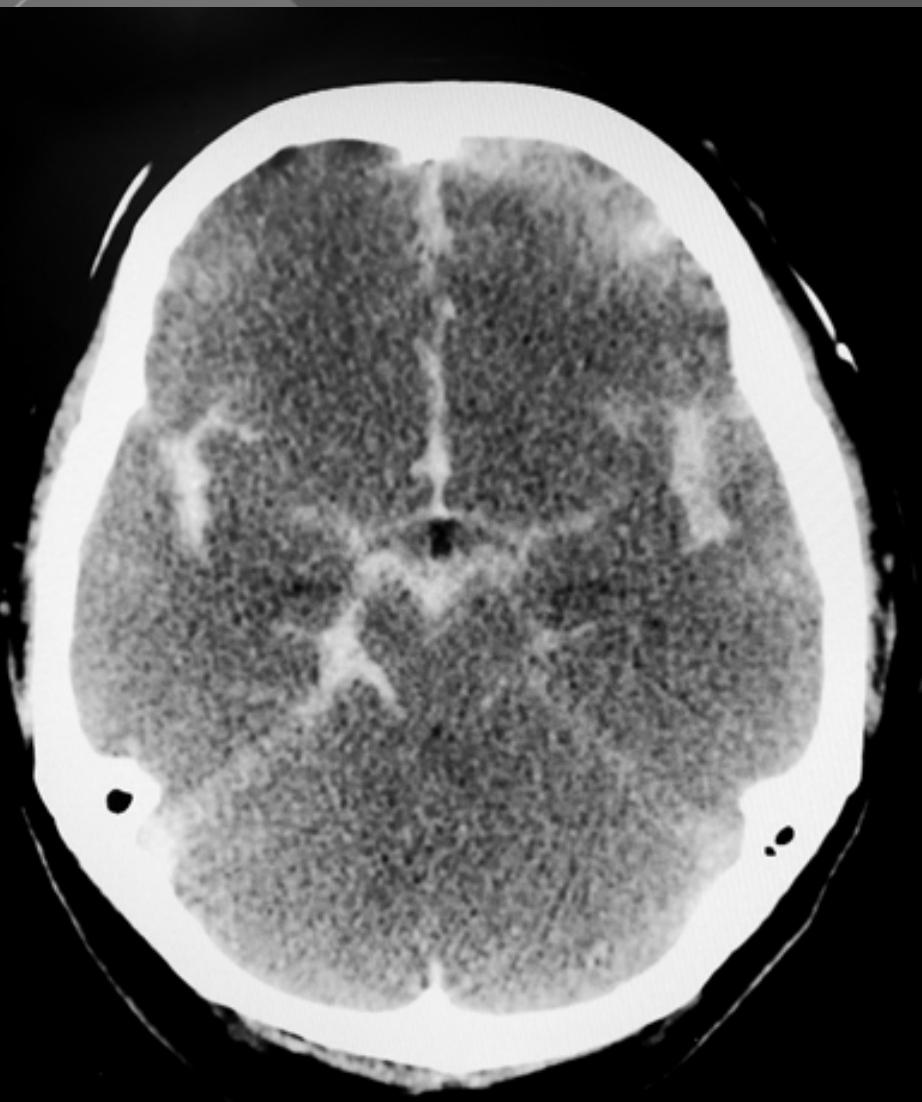


Genetica y epigenética del vasoespasmo y la ict en la hemorragia subaracnoidea aneurismática

II congreso anual ricor-ictus



Elisa Cuadrado-Godia, Isabel Fernández Pérez , Joan Jiménez Balado
Neurology Department &Neurovascular Research Group
Hospital del Mar Medical Research Institute (IMIM)
Barcelona(SPAIN)



ANEURISMAL SUBARACHNOID HEMORRHAGE

1

Epidemiology

5% all stroke

mean age 50-60y

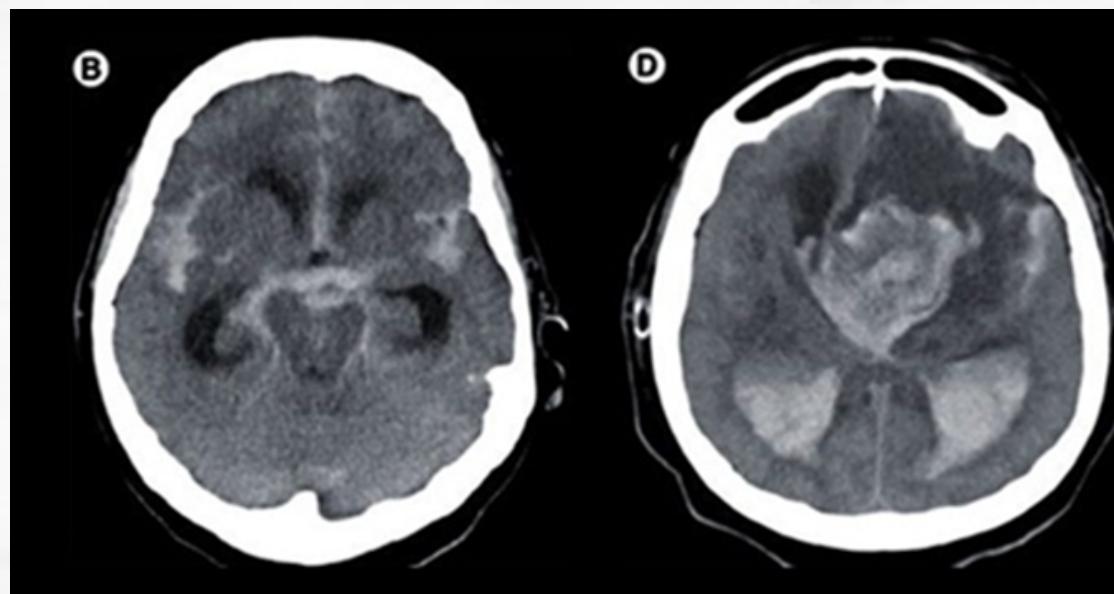
2

Prognosis

Mortality 30%

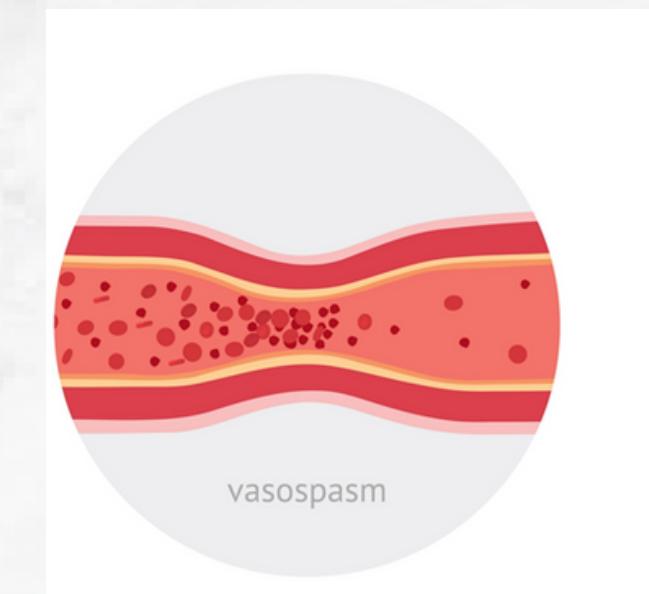
Improved in last decades

PROGNOSIS



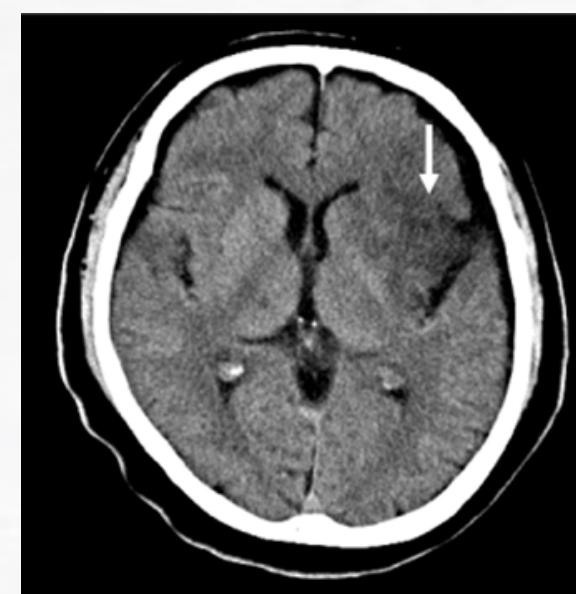
INITIAL BLEEDING

WFNS scale
Fisher scale



COMPLICATIONS

Rebleeding
Vasospasm
DCI

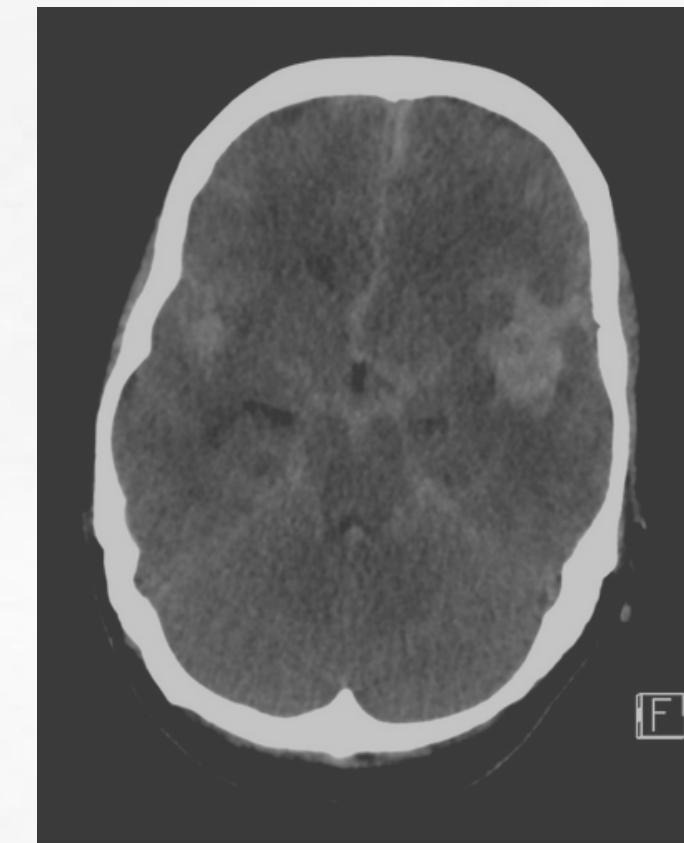


27 y female, right MCA ane, clipped



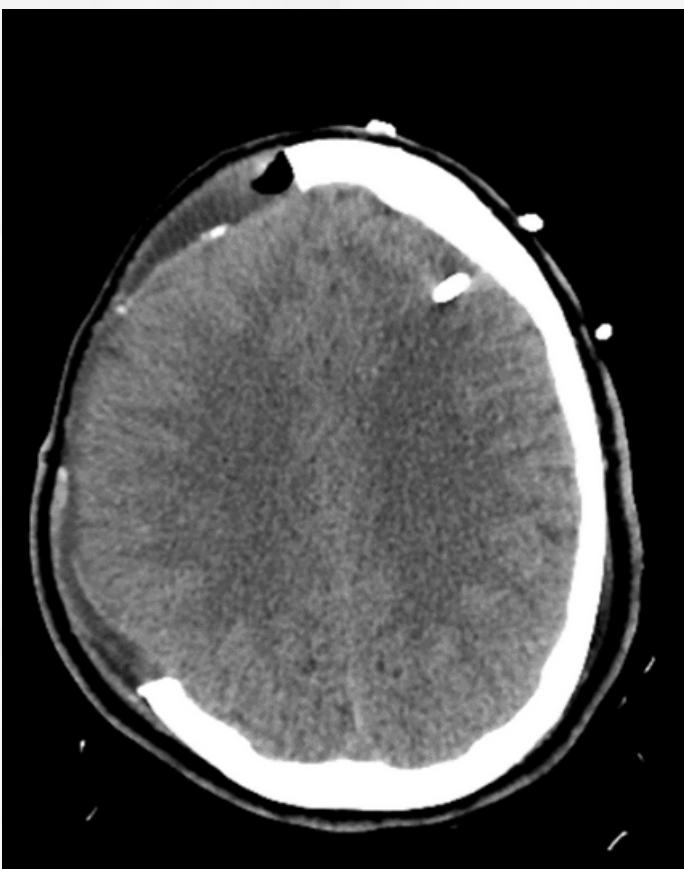
Severe Vasospasm

44y female, left MCA ane, coiled



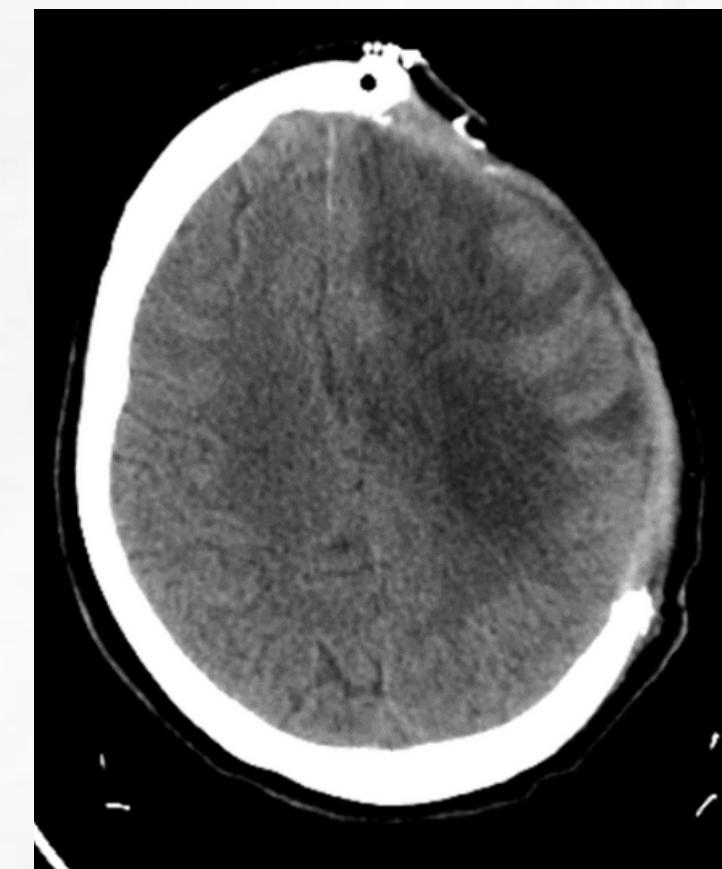
NO VASOSPASM

No DCI



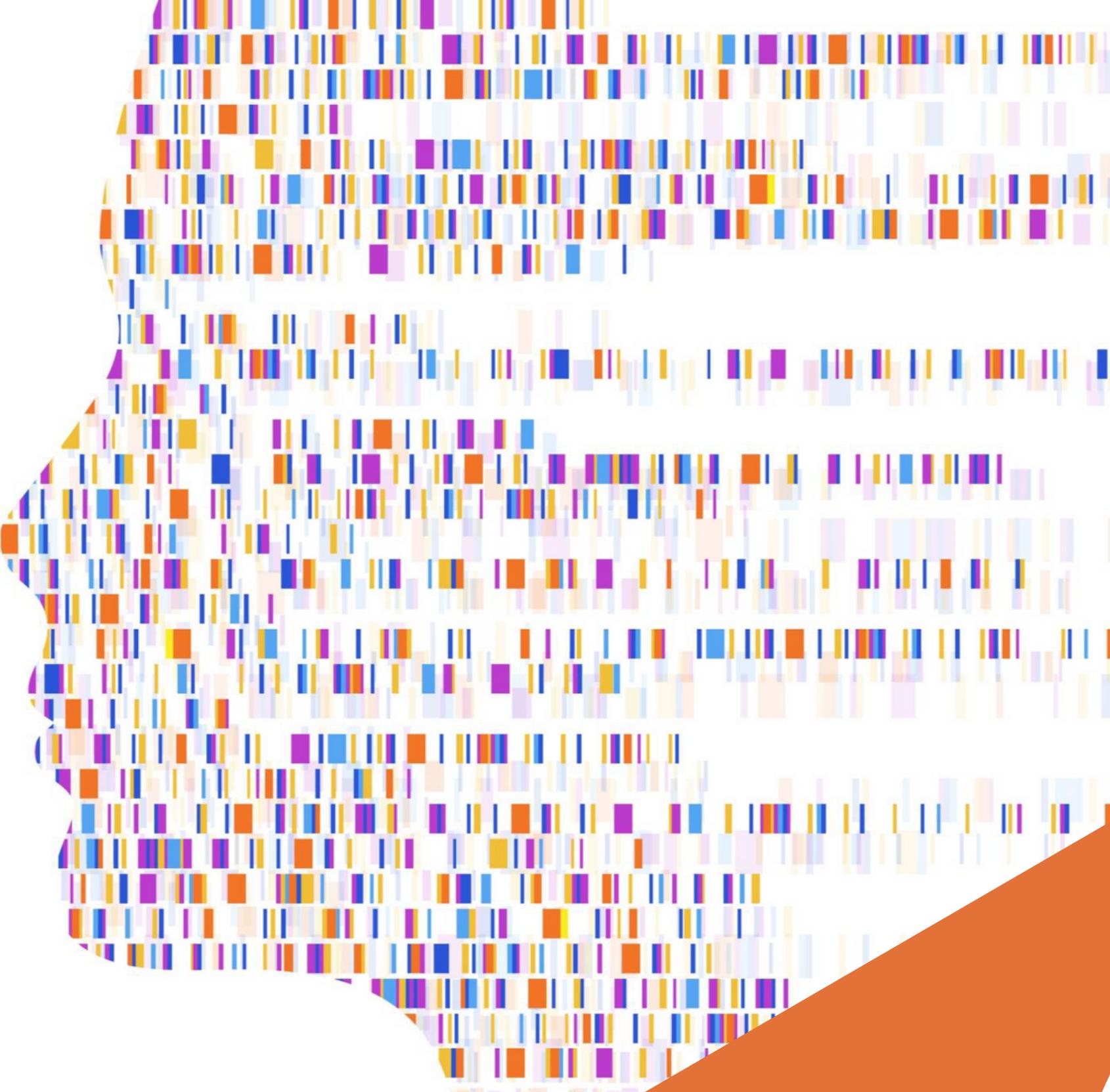
Severe Vasospasm 4rd day

DCI



NO VASOSPASM

*is there a genetic
or epigenetic
predisposition to
vasospasm or
DCI?*



GENETIC STUDIES

Candidate Gene

Inflammation
Endothelial dysfunction
Fibrinolytic system
Brain metabolism

GWAS

N= 245 aSAH, Rs999662 (SLC12A3) ---->TCD vasospasm in

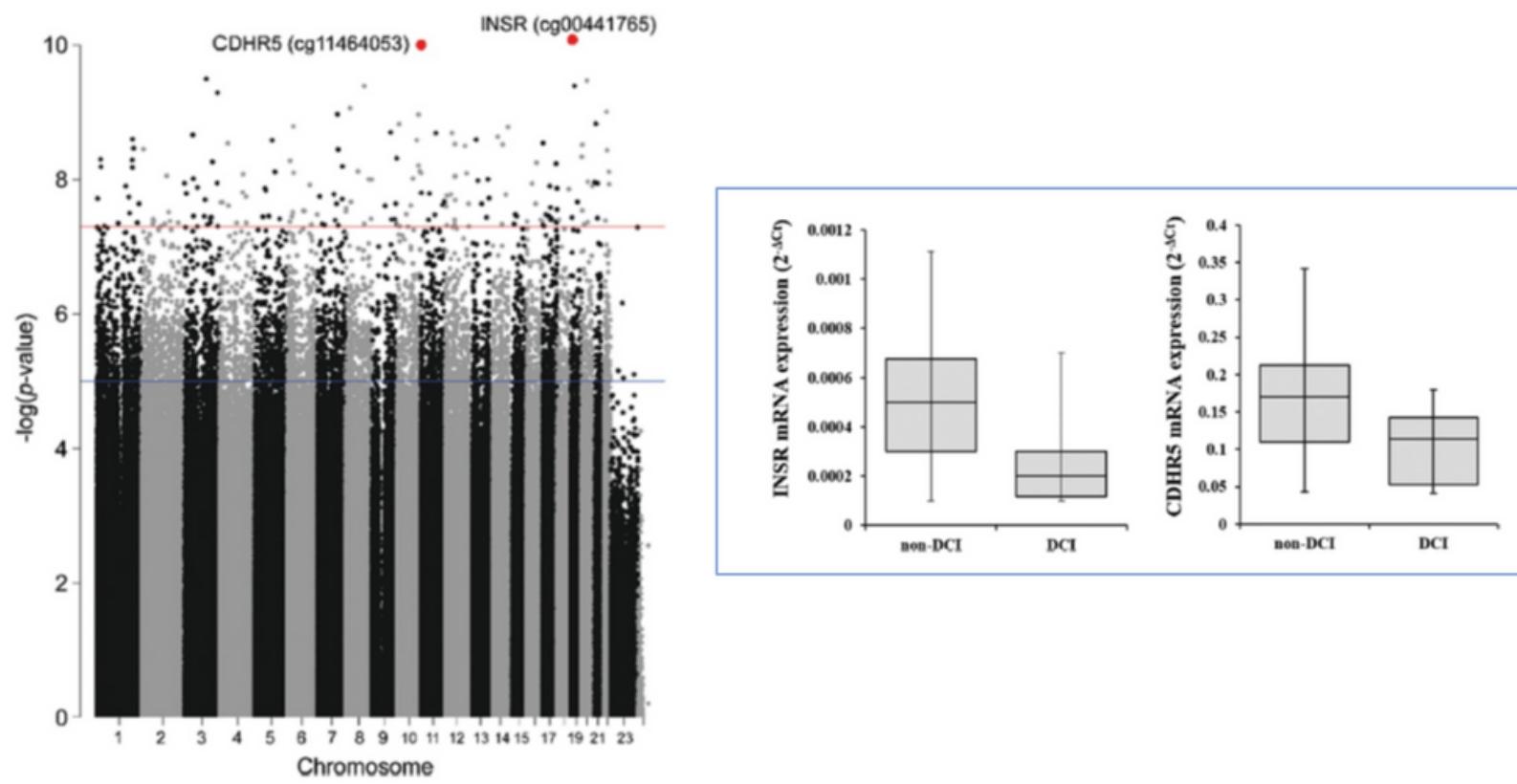
P = 3·39 10-8

N=190 aSAH, no association in GWAS

Ducruet AF. J Cer blood Flow & Metab2010,
Ladner T. Neurol Res Int 2013, Rosalind Lai. World Neurosurgery (2015),Jabbarly R. Neurosug rew 2018

EPIGENETIC STUDIES DCI

EWAS



EWAS

Kim BJ et al. *Sci. Reports* 2020

AIM OF OUR LINE OF RESEARCH

Identify **genetic variants** of vasospasm and/or DCI, through Genome-wide association (**GWAS**)

Identify **methylation alterations** in loci associated with vasospasm and/or DCI through Epigenome-Wide Association (**EWAS**)

Analyze the influence of the genetic sequence on DNA methylation at specific loci (**meQTLs**),

Create a **risk scale** for vasospasm/ DCI

GENETICS OF VASOSPASM AND DELAYED CEREBRAL ISCHEMIA IN ANEURYSMAL SUBARACHNOID HAEMORRHAGE

GENVADER



multicentric

inclusion

gwas

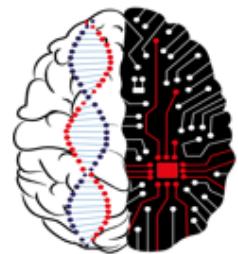
computing

1. Hospital del Mar Barcelona, Spain.
2. UCL Queen Square Institute of Neurology, University College London, UK.
3. University Medical Center Utrecht, The Nederlands.
4. University of Helsinki, Finland.
5. University of Geneva, Switzerland.
6. CHU Nantes, France.
7. CAN project investigators, France.

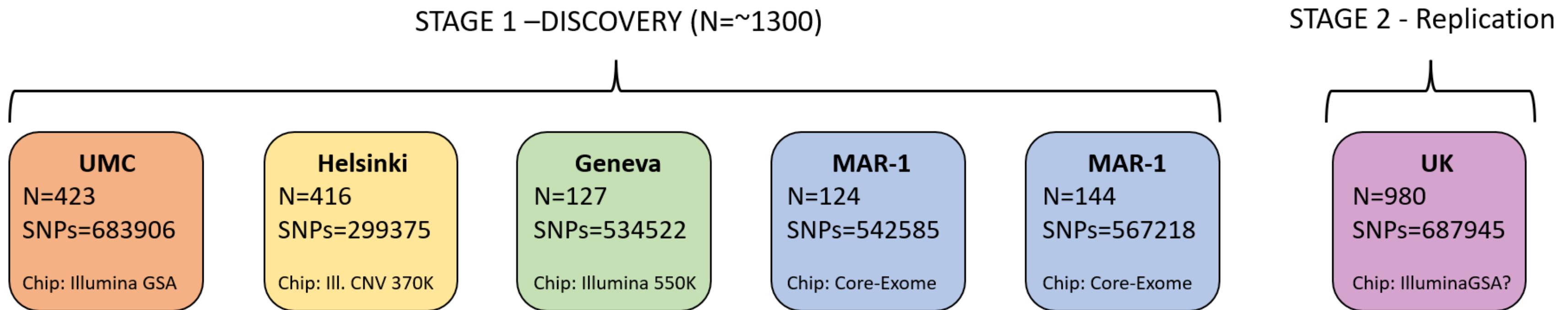
1. aSAH with data on VS and DCI
2. individual level whole-genome genotype data

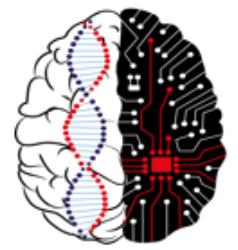
QC analyses
Controlling for population stratifications
Mixed Model GLM
VS-DCI separately

Poligenic risk score
Mediation analysis VS-DCI
Characterization of GWAS loci
Drug target enrichment



Cohorts description: Patients already included





Discovery QC's Diagram

- Relaxed cutoff and projection to ref pop.
- Filter SNPs CR < 98%
- Filter Individuals CR <98%
- Keep only biallelic SNPs
- Remove C/G A/T SNPs
- Keep CHR 1 to 22 and X
- Remove monomorphic SNPs
- Check Sex Discrepancy
- Delete Monomorphic SNPs
- Check Relatedness
- PCA → Exclude sample outliers
- HWE

QC's-Cohort

- Filter SNPs CR < 98%
- Filter Individuals CR <98%
- Delete Monomorphic SNPs
- Check Relatedness
- Filter MAC < 10
- Heterozygosity
- HWE
- PCA –

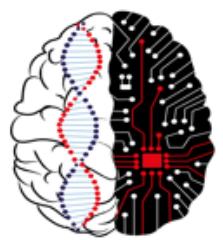
QC's-STRATA

- Filter SNPs R > 0.8
- Filter SNPs CR < 98%
- Filter Individuals CR <98%
- Check Relatedness
- Filter MAC < 10
- Filter MAF < 0,005
- Heterozygosity
- HWE

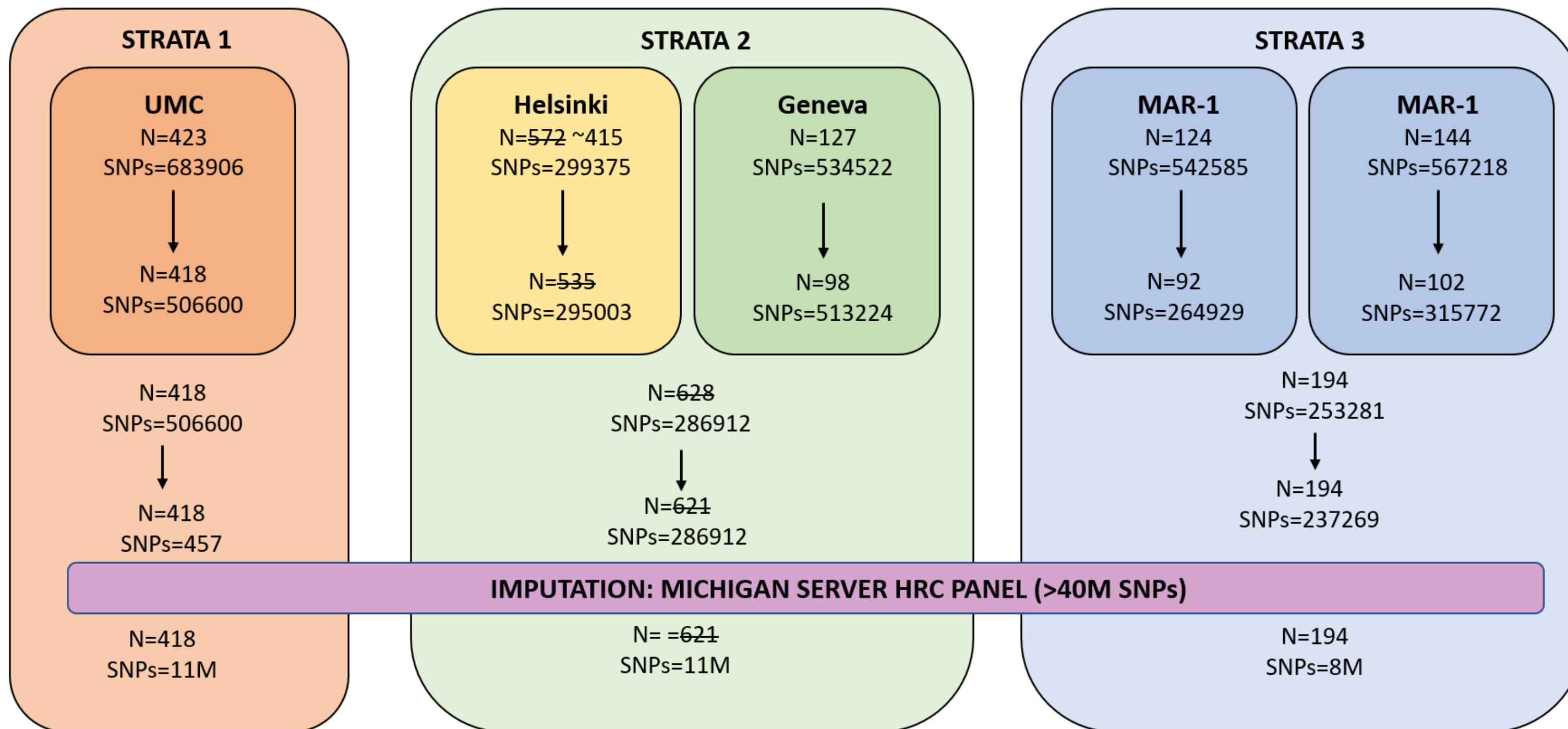
QC's-
IMPUTATION

Merge cohorts
with similar chip

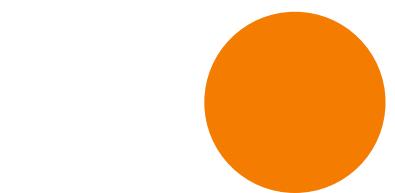
IMPUTATION
Michigan HRC



Cohorts after Merging

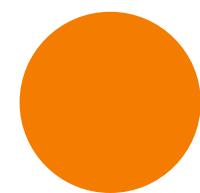


EPIGENETIC CHANGES IN ANEURISMAL SUBARACHNOID HEMORRAGE



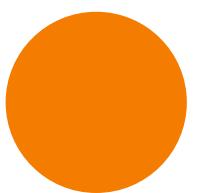
mocentric

1. Hospital del Mar
Barcelona, Spain



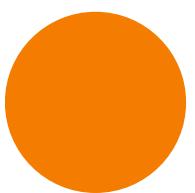
inclusion

aSAH with data on VS
and DCI



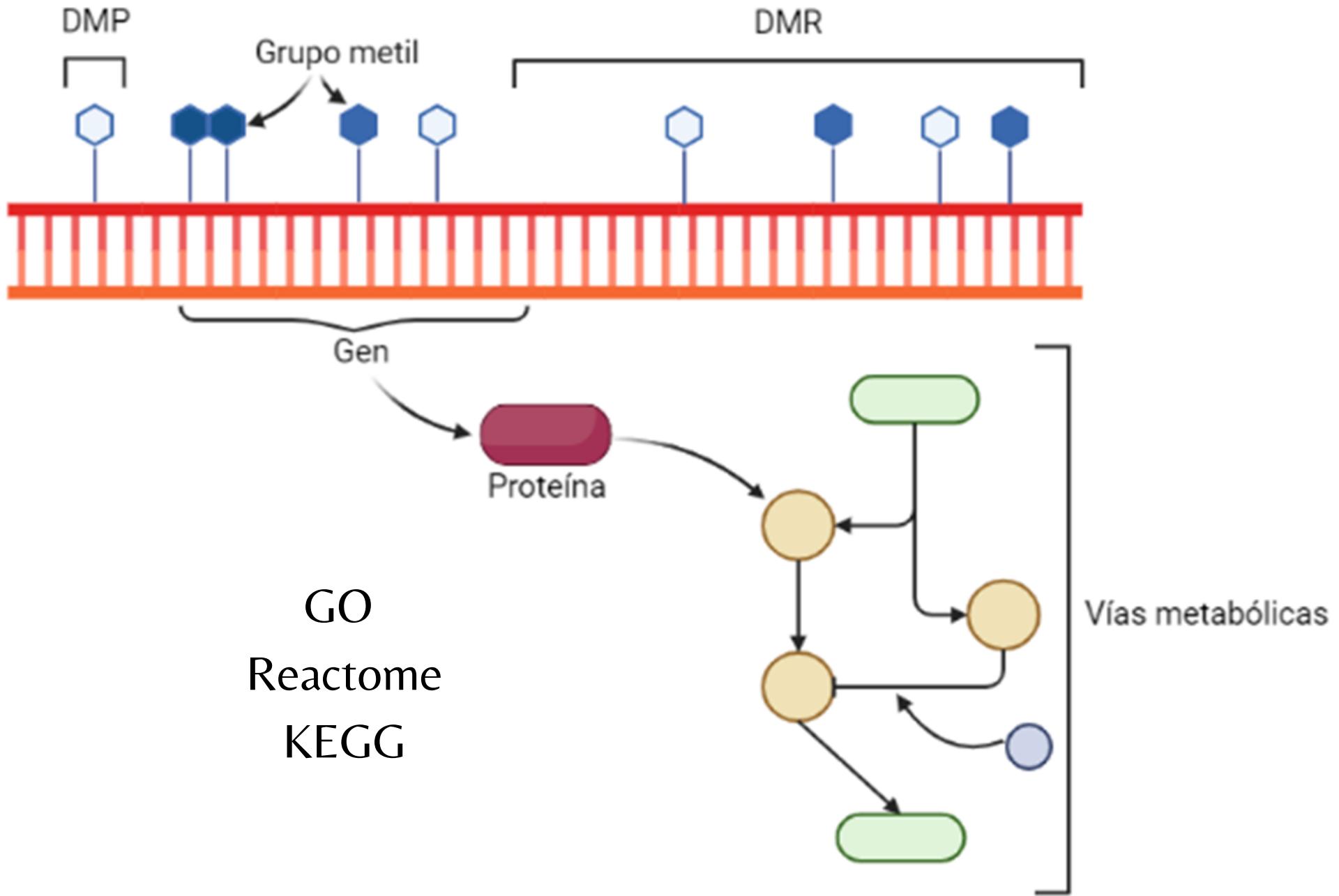
ewas

EPIC Illumina chip (>850K CpG s)
Differential methylation (DMP)
VS-DCI separately
Replication



computing

EPIGENETIC CHANGES IN ANEURISMAL SUBARACHNOID HEMORRAGE



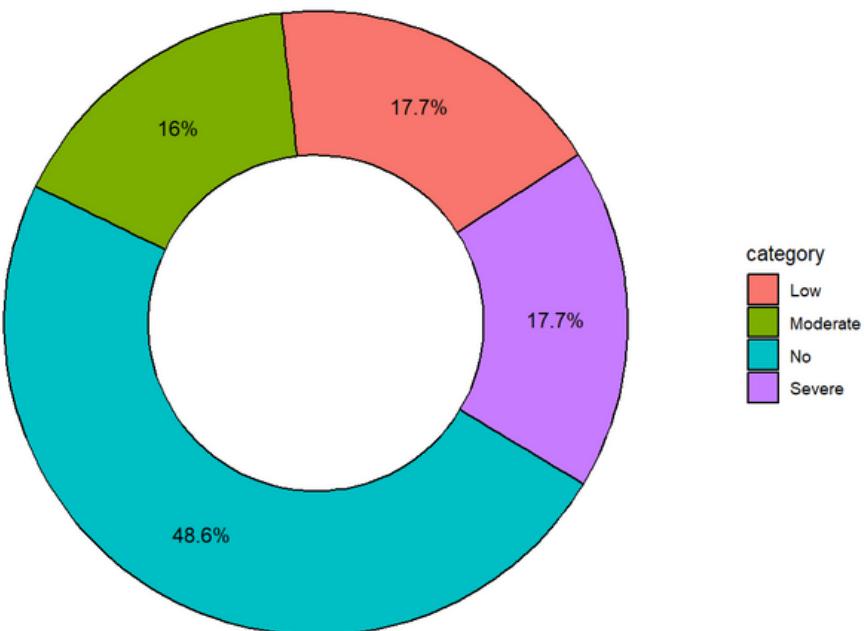
computing

RESULTS

	N=282
Edad	55.0 [17.0;88.0]
Sexo:	
Hombre	93 (33.0%)
Mujer	189 (67.0%)
Etnia:	
Caucásica	241 (85.5%)
Otra	41 (14.5%)
HTA	133 (47.2%)
Fumador	106 (37.6%)
Fisher:	
1	9 (3.19%)
2	23 (8.16%)
3	66 (23.4%)
4	184 (65.2%)
Hunt Hess:	
1	6 (2.13%)
2	152 (53.9%)
3	59 (20.9%)
4	32 (11.3%)
5	33 (11.7%)
Días hasta llegada	0.00 [0.00;32.0]

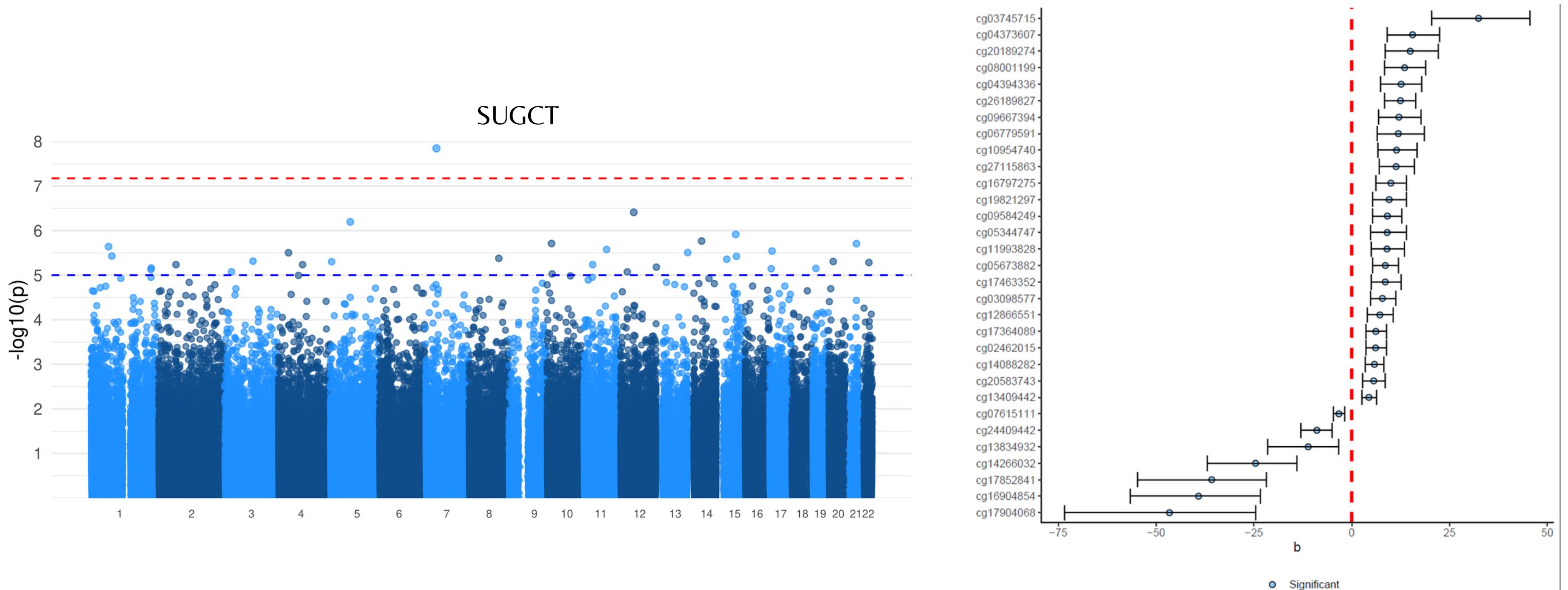
VS 145 (51.4%)

Proportion of Vasospasm Categories



DCI 71 (25.2%)

RESULTS VASOSPASM

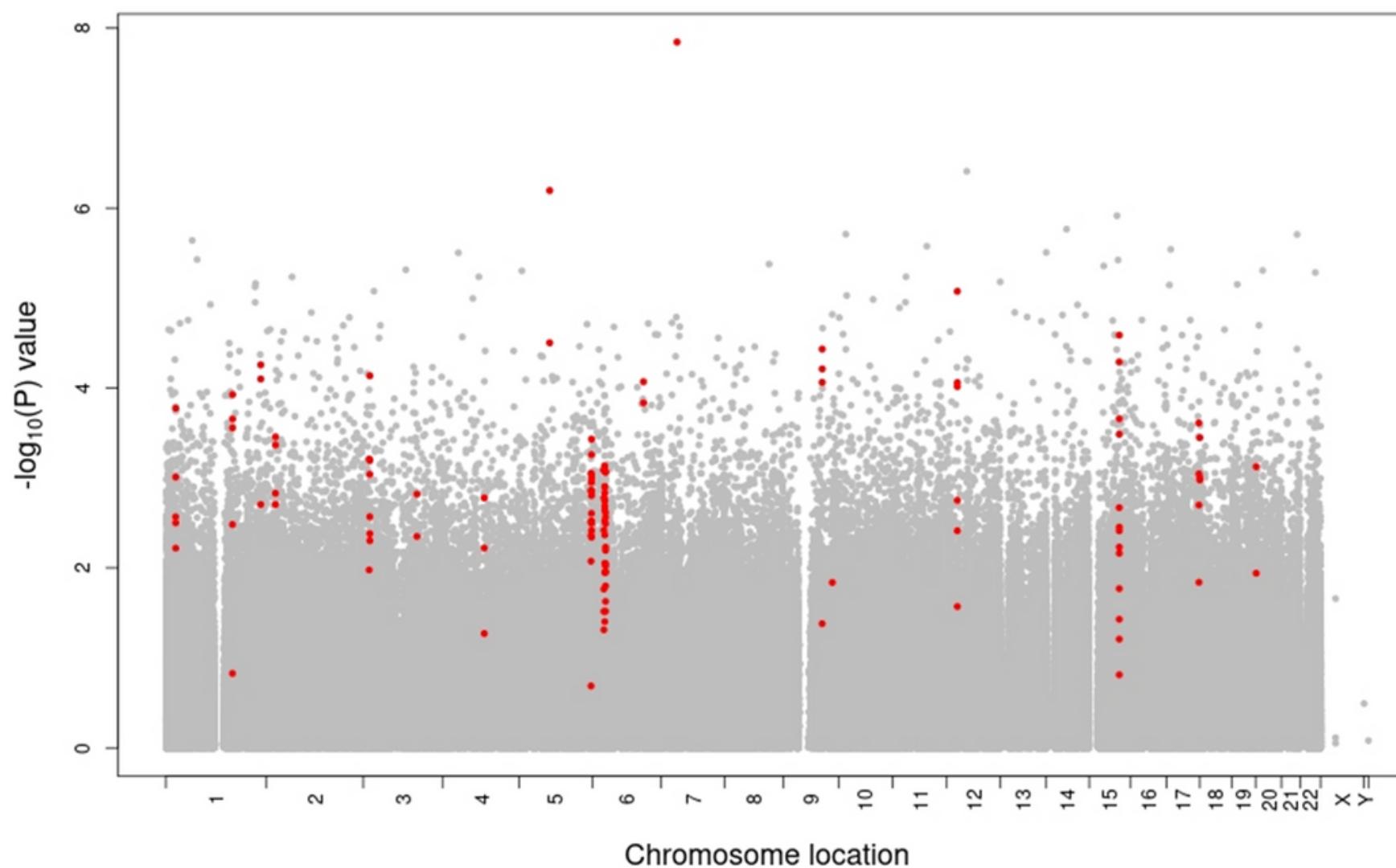


adjusted by age + sex + smoker + ethnicity + fisher

DMR

PATHWAYS

vasospasm



- GO
- Reactome
- KEGG

Óxido nítrico
endotelial

Estrés
oxidativo

Inmunidad e
inflamación

Apoptosis

- * HLA-B, HLA-DPA1, COL11A2 (chr 6)
- * COL23A1 (chr 5)

LIMITATIONS

Sample size
Ewas from blood
Need replication

STRENGTH

largest GWAS and EWAS
Exhaustive study and phenotyping
Pathophysiology



ecuadado@psmar.cat

join us !!

**Neurology Department & Neurovascular Research Group
Hospital del Mar Medical Research Institute (IMIM)
Barcelona(SPAIN)**



J.Jiménez Conde MD, PhD



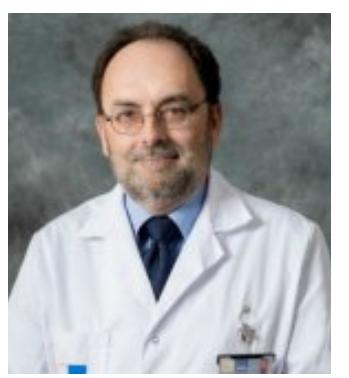
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Marta Vallverdú, MSc PhD



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Jaume Roquer MD, PhD



Antoni Suàrez , MD Adrià Macias, MD



Ana Rodríguez Campello MD, PhD



Eva Giralt MD, PhD



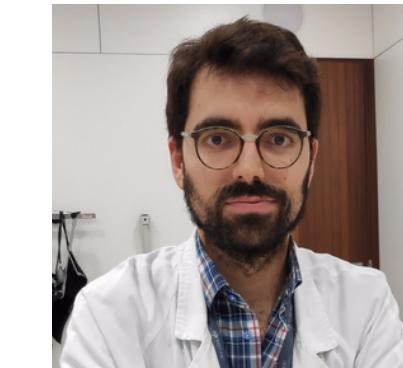
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Daniel Guisado Alonso, MD