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# PhD Open Days

## Genetic Modulation of Stroke in Children with Sickle Cell Anaemia

PhD Program in Biotechnology and Biosciences

Marisa Silva (marisa.silva@insa.min-saude.pt)

Sofia Vargas (INSA, DGH), Andreia Coelho (INSA, DGH), Joana Mendonça (INSA, DGH), Luís Vieira (INSA, DGH; Toxomics), Paula Kjöllerström (HDE), Raquel Maia (HDE), Rita Silva (HDE), Alexandra Dias (HFF), Teresa Ferreira (HFF), Anabela Morais (HSM), Isabel Mota Soares (HGO), João Lavinha (INSA, DGH; BioISI), Paula Faustino (INSA, DGH; ISAMB)

GOVERNO DE **PORTUGAL** 

**TÉCNICO** LISBOA

SINS SERVIÇO NACIONAL DE SAÚDE

## 1. Subject

# Sickle cell anaemia (SCA) is an autosomal recessive genetic disease, caused by a mutation in the HBB

- gene, which results in the synthesis of an abnormal haemoglobin (HbS)
- HbS polymerises inside red blood cells causing them to be sickle-shaped, fragile, rigid and adherent-prone to vessel walls (endothelium) and to other blood cells
- Several pathways also play a role in disease severity such as endothelium dysfunction, cell adhesion, nitric oxide metabolism and haemolysis (Fig.1)
- SCA most catastrophic complication: cerebral vasculopathy (CVA), namely stroke and silent cerebral infarcts (SCIs)
- Current therapeutic approaches for CVA:

 70 SCA children (>= 3 years old) Several degrees of CVA (stroke, SCI, Sample selection risk, normal) Database with clinical, imaging and laboratory data Candidate genes **Genotyping**/ **VCAM1** (endothelial activation) Sequencing • **ITGA4** (cell-endothelium adhesion) **NOS3** (endothelial nitric oxide synthesis)

### **3. Plan of Action**

HOSPITALAR DE LISBOA CENTRAL EPE

HRE

CENTRO HOSPITALAF LISBOA NORTE, EPE

BIOSISTEMS and Integrative

FCT

- Transcranial Doppler, magnetic resonance imaging (MRI) or computerized tomography (CT scan) for risk assessment and diagnosis
- Hydroxyurea and chronic blood transfusions
- Primary CVA prevention and prognosis are still not lacksquaresensitive enough to detect all potentially affected children nor to evaluate long term prognosis



#### 2. Research Question(s)



- How do we design more effective approaches for paediatric CVA prevention in SCA?
- Could genetic variants act as modulators of CVA severity and, if so, could they be used as sensitive/specific biomarkers?



Supervisors:



Dr. Paula Faustino (National Institute of Health Dr. Ricardo Jorge, Human Genetics Department; ISAMB) phdopendays.tecnico.ulisboa.pt Prof. Arsénio Fialho (IST, Bioengineering Department)