

















ITHACA Board Meeting

2020 December 10 – 12

Satellite Meetings : Friday Dec. 11th WG 7 Research

Follow-up on research collaborative call: Pai/OAFNS





Collaborative call – Pai OAFNS



 First call as soon as the collaborative call initiative was launched in May

Targeted gene(s)/phenotype under study: Pai/OAFNS syndrome, ideally after negative exome

Abstract: The Solve-RD project is aiming at identifying genes in diseases resistant to exome sequencing. Pai and OAFNS are overlapping syndromes, belonging to the unsolvable cohorts in solve-RD, taking into account that exomes have remained negative in a number of patients. Multiomics are proposed to identify the molecular bases of these entities. New samples are required, including blood, urine and saliva of the proband and parents, and ideally tissue of affected area if available.

Coordinating clinician: Pr Laurence Faivre

Institution (dept, hospital, City): CHU Dijon/France

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Specific requirements beyond clinical data and genotype data sharing:

- Re-analysis of DNA samples : N
- Resampling of patients No if the trio is available: Y
- Linked to a translational/basic research project? Y, Solve-RD



Collaborative call – Pai OAFNS



Pai syndrome and oculo-auriculo-fronto-nasal syndrome (OAFNS) hypotheses

Constitutional monogenic disease

Mosaic disease

Oligogenic disease

Coding or non-coding (regulatory) variation not detectable by short read sequencing

Variation not detectable in blood

Multiple genes belonging to the same pathway

→ Long read sequencing

→RNA-seq

→ Proteome

→ Epigenome

→ Tissue analysis

→ Multiomics data integration



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Collaborative call – Pai OAFNS



Analysis strategy for the Pai syndrome and OAFNS

Affected tissue available





- SR GS
- Epigenome (methylation DNA)



- LR GS
- RNAseq
- Proteome



- SR GS
- Epigenome (methylation DNA)
- Proteome (plasma)

No affected tissue available



- SR GS
- LR GS
- RNAseq
- Epigenome (methylation DNA)
- Proteome (plasma)

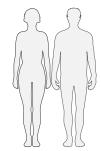


- SR GS
- Deep ES



+/- Deep ES

SR: short-read LG: long-read





- SR GS +/- LR GS
- RNAseq
- Epigenome (methylation)
- Proteome (plasma)



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(WG8) Work Groups / Satellite Meetings : Friday December 11th



	France	ITHACA
Pai/OAFNS included with tissue	4	0
Pai/OAFNS included with no tissue	4	0
Pai/OAFNS with no samples sent	5	8 (UK 2, Italy 1, Spain 1, Netherlands 1, Cyprus 1, New Zeland 2)
Replication cohort	5	3

