





# Actualités des DIH 2020



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CEREDIH, 11/12/2020







### PIDs' genes in 2020



### In 2020: PIDs includes over 450 known genes

### **PAX1 deficiency** (AR-SCID T-B+NK+)

(Paganini 2017, clin genet)



PAX1 member of the paired box (PAX) family of transcription factors and plays a critical role in pattern formation during embryogenesis.

**Otofaciocervical syndrome (OTFCS2) :** facial anomalies, cup-shaped low-set ears, preauricular fistulas, hearing loss, branchial defects, skeletal anomalies, mild intellectual disability.

2 patients from 1 kindred with an AR OTFCS2 and SCID with homozygous mutation in *PAX1* gene. c.1104C>A, p.(Cys368\*)

#### - PAX1 is essential for development and function of the human

thymus (Yamazaki et al. Sci Immunol 2020)

6 SCID pts from 3 kindred with ears abnormalities and homozygous PAX1 mutations c.463\_465del (p.Asn155del), c.1104C>A, (p.Cys368\*), c.439G>C, (p.Val147Leu)

Severe T lymphopenia, few TRECs, normal B cell, ~normal IgM, low IgA, 个 IgE,  $\downarrow$ TTL T cell deficiency not corrected by HSCT despite donor chimerism

Severe T cell ID characterized by a primary thymic defect: FOXN1 deficiency, complete DiGeorge and CHARGE syndromes

#### AR SCID T-B+NK+ caused by inositol-trisphosphate3-kinase B (ITPKB)



*ITPKB* gene encoded for one of 3 isoforms of <u>Inositol-trisphosphate 3-kinase</u> expressed in humans. It regulates immune cell function and is required for T and B cell development.

- <u>1 patient 6 months</u>: failure to thrive, recurrent skin abscess, pneumonia and oral thrush, at 6 months, the absence of a thymic shadow (chest radiograph) She died at 11 months (*Staphylococcal* sepsis)
- Profound T lymphopenia
- Neutropenia, anemia, thrombocytopenia. (bone marrow failure in mice model)

#### **<u>AR SLP76</u>** (AR SCID T-B+NK+, PN and platelets dysfunction)

Inherited SLP76 deficiency in humans causes severe combined immunodeficiency, neutrophil and platelet defects. (Lev et al. , JEM 2020)

<u>1 patient with CS parents: early-onset with</u> infections, AHAI at the age of 2 mo, recurrent skin abscesses and skin rash. BCGite, HSMG,

Respiratory distress at 3 mo,

generalized seizure at 4 mo multiple brain abscesses (Aspergillus) and CMV infection.

patient died of transplant-related complications 30 d.

T CD4 lymphopenia +  $\downarrow$ Trecs,

Combined T and B cell immunodeficiency with severe neutrophil defects and impaired platelet aggregation.

SLP76 molécule adaptatice TCR via LAT SLP76 is a key protein involved in TCR signalling and in other hematopoietic pathways Homozygous mutation C.957+1G>A ; K309FSx17 in the *SLP76* gene affects a donor splice site at the beginning of intron 14





#### **CDH17 deficiency** (AR-CID T-B-NK+)

(Smith et al. 2017, Blood adv)

Cadherin 17 encoded by the *CDH17* gene, member of the cadherin superfamily encoding calcium-dependent membrane-associated glycoproteins

<u>1 patient of 5 years-old</u>: sinopulmonary infections, EBV<sup>+</sup> B cell lymphoma (DLBCL)

**CDH17** compound heterozygous mutation:

Severe T lymphopenia, few TRECs, low T cell proliferation; B lymphopenia (low specific Ab), normal NK cells and function

*CDH17* is expressed in human thymic epithelial cells. *CDH17* mutations may be a rare cause of leaky SCID that can be corrected by HSCT.



c.1796+2 T>C / c.527A>G ; (p.N176S)

### MAN2B2 deficiency (AR-CID)

Defining a new immune deficiency syndrome: MAN2B2-CDG. (Verheijen J, al. JACI 2020)

MAN2B2 : mannosidase alpha class 2B member 2

- <u>1 patient</u> with CID childhood onset (2 weeks) recurrent pneumonia and thrush, small vessels vasculitis (3 mths), arthritis (9 mths), chronic diarrhea microcephaly, neurodevelopmental delay, thrombotic stroke
- No thymic shadow, ↓ T cells (↓ naïve, ↑ T<sub>EMRA</sub>), no TRECs, ↓ TCR proliferation, ↓ total B cells, ↑ plasmablasts, ↑ B <u>CD21IowCD38Iow</u> (54%), ~normal IgM, low IgA, ↑ IgE
- Homozygous *MAN2B2* mutation: <u>p.Asp38Asn</u>
- Mutation affecting both *N*-glycan synthesis and glycan degradation
- Congenital Disorder of Glycosylation (CDG)



### POLR3E deficiency (AR-CID)

A mutation in POLR3E impairs antiviral immune response and RNA polymerase III. (Ramanathan A, et al. PNAS 2020)

<u>**1** patient</u> with homozygous mutation in *POLR3E* gene: p.(D40H) :

Recurrent, systemic viral infections (fatal)

Langerhans cell histiocytosis

**↑** IFN-signature by virus-infected fibroblasts

Increased cell susceptibility to infection by CMV.

D Stalk POLR3H CRCP POLR3C, 3F, 3G DNA/RNA Active site complex POLR3A, 3B Termination complex POLR3D, 3E, 3K

Heterozygous inborn errors in several subunits of PolIII:

POL III is recruited to target genes by transcription factors that facilitate initiation and reinitiation of transcription.

Others PID with genetic defects affecting polymerases: *POLA1, POLA2, POLD1, POLD2, POLE1, POLE2, POLR3A, POLR3C, POLR3E, POLR3F* 

## **MCM10 deficiency** (AR-CID)

Human NK cell deficiency as a result of biallelic mutations in MCM10. (Mace EM, et al. J Clin Invest. 2020)

- **1** patient **16** months severe CMV infection, HLH-like,
- HSCT, died at 24 months (CMV)
- Mild lymphopenia  $\downarrow$  T<sub>CM</sub>, T<sub>EM</sub> cells,  $\downarrow$  T cells prolifs (PHA)
- $\downarrow \downarrow \downarrow \downarrow NK$  cells only CD56<sup>bright</sup> (absent CD56<sup>dim</sup> mature NK)
- ↓ NK function, IgG slightly decrease, IgA and M normal
- human NK cell deficiency (NKD): •
  - GATA binding protein 2 (GATA2-AD),
  - IFN regulatory factor 8 (*IRF8-AD*),
  - minichromosome maintenance 4 (MCM4-AR),
  - go-ichi-nisan complex subunit 1 (GINS1 AR)



c.1744C>T

c.1276C>T



Proband

RPA

CDC45

GINS

Trends in Genetics

MCM10, GINS1 and MCM4 deficiencies are phenocopies. Disorders of NK cell terminal maturation by abnormal requirement for the CDC45-MCM2-7-GINS (CMG) complex.

These NKD strengthens importance of the CMG complex and DNA replicative machinery in human NK cell functional maturation.

#### (Horev et al. 2015 J AM ACAD DERMATOL)

| IL-/ deficiency (AR-CID)                 |  | Patient 1 Patient 2 Patient 3 |        |         |             |     |
|--|--|-------------------------------|--------|---------|-------------|-----|
|  | Age, y   | 39                            | 46     | 51      | 51          | 63  |
| 5 adult patients from 3 kindreds         | Sex  | Female                        | Female | Male    | F           | N   |
| with CID phenotype                       | Cutaneous manifestations                       |                               |        |         |             |     |
| with cid plicitotype.                    | Deep palmoplantar warts                        | +                             | +      | +       | +           | +   |
| Verrucous skin HPV (3,7) infections (5), | Verrucous papules on the                       | +                             | +      | +       | +           | +   |
| Squamous cell carcinomas                 | dorsal aspect of the fingers<br>and toes       |                               |        |         |             |     |
| Cryptococcus neoformans meningitis       | Flat warts (tinea versicolor-<br>like lesions) | -                             | +      | +       |             |     |
| Severe T cell lymphopenia (CD4>CD8)      | Seborrheic keratosis-like                      | _                             | _      | +       |             |     |
|  | warts  |                               |        |         |             |     |
|  | Palmar hyperkeratosis and                      | —                             | -      | +       |             |     |
| Homozygous mutations of <i>IL7</i> gene: | diffuse thickening of the<br>dorsal hands      |                               |        |         |             |     |
| c.205A>T ( p.R69*), c.3G>A (p.Met1?)     | Nonmelanoma skin cancers                       | _                             | _      | + (SCC) |             |     |
|  | Cryptococcal meningitis                        | _                             | +      | +       |             |     |
| Mildor disease that II 7P deficiency     | T-cell parameters                              |                               |        |         |             |     |
| winder disease that it/k deficiency      | T cells $[/\mu L]$ (700-2100)                  | 166                           | 107    | 378     | 45          | 120 |
| (intact TSLPR signaling?)                | CD4 T cells [/µL] (300-1400)                   | 36                            | 31     | 48      | <b>45</b> 1 | 120 |
| Case 1 (B) Case                          | CD8 T cells [/µL] (200-900)                    | 108                           | 94     | 310     |             |     |
|  | Regulatory CD4 T cells                         | 3.8                           | 6.4    | 12.5    |             |     |
|  | [% of CD4 T cells] (2.9-8.1)                   |                               |        |         |             |     |



II 7 deficiency (up any

**Two cases of interleukin-7 (IL-7)-deficient generalized verrucosis.** (Kosumi H, et al. Clin Infect Dis. 2020), start at 10yo

Interleukin-7 plays an essential role in the development, survival, and homeostasis of T cells.

**CD4 deficiency** (AR-CID) (Rosa Anita Fernandes et al., Front Immunol 2019)

Extensive, refractory warts in <u>1 adult patient (45y)</u> Normal lymphocyte counts:  $\downarrow$  CD4+ T cells (<0.01 CD4+ T-cells/µl),  $\uparrow$  CD8+ T cells,  $\uparrow$  DN/TCR $\alpha\beta$ + T cells  $\uparrow$  B cells, normal NK cells, normal Ab production APC MHC II a2 B2 D1 Antigen CD4 TCR

Homozygous mutation: last bp of the 7 intron (g.6818420 G>A, c.1157-1G>A) caused two truncated forms of CD4 RNA/cDNA with <u>no CD4 expression</u> on blood Tcells, monocytes, and DCs.



### **MR1 deficiency** (AR-CID)

Absence of mucosal-associated invariant T cells in a person with a homozygous point mutation in MR1 (Howson LJ, et al. . Sci Immunol. 2020)

- <u>1 patient (31y):</u>
- Tattoo-associated persistent <u>HPV<sup>+</sup></u> <u>warts,</u> *Campylobacter* GI enteritis, VZV pneumoniae
- <u>No MAIT cells</u>; but expanded
   Vγ9/Vδ2+ T cell population,
   ↑ γδ ; mildly ↓ CD4, CD8 T cells, mostly memory

Homozygous mutation in MR1

gene : c.92 G>A; p.Arg31His



MAIT maturation is dependent on the major histocompatibility complex class Irelated protein (MR1) class 1b non-polymorphic molecule of the major histocompatibility system, which is expressed by different types of cells

### AR NFKB1 deficiency (AR CID)

#### Combined immunodeficiency caused by a novel homozygous *NFKB1* mutation.



- CID in a patient of 7 yo with failure to thrive, persistent EBV viremia and hepatitis, *Pneumocystis jirovecii* pneumonitis (7 mths), and generalized lymphadenopathy
- Progressive T cell lymphopenia with ↓↓ CD4 memory, ↑ CD8 T<sub>EM</sub>/<sub>TEMRA</sub>, normal B cell, ↓IgD+CD27+ memory B cells, ↑transitional B , defect specific Ab prod
- Homozygous mutation in NFKB1: c.2878G>A, p.Gly960Arg (G960R).
- This affected p105 phosphorylation and p50 formation on Ag and cytokine stimulation, as well as attenuating nuclear signal transmission.
- Blockade of NF-κB pathway signalling, resulting in aberrations in T- and B-cell maturation and function.

## **APRIL/TNFSF13 deficiency:** AR CVID

#### APRIL-dependent lifelong plasmacyte maintenance and immunoglobulin production in humans

(Tzu-Wen Yeh et al. JACI 2020)



A proliferation-inducing ligand (APRIL, TNF ligand superfamily), is secreted from myeloid cells and known to induce the differentiation of memory B cells to plasmocytes.

APRIL is critical for lifelong maintenance of plasmocytes to produce Ig in humans.

## **CTNNBL1 deficiency** (AR-CIVD)

(Disease-associated CTNNBL1 mutation impairs somatic hypermutation by decreasing nuclear AID (Marcel Kuhny, //Jordan S. Orange, Eric Meffre; JCI 2020)

CTNNBL1:  $\beta$ -catenin–like protein 1

Homozygous: c.1396A>G; p.M466V

1 patiente 15 yo with progressive hypogamma, autoimmune cytopenias, recurrent infections, hyperplastic GC's (hyperplastic GC reactions ↑ T follicular helper–like

Hypogamma, ↓ memory B cells;, expanded peripheral autoimmune/autoreactive CD19hiCD21-/loCD10-CD27- B cell subset, ↓ Tregs, Impaired CSR and SHM

| Immunoglobulin levels (mg/dL) | (before IVIL) |
|-------------------------------|---------------|
| lgG                           | 198           |
| IgA                           | 13.7          |
| lgM                           | 5.7           |
| lgE (IU/mL)                   | <4.0          |



#### Mutation M466V $\downarrow$ binding of CTNNBL1 to AID $\Rightarrow$ Less nuclear translocation of AID.

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### **AR FNIP1 deficiency** (AR-PAD)

Mutations of the gene *FNIP1* associated with a syndromic autosomal recessive immunodeficiency with cardiomyopathy and pre-excitation syndrome. (Niehues et al. Eu J I 2020)

FNIP1 forms a complex with the γ2 subunit of phosphorylated Adenosine monophosphate-activated protein kinase (AMPK)= Master sensor of energy consumption and activated during energy deprivation

**3 pts from 2 kindreds with homozyg. mutation in FNIP1 gene:** K1 exon 17 (c.3353G>A; p.Ser1118Asn K2 exon 12 (c.1289delA). p.His430Profs7\* **No FNIP1 protein expression** 

Fnip1 is essential for pre-B cell development and Myc-induced lymphomagenesis



#### Early onset (<6 months),

- ENT and lung infections viruses and bacteria, encapsulated bacteria (pneumonia, meningitis)
- Pulmonary: obstructive bronchitis, pneumonic abscess, bronchiectasis, fibrosis, cell-rich interstitial pneumonia.
-Gut infections (*Campylobacter, Rotavirus, Salmonella*)

- Cardiac: Wolff-Parkinson- White syndrome, metabolic myopathy, muscular hypotonia
- ↓ B cells , ↓ switched memory B cells and hypogammaglobulinemia (IgG, IgM), intermittent neutopenia

- BOM : B-cell maturation defect at the pre-B-cell stage with an enrichment of CD34-CD10+CD21Iow B-cell precursors

### AD IL6ST deficiency AD HIES (DN)

Dominant-negative mutations in human IL6ST underlie hyper-IgE syndrome.



- 12 pts from 8 Kds with AD-HIES : DN *IL6ST* mutations.
- Mutant alleles encode GP130 receptors lacking both the recycling motif and all 4 STAT3-recruiting tyrosine residues.
- Overexpression, the mutant proteins accumulate at the cell surface and are LOF + DN for cellular responses to IL-6, IL-11, LIF, OSM.
- The patients' cells respond poorly to IL-6 and IL-11.
- Patients with *STAT3* or *IL6ST* mutations display infectious and allergic manifestations of IL-6R deficiency and some of the skeletal abnormalities of IL-11R deficiency.

DN STAT3 and IL6ST mutations thus appear to underlie clinical phenocopies through impairment of the IL-6 and IL-11 response pathways.



-Y-P

STAT3

gp130

IL-6 receptor

JAK

mRNA

P-Y

HP-2

APKs

### AR IL6ST deficiency

- <u>Cytokine-selective defects in GP130</u>: Homozygous missense variants in *IL6ST* (c.1210A>T; p.Asn404Tyr and c.1493C>T; p.Pro498Leu) cause immunodeficiency with elevated IgE and craniosynostosis (Schwerd et al.2017)

- <u>Cytokine-complete defects in GP130</u>: Fatal Stuve-Weidemann syndrome, 5 pts from 3 kindreds  $\Rightarrow$  Absence of GP130 cytokine receptor signaling causes extended Stuve-Wiedemann syndrome. (Chen YH, et al. J Exp Med. 2020)

- Homozygous variants in *IL6ST* (c.841C>T;p.Arg281\*, c.1699+4A>G)
- osteoporosis, skeletal dysplasia,
- spontaneous fractures, hyperextensibility of joints, scoliosis
- lung dysfunction, renal abnormalities thrombocytopenia, dermatitis, eczema.
- Fatal outcome during between neonatal to childhood

Defective acute phase response.

Complete unresponsiveness to IL-6 family cytokines (IL-6, IL-11, IL-27, OSM, LIF)

Genetic syndrome caused by the <u>complete lack of signalling of a whole family of</u> <u>GP130-dependent cytokines</u> in humans and highlights the importance of the LIF signalling pathway in pre- and perinatal development.

### **AR ILGR deficiency** – AR HIES

Loss of the interleukin-6 receptor causes immunodeficiency, atopy, and abnormal inflammatory responses (Spencer et al, JEM 2019)









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# Impaired IL-6 immunity and HIES

#### AD; AR HIES

The nature of human IL-6

**JEM 2019** 

Anne Puel<sup>1,2</sup> and Jean-Laurent Casanova<sup>1,2,3,4,5</sup>

| Gene defect   | High<br>IgE | Eczema | Staphylococcal<br>disease | Inflammation | Extra-hematopoiet<br>features | ic IL-6<br>response            | References                                       |
|---|-------------|--------|---------------------------|--------------|-------------------------------|--------------------------------|--|
| TYK2 <sup>-/-</sup> (Japanese <b>AR</b><br>patient) | +           | +      | +                         | ND           | -1                            | -                              | Minegishi et al., 2006                           |
| TYK2 <sup>-/-</sup> (other patients)                | 2           | 192    |                           | +            |                               | +                              | Kreins et al., 2015                              |
| ТҮК2 <sup>р1104A/р1104A</sup>                       | -           | :00    | -                         | +            | -3                            | +                              | Boisson-Dupuis et al.,<br>2018                   |
| STAT3+/- AD   | +           | +      | +                         | -            | +                             | <u> </u>                       | Minegishi et al., 2007                           |
| ZNF341-/- AR  | +           | +      | +                         | +/-          | +/-                           | -                              | Béziat et al., 2018;<br>Frey-Jakobs et al., 2018 |
| IL6ST-/-a AR/AD                                     | +           | +      | +                         |              | + 6                           | Beziat <del>e</del> t al. 2020 | Schwerd et al., 2017                             |
| ILGR-/- AR  | +           | +      | +                         |              | <b>3</b>                      | -                              | Spencer et al., 2019                             |
| Anti-IL-6 autoantibodies                            | -           | -      | +                         | -            | <u>14</u> 3                   | -                              | Puel et al., 2008                                |
| Erbin+/- AD   | +           | +      | +                         | -            | +                             | -                              | Lyons et al., 2014                               |

HyperIgE phenotype: eczema, recurrent infections and elevated serum IgE:

STAT3 (AD), ERBB2IP (Erbin AD), IL6ST (AR, AD), CARD11 (AD), IL6R (AR), ZNF341 (AR), DOCK8 (AR), PGM3 (AR)

### AD Sec61α1 deficiency (AD-SCN)

#### Defective SEC61α1 underlies a novel cause of autosomal dominant severe congenital neutropenia (Van Nieuwenhove et al. JACI 2020): gene SEC61A1 c.A275G;p.Q92R



ER, endoplasmic reticulum; UPR, unfolded protein response, ★ missense mutations

# Specific mutations in *SEC61A1* cause SCN through dysregulation of the unfolded protein response (UPR).

### AR NOS2 deficiency (AR-CMV)

Fatal Cytomegalovirus Infection in an Adult with Inherited NOS2 Deficiency. (Drutman SB et al. N Engl J Med. 2020)

Severe susceptibility to CMV-induced disease, previously healthy man (51 yo) who after acute CMV infection had an onset of progressive CMV disease that led to his death 29 months later.

 $\downarrow$  CD4+ T cells;  $\downarrow$  NK cells, normal CD8+ T,  $\downarrow$  B cells



B CT of the Patient's Lungs at Presentation





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c.1436\_1437insT p.lle3911lefsTer26

### **AR CD16A deficiency** (AR-EBV)

Identification of the first cases of complete CD16A deficiency: Association with persistent EBV infection (A. Perez-Portilla et al. JACI 2020)

1 pt 11 yo and an extended workup revealed persistently high levels of EBER DNA in peripheral blood an multiple-organ infiltration with EBV-positive T lymphocytes including an IgA- and EBV-positive interstitial nephritis (CAEBV)

Normal NK cell numbers but ,  $\downarrow$  adaptive NK (CD56<sup>dim</sup>CD57<sup>hi</sup>); Impaired NK cell ADCC



The CD16A-deficient individuals had inherited a maternal chromosome where the FCGR3A gene with deletion and a paternal chromosome that encoded a hybrid FCGR3B/A gene C. Picard 11/12/2020

### **AR DEF6 deficiency** (AR-EBV)

#### DEF6 deficiency, a mendelian susceptibility to EBV infection, lymphoma and autoimmunity (B. Fournier, JACI 2020)

DEF6 is a specific guanine nucleotide exchange factor for the Rho GTPase Cdc42 and Rac1. DEF6 is involved in TCR signalling through its recruitment to the immune synapse.

A homozygous mutation in *DEF6* (WES): c.940C>, p.Q314\* in 4 members of a CS family with autoimmunity and susceptibility to EBV (EBV-associated lymphoproliferation and lymphoma) <u>without</u> extra-hematopoietic manifestations.



### AD CD48 deficiency (AD-EBV)

Recurrent inflammatory disease caused by a heterozygous mutation in CD48 (Volkmer B, JACI 2019)

1 pt age of 9 mths onward in a patient now 24yo fever, polymorphic rashes, SMG, Pancytopenia, 个 ferritinemia, 个TG, HLH-like, partial cytotoxic defect (T/NK cells)

 $\uparrow \uparrow HLA-DR+CD8 T cells; Mild \downarrow CD4, T cells;$ 

↑ immature Nkbright / ↓ mature NK cells, Hemophagocytosis in the bone marrow

The patient recovered spontaneously from these episodes (4 to 7 days); mild SMG and CD4 lymphopenia between episodes

**CD48 expression is needed to :** 

- (1) Control inflammation,
- (2) ensure correct NK cell maturation and function,
- (3) regulate target cell susceptibility to cytotoxicity.

CD48 : c.659C>A, p.S220Y (haploinsufficiency)







#### PID reveal the molecular requirements for effective host defense against EBV infection

Stuart G. Tangye, Sylvain Latour, Primary immunodeficiencies reveal the molecular requirements for effective host defense against EBV infection, Blood, 2020,



# AD BPIFA1/SPLUNC1 (AD-DN: Meningo)

A Rare Mutation in SPLUNC1 Affects Bacterial Adherence and Invasion in Meningococcal disease. (Mashbat B et al. Clin Infect Dis. 2020)

- 3 pts from 2 kindreds: Invasive meningococcal disease (*Neisseria meningitidis*) and heterozygous *BPIFA1 mutation:* c.65G>A, p.(Gly22Glu)
- The dominant negative (DN) mutant recombinant *BPIFA1* (p.Gly22Glu) showed reduced antibiofilm activity, increased meningococcal adhesion, and increased invasion of cells, compared with WT *BPIFA1* cells.



Conclusions: A mutation in *BPIFA1* affecting mucosal attachment, biofilm formation, and invasion of mucosal epithelial cells is a new genetic cause of meningococcal disease.

### **AR IFNG deficiency** (AR-MSMD)

Inherited human IFNgamma deficiency underlies mycobacterial disease. (Kerner G, Rosain J et al. J Clin Invest. 2020)

#### 2 cousins: disseminated mycobacterial disease (BCGosis) homozygous mutation in *IFNG* gene:

c.354\_357del; p.(T119Ifs4\*), Loss of expression and loss of function.

Patients T and NK cells from also failed to produce and secrete detectable amounts of IFN-γ

The patients' herpesvirus Saimiri–immortalized T lymphocytes: No IFN-y production

Biological phenotype rescued by retrotransduction with WT *IFNG* cDNA.



## AR TBX21 (Tbet) deficiency (AR-MSMSD)

Human T-bet governs innate and innate-like adaptive IFN-γ immunity against mycobacteria. (Yang R //Casanova JL. Cell. 2020)

- **1 MSMD patient** : p.Glu156\_Met157delins homozygote; ↓ IFN-γ producing T cells
- CD4 and CD8 T cells, naïve and memory subsets normal %
- ↓ CXCR3<sup>+</sup> Th1 cells,
- ↑ immature NK, ↓ mature NK
- $\downarrow$  NKT,  $\downarrow$  MAIT, V $\delta$ 2<sup>+</sup>  $\gamma\delta$  T cells



Impairment of development and fonction: T CD4, NK, NKT, MAIT, γ/δ2, γ/δ T cells No IFNγ production by T cells

Nature Reviews | Immunology

The T-box transcription factor T-bet (Tbx21) is a key regulator of type 1-like immunity, playing critical roles in the establishment and/or maintenance of effector cell fates in T and B lymphocytes, as well as DC and NK cells.

### AR SNORA31 deficiency (AR-HSE)

Human SNORA31 variations impair cortical neuron-intrinsic immunity to HSV-1 and underlie herpes simplex encephalitis (Lafaille FG, et al. Nat Med. 2019)

*SNORA31,* encoding a small nucleolar RNA of the H/ACA class snoRNA31 function: a guide RNA directing the chemical modification of target uridine residues into pseudouridine in rRNA and small nuclear RNA28

**5 patients with Herpes Simplex Encephalitis (HSE):** n.36T>C, n.75C>G, n.96T>G, n.111T>C

Transcriptome analysis of SNORA31-mutated neurons revealed normal responses to TLR3 and IFN- $\alpha/\beta$  stimulation but abnormal responses to HSV-1



Deficient SNORA31 iPSC's cortical neurons are susceptible to HSV1 infection.

It is a new genetic aetiology and immunological mechanism of HSE, involving a disruption of CNS cortical neuron intrinsic immunity to HSV-1 with maintenance of the TLR3 and IFNAR1/IFNAR2 response pathways.

### AD MAPK8 deficiency (AD-CMC)

Chronic mucocutaneous candidiasis and connective tissue disorder in humans with impaired JNK1-dependent responses to IL-17A/F and TGF-beta (Li J, Ritelli M,// Puel A et al. Sci Immunol. 2019)

Chronic Mucocutaneous Candidiasis (CMC) and connective tissue disorder (similar to Ehlers-Danlos syndrome)

AD mutation in MAPK8 gene (haploinsufficiency): c.311+1G>A, 3 pts

**MAPK8** gene encodes JNK1, one of the 3 members of the JNK family. This protein is a component of the mitogen-activated protein kinase (MAPK) pathway

- ↓ functional Th17 cells *ex vivo, in vitro*
- ↓ responses of fibroblasts to IL-17A, IL-17F
- **\downarrow** TGFβ signalling



The integrity of the human JNK1 pathway is essential for IL-17A/F-dependent mucocutaneous immunity to Candida and for the TGF-β-dependent homeostasis of connective tissues.

### **Susceptibility to COVID-19:**

Inborn errors of type I IFN immunity in patients with life-threatening COVID-19 (Zhang et al., Science 2020)

3.5% of patients with life threatening COVID-19 pneumonia had :

- known: <u>AR</u> IRF7 and IFNAR1 deficiencies or <u>AD</u> TLR3, TICAM1, TBK1 and IRF3 deficiencies
- or new AD UNC93B1, IRF7, IFNAR1, and IFNAR2 deficiencies genetic defects (Haplo, 1 DN)



Genes involved in the TLR3- and IRF7-dependent induction and amplification of type I IFNs

This discovery reveals essential roles for both the double-stranded RNA sensor TLR3 and type I IFN cell-intrinsic immunity in the control of SARS-CoV-2 infection.

| Gene    | Inheritance | Genetic form | Genotype                | Gender | Age [years] | Ancestry/residence    | Outcome  |
|---------|-------------|--------------|-------------------------|--------|-------------|-----------------------|----------|
| TLR3    | AD          | Known        | p.Ser339fs/WT           | М      | 40          | Spain                 | Survived |
| TLR3    | AD          | Known        | p.Pro554Ser/WT          | M      | 68          | Italy                 | Survived |
| TLR3    | AD          | Known        | p.Trp769*/WT            | M      | 77          | Italy                 | Survived |
| TLR3    | AD          | Known        | p.Met870Val/WT          | M      | 56          | Colombia/Spain        | Survived |
| UNC93B1 | AD          | New          | p.Glu96*/WT             | M      | 48          | Venezuela/Spain       | Survived |
| TICAM1  | AD          | Known        | p.Thr4lle/WT            | M      | 49          | Italy                 | Survived |
| TICAM1  | AD          | Known        | p.Ser60Cys/WT           | F      | 61          | Vietnam/France        | Survived |
| TICAM1  | AD          | Known        | p.Gln392Lys/WT          | F      | 71          | Italy                 | Deceased |
| TBK1    | AD          | Known        | p.Phe24Ser/WT           | F      | 46          | Venezuela/Spain       | Survived |
| TBK1    | AD          | Known        | p.Arg308*/WT            | M      | 17          | Turkey                | Survived |
| IRF3    | AD          | Known        | p.Glu49del/WT           | F      | 23          | Bolivia/Spain         | Survived |
| IRF3    | AD          | Known        | p.Asn146Lys/WT          | F      | 60          | Italy                 | Survived |
| IRF7    | AR          | Known        | p.Pro364fs/p.Pro364fs   | F      | 49          | Italy/Belgium         | Survived |
| IRF7    | AR          | Known        | p.Met371Val/p.Asp117Asn | M      | 50          | Turkey                | Survived |
| IRF7    | AD          | New          | p.Arg7fs/WT             | M      | 60          | Italy                 | Survived |
| IRF7    | AD          | New          | p.Gln185*/WT            | M      | 44          | France                | Survived |
| IRF7    | AD          | New          | p.Pro246fs/WT           | M      | 41          | Spain                 | Survived |
| IRF7    | AD          | New          | p.Arg369GIn/WT          | М      | 69          | Italy                 | Survived |
| IRF7    | AD          | New          | p.Phe95Ser/WT           | M      | 37          | Turkey                | Survived |
| IFNAR1  | AR          | Known        | p.Trp73Cys/Trp73Cys     | M      | 38          | Turkey                | Survived |
| IFNAR1  | AR          | Known        | p.Ser422Arg/Ser422Arg   | M      | 26          | Pakistan/Saudi Arabia | Deceased |
| IFNAR1  | AD          | New          | p.Pro335del/WT          | F      | 23          | China/Italy           | Survived |
| IFNAR2  | AD          | New          | p.Glu140fs/WT           | F      | 54          | Belgium               | Survived |

Table 1. Disease-causing variants identified in patients with life-threatening COVID-19.

(Zhang et al., Science 2020)

### **XL-TLR7 deficiency** (XL-Covid19)

#### Presence of Genetic Variants Among Young Men With Severe COVID-19



The 4 male patients mean age of 26 years (range: 21-32), with severe COVID-19 infection, requiring mechanical ventilation.

WES : loss-of-function hemizygous variants of the X-chromosomal *TLR7* gene: F1:c.2129\_2132del; p.[Gln710Argfs\*18] ; F2 (c.2383G>T; p.[Val795Phe].

Primary peripheral blood mononuclear cells from the patients, significantly decreased mRNA expression of *IRF7*, *IFNB1*, and *ISG15* on stimulation with the TLR7 agonist imiquimod The production of IFN-γ was decreased in patients in response to stimulation with imiquimod. = IFN-γ production defect

### **AR TET2 deficiency** (AR-ALPS-like)

Germline TET2 loss of function causes childhood immunodeficiency and lymphoma.

(Stremenova Spegarova J, et al. Blood. 2020)

3 children with an immune dysregulation syndrome, susceptibility to infection: lymphadenopathy, HSMG, developmental delay, autoimmunity (cytopenia), lymphoma of B-cell (n:2, 1 EBV) or T-cell (n:1).

All 3 showed early autologous T-cell reconstitution after HSCT.



Mutated TET2 protein was absent or enzymatically defective for 5hydroxymethylating activity, resulting in whole-blood DNA hypermethylation.

### AD SOCS1 deficiency (AD autoimmunity)

Early-onset autoimmunity associated with SOCS1 haploinsufficiency (Hadjadj et al. Nature Com 2020) and several others reports

c.368C>G,p.(P123R); c.24delA,p.(A9fs\*76); c.64C>T, p.(R22W); c.460T>C, p.(Y154H c.476\_480dupGCCGC,p.M161fs\*46

10 patients (5 families) with early onset autoimmune manifestations: heterozygous (AD) germline loss-of-function mutations in the *SOCS1* gene (incomplete penetrance)

Early onset severe multisystemic autoimmunity (flared in context of infection-induced inflammation):

- ITP, AIHA, Evan's syndrome
- SLE, GN, psoriasis, arthritis, thyroiditis.....
- Lymphoma, syndrome lymphoproliferatif
- Recurrent bacterial infections,
- ↓ marginal zone B cells
- $\downarrow$  switched memory B cell
- ↑ CD21Iow CD38Iow B-cell
- **↑**plasma B-cell activating factor (BAFF) levels

Auto-antibodies,  $\checkmark$  Treg and suppressive fct



The intracellular protein SOCS1 downregulate cytokine signalling by inhibiting the JAK-STAT pathway

SOCS1 insufficiency provide mechanistic insight into the development of autoimmunity induced by cytokine hypersensitivity in immune cells following the loss of a downregulatory element.

#### **AR GIMAP5 and GIMAP6 deficiencies** (AR autoimmunity)

Loss of GTPase of immunity-associated protein 5 (Gimap5) promotes pathogenic CD41 T-cell development and allergic airway disease (Patersson et al. JACI 2019)

A young adult patient with history of impaired viral immunity, auto-immunity. SMG

Progressive T-cell lymphopenia (CD4 > CD8) loss of naïve T-cells, particularly CD4 T-cell Abnormal T-cell function *in vitro*.

WES a homozygous mutation Leu204Pro in gene encoding GTPase IMAP Family Member 5 (*GIMAP5*)

Absent protein expression in T-cells. Impaired survival of T-cells that partially could be rescued *in vitro* by lithium.

GIMAP-5 deficiency is a disorder of immunodysregulation/immune-senescence.

A Human Case of GIMAP6 Deficiency: A Novel Primary Immune Deficiency. (Shadur B, et al. Eur J Hum Genet. In press)

2 siblings with sinopulmonary infections mild hepatosplenomegaly, Patient with a lupus-like PID, SMG Autoimmunity.

#### ~near normal T cells and subsets

- ↑ DN ab T cells
- **T** cell apoptosis

   **near normal B cells/subsets CD21low B cells, Normal Ig (GAM)**

GIMAP6 is a novel PID-associated gene = cell-specific regulator of autophagy.

(GIMAPs) GTPase of the immunity associated protein family are a protein family small GTPases. Immunological functions, such as thymocyte development, apoptosis of peripheral lymph. and T helper cell differentiation

#### **AD PTPN2 Deficiency in Early-Onset Intestinal Autoimmunity**

Loss-of-Function Mutation in PTPN2 Causes Aberrant Activation of JAK Signaling Via STAT and Very Early Onset Intestinal Inflammation. (Parlato M, et al. Gastroenterology. 2020)

- a 3-year-old girl (P1) born from healthy non-consanguineous parents presented with severe chronic secretory diarrhea and eczema since the age of 3 months.
- De novo a heterozygous missense variant in *PTPN2* gene: c.646T>G, p. Cyst216Gly, haploinsufficiency
- PTPN2 encodes an ubiquitous non-receptor protein tyrosine phosphatase that exerts a <u>negative feedback</u> on the JAK-STAT pathway



#### AD RIPK1 deficiency (AD CID-autoinflammation)

A dominant autoinflammatory disease caused by non-cleavable variants of RIPK1. (Tao P et al. Nature. 2020)

Mutations that prevent caspase cleavage of RIPK1 cause autoinflammatory disease.

(Lalaoui N et al. Nature. 2020)

**RIPK1** is a key regulator of innate immune signalling pathways = CRIA syndrome

Heterozygous missense mutations D324N, D324H, D324V, D324Y, D138N (DN)  $\Rightarrow$  prevent caspase cleavage of RIPK1 in humans

 Early onset (<1month): 12 pts from 5 kindreds: early-onset periodic fever syndrome (3-5days) and severe intermittent lymphadenopathy, HSMG, ulcers, tonsillitis, arthralgia, GI features,

↑inflamm markers, ↑ pro-inflamm cytokines/gene signature;
 with responsive to Tocilizumab (IL-6 inhibitor), not to IL1/TNF blockade.
 ↑ counts of both double-negative T cells and naive B cells. ↓ T CD4 naïf and T CD8



### AD CDC42 deficiency (AD-auto-inflammation)

A novel disorder involving dyshematopoiesis, inflammation, and HLH due to aberrant CDC42 function ( (Tam et al. JEM 2019), + several others reports.....)

<u>Neonatal onset:</u> pancytopenia, fever, rash, HSMG, HLH, myelofibrosis, multisystemic inflammatory disease

> 15 patients

Recurrent infections; enterocolitis, diarrhea with intestinal bleedings, neurodevelopmental delay (+/-). Treated: Anakinra/IFNγ mAb (IL-1 and IFNγ neutralization) and HSCT.

- **↑**Serum levels of IL-1, IL-18, IFNγ, ferritin, sCD25, CRP
- $\downarrow$  NK function (cytox),

↑ central memory CD4+ cells with  $\downarrow$  T naive CD31+ cells, ↑ Tregs with high memory phenotype

 $\uparrow$  B cell switched memory cells, plasma cells and autoreactive B cells; and  $\downarrow$  transitional B cells,

 $\downarrow$  PN, DC and monocytes

<u>c.556C>T ; (p.R186C),</u> c.563G>A; (p.C188Y), c.576A>C; (p.\*192C\*24), c.242G>A; p.Cys81Tyr

Mutations haplo-insuf. affects actin function. The mutated protein impaired interaction with regulators and effectors, (RhoGDI, IQGAP1, WASP), leading to aberrant subcellular localization, actin cytoskeleton rearrangement, and reduced migration.



#### Autoinflammatory syndrome : CDC42 cause a novel IL-1 inhibition – responsive

CDC42 encodes a small Rho family GTPase that regulates multiple signalling pathways controlling cell polarity and migration, actin polarization, cytoskeletal architecture, endocytosis, and cell-cycle progression.

### **<u>AR HEM1/NCKAPL1 deficiency</u>** (AR auto-inflammation)

HEM1 deficiency disrupts mTORC2 and F-actin control in inherited immunodysregulatory disease

(Cook et al., Science 369, 202–207 2020); and several other reports

>10 pts from 8 families :

PID with atopy, lymphoproliferation, hyperinflammation Recurrent URTI, skin rashes /abscesses, ulcers, SLE-like, lymphadenopathy, fever, HLH-like

Normal T cell numbers ,  $\uparrow T_{CM}$ , exhausted cells;

 $\downarrow$  T cell proliferation,

Normal B cells and naïve/memory subsets;

 $\uparrow$  CD21<sup>low</sup> cells, Normal/  $\downarrow$  Ig levels ,  $\uparrow$  IgE autoAbs, cytoskeletal defects; immature NK cells but intact function,

Cytokine overproduction (Th1) anti-dsDNA Abs,



WAVE regulatory complex (WRC)

Human patients with immunodeficiency and immune hyperactivation with LOF mutations in *NCKAP1L*, the gene encoding HEM1, disrupt WRC-mediated actin polymerization and abrogate mTORC2 activation of AKT.

WRC deficiency caused by the absence of the WRC hematopoietic cell–specific subunit HEM1 results in a human disorder with <u>severe immune dysregulation and recurrent infections.</u>

#### Inborns errors of immunity affecting the actin cytoskeleton



#### **XL-UBA1** (XL somatic mutation - autoinflammation)

Somatic Mutations in UBA1 and Severe Adult-Onset Autoinflammatory Disease. (Beck DB et al. N Engl J Med. 2020)

- 25 male patients 64yo (45-80) with adult onset treatment-refractory inflammatory synd.
- Fevers, cytopenias, dysplastic bone marrow, neutrophilic cutaneous and pulmonary inflammation
- Inflammatory syndrome (relapsing polychondritis, Sweet's syndrome, polyarteritis nodosa, giant-cell arteritis) or a hematologic condition (MDS, multiple myeloma)
- Often fatal



(Met41Val, Met41Thr, Met41Leu) Mutations > half the hematopoietic stem cells, including peripheral-blood myeloid cells but <u>not</u> lymphocytes or fibroblaste

| VEXAS syndrome    |
|-------------------|
| Vacuoles,         |
| E1 enzyme,        |
| X-linked,         |
| Autoinflammatory, |
| Somatic           |

Mutant peripheral-blood cells  $\downarrow$  ubiquitylation and activated innate immune pathways.

### **AR STAT2 GOF**

c.443G>A, p.R148Q

Homozygous STAT2 gain-of-function mutation by loss of USP18 activity in a patient with type I interferonopathy (Gruber et al. JEM 2020)

1 patient from CS parents with early-onset, severe autoinflammation evocative of severe type I interferonopathy. Skin ulcerations, seizures, cerebral calcifications, and ultimately respiratory failure and death

less myeloid and NK cells and more B and T cells, mRNA levels of four ISGs (*IFIT1, IFI27, RSAD2, ISG15*) all were elevated >1000-fold / healthy donor

The defect in USP18 trafficking to the receptor and explains both the GOF and recessive inheritance of STAT2 R148Q





#### **AR RCH1 deficiency** (AR-autoinflammation - HLH)

A human immune dysregulation syndrome characterized by severe hyperinflammation with a homozygous nonsense Roquin-1 mutation. (Tavernier SJ, et al. Nat com 2019)

One patient 18-year-old male, who was referred at age 11 suffering from hyperinflammation clinically ~ hemophagocytic lymphohistiocytosis (HLH)

WES : a homozygous nonsense R688X *RC3H1* mutation

RC3H1 encodes Roquin-1 a post-transcriptional repressor of immune-regulatory proteins such as ICOS, OX40, CTLA4, Rel and TNF.

The R688X variant reveals a phenotypic immune cell activation, hypercytokinemia and disease development (similar mice model)



R688X Roquin-1 fails to localize to P-bodies and interact with the CCR4-NOT deadenylation complex, impeding mRNA decay and dysregulating cytokine production.

The results suggest that impaired Roquin-1 function provokes hyperinflammation by a failure to quench immune activation.

#### Biallelic *RIPK1* mutations in humans cause severe **AR-CID** immunodeficiency, arthritis, and intestinal inflammation P4 P3 P1.P2 153 230 289 312 Delphine Cuchet-Lourenco<sup>1\*</sup>, Davide Eletto<sup>1\*</sup>, Changxin Wu<sup>1\*+</sup>, Vincent Plagnol<sup>2</sup>, Olivier Papapietro<sup>1</sup>, 671 James Curtis<sup>1</sup>, Lourdes Ceron-Gutierrez<sup>3</sup>, Chris M. Bacon<sup>4,5</sup>, Scott Hackett<sup>6</sup>, Badr Alsaleem<sup>7</sup>, Mailis Maes<sup>1</sup>, Miguel Gaspar<sup>1</sup>, Ali Alisaac<sup>1,8</sup>, Emma Goss<sup>1</sup>, Eman AlIdrissi<sup>9</sup>, Daniela Siegmund<sup>10</sup>, Harald Wajant<sup>10</sup>, Dinakantha Kumararatne<sup>3</sup>, Mofareh S. AlZahrani<sup>9</sup>, Peter D. Arkwright<sup>11</sup>, Kinase domain Intermediate domain Death domain Mario Abinun<sup>12</sup>, Rainer Doffinger<sup>3</sup>, Sergey Nejentsev<sup>1</sup><sup>‡</sup> p.Y289X, Science 2018 p.N230MfsX8, **Receptor Interacting Serine/Threonine Kinase 1 (RIPK1) = master regulator of Deletion ex 4** • signaling pathways leading to inflammation + cell death

- 4 pts from 3 kindreds: homozygous (AR) mutations *RIPK1*: 1<sup>st</sup> month onset
- Recurrent infections (CMV, VZV, RSV, M.avium, bacteria, Aspergillus),
- Chronic lung disease DDB, Early-onset severe IBD and progressive polyarthritis.
- Lymphopenia (T CD4, T CD8, <u>NK</u>), pro-inflammatory cytokines production  $\downarrow \downarrow$



#### Actualités 2020

#### Combined immunodeficiency (CID):

- AR PAX1 deficiency (SCIDT-B+NK+)
- AR ITPKB deficiency (SCIDT-B+NK+)
- AR SLP76 (SCIDT-B+NK+, PN and platelets dysfunction)
- AR CDH17 deficiency (CID T-B-NK+)
- AR MAN2B2 deficiency (CID-CDG)
- AR POLR3E deficiency (CID-virus)
- AR IL7 deficiency (CID-HPV)
- AR MCM10 deficiency (CID-HPV)
- AR CD4 deficiency (CID-HPV)
- AR MR1 deficiency (CID-HPV)
- AR NFKB1 deficiency (CID)

#### **B cell deficiency** (PAD):

- AR TNFSF13 deficiency (APRIL, CVID)
- AR CTNNBL1 deficiency (CIVD)
- AR FNIP1 deficiency (PAD, WPW)

#### HyperIgE syndrome :

- AD (DN) IL6ST deficiency (partial gp130 def.)
- AR IL6ST deficiency (complete gp130 def.)

#### **Phagocytes deficiency :**

- AD (HI) Sec61α1 deficiency (SCN)

#### Mendelian predisposition:

- AR NOS2 deficiency (CMV)
- AR CD16A deficiency (EBV)
- AR DEF6 deficiency (EBV)
- AD (HI) CD48 deficiency (EBV)
- AD (DN) BPIFA1/SPLUNC1 (Neisseria)
- AR IFNG deficiency (vaccin: BCG, MSMD)
- AR TBX21 (Tbet) deficiency (MSMSD)
- AR SNORA31 deficiency (HSE)
- AD (HI) MAPK8 deficiency (CMC

#### **Susceptibility to COVID 19:**

- AD (HI) UNC93B1 deficiency
- AD (HI) IRF7 deficiency
- AD (HI/ DN) INFAR1 deficiency
- AD (HI) IFNAR2 deficiency
- XL TLR7 deficiency

#### Actualités 2020 suite

#### **Autoimmunity:**

- AR TET2 deficiency (ALPS-like)
- AD (HI) SOCS1 deficiency
- AR GIMAP5 deficiency
- AR GIMAP6 deficiency
- AD (HI) PTPN2 deficiency

#### **Autoinflammation:**

- AD (HI) CDC42 deficiency
- AR HEM1/NCKAPL1 deficiency
- XL UBA1 (somatic mutation)
- AR STAT2 GOF
- AR RCH1 deficiency
- AD (DN) RIPK1 deficiency

#### **ESID 2020**



#### **Combined immunodeficiency (CID):**

- XL SASH3 deficiency (ENT, VZV, HSV, HPV, DID, NHL)
- AR ITPR3 deficiency (CID EBV)

#### **B cell deficiency (PAD):**

- AR BOB1 (POUZAF1) deficiency (agamma)

#### **Phagocytes deficiency :**

- AR DBF4 deficiency (SCN)

#### **Dysregulation:**

- AR RHOG deficiency (FHL)
- AD TRAF3 (haploinf) lymphoïde hyperplasia

#### **Autoinflammation:**

- AD NLRP6
- XL TLR8 GOF (mosaic)

### **XL SASH3 deficiency** (XL-CID)



**SASH3 :** SAM and SH3 domain containing 3 May function as a signaling adapter protein in lymphocytes

- 11 patients from 10 kindreds (p.R245X, p.R288X, p.R347C, p.I1227T...)
- CID: ENT, Viral infections: severe VZV (recurrent), HSV (chronic), HPV,
  - Autoimmunity : DID; thyroïde, cytopenia....
    - Eczema; inflammatory, granuloma, Lymphoma NHL
- Lymphopenia
- $\downarrow$  CD4 and CD8 naïfs,  $\downarrow$  B,  $\downarrow$  NK,  $\downarrow$  TTL, abnormal T cell repertory
- **↓** TCR signalling (phospho)

Hemizygous mutation caused Impairement of TCR signalling pathway (prolif  $\downarrow$ ,  $\uparrow$  apoptosis).

• SASH3, also called SLY/SLY1 (SH3-domain containing protein expressed in lymphocytes), is expressed exclusively in lymphocytes and is essential in the full activation of adaptive immunity

### AR ITPR3 deficiency (AR-CID)



the type 3 isoform of the inositol 1, 4, 5-trisphosphate receptor (ITPR3)

- 3 patients CID Infections, Leyomosarcome EBV (1), inflammation, granuloma
- Arg1850, Tyr2033, Arg2524Cys, Phe1618
- ↓ calcic flux, with the exception of one patient with stop gain mutation ↑ calcic flux



ITPR3 (or IP3R3) is a Ca<sup>2+</sup> channel enriched at ER-mitochondria contact sites Realize where it provides Ca<sup>2+</sup> for transport into the mitochondria

### AR BOB1 (POUZAF1) deficiency (AR agamma)







Patient from consanguinous kindred with homozygous null mutation in *POUZAF1* gene:

Infection (lung),

neurologic disaese with  $\checkmark$  myelinisation white matter and  $\uparrow$  cells in CSF

- B cells normal count , but  $\checkmark \checkmark$  B CD27+ cells
- No plasmablasts
- Agammaglobulinemia
- Defect production of specific Abs

BCR impairment and defective class switch

Mice lacking the lymphocyte-specific transcription factor Bob1 fail to generate germinal centers and a robust lg responses.

C. Picard 11/12/2020

19TH BIENNIAL MEETING OF

THE EUROPEAN SOCIETY FOR IMMUNODEFICIENCIES

# AR DBF4 deficiency (AR-SCN)



- 1 patient c.627A>C; p. Lys209Asn homozygote
- Infections, dysmorphia, Neutropenia
- BOM: Arrest at promyelocytes stage
- DDK deficiency

CDC7-DBF4 kinase (DDK) initiates DNA replication in eukaryotes by activating the replicative MCM helicase





# **AR** *RHOG* deficiency , HLH

RhoG regulates gene expression and the actin cytoskeleton in lymphocytes

Patients with HLH manifestations with biallelic mutations in *RHOG* gene that caused abnormalities of cytotoxic vesicles for the membrane docking

RhoG has a role for the Munc13-4 docking at cellular membrane



ESID

*RhoG* (Ras homology Growth-related) (or ARGH) is a small (~21 kDa) monomeric GTP-binding protein (G protein), and is an important component of many intracellular signalling pathways.

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## **XL** TLR8 GOF (mosaic)

- 6 male patients with hyper inflammation, infections, syndrome Lympho-Proliferatif
- Hemizygous mutations in *TLR8* gene: P432L, F494L in 30% of cells = Mosaicism

#### Neutropenia, B cell defect

**↑**CD8 and  $T_{Temra}$ , exhausted cells, ↓ B cells and ↓ ↓ B memory ;

Hypogammaglobulinemia, **↑**pro-inflammatory cytokines



# ESID 2024: Marseille





AD TRAF3 (haploInf) lymphoïde hyperplasia: Hyperplasia tonsilar, tongue

AD NLRP6 p.R653G: skin and liver inflammation with increased IL-1b , IL-18, NFKB activation

SHARPIN c.220dupC; p.L174PfsX86: Polyarthritis, colitis, hepatic glycogenolysis