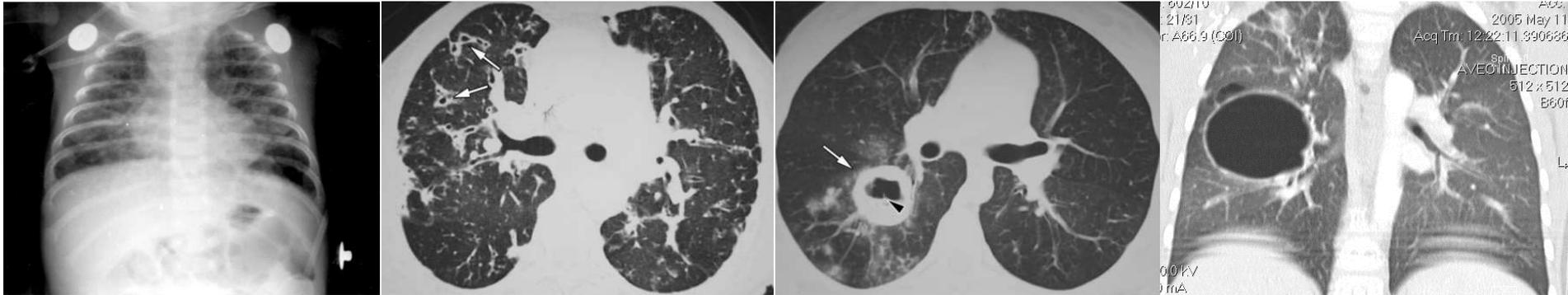


Susceptibilité génétique aux infections



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Classification PIDs

1. **Combined T and B cell immunodeficiencies**
2. **Antibody immunodeficiencies (B cell immunity)**
3. **Phagocytes deficiencies (PN, mono, Macro, DC)**
4. **Opsonisation deficiencies (complement)**
5. **Defects in innate immunity**
6. **Diseases of immune dysregulation**
7. **Auto-inflammatory disorders**

Defects of Cell-Mediated Immunity

clinical signs

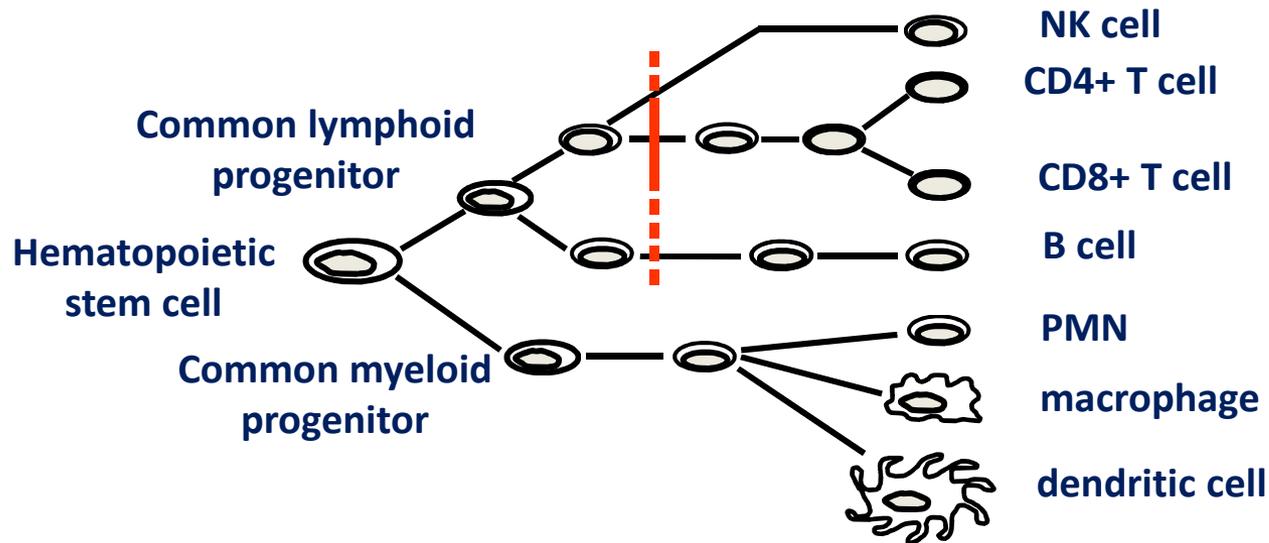
- Recurrent severe infections (bacteria, viruses, fungi, parasite)
 - Often caused by opportunistic pathogens (*P.jirovici*, *BCGitis...*)
- Failure to thrive – particularly prominent
- Chronic diarrhea
- Graft versus Host Disease (materno-foetal or after blood transfusion).



Combined immunodeficiency

T lymphopenia cell deficiency with or without B and/or NK cells

- If T cells < 500 cells/mm³ : Severe combined immunodeficiency
- Usually, diagnostic during childhood



CID and genetic defects

Up 60 (25 for SCID) genes to be studied (expanding year after year):

1. Enzymatic disorders: *ADA, AK2, PNP*
2. Cytokine signaling pathway defects: *IL2RG, JAK3, IL7RA, IL2RA,*
3. VDJ disorders: *RAG1, RAG2, DCLRE1C, DNAPKC, DNA ligase IV, NHEJ1*
4. TCR signaling pathway disorders: *CD45, CD3D, CD3E, CD3Z, CD3G, CD8A, ZAP70, LCK, ORAI1, STIM1, MAGT1, TRAC, UNC119*
5. Thymus differentiation defects: *CORO1A, TBX1, FOXP1*
6. Switch defects: *TNFSF5, TNFRSF5*
7. Class II and I deficiencies: *C2TA, RFXANK, RFX5, RFXAP, TAP1, TAP2, TAPBP*
8. Others: *STAT5B, ITK, CD27, DOCK8, MST1/STK4, WASP, FOXP3, IKBA, RHOH, CARD11*
9. DNA repair defects: *ATM, MRE11, NBS1, BLM, DNMT3B, ZBTB24, PMS2, RNF168, RMRP, SMARCA1*
10. Dyskeratosis: *DKC1, TERC, TERT, NOP10, NHP2, TIN2, C16orf57, TCAB1*

Humoral Defects

- X-linked agammaglobulinemia (XLA) = Bruton
- AR agammaglobulinemia (*μ heavy chain, I5, Iga, Igb, BLNK*)
- Switch deficiency syndromes (AID and UNG)
- Common variable immunodeficiency (CVID)
- Selective IgG subclass deficiency
- Specific polysaccharide antibody deficiency (SPAD)

Clinical signs

Bacterial infections (lung and ENT).

Bronchiectasis, sinusitis

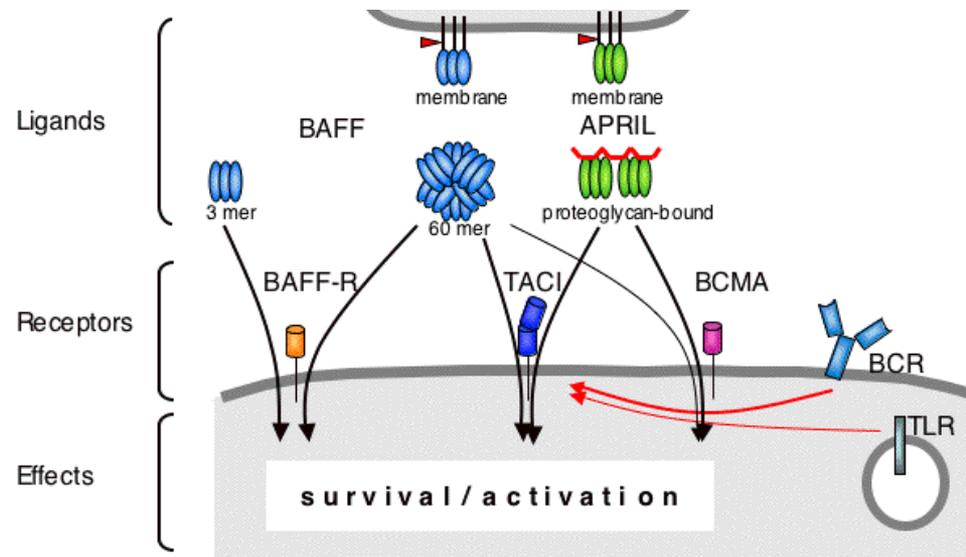
Chronic Gastroenteritis

+/-Autoimmune Disease

Common Variable Immunodeficiency (CVID)

- Frequency 1/30-50 000, M = F, 15 to 30 years, family forms: 20%
- Heterogeneous syndrome with lack of Ig production :
 - Failure to produce Ig: IgG + another IgA or M)
 - Presence of B cells > 1%
 - Exclusion of other known genetic defects: XLA (Bruton), Thymoma (Good)
Hyper-IgM (CD40L, CD40, AID, UNG)
XLP (Purtilo)

- Genetic etiologies: rare, ICOS, TACI, BAFF-R, CD19, CD20, CD81



Common Variable Immunodeficiency (DICV)

- **Infections:**
 - ENT and lung (bronchiectasia) (*H. influenzae*, *S. pneumoniae*, *S. aureus*)
 - Sepsis (*Pseudomonas*, *S. pneumoniae*, *H. influenzae*)
 - Chronic diarrhea (*Giardia lamblia*, Salmonelles, *Campylobacter*)
- **Proliferation**
 - lymphoid proliferations
 - granulomatosis
 - cancer
- **Autoimmunity**
 - cytopenia
 - rheumatoid

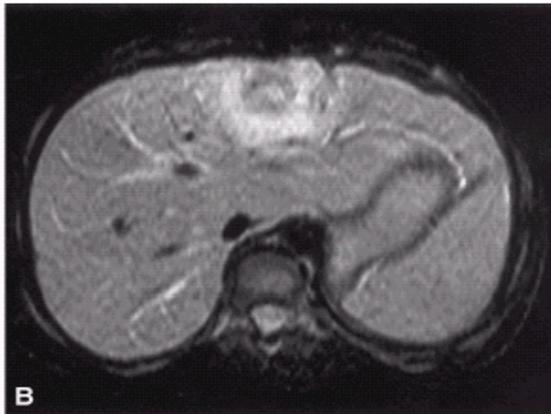
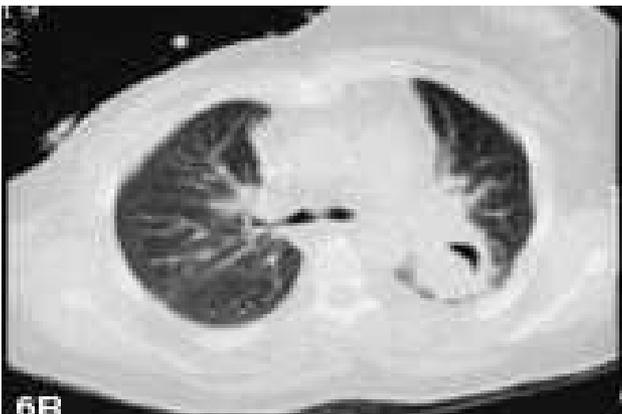


DICV et manifestations cliniques

(Chapel et al. Blood 2008, n= 389 ESID)

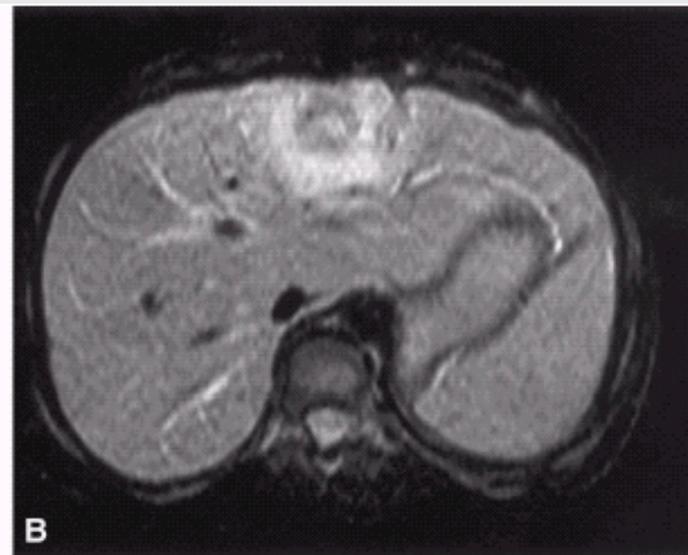
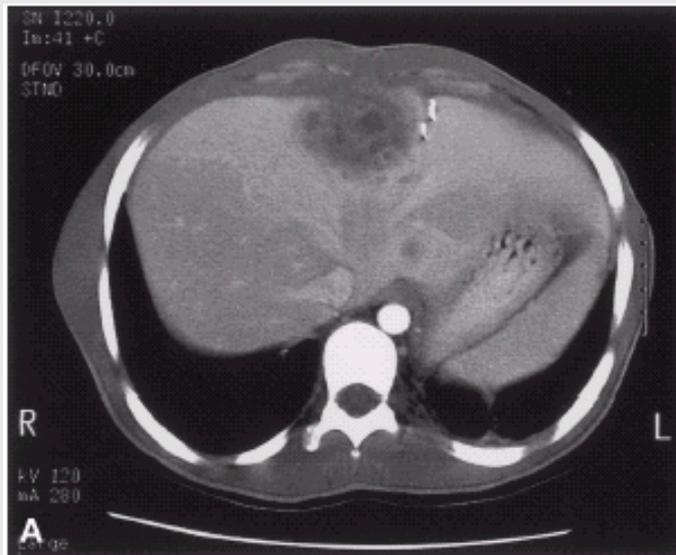
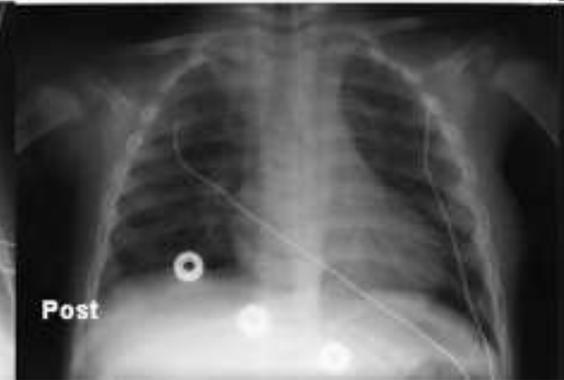
Chronic granulomatous disease (CGD)

- Inherited disorder of phagocytic cells
- Frequency: 1/220 000
- Recurrent tissular bacterial (*S.aureus*, enterobacteria..) and fungal (Aspergillosis+++, *C. albicans*+) infections
- Associated Symptoms
 - Excessive Granulomata – often in GI tract (IBD like diseases)
 - Chorioretinitis
- Treatment (Antibacterial and antifungal prophylaxis) +/- HSCT

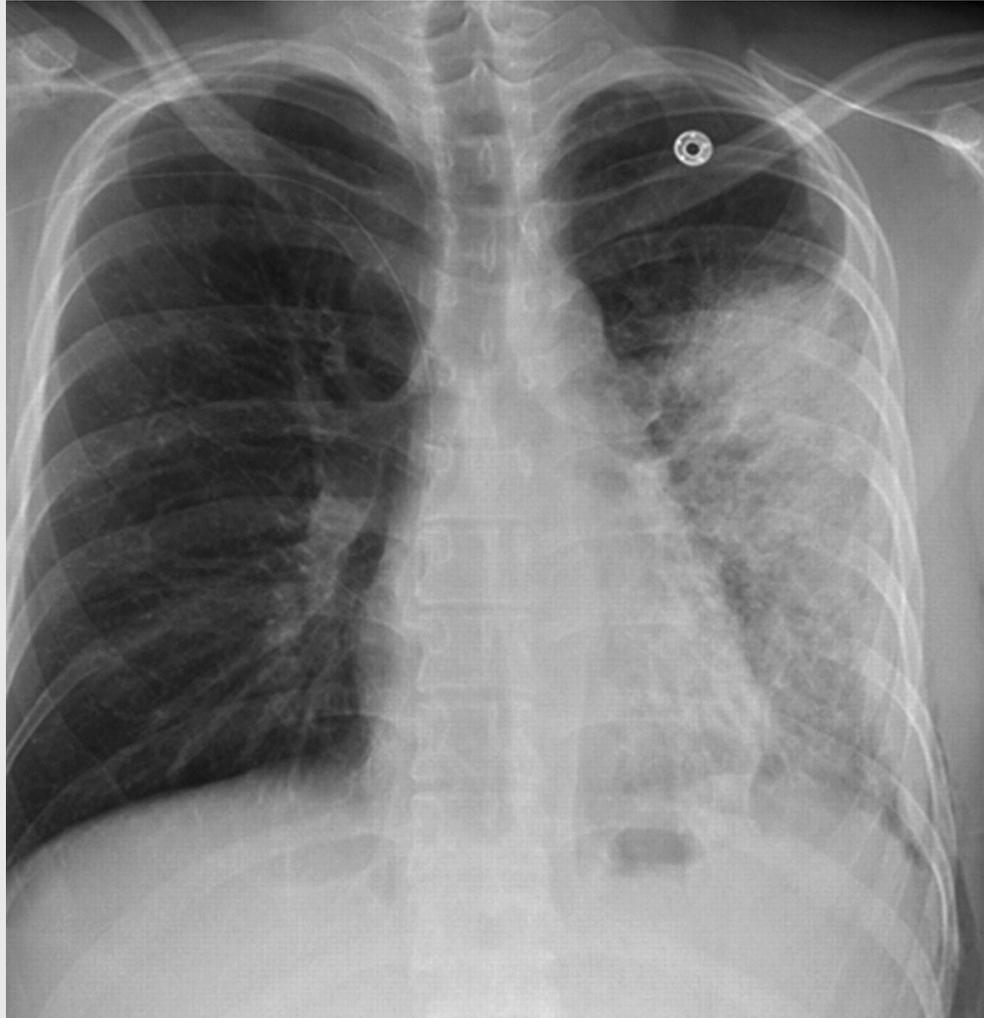


Chronic granulomatous disease (CGD)

- **Pneumonia: *Aspergillus* (41%), bacteria**
- **Abscess:**
 - Subcutaneous: *Staphylococcus spp* (27%)
 - Hepatic: *Staphylococcus spp* (50%)
 - Lung: *Aspergillus* (23%)
 - Brain: *Aspergillus* (58%)
- **Adenitis: *Staphylococcus spp* (26%), *Salmonella*, *Actinomyces*, BCG**
- **Osteomyelitis: *Serratia* (29%), *Aspergillus* (22%)**
- **Septicaemia: *Salmonella* (18%), *Burkholderia* (12%), *Candida* (11%)**
- **Meningitis: *Candida* (20%)**
- ***Aspergillus* is responsible for 1/3 of death**



—22-year-old man with history of chronic granulomatous disease presenting with fever and productive cough for 10 days.

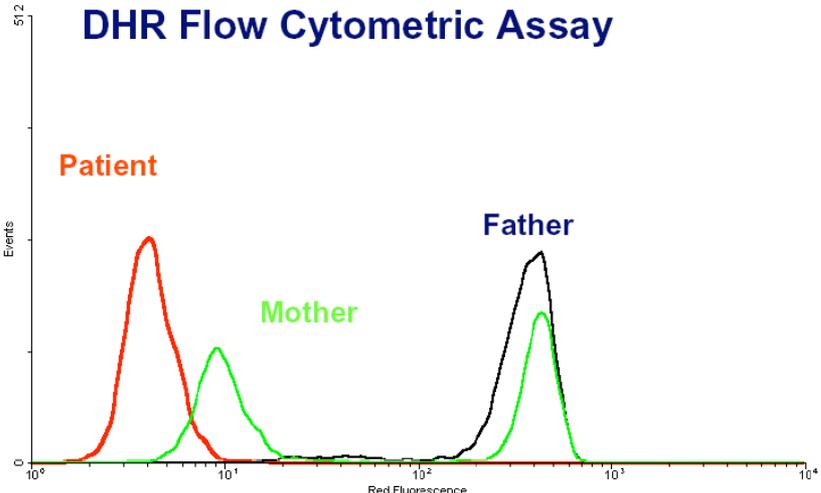


Godoy M C B et al. AJR 2008;191:1570-1575



5 genes: XR (*CYBB*, gp91phox) = 67%
AR (*NCF1*, p47phox) = 33%
AR (*CYBA*, p22phox) = 5%
AR (*NCF2*, p67phox) = 5%
AR (*NCF4*, p40phox) = 1 pt

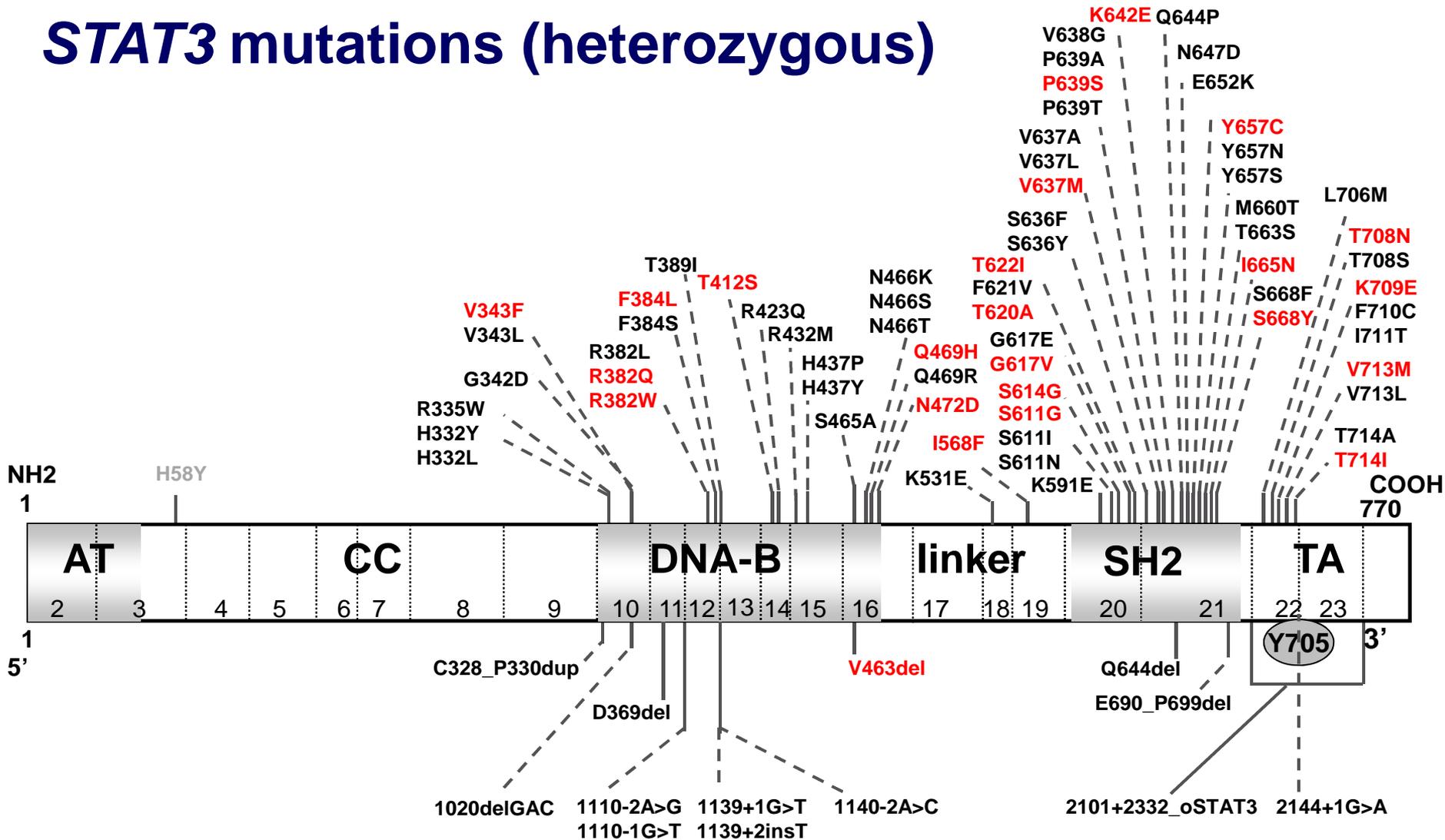
Nitroblue tetrazolium reduction (NBT)



HyperIgE syndromes - introduction

- **HyperIgE (>2000IU/ml in adult) and recurrent infections**
- **Genetic heterogeneity:**
 - **Autosomal dominant form**
 - **AD-HIES or Job** (Davis et al Lancet 1966) or **Buckley syndrome** (Buckley et al. 1972 Pediatrics)
 - **Autosomal recessive forms**
 - **TyK2 genetic defect (one patient)** (Minegishi et al. 2006, Immunity)
 - **AR-HIES + T cellular immunodeficiency** (Renner et al. 2004, J Ped)
 - ⇒ **DOCK8 deficiency** (Zhang et al. 2009, NEJM)

STAT3 mutations (heterozygous)



n=60 French patients, 24 mutations, 13 news

55% mutation located in the DNA binding domain (DBD),

32% in the SH2 domain,

2% in the linker domain and 10% in the TA domain

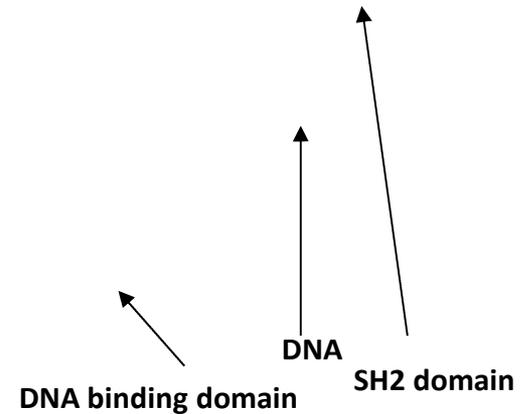
(Chandesris et al. 2012 Medicine)

STAT3 (transcriptional factor)

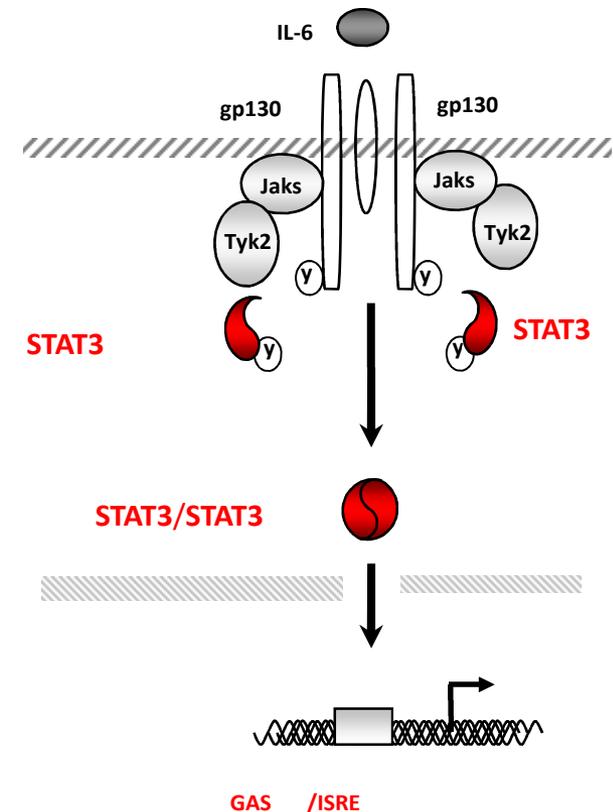
- **STAT3 is involved in responses to multiple cytokines:**

- IL-6 receptor superfamily (gp-130) :
IL-6, IL-11, IL-27, IL-31,
OSM, CNTF, LIF and CT-1
- IL-10
- IL-2 family cytokines: IL-2, 7, 9, 15 and 21

- **STAT3 nuclear translocation**
Acute-phase protein genes (APP genes)



(Becker et al. 1998, Nature)



STAT3 deficiency-clinical phenotype



Dysmorphia variable

(B.Grimbacher, NEJM 1999)

Hyperextensibility

Retained primary teeth



Fractures
+ Osteoporosis

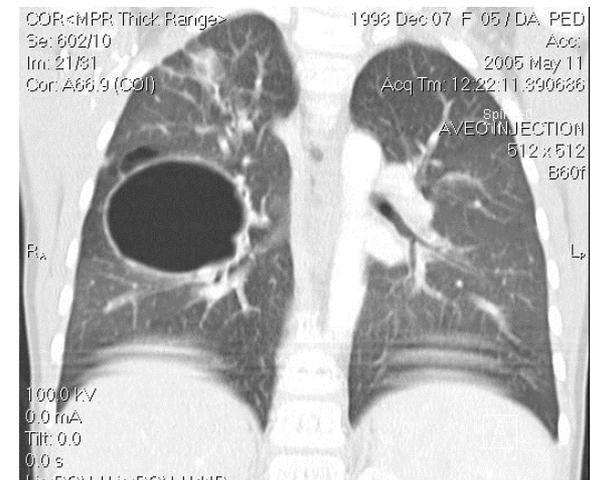
STAT3 deficiency-clinical phenotype

eosinophilic spongiotic dermatitis

Candidiasis (80%)

recurrent cold skin abscess

(C. Bodemer)



**Recurrent bacterial
Pneumonia and II cyst
Formation (**Asperillosis**)**

Bacterial pneumonia 92% of patients

Mean age of first pneumonia = 39.5 months (range 0 to 192 months)

3 episodes per patient (range 1 to 30 episodes)

1 SDRA woman 20 yo

1 child of 14 months died (*S.pneumoniae*)

Acute pneumonia Germs :

***S. pneumoniae* 30%**

***S. aureus* 29%**

***Haemophilus influenzae* 13%,**

Lung-complications

Sequellae secondary to previous bacterial

infections = 68% of patients:

mean = 21 years (range 1 to 45 years)

- **Bronchiectasis in 66% of patients**
- **Pneumatoceles in 53% of patients**
- **Broncho-pleural fistulas, Hemoptysis, pneumothorax, Tracheal stenosis .**

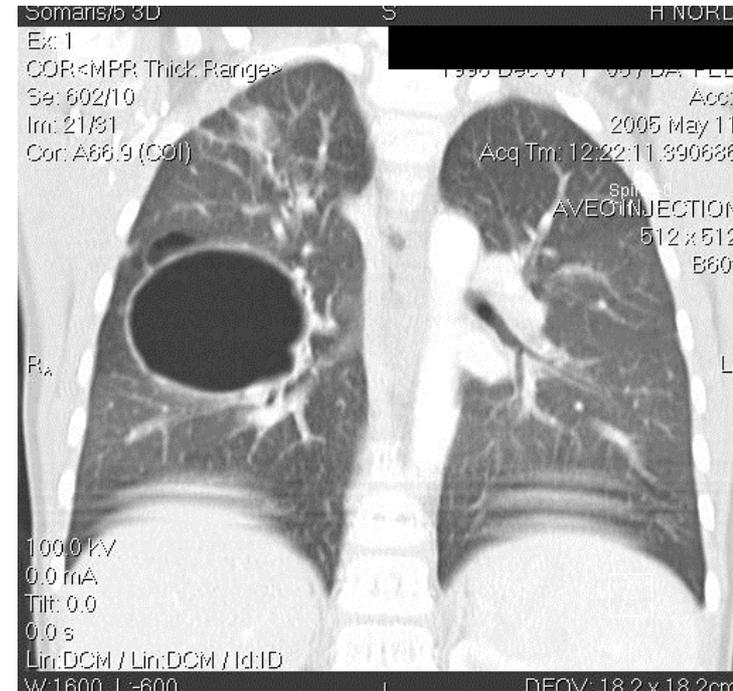
Lobectomies (range 1 to 3) in 22% of patients

median age of 10 years (range 1 to 17 years)

***Aspergillus spp.*-related diseases were observed in**

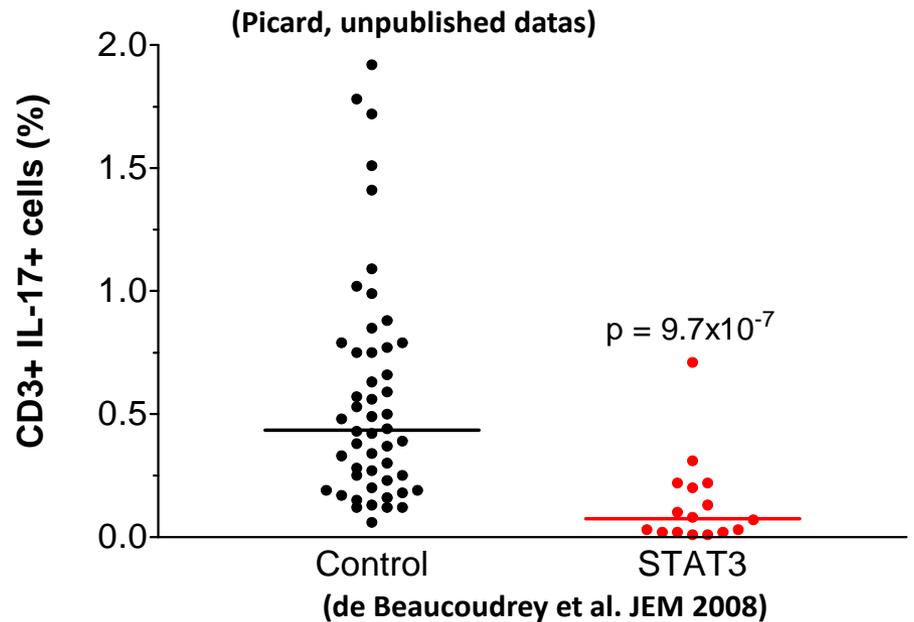
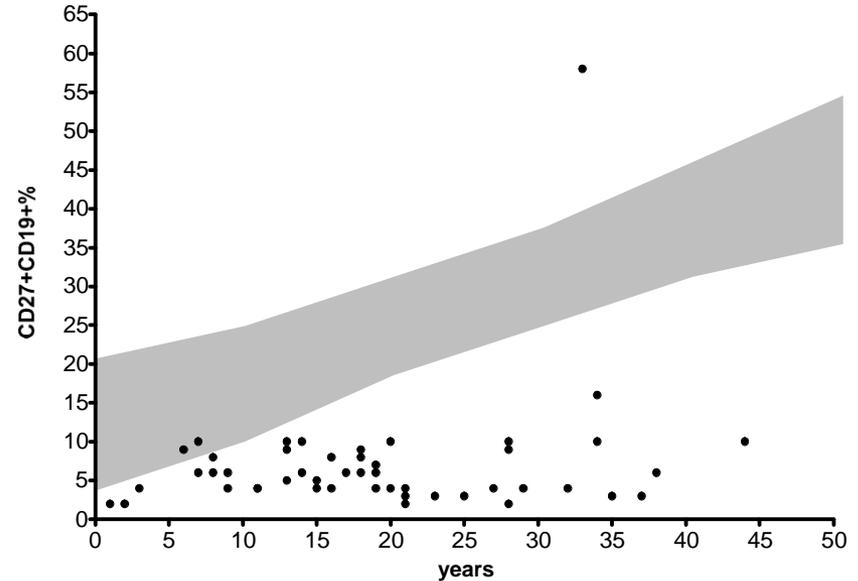
22% patients : Aspergilloma : 4 patients

12.7 years (range 10 to 16 years)



STAT3 deficiency-Immunology

- Normal phagocytic explorations
- PEO > 600 cell/ μ l : 75% patients
- B immunity :
 - B memory cell lymphopenia
 - IgE : Increased in all patients
 - Normal or \uparrow of IgG, A and M
- T immunity:
 - Th17 lymphopenia
 - TTL : normal



Summary immunology-STAT3+/-deficiency

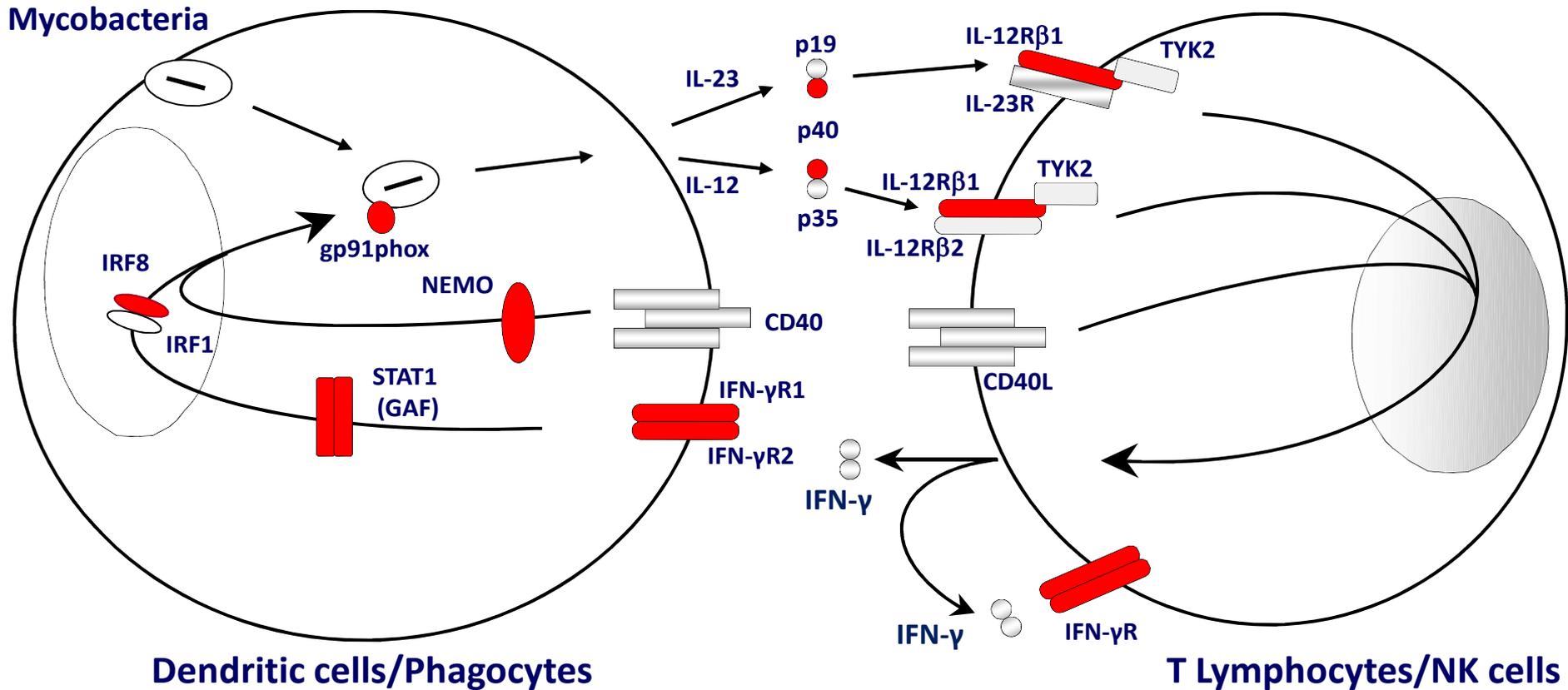
- **Bacterial susceptibility of STAT3+/- deficient patients:**
 - B cell memory deficiency (impairment of IL-21 signalling pathway)
 - Defect in differentiation of naive B cells in plasma cells (IL-21 sig. pathway)
 - Defect in inflammatory response (impairment of IL-6 signalling pathway)
 - TH17 lymphopenia
- **Skin & mucosal immunity to *C. albicans* is controlled by IL-17-producing T cells and cytokines IL-17A, IL-17F, IL-22.**
- **Pneumocyst formation ⇒ Role of IL-6-dependent gp130/STAT3 pathway in repair of the brochiolar epithelium**

(mice specific deletion of Stat3 in lung epithelial cells, Kida et al. Am J Physiol Lung Cell Mol Physio 2005)

Conclusions

- **Infections are the main clinical signs of primary immunodeficiency**
- **Several PIDs developed pneumonia :**
 - **Interstitial pneumonia → CID**
 - **Recurrent bacterial infection → B cell deficiency**
 - **Fungal or bacterial abscess → CGD**
 - **lung complications such as bronchiectasia or pneumatoceles**

Mendelian susceptibility to mycobacterial disease: 8 genes so far



First genetic etiology discovered in 1996: Newport et al. *N Engl J Med* 1996.

Jouanguy E, et al. *N Engl J Med* 1996.

Courtesy of S.Boisson-Dupuis

Inherited disorders of the IL-12/23-IFN γ loop

Gene	Cellular exp.	Deficiency	Inheritance	Frequency
<i>IFNGR1</i>	-	Complete	AR	~30%
	+	Complete	AR	
	+	Partial	AR	
	+++	Partial	AD	
<i>IFNGR2</i>	-	Complete	AR	< 5%
	+	Complete	AR	
	+	Partial	AR	
<i>STAT1</i>	-	Complete	AR	< 5%
	+	Partial	AD	
	+	Partial	ADomissive	
<i>IL12RB1</i>	-	Complete	AR	~40%
	+	Complete	AR	
<i>IL12B</i>	+	Complete	AR	~14%
<i>NEMO</i>	+	Partial	XR	<3%
<i>CYBB</i>	+	Partial	XR	< 2%
<i>IRF8</i>	-	Complete	AR	< 2%
	+	Partial	AD	

Inherited disorders of the IL-12/23-IFN γ loop: IFN- γ R1 deficiency

IFN- γ R1 deficiency	AR Complete deficiency (n=22)	AD Partial deficiency (n=38)
IFN- γ signalization	none	residual
Asymptomatic pts	no	yes, 8%
BCG infection	100% of vaccinated	73% of vaccinated
EM infection	77%	82% (mostly, <i>M. avium</i>)
Salmonella infection	14%	5%
Infection localization	Mostly disseminated	Bones (80%)
Mortality	64%	5%

(Dorman et al. Lancet 2004)

Inherited disorders of the IL-12/23-IFN γ loop: IL-12R β 1 and IL-12p40 deficiencies

	AR IL-12R β 1 deficiency (n=141)	AR IL-12p40 deficiency (n=49)
Cellular phenotype	No response to IL-12	No IL-12 production
Asymptomatic pts	Yes, 6%	Yes, 10%
BCG infection	78% of vaccinated	97.5% of vaccinated
EM infection	16%	4%
Salmonella infection	40%	22%
Infection localization	Lymph nodes +/- skin (vasculitis)	Lymph nodes
Mortality	29.7%	28.6%

(de Beaucoudrey et al. Medicine 2010)

(Prando et al. Medicine in press)

Mendelian susceptibility to mycobacterial diseases

- ✓ Importance of IL-12/23-IFN γ signaling pathway in anti-mycobacterial immunity
- ✓ Correlation signaling by IFN γ and disease severity

AR IL-12R β 1 deficiency is also a genetic etiology of severe TB