



**Instituto de Bioquímica Médica  
(Laboratório de Imunologia Tumoral  
e Laboratório de Biologia molecular  
e Instituto de Ciências  
Biomédicas/UFRJ)**



**Centro de Transplante de Medula Óssea  
Hospital de Clínicas de Curitiba/UFPR**

# **Fanconi Anemia: immune deficiency and susceptibility to cancer**



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**This work was supported by CNPq and FAPERJ,  
G.A.B. was supported by CAPES, Brazil**



# Fanconi Anemia



**Guido Fanconi, MD - 1927**



Cytogenetic test supports the clinical diagnosis

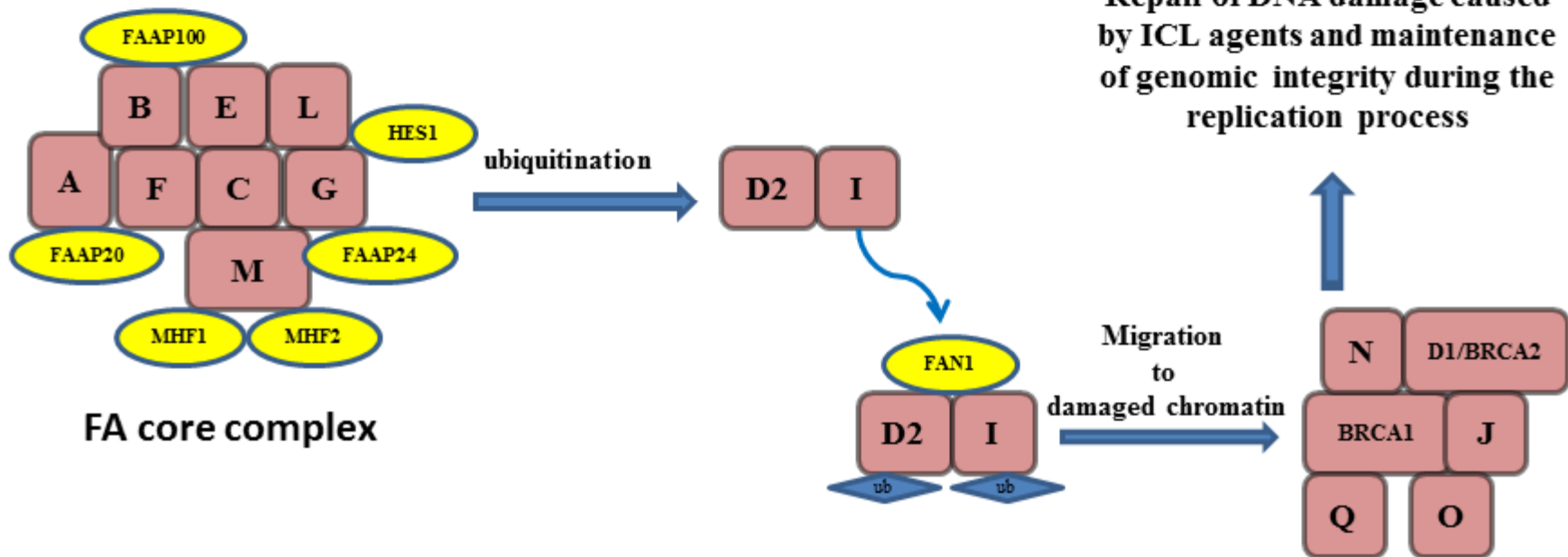


**Table 1. Summary of the Fanconi Anemia pathway, genes, proteins, and functions.**

Complementation Group	Gene	Chromosome location	Protein weight (kD)	Motifs	Required for D2 monoubiquitination
A	<i>FANCA</i>	16q24.3	163	NLS, NES	yes
B	<i>FANCB</i>	Xp22.31	95	NLS	yes
C	<i>FANCC</i>	9q22.3	63	none	yes
D1	<i>FANCD1/BRCA2</i>	13q12.13	380	BRC repeats	no
	<i>FANCD2</i>	3p25.3	155/162	none	yes
E	<i>FANCE</i>	6p21.22	60	NLS	yes
F	<i>FANCF</i>	11p15	42	none	yes
G	<i>FANCG/XRCC9</i>	9p13	68	TPRs	yes
I	<i>FANCI/KIAA1794</i>	15q25-26	146	none	yes
J	<i>FANCI/BRIP1/BACH1</i>	17q22-24		ATPase/helicase	no
L	<i>FANCL/PHF9</i>	2p16.1	43	E3 ligase	yes
M	<i>FANCM</i>	14q21.3	250	ATPase, DNA translocase	
N	<i>FANCN/PALB2</i>	16p12	130	WD40	no
O	<i>FANCO/RAD51C</i>	17q23	37	RAD51 paralog	no
P	<i>FANCP/SLX4</i>	16p13.3	268	endonuclease	no
Q	<i>FANCP/XPF4/ERCC4</i>	16p13.12	100	endonuclease	no



# The FA-BRCA pathway



# Multifunctionality of the FA proteins

## FA DNA repair pathway

FANCA  
FANCB  
FANCC  
FANCE  
FANCF  
FANCG/XRCC9  
FANCL/POG  
FANCM

FA Core complex.  
Monoubiquitinates  
FANCD2 and FANCI  
in response to DNA  
damage

FANCD2  
FANCI

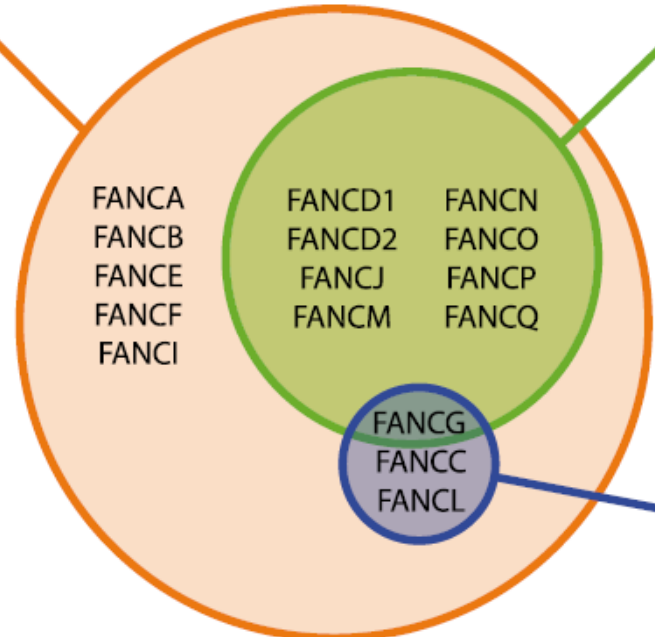
FA Core complex substrates.  
Monoubiquitinated FANCD2  
is required for incisions,  
TLS and HR

FANCP/SLX4  
FANCO/XPF

Nuclease complex.  
Required for incision(s) at the  
site of crosslinked DNA

FANCD1/BRCA2  
FANCI/BRIP1  
FANCI/PALB2  
FANCI/RAD51C

Homologous recombination.  
BRCA/RAD51-mediated  
repair.



## Other functions in DNA repair

FANCG/XRCC9  
FANCD2  
FANCD1/BRCA2

Homologous recombination.  
They form a complex with  
XRCC3.

FANCM

Interacts with BLM

FANCP/SLX4

Interacts with MUS81/EME1 and  
SLX1. Holliday junction resolvase  
Telomere maintenance

FANCO/XPF

NER and some HR transactions.

FANCD1/BRCA2  
FANCI/BRIP1  
FANCI/PALB2  
FANCI/RAD51C

DBS repair.

FANCI/BRIP1

G quadruplet resolution.

## Other functions

FANCC

Modulates PKR, STAT1, Hsp70

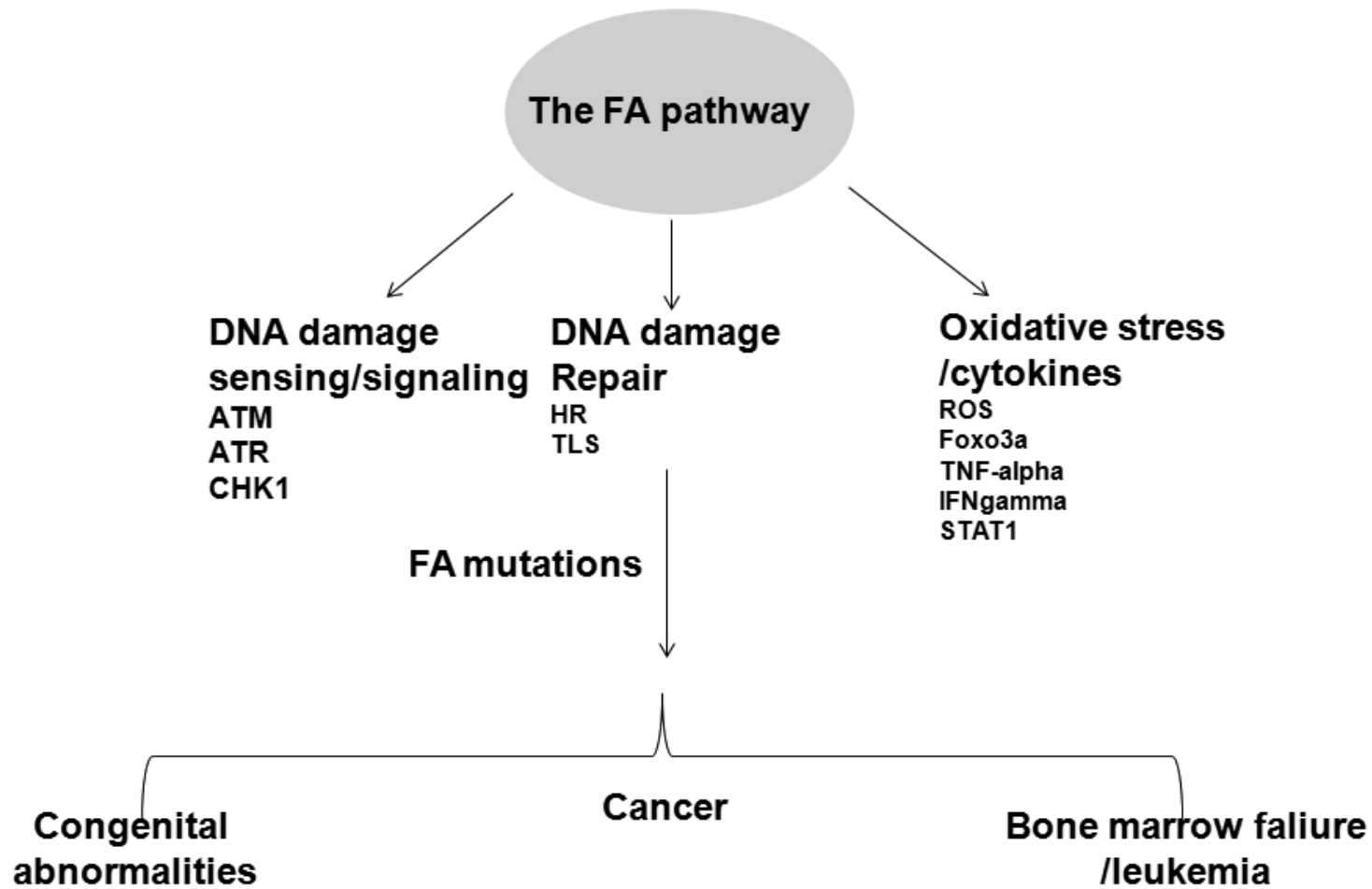
FANCC  
FANCG

Interact with REDOX factors

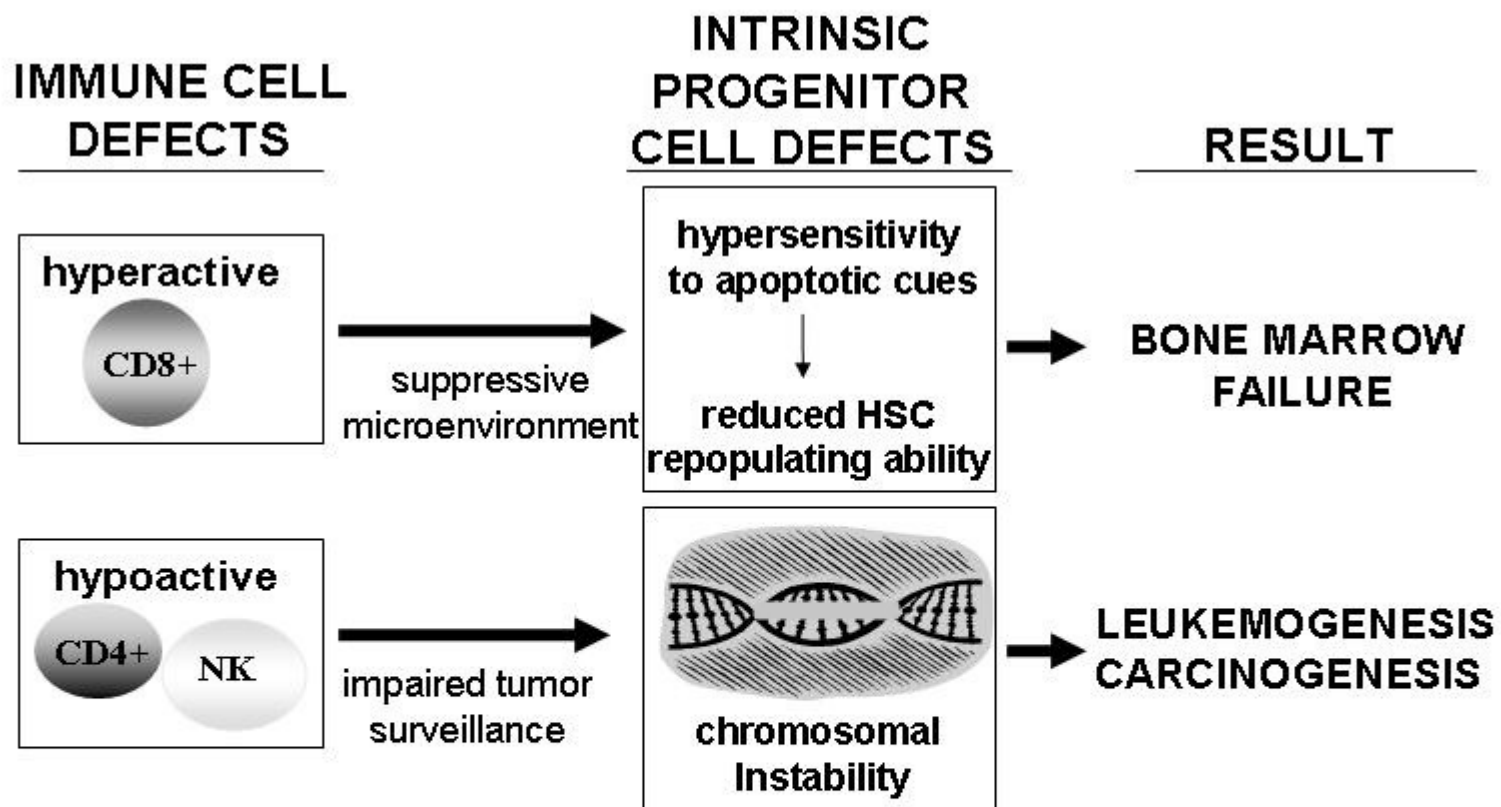
FANCL

Polyubiquitinates  $\beta$ -catenin





# Cancer and Immunology in Fanconi anemia







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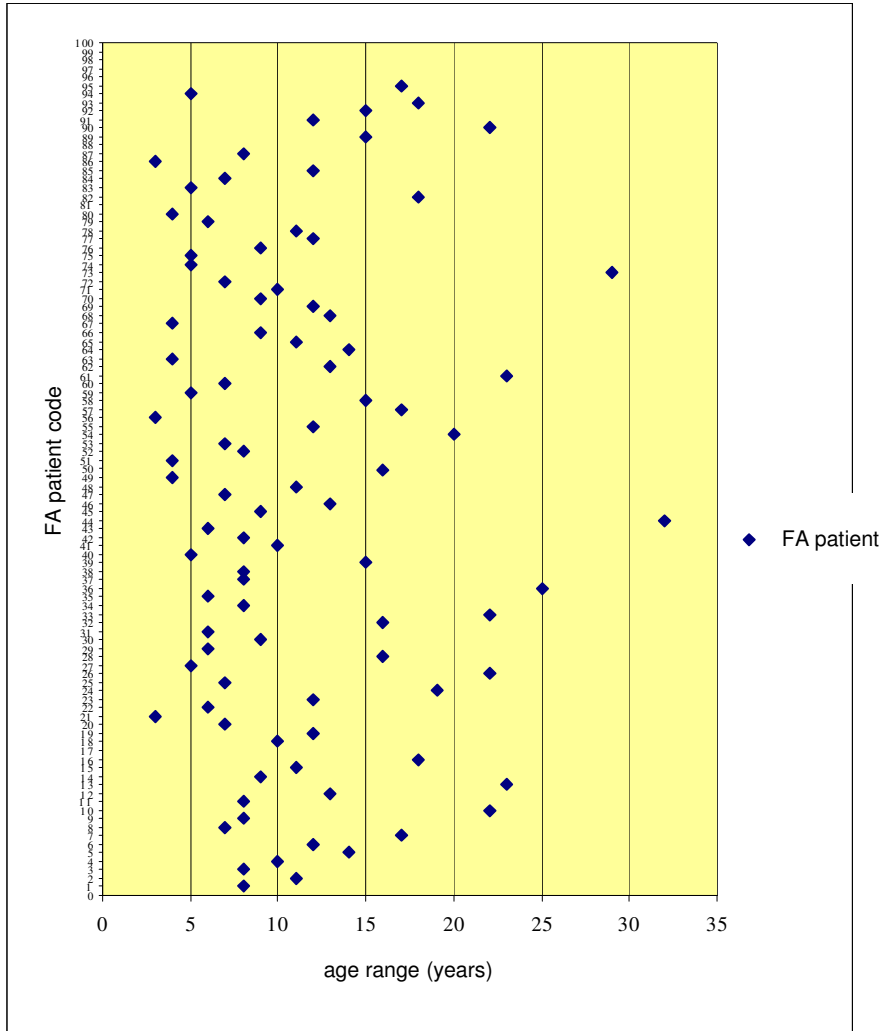
**E-mail: [magrajusto@hotmail.com](mailto:magrajusto@hotmail.com)**

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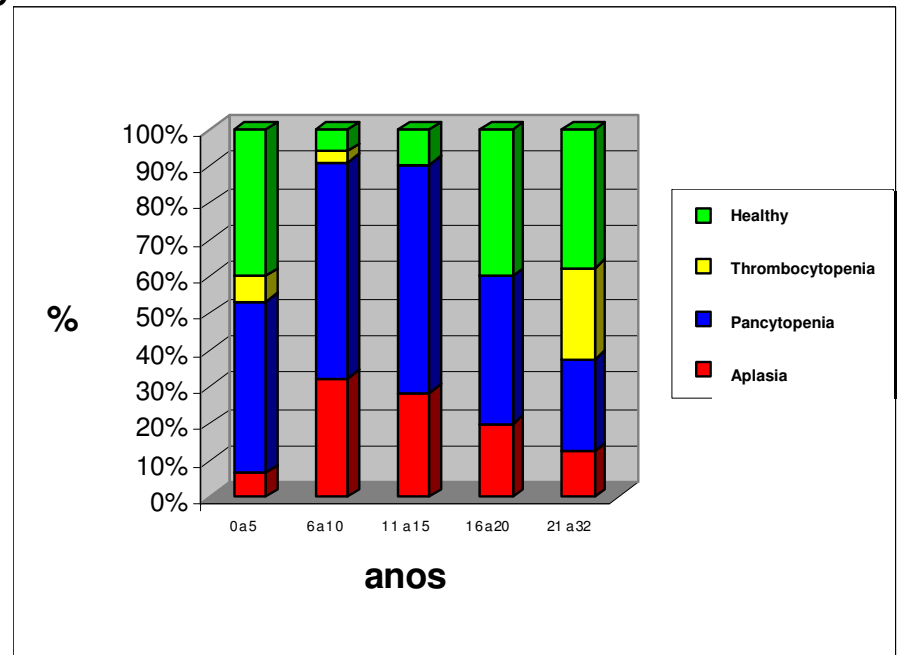


# Chart of FA patients by age and Hematological clinical data

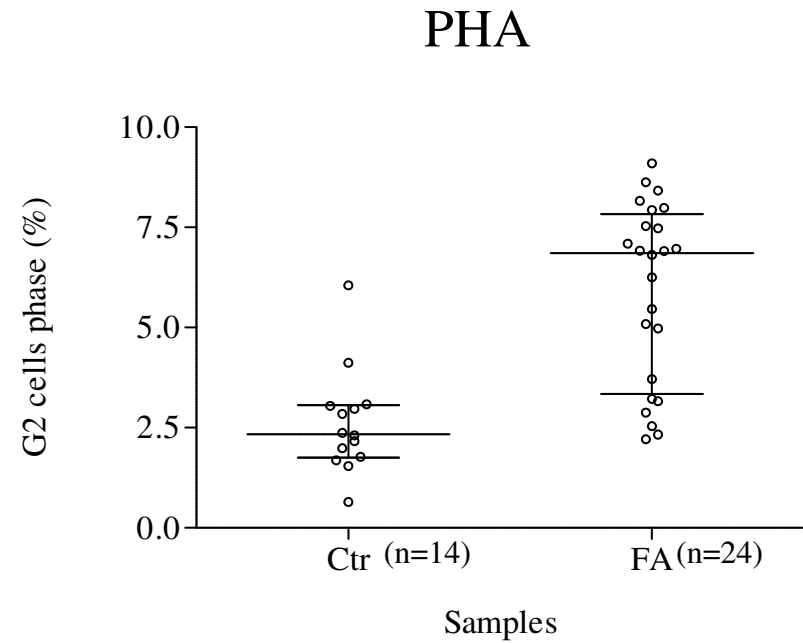
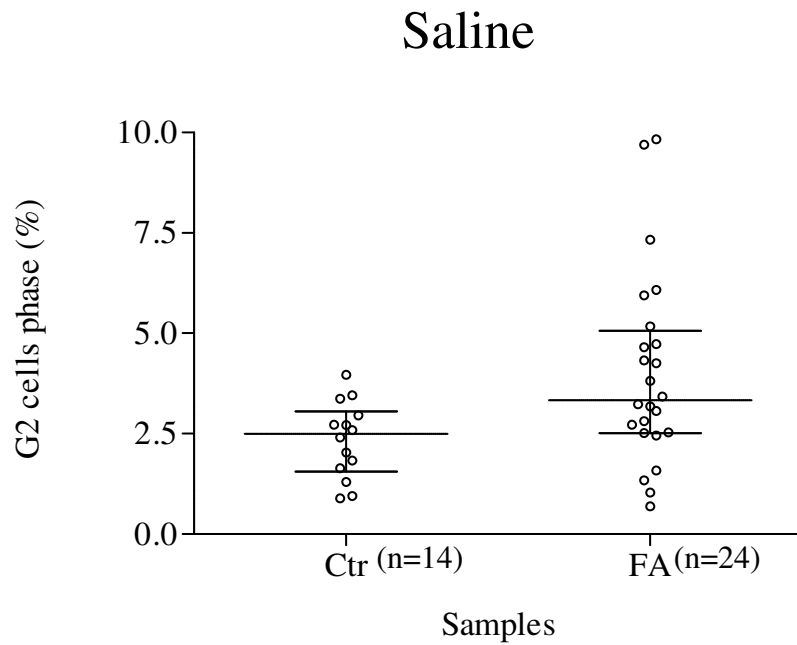
A



B

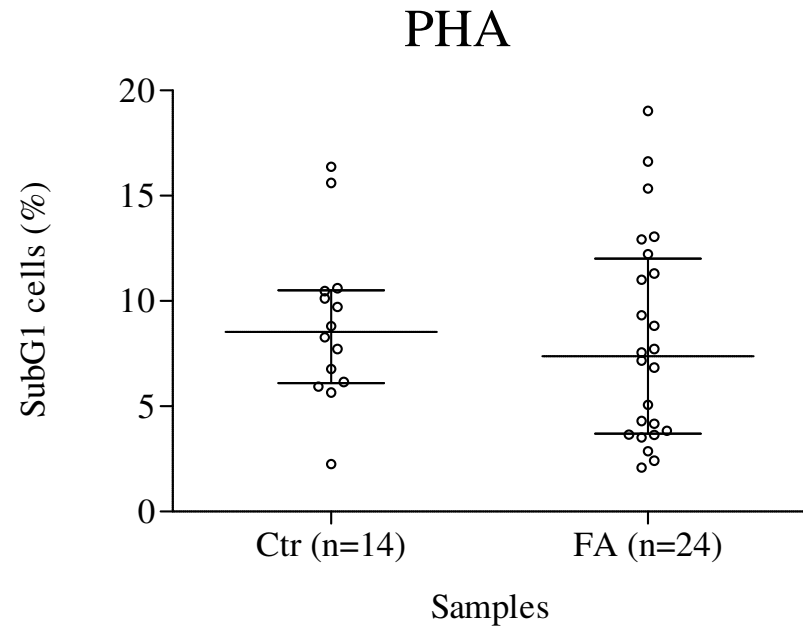
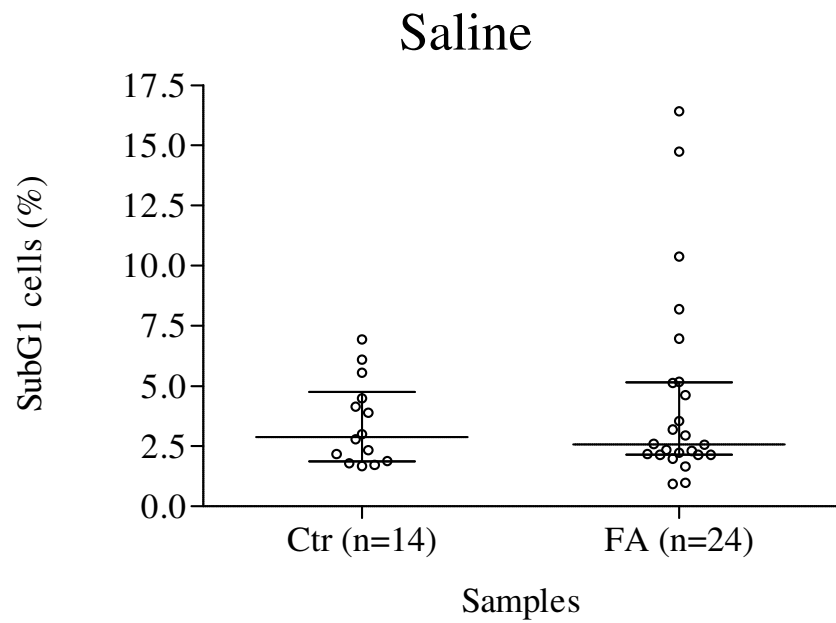


# FA cells in the G2 cell cycle phase



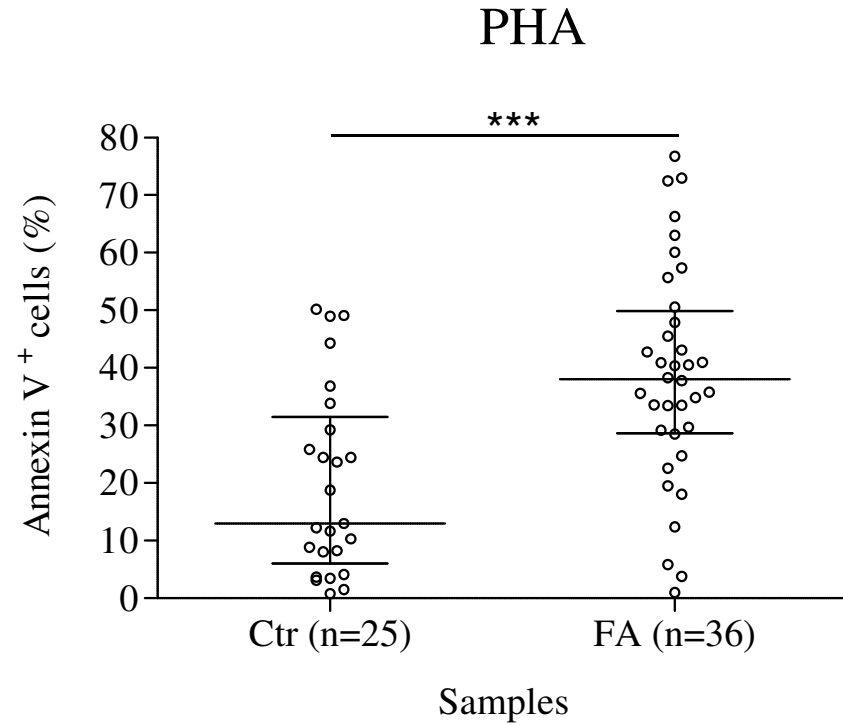
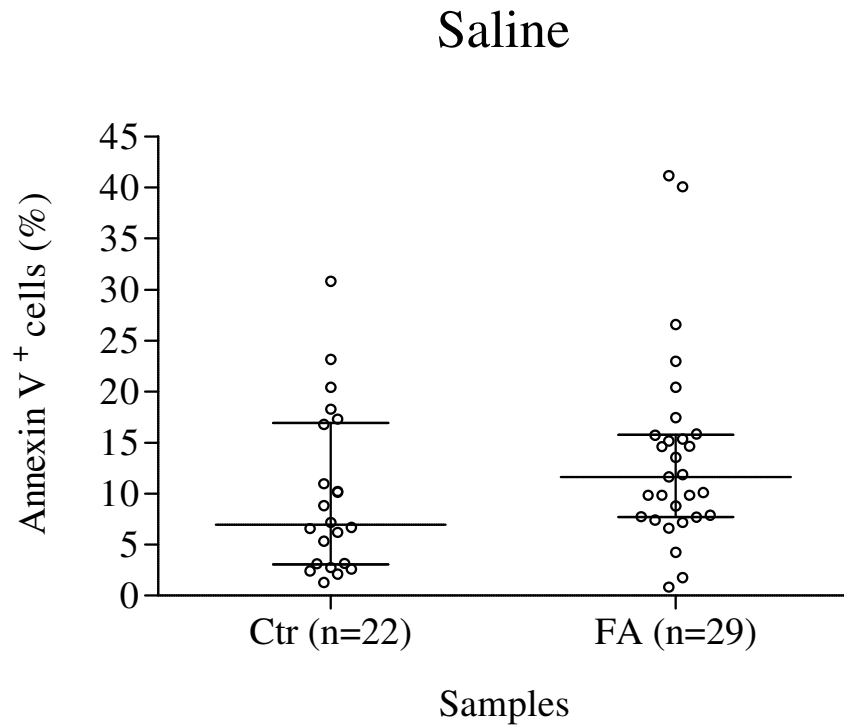
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# Percentage of lymphocytes on Sub-G1 cell cycle phase

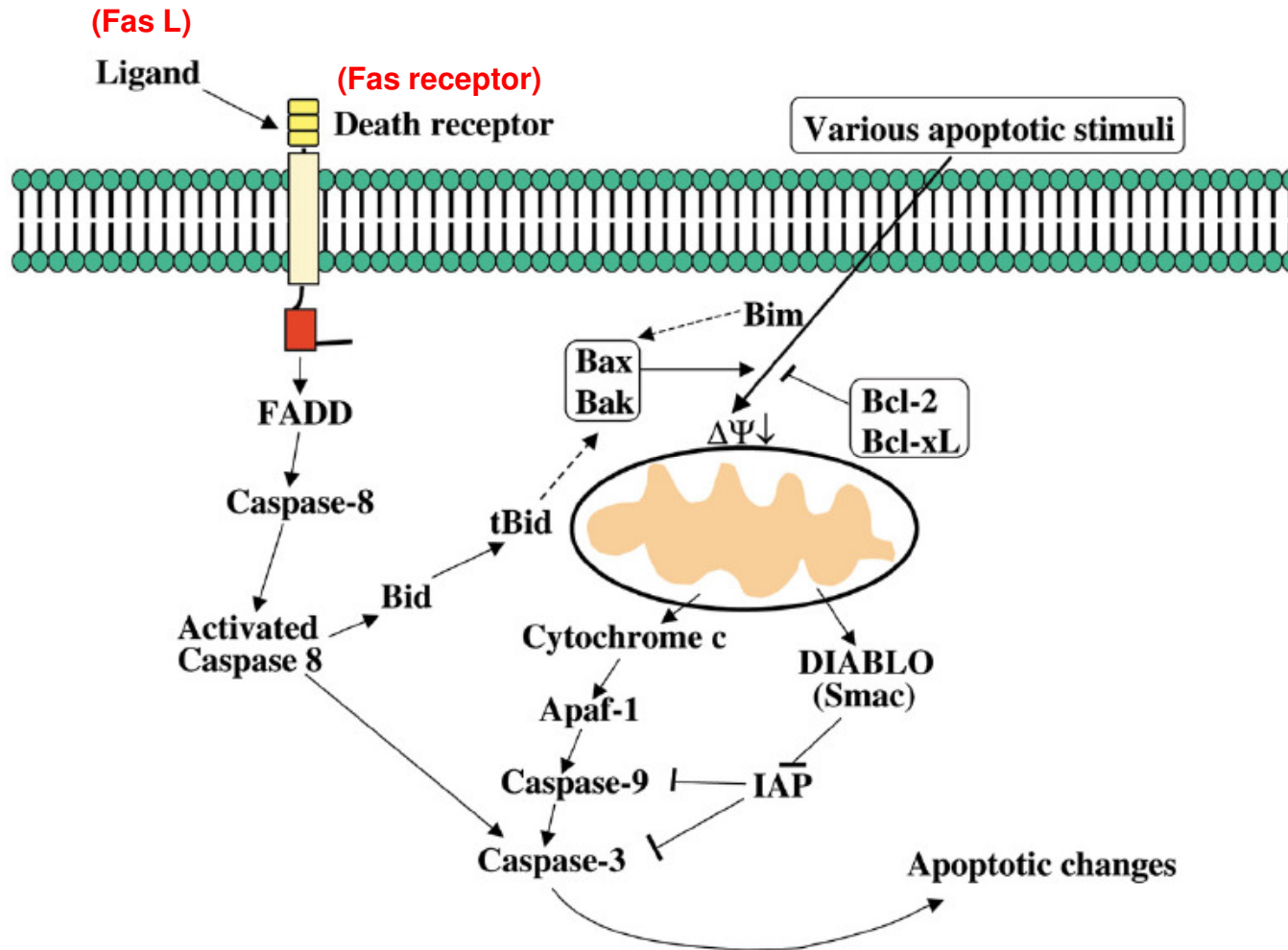


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## Percentage of cells on spontaneous and PHA activated induced apoptosis



# Apoptosis in peripheral lymphocytes



## Increased Fas expression and relation to increased apoptosis

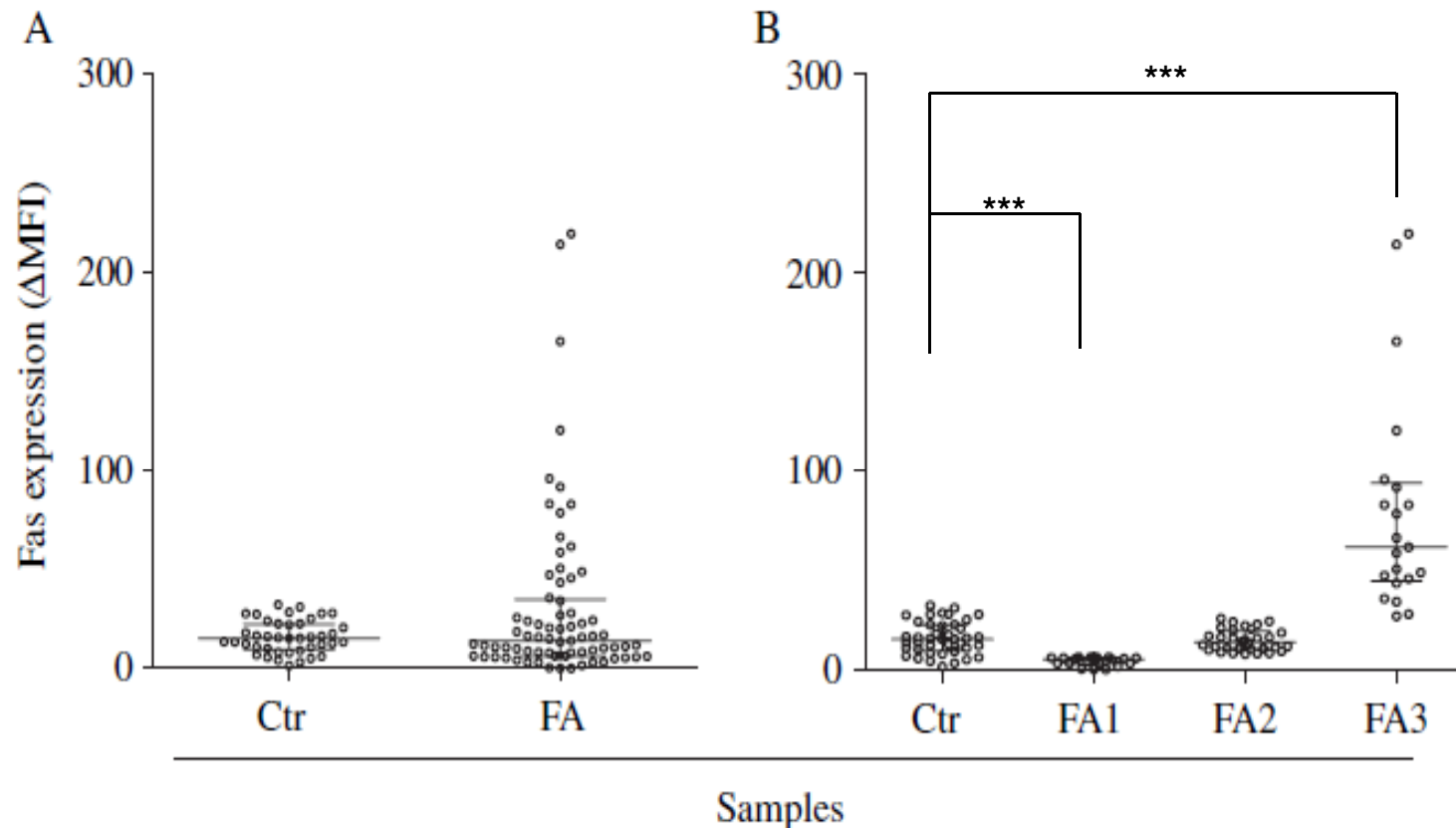


Fig. 1. Expression of Fas receptor in PBMC from FA patients and control samples.



## Apoptosis on the Fas subgroups

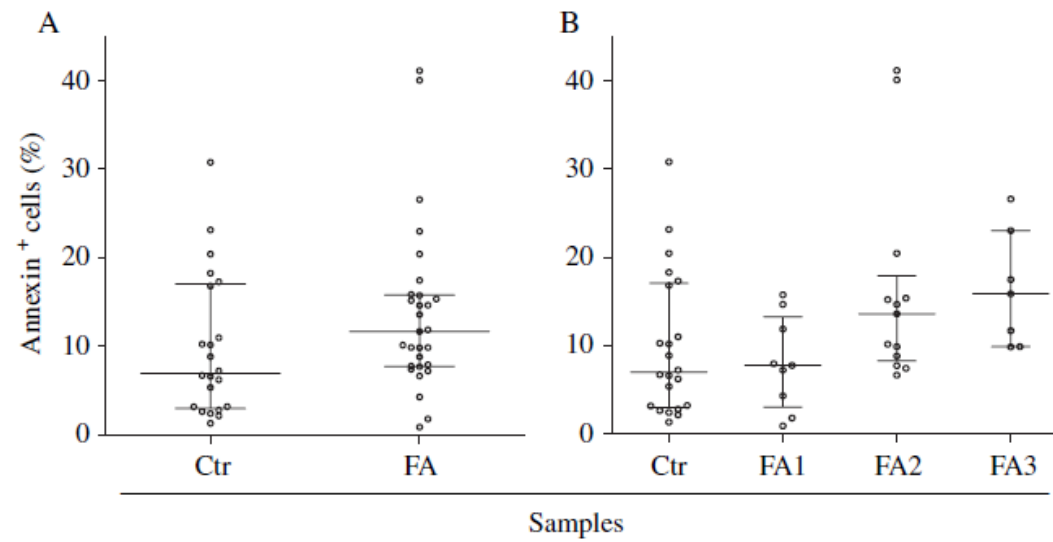


Fig. 4. Spontaneous apoptosis in 24 h PBMC cultures from FA patients and control samples.

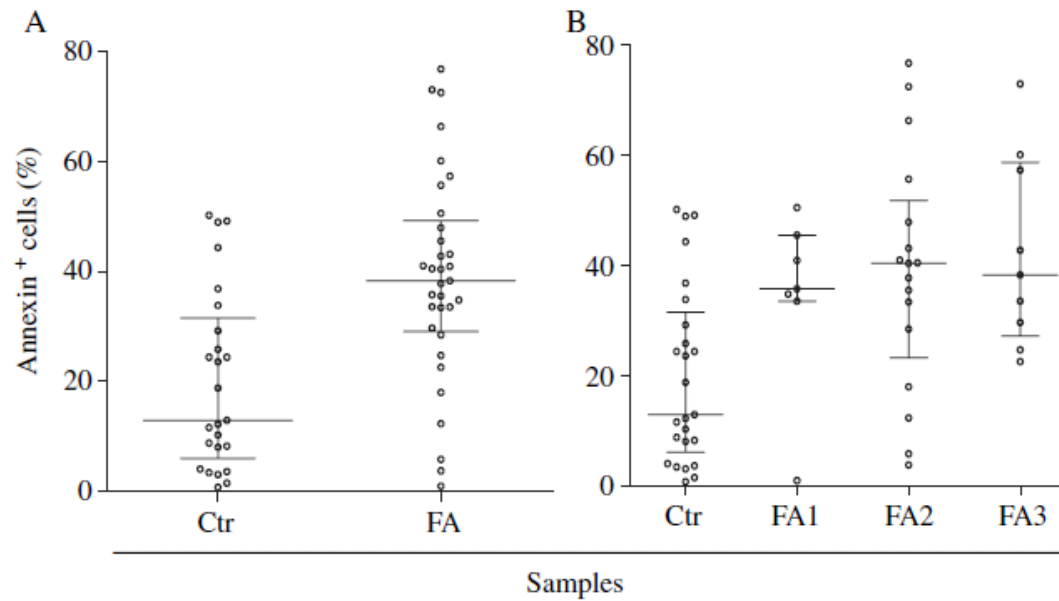
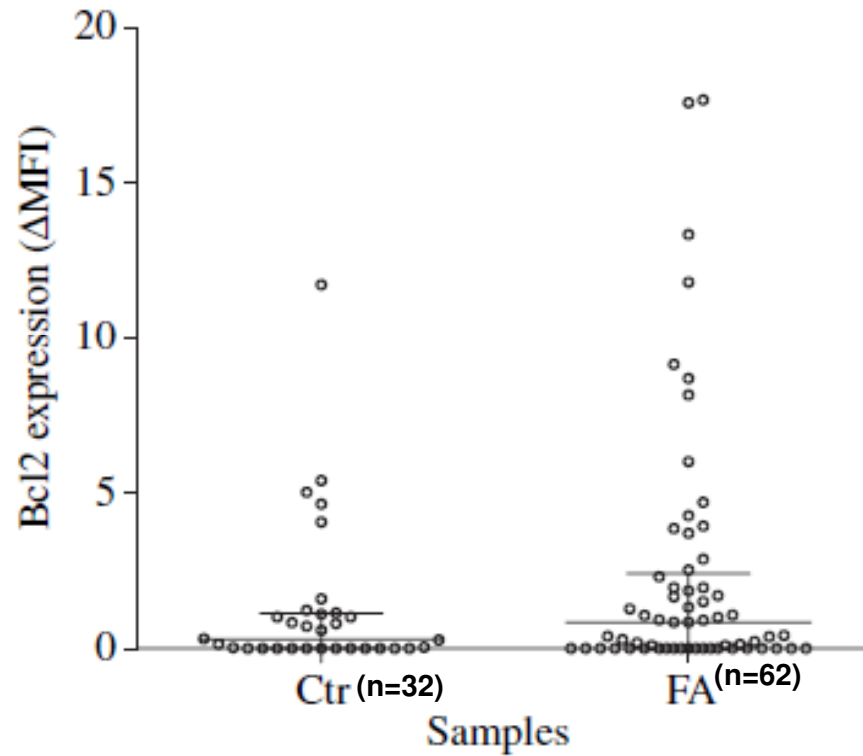


Fig. 5. Activation-induced apoptosis in 24 h PBMC cultures stimulated with 5 µg/mL PHA from FA patients and control samples.





## Normal Bcl-2 expression on Fanconi anemia patients



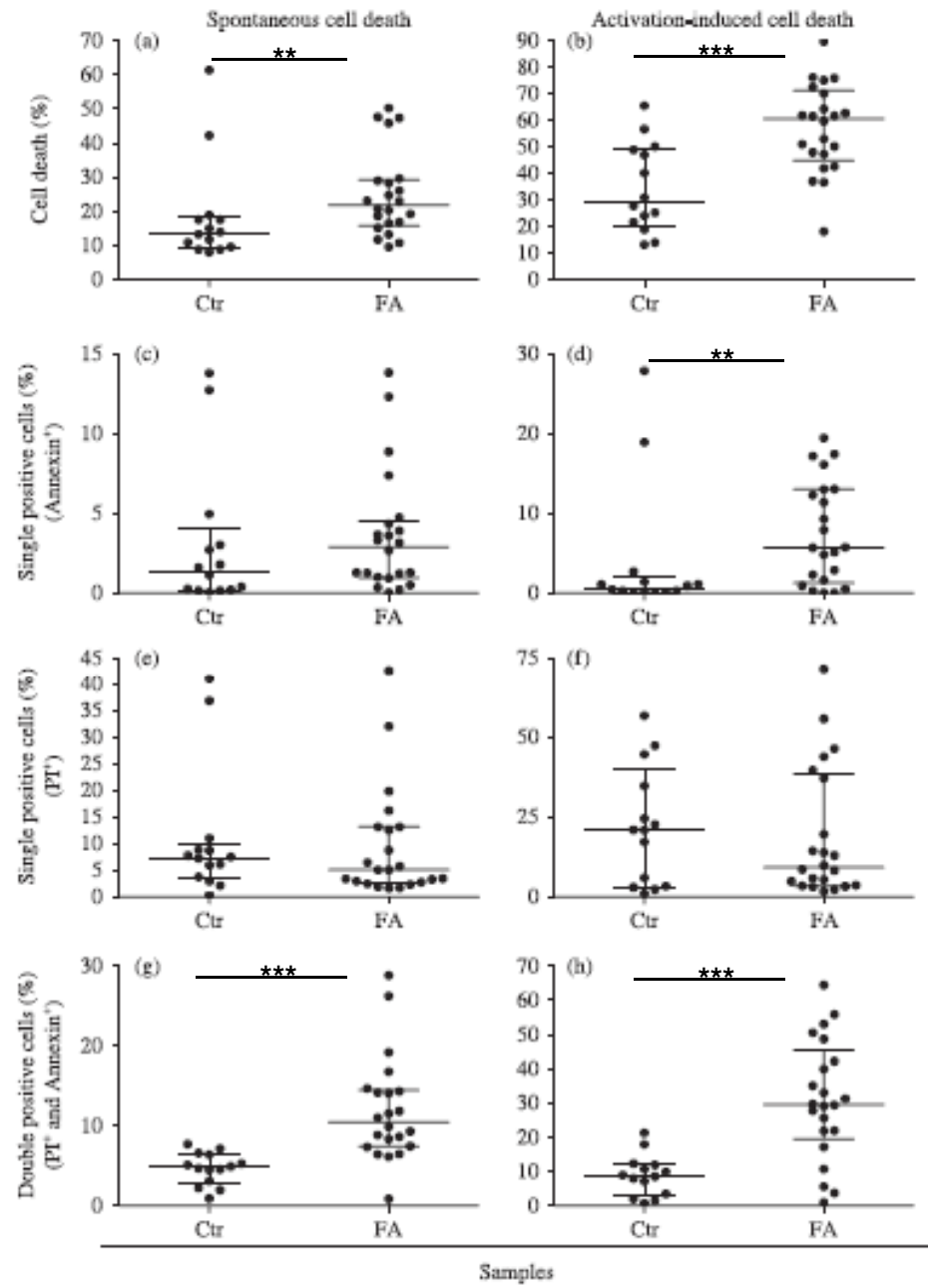


Figure 3. Percentage of cell death in 24 h lymphocyte cultures from 26 FA patients and 14 control samples.



## Bax expression on Fanconi anemia patients

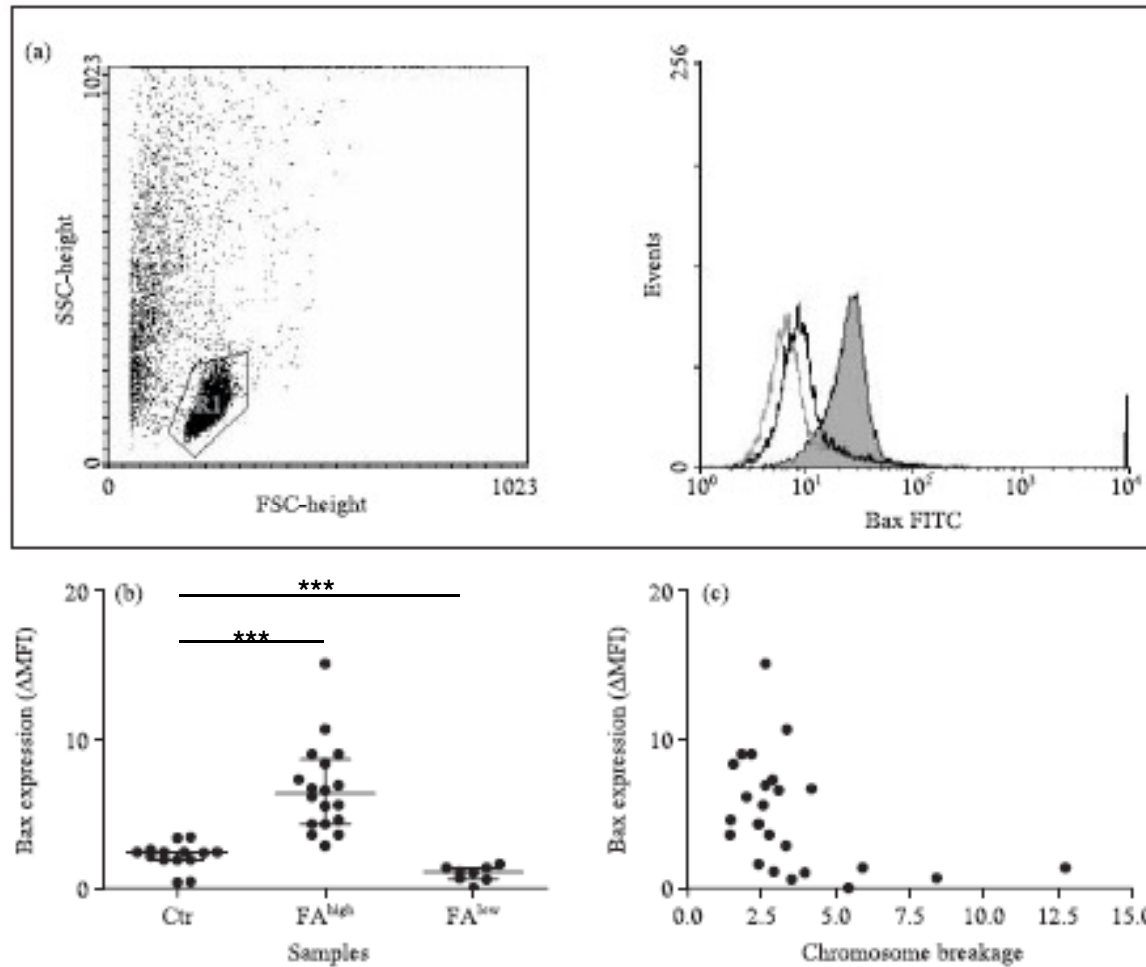
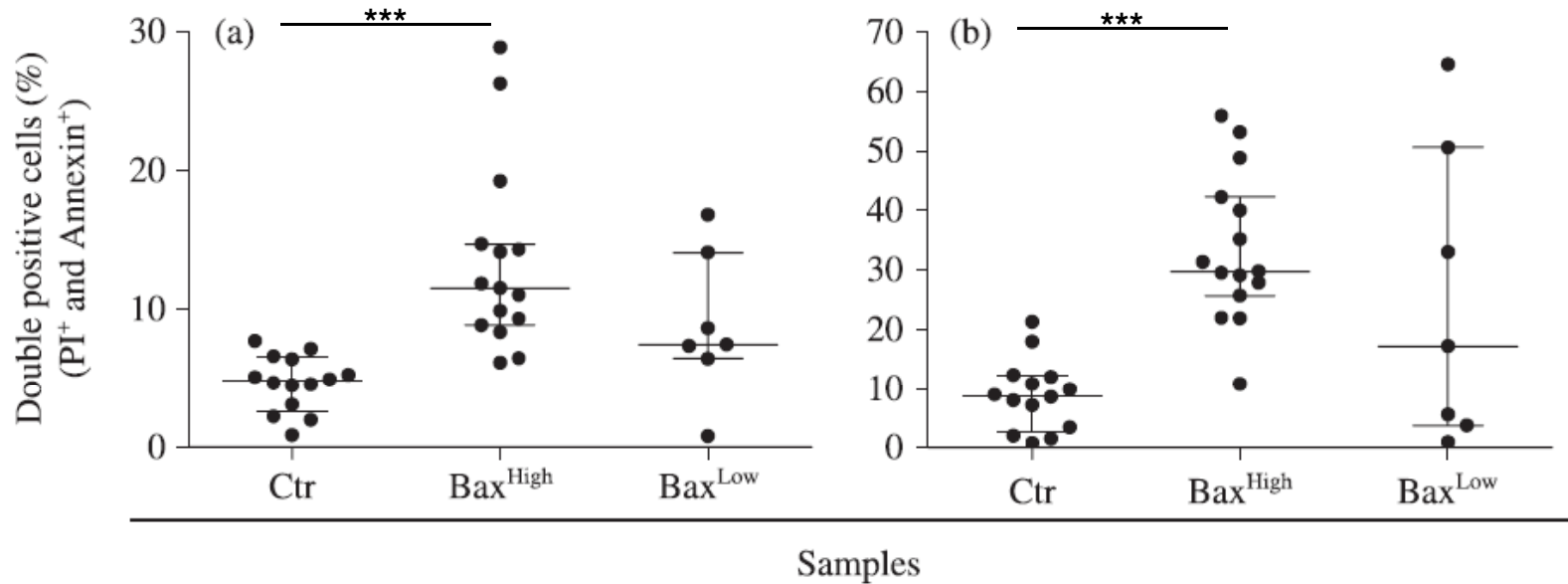


Figure 1. Intracellular Bax expression in permeabilized lymphocytes from Fanconi anaemia (FA) patients



## Direct relation between high Bax expression and increased apoptosis in Fanconi anemia



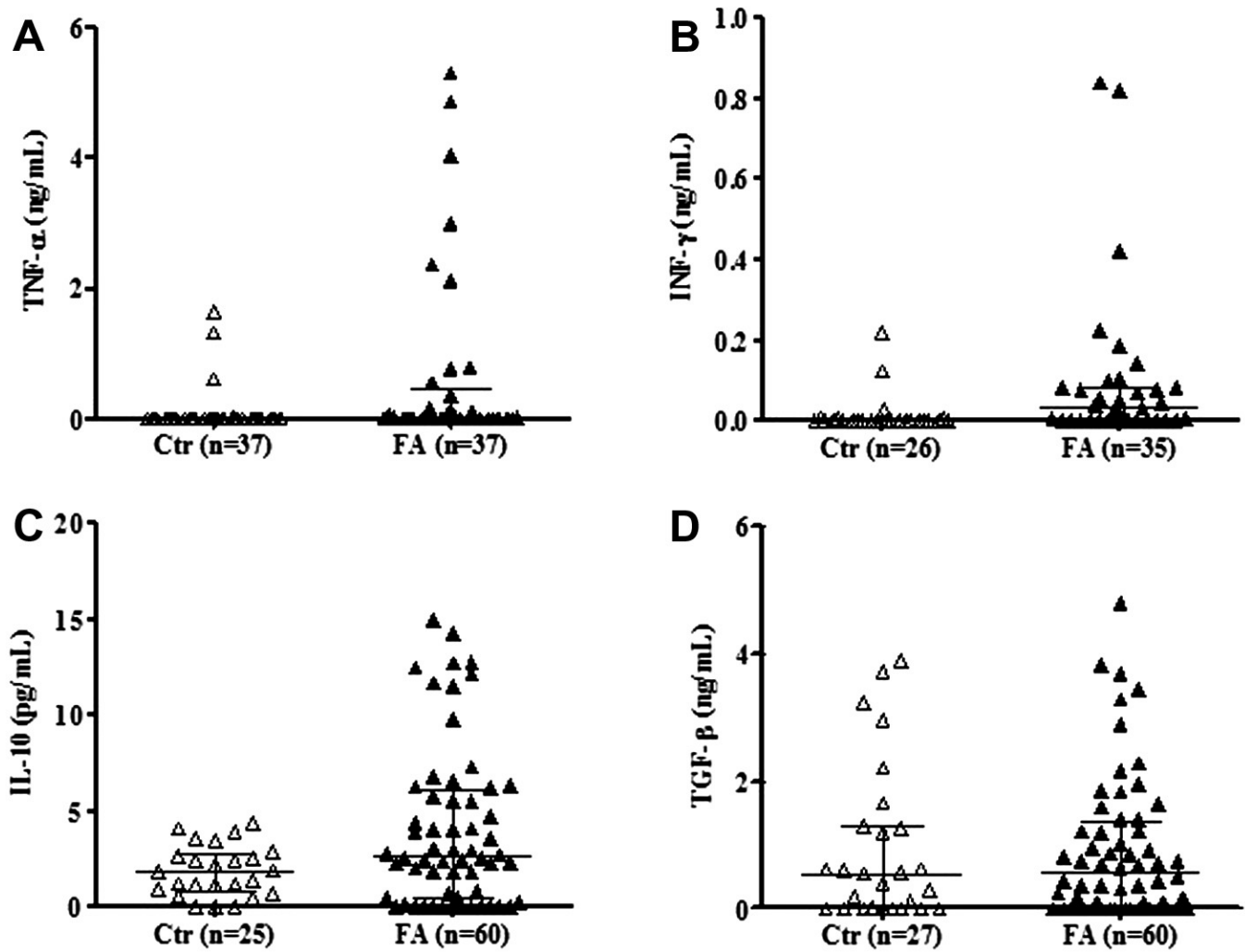
**Figure 4. Percentage of secondary necrosis in the Bax subgroups (high and low).** Scatter plots represent: (a) spontaneous cell death; (b) activation-induced cell death. Ctr = control samples ( $n = 14$ ); Bax<sup>high</sup> ( $n = 15$ ), Bax<sup>low</sup> ( $n = 7$ ).



## Conclusions

- The results, obtained with samples from 26 FA patients, confirm that lymphocytes of the majority of FA patients are more susceptible to cell death, especially activation-induced, that the death process has features of both necrosis and apoptosis, and that this susceptibility is associated with increased Bax expression.
- The results also suggest that the mitochondrial pathway is involved in the majority of FA samples.
- The extrinsic pathway that depends on the activation of death receptors is also involved by the increased expression of Fas receptor, but Bax showed to be a better indicator of apoptosis than the Fas receptor in lymphocytes of FA patients.
- Despite this apparent increased susceptibility of peripheral lymphocytes to apoptotic induction, no correlation could be observed between these proteins levels (Bax and Fas) and the various haematological parameters or androgen therapy.

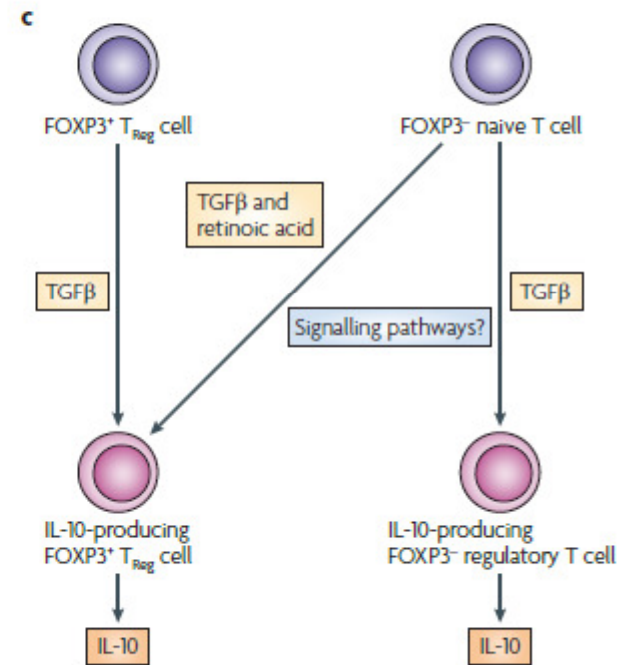
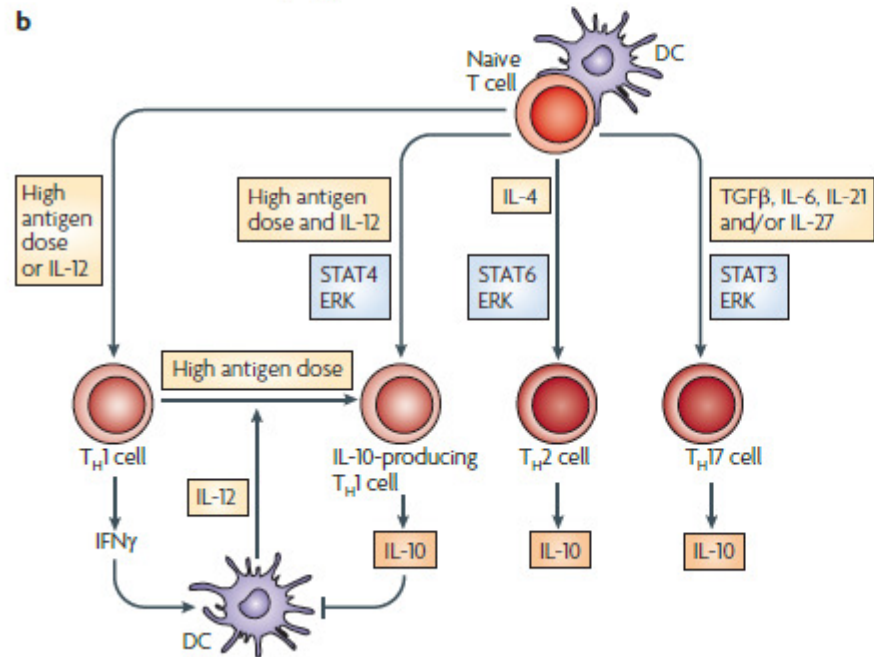
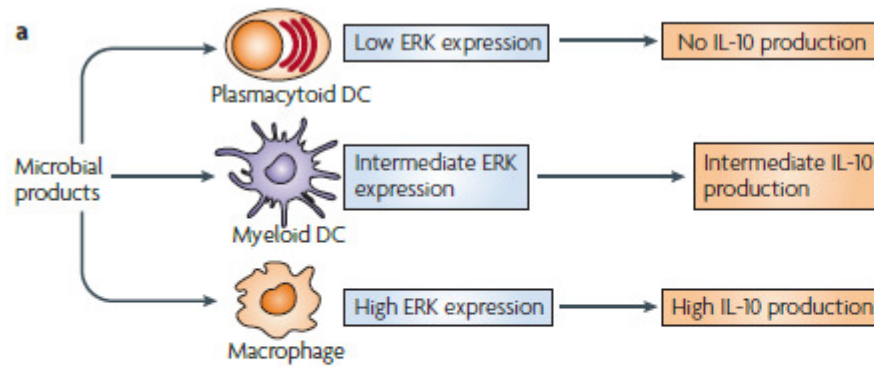




**Fig. 1.** Plasmatic levels of cytokines in the plasma from FA patients' and healthy controls' (Ctr) samples by ELISA assay



# Interleukin-10 expression in the immune system



## Cytokines and Hematological Clinical features

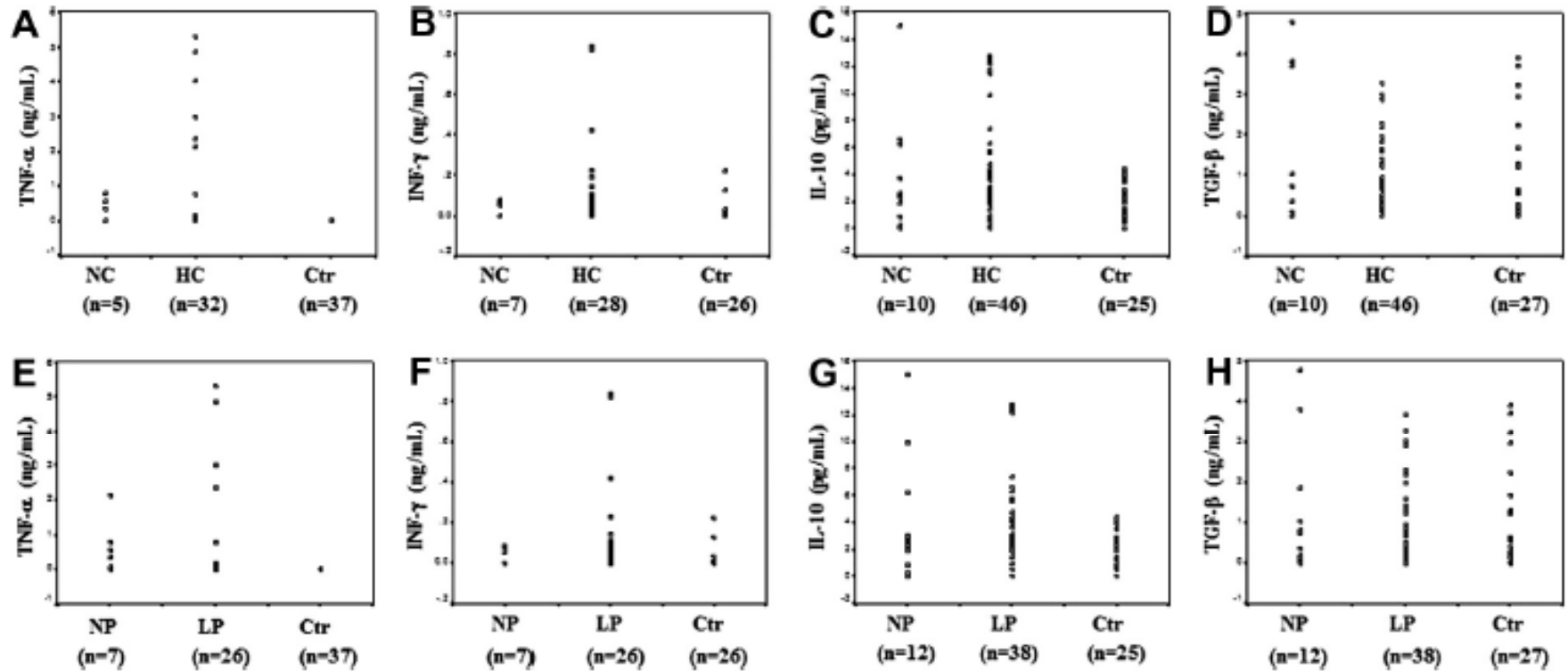


Fig. 2. Variation of FA patients plasmatic levels of cytokines according to hematological clinical features and platelet levels.



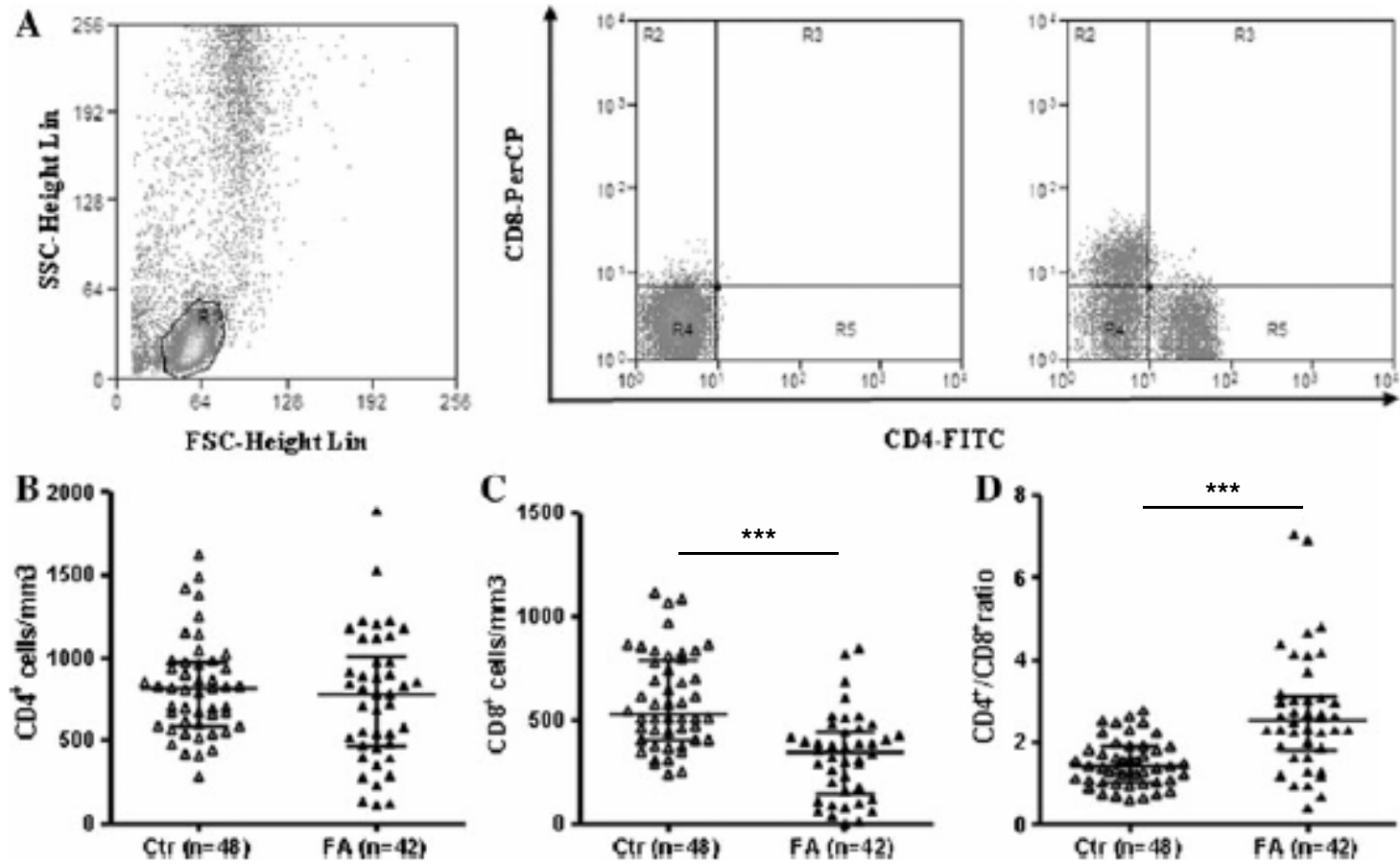


## Conclusions

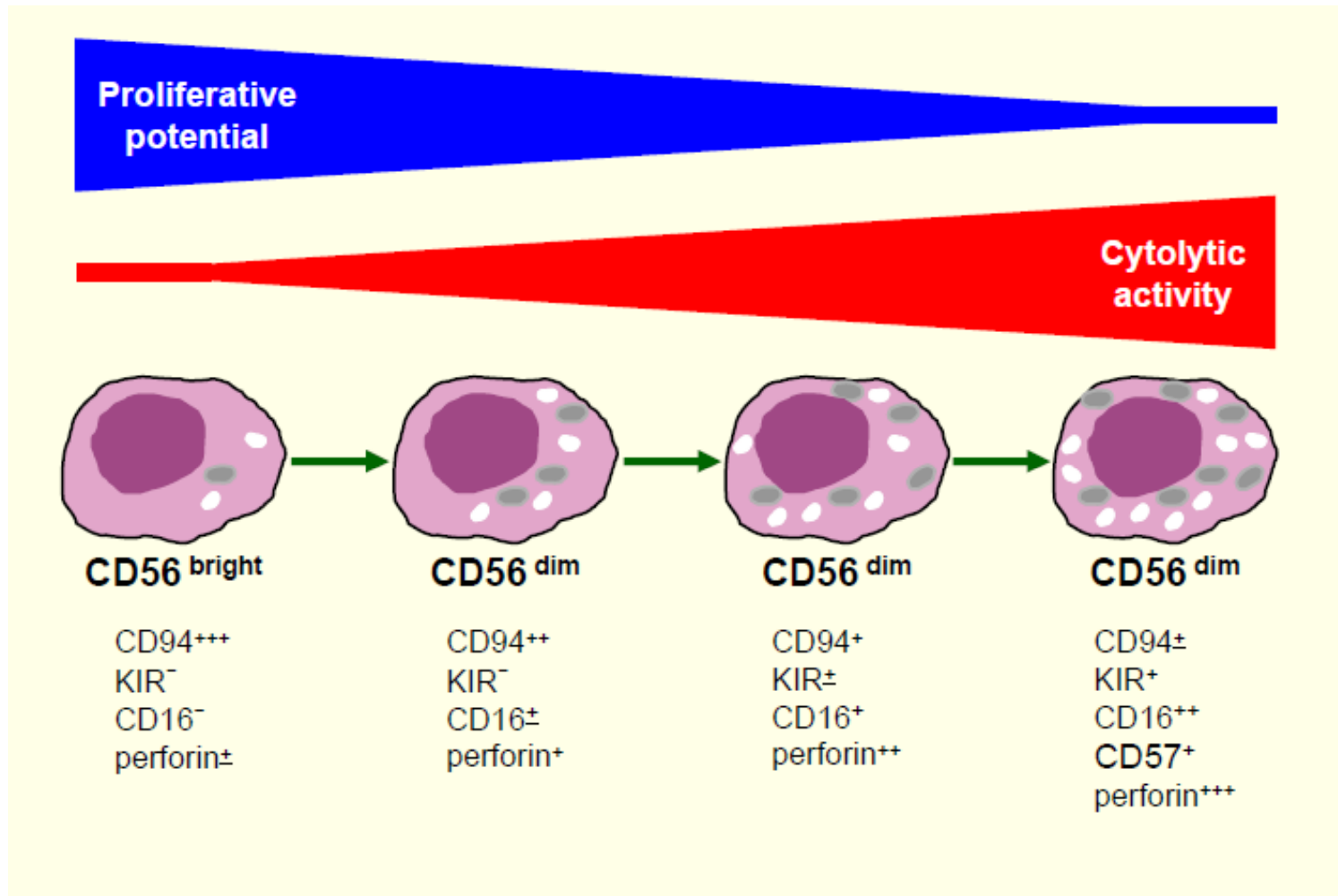
- increased plasma levels of TNF- $\alpha$  and INF- $\gamma$  were observed in 24% and 23% of the patients, respectively, without a correlation between the levels of the two cytokines, suggesting independent phenomena
- Elevated IL-10 plasma levels were also observed in 25% of the FA patients, but no correlation was seen between IL-10 and IFN-  $\gamma$
- Our data suggest that augmented pro-inflammatory cytokines' levels are present together with bone marrow hypocellularity, a feature that was not observed with IL-10 or TGF-  $\beta$
- Levels of TGF- $\beta$  showed a high variation among the healthy controls and were within the normal range in FA samples, but correlated with IL-10 plasma levels



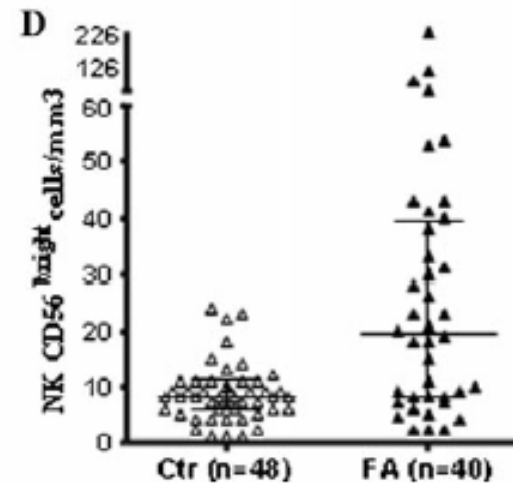
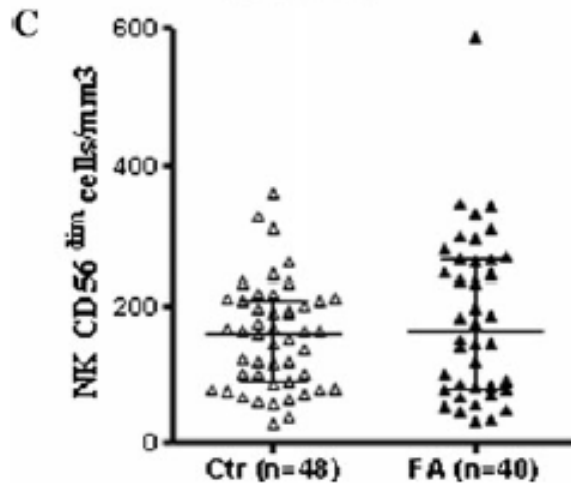
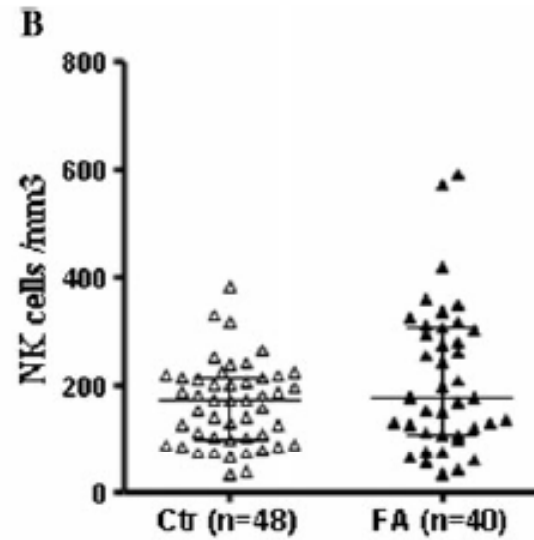
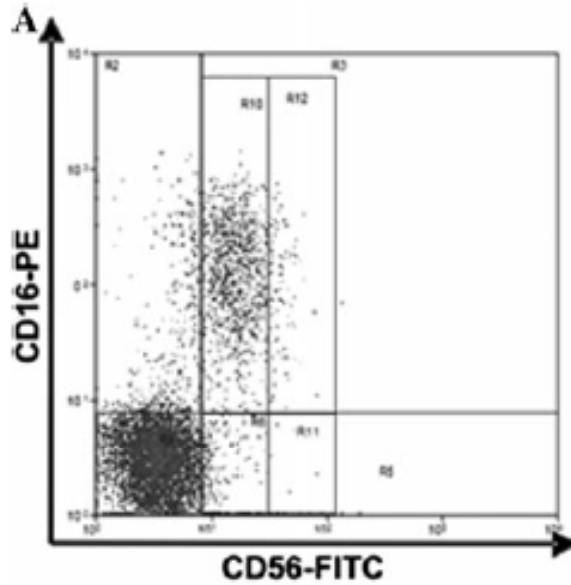
## PBMC Lymphocyte populations of FA patients



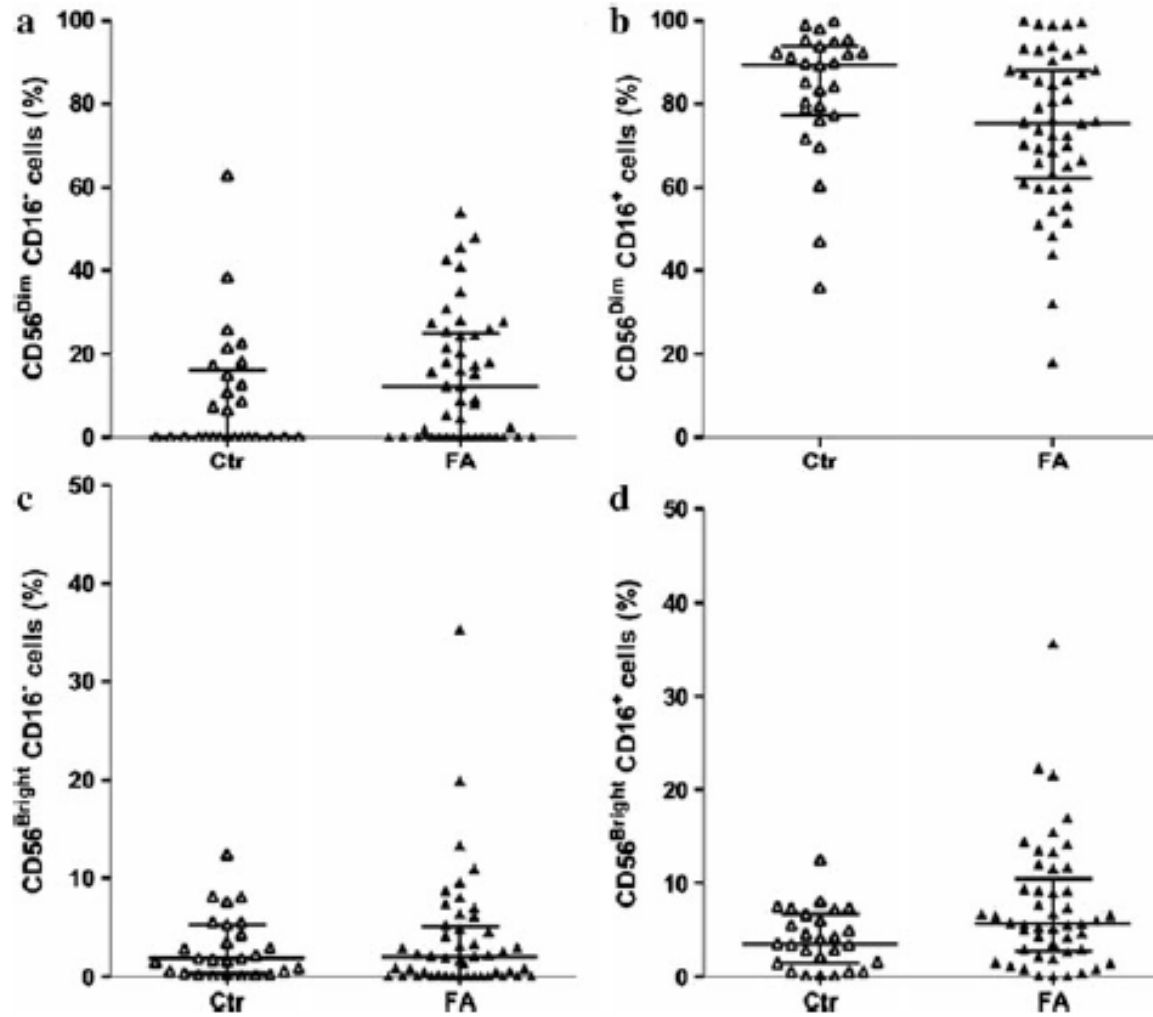
peripheral blood CD56<sup>bright</sup> give rise to CD56<sup>dim</sup> NK cells.



# Natural Killer cells and its main subsets CD56<sup>dim+</sup> and CD56<sup>bright+</sup>



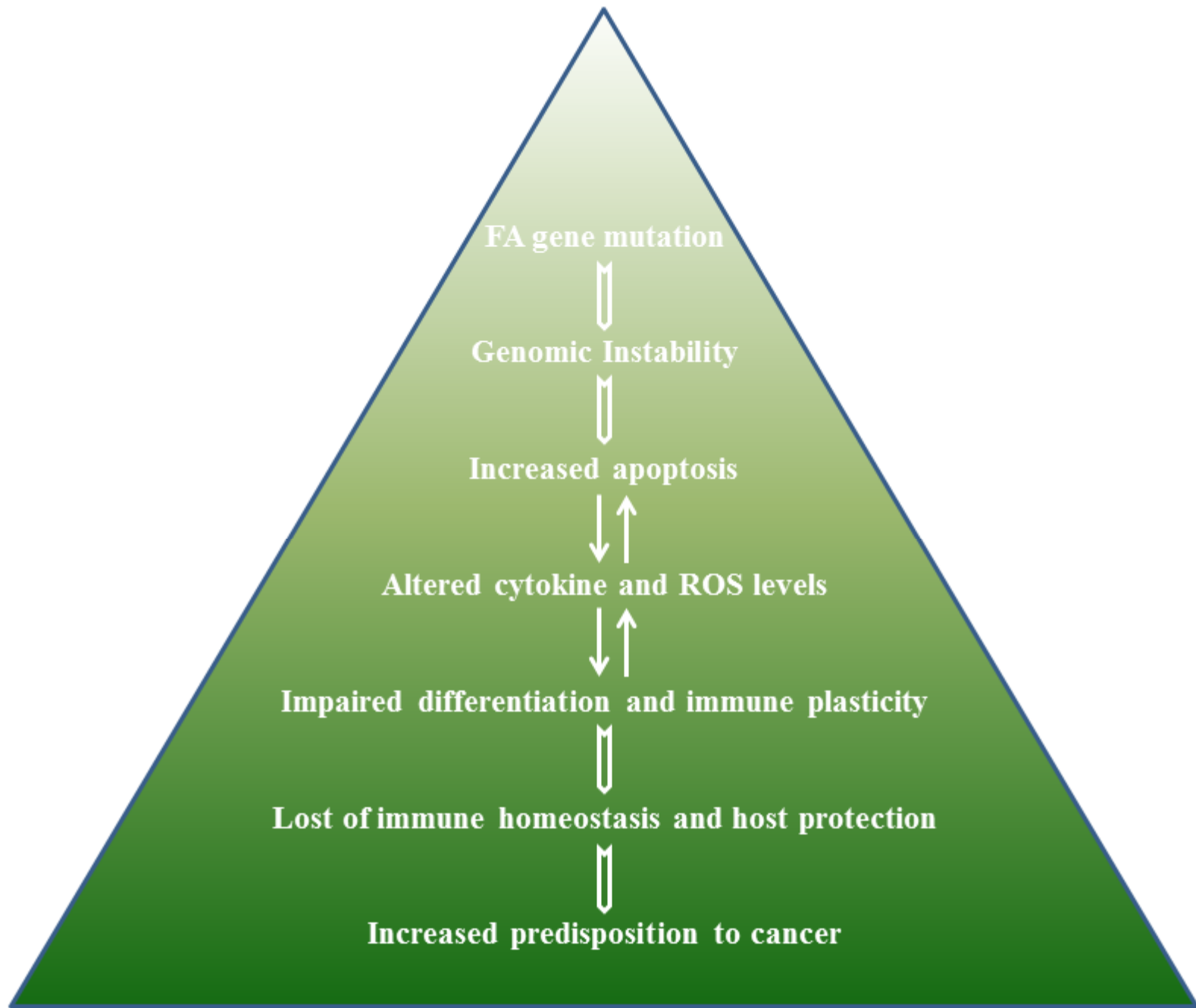
# CD56<sup>dim+</sup> and CD56<sup>bright+</sup> Natural Killer cells based on CD16 expression



## Conclusions

- A decrease in the number of cytotoxic CD8<sup>+</sup> T cells and CD56<sup>dim</sup>CD16<sup>+</sup> NK cells, was observed, but not in the number of CD4 T cells
- the diminished number of CD8 T cell lymphocytes observed in this work suggests that it may lead to higher rates of vulnerability to infections observed in FA patients
- Total NK levels were within the normal range, but there was an imbalance between its main cell subsets CD56<sup>bright</sup> and CD56<sup>dim</sup>; with the prevalence of the more undifferentiated population NK CD56<sup>bright</sup>
- Our results showed that additionally to a defect in the cytotoxic response, FA cells seem to present a defective differentiation of NK cells in their subpopulations:
  - CD56<sup>bright</sup> NK cells were increased in patients with bone marrow hypocellularity,
  - decreased levels of NK CD56<sup>dim</sup>CD16<sup>+</sup> cells were observed in patients with normal hematological clinical features.







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Thank you!!!

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