# PIASTRINOPENIE COSTITUZIONALI «PERICOLOSE»: QUANDO L'APPARENZA INGANNA

Sistema Socio Sanitario





ATS Pavia

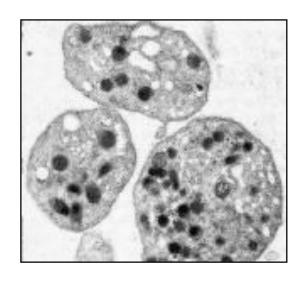
Federica Melazzini, MD PhD

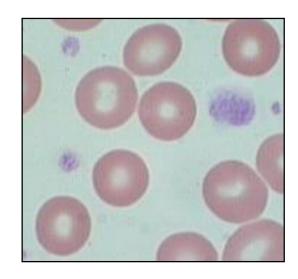
SC Medicina Generale I. Fondazione IRCCS Policlinico San Matteo

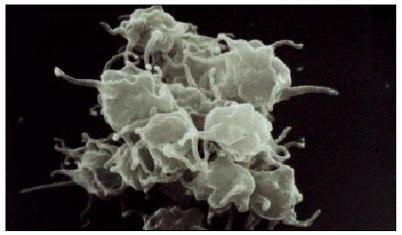


# **SUMMARY**

- Inherited thrombocytopenias: past & present
- Cases
- Predisposing Syndromes
- Discussions & Conclusions



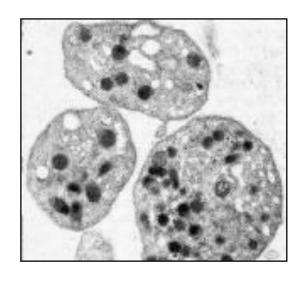


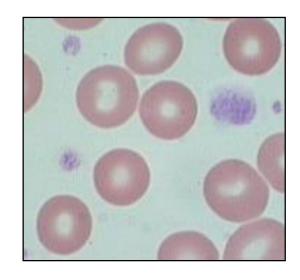


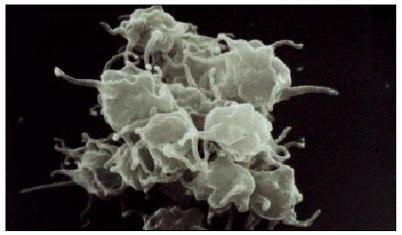


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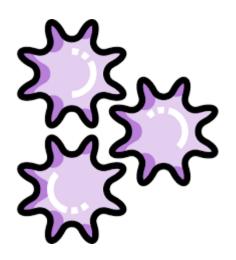






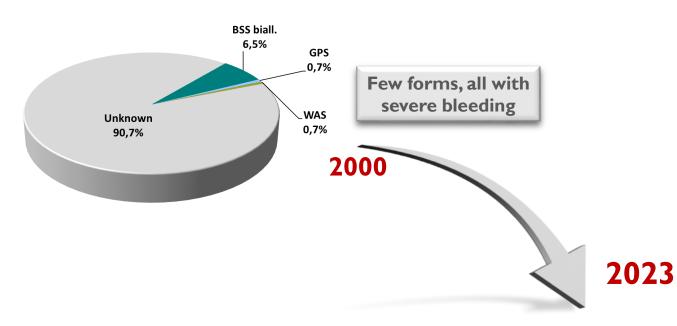
# INHERITED THROMBOCYTOPENIAS

- Rare disorders
- Reduced platelet number ( $<150 \times 10^9/L$ )
- Mainly primary haemostasis defects
- Bleeding tendency not always correlates with platelet count
- Often misdiagnosed (unnecessary therapies)
- Genetically heterogeneous though often with similar phenotypes



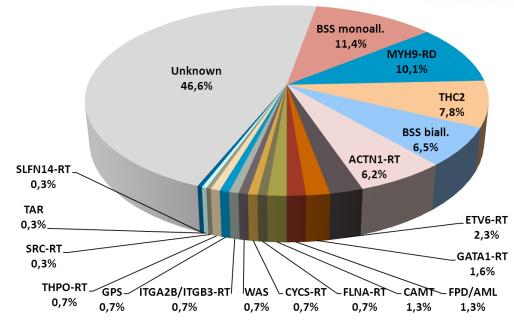








Many forms. Mild or no bleeding in most cases
Risk of developing additional disorders



# **«OLD»** CLASSIFICATION FOR ITS

Inheritance pattern

Autosomal dominant
Autosomal recessive
X-linked
De novo mutations

Platelet size

Giant/large platelets
Normal sized platelets
Small platelets

Mechanism of thrombocytopenia

Defects in Mk differentiation

Defects in Mk maturation

Defects in platelet release

Shortened platelet survival

Presence of additional symptoms

Syndromic forms

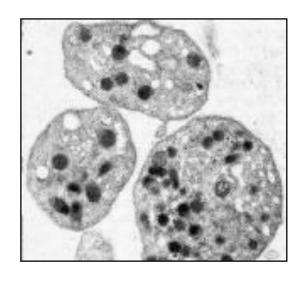
Non syndromic forms

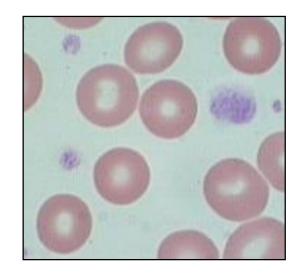
# **«NEW»** CLASSIFICATION FOR ITS

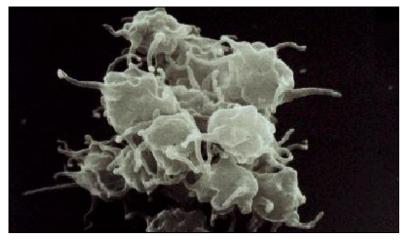
**ONLY OTHER PREDISPOSING SYNDROMES** thrombocytopenia **Congenital Defects** • Brain defects (JBS, FLNA-RT) • Facial dysmorphism • Deafness (CTRUS) (TAR, JBS) • Hearth defect (TAR, JBS) Kidney defects • Enlarged spleen (TAR, JBS) (GPS, GATA1-RD) Bowel defect • Upper limb defects (TAR, JBS, WAS) (TAR, CTRUS) • Genitalia defects (JBS) • Lower limb defects (TAR) • Tendon xanthomas (STSL)

# **SUMMARY**

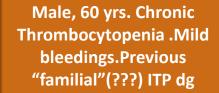
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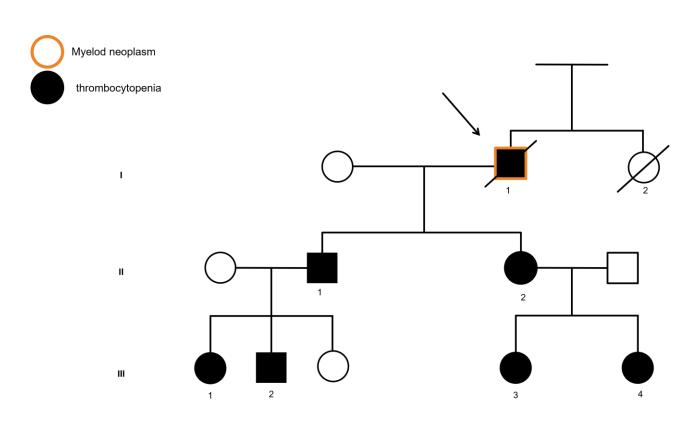


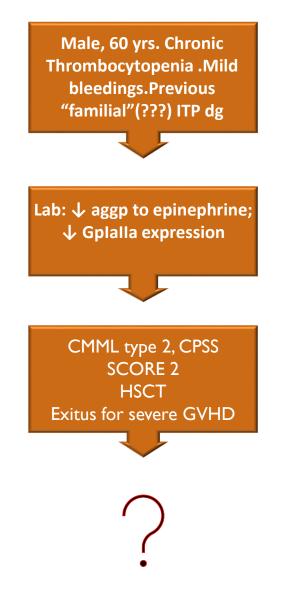


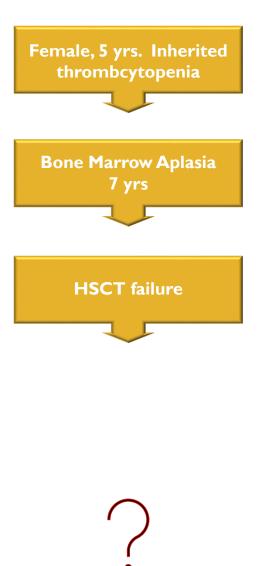
CMML type 2, CPSS SCORE 2 HSCT

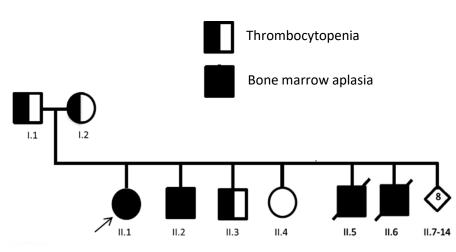
Exitus for severe GVHD

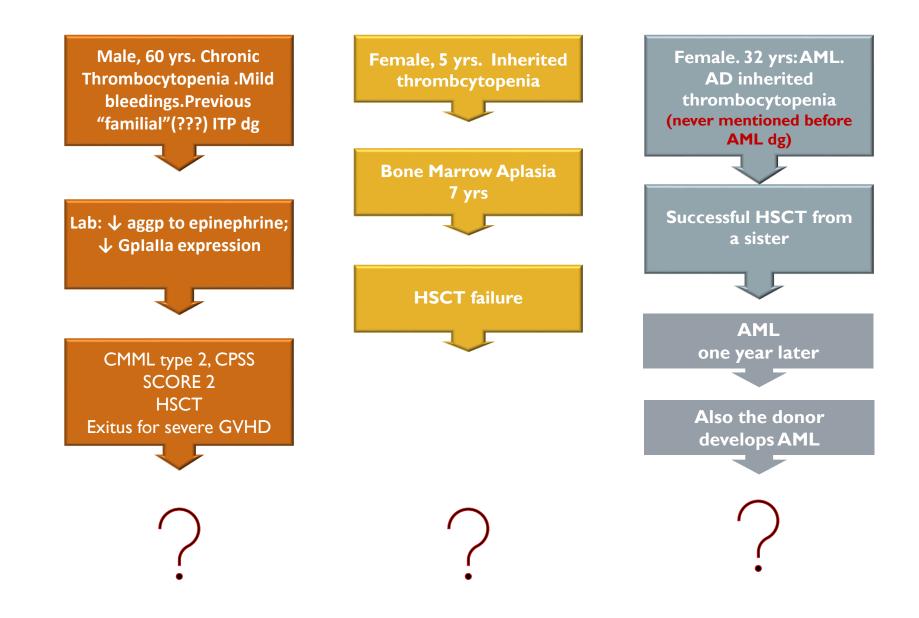


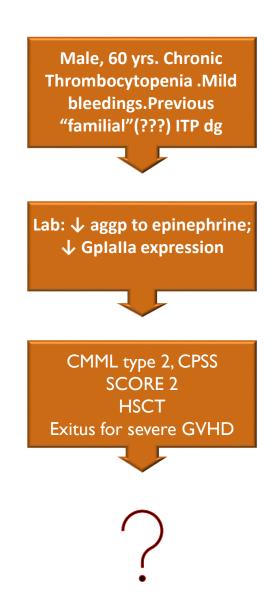


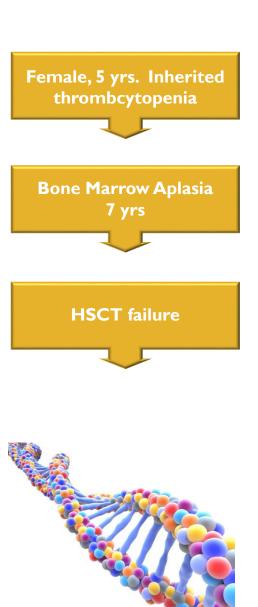


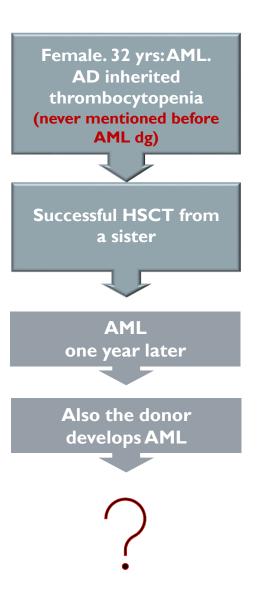


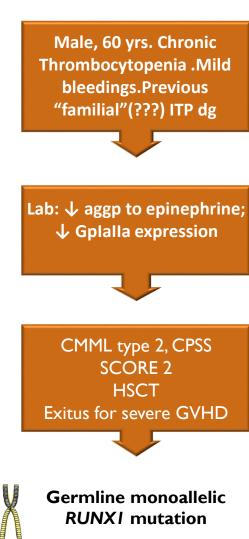


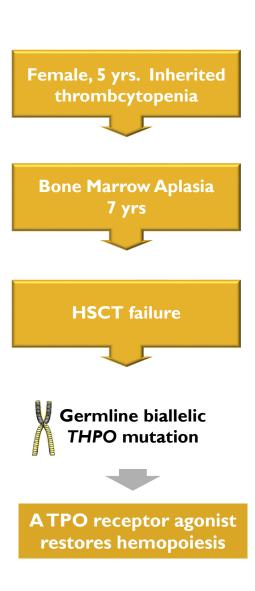


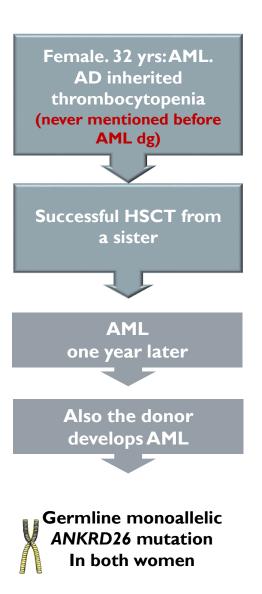






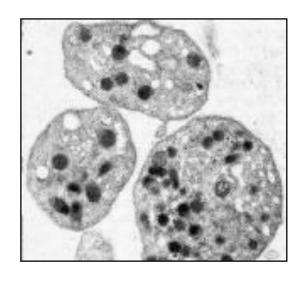


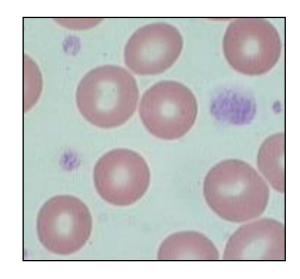


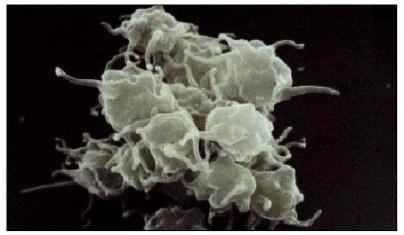


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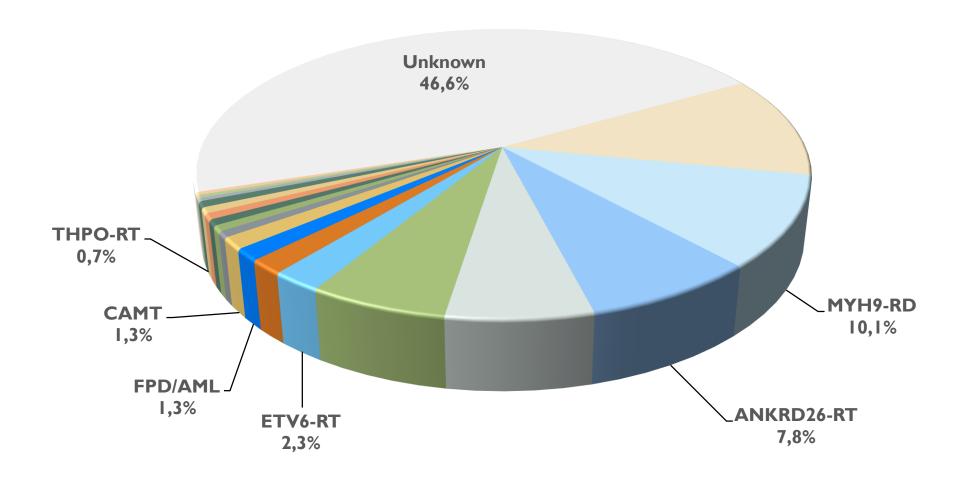


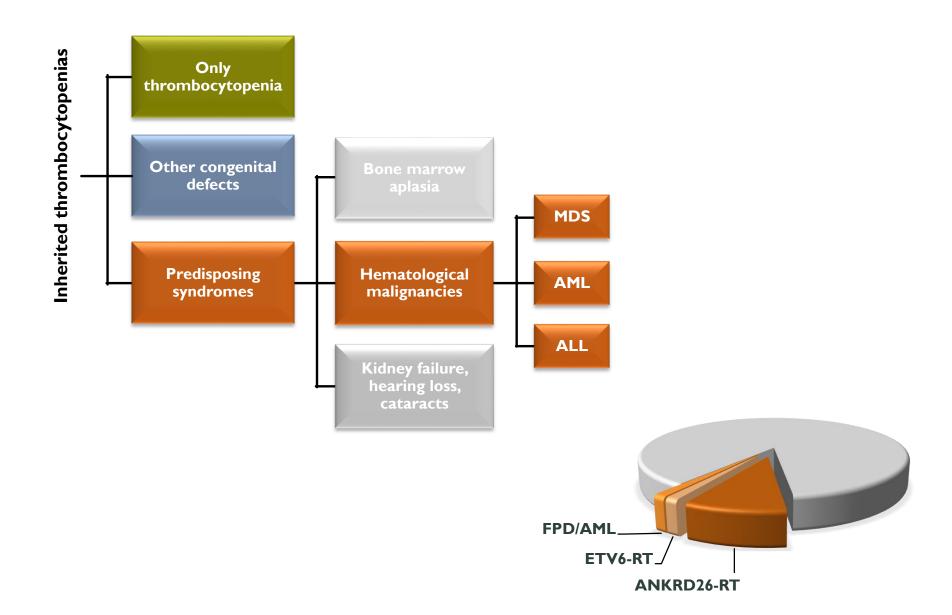






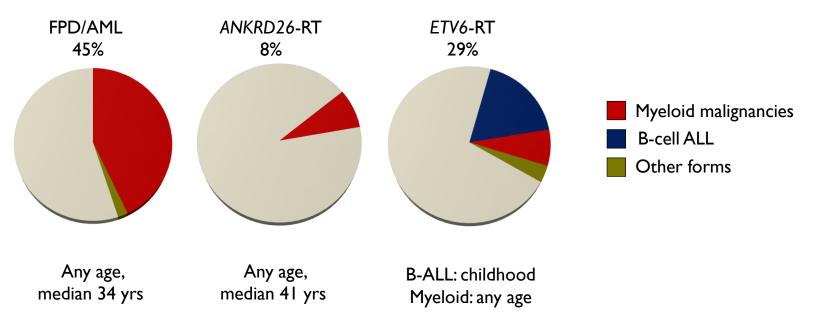
# 45% of patients with known inherited thrombocytopenias are at risk of developing additional disorders





	ANKRD26-RT	FPD/AML	ETV6-RT
Gene	ANKRD26	RUNXI	ETV6
Transmission	AD	AD	AD
Relative frequency (% of known IT)	18%	3%	5%
Thrombocytopenia	Mild/moderate	Mild/absent	Mild
Platelet size	Normal	Normal	Normal
Platelet Function	Normal	Abnormal	Normal
Bleeding tendency	Absent/mild	Absent/moderate	Absent/mild

# % of patients with hematological malignancies



#### THE UPDATED WHO CLASSIFICATION OF HEMATOLOGICAL MALIGNANCIES

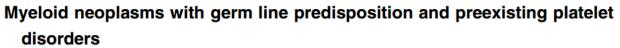
The 2016 revision to the World Health Organization classification of myeloid neoplasms and acute leukemia

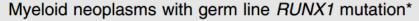
### Myeloid neoplasms with germ line predisposition

# Myeloid neoplasms with germ line predisposition without a preexisting disorder or organ dysfunction

AML with germ line CEBPA mutation

Myeloid neoplasms with germ line DDX41 mutation\*





Myeloid neoplasms with germ line ANKRD26 mutation\*

Myeloid neoplasms with germ line ETV6 mutation\*

### Myeloid neoplasms with germ line predisposition and other organ dysfunction

Myeloid neoplasms with germ line GATA2 mutation

Myeloid neoplasms associated with BM failure syndromes

Myeloid neoplasms associated with telomere biology disorders

JMML associated with neurofibromatosis, Noonan syndrome or

Noonan syndrome-like disorders

Myeloid neoplasms associated with Down syndrome\*

<sup>\*</sup>Lymphoid neoplasms also reported.

We do not know why patients with FDP/AML, ANKRD26-RT and ETV6-RT are prone to hematological malignancies.

However, these disorders have some common features:

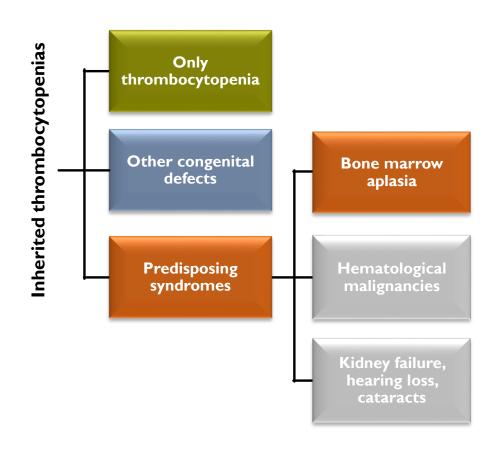
**RUNXI** and **ETV6** are transcription factors and mutations result in loss of transcriptional repression. Mutation in **ANKRD26** affect the 5'UTR of the gene and result in loss of transcriptional repression by RUNXI.

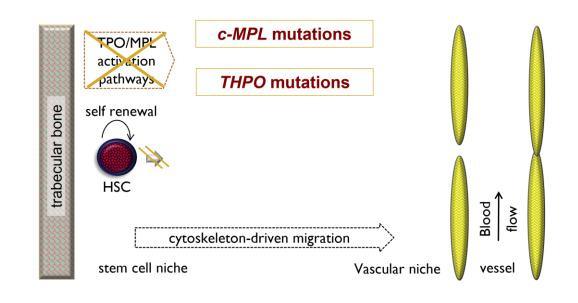
The balance between proliferation and differentiation of hemopoietic progenitors is deranged: increased proliferation.

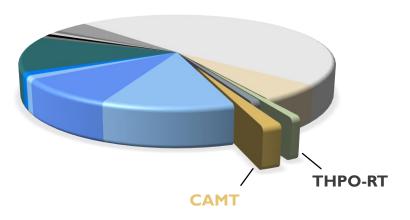
Penetrance of malignancies is incomplete: other factors are required for oncogenesis.

### **RECOMMENDATIONS**

- Bone marrow examination with cytogenetic analysis at diagnosis
- > Complete blood count and clinical examination at regular intervals (each year?)
- Any time significant changes in the blood counts are identified: repeat bone marrow examination (and cytogenetic analysis)
- In case patients develop hematological malignancies, HSCT is an important option. If a related donor is available, exclude that he is affected too by the same disorder: transplanting from affected relatives exposes to the risk of developing again a new hematological malignancy





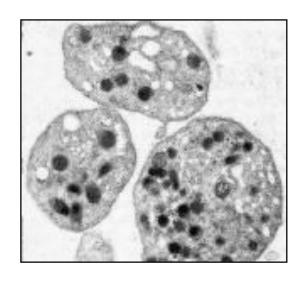


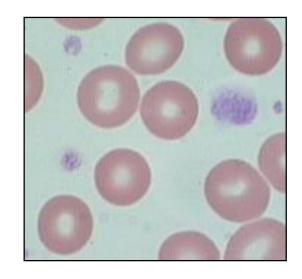
	CAMT	CAMT variant
Gene	MPL	ТНРО
Frequency	Low	Low
Transmission	AR	AR
Thrombocytopenia	Severe	Severe/moderate
Platelet size	Normal	Normal
Evolution into bone marrow aplasia	Infancy	Infancy/young adult
HSCT	Effective	Not effective

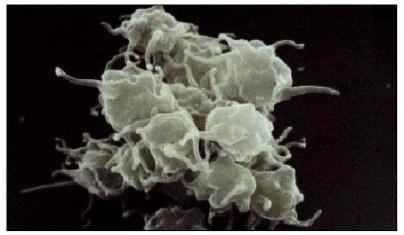
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# **DISCUSSION**

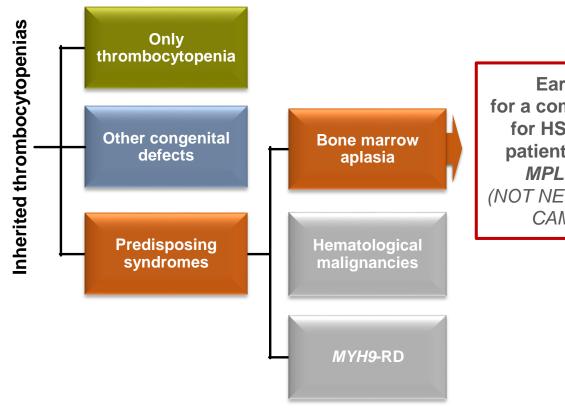
✓ Why look for a molecular diagnosis of an inherited thrombocytopenia?

✓ How to reach the molecular diagnosis of an inherited thrombocytopenia?

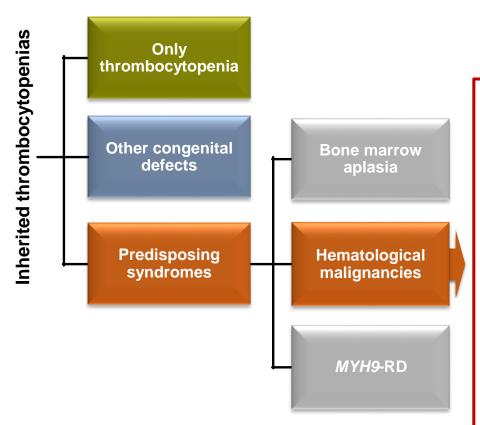
# **DISCUSSION**

✓ Why look for a molecular diagnosis of an inherited thrombocytopenia?

✓ How to reach the molecular diagnosis of an inherited thrombocytopenia?



Early search
for a compatible donor
for HSCT in CAMT
patients caused by
MPL mutations
(NOT NECESSARY FOR
CAMT variant)



#### **AT DIAGNOSIS**

- Genetic counseling
- Physical examination
- CBC with differential
- Blood smear examination
- Bone marrow examination with cytogenetic analysis

#### **EVERY 6-12 MONTHS**

- Physical examination
- \* CBC with differential
- Blood smear examination



#### any change

- \* Bone marrow examination with cytogenetic analysis
  - 1

#### evolution towards leukemia

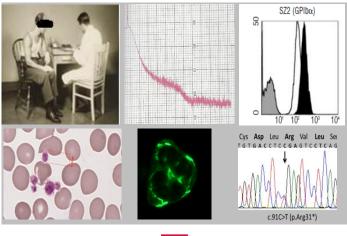
 In case of HSCT from a related donor, mutational screening

# **DISCUSSION**

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✓ How to reach the molecular diagnosis of an inherited thrombocytopenia?

SINGLE step diagnostic approach

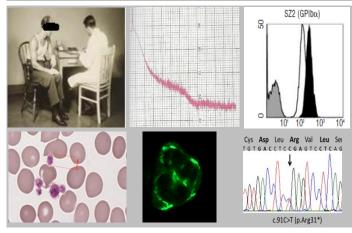




### Time consuming

Performed in many non-specialized laboratories

# **SINGLE step diagnostic approach**





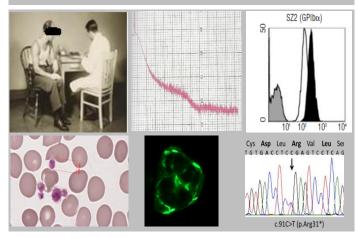
### Time consuming

Performed in many non-specialized laboratories

### **SINGLE step diagnostic approach**



- ✓ Targeted NGS testing
- √ Whole exome sequencing (WES)
- Whole genome sequencing (WGS)





### Time consuming

Performed in many non-specialized laboratories

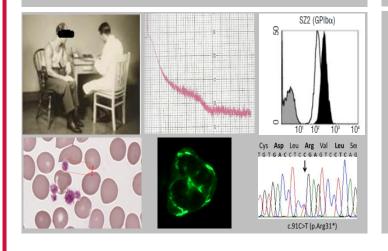
### SINGLE step diagnostic approach



Available only in a few specialized laboratories

Quick if patient has known mutations

<u>Discriminating between pathogenic and</u> <u>non-pathogenetic variants may be a</u> <u>major problem</u>



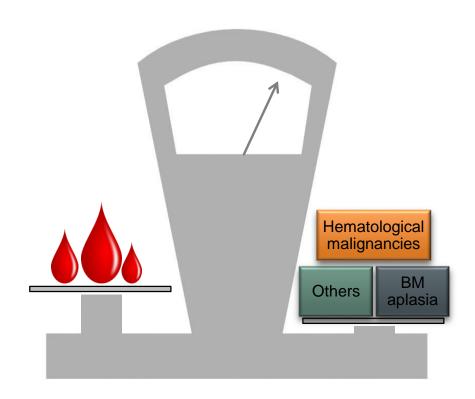
### SINGLE step diagnostic approach



**Targeted NGS testing** 

Next generation sequencing and clinical-laboratory approach are mutually supportive and their combination offers the best chance of reaching the right diagnosis

# CONCLUSION



Bleeding is not the biggest risk for most patients with ITs

A multidisciplinary approach is required to reach the correct molecular diagnosis to arrange the most appropriate treatment and follow-up for all patients

# **GRAZIE!**

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