



EUROPEAN  
HEMATOLOGY  
ASSOCIATION



MADRID  
22<sup>ND</sup> CONGRESS  
JUNE 22-25 | 2017

European Hematology Association

# Novità nelle MDS

***Matteo G Della Porta***

*Cancer Center*

*IRCCS Humanitas Research Hospital*

*& Humanitas University*

*Rozzano – Milano, Italy*

[matteo.della\\_porta@hunimed.eu](mailto:matteo.della_porta@hunimed.eu)

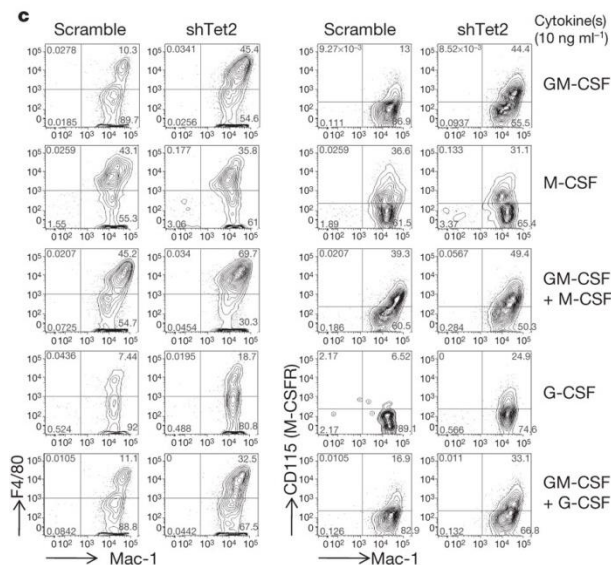
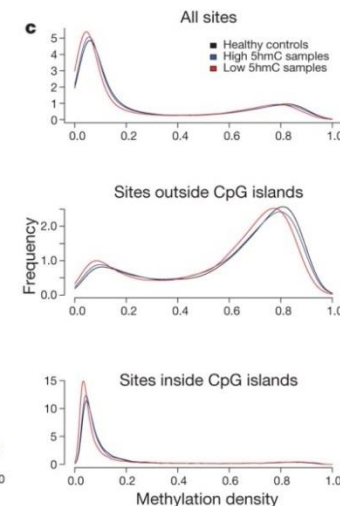
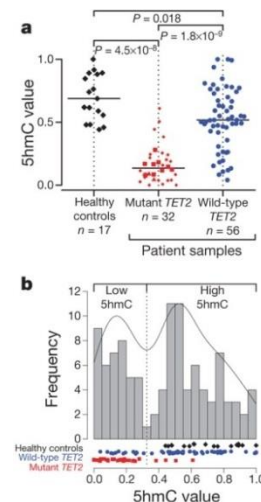
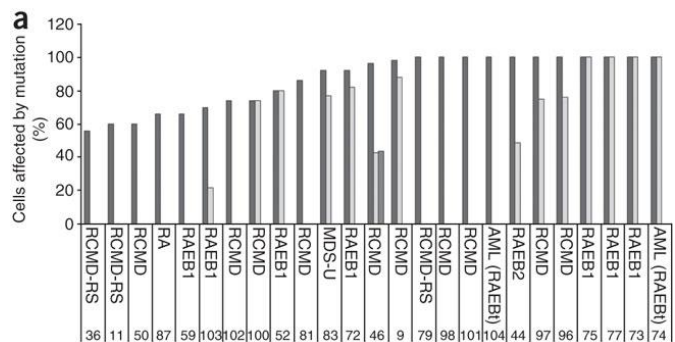
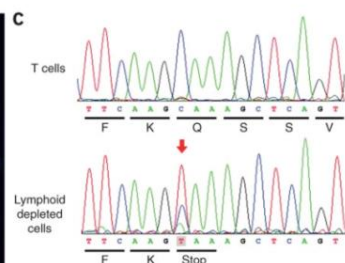
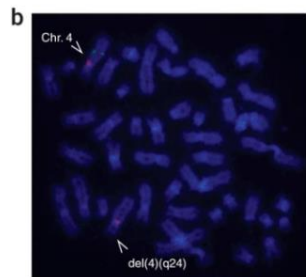
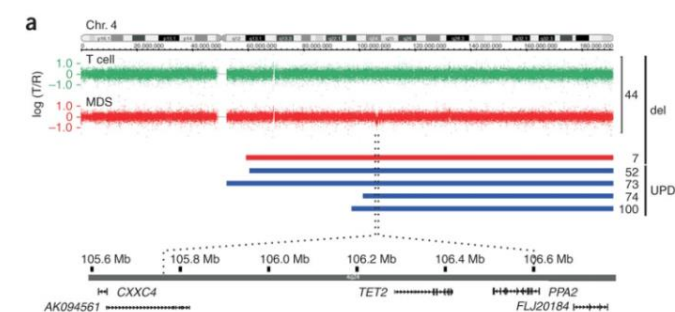
**HUMANITAS**  
RESEARCH HOSPITAL

**HUMANITAS**  
UNIVERSITY

# Outline

- ARCH
- Predictive value of somatic mutations
- Treatment of anemia

# Acquired mutations in *TET2* in myeloid neoplasms

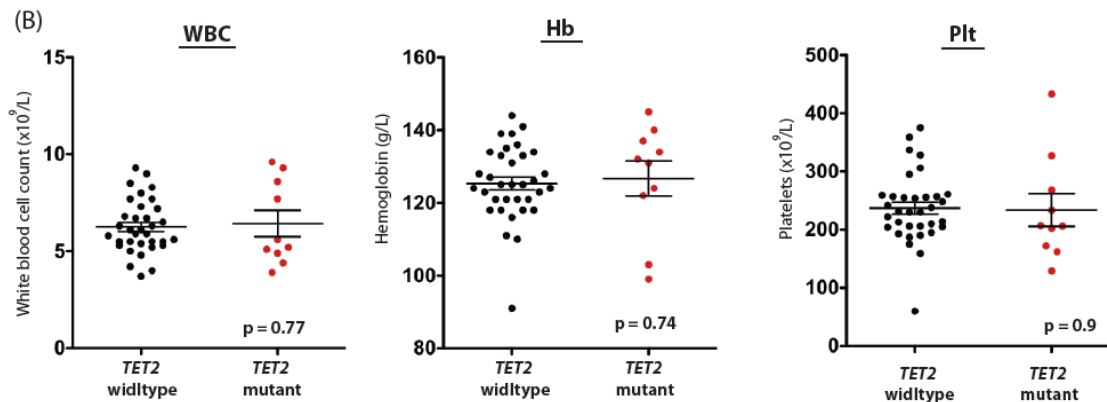
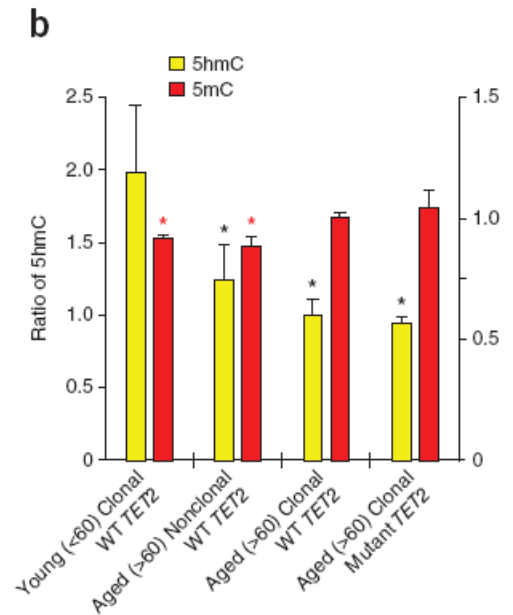


# Recurrent somatic *TET2* mutations in normal elderly individuals with clonal hematopoiesis

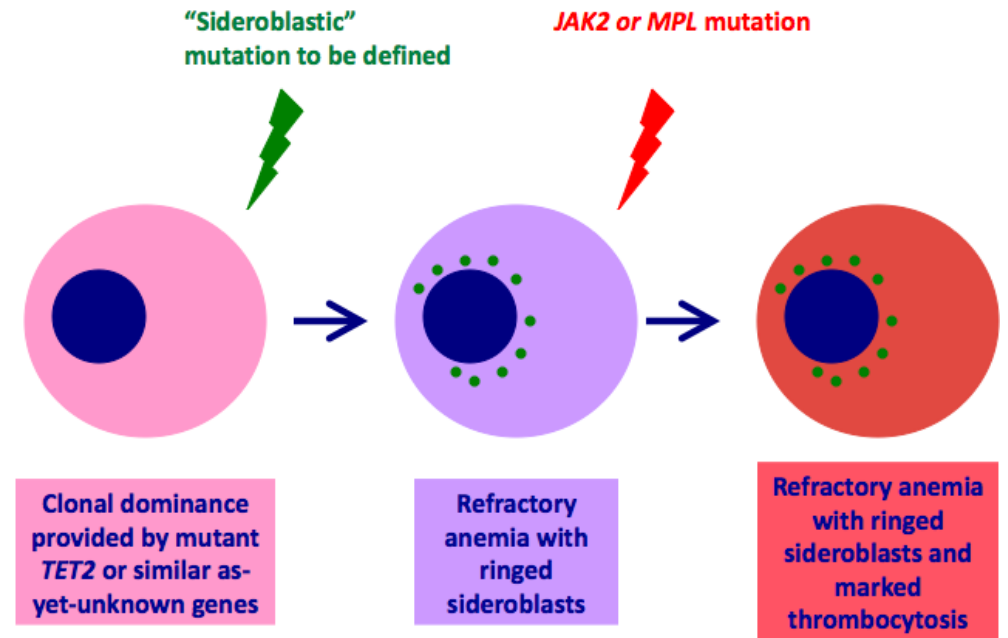
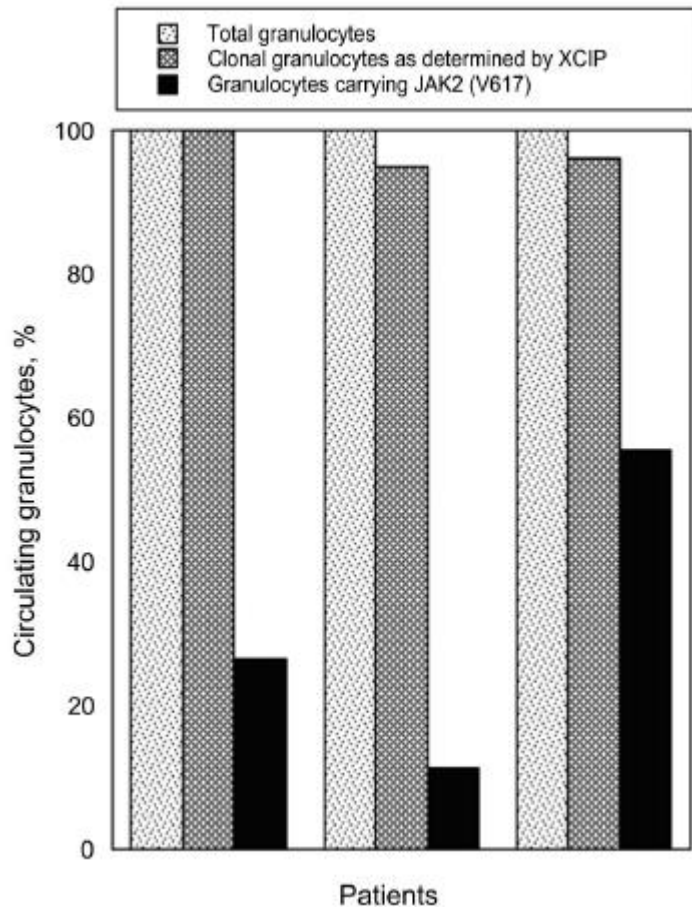
**Table 1** *TET2* somatic mutations found in normal elderly individuals ( $n = 10$ )

Nucleotide substitution <sup>a</sup>	Amino-acid substitution	Chromosome	Position
c.286_298delCGCAC AGTTAGTG	p.Arg96Asnfs*12	4	106155385
c.1330delA	p.Thr444Hisfs*6	4	106156429
c.1348delA	p.Lys450Lysfs*2	4	106156447
c.1547delC	p.Pro516Hisfs*16	4	106156646
c.1630C>T	p.Arg544*	4	106156729
c.3311_3312insAT	p.Phe1104Leufs*3	4	106158411
c.3991A>C	p.Thr1331Pro	4	106182952
c.5200delA	p.Met1734Leufs*11	4	106196867
c.5575insT	p.Ile1859tyrfs*16	4	106197239
c.5725G>T	p.Glu1909*	4	106197392

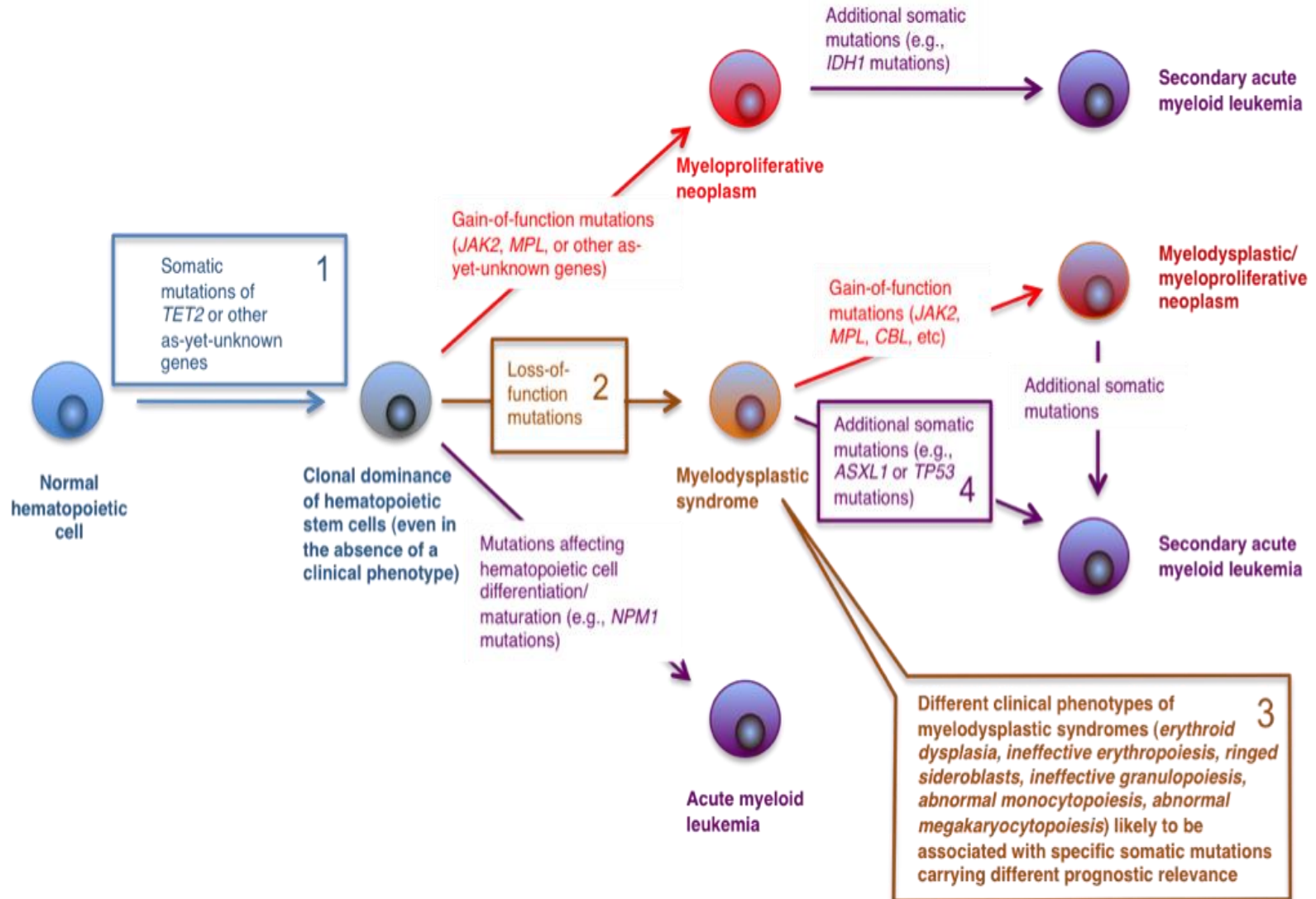
<sup>a</sup>The reference sequence used to annotate *TET2* mutations was [NM\\_001127208](#).



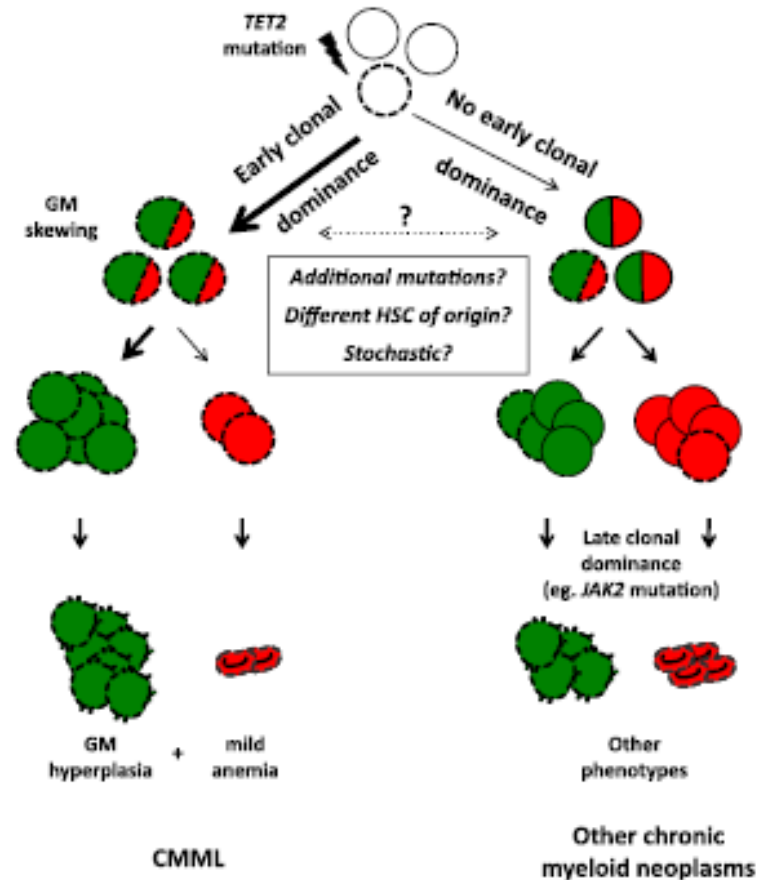
# Molecular and clinical features of refractory anemia with ringed sideroblasts associated with marked thrombocytosis



# Molecular pathogenesis of Myeloid Neoplasms

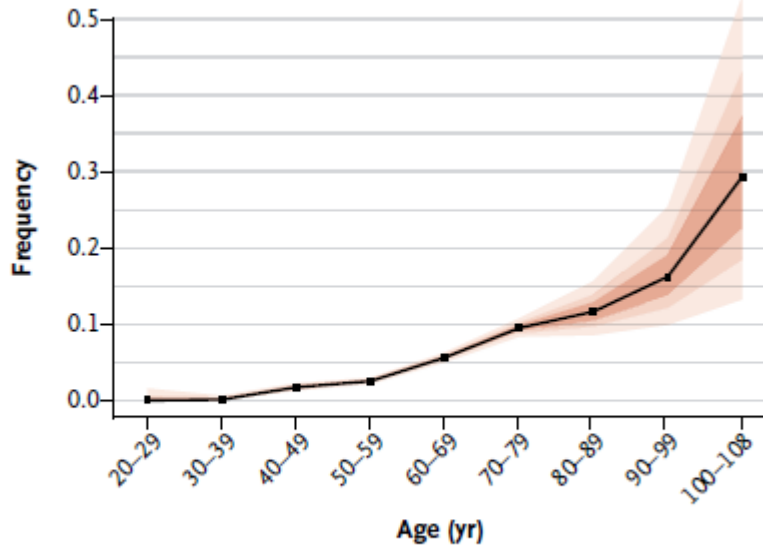


# Clonal architecture of chronic myelomonocytic leukemias

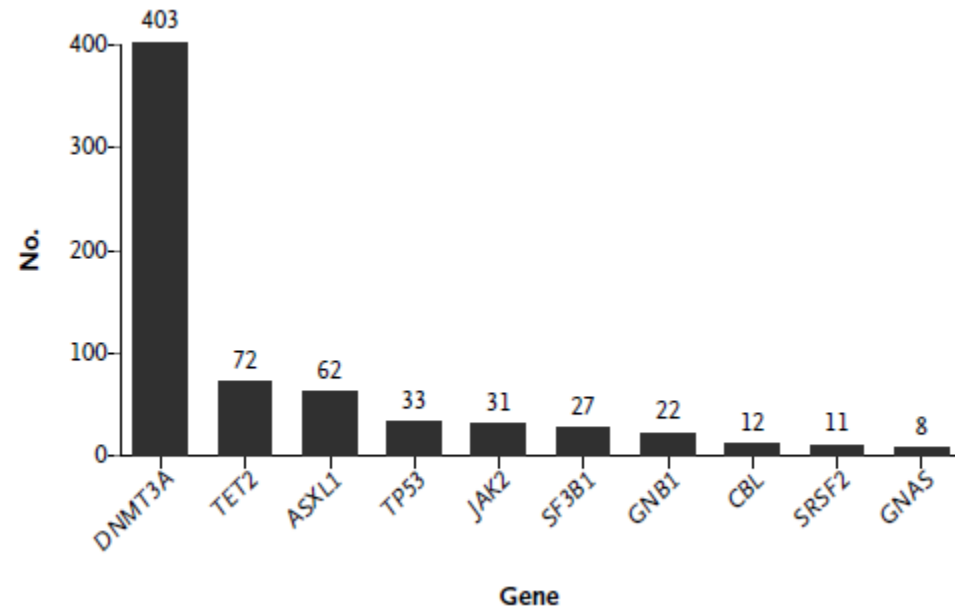


# Age-Related Clonal Hematopoiesis

Prevalence of Somatic Mutations, According to Age.



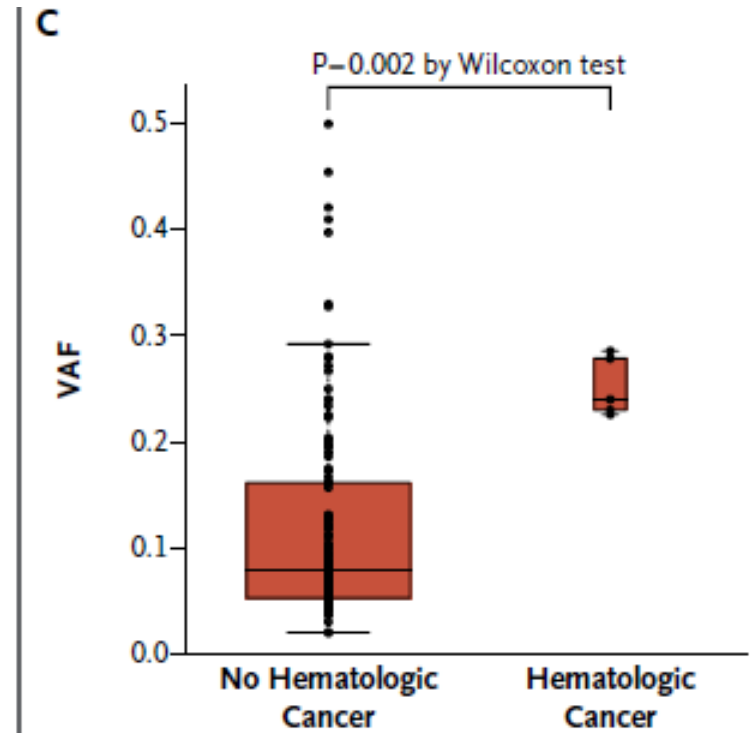
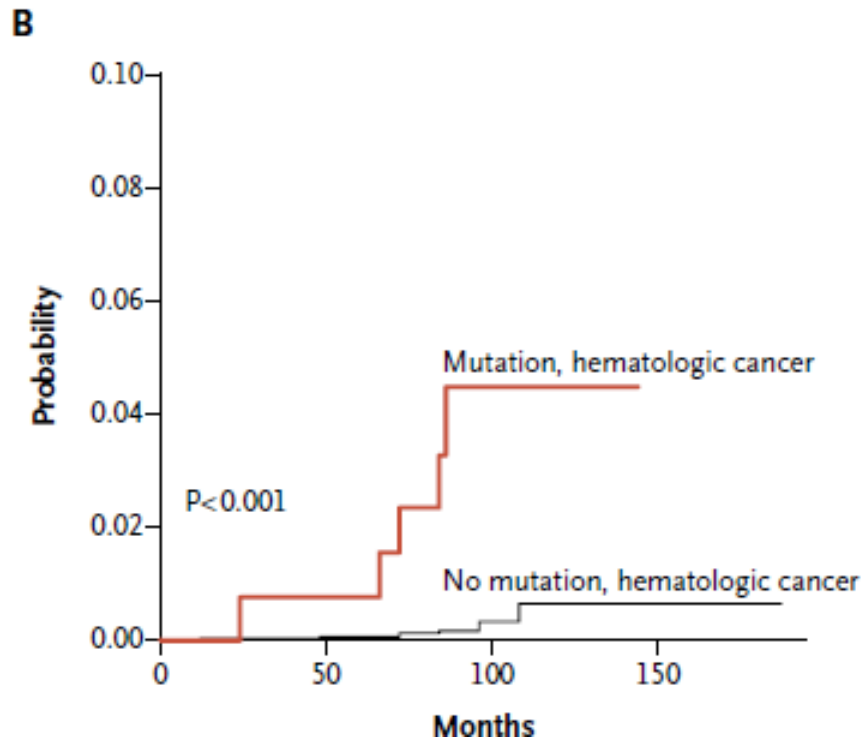
Characteristics of Candidate Somatic Variants.



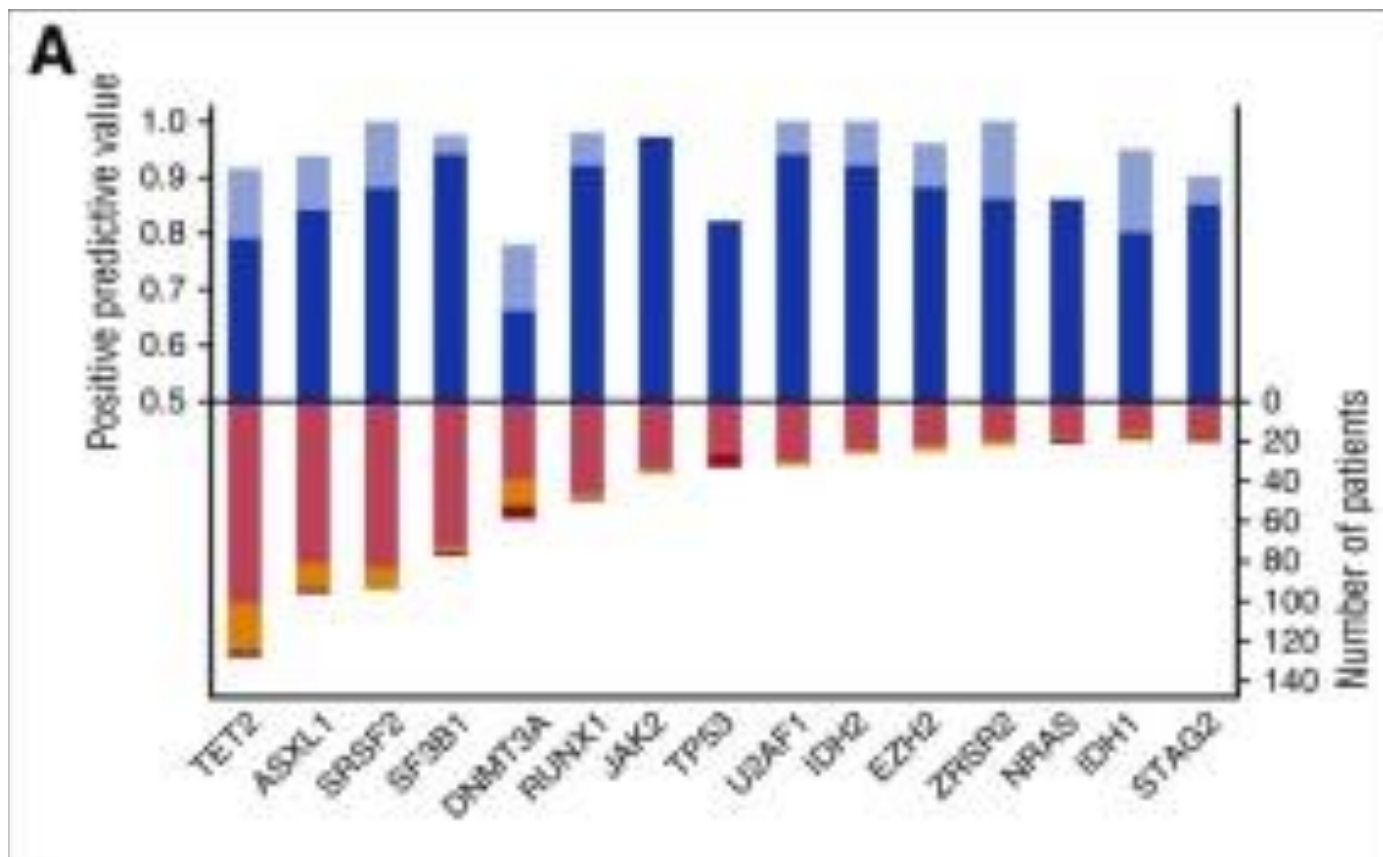
No. with Mutation	0	1	50	138	282	219	37	14	5
Total	240	855	2894	5441	5002	2300	317	86	17



# ARCH is associated with an increase in the risk of hematologic cancer (HR 11.1)



# Clinical significance of somatic mutation in unexplained blood cytopenia



# Clonal Hematopoiesis and Risk of Atherosclerotic Cardiovascular Disease.

- Case control analysis (8000 subjects)
- Carriers of CHIP had a risk of coronary heart disease and infarction that were 1.9 and 4.0 times as great as in non-carriers
- Hypercholesterolemia-prone mice engrafted with BM from homozygous or heterozygous Tet2 knockout mice had larger atherosclerotic lesions
- Macrophages from Tet2 knockout mice showed elevated expression of several chemokine and cytokine genes that contribute to atherosclerosis

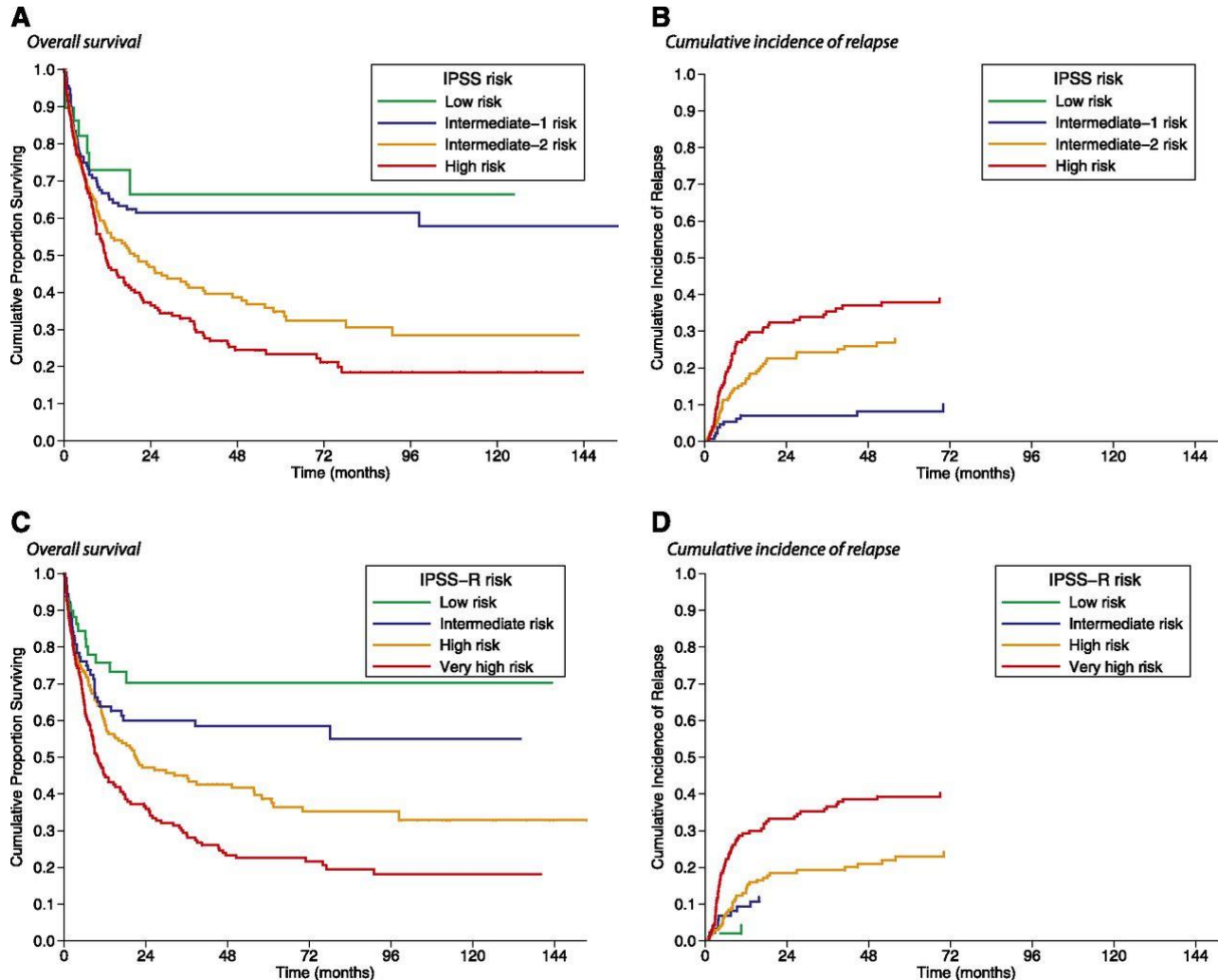
# ARCH - summary

- ARCH is associated with increased risk of hematological malignancies
- In ARCH carriers, type of mutation, VAF, co-mutations are major determinant of the individual risk of developing myeloid cancers
- ARCH is associated with cardiovascular risk (by inducing inflammation)

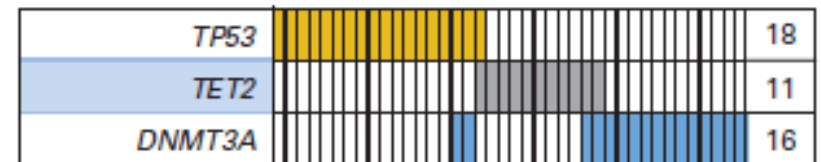
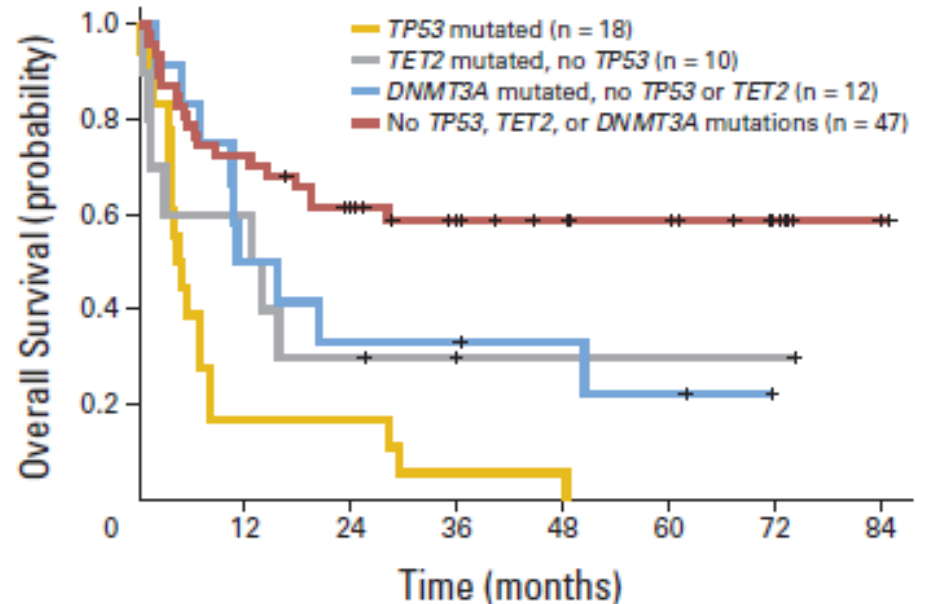
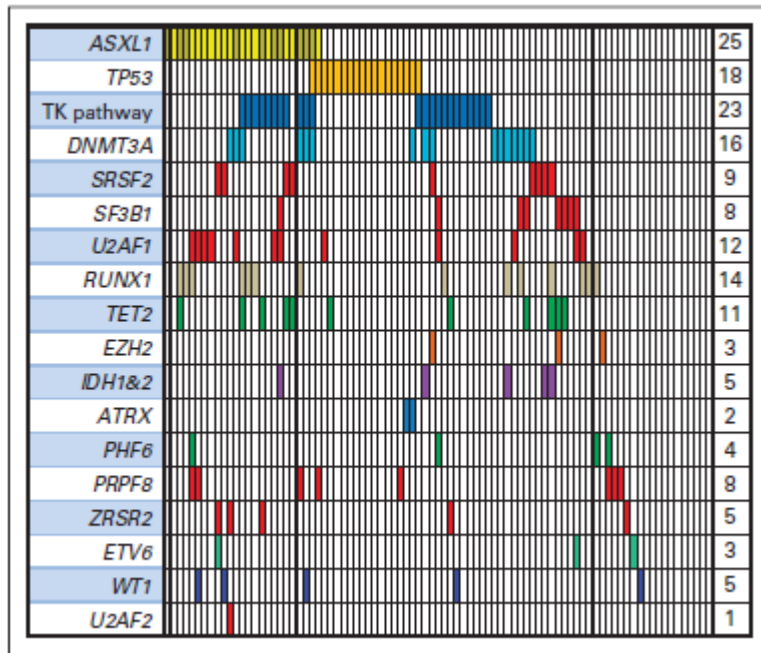
# Outline

- ARCH
- Predictive value of somatic mutations
- Treatment of anemia

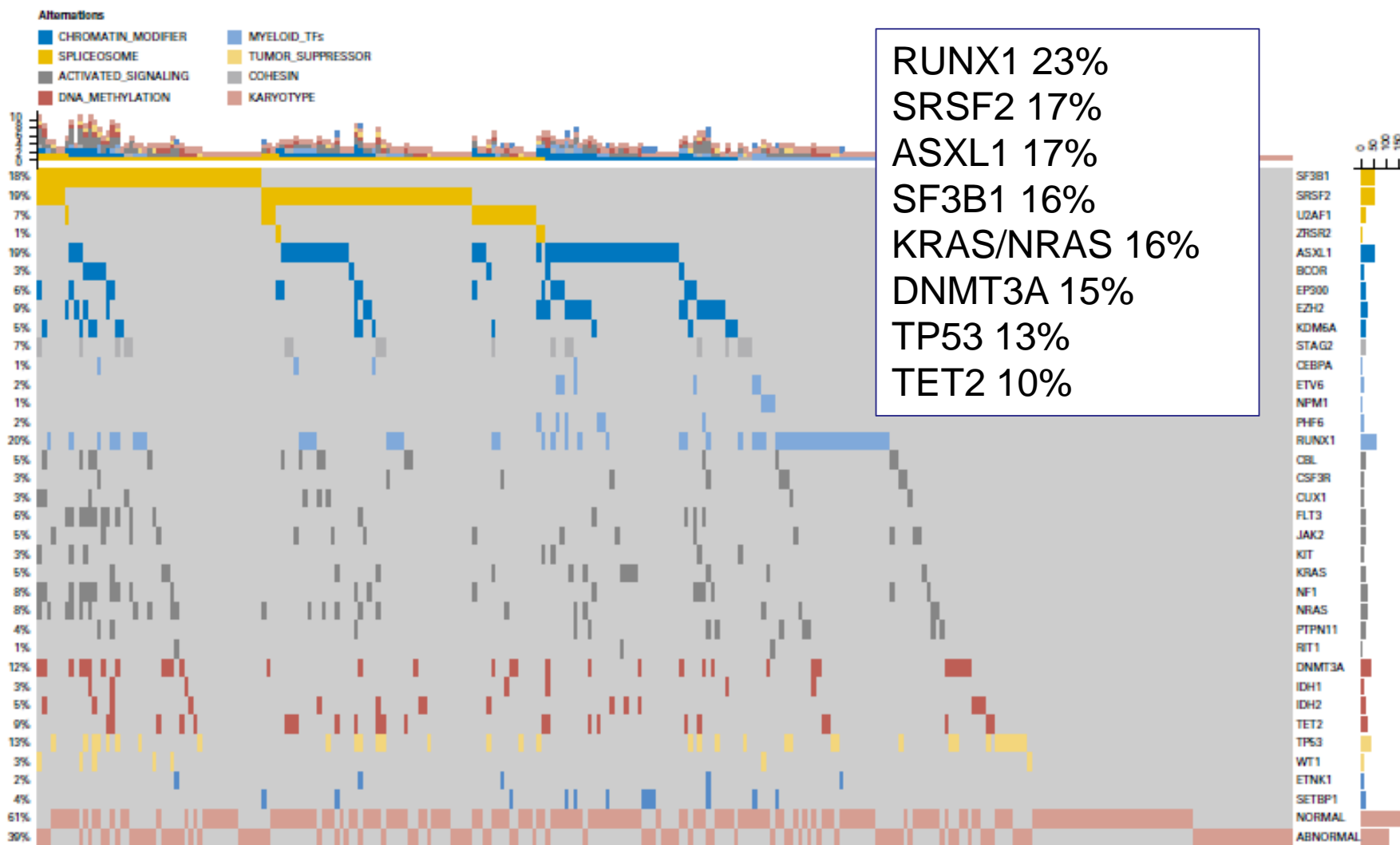
# Kaplan-Meier analysis of survival and cumulative incidence of relapse following allogeneic HSCT in MDS patients stratified according to IPSS or IPSS-R risk.



# Somatic Mutations Predict Poor Outcome in Patients With MDS After Hematopoietic Stem-Cell Transplantation

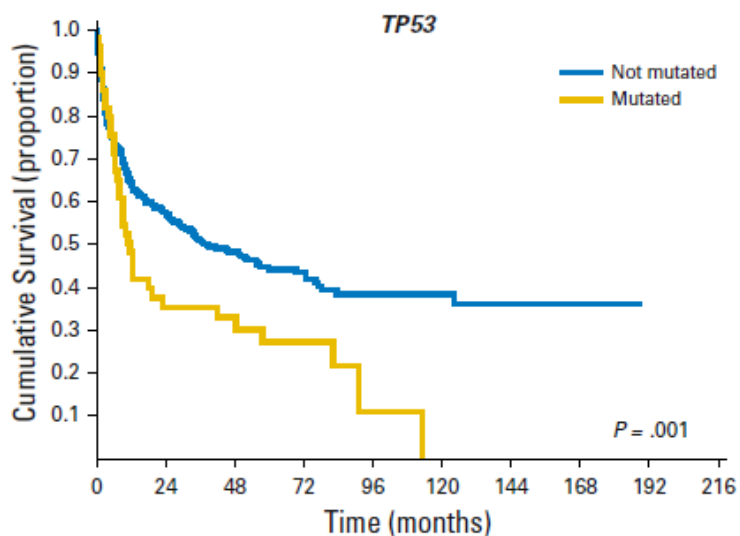
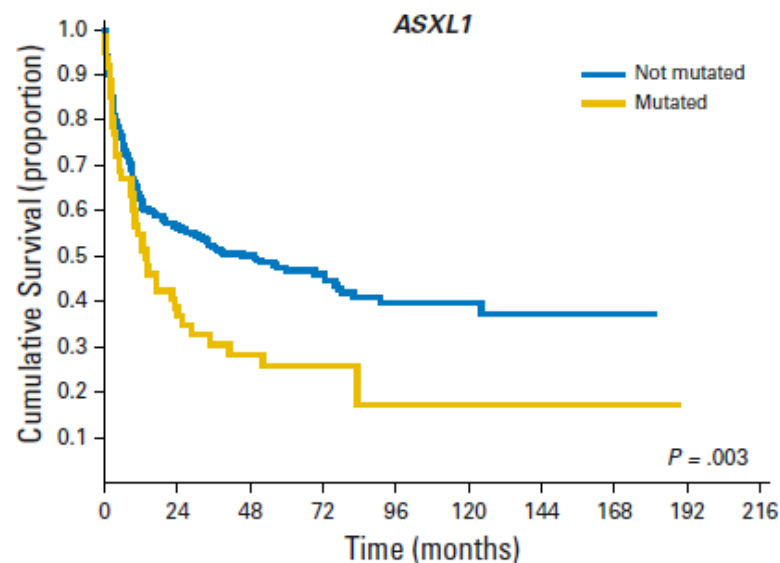
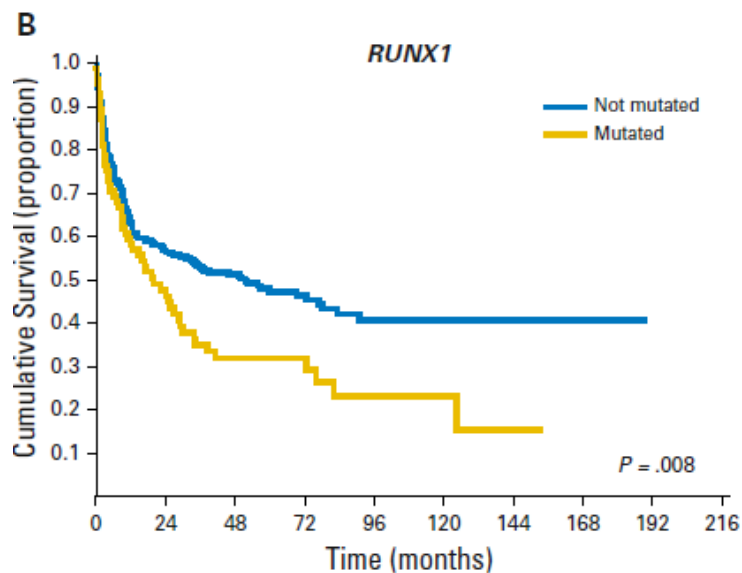


# Mutation patterns observed in MDS treated with allo-HSCT





# Relationship between type of oncogenic mutations and overall survival of MDS receiving allo-HSCT



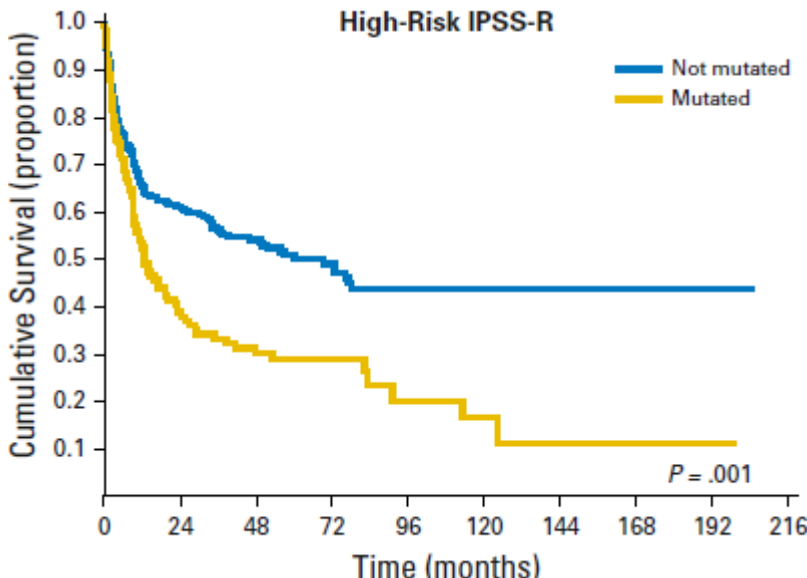
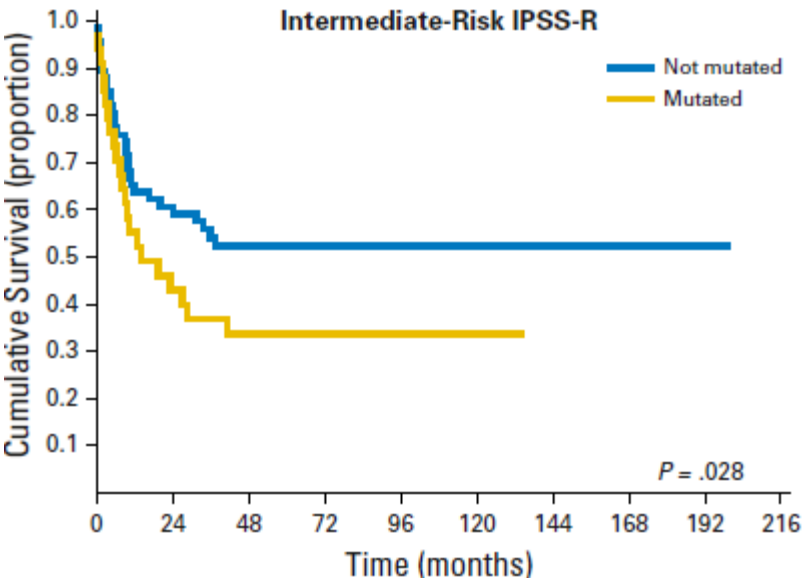
Multivariable analysis				
MDS patients	Probability of relapse		Overall Survival	
Variable	HR	P	HR	P
<i>ASXL1</i>	1.89	.003	1.72	.008
<i>RUNX1</i>	1.67	.02	1.59	.035
<i>TP53</i>	1.90	.019	1.82	.022

# Mutation Pattern at Disease Relapse After HSCT in Patients With MDS and MDS/AML

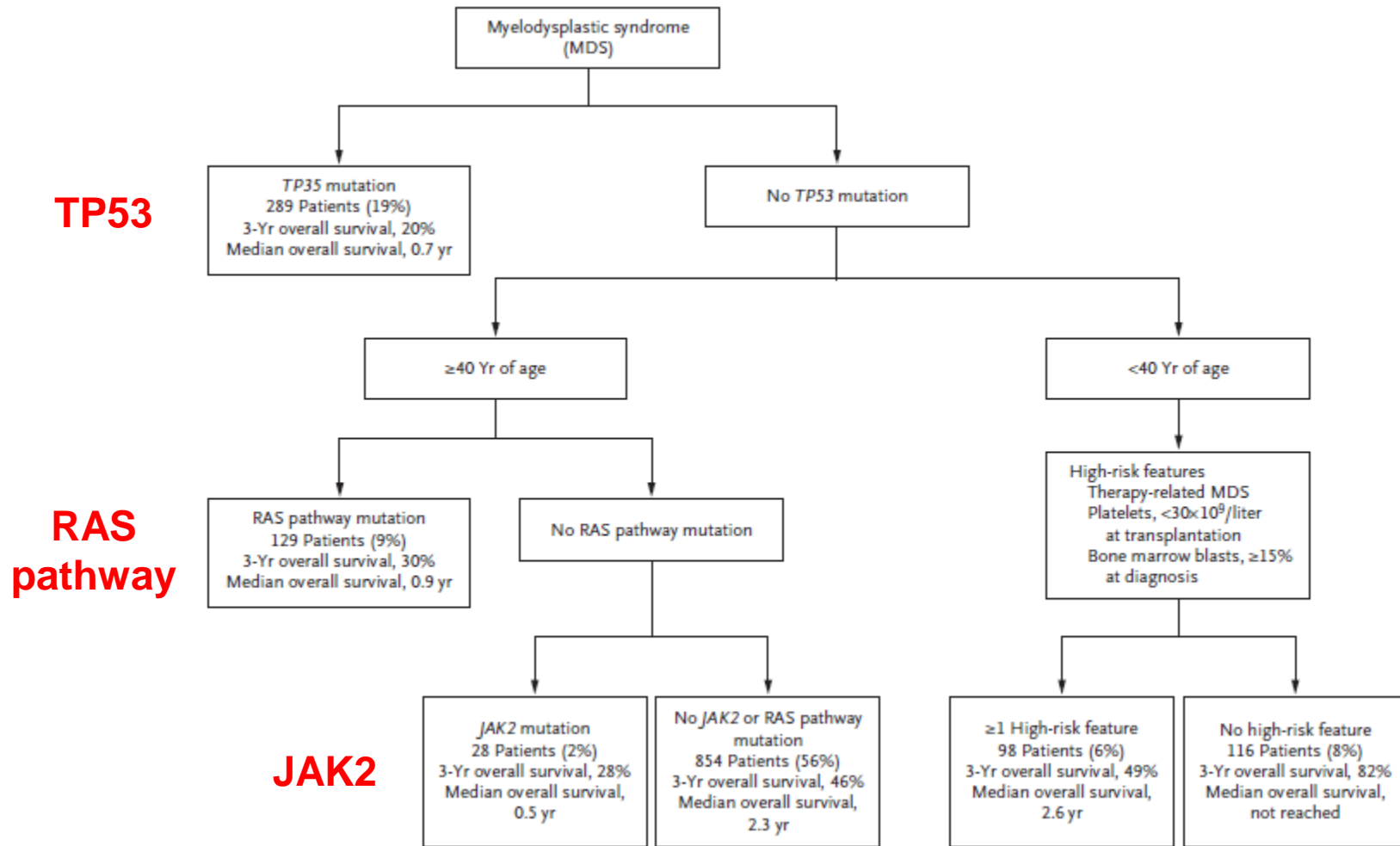


Patient	WHO Category (before HSCT)	Founding Clone (before HSCT)	Clonal Evolution (disease relapse)
GITMO 1	RAEB-2	<i>PTPN11</i>	Founder clone recurs
GITMO 2	MDS/AML	<i>NPM1</i>	Founder clone recurs
GITMO 3	RAEB-1	<i>RUNX1</i>	Founder clone recurs
GITMO 4	RAEB-2	<i>DNMT3A</i>	A subclone expands ( <i>IDH1</i> )
GITMO 5	RAEB-1	<i>STAG2</i>	Founder clone recurs
GITMO 6	MDS/AML	<i>SRSF2</i>	Founder clone recurs
GITMO 7	RAEB-2	<i>EZH2</i>	A subclone expands ( <i>RUNX1</i> )
GITMO 8	RCMD	<i>SRSF2</i>	Founder clone recurs
GITMO 9	RAEB-2	<i>SRSF2</i>	Founder clone recurs

# Clinical Impact of Somatic Mutations in Patients With MDS Receiving HSCT, Stratified According to IPSS-R

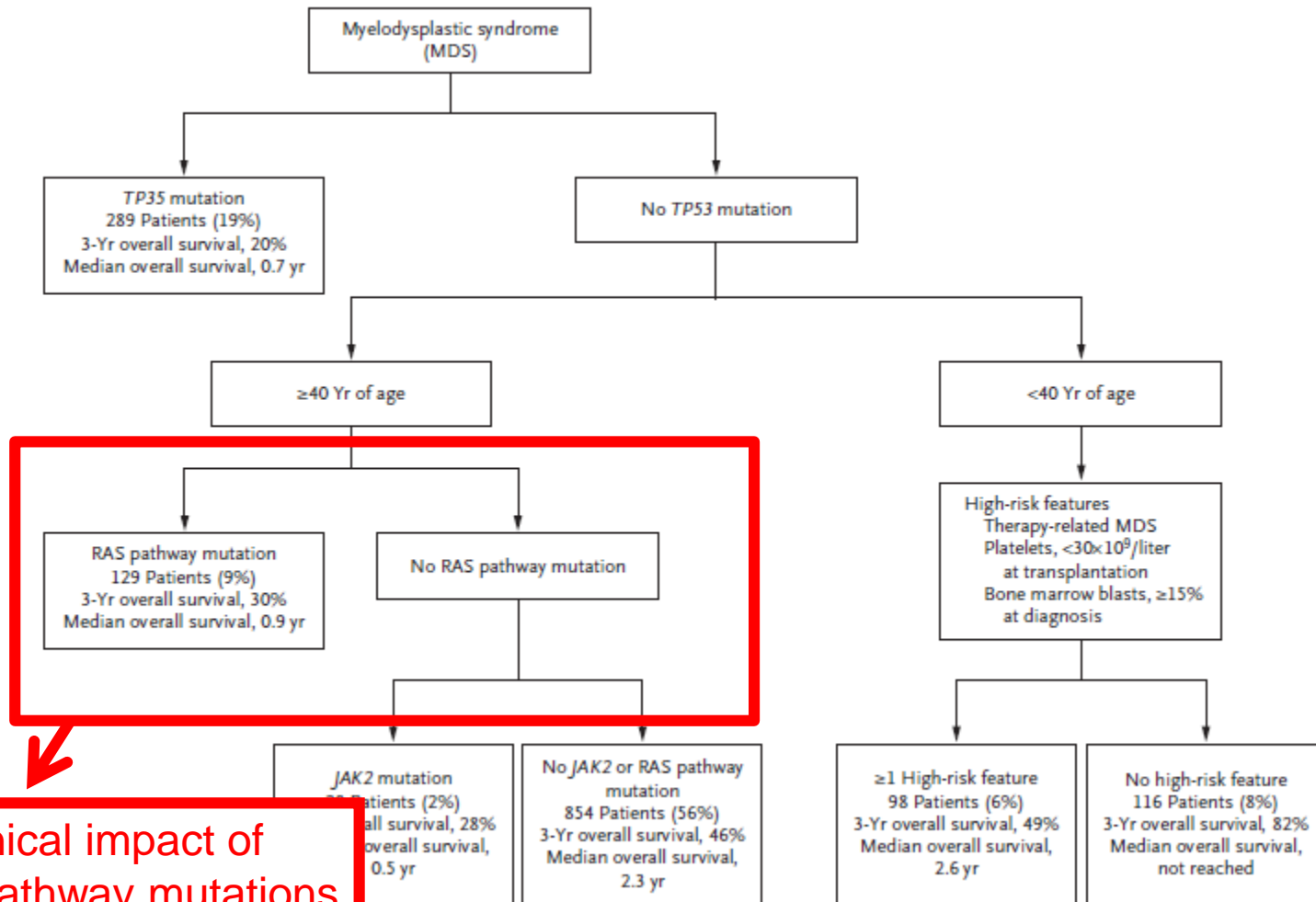


# Prognostic Mutations in Myelodysplastic Syndrome after Stem-Cell Transplantation



Lindsley, RC et al. N Engl J Med 2017;376:536-47.

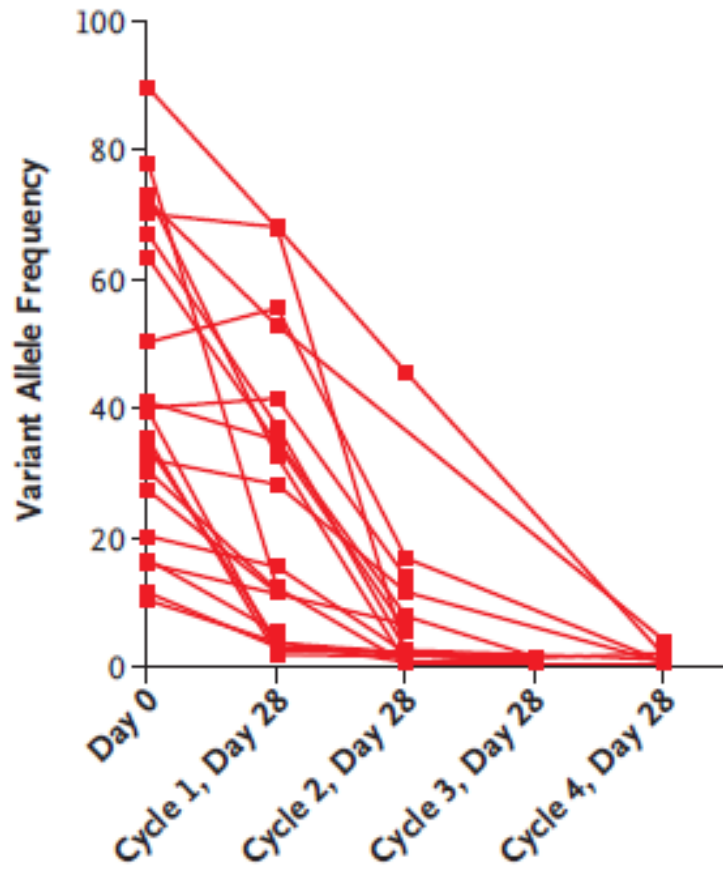
# Prognostic Mutations in Myelodysplastic Syndrome after Stem-Cell Transplantation



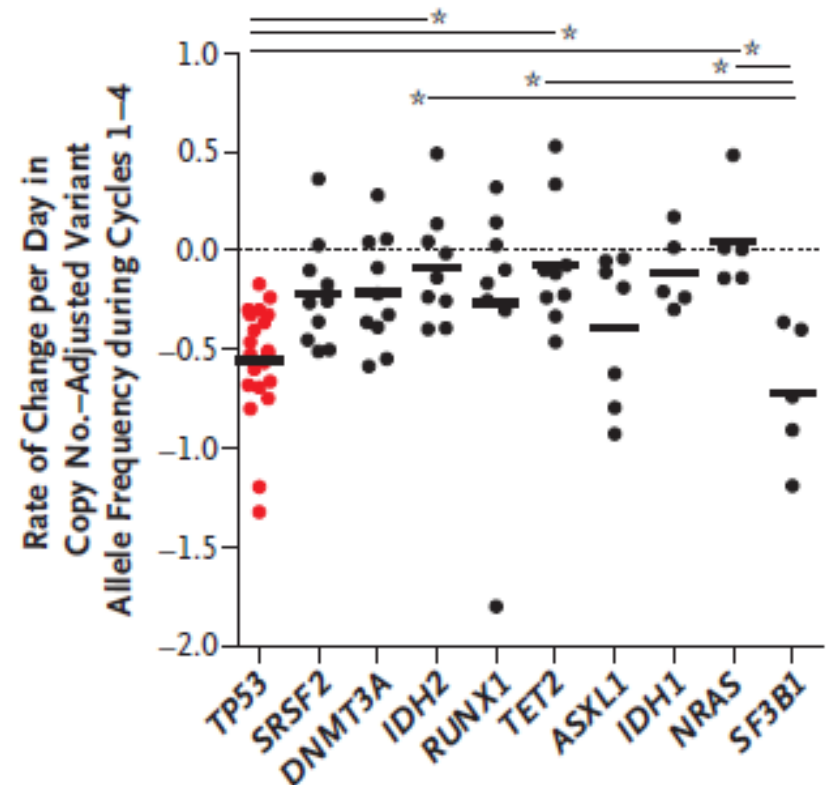
Clinical impact of RAS pathway mutations limited to MDS/MPN

# TP53 and Decitabine in Acute Myeloid Leukemia and Myelodysplastic Syndromes

**D** Clearance of *TP53* Mutations



**E** Clearance of Mutations



# Predictive value of somatic mutations - Summary

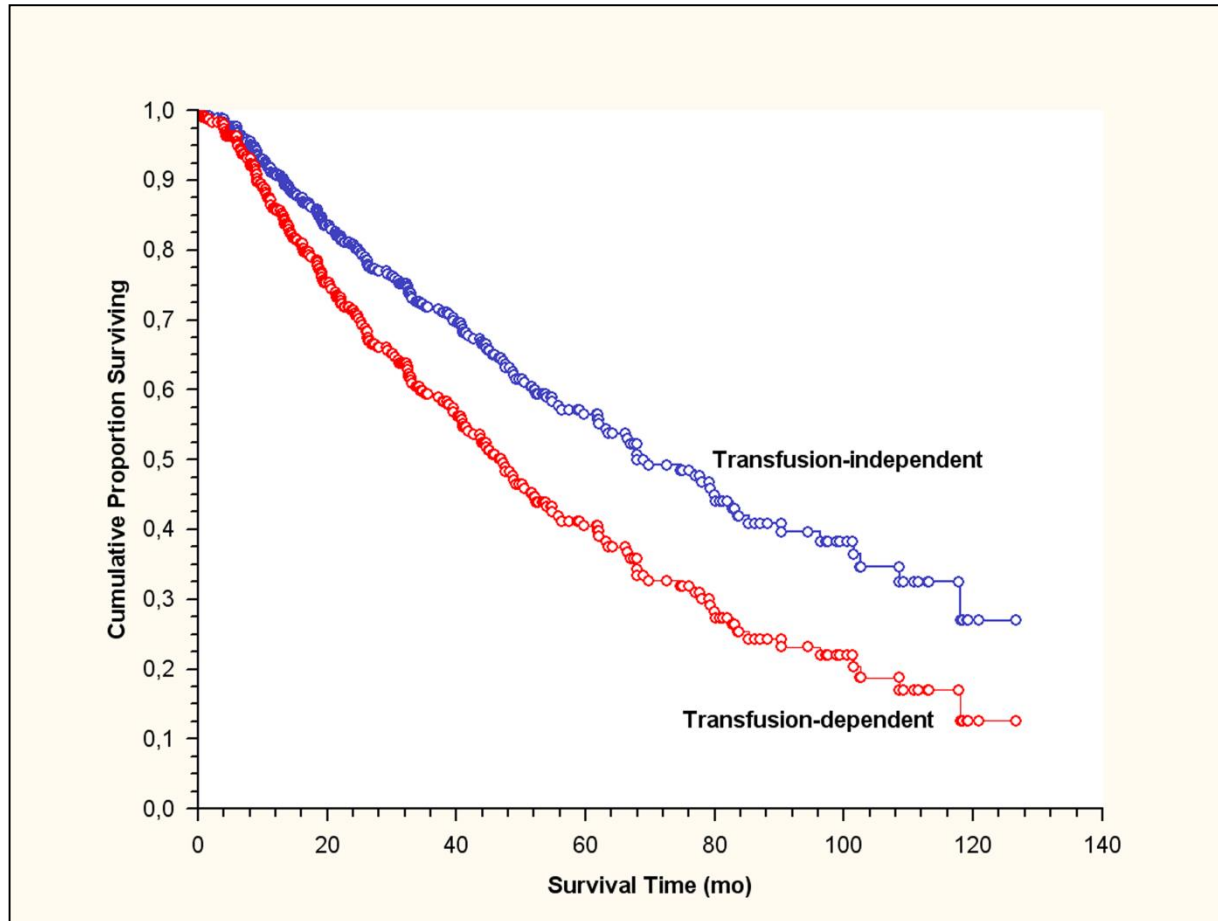
- The definition of molecular basis of MDS is expected to improve diagnosis, prognostic assessment and clinical decision-making
- Mutation screening may affect clinical decision making in transplantation (TP53 mutations are associated to a high probability of disease relapse)
- Molecular biomarkers will be a solid basis for the implementation of personalized medicine programs in hematology

# Outline

- ARCH
- Predictive value of somatic mutations
- Treatment of anemia

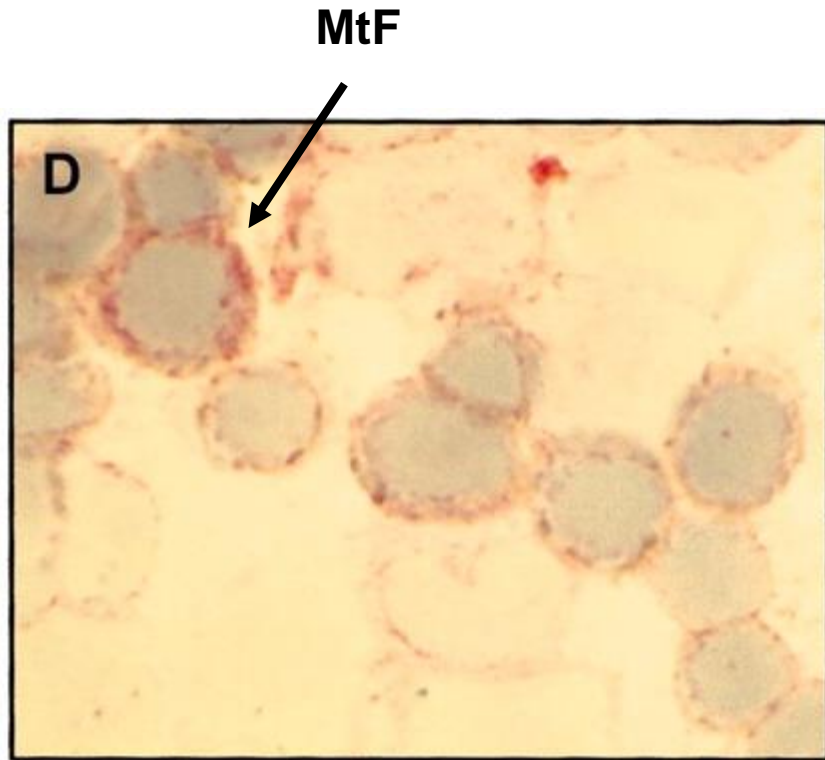


# Survival of MDS patients according to transfusion-dependency



# Refractory Anemia with Ring Sideroblasts

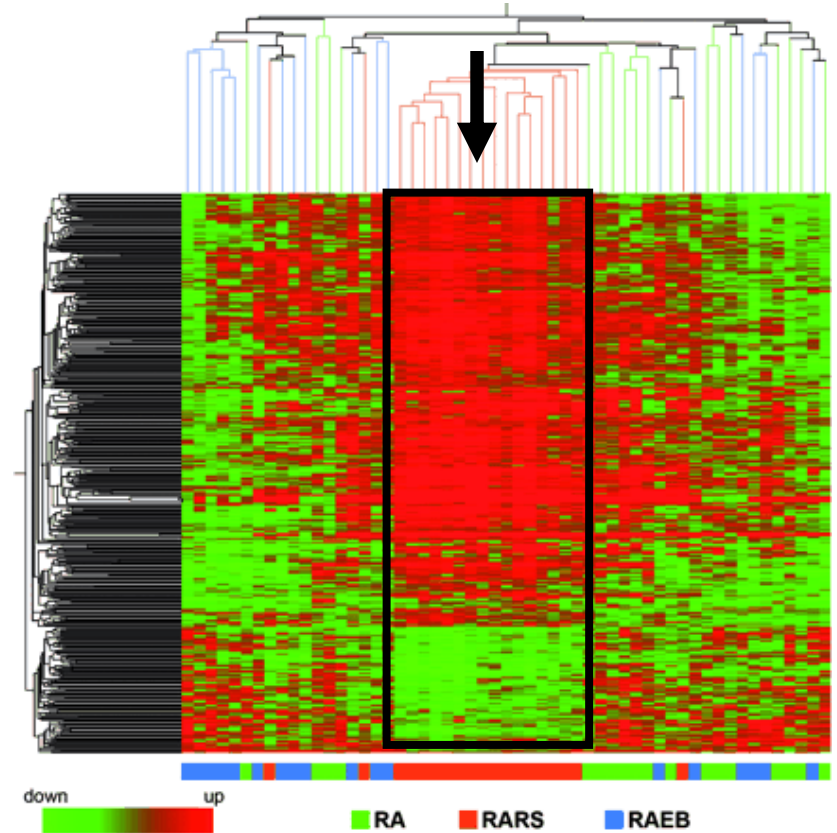
Mitochondrial Ferritin (MtF)



Iron accumulation in ringed sideroblasts is in the form of MtF

*Blood. 2003;101:1996-00*

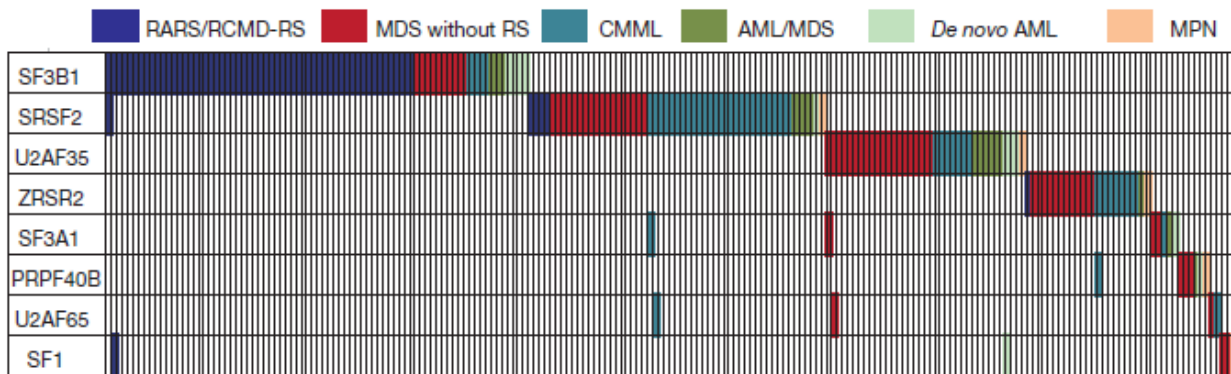
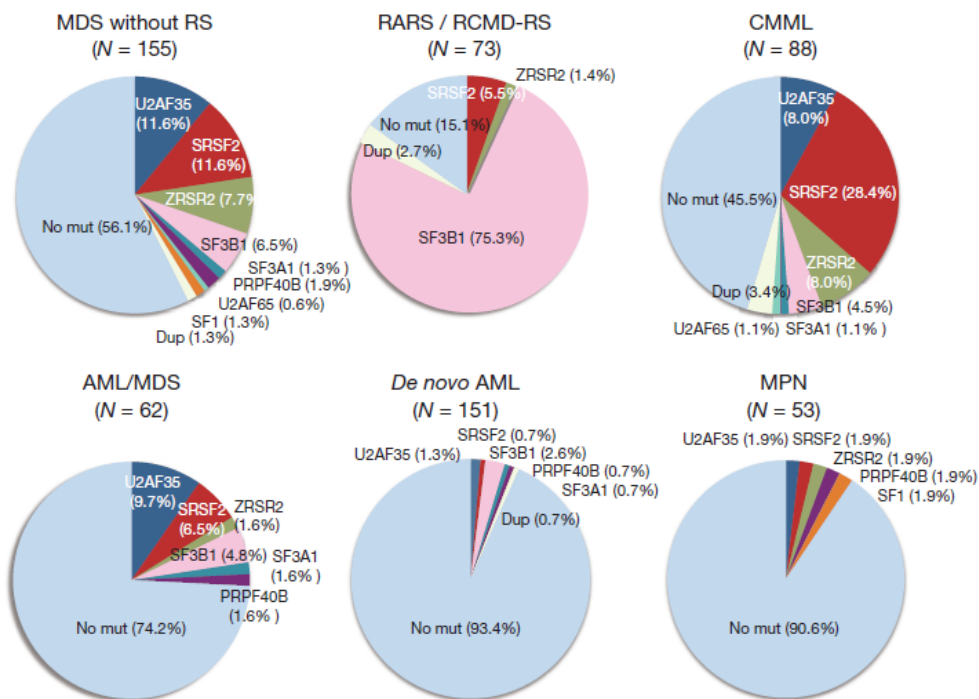
Gene Expression Profile



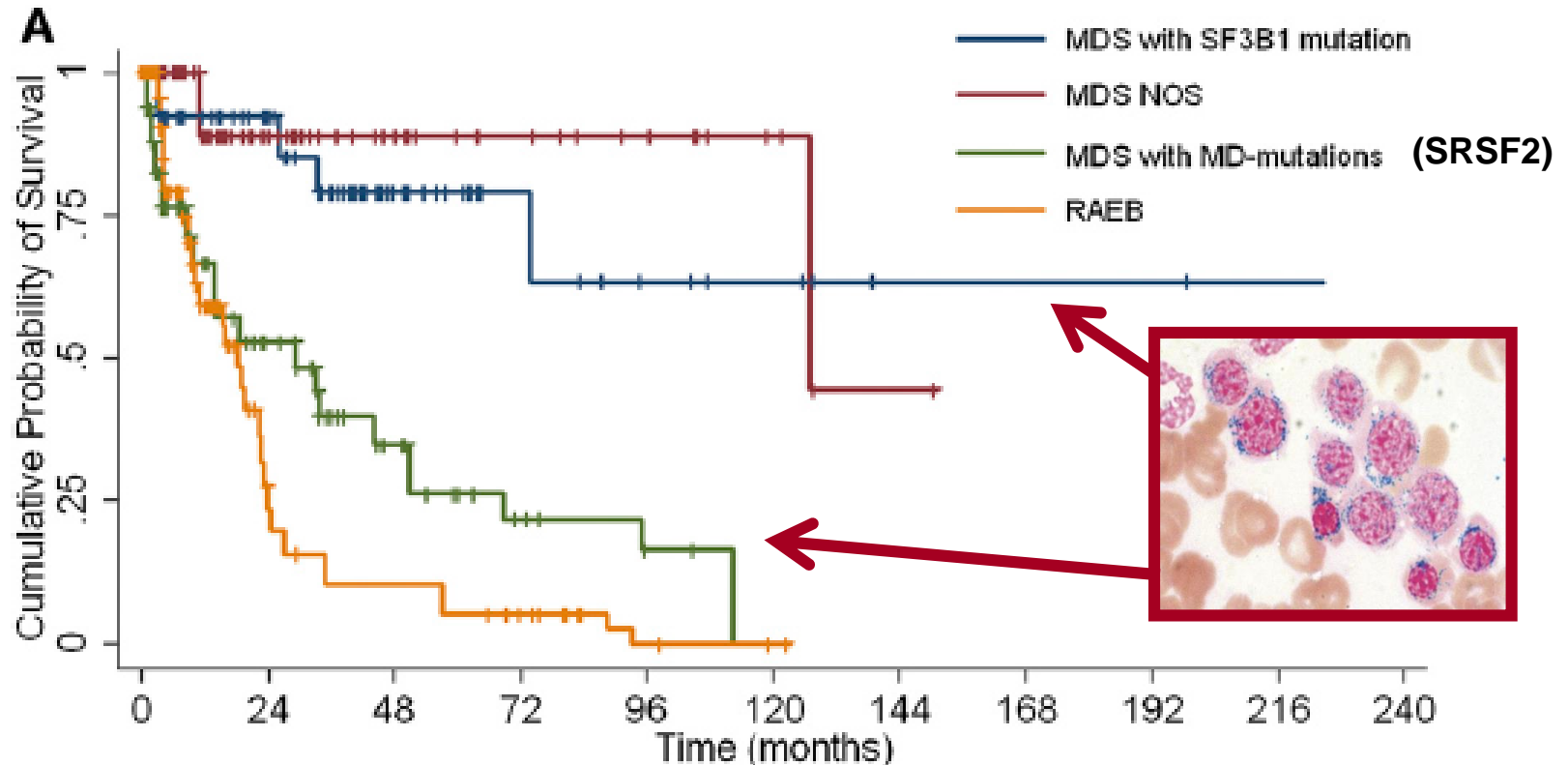
Up-regulation of genes involved in heme synthesis (*ALAS2*)

*Blood. 2006;108:337-45*

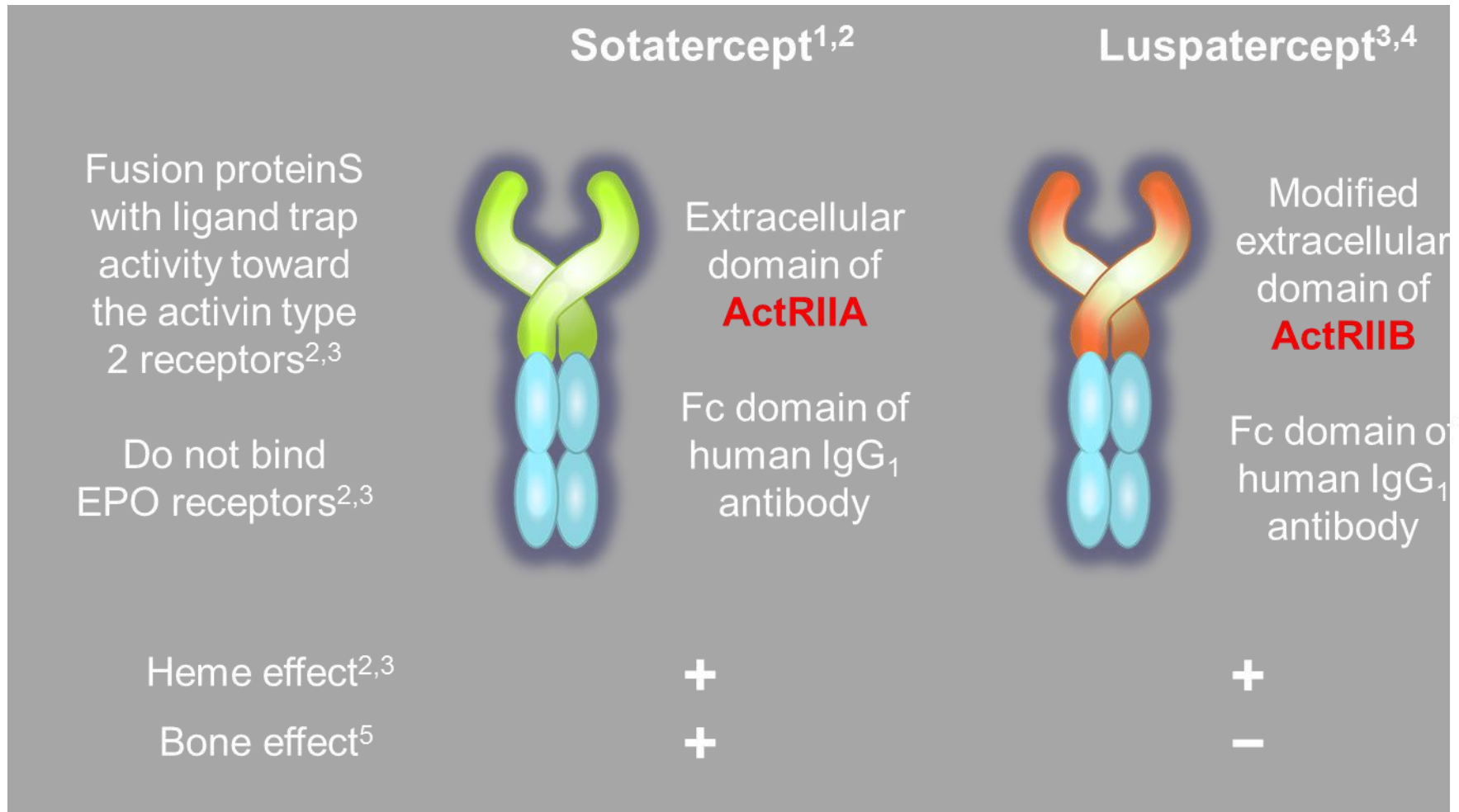
# Frequent pathway mutations of splicing machinery in myelodysplasia



# Driver somatic mutations identify distinct disease entities within myeloid neoplasms with myelodysplasia



# Sotatercept and Luspatercept: Novel Ligand Traps for TGF- $\beta$ Superfamily Ligands



1. Komrokji R, et al. *Blood*. 2014;124(21) [poster presentation; abstract 3251]. 2. Carrancio S, et al. *Br J Haematol*. 2014;165(6):870-882. 3. Suragani R, et al. *Nat Med*. 2014;20(4):408-414. 4. Platzbecker U, et al. *Blood*. 2014;124(21) [oral presentation; abstract 411]. 5. Iancu-Rubin C, et al. *Exp Hematol*. 2013;41(12):155-166.e17.

# Rationale for Luspatercept in Anemia

- SMAD2/3 is constitutively activated in the hematopoietic progenitors, resulting in ineffective erythropoiesis
- In preclinical murine models, luspatercept
  - Promoted maturation of late-stage erythroid precursors in vivo
  - Increased RBC, hematocrit, and Hb levels in a dose-dependent manner
- RAP-536, a murine version of luspatercept, prevented or reduced anemia in different murine anemia models, including MDS and  $\beta$ -thalassemia
- In a phase I clinical trial in healthy post-menopausal women
  - Luspatercept stimulated RBC production and increased Hb levels at effective dose levels

# Higher response rates were observed in patients with RS, lower EPO levels, and SF mutations

Subgroup n (%)	IWG HI-E Response Rate	RBC-TI Response Rate
<b>All</b>	24 of 49 (49)	14 of 40 (35)
RS+	22 of 40 (55)	12 of 31 (39)
RS-	2 of 7 (29)	2 of 7 (29)
SF3B1 mutation	18 of 30 (60)	9 of 24 (38)
Any SF mutation	20 of 36 (58)	13 of 29 (45)
EPO < 200 U/L	16 of 25 (64)	10 of 18 (56)
EPO 200–500 U/L	4 of 11 (36)	3 of 9 (33)
EPO > 500 U/L	4 of 13 (31)	1 of 13 (8)
Prior ESA	16 of 35 (46)	10 of 29 (35)
ESA naïve	8 of 14 (57)	4 of 11 (36)

EPO, erythropoietin; ESA, erythropoietin stimulating agent ; RS, ring sideroblasts; SF, splicing factor; SF3B1, Splicing Factor 3b, Subunit 1.

# Treatment of anemia - Summary

- The definition of molecular basis of MDS is expected to improve diagnosis, prognostic assessment and clinical decision-making
- Luspatercept is an effective treatment options for patients with anemia due to ineffective erythropoiesis, especially in those with SF3B1 mutation
- Molecular biomarkers will be a solid basis for the implementation of personalized medicine programs in hematology



# Spunti discussione

- Early detection: Implicazioni cliniche
- Implementazione clinica dell'uso delle mutazioni somatiche per la diagnosi, il prognostic assessment e la decisione terapeutica a livello individuale
- Sviluppo di programmi di medicina personalizzata nelle MDS