


Unmet challenges in high risk
hematological malignancies:
from benchside to clinical practice

HUMANITAS
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How I treat high-risk MDS

Matteo G Della Porta

Cancer Center

IRCCS Humanitas Research Hospital

& Humanitas University

Milano, Italy

matteo.della_porta@hunimed.eu

Clinical decision making in MDS - Critical issues

- How we can define HIGH-RISK MDS?

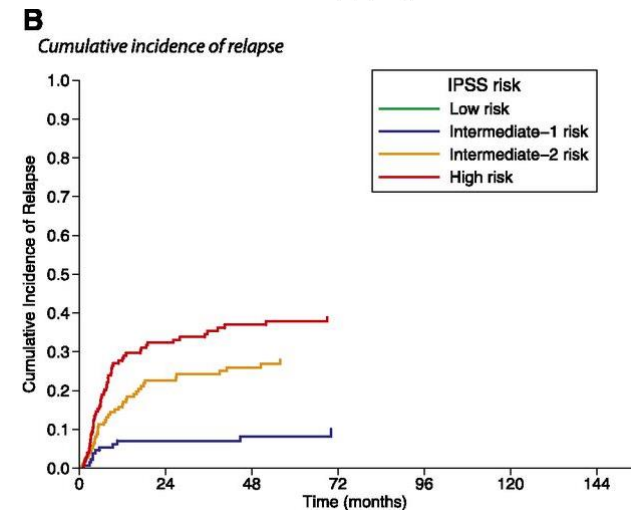
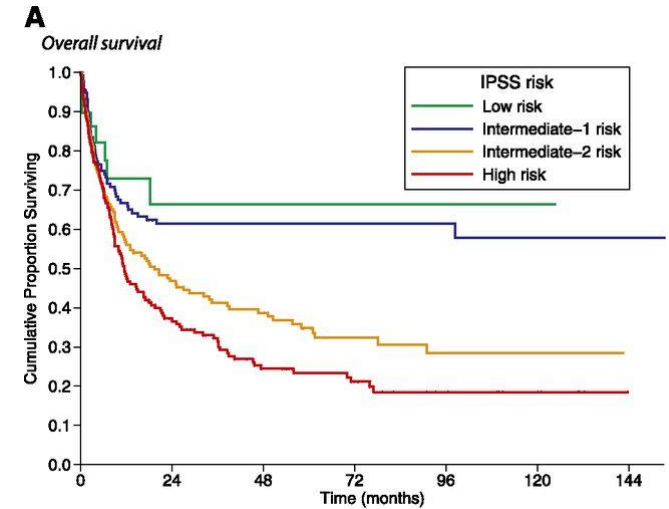
International Prognostic Scoring System (IPSS) for MDS

Variable	0	0.5	1	1.5	2
BM blasts %	<5	5-10	-	11-20	21-30
Karyotype*	Good	Intermediate	Poor		
Cytopenias°	0/1	2/3			

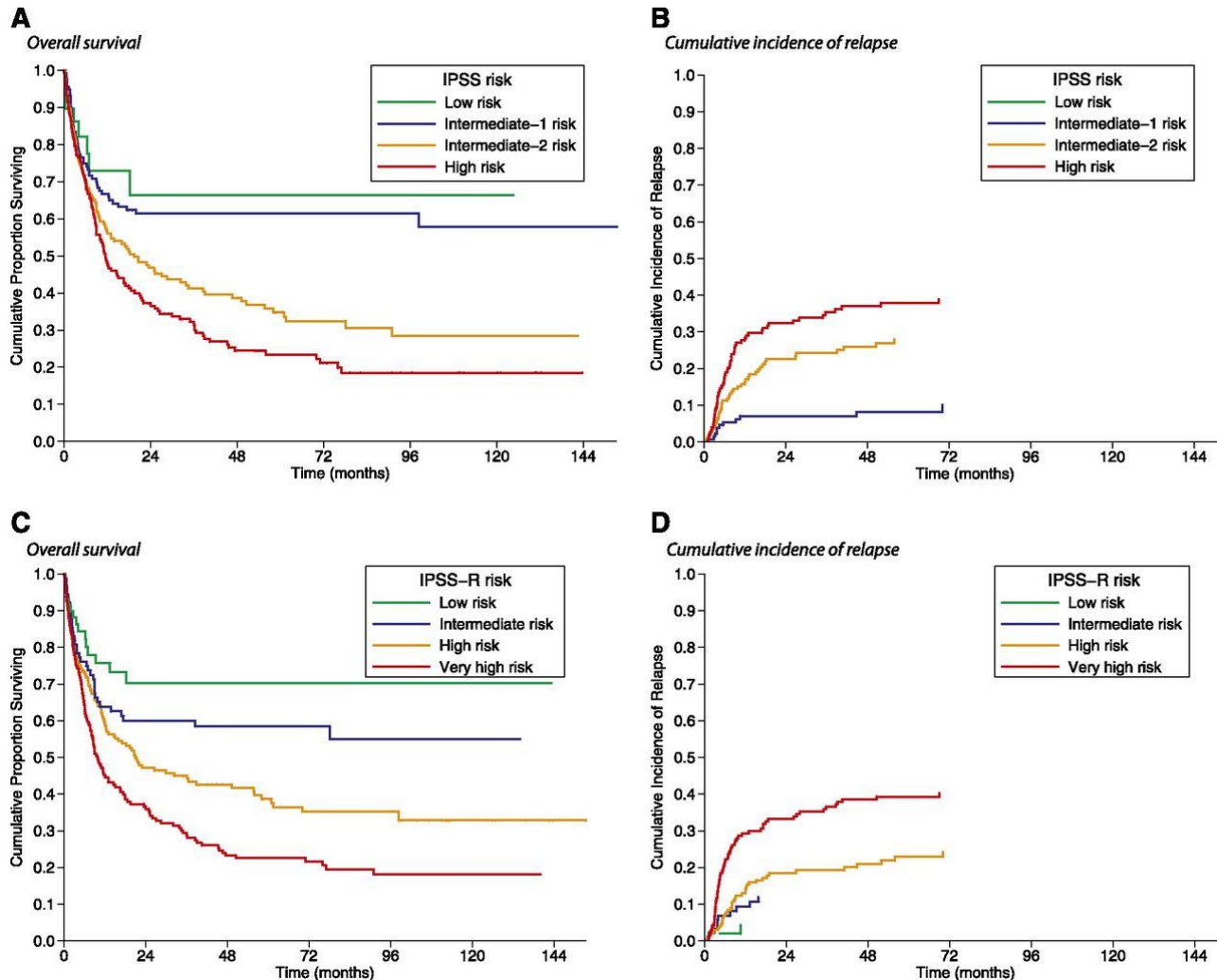
* *Good*: normal, -Y, del(5q), del(20q); *Poor*: complex, chromosome 7 anomalies; *Intermediate*: other abnormalities.

° Hemoglobin < 10 g/dL, absolute neutrophil count < 1,500/ μ L, platelet count < 100,000/ μ L.

Scores for risk groups are as follows: Low, 0; INT-1, 0.5-1.0; INT-2, 1.5-2.0; and High, 2.



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



Della Porta MG et al. *Blood* 2014;123:2333-2342
Della Porta MG et al. *Leukemia*. 2015 ;29:1502-13.

ASH 2017 - Somatic Mutations in MDS Predict Prognosis Independent of the IPSS-R (Analysis by IWG-PM)

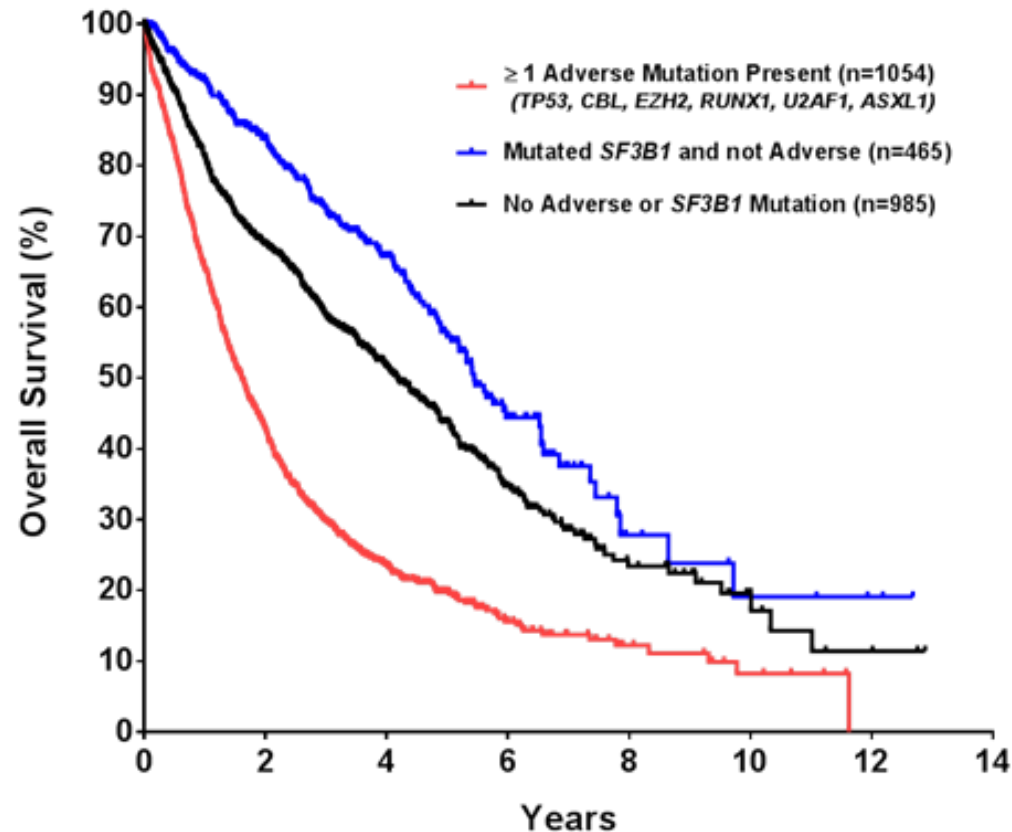


Figure 2: Kaplan-Meier curve of overall survival in years for the 2504 patients with sequence results for *SF3B1* and all six adverse genes (*TP53*, *CBL*, *EZH2*, *RUNX1*, *U2AF1*, and *ASXL1*).

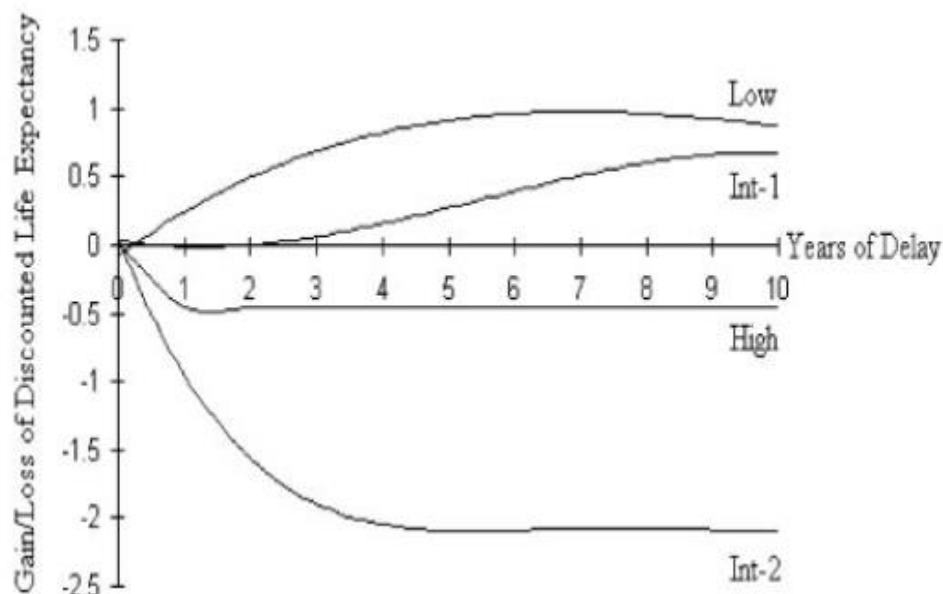
Transplantation decision making in MDS - Critical issues

- Which tools are available for transplant decision making?
- How we can define optimal timing of transplantation in individual patient?
- What is the clinical relevance of somatic mutations in transplantation decision making in MDS?

Transplantation strategy according to IPSS

Discounted life expectancy, in years, for alternative transplantation strategies

Patients, by IPSS risk group	Transplantation at diagnosis	Transplantation at AML progression
All patients		
Low	6.51	7.21
Int-1	4.61	5.16
Int-2	4.93*	2.84
High	3.20*	2.75
Patients younger than 40 y		
Low	5.62	10.21*
Int-1	2.48	10.21*
Int-2	1.65*	1.53
High	—	—



Transplantation policy according to IPSS-R

Patient AGE

	<i>delay time (months)</i>	<i>40</i>	<i>50-55</i>	<i>>60</i>
Years of life expectancy under policy 1: IPSS-R Low	0	16.4	16.1	15.1
	12	17.3	16.8	15.4
	24	17.9	17.3	15.6
	48	18.5	17.7	15.7
	60	18.7	17.9	15.7
Years of life expectancy under policy 2: IPSS-R intermediate	0	19.3	18.1	15.9
	12	17.9	17.1	14.9
	24	17.1	16.4	14.5
	48	16.3	15.7	14.2
	60	16.0	15.5	13.9

Optimal timing of alloSCT

gain of life expectancy:

- 5.3 y pts <50y
- 4.7 y pts 60 y
- 2.8 y pts 65 y

Transplantation policy according to IPSS vs. IPSS-R

	<i>IPSS-based policy*</i>	<i>IPSS-R</i>	<i>%</i>	<i>IPSS-R based policy **</i>
IPSS Low	<i>Delayed</i>	Very low	37	<i>Delayed</i>
		Low	50	<i>Delayed</i>
		Intermediate	13	<i>Immediate</i>
		High	-	
IPSS Intermediate-1	<i>Delayed</i>	Very low / Low	48	<i>Delayed</i>
		Intermediate	40	<i>Immediate</i>
		High	11	<i>Immediate</i>
		Very high	1	<i>immediate</i>

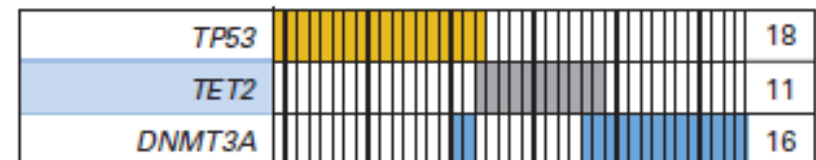
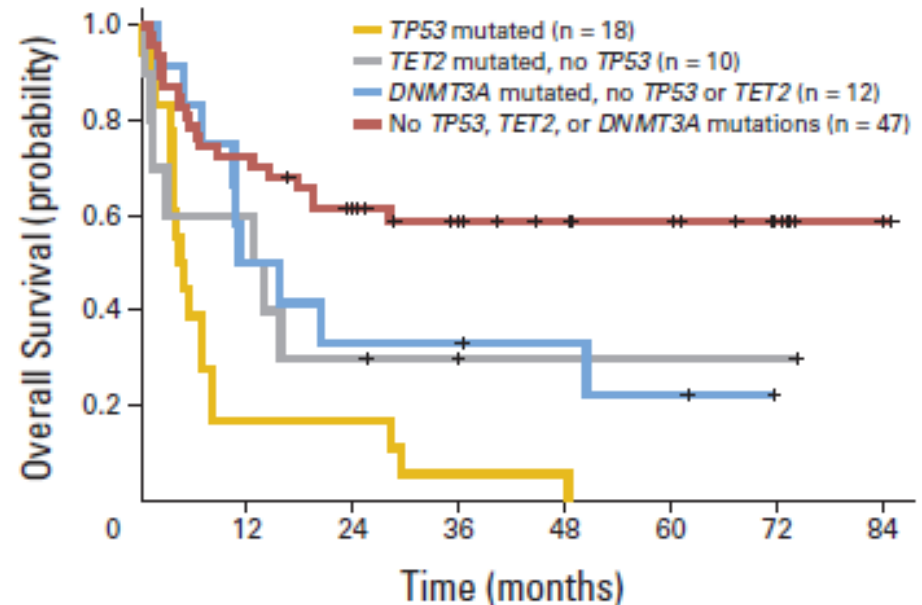
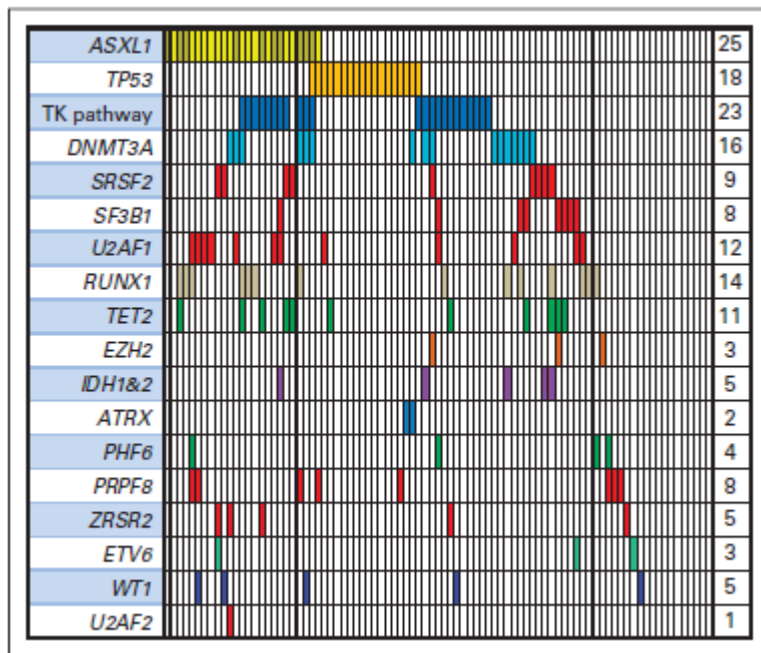
* Cutler CS et al. Blood 2004;104(2):579-85.

** Della Porta MG et al. Leukemia. 2017 Apr 7. doi: 10.1038/leu.2017.88

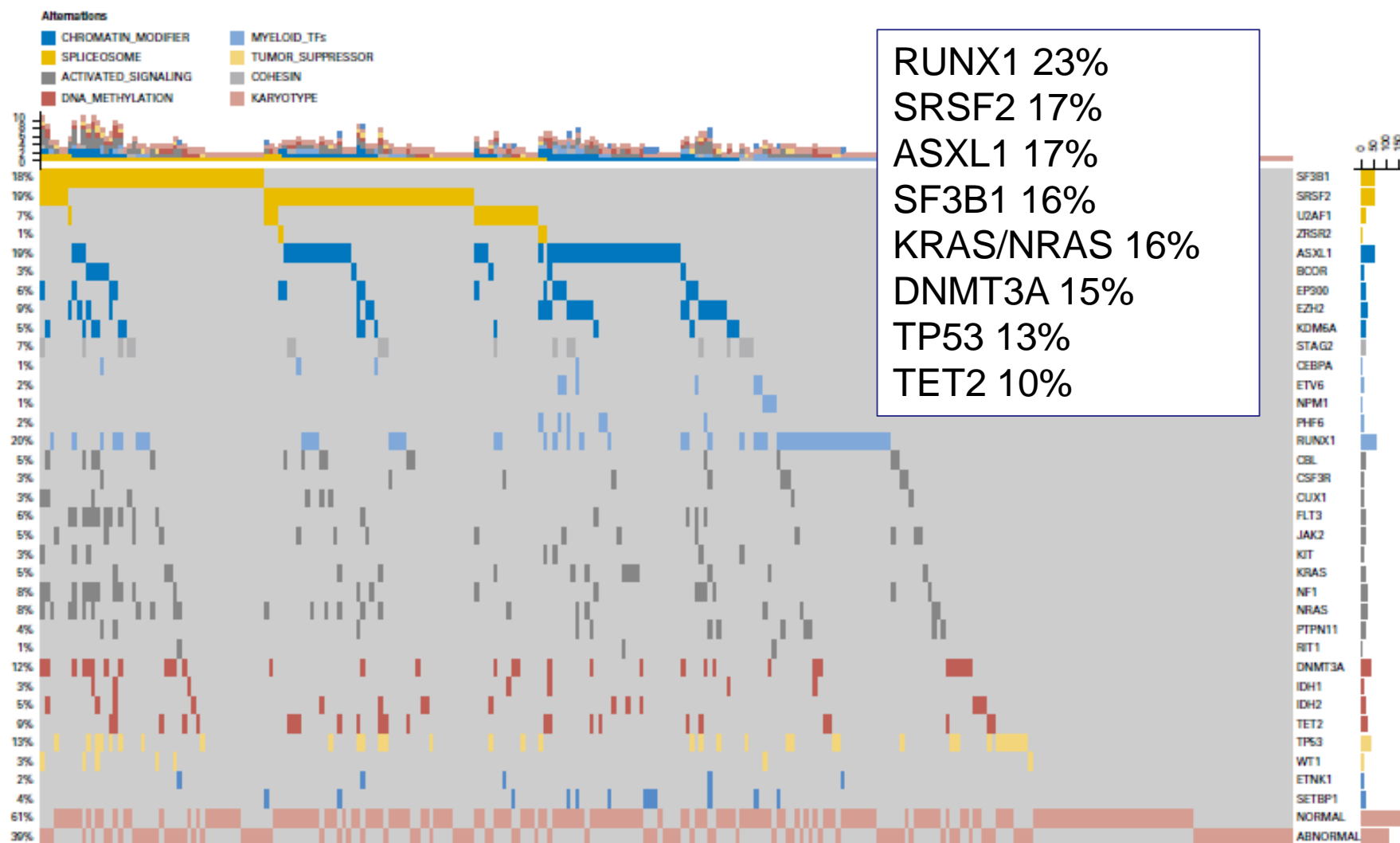
Transplantation decision making in MDS - Critical issues

- What is the clinical relevance of somatic mutations in transplantation decision making in MDS?

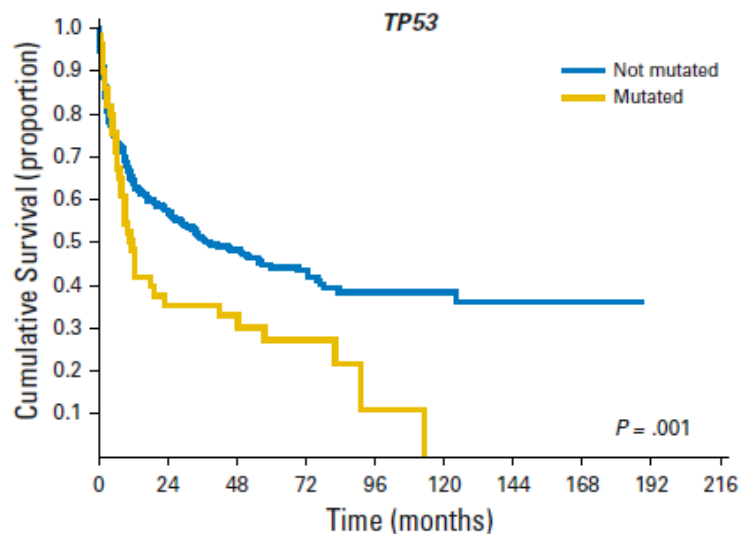
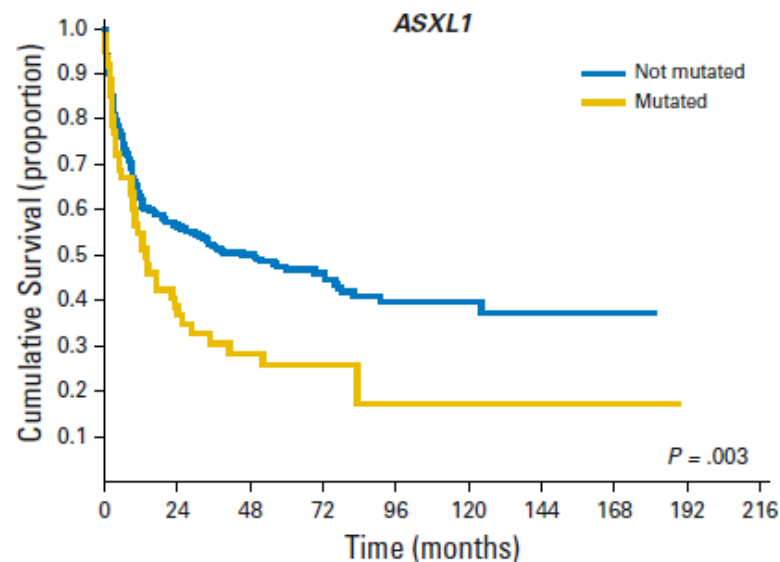
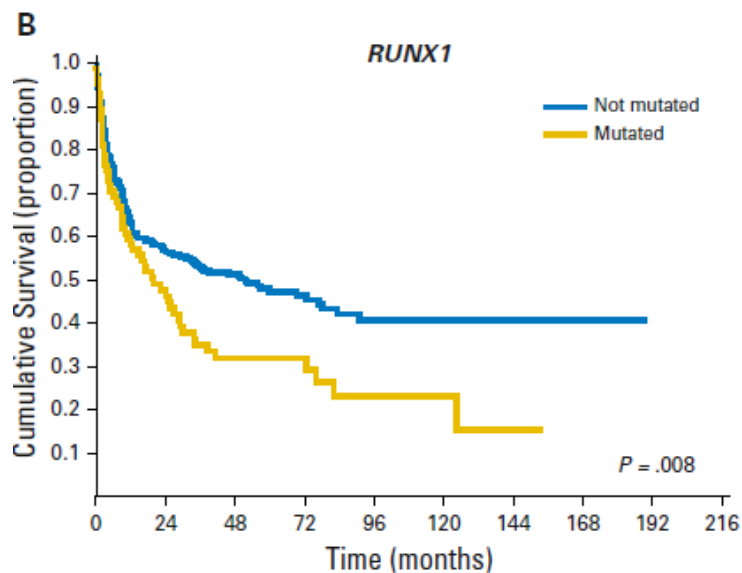
Somatic Mutations Predict Poor Outcome in Patients With MDS After Hematopoietic Stem-Cell Transplantation



Mutation patterns observed in MDS treated with allo-HSCT

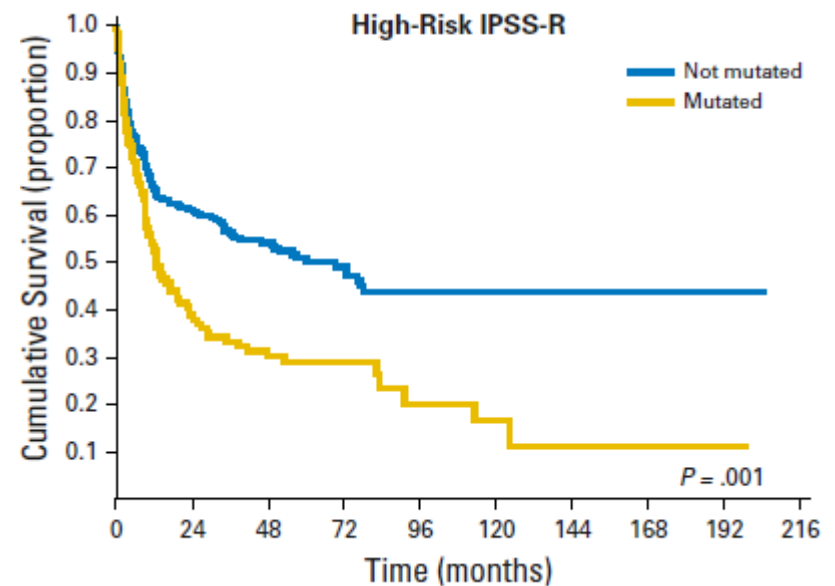
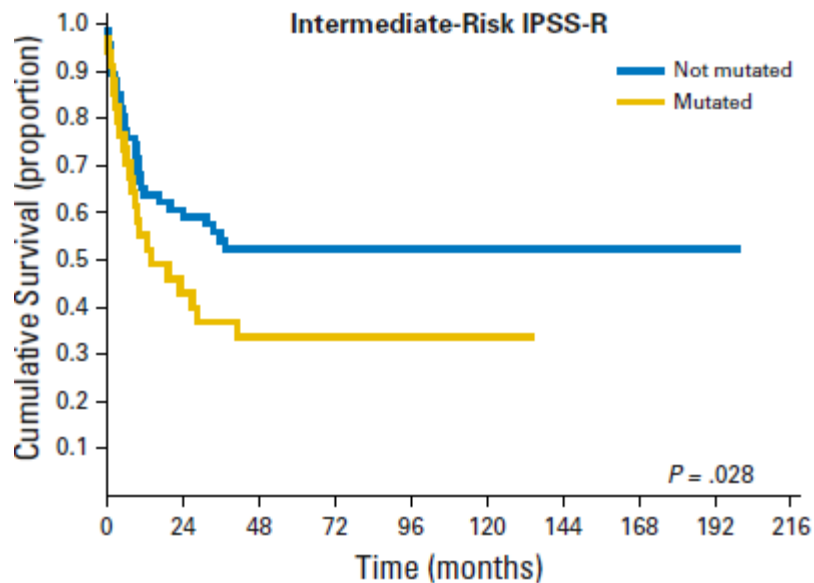


Relationship between type of oncogenic mutations and overall survival of MDS receiving allo-HSCT



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

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

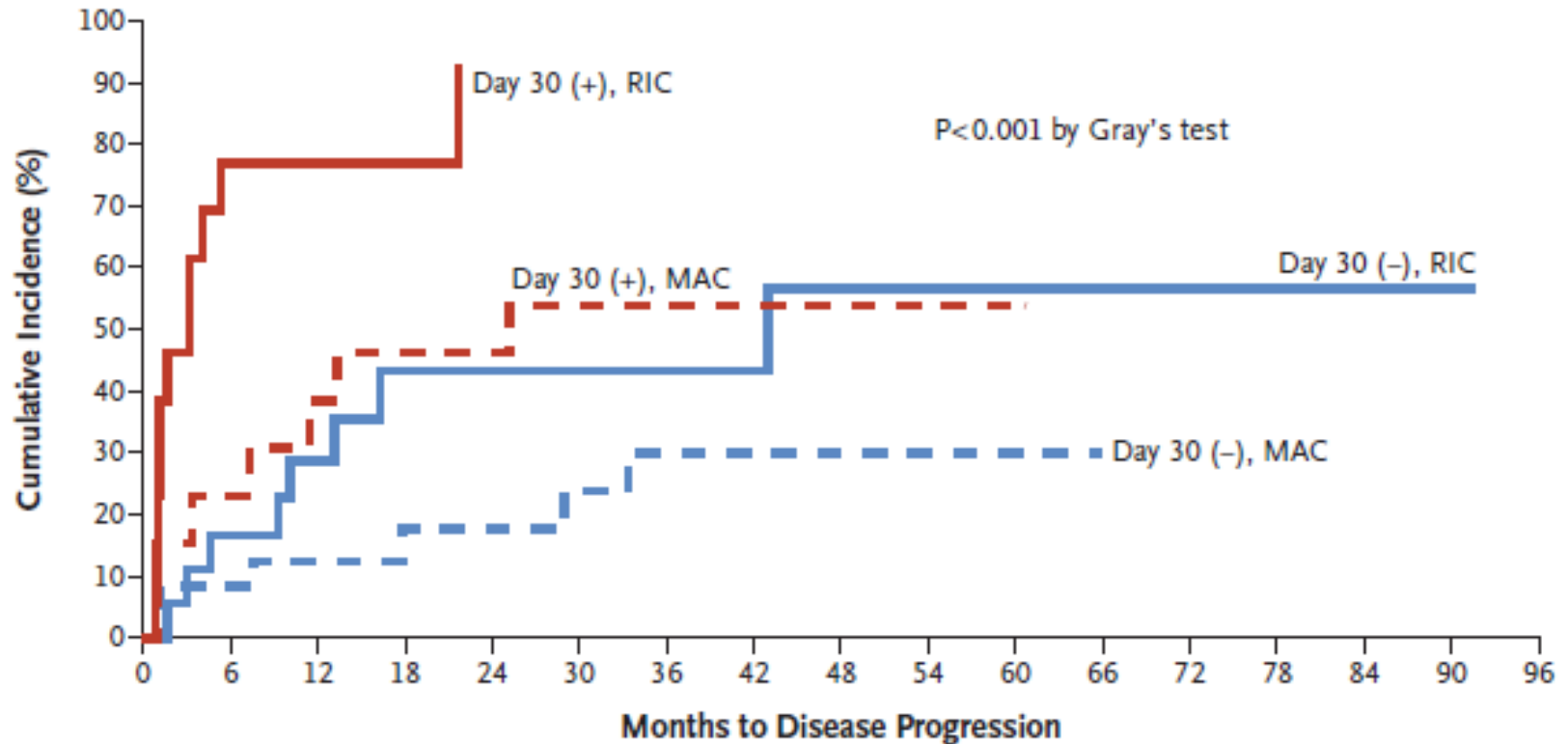


Mutation Pattern at Disease Relapse After HSCT in Patients With MDS and MDS/AML



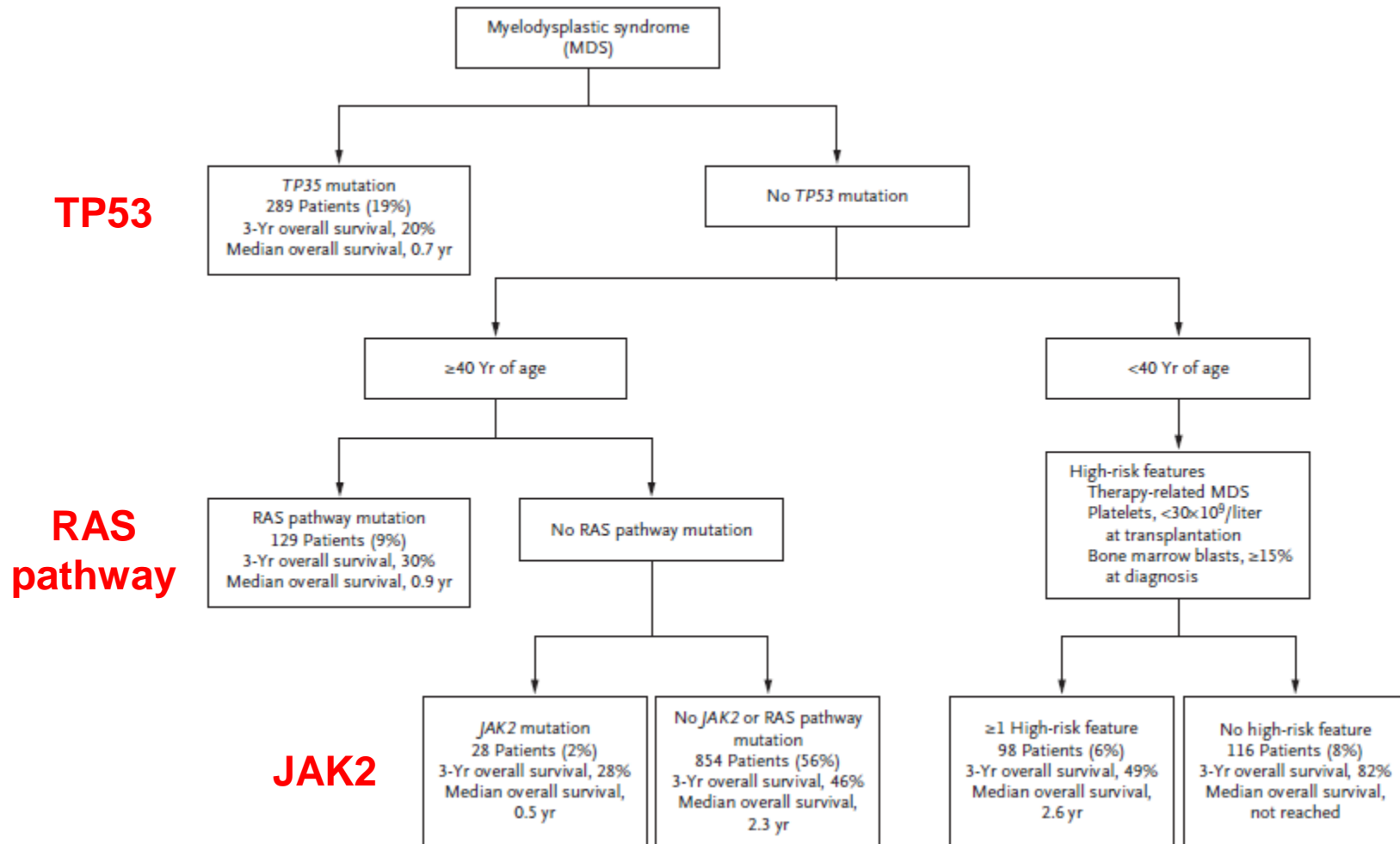
Patient	WHO Category (before HSCT)	Founding Clone (before HSCT)	Clonal Evolution (disease relapse)
GITMO 1	RAEB-2	<i>PTPN11</i>	Founder clone recurs
GITMO 2	MDS/AML	<i>NPM1</i>	Founder clone recurs
GITMO 3	RAEB-1	<i>RUNX1</i>	Founder clone recurs
GITMO 4	RAEB-2	<i>DNMT3A</i>	A subclone expands (<i>IDH1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

Mutation Clearance after Transplantation for Myelodysplastic Syndrome



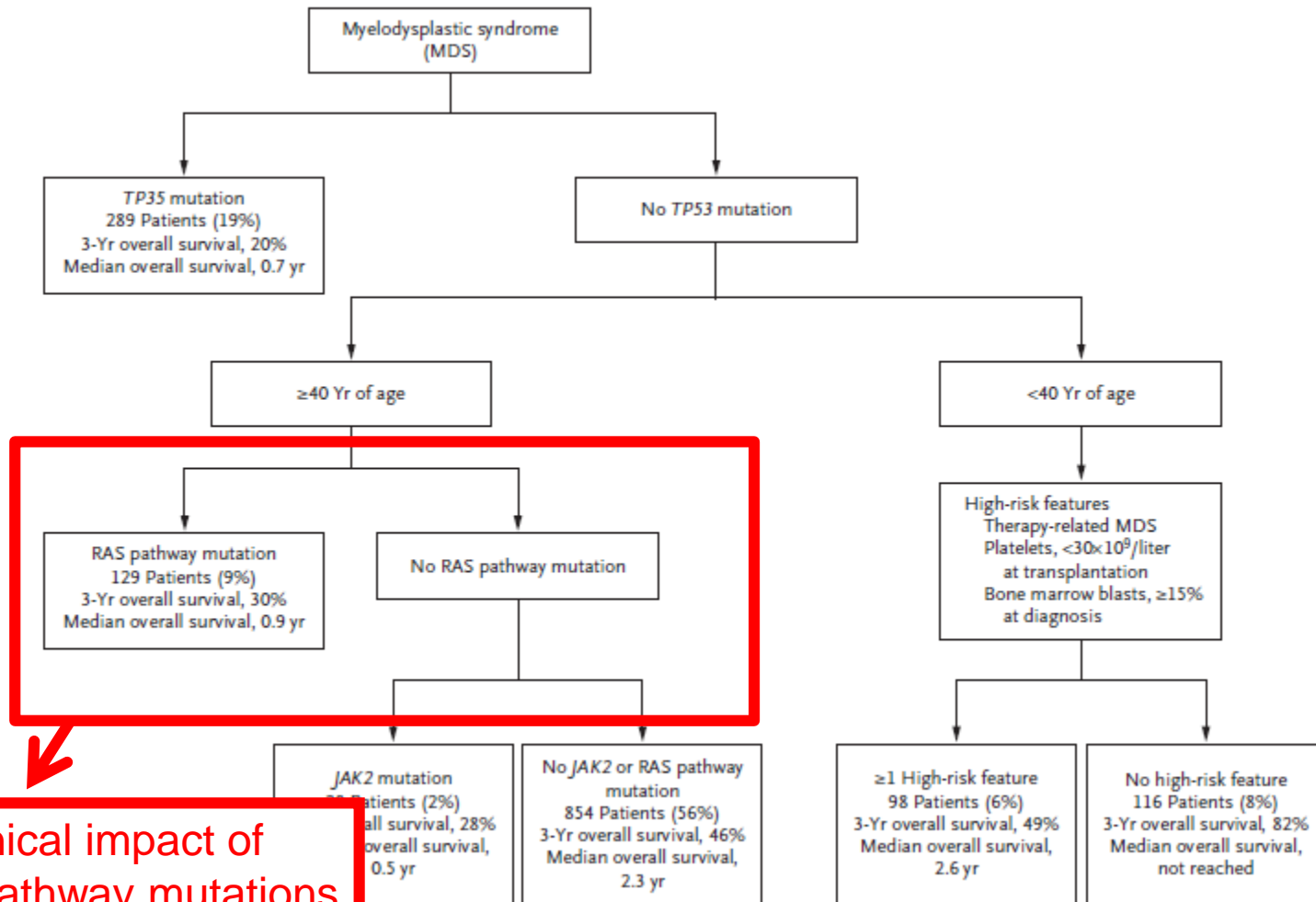
N Engl J Med 2018;379:1028-41.

Prognostic Mutations in Myelodysplastic Syndrome after Stem-Cell Transplantation



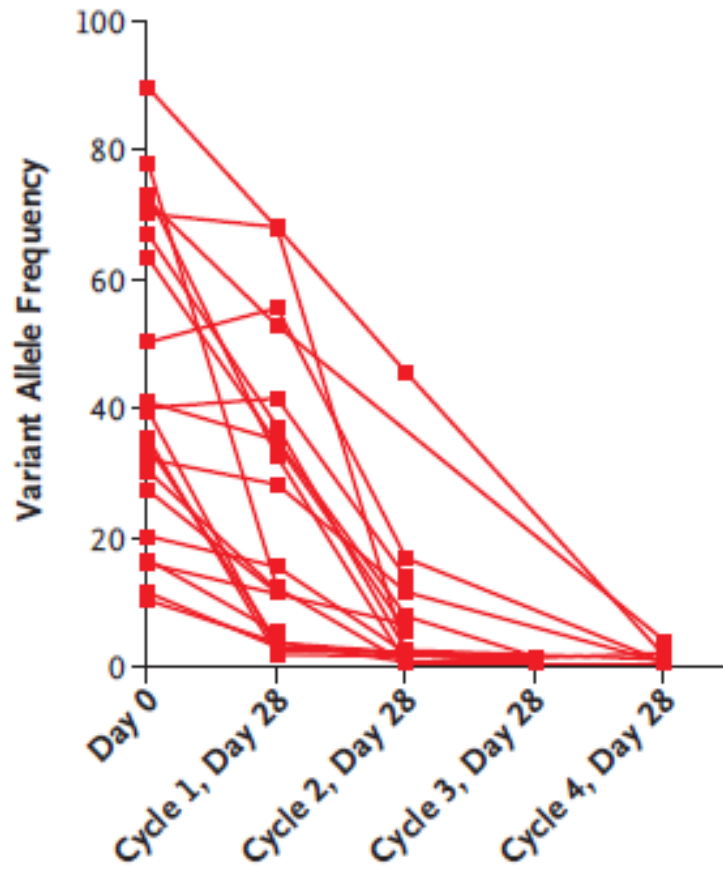
Lindsley, RC et al. N Engl J Med 2017;376:536-47.

Prognostic Mutations in Myelodysplastic Syndrome after Stem-Cell Transplantation

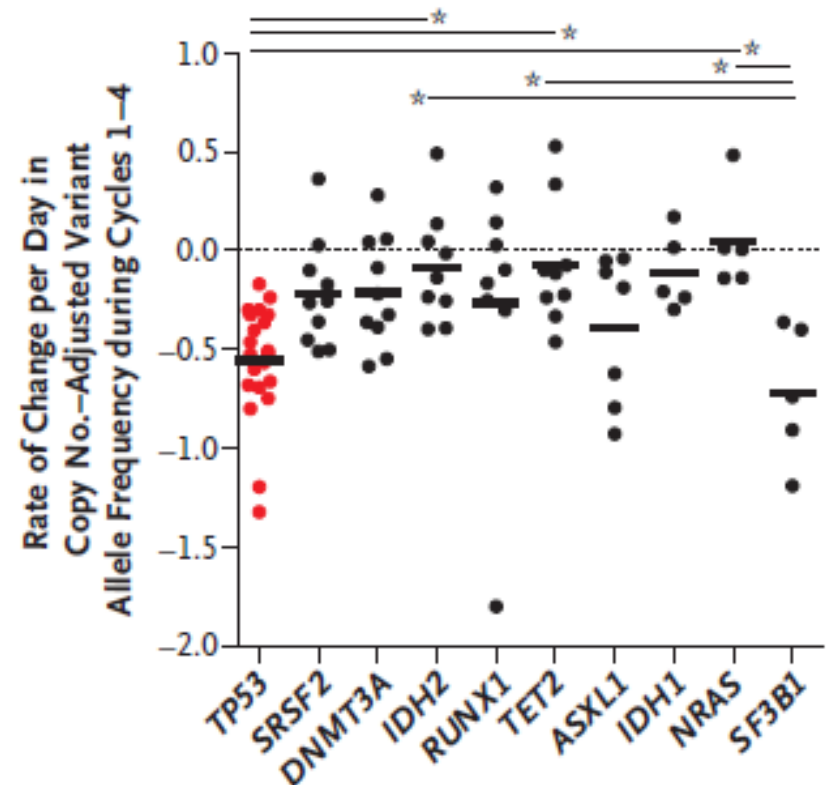


TP53 and Decitabine in Acute Myeloid Leukemia and Myelodysplastic Syndromes

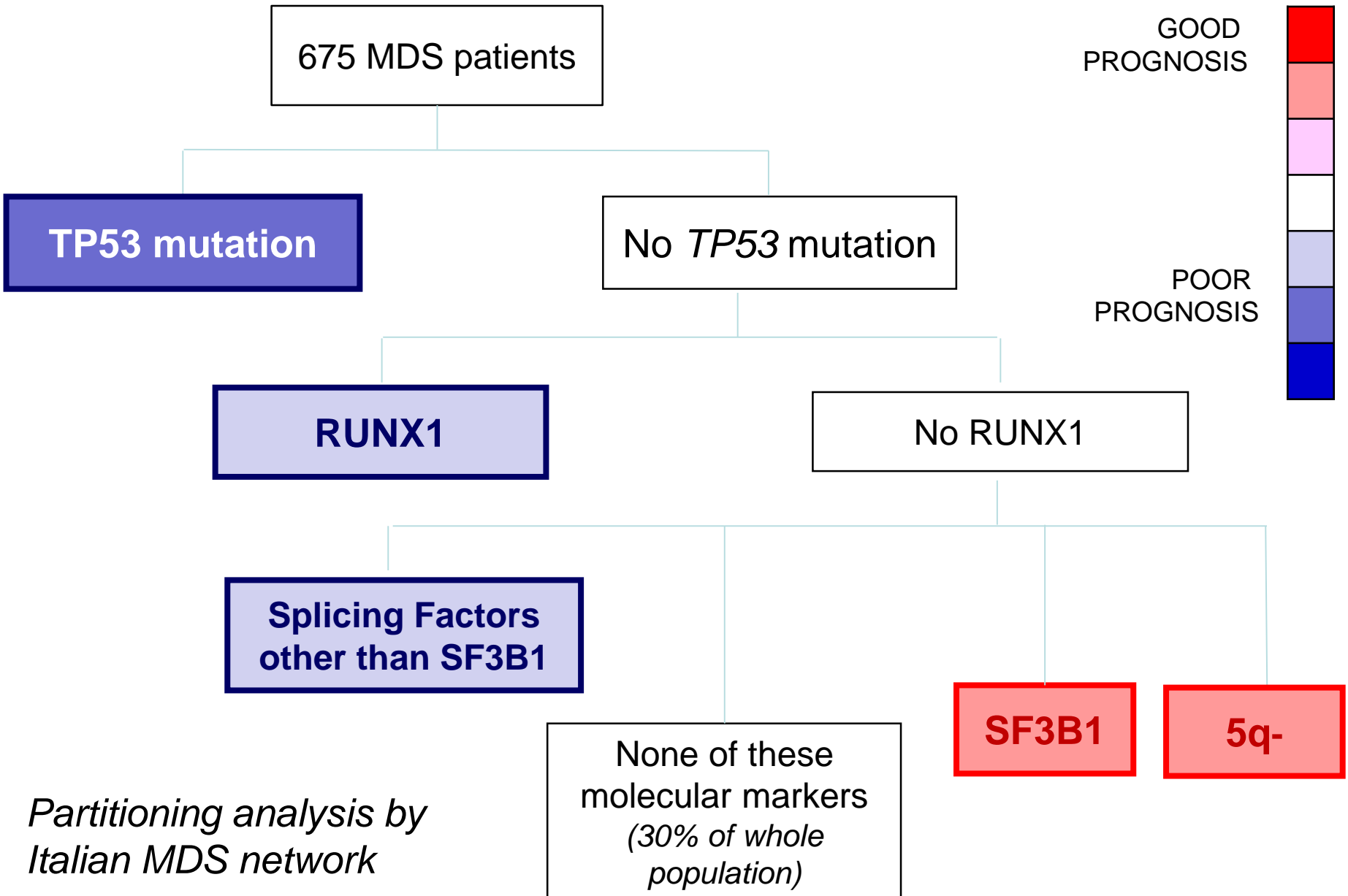
D Clearance of *TP53* Mutations



E Clearance of Mutations



Genotype-based transplant strategy in MDS



SUMMARY

- The implementation of IPSS-R is expected to result in a more effective prognostic assessment among patients with early disease stage
- Mutation screening provides relevant prognostic information at individual patient level
- According to a IPSSR-based transplantation strategy, maximal life expectancy was obtained when delaying allo-HSCT after progression to the intermediate risk score.
- Mutation screening may affect clinical decision making in transplantation (TP53 mutations are associated to a high probability of disease relapse)

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