

Understanding your MDS risk score

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Case 1 - Mary

- 66 yo woman with anemia and low white blood cell count
- She has a bone marrow biopsy and is told she has myelodysplastic syndrome with excess blasts-1
- What does this mean?
- How can we try to determine how this might behave?



Canadian and Global Incidence of MDS

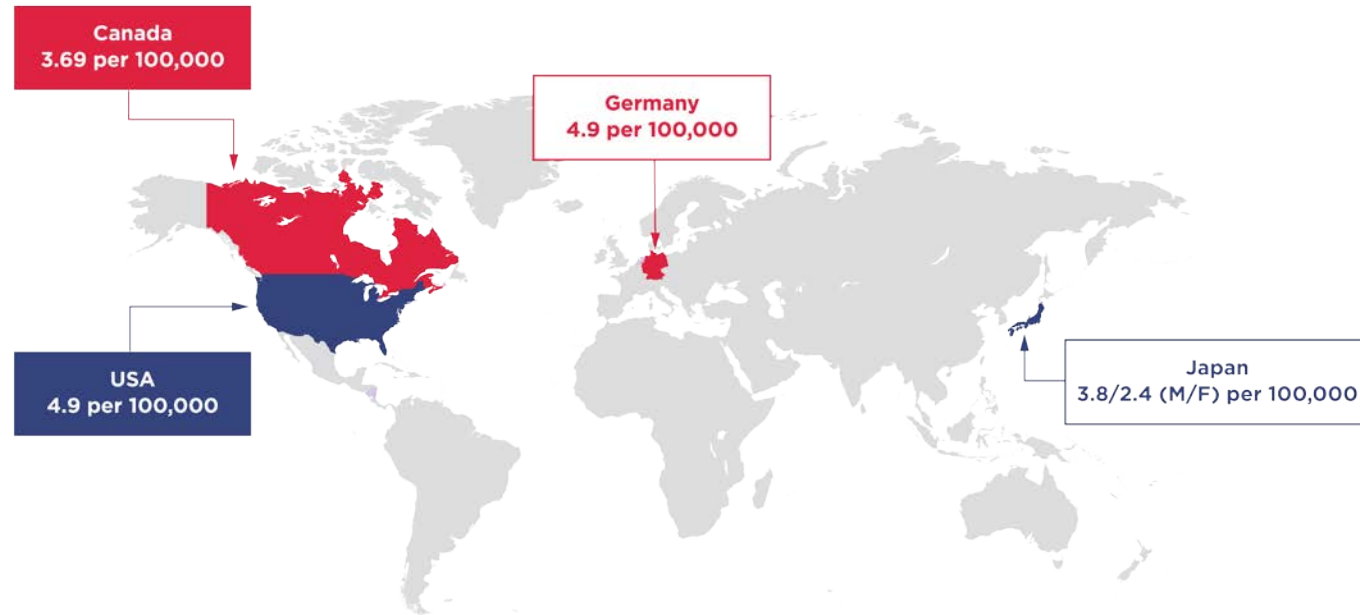
> 350,000 patients have MDS worldwide¹

In Canada, it is estimated that there are 10,000 – 40,000 patients who have been diagnosed with MDS²

- Approximately 1,800 – 5,900 new cases are diagnosed each year in Canadians >65 years²
-

The highest incidence rates are found in the U.S, Japan, and Germany³

ANNUAL INCIDENCE OF MDS AROUND THE WORLD

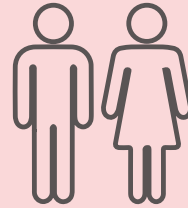


MDS, myelodysplastic syndromes.

¹Research and Markets. Global Myelodysplastic Syndrome (MDS) Market 2017-2027: Prevalence Forecast, Licensing and Acquisition Deals & Drug-Specific Revenue Forecasts. Available at: <https://www.globenewswire.com/news-release/2018/01/22/1298306/0/en/Global-Myelodysplastic-Syndrome-MDS-Market-2017-2027-Prevalence-Forecast-Licensing-and-Acquisition-Deals-Drug-Specific-Revenue-Forecasts.html>. ²Leukemia & Lymphoma Society of Canada. Blood Cancer in Canada. 2016. ³Slack J, et al. Journal of Applied Laboratory Medicine. 2018;03:378-383.

DEMOGRAPHICS OF MDS

The incidence of MDS varies depending on:



Sex

MDS affects more males than females¹



Race

MDS affects individuals of all races but the incidence rate is highest in White individuals¹



Age

The median age of diagnosis for MDS is 76 years²
Incidence rates increase drastically in individuals >80 years³

*Based on data from the United States
MDS, myelodysplastic syndromes.

¹Rollison DE, et al. Blood. 2008;112;45-52. ²Wells R, et al. Canadian Medical Association Journal. 2016;188(10):751. ³SEER Cancer Statistics Review 1975-2016. Section 30. National Cancer Institute. Available at https://seer.cancer.gov/csr/1975_2016/results_merged/sect_30_mds.pdf#search=mds.

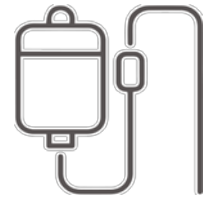
RISK FACTORS FOR DEVELOPING MDS

Most cases of MDS occur *de novo* but a number of risk factors have been identified:



Environmental/ Lifestyle

Occupational benzene exposure (e.g., petrochemical industry), or through cigarette smoke (increases MDS risk for as long as 15 years from time of quitting)^{1,2}



Therapy-Related*

Therapy-related MDS (t-MDS) constitutes **10 – 15%** of all MDS cases³
T-MDS is typically considered higher-risk, with greater resistance to treatment, and worse prognosis⁴



Other

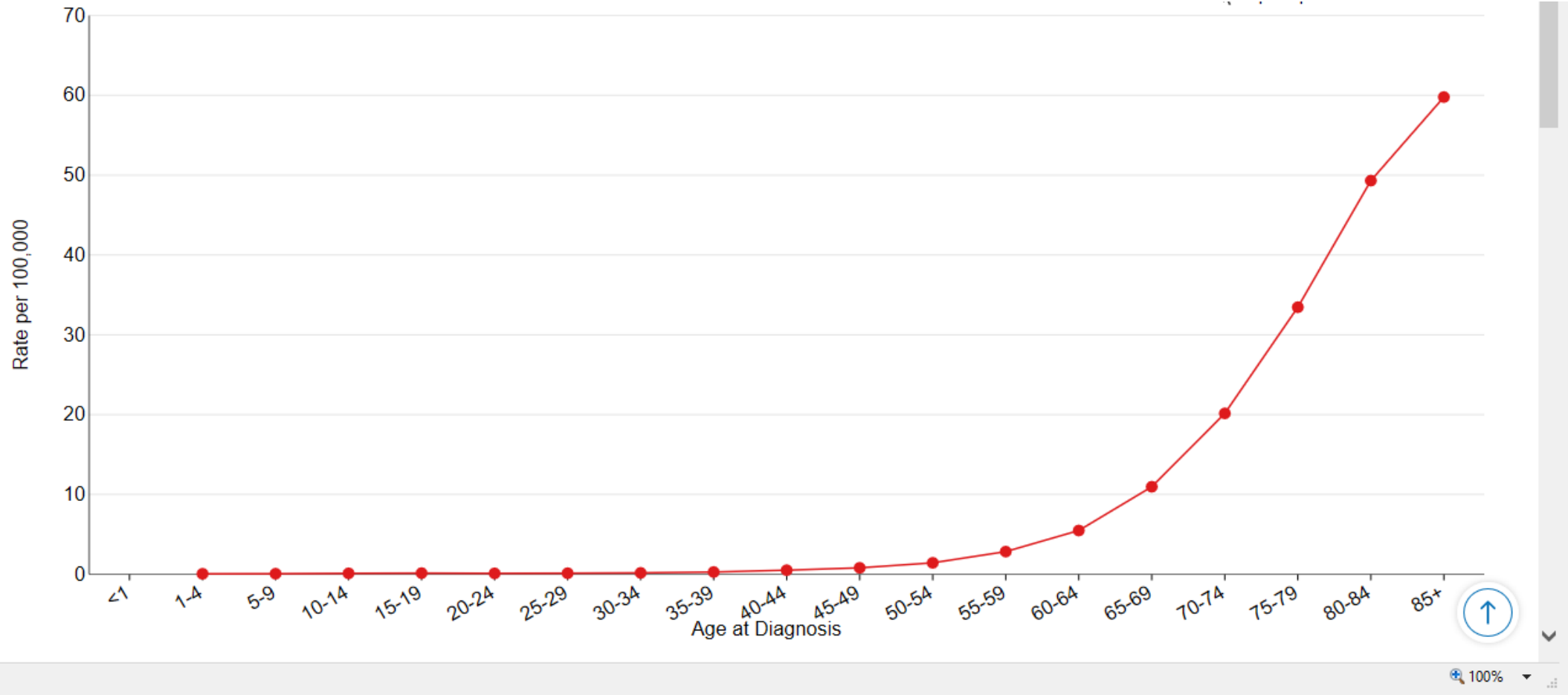
Family history of hematopoietic cancers, certain inherited blood disorders[†], and a BMI >25 increase an individual's risk of developing MDS^{1,3,5}

*Developed after chemotherapy and/or radiotherapy treatment for other cancers.[†]Such as Fanconi anemia, congenital neutropenia.

BMI, body mass index; MDS, myelodysplastic syndromes; t-MDS, therapy-related MDS.

¹Strom SS, et al. Leukemia. 2005;19:1912-1918. ²Schnatter AR, et al. J Natl Cancer Inst 2012. 104:1724-1737. ³Steensma DP. Mayo Clinic Proceedings. 2015;90:969-983. ³Fukumoto JS, Greenberg PL. Critical Reviews in Hematology Oncology. 2005;56:179-192. ⁵Ma X, et al. American Journal of Epidemiology. 2009;169:1492-1499.

Incidence rates of MDS by age at diagnosis, SEER database 2013-2017



Environmental exposures & High risk professionals¹

Environmental exposures

Professional	Adjusted OR	95% CI	P
Petrol	2.5	(0.9–7.7)	0.8
Aromatic polycyclic hydrocarbons	1.8	(0.7–4.6)	0.93
Exhaust gases	1.0	(0.5–1.9)	0.27
Dyes	2.3	(0.5–13.9)	0.77
Plastic fumes and dusts	3.5	(0.7–34.5)	0.09
Wood dusts	1.0	(0.3–3.3)	0.71
Pesticides	3.2	(1.1–11.2)	0.21
Fertilizers	2.9	(1.2–8)	0.96
Cereal dust	2.6	(1.2–6)	0.58
Poultry	15	(2.3–631.5)	0.1
Cotton and flax dusts	3.4	(1.6–8.2)	0.013
Radiation	2.25	(1.1 - 4.7)	NR

High risk professionals

Professional	Adjusted OR	95% CI	P
Agricultural workers	3.66	(1.9–7.0)	0.0001
Textile operators	3.66	(1.7–7.9)	0.001
Health professionals	10.0	(2.1–48.7)	0.004
Living next to an industrial plant	2.45	(1.5–4.1)	0.007
Commercial and technical sales representatives	4.45	(1.4–14.6)	0.013
Smoking	1.74	(1.1–2.7)	0.015
Machine operators	2.69	(1.2–6.0)	0.015
Oil use	1.10	(1.0–1.2)	0.029

1. Occupational and environmental risk factors of the myelodysplastic syndromes in the North of France, C Nisse et al, Br J Haematol, 2001 Mar;112(4):927-35

COMMON SIGNS AND SYMPTOMS OF MDS

MDS can be asymptomatic and discovered during routine blood work¹

Low blood counts associated with MDS may manifest as signs and symptoms leading to a diagnosis¹

Common Signs and Symptoms of MDS ^{1,2}
Fatigue, weakness
Loss of sense of well-being
Pale
Shortness of breath with exertion
Frequent severe infections
Easy bruising and/or bleeding (e.g., nosebleeds, gum bleeds)
Weight loss
Loss of appetite
Fever

MDS, myelodysplastic syndromes.
¹Mayo Clinic. Myelodysplastic Syndromes. Available at: <https://www.mayoclinic.org/diseases-conditions/myelodysplastic-syndrome/symptoms-causes/syc-20366977>. ²Canadian Cancer Society. Myelodysplastic Syndromes. Available at: <https://www.cancer.ca/en/cancer-information/cancer-type/leukemia/leukemia/myelodysplastic-syndromes/?region=on>.

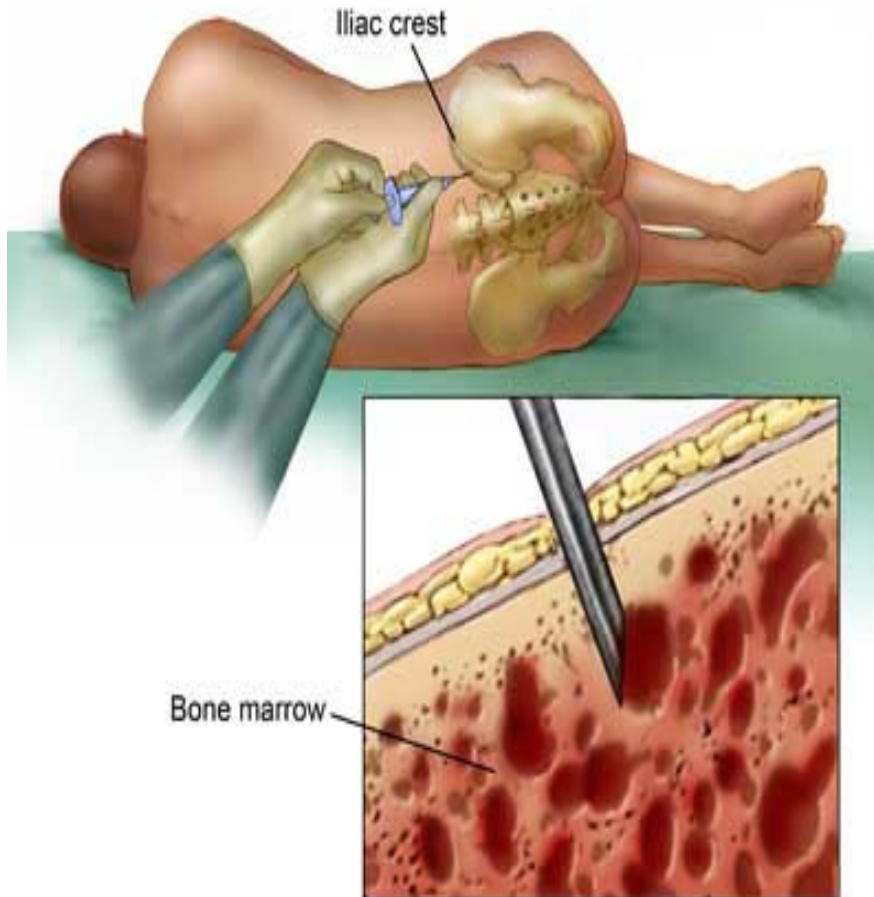
SYMPTOMS ASSOCIATED WITH LOW BLOOD COUNTS

MDS is characterized by cytopenias

- **Anemia** is the most common cytopenia in MDS
- **Thrombocytopenia (low platelets)** is observed in ~50% of patients with lower risk MDS
- Neutropenia (low neutrophils, a type of white blood cell) is observed in ~50% of patients with MDS

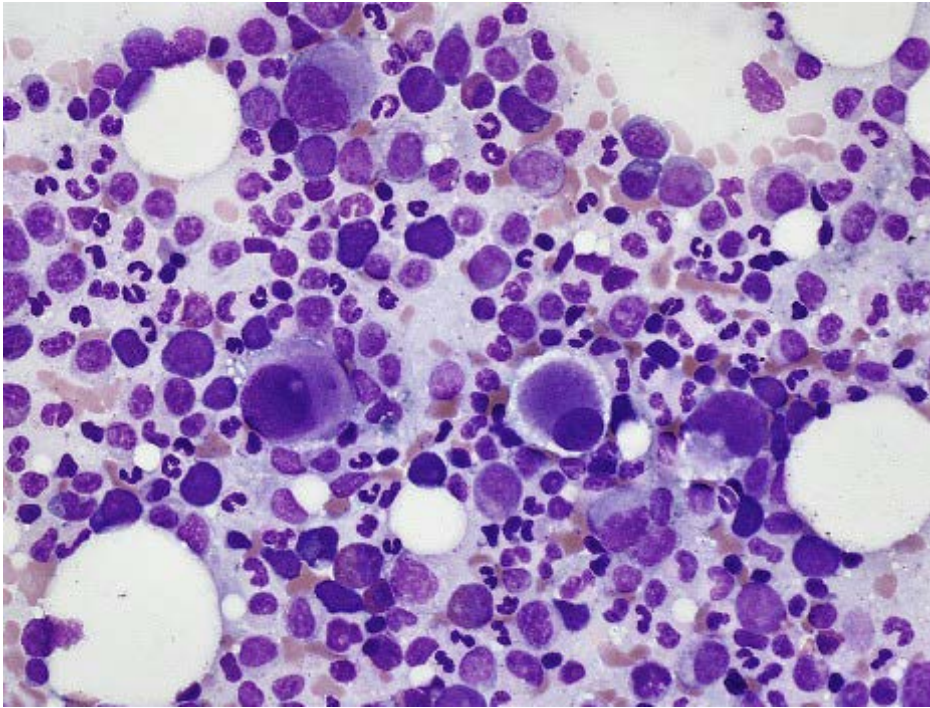
Anemia	Thrombocytopenia	Neutropenia
Fatigue	Petechiae	Fever
Weakness	Easy or excessive bruising	Frequent, severe or unusual infections
Lightheadedness	Abnormal or excessive bleeding from nose or mouth	
Shortness of breath	Unusually heavy menstrual flow	
Rapid heartbeat	Blood in urine or stool	
Pallor	Bleeding in the gastrointestinal tract	
Chest pain	Neurological symptoms including headache, confusion or balance difficulties in severe cases	
Confusion		

Bone Marrow Biopsy

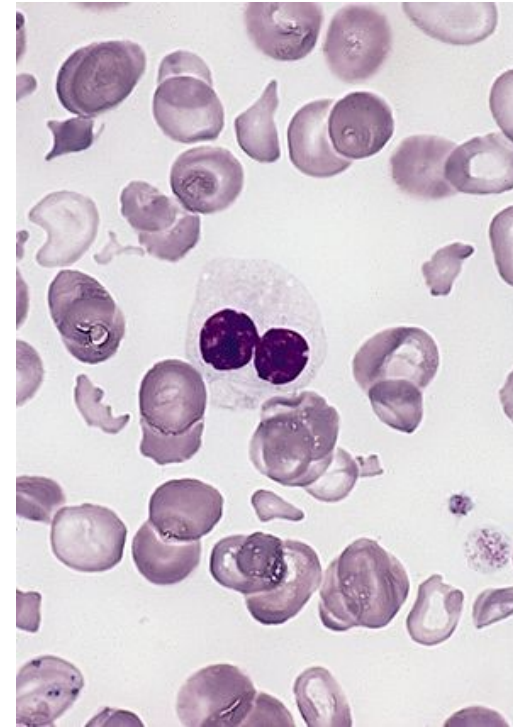


- Examination of marrow from posterior superior iliac crest by:
 - Microscopy
 - Stains for immunohistochemistry
 - Flow cytometry
 - Chromosome testing
 - Molecular testing
 - Usually acquired mutations but some are inherited

Bone marrow examination

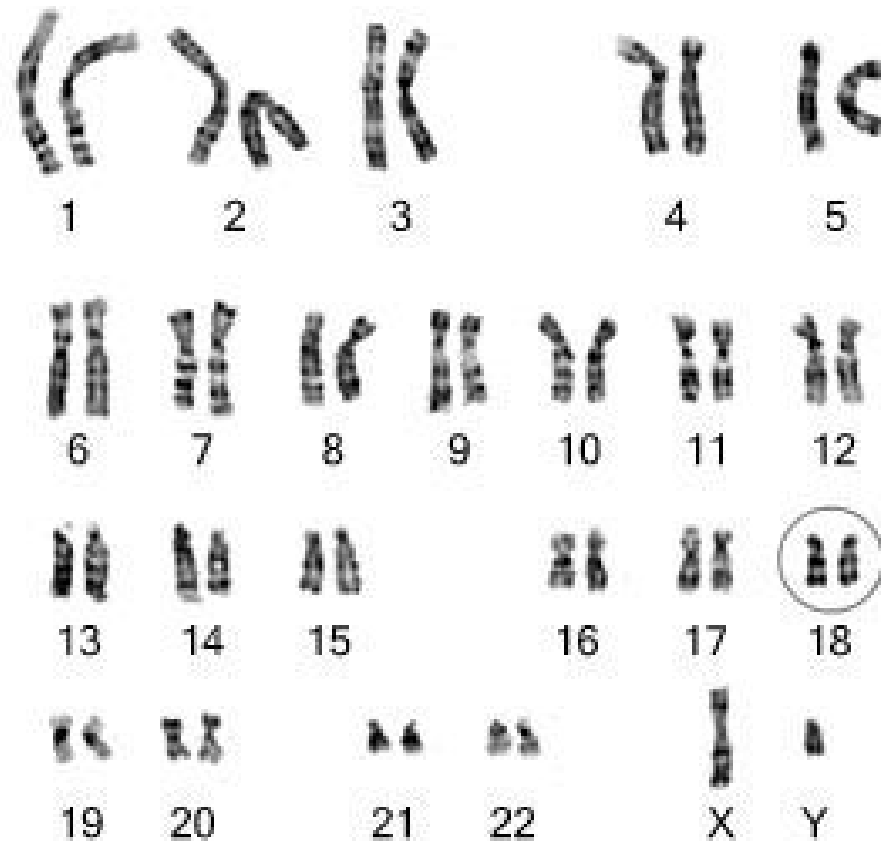


Bone marrow biopsy

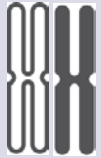


Bone marrow aspiration in
patient with MDS

Chromosome analysis (Karyotype)



DIAGNOSING MDS - CYTOGENETIC ABNORMALITIES



Cytogenetic features significantly impact prognosis and treatment, and can be used to confirm MDS diagnosis in cases where morphologic changes are subtle¹⁻⁴



~50%

of *de novo* MDS patients have ≥ 1 cytogenetic abnormality⁵



The majority of chromosomal aberrations in MDS are unbalanced, resulting in the loss or gain of chromosomal material⁶



The loss or gain of chromosomes impacts the production of gene products that are important in MDS-related pathways⁷

MDS, myelodysplastic syndromes.

¹Greenberg P, et al. Blood. 1997;89(6):2079-2088. ²Greenberg P, et al. Blood. 2012;120(12):2454-2465. ³Fenaux P, et al. Blood. 2011;118:3765-3776. ⁴Expert opinion. ⁵Haase D, et al. Blood. 2007;110:4385-4395. ⁶Ogawa S. Blood. 2019;133:1049-1059. ⁷Sperling AS, et al. Nature Reviews Cancer. 2017;17:5-19.

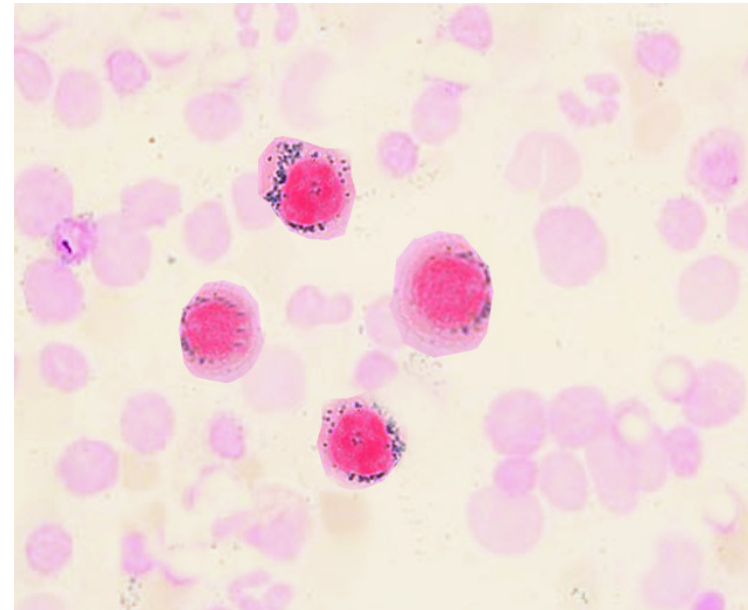
2016 WHO Classification

Myelodysplastic Syndromes:

- Myelodysplastic syndrome with single lineage dysplasia
- Myelodysplastic syndrome and multilineage dysplasia
- Myelodysplastic syndrome with ringed sideroblasts
 - Single lineage dysplasia
 - Multilineage dysplasia
- Myelodysplastic syndrome with excess blasts
 - MDS-Excess blasts-1 (5-9%)
 - MDS Excess blasts-2 (10-19%)
- Myelodysplastic syndrome with isolated del(5q)
- Myelodysplastic syndrome unclassifiable

Myelodysplastic/myeloproliferative neoplasms:

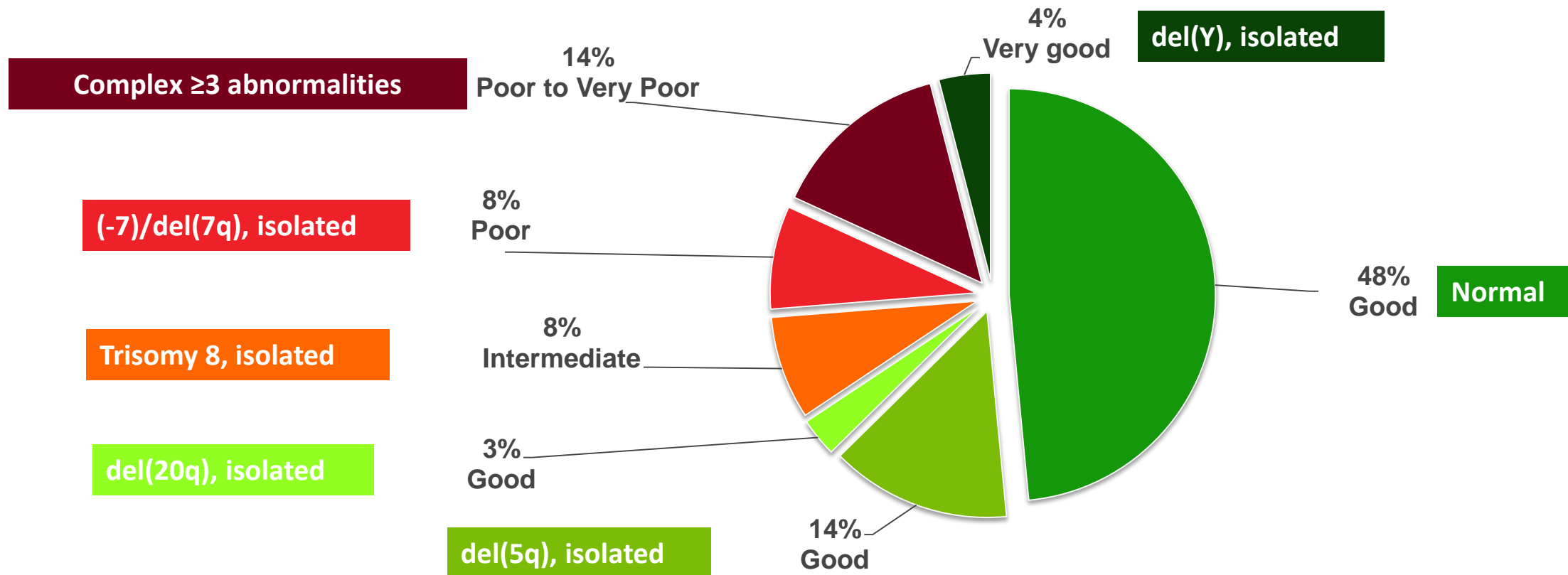
- Chronic myelomonocytic leukemia
- Others



Ringed sideroblasts

MOST COMMON CYTOGENETIC ABNORMALITIES

Incidence and Prognosis of Cytogenetic Abnormalities in MDS*^{1,2}



*Primarily in de novo MDS (93.3% primary vs. 6.7% secondary).

MDS, myelodysplastic syndromes.

¹Haase D, et al. Blood. 2007;110:4385-4395. ²Sperling AS, et al. Nature Reviews Cancer. 2017;17:5-19.

Clinical features that affect prognosis of MDS

- Blast count in the bone marrow
- Cytogenetics risk group
- Degree of low red cells, platelets and neutrophils
- Other features:
 - Secondary MDS (after previous chemo or radiation)
 - Molecular risk factors
 - Fibrosis in the marrow biopsy
 - Transfusion dependency

IPSS-R

Revised International Prognostic Scoring System

Parameter	Categories and Associated Scores				
Cytogenetics	Very good	Good	Intermediate	Poor	Very Poor
	0	1	2	3	4
Marrow Blasts	≤2%	>2 - <5%	5 - 10%	>10%	
	0	1	2	3	
Hemoglobin (g/L)	≥100	80 - <100	<80		
	0	1	1.5		
Absolute neutrophil count (x 10 ⁹ /L)	≥0.8	<0.8			
	0	0.5			
Platelet count (x 10 ⁹ /L)	≥100	50 - 100	<50		
	0	0.5	1		

Very good = del(11q)

Good = Normal, del(5q), del(12p), del(20q), double including del(5q)

Intermediate = del(7q), 8, 19, i(17q), any other single or double independent clones

Poor = inv(3)/t(3q)/del(3q), double including 7/del(7q), complex: 3 abnormalities

Very poor = Complex: 3 abnormalities

Link to online R-IPSS calculator: <https://www.mds-foundation.org/advanced-calculator/>

IPSS-R Clinical Outcomes

Risk Category	Risk Score	Median Survival	Time for 25% to Progress to AML
Very Low	1.5	8.8 years	Not Reached
Low	1.5-3	5.3 years	10.8 years
Intermediate	3-4.5	3.0 years	3.2 years
High	4.5-6	1.6 years	1.4 years
Very high	6	0.8 years	0.73 years

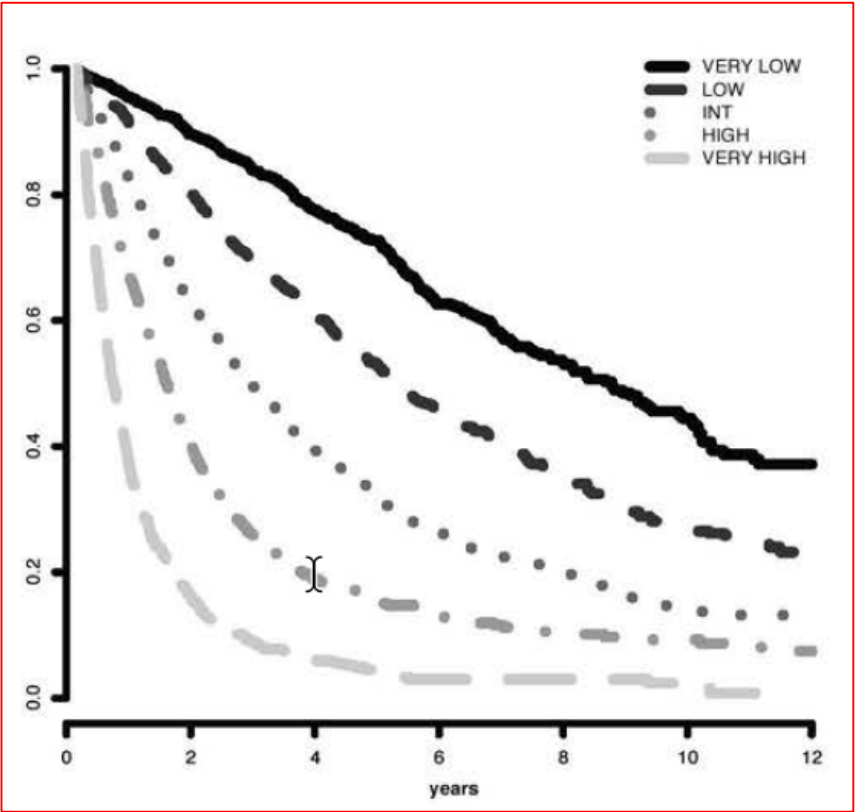
IPSS-R to be used at the time of diagnosis only. It is not applicable for therapy-related MDS. Outcomes are relevant in the absence of therapy.

Survival and AML Progression: IPSS-R¹

Survival and AML Progression

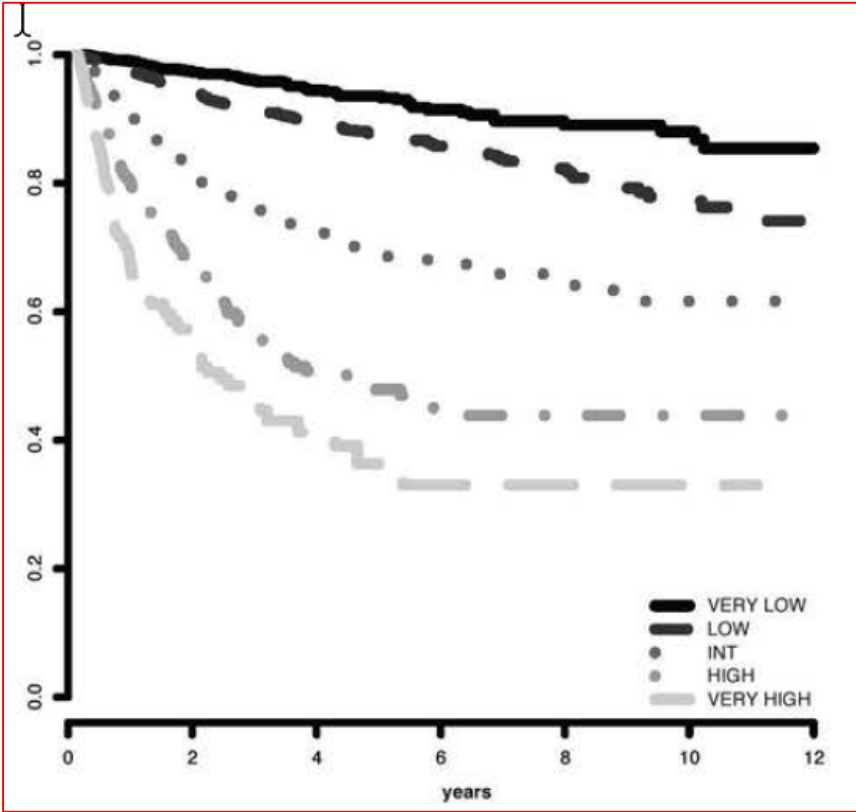
Survival based on IPSS-R prognostic risk-based categories

Patients
alive



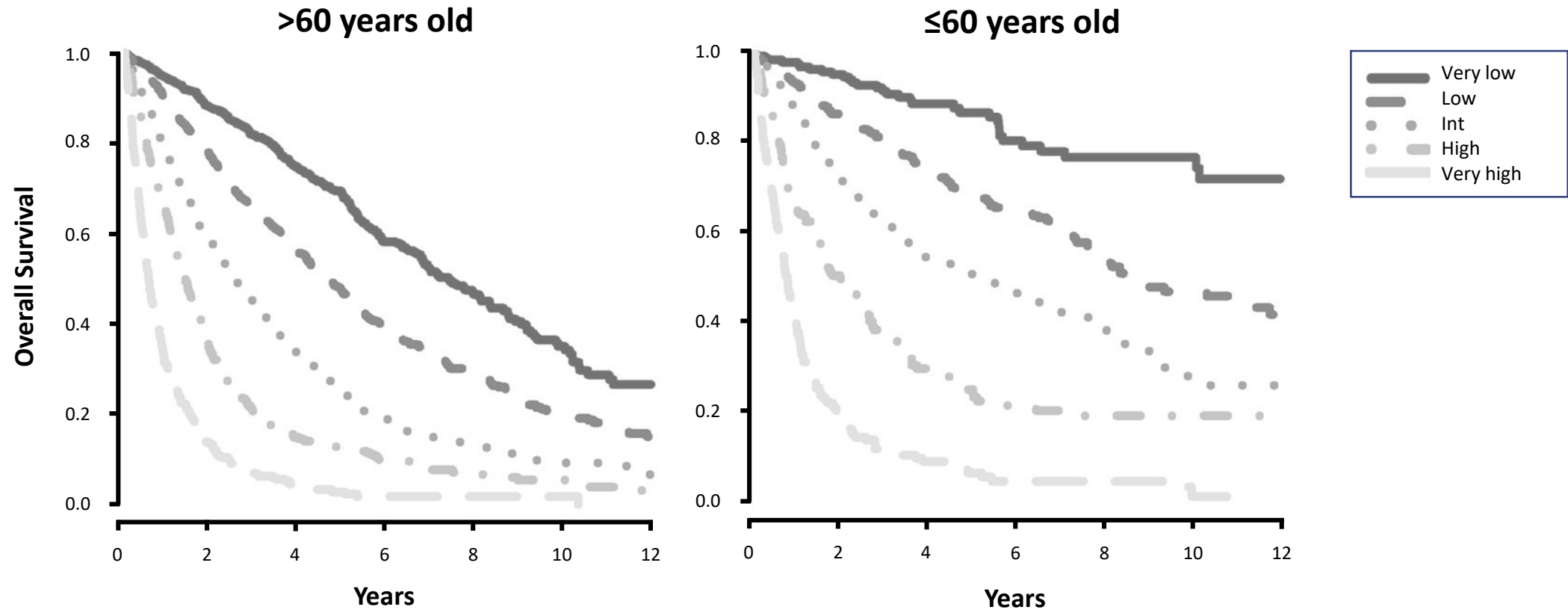
AML evolution based on IPSS-R prognostic risk-based categories

Patients
free of AML



OVERALL SURVIVAL ACCORDING TO IPSS-R SCORE

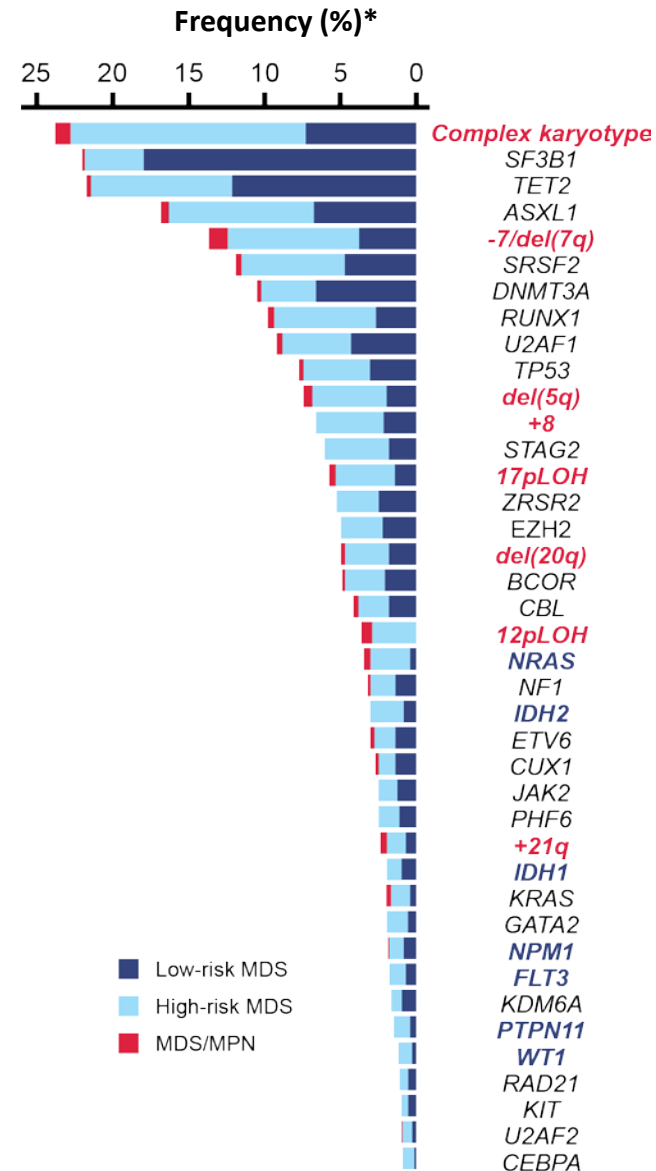
Overall Survival According to IPSS-R Score in Patients with MDS Aged >60 Years and ≤60 Years¹



DRIVER ALTERATIONS IN MDS

- >30 driver genes involved in the pathogenesis of MDS have been identified
- A typical MDS patient has a median of 2 or 3 driver mutations
- High-risk MDS tends to show higher numbers of driver mutations than lower-risk MDS

*Frequencies of major driver mutations and CNAs are plotted, combining data from 7 publications
 AML, acute myeloid leukemia; CNA, chromosomal and copy-number abnormalities; MDS, myelodysplastic syndromes; MPN, myeloproliferative neoplasms
 Ogawa S. Blood. 2019;133(10):1049-1059



MDS is Heterogeneous

- DNA damage and associated evolution of clonal cytogenetic abnormalities
 - Frequent abnormalities in DNA methylation and epigenetics

Signalling	Epigenetic	mRNA Splicing	Transcription	Cohesin genes	DNA damage
Jak2 CBL NRAS PTPN11 BRAF GNAS	TET2 DNMT3A IDH1/2 EZH2 ASXL1	SF3B1 SRSF2 U2AF1/2 ZRSR2	RUNX1 ETV6 BCOR CEBPA GATA2	STAG2 SMC1A SMC3 RAD21	P53

MDS chromosomal abnormalities & Gene mutations¹

Overall Survival based on number of mutations¹

17 genes sequenced in 1996 patients with OS data

ASXL1

CBL

DNMT3A

ETV6

EZH2

IDH1

IDH2

JAK2

KRAS

NPM1

NRAS

RUNX1

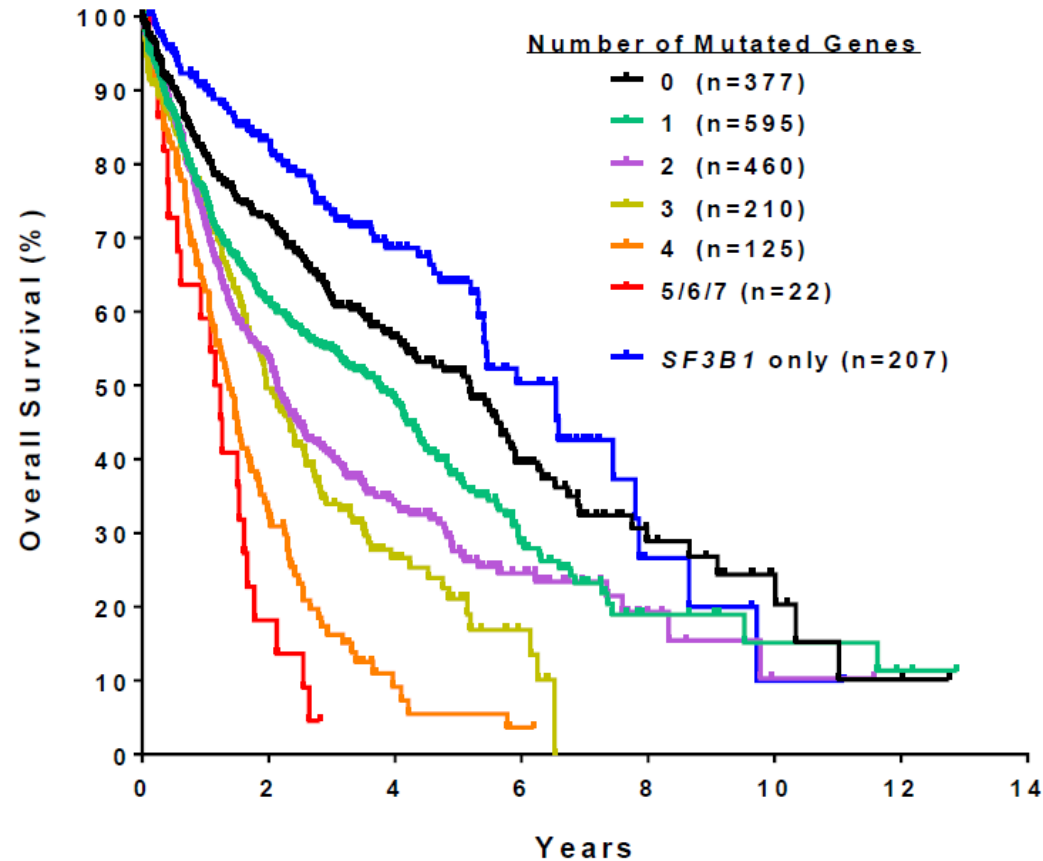
SRSF2

TET2

TP53

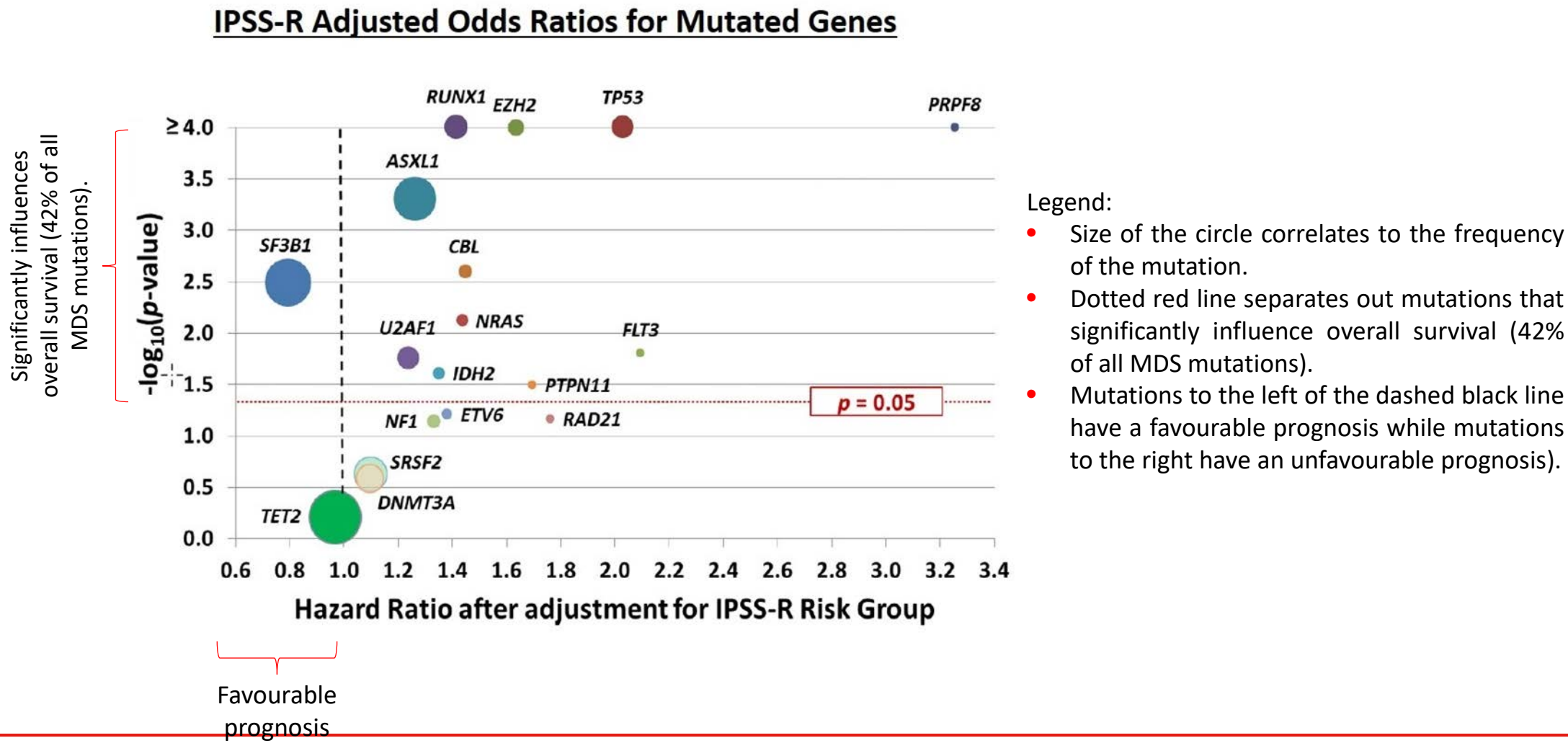
U2AF1

SF3B1



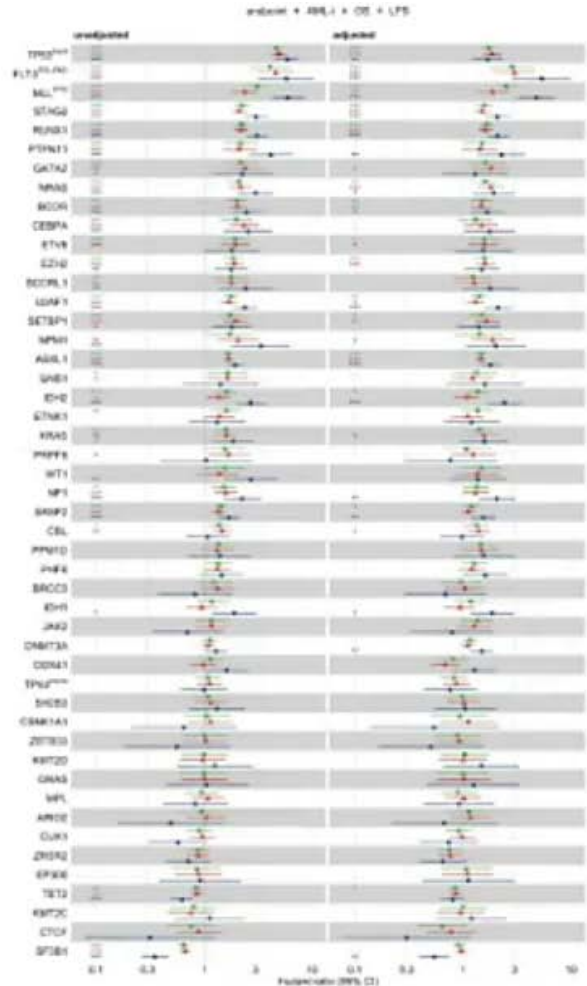
Some gene mutations are good prognosis and others are poor prognosis

IPSS-R Adjusted Odds Ratios for Mutated Genes¹



1. Integrating Molecular Findings into the Diagnosis and Prognosis of MDS, Rafael Bejar, Biological and Clinical Advances in MDS, San Diego, CA, December 2nd, 2016; Link: <https://www.mds-foundation.org/wp-content/uploads/2016/12/Bejar-IntegratingMolecularFindings.pdf>; Accessed on 22nd Dec, 2020

Molecular IPSS score (IPSS-M) for MDS



Association between gene mutations and clinical endpoints

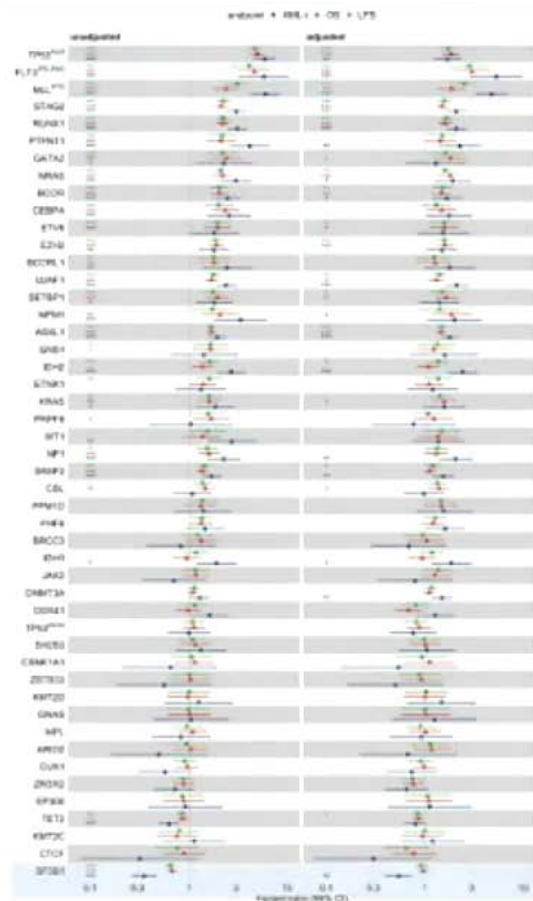
Leukemia free survival (LFS)

Overall survival (OS)

AML transformation (AML-t)

Adjusting for age, sex, MDS type (primary vs. therapy-related), and IPSS-R raw score,

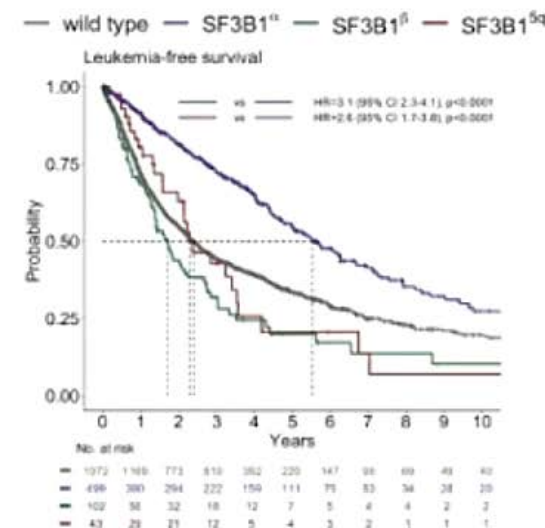
14, 16 and 15 genes were significantly associated with adverse outcomes for the three endpoints, respectively.



SF3B1 mutations were associated with favorable outcomes.

But this association was modulated by its **pattern of co mutations**.

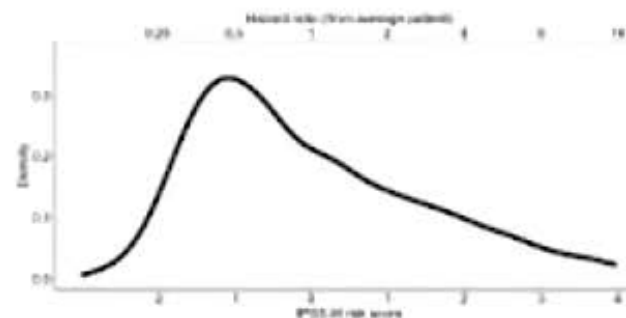
1. ***SF3B1*^{5q} (7%)**
Concomitant isolated del(5q)
2. ***SF3B1*^β (15%)**
Co-occurrence of mutations in *BCOR*, *BCORL1*, *RUNX1*, *NRAS*, *STAG2*, *SRSF2*
3. ***SF3B1*^α (78%)**
Any other *SF3B1* mutations.



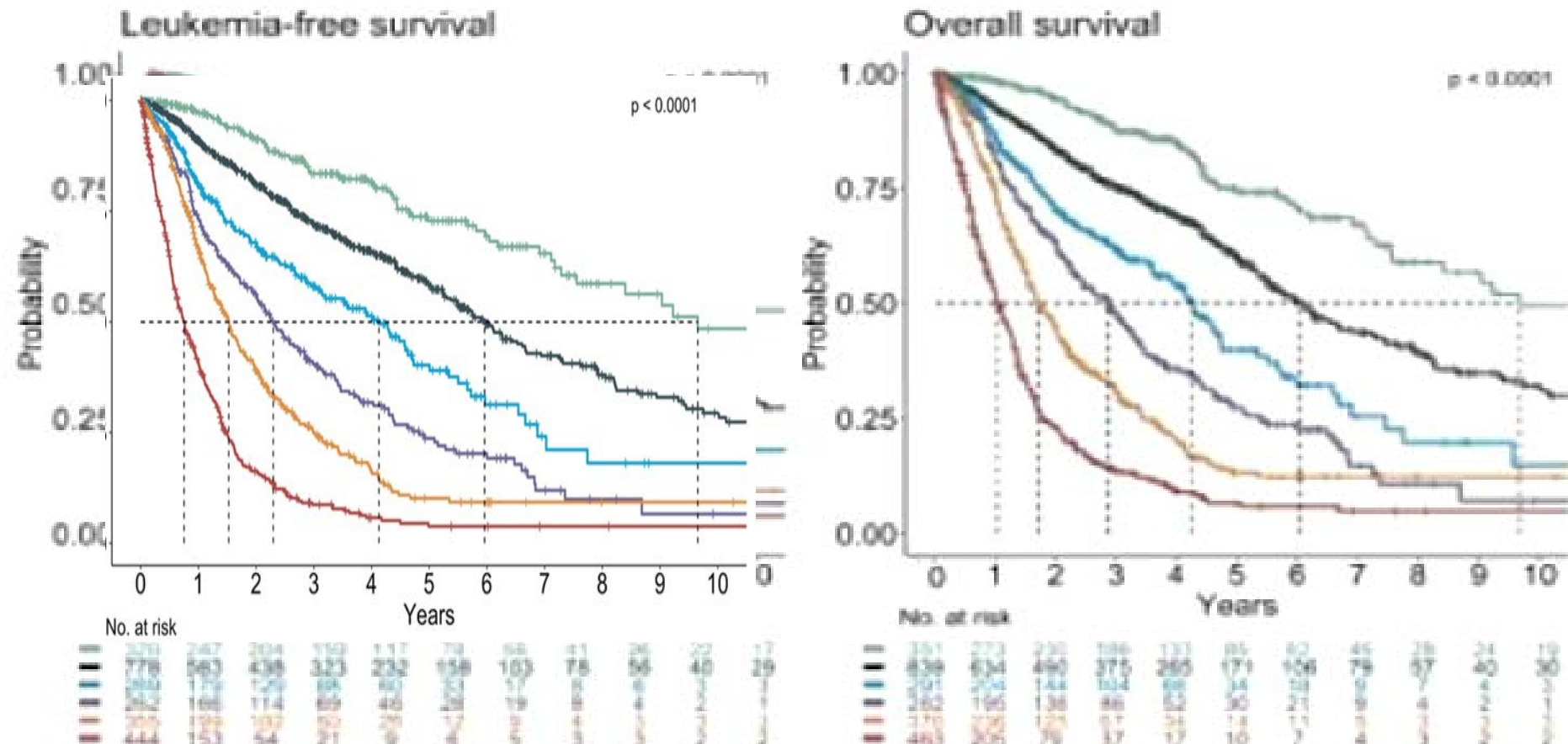
Malcovati et al. Blood 2011;2015;2021
Meggendorfer et al. Haematologica 2017

Category	Variable	Multivariable model: hazard ratio ^a (95% CI)	Weight w	Scaling σ^{mean}
confounder	Age, in years	1.23 (1.05 - 1.43)	N/A	N/A
	Sex/Male	1.22 (1.06 - 1.41)	N/A	N/A
	Type/Secondary/Therapy-related	1.26 (1.10 - 1.46)	N/A	N/A
clinical	% Bone Marrow Blasts, in %	1.42 (1.30 - 1.55)	0.352	0.922
	% mat(Platelets,250), in x10 ⁹ /L	0.89 (0.72 - 0.89)	-0.222	1.41
	Hemoglobin, in g/dL	0.84 (0.81 - 0.88)	-0.171	9.87
cytogenetics	IPSS-R category vector ^a	1.33 (1.21 - 1.47)	0.267	1.380
gene main effects 17 variables, 15 genes	TP53 ^{mut}	3.27 (2.38 - 4.48)	1.18	0.0710
	BCR ^{TR}	2.22 (1.49 - 3.32)	0.706	0.0247
	FLT3 ^{ITD}	2.22 (1.11 - 4.45)	0.706	0.0108
	SP3B ^{WT}	1.68 (1.03 - 2.68)	0.504	0.0168
	NPM1	1.54 (0.78 - 3.02)	0.430	0.0112
	RUNX1	1.53 (1.23 - 1.89)	0.423	0.126
	KRAS	1.52 (1.05 - 2.20)	0.417	0.0360
	ETV6	1.48 (0.98 - 2.23)	0.391	0.0216
	IDH2	1.46 (1.05 - 2.02)	0.379	0.0426
	CBCL	1.34 (0.99 - 1.82)	0.296	0.0473
	EZH2	1.31 (0.86 - 1.75)	0.270	0.0588
	UGAF1	1.28 (1.01 - 1.61)	0.247	0.0866
	SRSF2	1.27 (1.03 - 1.56)	0.239	0.158
	DNM3A	1.25 (1.02 - 1.53)	0.221	0.181
	ASXL1	1.24 (1.02 - 1.51)	0.213	0.252
	KRAS	1.22 (0.84 - 1.77)	0.202	0.0271
	SP3B ^{WT}	0.92 (0.74 - 1.14)	-0.3764	0.186
gene residuals ^b 1 variable, 15 genes	res(Nres,2) Possible values are 0,1 or 2	1.26 (1.12 - 1.42)	0.331	0.368

$$\text{score} = \frac{1}{\log 2} \sum_{\text{variables } j} w_j (x_j - x_j^{\text{mean}})$$



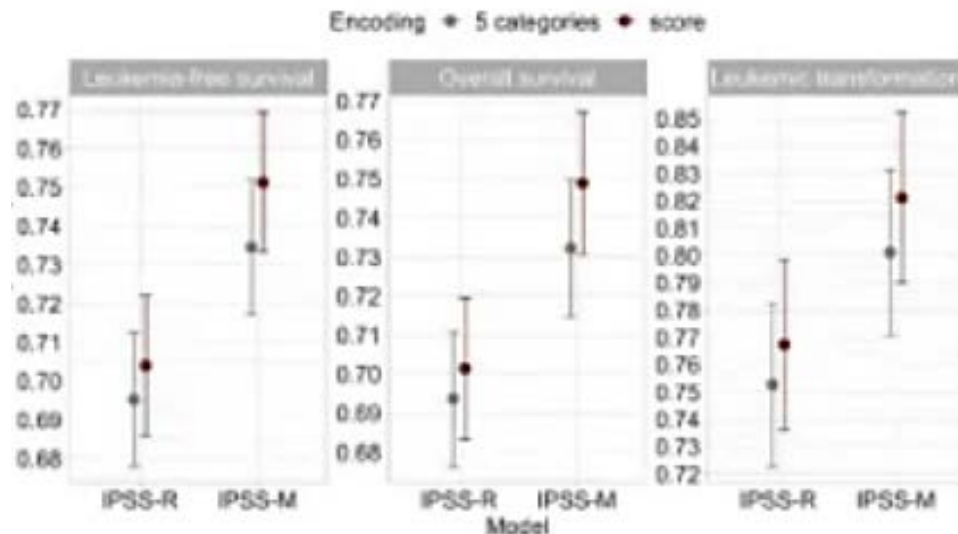
IPSS-M — Very Low — Moderate Low — High
 — Low — Moderate High — Very High



Prognostic separation of the IPSS-M risk categories

Better prediction of outcomes with incorporation of molecular mutations and clinical factors

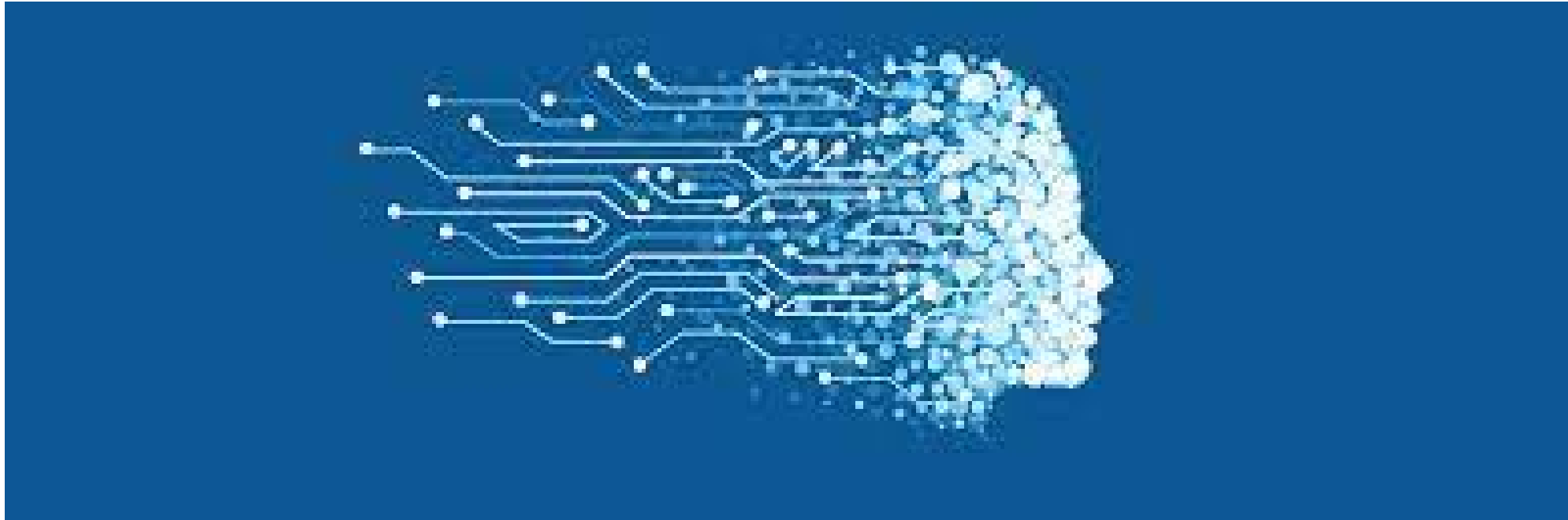
Improved prognostic discrimination



Web calculator that helps look at mutations and computational status and incorporate these with clinical features to improve prognostication

Approximately 46% of 1,223 patients were reclassified to a different risk group

Machine Learning may help incorporate MDS molecular risk factors into prognostic scores



GENOMED4ALL: Improving ...

Login to Concerto

FileEditViewFavoritesToolsHelp

https--iriss.ualgary

Suggested Sites

desktop

https--iriss ucalgary

GENOMED4ALL: IMPROVING MDS CLASSIFICATION AND PROGNOSIS BY AI

SPONSOR

Istituto Clinico Humanitas (Other)

OVERALL STATUS

Active, not recruiting

CT.GOV ID

NCT04889729

COLLABORATOR

(none)

8,200

ENROLLMENT

1

LOCATION

45.6

ANTICIPATED DURATION (MONTHS)

179.9

PATIENTS PER SITE PER MONTH

STUDY DETAILS

STUDY DESCRIPTION

BRIEF SUMMARY

Save Item

View History of Changes

STUDY DETAILS

STUDY DESCRIPTION

STUDY DESIGN

ARMS AND INTERVENTIONS

OUTCOME MEASURES

ELIGIBILITY

CONTACTS AND LOCATIONS

MORE INFORMATION

No Results Posted

100%

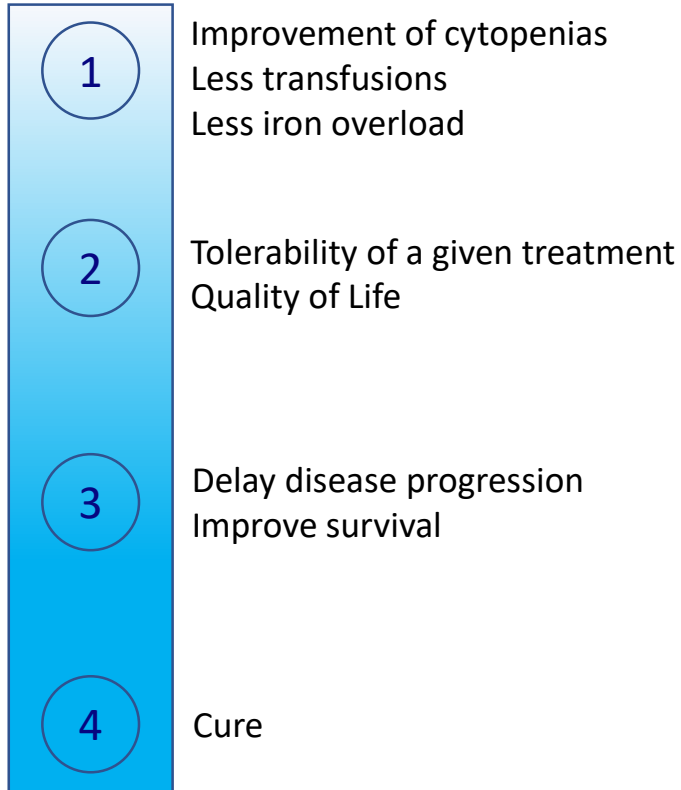
How do we use this information to guide care?



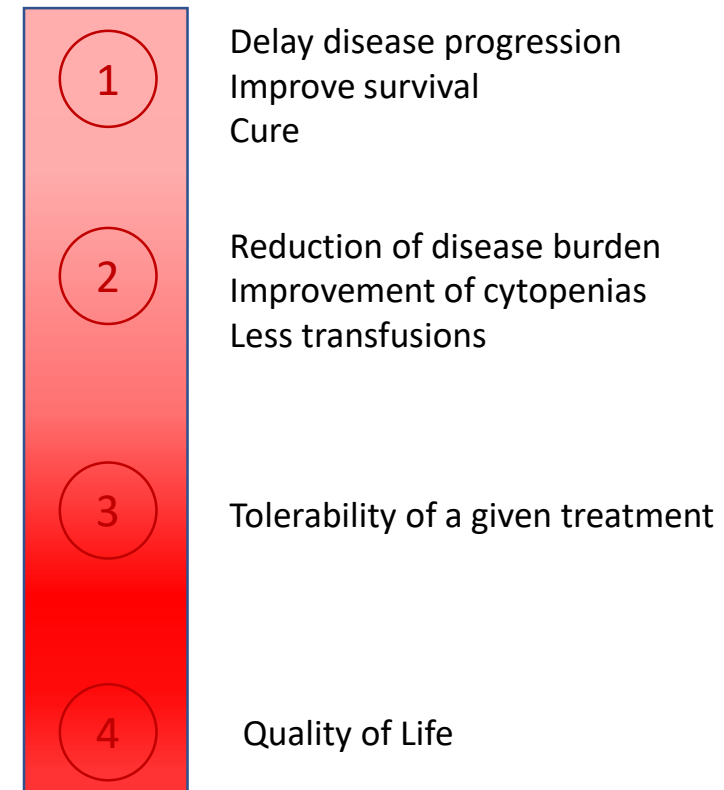
Goals of Therapy

Goals of therapy¹

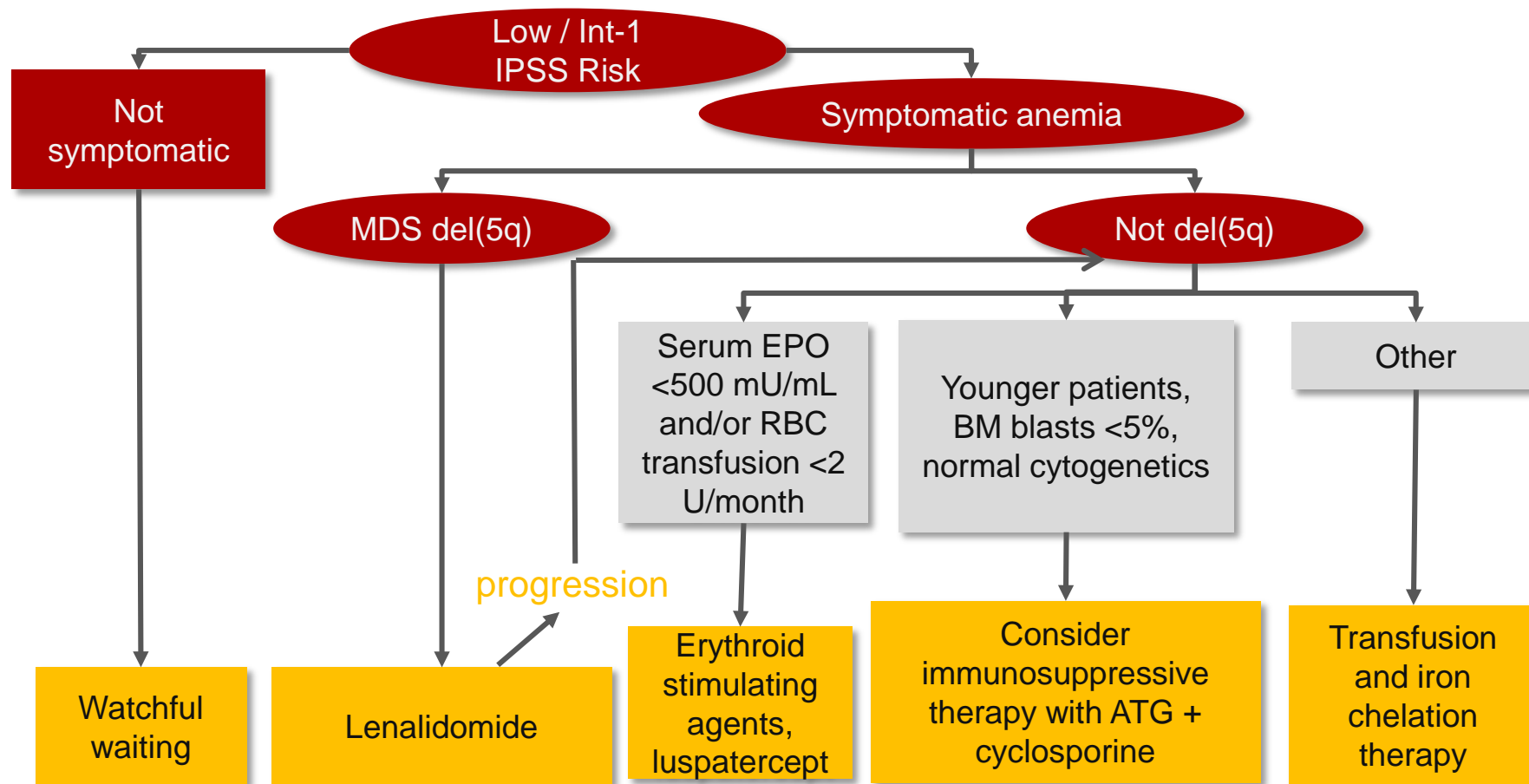
Priorities in Low-Risk MDS



Priorities in High-Risk MDS

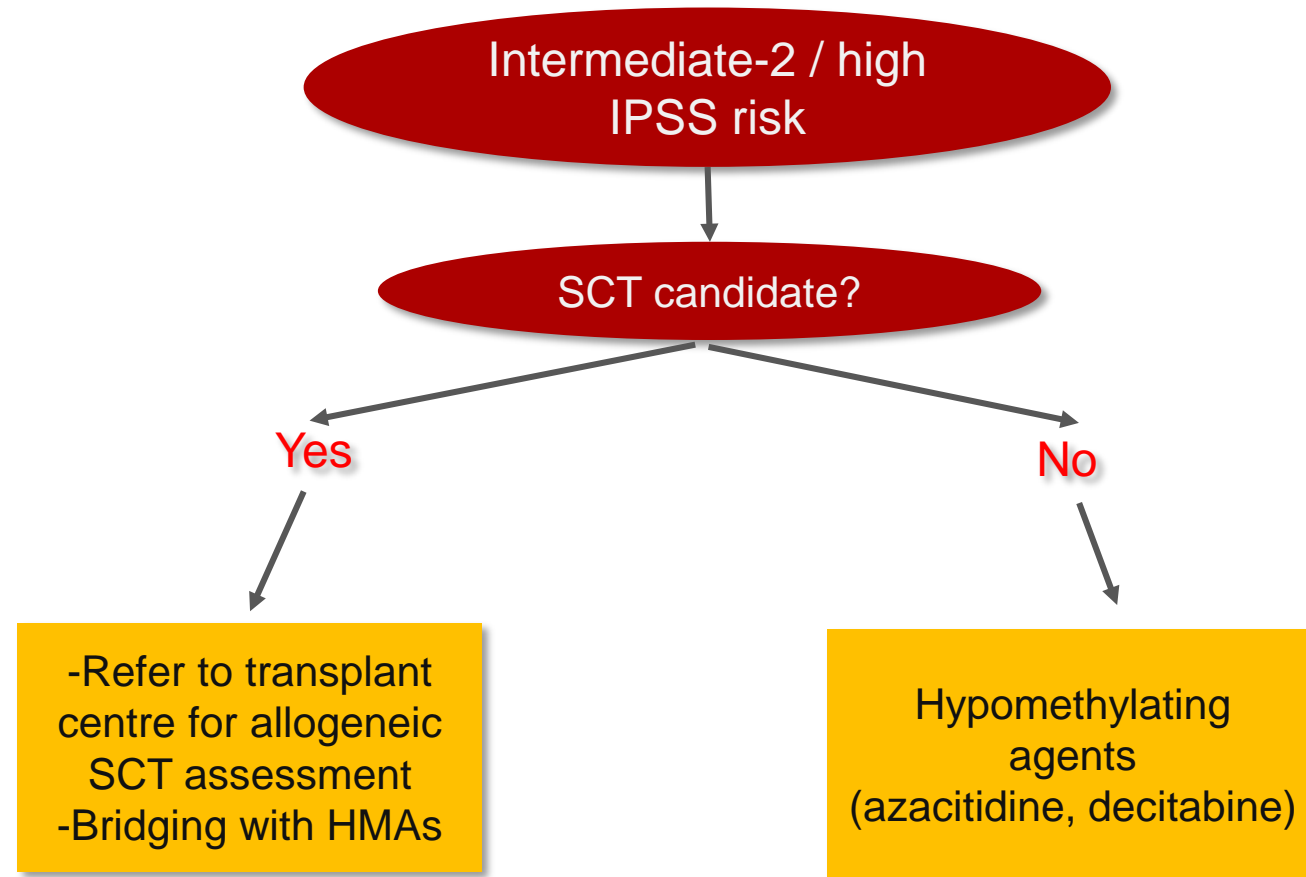


Recommendations for patients with low / intermediate-1 IPSS risk



ATG, anti-thymocyte globulin. BM, bone marrow. EPO, erythropoietin.
G-CSF, granulocyte colony-stimulating factor. IPSS, International Prognostic Scoring System. RBC, red blood cell

Recommendations for patients with intermediate-2 / high IPSS risk



SCT, stem cell transplant.

Between a rock and a hard place

Transplantation is currently the only curative therapy for myelodysplastic syndromes



**Median age at diagnosis is 65-70
Toxicity of transplantation can be prohibitive**

Questions?

