



Terapia della mielofibrosi idiopatica e secondaria- Inquadramento dell'argomento

Giovanni Barosi

Centro per lo Studio della Mielofibrosi

Fondazione IRCCS Policlinico S. Matteo, Pavia

Progetto Ematologia-Romagna. Cesena 16 settembre 2017

Primary myelofibrosis today (2017)

- ✓ A changing disease
- ✓ A molecular disease
- ✓ A disease in search of a new treatment paradigm

Italian diagnostic criteria for myelofibrosis (*Barosi et al, BJH 1998*)

Necessary criteria

1. *Diffuse bone marrow fibrosis*
2. *Absence of Philadelphia chromosome or BCR-ABL rearrangement*

Optional criteria

1. *Splenomegaly of any grade;*
2. *Anisopoikilocytosis with tear-drop erythrocytes*
3. *Presence of circulating immature myeloid cells*
4. *Presence of circulating erythroblasts*
5. *Presence of clusters of megakaryoblasts and anomalous megakaryocytes in bone marrow sections*
6. *Myeloid metaplasia.*

WHO diagnostic criteria for myelofibrosis (2016)

Prefibrotic/early stage

Major criteria

1. *Megakaryocyte proliferation and atypia, without reticulin fibrosis grade >1, accompanied by increased age-adjusted BM cellularity, granulocytic proliferation, and (often) decreased erythropoiesis*
2. *Not meeting WHO criteria for other myeloid neoplasms*
3. *JAK2, CALR, or MPL mutation, or presence of another clonal marker*

Minor criteria

1. *Presence of ≥ 1 of the following: anemia not attributed to comorbid condition; leukocytosis $\geq 11 \times 10^9/L$, palpable splenomegaly; lactate dehydrogenase level above the upper limit of the institutional reference range*

Fibrotic stage

Major criteria

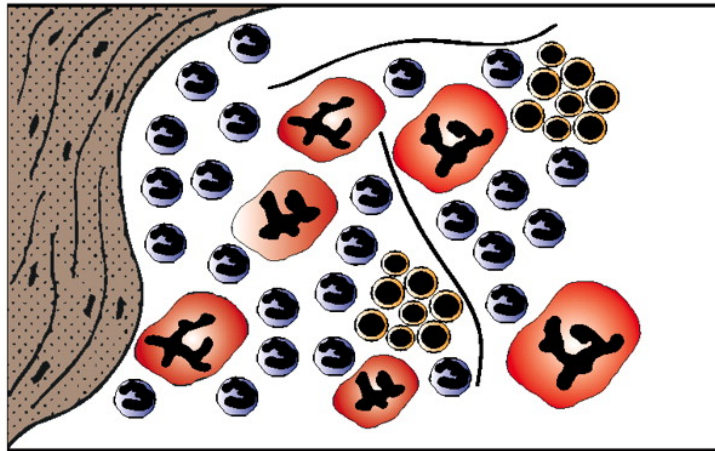
1. *Megakaryocyte proliferation and atypia, accompanied by reticulin and/or collagen fibrosis grade 2 or 3*
2. *Not meeting WHO criteria for other myeloid neoplasms*
3. *JAK2, CALR, or MPL mutation, or presence of another clonal marker*

Minor criteria

1. *Presence of ≥ 1 of the following: anemia not attributed to comorbid condition; leukocytosis $\geq 11 \times 10^9/L$, palpable splenomegaly; lactate dehydrogenase level above the upper limit of the institutional reference range*

ET

- no or only slight increase in age-matched cellularity
- no significant increase in granulo- and erythropoiesis
- prominent large to giant mature megakaryocytes with hyperlobulated or deeply folded nuclei, dispersed or loosely clustered in the marrow space
- no or very rarely minor increase in reticulin fibers



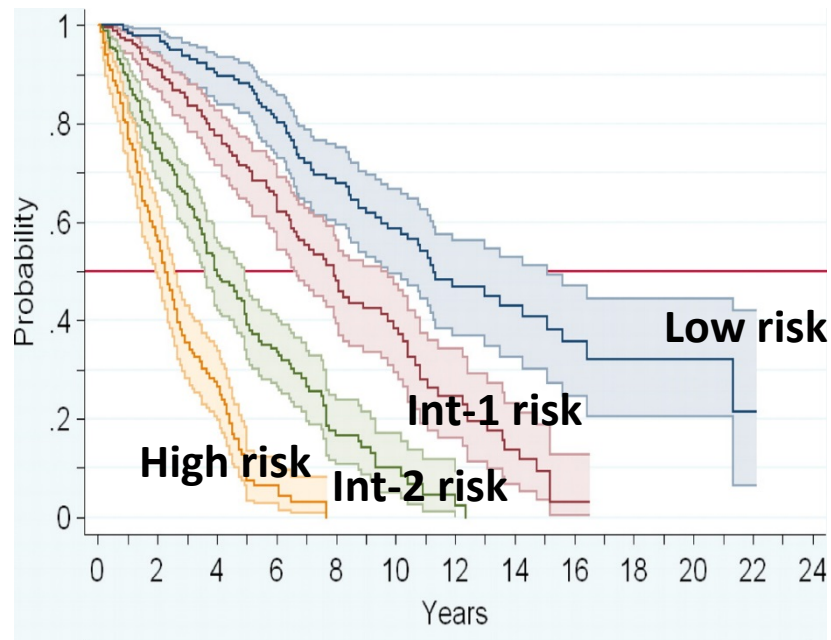
PMF (early-prefibrotic stage)

- marked increase in age-matched cellularity
- pronounced proliferation of granulopoiesis and reduction of erythroid precursors
- dense or loose clustering and frequent endosteal translocation of medium sized to giant megakaryocytes showing hyperchromatic, hypolobulated, bulbous, or irregularly folded nuclei and an aberrant nuclear/cytoplasmic ratio
- no or no significant increase in reticulin fibers

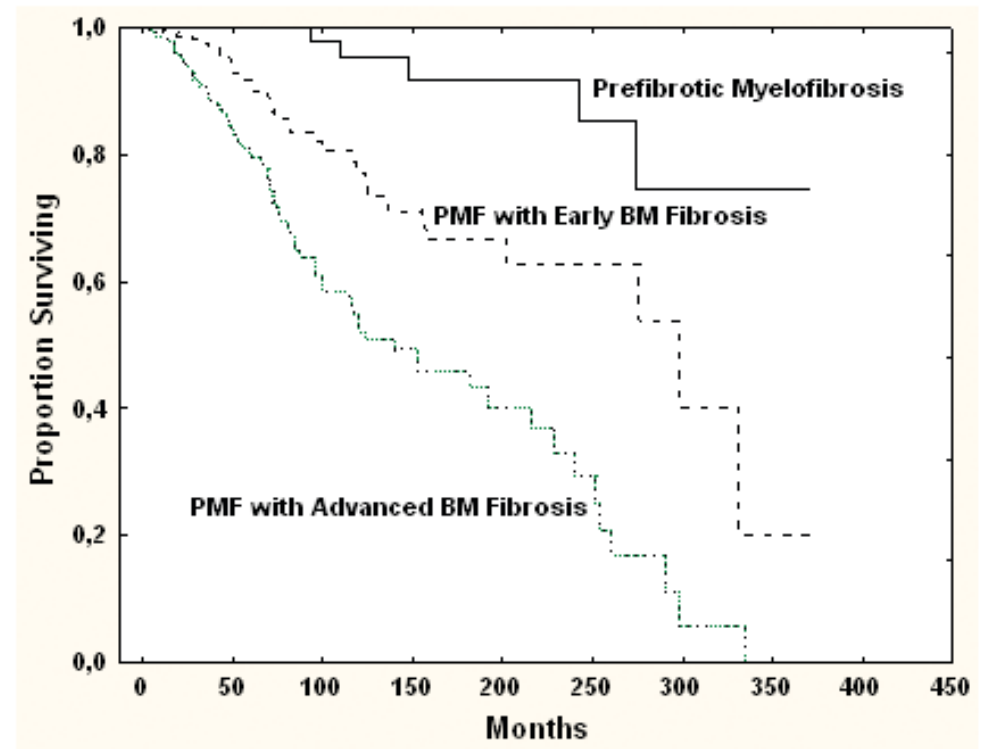


● Megakaryopoiesis; ● Granulopoiesis; ● Erythropoiesis; ʘ Reticulin fibers

Overall survival of PMF

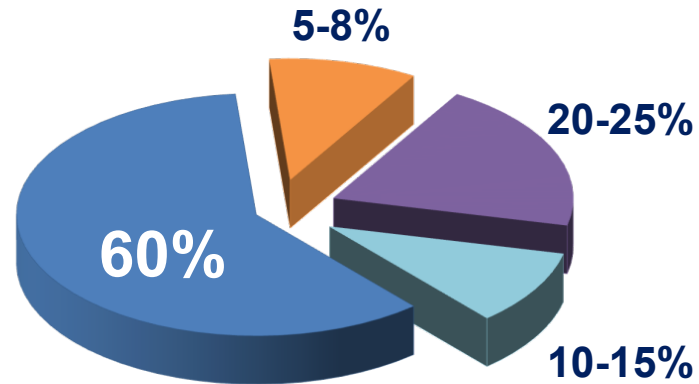


Cervantes et al. Blood 2009



Barosi et al. Plos One 2012

Phenotypic Driver Mutations in MPN



 **JAK2 V617F**

 **MPL (W515X)**

 **CALR mut**

 **Unknown ("Triple Negative")**

Additional, Not-driver, Somatic Mutations

Gene	Chromosome location	PV (%)	ET (%)	MF (%)	Blast phase (%)
<i>TET2</i>	4q24	10-16	4-5	7-17	17-32
<i>IDH1/2</i>	2q33.3 / 15q26.1	2	1	4	9-22
<i>DNMT3A</i>	2p23	3-7	<1	2-15	14-17
<i>EZH2</i>	7q36.1	3	<1	7-13	---
<i>ASXL1</i>	20q11.1	2-7	0-3	13-32	18-33
<i>SRSF2</i>	17q25.1	---	---	≈15%	≈20%
<i>SF3B1</i>	2q33.1	---	---	7%	---
<i>CBL</i>	11q23.3	rare	rare	6%	---
<i>TP53</i>	17p13.1	---	---	4%	27%
<i>U2AF1</i>	21q22.3	---	---	16%	---

Vainchenker W et al, Blood. 2011; 18;118(7):1723-35;
 Vannucchi AM et al, Leukemia 2013; 27:1861-9.

Clinical Scores for Risk Stratification in PMF

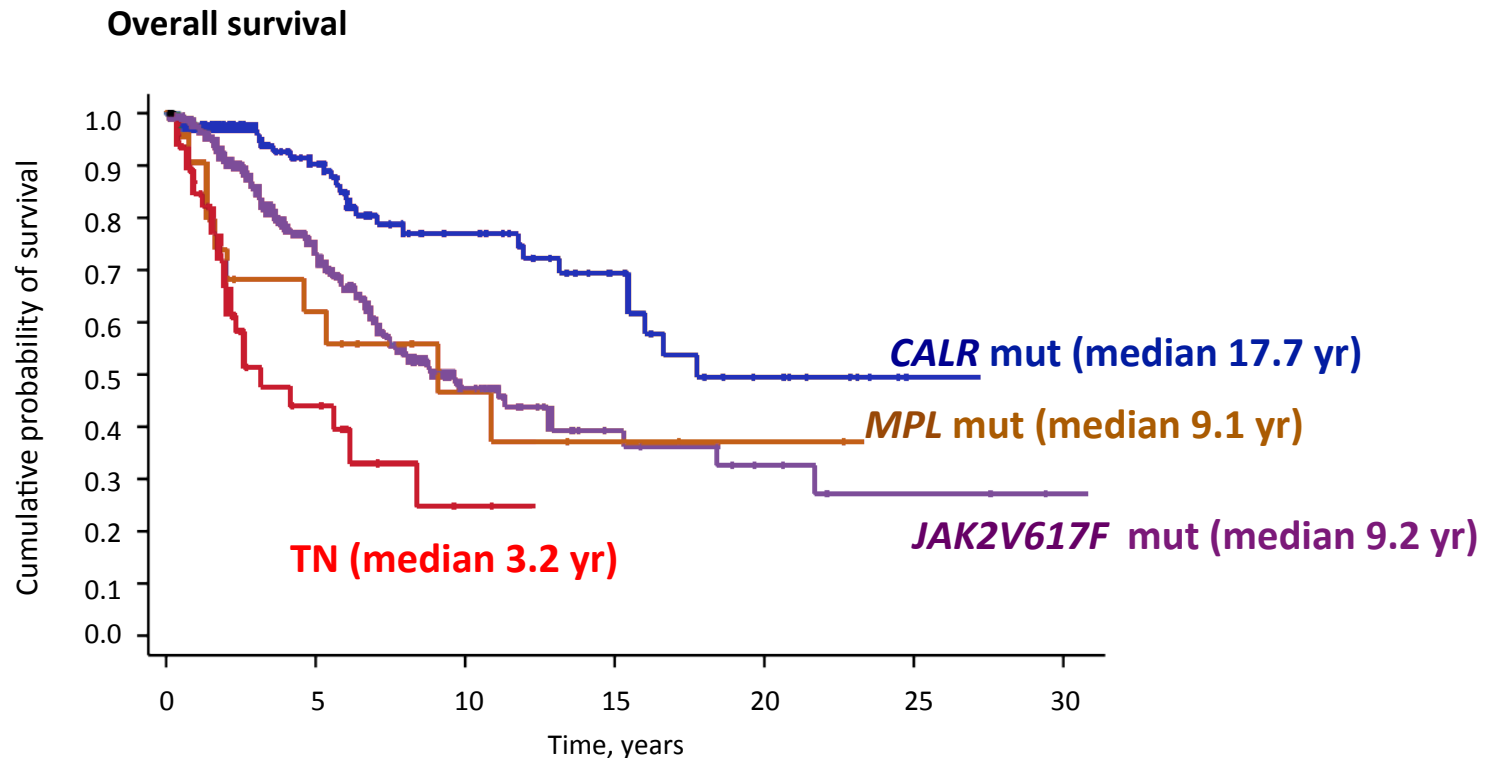
Variable	IPSS	DIPSS	DIPSS-plus
Age >65 y	√	√	If DIPSS: Low= 0 Int-1= 1 Int-2=2 High= 3
Constitutional symptoms	√	√	
Hemoglobin <10 g/dL	√	√	
Leukocyte count >25x10 ⁹ /L	√	√	
Circulating blasts ≥1%	√	√	
Platelet count <100x10 ⁹ /L			√
RBC transfusion need			√
Unfavorable karyotype +8,-7/7q-,i(17q),inv(3), -5/5q-,12p-, 11q23 rearr.			√

Cervantes F, et al. *Blood*. 2009;113:2895-901

Passamonti F, et al. *Blood*. 2010; 115:1703-8

Gangat N, et al. *J Clin Oncol*. 2011; 29:392-7

Phenotype Driver Mutations Have a Strong Prognostic Impact in PMF



Study	Hazard Ratio	HR	95%-CI	W(fixed)
Andriakovics		6.30	[0.71; 55.45]	1.7%
Nangalia		1.00	[0.09; 11.59]	1.3%
Rumi		2.29	[1.58; 3.33]	57.6%
Tefferi		2.61	[1.66; 4.10]	39.3%
Fixed effect model		2.43	[1.83; 3.22]	100%

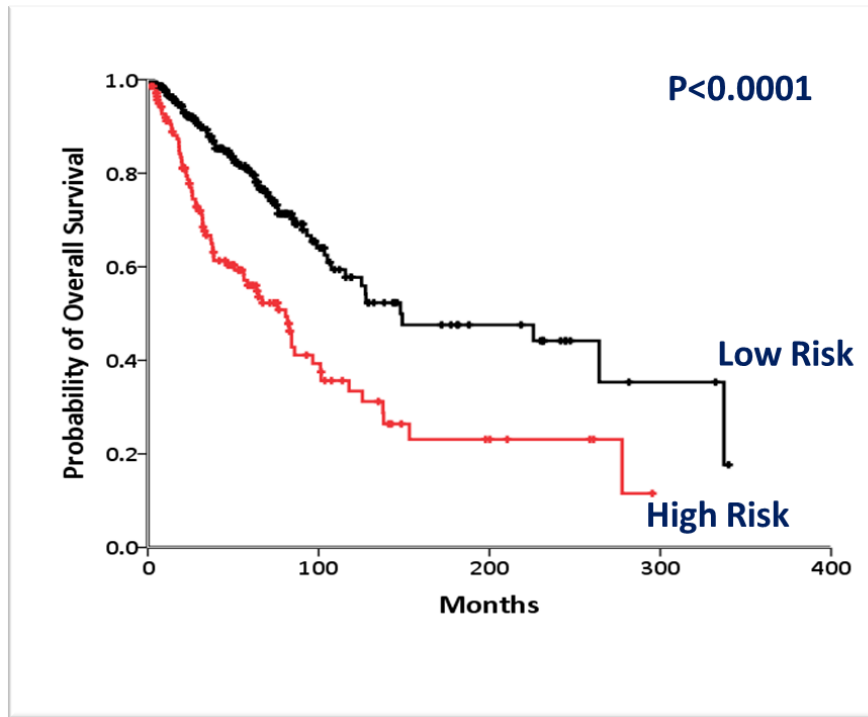
Heterogeneity: I-squared=0%, tau-squared=0, p=0.6942

JAK2 mutated patients had shorter overall survival compared with those CALR⁺ (Meta-analysis combined hazard ratio, 2.43; 95% CI, 1.83-3.22; P= < .001).

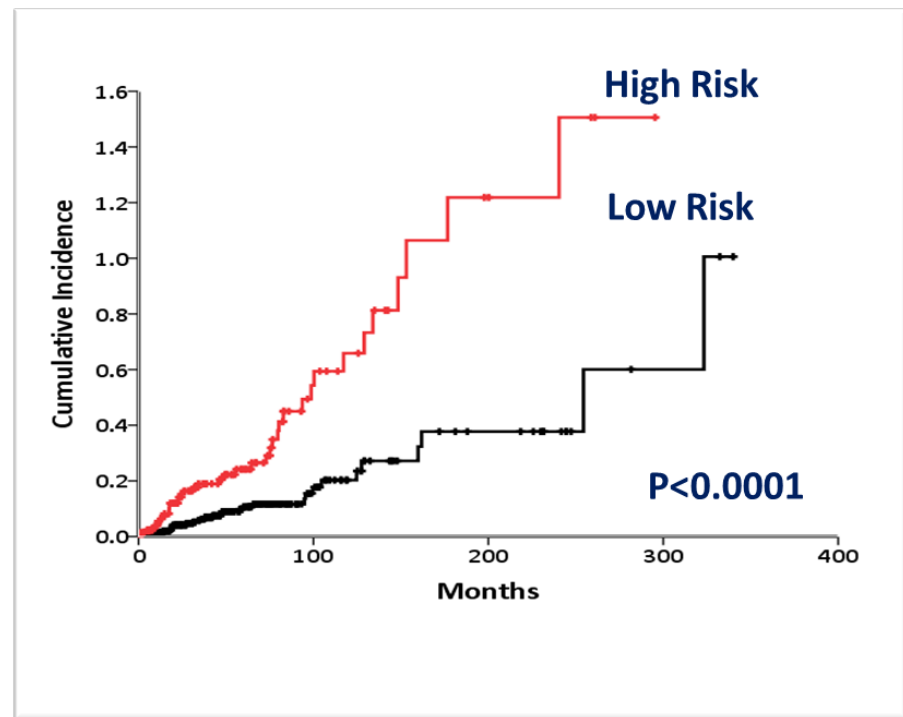
High Molecular Risk Prognostic Category

harboring ≥ 1 mutation in any one of *ASXL1*, *EZH2*, *SRSF2*, *IDH1/2*

Overall Survival



Blast Transformation



- A HMR status is associated with reduced OS and increased risk of blast transformation in PMF patients independent of IPSS/DIPPS-plus

Genetically driven prognostic model in PMF

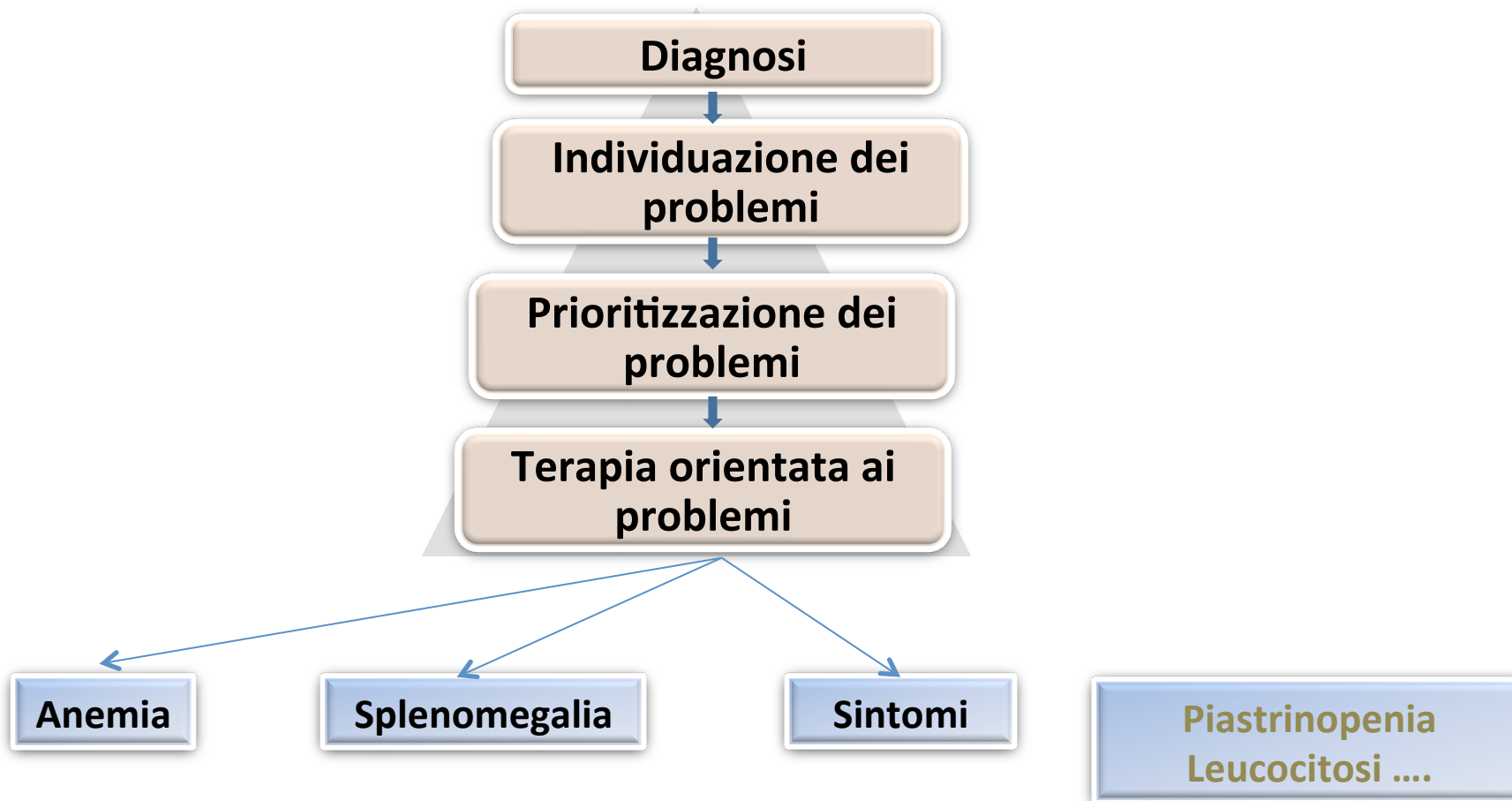
	MIPSS	GIPSS
Age >65	1.5	2
Constitutional symptoms	0.5	No
Hemoglobin <10 g/dL	0.5	No
Platelets < 200 x10⁹/L	1	No
Triple negative	1.5	2
JAK2 or MPL mutation	0.5	2
ASXL1 mutation	0.5	1
SRSF2 mutation	0.5	1
CALR Type 2-Type 2 like	No	2
Unfavorable cytogenetics	No	3 for very high risk; 2 for high risk

MIPSS = Mutation-Enhanced International Prognostic Scoring System (Vannucci et al, Blood 2014;124:405)

HMR: How Many Patients Would be Reclassified?

IPSS Risk Categories	<i>ASXL1</i> N. (%)	<i>EZH2</i> N. (%)	<i>SRSF2</i> N.(%)	<i>IDHs</i> N. (%)	N (%) Of HMR patients
LOW	24/162 (14.8%)	6/165 (3.6%)	7/151 (4.6%)	2/157 (1.3%)	35/166 (21.1%)
INT- 1	28/142 (19.7%)	6/143 (4.2%)	6/136 (4.4%)	6/142 (4.2%)	34 /146 (23.4%)
INT- 2	23/100 (23.0%)	4/99 (4.0%)	9/97 (9.3%)	2/96 (2.1%)	31 /104 (29.8%)
HIGH	27/65 (41.5%)	8/66 (12.1%)	16/63 (25.4%)	1/60 (1.7%)	39/68 (57.3%)

La terapia della mielofibrosi



How to change the therapy paradigm of MF?

- ✓ Targeting new molecular pathways/mechanisms
- ✓ Extending the use of allo-SCT
- ✓ Individualizing the use of available therapies

La medicina di precisione – un nuovo paradigma

Strategie di prevenzione e trattamento che considerano la individuale variabilità genetica e molecolare