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**Settima  
Giornata Fiorentina  
dedicata ai pazienti con  
malattie mieloproliferative  
croniche**

**Sabato 13 Maggio 2017**

**CRIMM**  
**Centro di Ricerca e Innovazione per le**  
**Malattie Mieloproliferative**  
**AOU Careggi**



UNIVERSITÀ  
DEGLI STUDI  
FIRENZE



## **Intro alla patologia**

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## La mielofibrosi

- ✓ Una malattia in evoluzione
- ✓ Una malattia (qualche volta) difficile da diagnosticare
- ✓ Una malattia molecolare
- ✓ Una malattia dalle scelte (terapeutiche) condivise

# **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

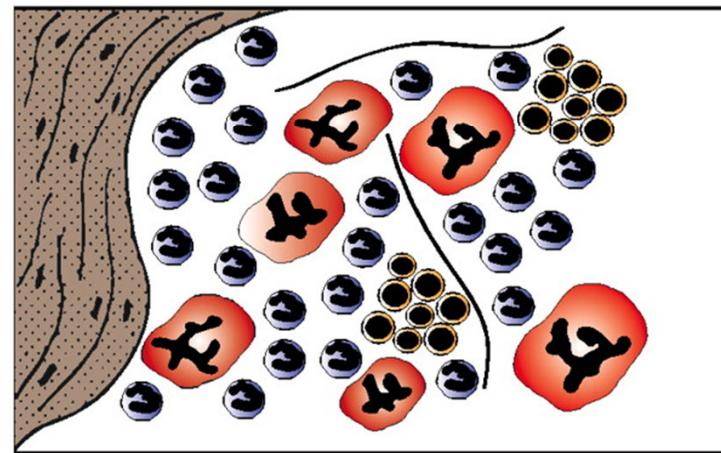
1. *Megakaryocyte proliferation and atypia, accompanied by reticulin and/or collagen fibrosis grade 2 or 3*
2. *Not meeting WHO criteria for other myeloid neoplasms*
3. *JAK2, CALR, or MPL mutation, or presence of another clonal marker*

### 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*

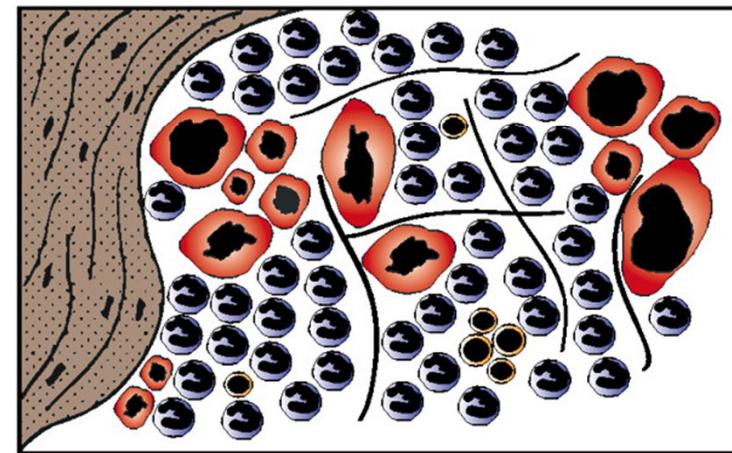
## ET

- 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



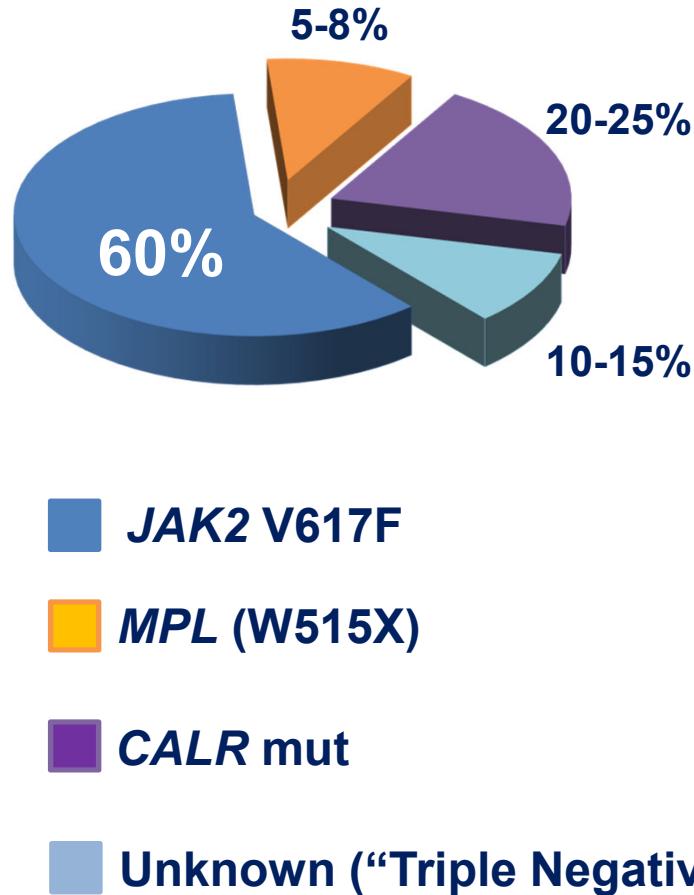
## PMF (early-prefibrotic stage)

- 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; ✕ Reticulin fibers

# Phenotypic Driver Mutations in MPN



Klampfl T, et al. NEJM 2013;369(25):2379-90; Nangalia J, et al. NEJM 2013;369(25):2391-405.

# Additional, Not-driver, Somatic Mutations

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

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

# **La medicina di precisione – un nuovo paradigma**

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

# Medicina di precisione nella mielofibrosi

1. Usare i marcatori molecolari (mutazioni) per migliorare la prognosi
2. Usare i marcatori molecolari (mutazioni) per scegliere la terapia

# Clinical Scores for Risk Stratification in PMF

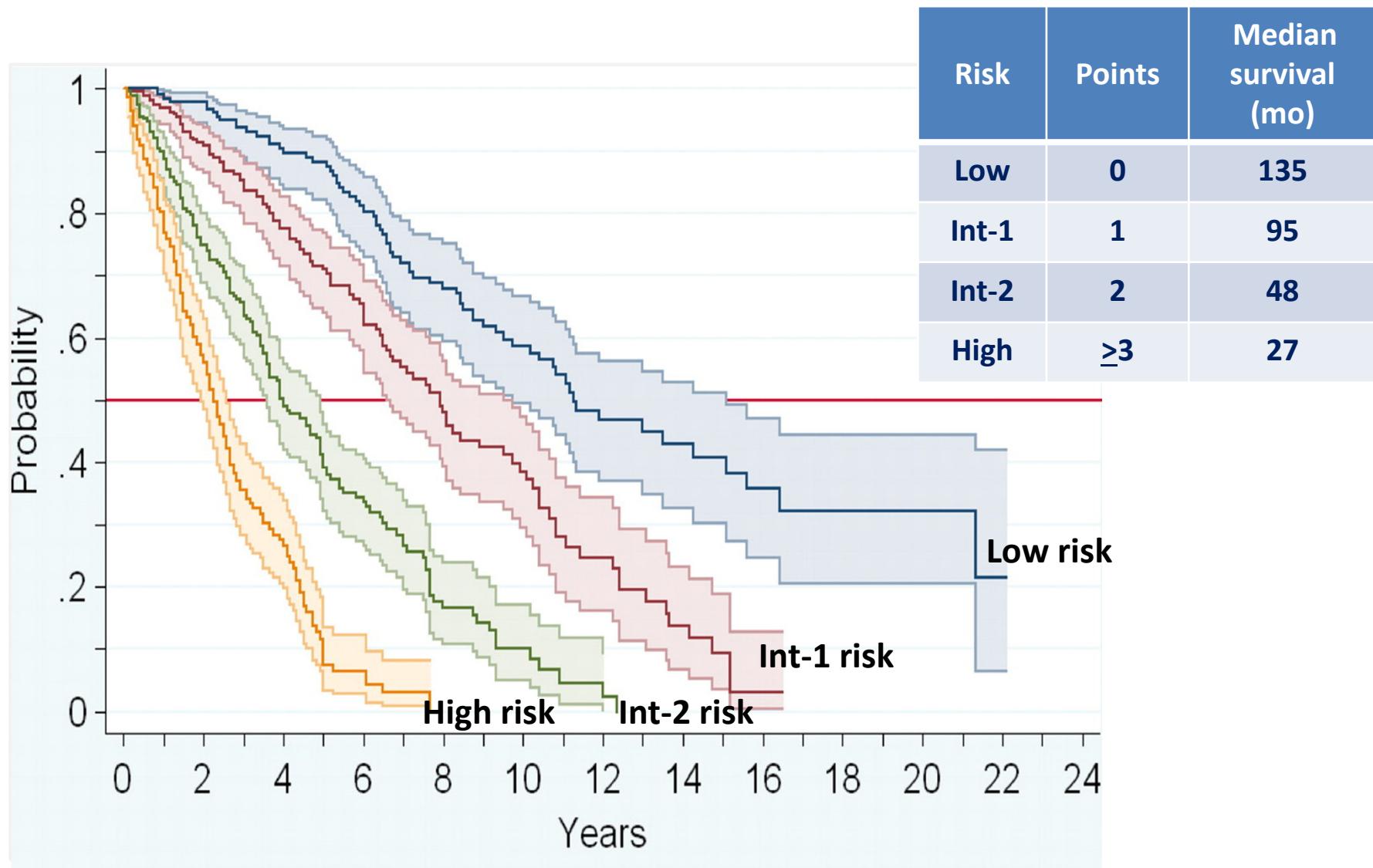
Variable	IPSS	DIPSS	DIPSS-plus
Age >65 y	✓	✓	If DIPSS:
Constitutional symptoms	✓	✓	Low= 0
Hemoglobin <10 g/dL	✓	✓	Int-1= 1
Leukocyte count >25x10 <sup>9</sup> /L	✓	✓	Int-2=2
Circulating blasts $\geq$ 1%	✓	✓	High= 3
Platelet count <100x10 <sup>9</sup> /L			✓
RBC transfusion need			✓
Unfavorable karyotype +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

# International Prognostic Scoring System-IPSS



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

# Genetically driven prognostic model in PMF

	MIPSS	GIPSS
<b>Age &gt;65</b>	1.5	2
<b>Constitutional symptoms</b>	0.5	No
<b>Hemoglobin &lt;10 g/dL</b>	0.5	No
<b>Platelets &lt; 200 x10<sup>9</sup>/L</b>	1	No
<b>Triple negative</b>	1.5	2
<b>JAK2 or MPL mutation</b>	0.5	2
<b>ASXL1 mutation</b>	0.5	1
<b>SRSF2 mutation</b>	0.5	1
<b>CALR Type 2-Type 2 like</b>	No	2
<b>Unfavorable cytogenetics</b>	No	3 for very high risk; 2 for high risk

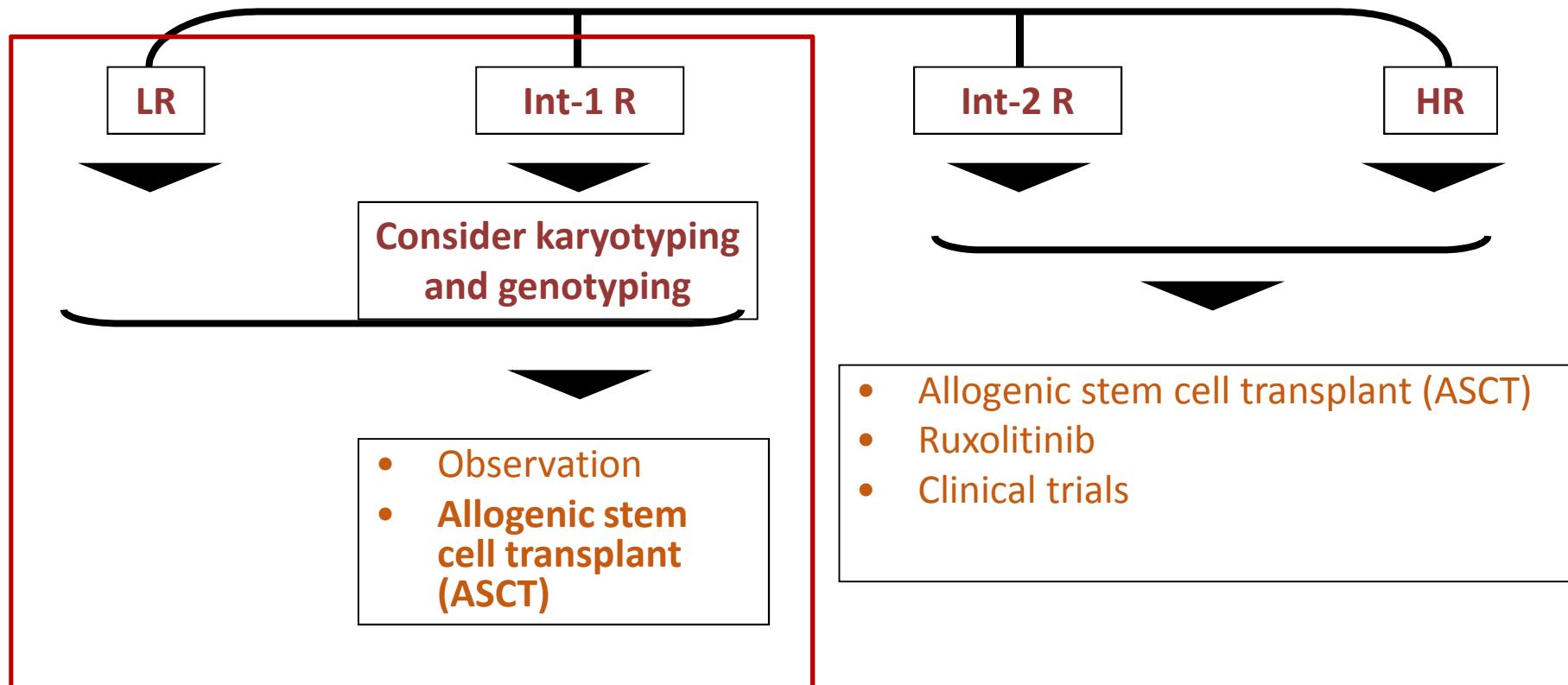
MIPSS = Mutation-Enhanced International Prognostic Scoring System (Vannucci et al, Blood 2014;124:405)

# HMR: How Many Patients Would be Reclassified?

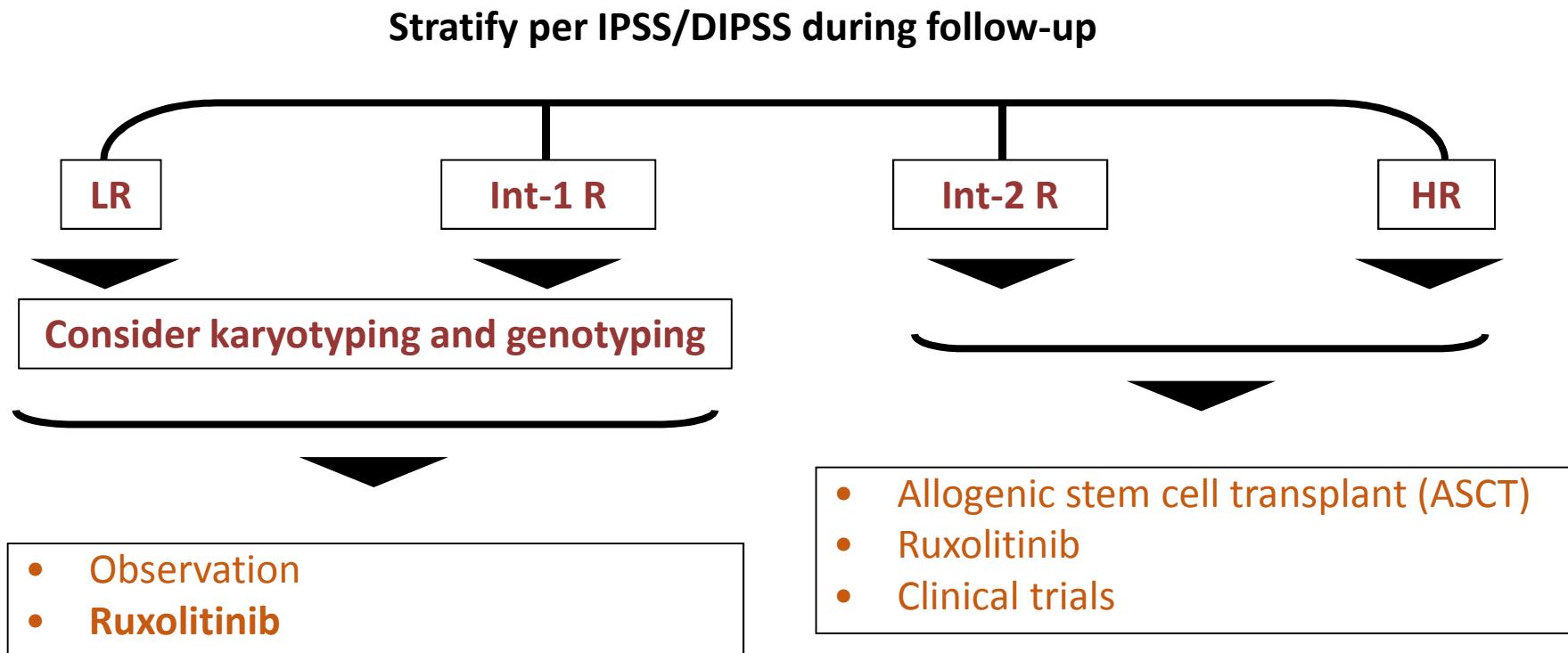
IPSS Risk Categories	ASXL1 N. (%)	EZH2 N. (%)	SRSF2 N. (%)	IDHs N. (%)	N (%) Of HMR patients
<b>LOW</b>	24/162 (14.8%)	6/165 (3.6%)	7/151 (4.6%)	2/157 (1.3%)	<b>35/166 (21.1%)</b>
<b>INT- 1</b>	28/142 (19.7%)	6/143 (4.2%)	6/136 (4.4%)	6/142 (4.2%)	<b>34 /146 (23.4%)</b>
<b>INT- 2</b>	23/100 (23.0%)	4/99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

# Personalized approach to MF: HSCT for DIPSS INT-1 disease

Stratify per IPSS/DIPSS during follow-up

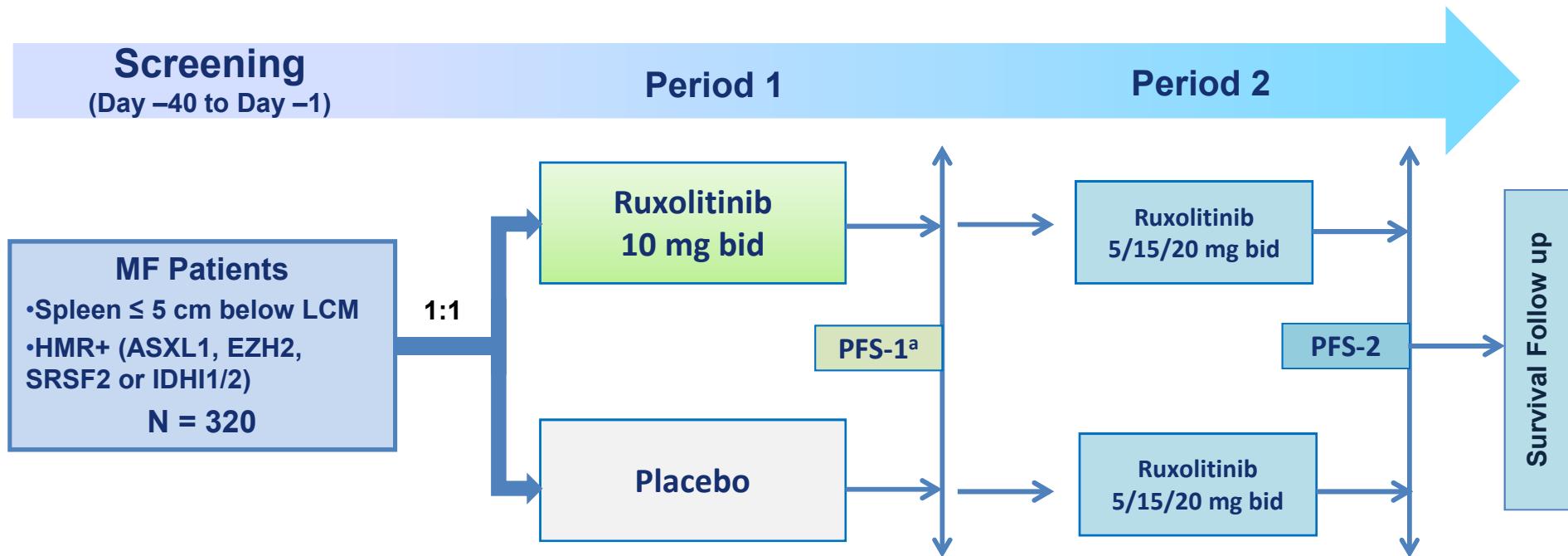


# Personalized approach to MF: Ruxolitinib for early phase disease



# Re-THINK: Trial Design

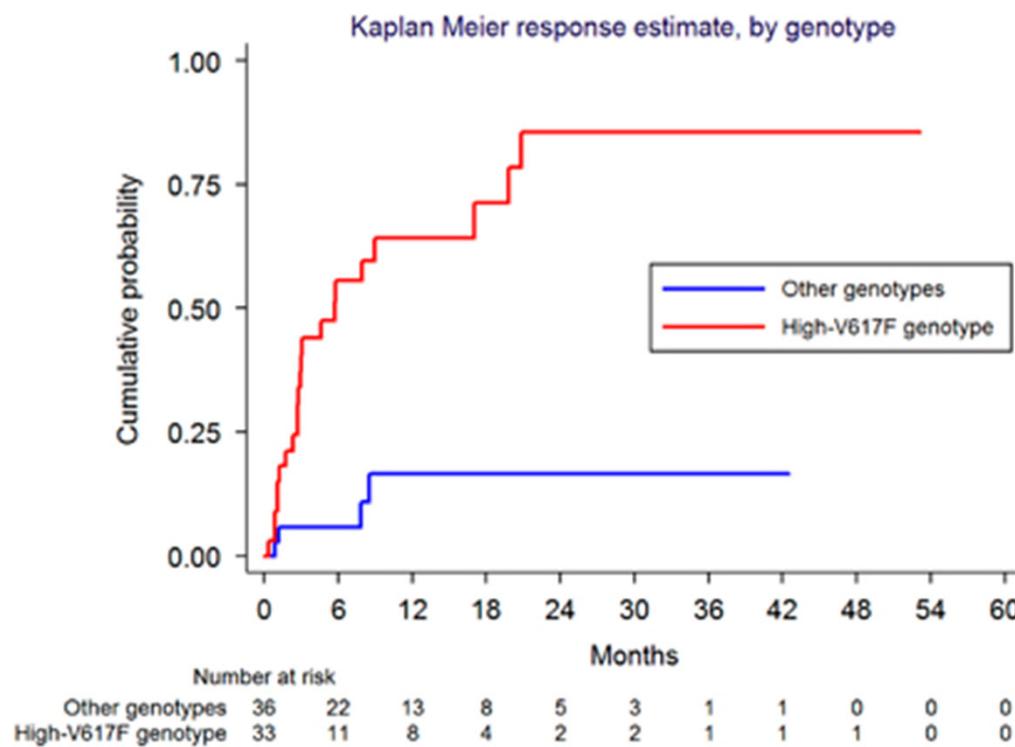
- ReTHINK is a randomized, double-blind, placebo-controlled, multi-center, phase 3 study of the efficacy and safety of ruxolitinib in patients with early MF and HMR mutations



<sup>a</sup> If progression is achieved by spleen or symptoms.

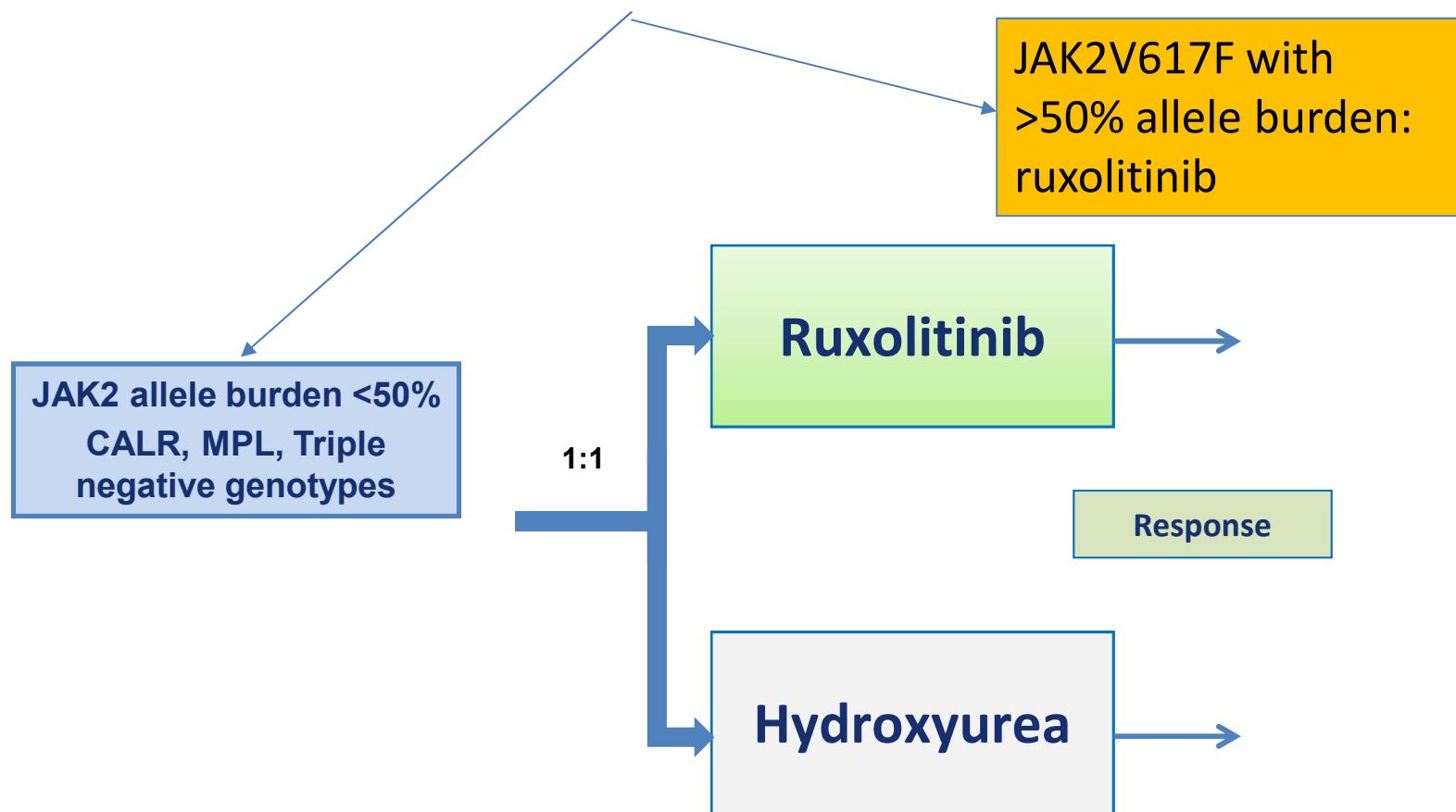
$JAK2^{V617F}$  allele burden  $\geq 50\%$  is associated with response to ruxolitinib in persons with MPN-associated myelofibrosis and splenomegaly requiring therapy

*Leukemia* (2016) 30, 1772–1775; doi:10.1038/leu.2016.45

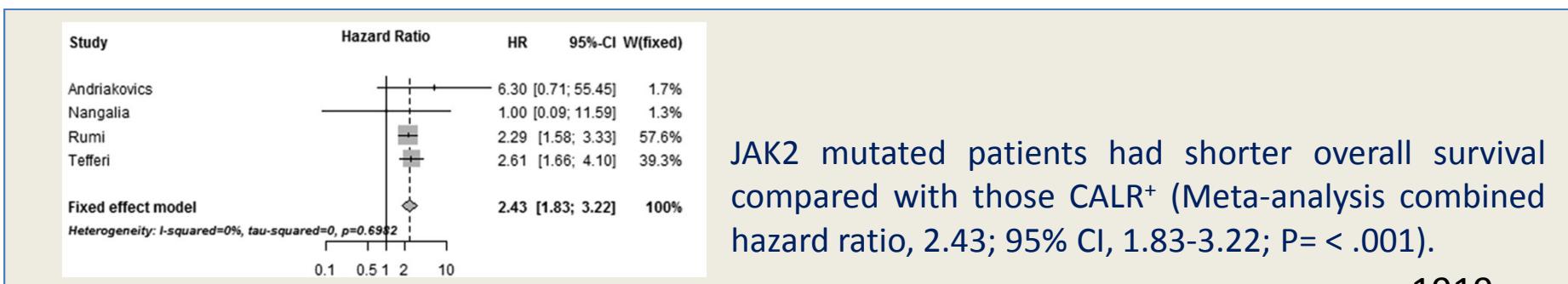
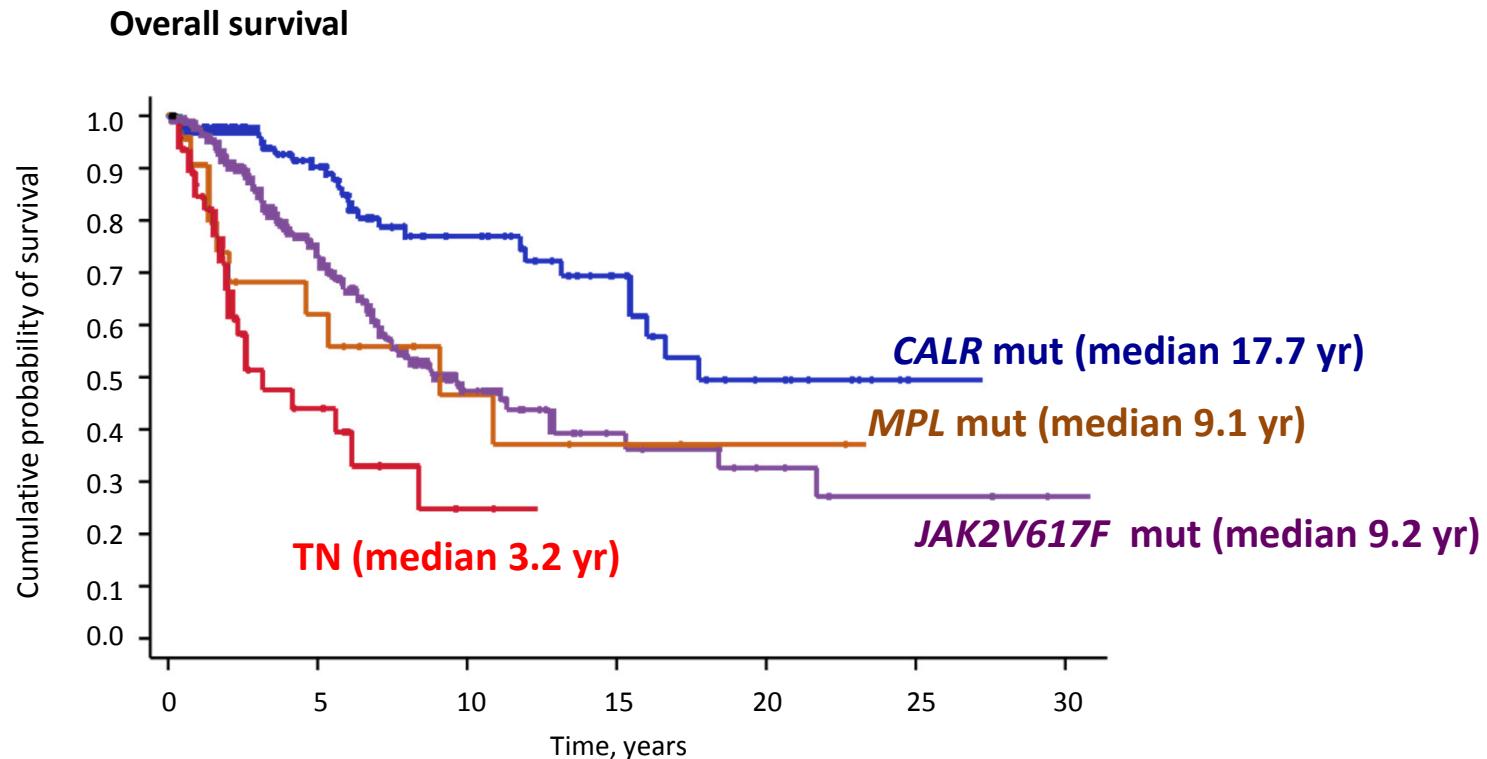


# Trial Design Hypothesis

Splenomegaly in need of therapy (treatment naive)



# Phenotype Driver Mutations Have a Strong Prognostic Impact in PMF

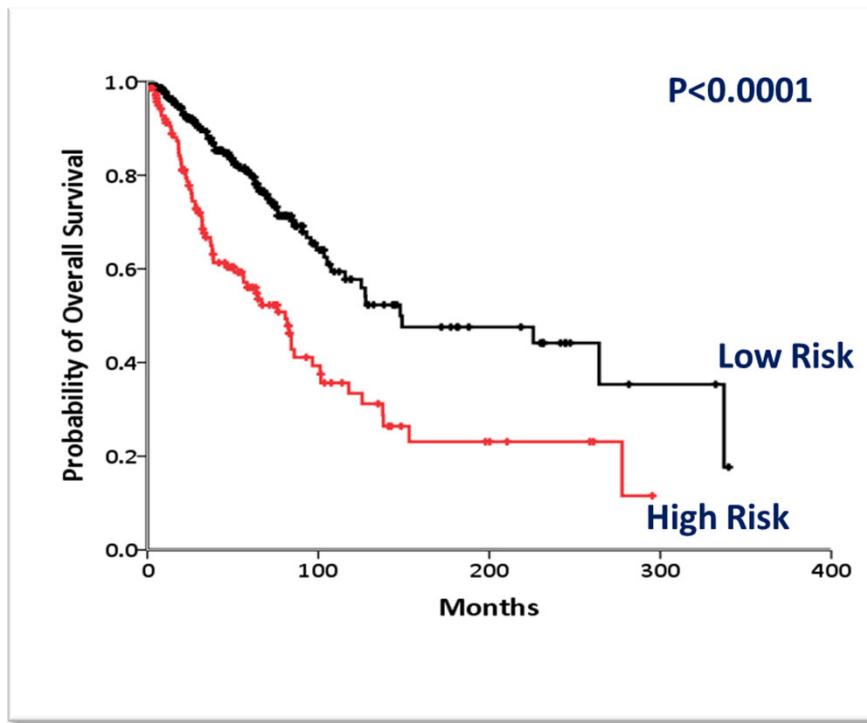


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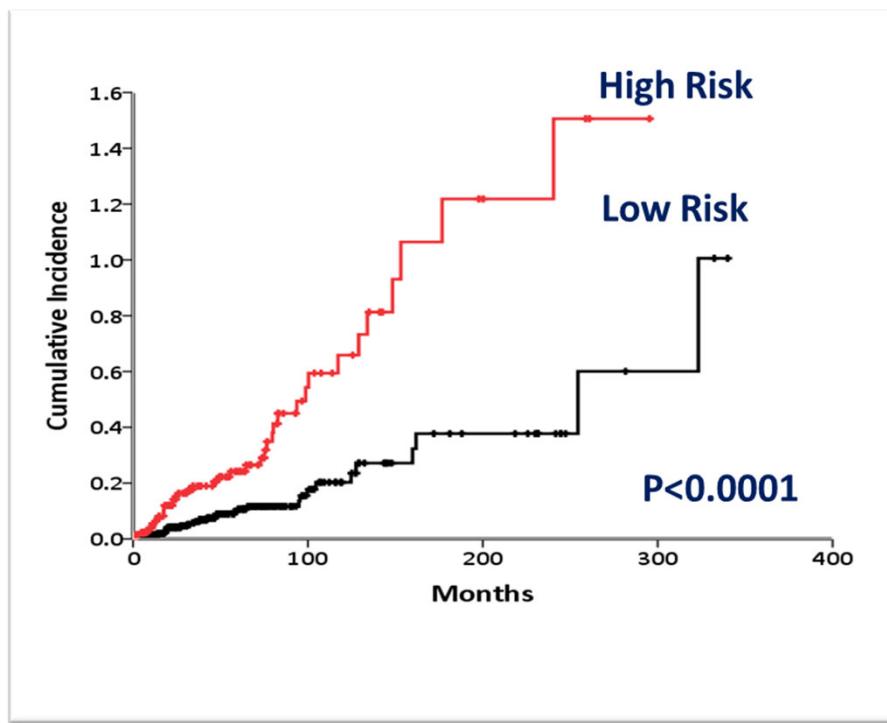
# High Molecular Risk Prognostic Category

harboring  $\geq 1$  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

# Trial Design Hypothesis

