

***PV ed ET alla diagnosi: score prognostici e  
fattori biologici***



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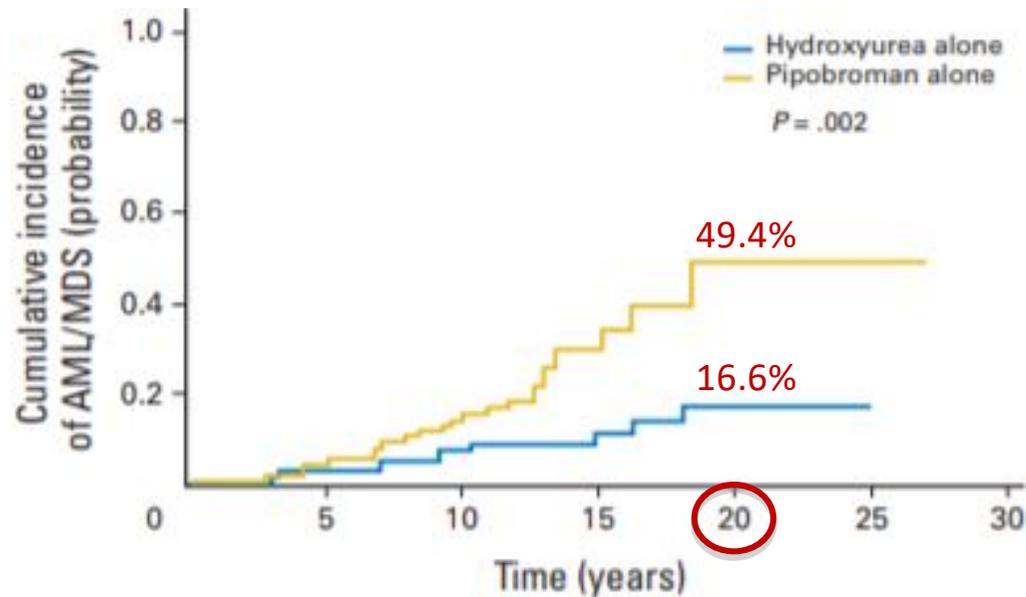
# Disclosures

- Speaker bureau: Novartis, Celgene
- Advisory boards: Celgene, Novartis, Roche, Janssen, Incyte, Abbvie.

## Events in PV, results from a meta-analysis

- Incidence rate of thrombosis:
  - Median value: 3.3% persons/year
  - 1.9% at 60 years with no history of thrombosis (low risk patients)
  - 6.8% at a median age of 80 years
- Risk of myelofibrosis:
  - 0.9%, 5% and 34% at 1, 5 and 10 years
  - The odds increase on average 6% for each year of age
- Risk of blast phase
  - 0.4%, stable incidence over time

# Pipobroman/Hydroxyurea and AML occurrence in PV



- PV patients treated with HU alone (N=94 for 12 yrs) and PB alone (N=130 for 9.5 yrs)

## Abnormal karyotype impacts on AML-free survival in PV

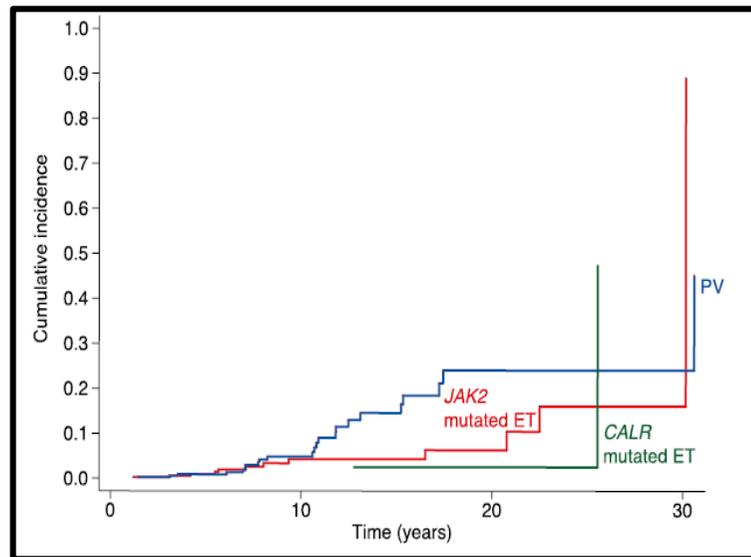
Predictors of leukemia-free survival	N	Univariate p-values	Age-adjusted p-values	Multivariate p-values (HR; 95% CI) <sup>a</sup>	Multivariate p-values (HR; 95% CI) <sup>b</sup>
Age	1545	0.0002			
Age > median (61y)	1545	0.005		0.007 (2.2;1.3-4.1)	0.004 (6.3;1.8-22)
Abnormal karyotype	631	0.03	0.03		0.03 (3.9;1.2-13.1)
WBC ≥ 15X10 <sup>9</sup> /L	1545	0.0003	0.0002	0.0004 (2.9;1.6-5.2)	0.01 (3.9;1.3-11.6)

Post-PV AML:  
50/1545 (3%) at  
a median time of  
10.8 yrs

<sup>a</sup> karyotype excluded as a covariate; <sup>b</sup> Karyotype included as a covariate

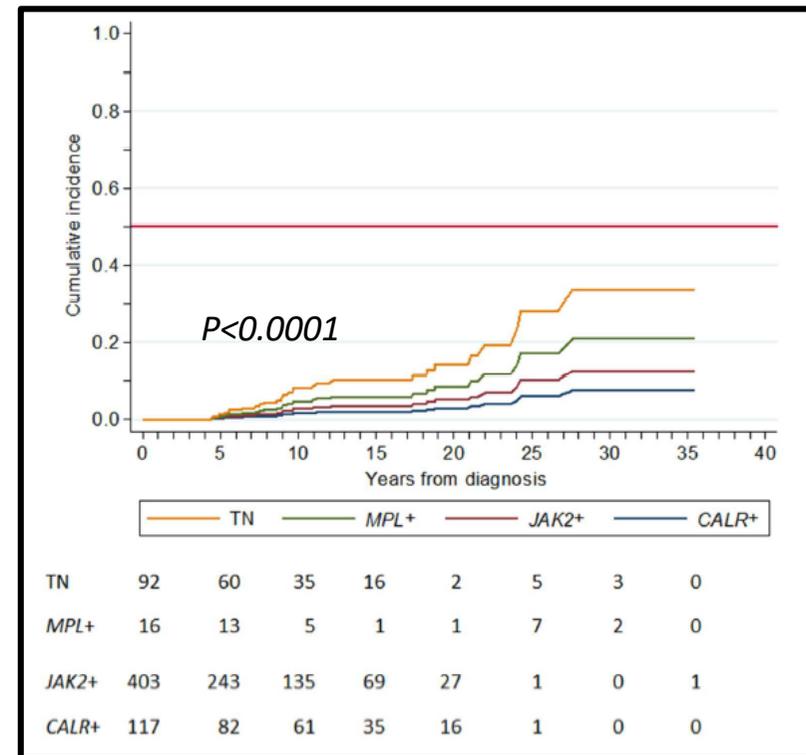
# Driver mutations' are predictive of post-MPN AML

Median follow-up: 5.2 years



*Rumi et al., Blood. 2014*

Median follow-up: 7.2 years



*Alvarez-Larran et al., BJH. 2017*

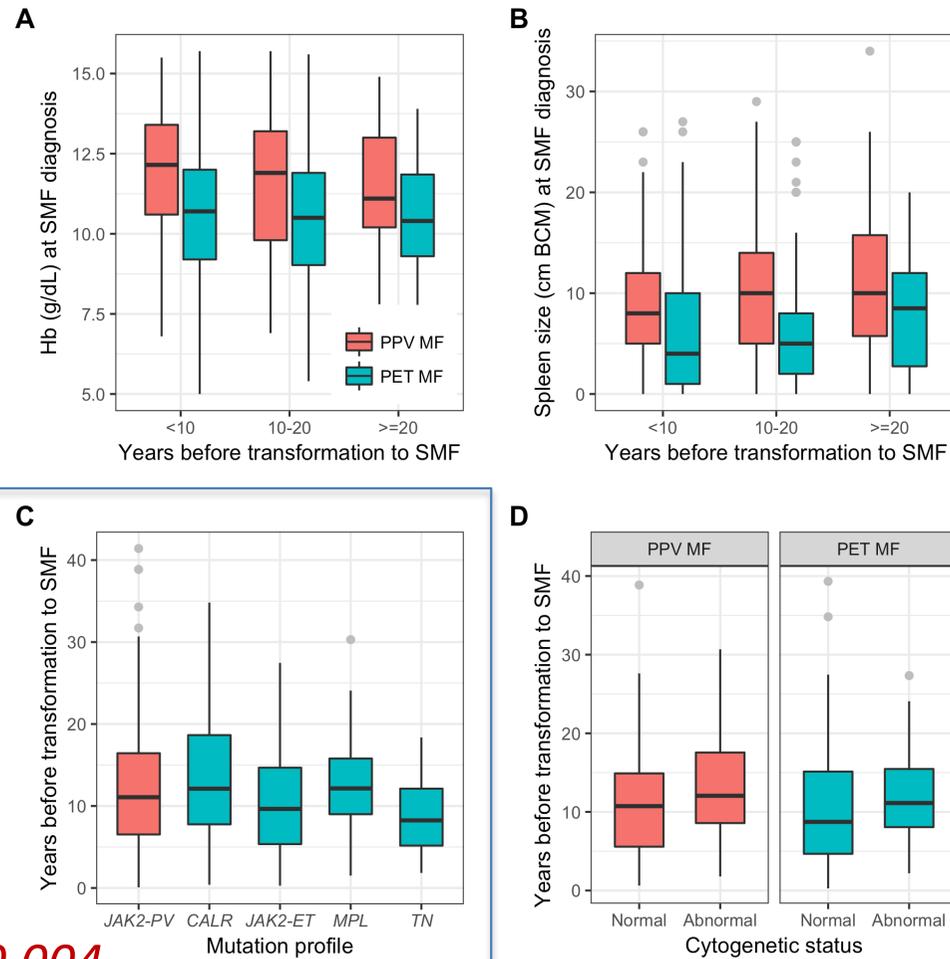
# Non-driver mutations' are predictive of post-MPN AML



→ HR 12.2 (95% CI 2.6-57.1,  $P=0.001$ ) in multivariate analysis

- 13.5% risk of AML transformation at 15 years from first sample
- *ASXL1*, *TP53*, *SRSF2*, *IDH1/2*, *RUNX1*

# Driver mutations and time to progression in PPV/PET MF



# NGS-enhanced prediction of survival in PV and ET

## ET

- *SRSF2/SF3B1* mutations (HR 2.8)
- Age >60 years (HR 6.7)
- Male sex (HR 1.8)

## PV

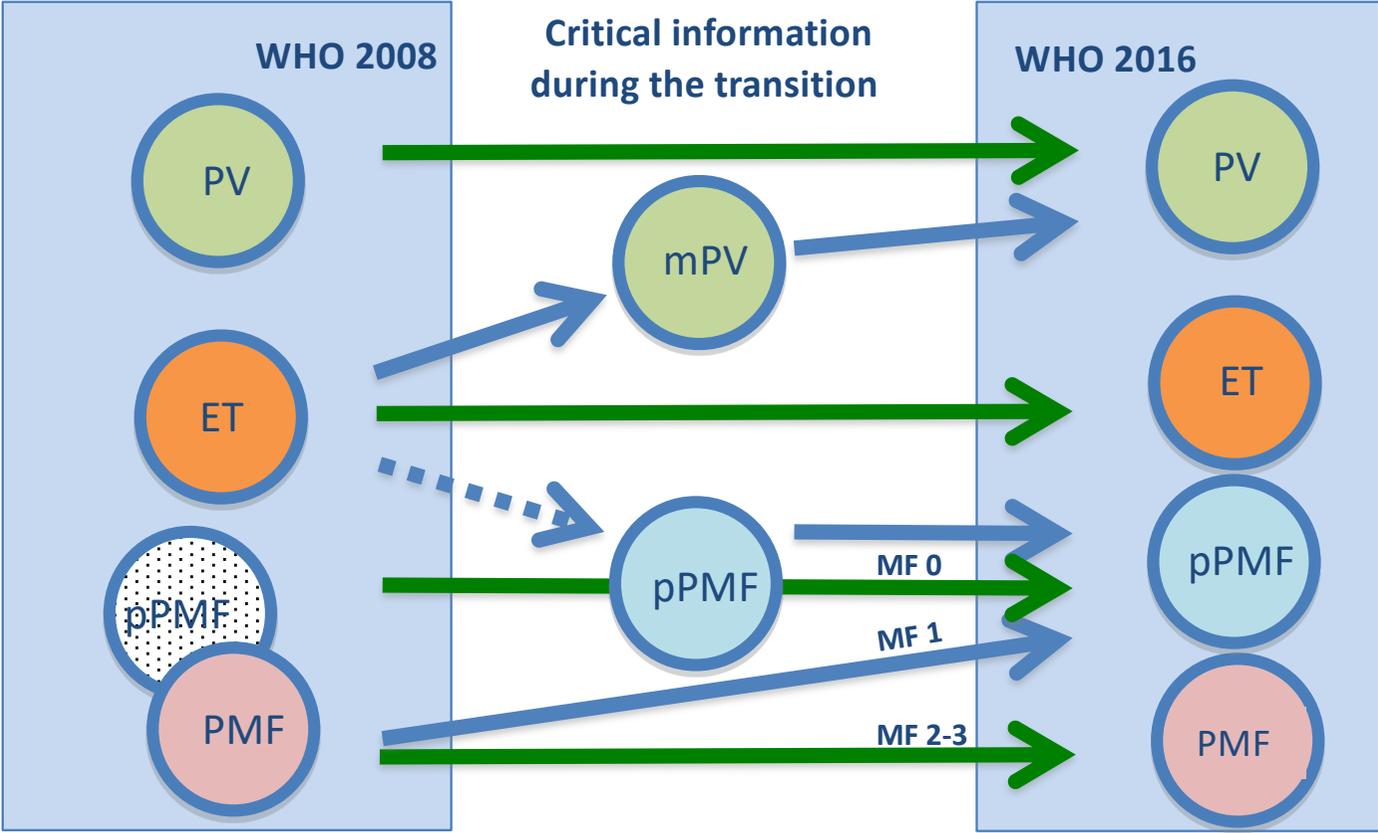
- *SRSF2* mutations (HR 7.0)
- Age >67 yrs (HR 5.7)
- Leukocytes  $\geq 11 \times 10^9/l$  (HR 2.4)
- Abnormal karyotype (HR 2.1)

***ET and PV (N=906)***



## ***Essential thrombocythemia***

# What is ET: WHO 2008 → 2017



# JAK2 assessment outperforms the two-tiered risk stratification for thrombosis: the IPSET-t model

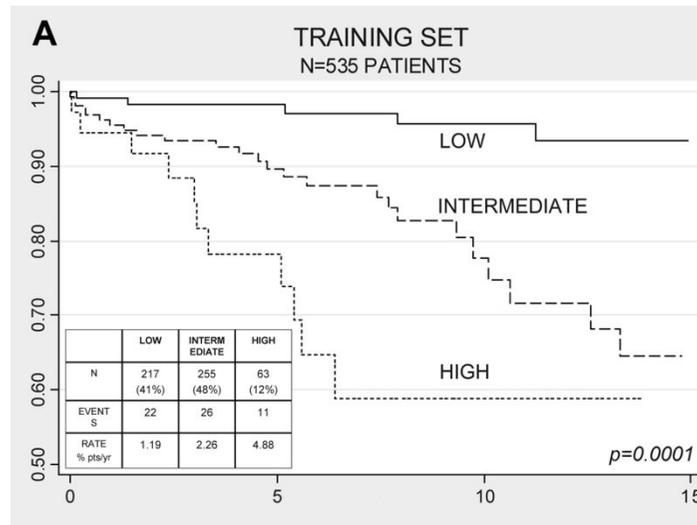
(891 WHO-defined ET, follow-up 6.2 years)

- **Prognostic factors**

- Age > 60 years (1 point)
- Thrombosis history (2 points)
- Cardiovascular risk (1 point)
- JAK2 V617F (2 points)

- **IPSET Model**

- Low risk (< 2 points): 1.03% p/y
- Intermediate risk (2 points): 2.35% p/y
- High risk (> 2 points): 3.56% p/y



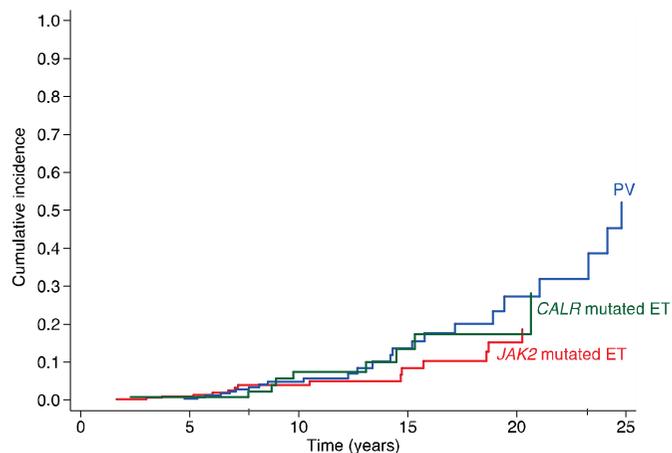
*Barbui et al. Blood 2012 Dec 20;120(26):5128-33*

**NO prognostic advantage of CALR mutation over IPSET-t in thrombosis prediction**

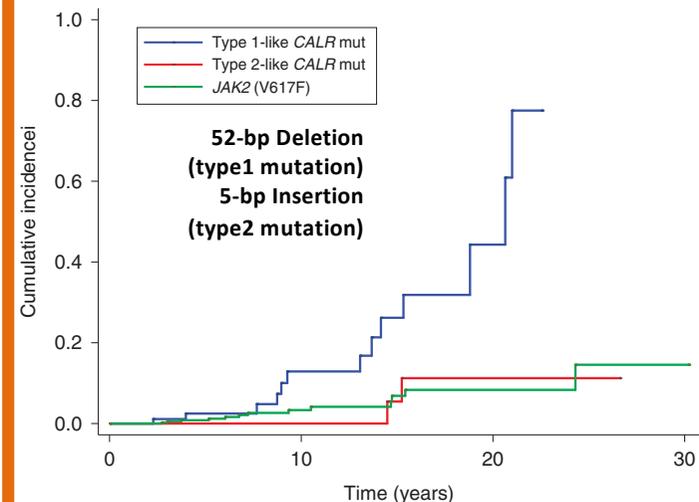
*Finazzi G. et al Blood. 2014 Oct 16;124(16):2611-2*

# Prediction of post-ET MF evolution

## Progression to post-ET MF per genotype



## Progression to post-ET MF per CALR



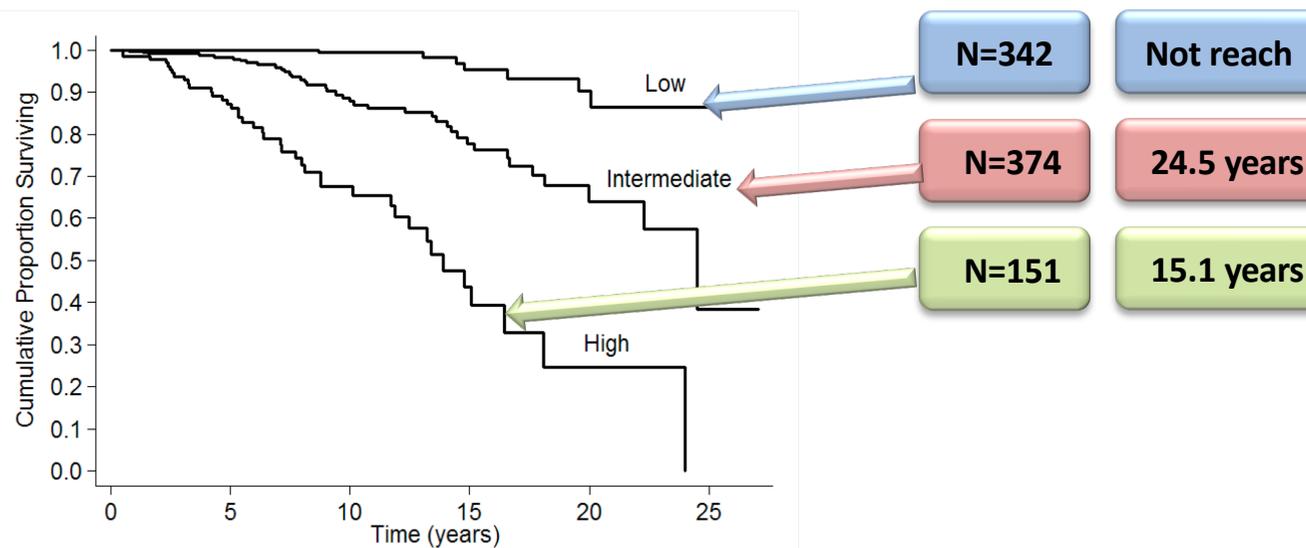
- In the MYSEC study on post-ET MF, *CALR* T1/T1-like genotype was more frequent (61%) than *CALR* T2/T2-like
- In the Mayo series, no specific association with post-ET MF have been disclosed among *CALR* sutypes

# Survival of WHO-based ET: the IPSET model

Risk factors	
Age > 60 years	2 points
Prior thrombosis	1 point
WBC > 11 x10 <sup>9</sup> /L	1 point

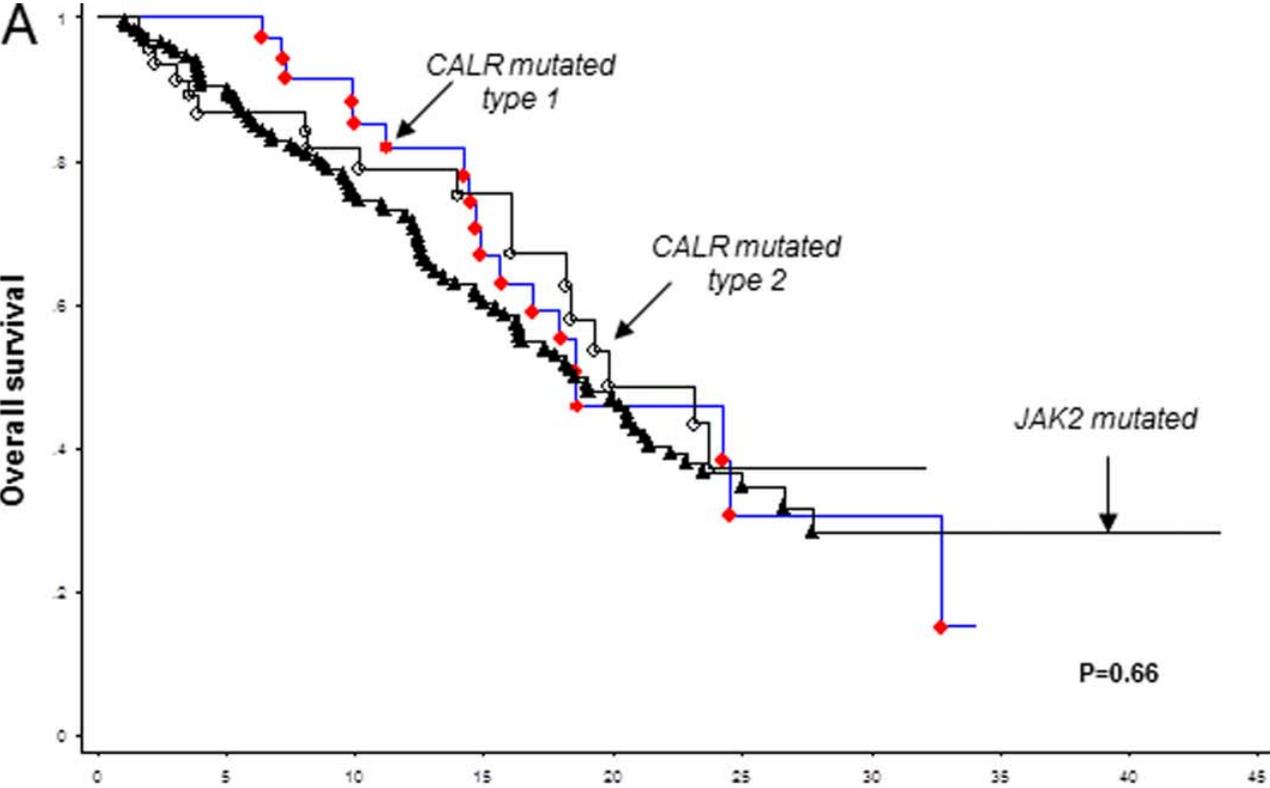


Risk Categories/score	
LR	0
Int-1	1-2
HR	3-4



Number at risk	0	5	10	15	20	25
Low	342	211	123	63	25	3
Intermediate	374	223	109	51	15	2
High	151	84	32	10	2	0

# No impact of genotype on survival in ET

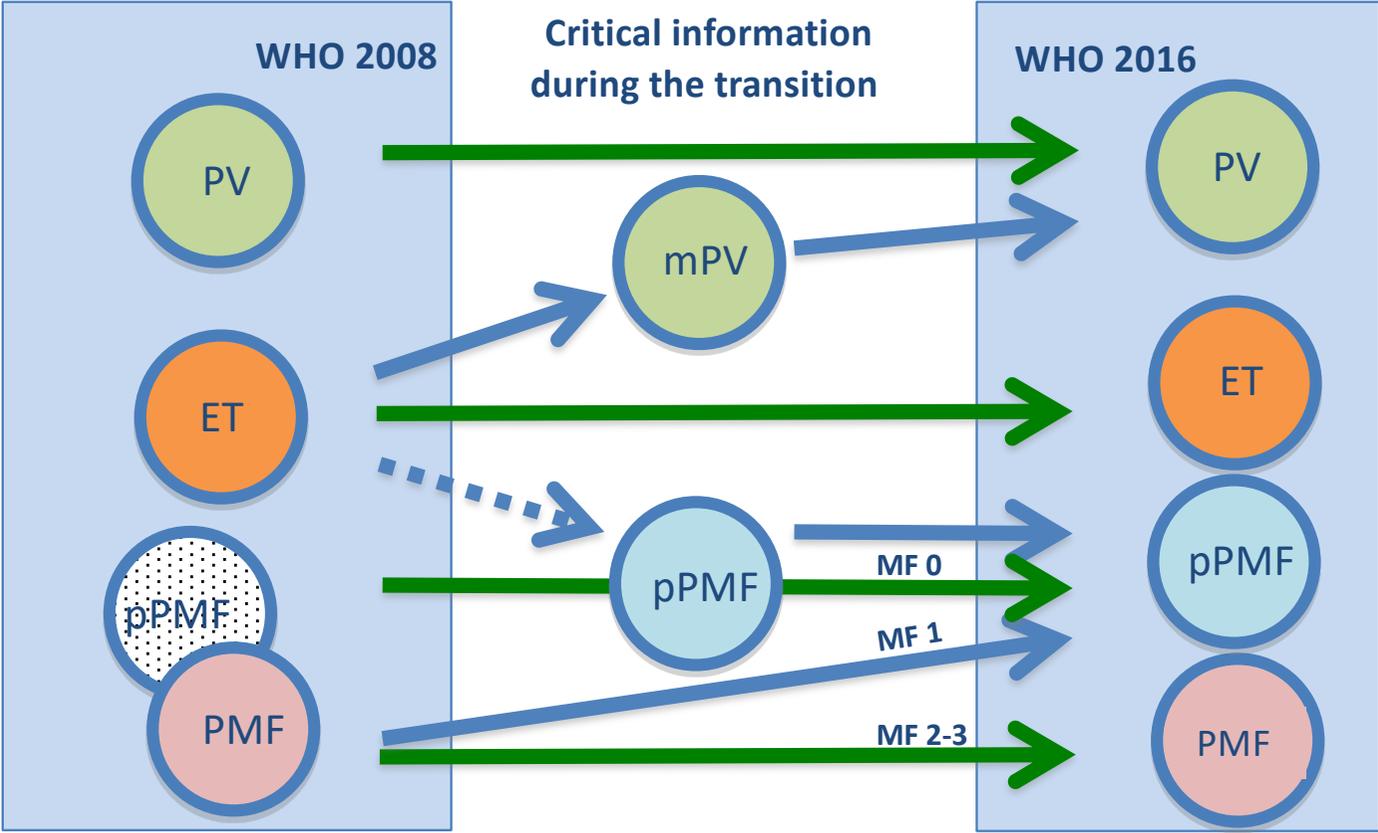


Tefferi et al, Am J Hematol. 2014 Aug;89(8):E121-4

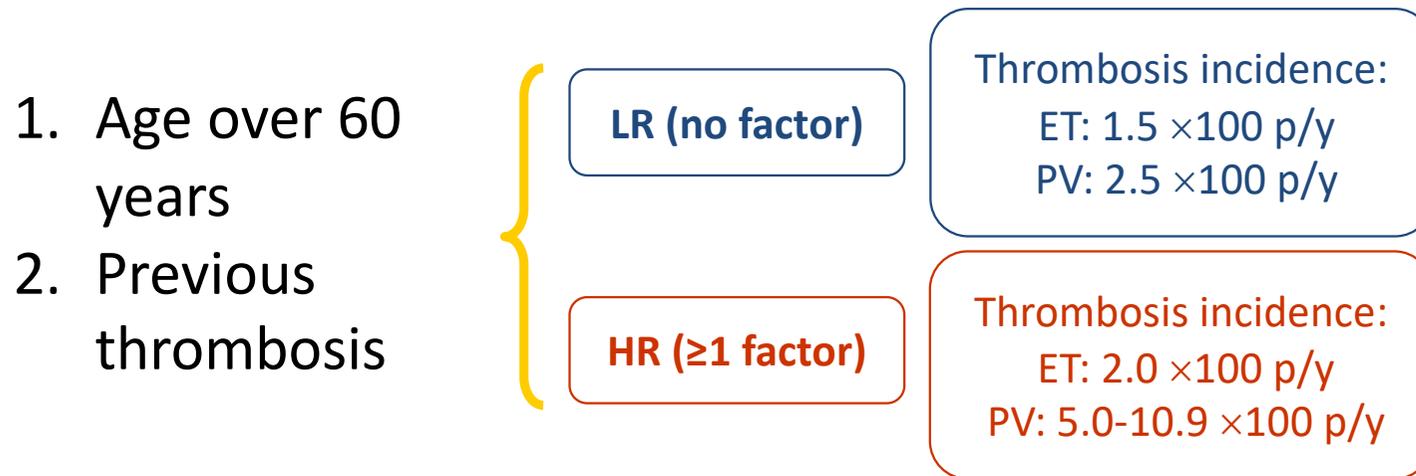


***polycythemia vera***

# What is PV: WHO 2008 → 2017



# Conventional prediction of thrombosis in PV



Cardiovascular risk factors to be corrected/prevented

Extreme thrombocytosis (PLT count  $>1,500 \times 10^9/L$ ) implies a higher risk of bleeding

# Prediction of post-PV MF evolution

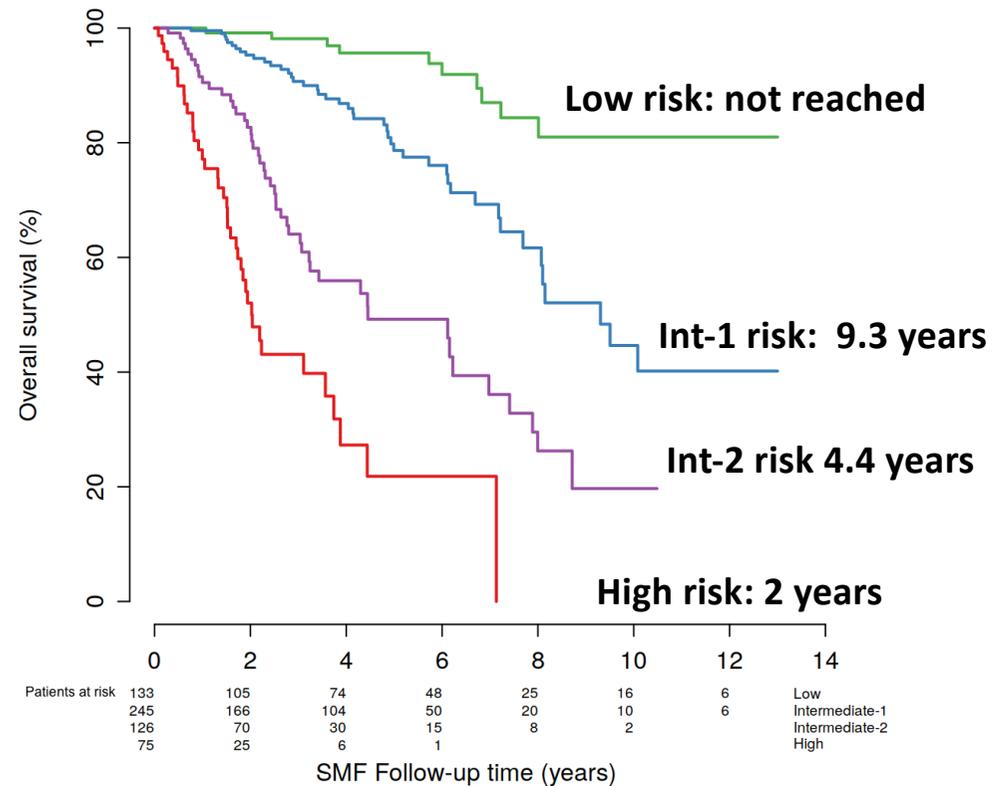
- SMF occur in 12-21% of PV
- Potential predictors of SMF evolution:
  - Detection of BM fibrosis in BM biopsy of PV at diagnosis
  - Some clinical parameters at diagnosis of PV (leukocytosis, splenomegaly, higher *JAK2* allele burden) or during PV course (HU-resistant splenomegaly)
  - *SRSF2*, *DNMT3A* mutations in PV; -2518 A/G polymorphism of MCP-1 (monocyte chemoattractant protein-1)
  - Abnormal cytogenetics

*Tefferi et al. Blood. 2014 Oct 16;124(16):2507-13; Barbui et al, J Clin Oncol. 2011 Aug 10;29(23):3179-84; Passamonti et al, Blood 2008; Tefferi et al; Leukemia 2013; Alvarez-Larran et al; Br J Haematol. 2016 Mar;172(5):786-93; Passamonti et al, Blood 2006; Passamonti et al Leukemia Sep;24(9):1574-9; Rumi et al., Blood 2013; Rumi E et al, Blood 2014;123(10):1544-51; Pietra et al. Leukemia. 2016 Feb;30(2):431-8; Passamonti et al. Leukemia. 2017 Jan 3. doi: 10.1038/leu.2016.351; Elala et al. Am J Hematol. 2016 May;91(5):503-6; Lundberg et al, Blood 2014; Bartels et al, Leukemia 2017; Masselli et al, Leukemia 2018*

# Prediction of survival in post-PV/ET MF

## The integrated clinical-molecular model MYSEC-PM

Covariates	Points
Age, years	0.15
Hemoglobin <11 g/dL	2
Platelet < 150 x10 <sup>9</sup> /L	1
Circulating blast cells ≥ 3%	2
<i>CALR</i> -unmutated genotype	2
Constitutional symptoms	1



# HCT level is critical in PV management: the CYTO-PV trial

365 *JAK2*-pos. PV pts treated with phlebotomy ± HU

R  
1:1

Treated to Hct <45%  
n = 182

Treated to Hct 45-50%  
n = 183

Primary endpoint achieved, n (%):

5\* (2.7)

18\* (9.8)

Secondary endpoint achieved, n (%):

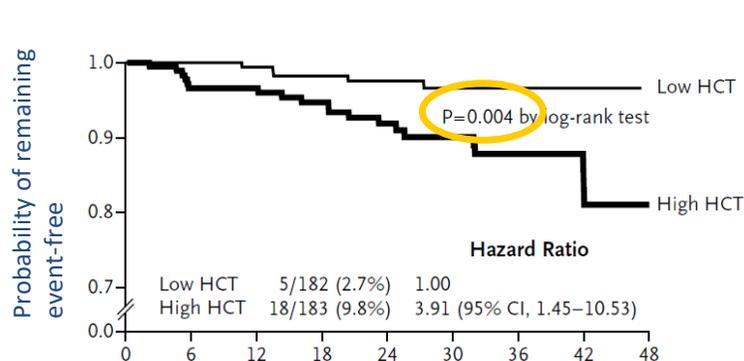
8 (4.4)

20 (10.9)

**Primary composite end point:** Death from CV cause or major thrombotic events (stroke, acute coronary syndrome, transient ischemic attack, PE, abdominal thrombosis, DVT, or peripheral arterial thrombosis)

**Secondary end point:** Primary end point + superficial-vein thrombosis

- 0 deaths in the Hct <45% arm,
- 4 deaths in the Hct 45-50% arm



Marchioli R et al, *N Engl J Med* 2013;368(1):22-33

# Inadequately controlled PV is a predictive factor

- Recorded in 137 patients (15.4%):
  - Need for phlebotomies (3.3%)
  - Uncontrolled myeloproliferation (1.6%)
  - Failure to reduce massive splenomegaly (0.8%)
  - Cytopenia at the lowest HU-dose to achieve response (1.7%)
  - Extra-haematological toxicity (9%)
- **Cytopenia affected survival, progression to MF, AML**
- **Splenomegaly affected MF**

## Conclusions

- Driver mutations enter PV and ET process for diagnosis.
- *JAK2*, *CALR*, *MPL* variants can identify patients with different risk of thrombosis, and myelofibrosis.
- AML occurrence is more disease-based than expected.