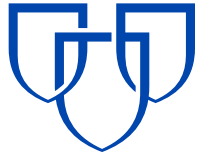


# Disclosures for Ayalew Tefferi

Principal investigator role	Janssen, Geron, Celgene, Sanofi-Aventis, Gilead Sciences, Incyte
Employee	None
Consultant	None
Major Stockholder	None
Speakers' Bureau	None
Scientific Advisory Board	None

**Presentation includes discussion of the following off-label use of a drug or medical device: Hydroxyurea, Interferon-alpha, Busulfan, Thalidomide, Lenalidomide, Pomalidomide, Ruxolitinib, Androgen preparations, Erythropoiesis stimulating agents**

MAYO  
CLINIC



**Myeloproliferative neoplasms other than PV, ET and PMF:**  
**CMML: chronic myelomonocytic leukemia**  
**Mastocytosis**  
**Eosinophilic disorders**  
**CNL: chronic neutrophilic leukemia**

Ayalew Tefferi, MD  
Professor of Medicine and Hematology  
Mayo Clinic College of Medicine

# Objectives

- Disease definitions
- Diagnosis
- Current prognostication
- Treatment

# 2016 WHO Classification of Myeloid Malignancies

Acute Myeloid Leukemia (AML)

Myelodysplastic Syndromes (MDS)

Myeloproliferative Neoplasms (MPN)

MDS/MPN overlap

Myeloid/Lymphoid neoplasms with eosinophilia and *PDGFR/FGFR1/PCM1-JAK2* mutation

CMML

Molecularly-defined eosinophilia

Chronic Myeloid Leukemia (CML)  
*BCR-ABL1*  
100% mutated

Chronic Neutrophilic Leukemia (CNL)  
*CSF3R*  
80-100% mutated

Chronic Eosinophilic Leukemia Not Otherwise Specified (CEL-NOS)

Polycythemia vera (PV)

Essential Thrombocythemia (ET)

Primary Myelofibrosis (PMF)

MPN Unclassifiable (MPN-U)

CNL

Vs. HES

*The JAK2/CALR/MPL mutated MPNs*

97% *JAK2* V617F  
3% other *JAK2* mutations

60% *JAK2* mutated  
22% *CALR* mutated  
3% *MPL* mutated  
15% triple-negative

60% *JAK2* mutated  
23% *CALR* mutated  
7% *MPL* mutated  
10% triple-negative

*Mastocytosis no longer under the WHO MPN category*

# Acquired eosinophilia

Secondary

- Drugs
- Infections
  - Parasites
- Allergy
- Inflammations
  - Kimura's
  - CSS
  - Well's
- Neoplasia
  - Hodgkins
  - NHL
  - Solid tumor

Primary

Clonal

**Cytogenetic, molecular or bone marrow morphologic evidence of an otherwise defined myeloid malignancy**

- PDGFRA/PDGFRB* mutated (imatinib sensitive)
- FGFR1* rearranged (urgent transplant)
- PCM1-JAK2* (ruxolitinib)

- Abnormal karyotype, non-specific (CEL-NOS)
- Excess blasts (CEL-NOS)

-AML/ALL/CML/MDS/CMML/SM/unclassifiable

Idiopathic

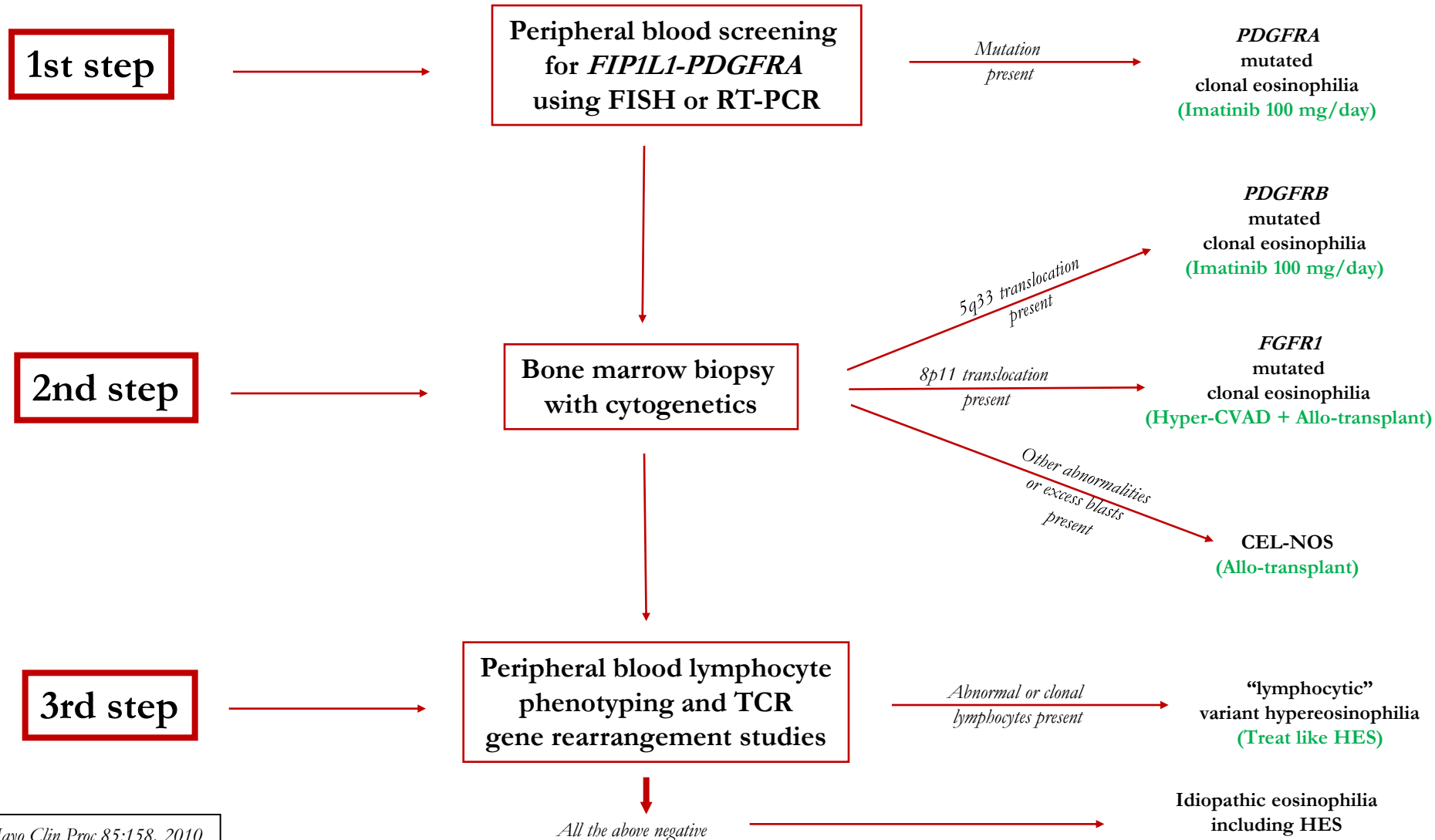
Neither reactive nor clonal

HES

if AEC > 1500/micL  
x 6 months and  
organ damage

20% of "HES" pts may display a T cell clone/abnormal phenotype (lymphocytic variant hyper-eosinophilia)

# Primary eosinophilia diagnostic algorithm



# Hyper-eosinophilic syndrome/idiopathic eosinophilia

98 Mayo Clinic patients with WHO-defined HES/IH (*Leukemia* 2016;30:1924)

NGS revealed 11% harbored pathogenic mutation;

*TET2*=3, *ASXL1* =2, *KIT*=2, and *IDH2*, *JAK2*, *SF3B1* and *TP53*=1 each.

15% harbored a variant of unknown significance (VUS);

*TET2*=8, *ASXL1*=2, *SETBP1*=2, and *CALR*, *CEBPA* and *CSF3R*=1 each.

**NO DIFFERENCE IN MUTATED VS NON-MUTATED IN PHENOTYPE**  
**MUTATED PATIENTS HAD INFERIOR SURVIVAL IN UNIVARIATE ANALYSIS**

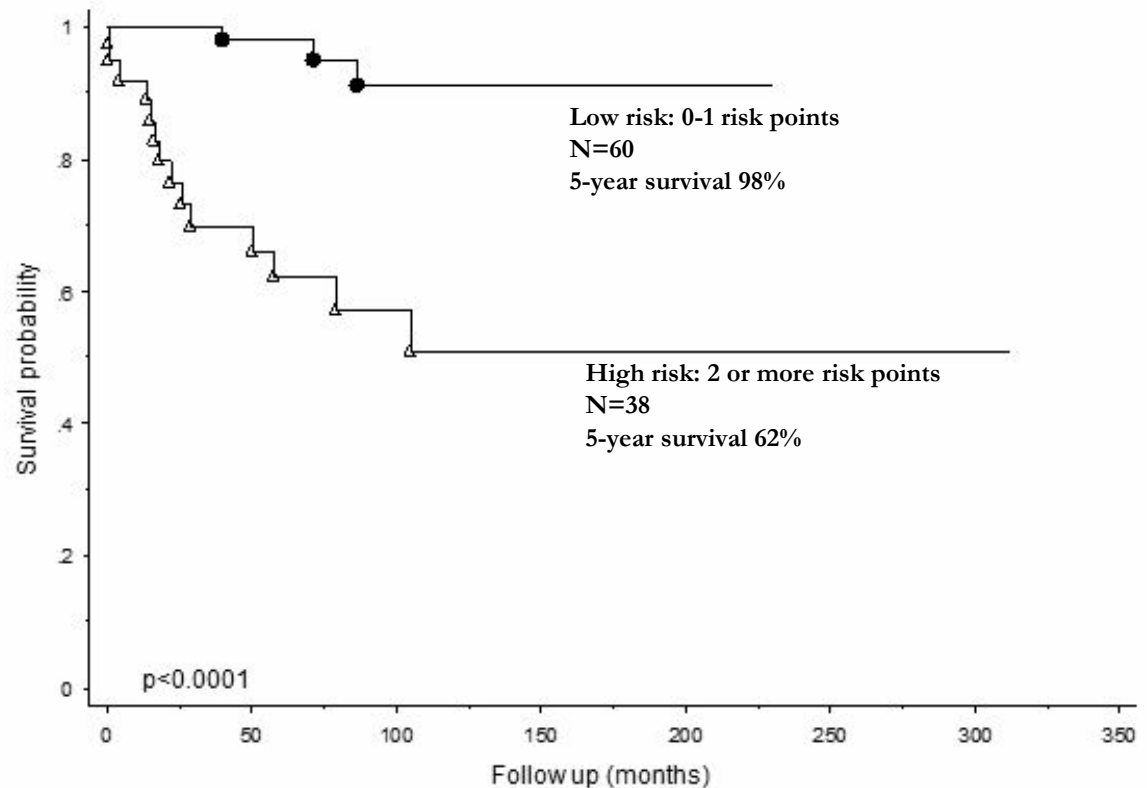
Risk factors for survival:

*Hepatosplenomegaly (3 points)*

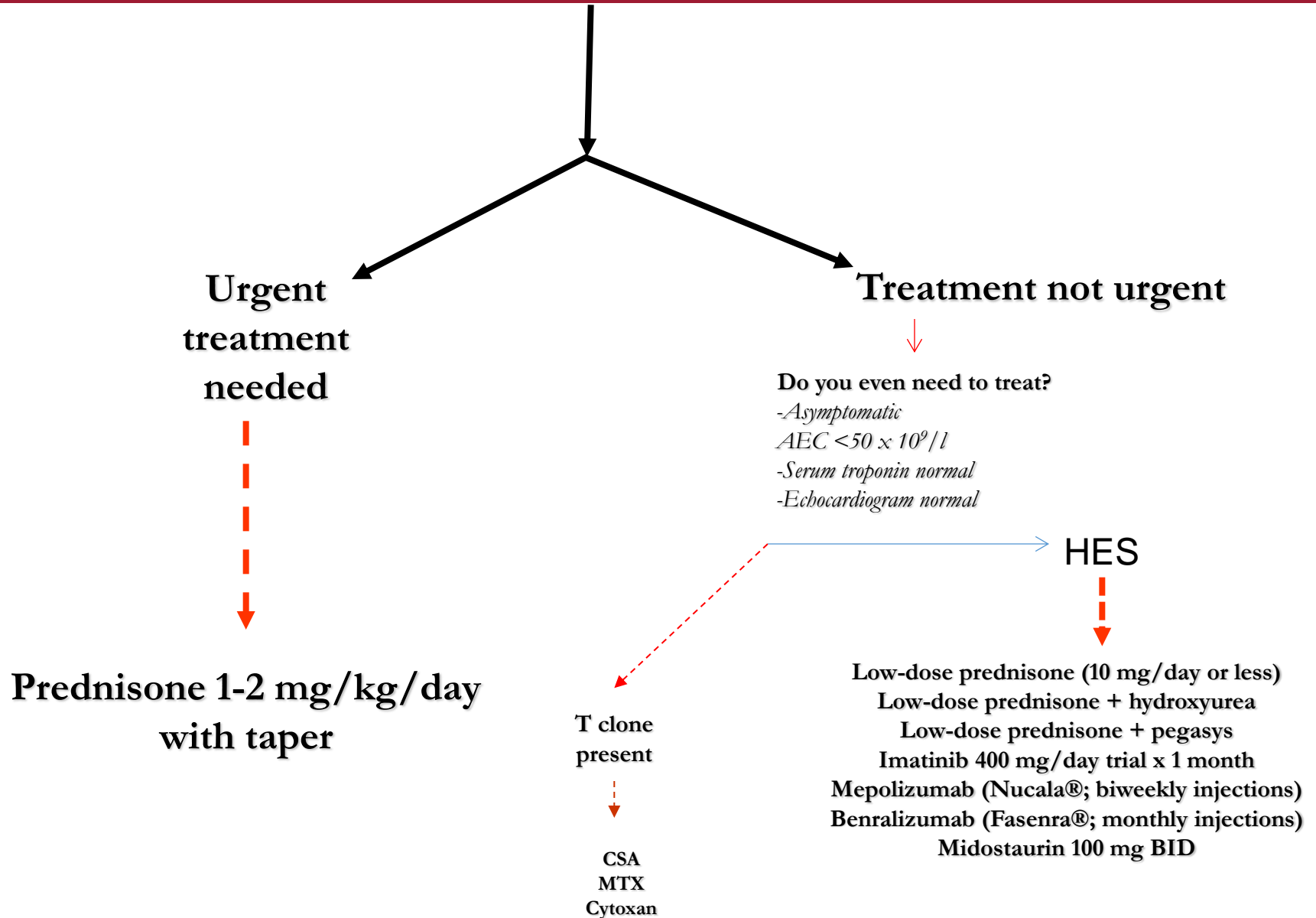
*Advanced age (2 points)*

*Hgb <10 g/dl (one point)*

*Cardiac involvement (one point)*

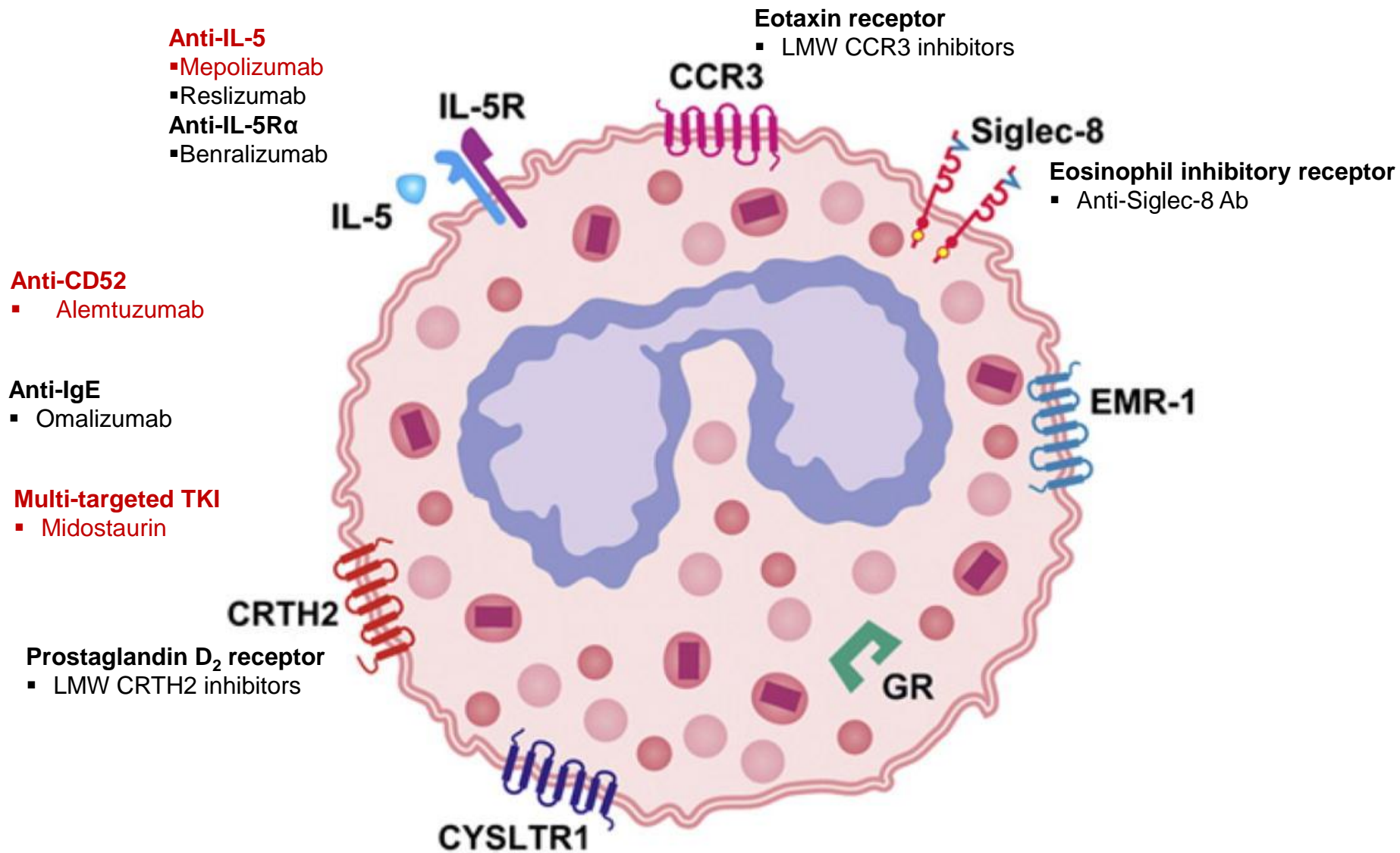


# HES treatment algorithm





# Novel targeted therapies for eosinophilic disorders



# When should you suspect mastocytosis?

- Urticaria pigmentosa

- Mast cell mediator symptoms

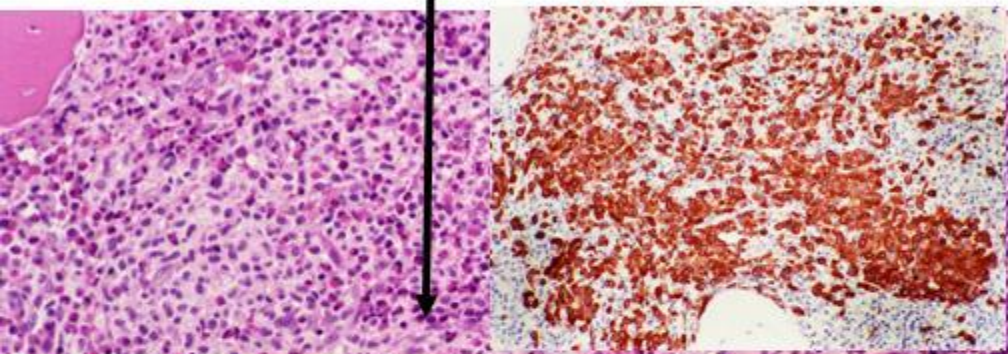
- Anaphylactoid
- Diarrhea
- Flushing/urticaria



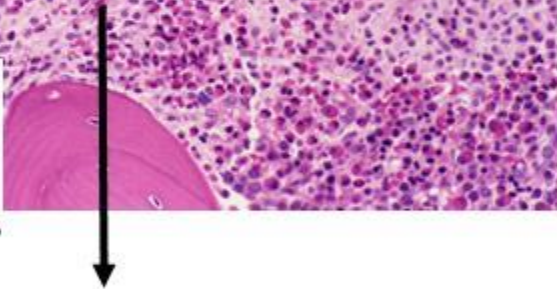
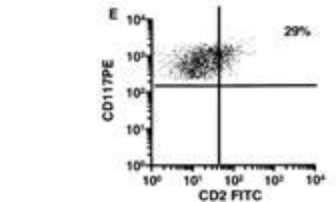
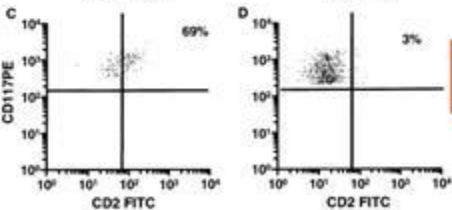
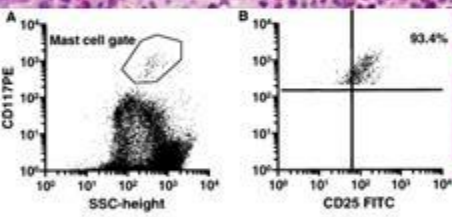
- Osteopenia/unexplained fractures

# Diagnostic Evaluation in Systemic Mastocytosis

↓  
**serum tryptase**



**Bone marrow biopsy with tryptase stains**



**Bone marrow mast cell flow cytometry**

**Normal mast cells —  
 CD117+, CD25-, CD2-**

**Abnormal mast cells —  
 CD117+, CD25+, CD2 ±**

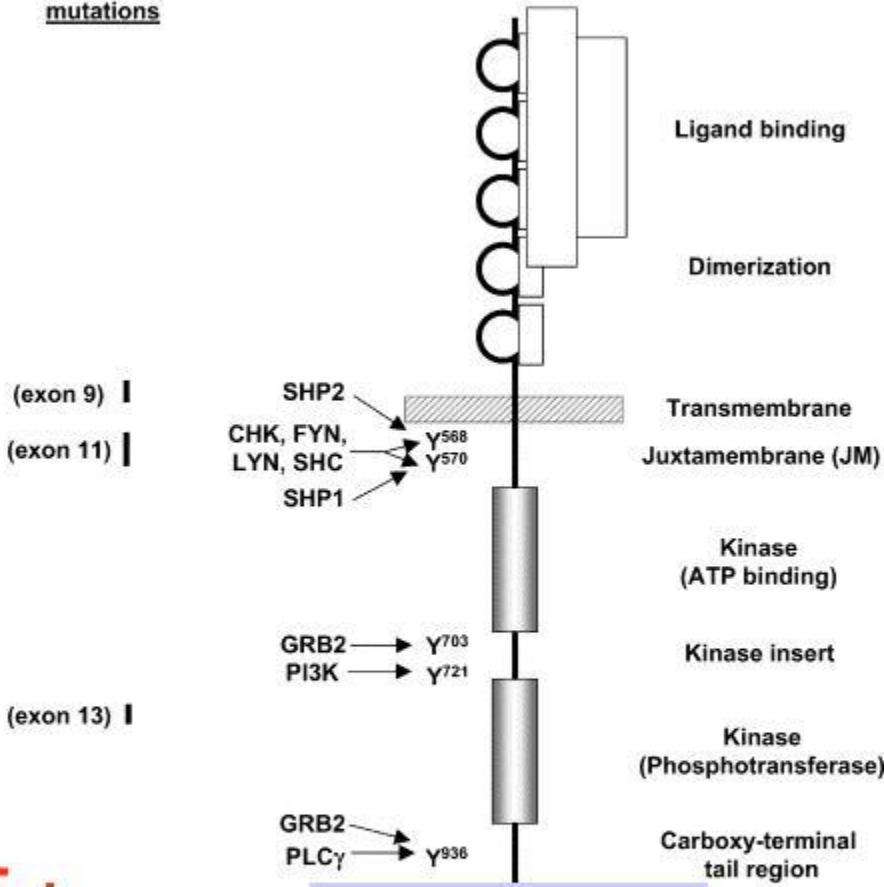
Pardanani et al. BJH 2003;120:691

Regions with activating mutations

Phospho-Tyr binding proteins

Structure

Domains



**±C-kit mutations  
 Asp816Val (kinase domain)  
 Val560Gly (JM domain)**

Sattler & Salgia, Leukemia Research, 2004

# Practical classification of mast cell disease

1

Cutaneous mastocytosis  
(skin-only disease)



*Both can manifest  
mast cell mediator  
release symptoms*



2



Systemic mastocytosis (SM)

i

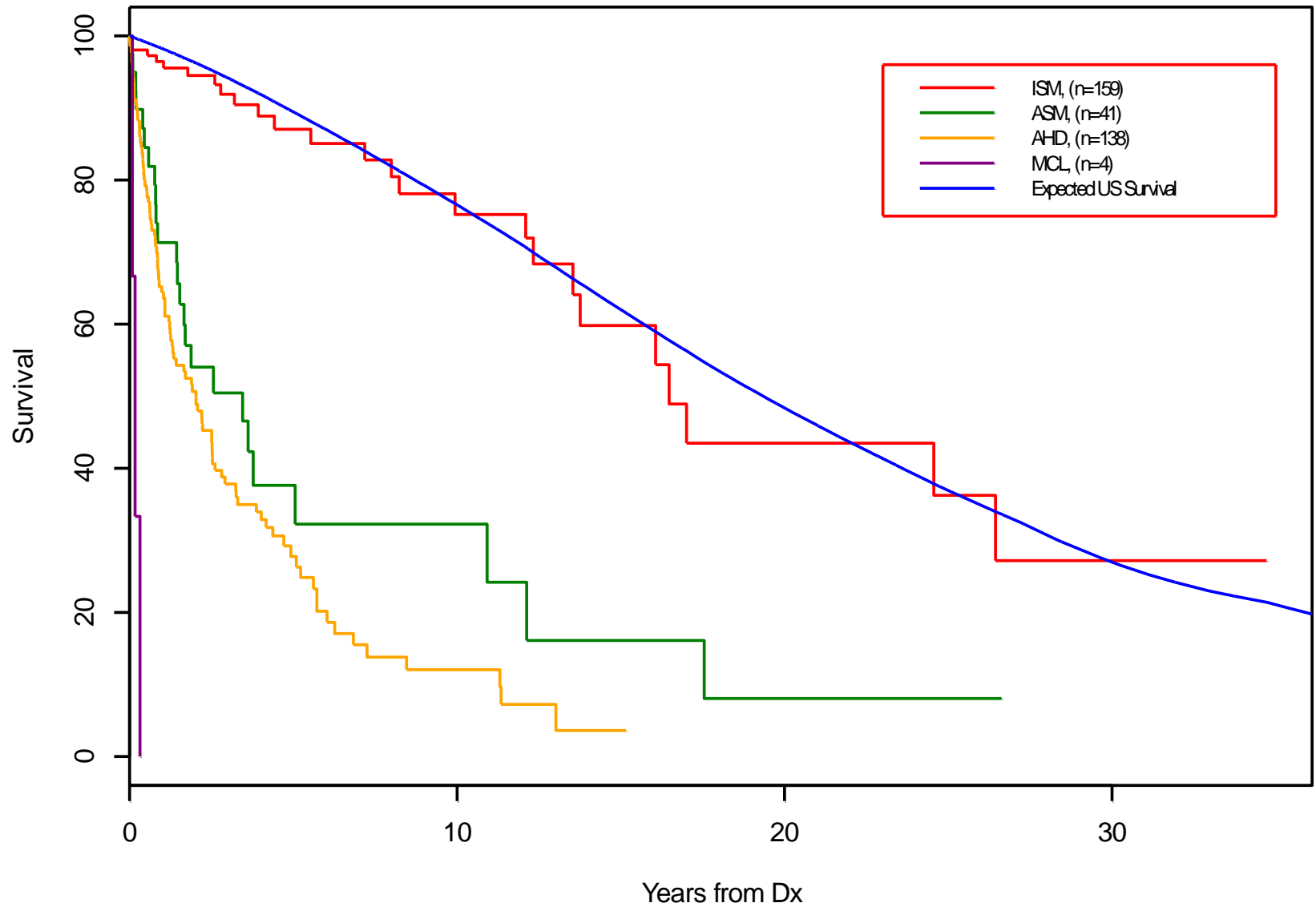
Indolent SM

ii

Aggressive SM (cytopenia, bone disease, organomegaly, etc.)

1. SM without associated 2<sup>nd</sup> myeloid neoplasm
2. SM with associated 2<sup>nd</sup> myeloid neoplasm
3. Mast cell leukemia

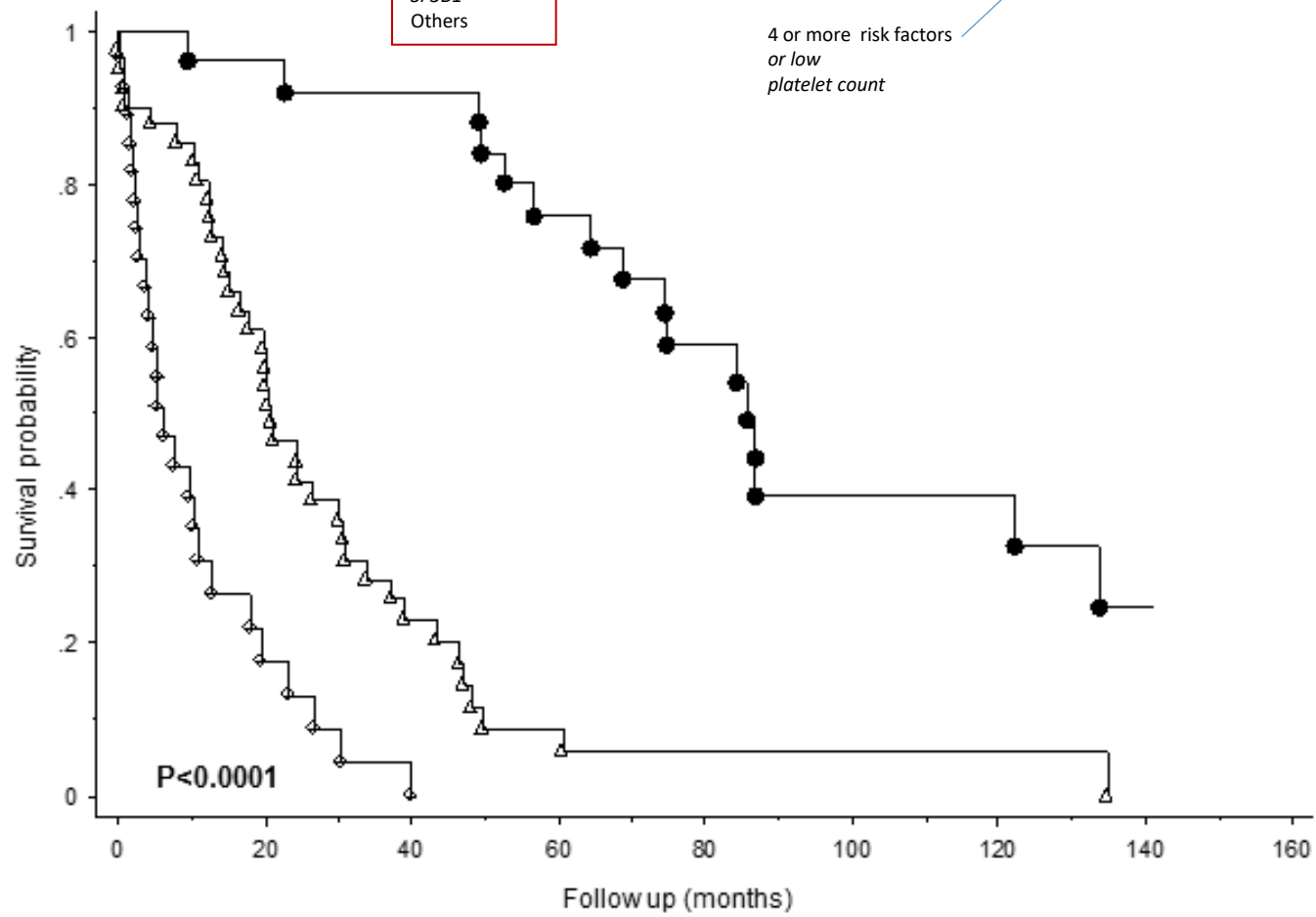
**Survival for 342 systemic mastocytosis patients classified by disease type compared with the expected age and gender matched US Population's survival**



# Mutation-augmented prognostic scoring system (MAPSS) in 94 patients with advanced mastocytosis

- ISM (n=44)**  
KIT 73%  
TET2 7%  
No other mutations
- ASM (n=25)**  
KIT 84%  
TET2 20%  
ASXL1 16%
- AHN (n=80)**  
KIT 75%  
TET2 45%  
ASXL1 26%  
CBL 19%  
JAK2  
DNMT3A  
U2AF1  
RUNX1  
SF3B1  
Others

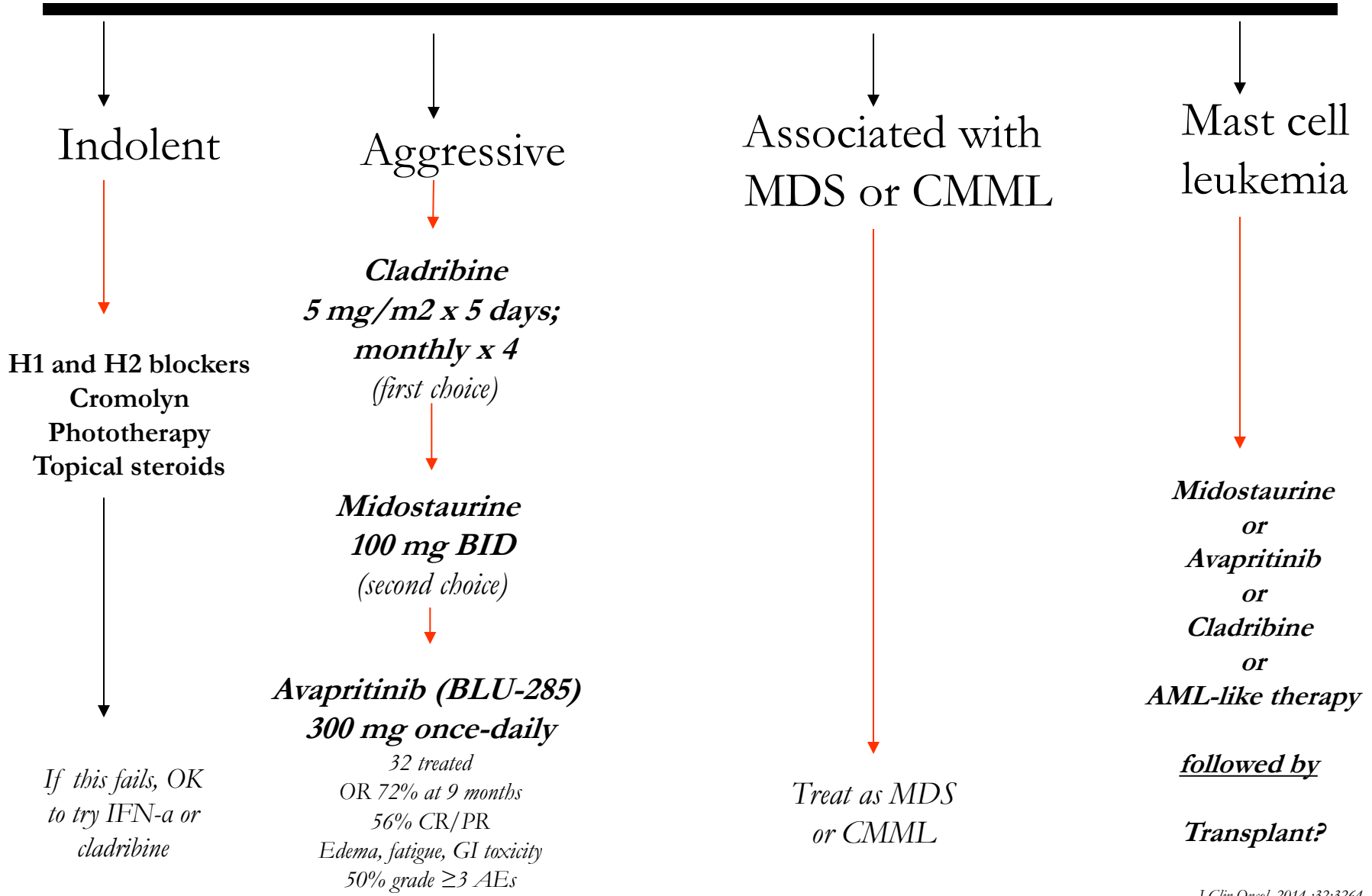
- One risk factor other than platelet count → ● Low-risk: n=26 (16 events), median OS=86 months
- 2-3 risk factors or low platelet count → △ Intermediate-risk: n=41 (38 events), median OS=21 months
- 4 or more risk factors or low platelet count → ◇ High-risk: n=27 (25 events), median OS=5 months



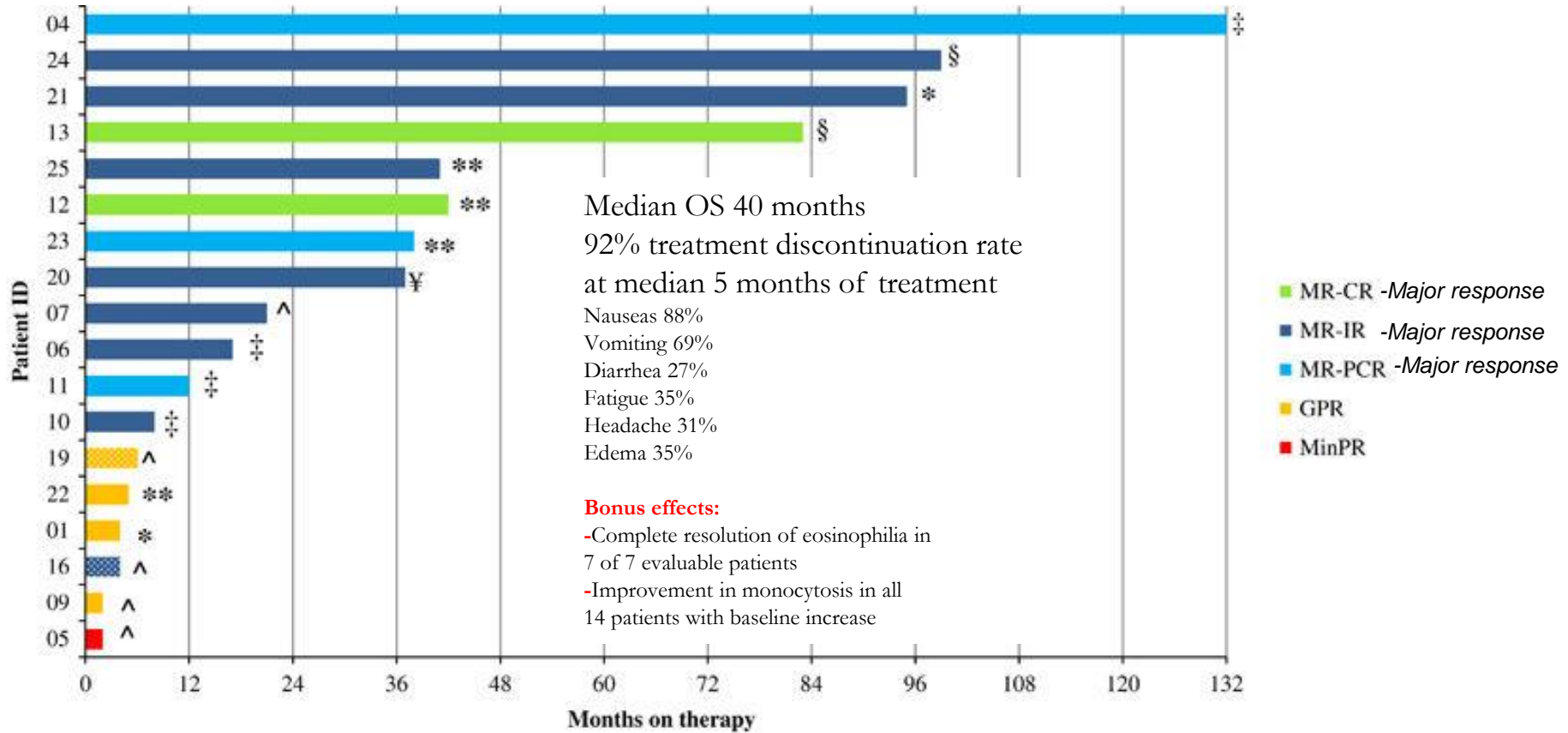
## Risk factors:

- Platelet count <150
- albumin <3.5
- age >60
- ASXL1/CBL mutated
- Hgb <10

# Treatment for Systemic mastocytosis



Phase-2 study of midostaurin 10-year follow-up (N=26; responders = 18 (69%; major response 50%)  
 SM-AHN = 17 (13 responders); ASM = 3 (1 responder), MCL = 6 (4 responders)



Study status/reason for discontinuation (# of patients)<sup>†</sup>

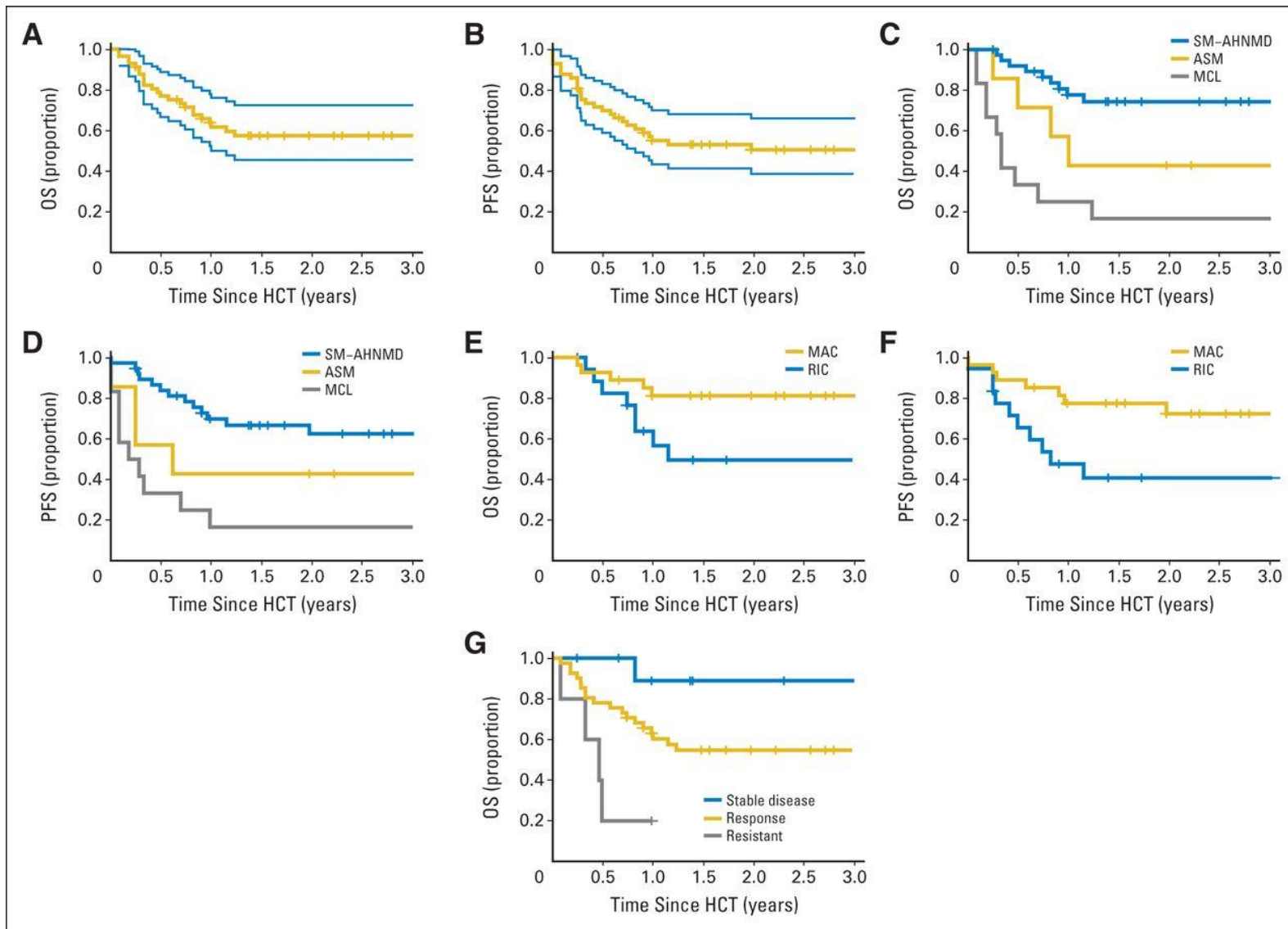
- § Continuing treatment (n=2)
- \* Adverse event: Grade 3 or 4 thrombocytopenia (n=2)
- \*\* Serious adverse event: sepsis (n=3), inflammatory mixed neuropathy/myopathy and altered mental status (n=1)
- † Withdrew consent (n=1)
- ^ Unsatisfactory therapeutic effect; discontinued per investigator discretion (n=5)
- ‡ Disease progression (n=4)

Median duration of treatment (months)	
median	19
range	2-132

<sup>†</sup> Data through 3/1/2017; best response at any time on therapy  
 Solid bars are *KIT* D816 mutation-positive, two patterned bars are *KIT* D816 mutation-negative



**Allogeneic hematopoietic stem-cell transplantation (alloHCT or HCT) outcomes in 57 patients with advanced systemic mastocytosis (SM): 38 SM-AHNMD; 12 MCL and 7 aggressive SM.**



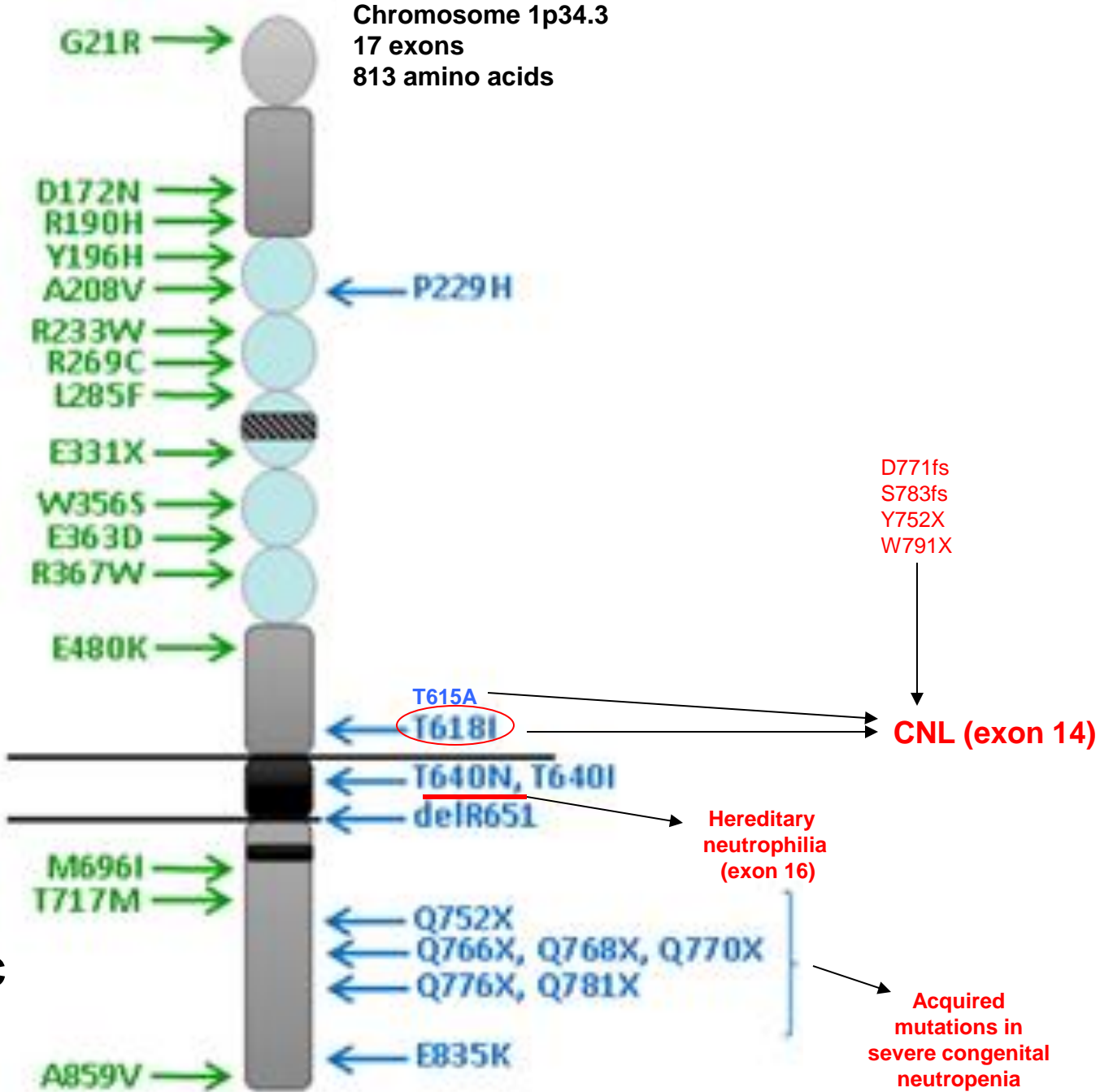
**CSF3R mutations**

Chromosome 1p34.3  
17 exons  
813 amino acids

**Extracellular domain**

*Membrane proximal*

**Cytoplasmic domain**



## WHO diagnostic criteria for CNL

1. Leukocytosis  $\geq 25 \times 10^9/L$ 
  - Neutrophils plus bands  $\geq 80\%$
  - Neutrophil precursors  $< 10\%$
  - Myeloblasts rarely observed
  - No dysgranulopoiesis
2. Hypercellular bone marrow
  - Neutrophil granulocytes increased
  - Neutrophil maturation normal
  - Myeloblasts  $< 5\%$
3. Not meeting WHO criteria for BCR-ABL1+ CML, PMF, PV or ET
4. No PDGFRA/PDGFRB/FGFR1/PCM1-JAK2
5. CSF3RT618I or other activating CSF3R mutation

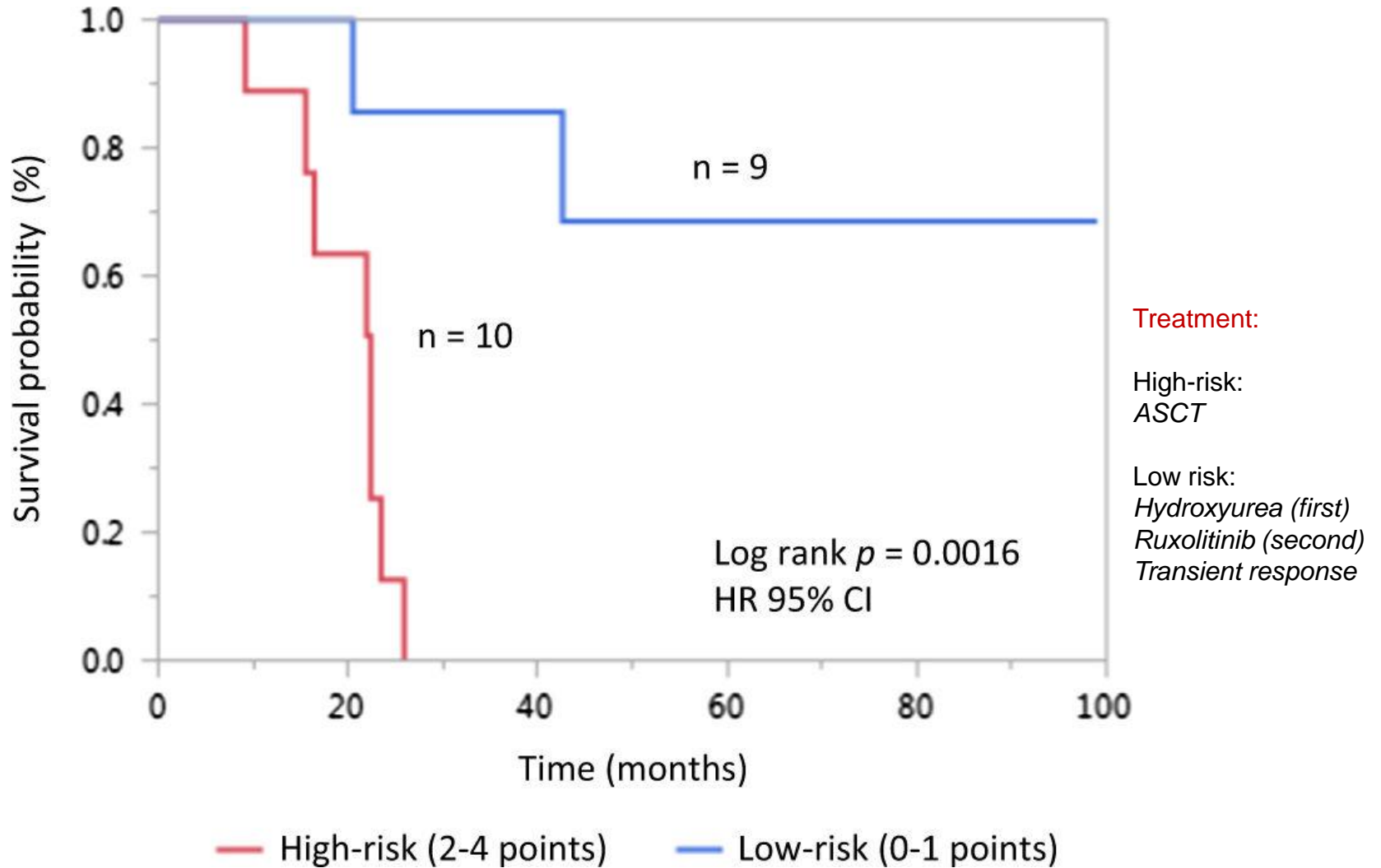
or

no identifiable cause of reactive neutrophilia

## WHO diagnostic criteria for aCML

- Leukocytosis with  $\geq 10\%$  precursors
- Dysgranulopoiesis
- Basophils  $< 2\%$
- Monocytes  $< 10\%$
- Hypercellular bone marrow with dysgranulopoiesis
- $< 20\%$  blasts in the blood and bone marrow
- No PDGFRA/PDGFRB/FGFR1/PCM1-JAK2
- Not meeting WHO criteria for BCR-ABL1+ CML, PMF, PV or ET

## Risk-stratified Kaplan–Meier survival curves for 19 *CSF3R*-mutated CNL patients.



Risk points: platelet count  $<160 \times 10^9/L$  (2 points); leukocyte count  $>60 \times 10^9/L$  (1 point); *ASXL1* mutation (1 point)

# CMML

## 2016 WHO Diagnostic Criteria

- AMC  $>1 \times 10^9$  /L and monocytes  $>10\%$
- Dysplasia in one or more myeloid lineages
- Not meeting WHO criteria for CML, PV, ET or MF
- No *PDGFRA/PDGFRB/FGFR1/PCM1-JAK2*
- $<20\%$  blasts in the blood and BM

In the absence of dysplasia, a diagnosis of CMML can still be made if:

- An acquired clonal cytogenetic or molecular genetic abnormality can be documented.  
these include *ASXL1*, *TET2*, *SRSF2* and *SETBP1* mutations
- Monocytosis has persisted for  $>3$  months
- Other causes of **reactive** monocytosis have been ruled out

# Differential diagnosis of Monocytosis

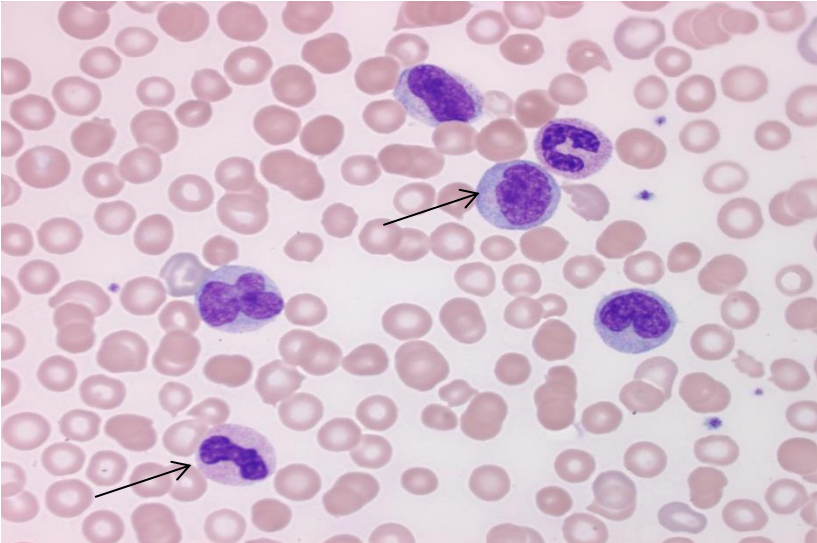
## Reactive

1. Viral infections.
2. Recovering bone marrow.
3. Connective tissue disorders.
4. Sarcoidosis
5. Tuberculosis, Brucellosis, Leishmaniasis.
6. SABA.

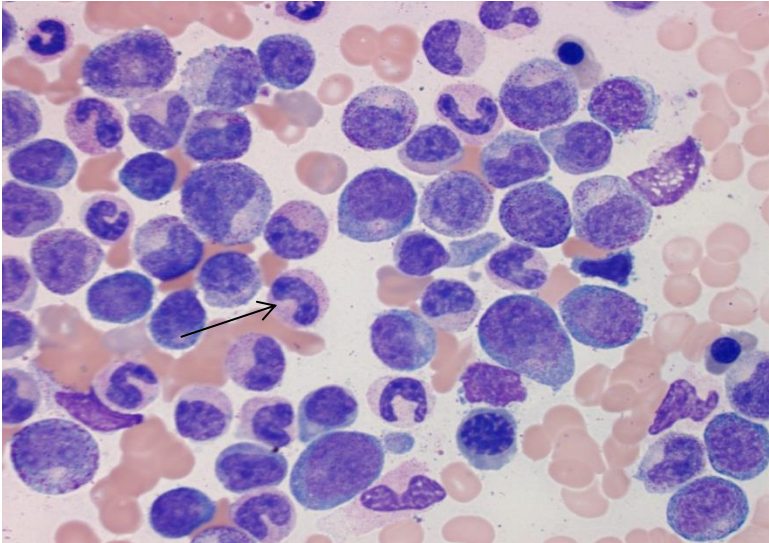
## Clonal

1. CMML
2. JMML
3. AML with monocytic differentiation.
4. MDS/MPN overlap syndromes- unclassifiable.

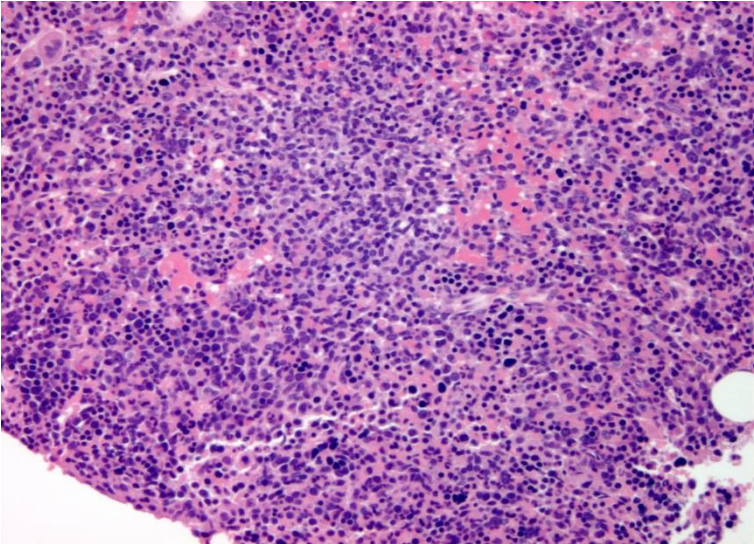
# CMML- Peripheral Blood and Bone Marrow Findings



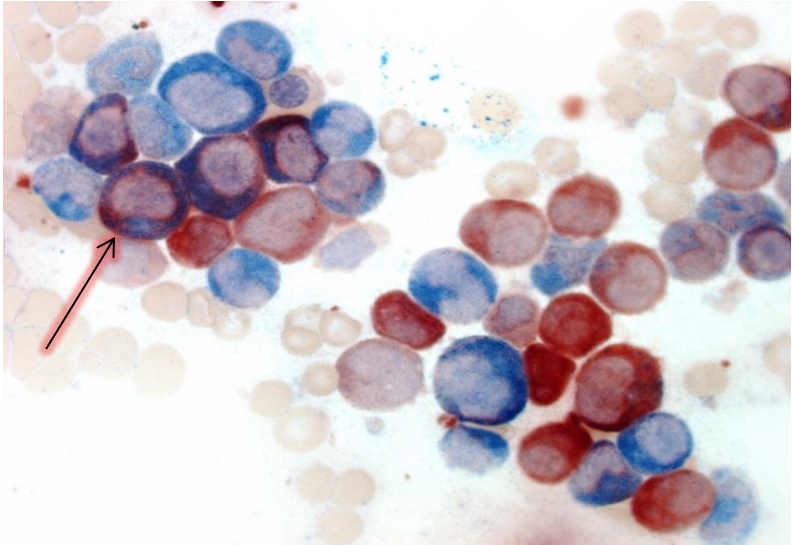
Peripheral Blood Smear



Bone Marrow Aspirate



Core Biopsy



Dual Esterase Stain

# WHO CMML Subcategories

## CMML-0

- Blasts + promonocytes <2% in PB.
- Blasts + promonocytes <5% in BM.

## CMML-1

- Blasts + promonocytes 2-4% in PB.
- Blasts + promonocytes 5-9% in BM

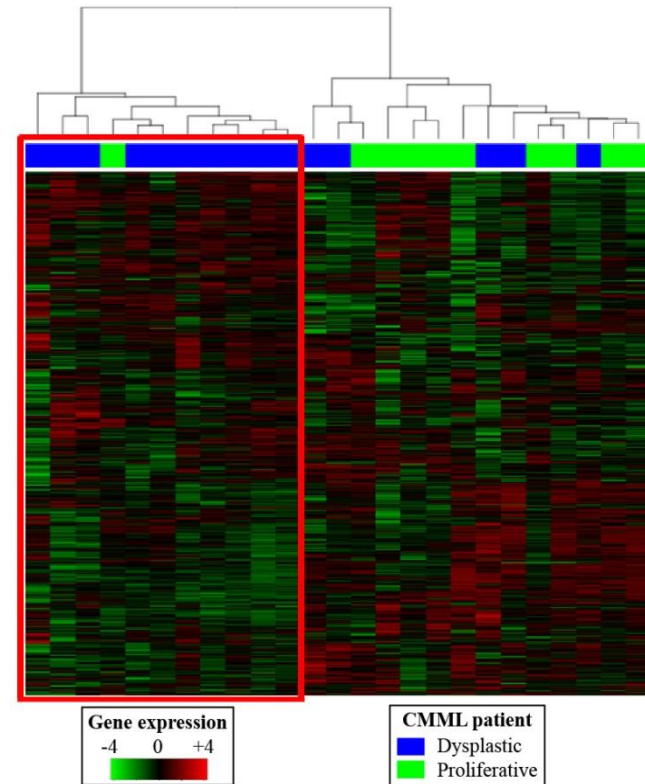
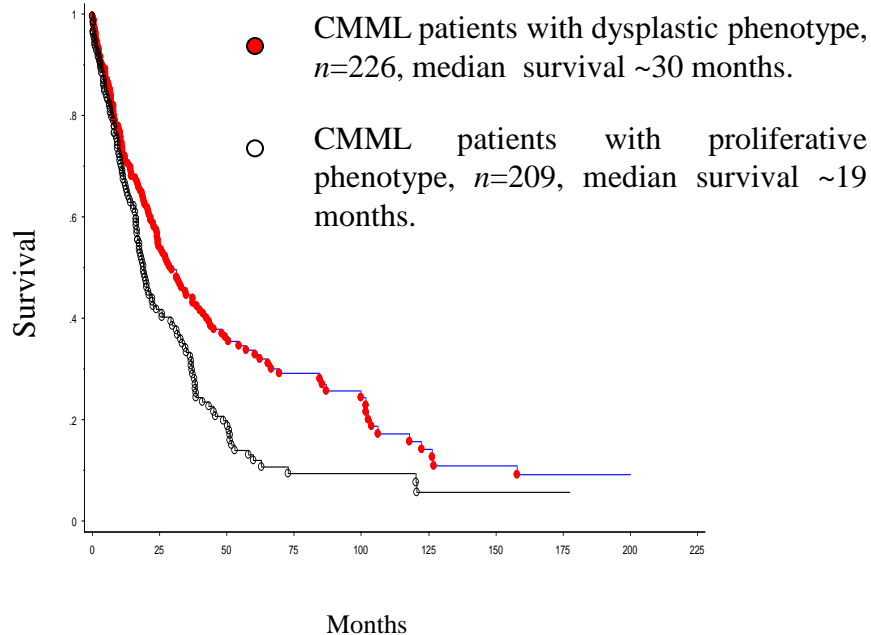
## CMML-2

- Blasts + promonocytes 5-19% in PB.
- Blasts + promonocytes 10-19% in BM.
- Presence of **Auer rods**, irrespective of blast count.



# Proliferative vs Dysplastic CMML

Survival data for 435 patients with WHO defined chronic myelomonocytic leukemia stratified by “Proliferative” versus “Dysplastic” sub-types.



**Figure 2:** Heat map of an unsupervised differential gene expression profile in peripheral blood CMML samples, demonstrating two predominantly unique clusters, segregating dysplastic (cluster 1) from proliferative (cluster 2) CMML subtypes.

# CMML – Genomics

- Epigenetic regulators – TET2 (~60%), *IDH1*, *IDH2*, *DNMT3A*
- Chromatin modeling – ASXL1 (~40%), *EZH2*
- Spliceosome components – SRSF2 (~45%), *SF3B1*, *U2AF1*, *ZRSR2*
- Transcription factors – *RUNX1* (~15%)
- Signal pathways – *JAK2*, *KRAS*, *NRAS*, *CBL*, *PTPN11* (RAS pathway ~30%)
- Others – *SETBP1* (~15%), *PHF6*, *BCOR*, *Tp53*

**> 90% CMML patients have ≥ 1 somatic mutations**

# Cytogenetic abnormalities in CMML

- Cytogenetic abnormalities seen in 20-40% of cases
- Most common are +8, chromosome 7 abnormalities and 12p deletions

## Spanish Cytogenetic Risk Stratification.

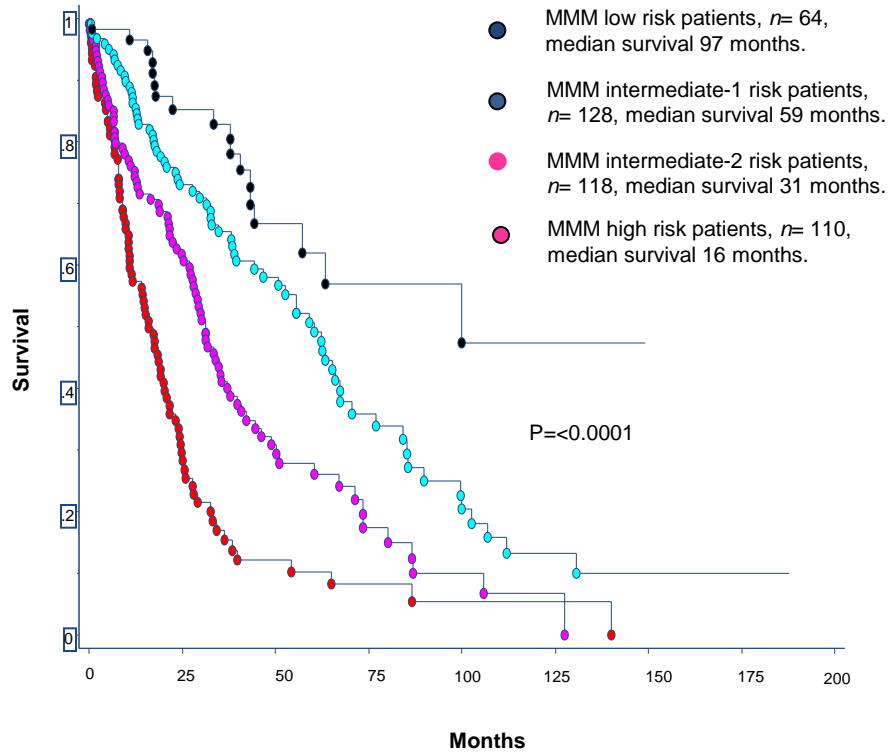
- Low: Normal, -Y
- Intermediate: all others
- High: +8, chromosome 7 abnormalities and complex changes

**5 year OS- 35%, 26% and 4%.**

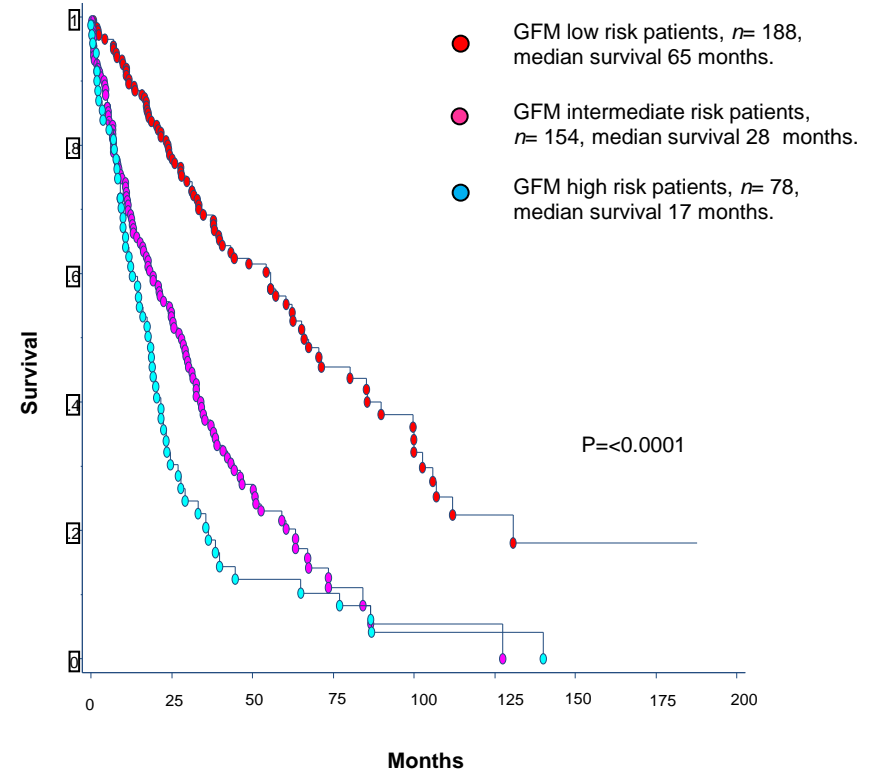
# CMML Prognostic Models

Variables	<u>Mayo Model</u>	<u>Molecular Mayo Model</u>	<u>GFM Model</u>	<u>CPSS-Molecular</u>
HB < 10 gm/dl	+	+	+	Red blood cell transfusion dependence
High WBC			+ (>15)	+ (>13)
AMC >10	+	+		
Platelets < 100	+	+	+	
Circulating IMC	+	+		
BM blasts				+ (>5%)
Age >65			+	
Cytogenetic risk groups				+
Molecular genetics		<i>ASXL1</i>	<i>ASXL1</i>	<i>ASXL1/ NRAS/ RUNX1 and SETBP1</i>

**Survival data for 420 patients with  
CMML stratified by the Molecular Mayo Model.**



**Survival data for 420 patients with  
CMML stratified by the GFM Model.**



# CMML Therapeutics

## Supportive care

- Transfusions
- Hydroxyurea
- ESA
- Iron chelation therapy

## Directed Therapies

- Hypomethylating agents
- Allogeneic SCT
- Clinical trials

# Hypomethylating (HMA) Agents in CMML

Study	N	Median Age (years)	Phase	Drug used	Response rates (%)	Median survival (months)
Wijermans 2008	31	71	II	Decitabine	CR-10 PR-16 HI-19	15
Costa 2011	38	70	II	Azacitidine	CR-11 PR- 3 HI- 25	12
Braun 2011	39	71	II	Decitabine	CR-10 PR-20 HI-8	18
Thorpe 2012	10	66	II	Azacitidine	CR-20 HI-40	NR
Ades 2013	76	70	II	Azacitidine	CR-17 PR-1 HI-17	29
Wong 2013	11	65	II	Azacitidine	CR-9 PR-9 HI-9	17
Fianchi 2013	31	69	II	Azacitidine	CR-45 PR-3 HI-6	37

## Role for Allogeneic SCT in CMML

Study	N	Age median (range)	Donor Source	Conditioning regimen	Relapse rate and TRM	Outcome OS & DFS
Kroger 2002	50	44 (19-61)	MRD-43 MUD-7	MAC- 50	RR-28% TRM-52%	5 yr DFS-18% 5 yr OS-21%
Symeonidis 2010	283	50	MRD-160 MUD-85	MAC-152 RIC-87	RR-25% TRM-37%	5 yr DFS-38% 5 yr OS-42%
Eissa 2011	85	51 (21-66)	MRD-38 MUD-47	MAC- 58 RIC- 27	RR-27% TRM-35%	10 yr DFS-40% 10 yr OS-40%
Park 2013	73	53 (27-66)	MRD-41 MUD-32	MAC- 30 RIC- 43	RR-35%	3 yr DFS-29% 3 yr OS-32%



# Novel Agents- Clinical Trials.

- MEK inhibitors.
- Hedgehog pathway inhibitors.
- GM-CSF monoclonal antibody (KB003).
- Neddylation inhibitors.
- MAP kinase inhibitor.
- P38/Tie-2 inhibitor.
- Aminopeptidase inhibitors.