

MDS/MPN overlap sd

Focus on CMML

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Learning objectives – Focus on CMML

- When should you think of CMML?
- How do you confirm the diagnosis?
- How do you estimate the risk?
- How do you decide on treatment, including transplantation?
- How do you manage non-transplant candidates?



MDS/MPN - Diagnostic requirements

- Ineffective hematopoiesis
 - Cytopenia
- Myeloproliferation
 - WBC, monocytes, platelets
- BCR-ABL negativity
 - Mandatory exclusion



MDS/MPN - WHO classification

WHO 2016	WHO 2022
CMML	CMML
Atypical CML	MDS/MPN with neutrophilia
MDS/MPN Ring Sideroblasts - Thrombocytosis	MDS/MPN with SF3B1 mutation and thrombocytosis
MDS/MPN-Unclassifiable	MDS/MPN NOS



How rare are MDS/MPN?

Cancer type	New cases/y (2022)*	
Breast	11302	Very frequent
MDS	950	Frequent in hematology
CMML	130	Very rare
Other MDS/MPN	< 5	Ultra rare



Mister M, 72 yo

- No symptoms
- Fit patient
- Normal clinical exam
- Routine blood analysis

Parameter	Apr 2025	Ref Range
WBC $\times 10^9/L$	12.0	4.5–11.0
Absolute Monocytes	1.2	0.2–0.8
Monocyte %	23%	2–8%
Neutrophils	10.8	1.8–7.7
Hb g/dL	11.2	13.5–17.5
MCV fL	101	80–100
Platelets $\times 10^9/L$	142	150–400
Peripheral Blasts	0	0%
LDH U/L	270	140–280

Step 1: Confirm persistence and exclude reactive causes

- **repeat the CBC in 3 months**
- **exclude common reactive causes**
 - **Infections**
 - **Inflammatory/autoimmune diseases**



Mister M, 72 yo, 1 year later

- Fatigue
- Weight Loss
- Splenomegaly

Parameter	Apr 2025	Apr 2026	Ref Range
WBC × 10 ⁹ /L	12.0	28.5	4.5–11.0
Absolute Monocytes	1.2	4.6	0.2–0.8
Monocyte %	23%	26%	2–8%
Neutrophils	10.8	17.5	1.8–7.7
Hb g/dL	11.2	8.4	13.5–17.5
MCV fL	101	104	80–100
Platelets × 10 ⁹ /L	142	68	150–400
Peripheral Blasts	0	4%	0%
LDH U/L	270	420	140–280

CMML symptoms and clinical features

Symptoms	No symptoms	~20–30%
	Fatigue	~50–70%
	Weight loss	~10–20%
	Night sweats	~10–20%
	Fever (non-infectious)	~5–10%
Organomegaly	Splenomegaly	~25–50%
	Hepatomegaly	~10–20%
	Lymphadenopathy	~5–10%



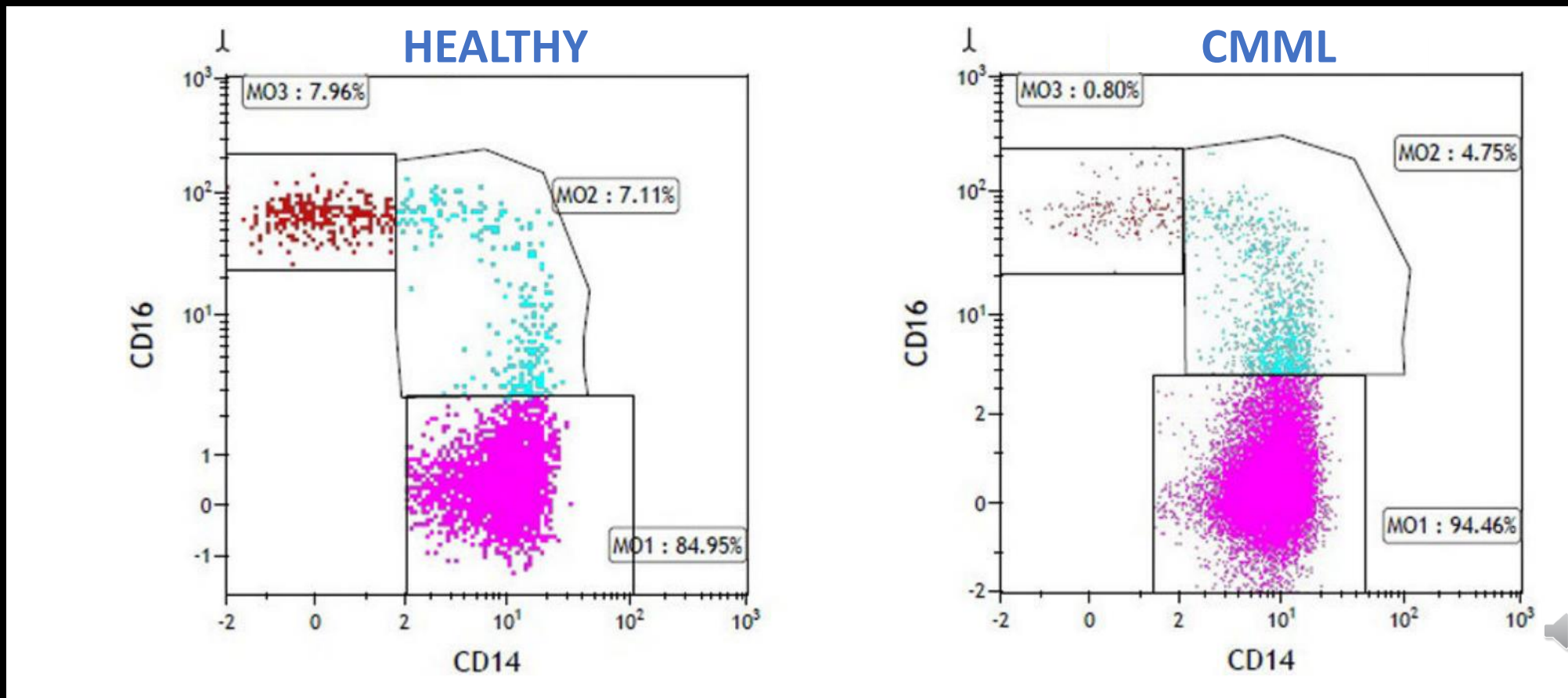
CMML-associated conditions

- **1. Immune dysregulation / inflammatory disorders (20–30%)**
 - Vasculitis (leukocytoclastic, large vessel...)
 - Inflammatory arthritis
 - Immune cytopenias (AIHA, ITP)
 - Serositis
 - Skin manifestations (Sweet syndrome...)
- **2. Clonal / associated hematologic conditions**
 - Systemic mastocytosis associated with CMML (SM-AHN)
- **3. Organ damage related to monocytosis**
 - Lysozyme nephropathy
 - Tissue infiltration (skin: leukemia cutis, lung, GI)



Step 2: Peripheral blood monocyte cytometry

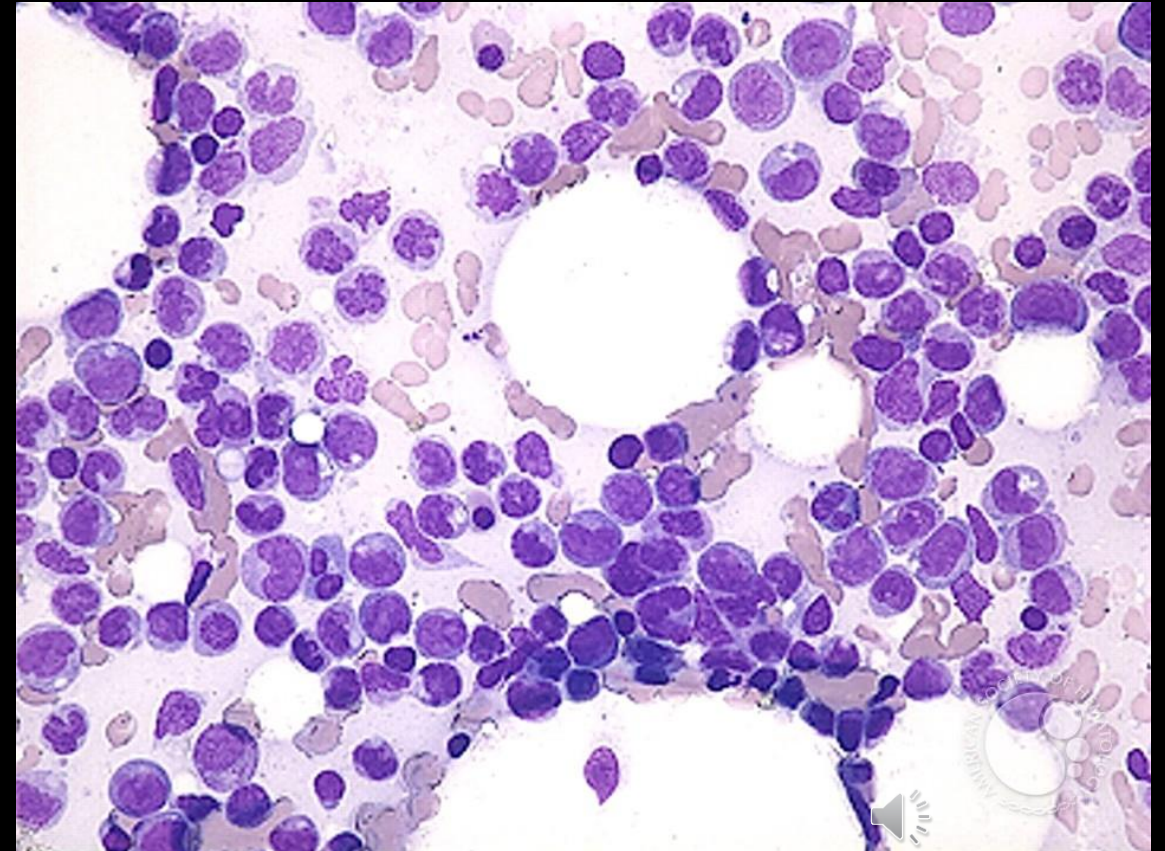
Classical monocytes CD14+ CD16- $\geq 94\%$
CMML diagnosis: se 90–92% sp 94-95%



Step 3: When to perform bone marrow and genetic testing

Indications

- Persistent unexplained monocytosis
- Cytopenias
- Circulating blasts
- Splenomegaly
or constitutional symptoms
- >94% classical monocytes



Step 4: Classification - Diagnostic criteria (WHO 2022)

Prerequisite criteria

1. **Persistent** absolute ($\geq 0.5 \times 10^9/L$) and relative ($\geq 10\%$) peripheral blood monocytosis.
2. **Blasts** constitute $< 20\%$ of the cells in the peripheral blood and bone marrow.^a
3. Not meeting diagnostic criteria of chronic myeloid leukaemia or other myeloproliferative neoplasms^b
4. Not meeting diagnostic criteria of myeloid/lymphoid neoplasms with tyrosine kinase fusions.^c

Supporting criteria

1. **Dysplasia** involving ≥ 1 myeloid lineages.^d
2. Acquired clonal cytogenetic or molecular abnormality.
3. Abnormal partitioning of peripheral blood monocyte subsets.^e

Requirements for diagnosis

- Pre-requisite criteria must be present in all cases.
- If monocytosis is $\geq 1 \times 10^9/L$: one or more supporting criteria must be met.
- If monocytosis is ≥ 0.5 and $< 1 \times 10^9/L$: supporting criteria 1 and 2 must be met.

Persistent ≥ 3 months

Blasts = myeloblasts, monoblasts and promonocytes

= No MPN BM and/nor high burden MPN-mutations (BCR-ABL, JAK2, CALR or MPL)

If eosinophilia, No PDGFRA and PDGFRB

$\geq 10\%$ Dysplasia in at least 1 lineage

CD14+CD16neg $\geq 94\%$ without AI or infl^o



CMML diagnostic criteria (ICC 2022)

Same same but different

Monocytosis defined as monocytes $\geq 0.5 \times 10^9/L$ and $\geq 10\%$ of the WBC


Cytopenia (thresholds same as MDS)*

Blasts (including promonocytes) $< 20\%$ of the cells in blood and bone marrow

Presence of clonality: abnormal cytogenetics and/or presence of at least one myeloid neoplasm associated mutation of at least 10% allele frequency†

In cases without evidence of clonality, monocytes $\geq 1.0 \times 10^9/L$ and $> 10\%$ of the WBC, and increased blasts (including promonocytes),‡ or morphologic dysplasia, or an abnormal immunophenotype consistent with CMML would be required for its diagnosis.

International Consensus Classification of Myeloid Neoplasms and Acute Leukemias: integrating morphologic, clinical, and genomic data

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Bone marrow examination with morphologic findings consistent with CMML (hypercellularity due to a myeloid proliferation often with increased monocytes), and lacking diagnostic features of acute myeloid leukemia, MPN or other conditions associated with monocytosis§

No *BCR::ABL1* or genetic abnormalities of myeloid/lymphoid neoplasms with eosinophilia and tyrosine kinase gene fusions

Features

WHO 2022

ICC 2022

Monocyte threshold	AMC $\geq 0.5 \times 10^9/L$ and monocytes $\geq 10\%$ of WBC differential	
Duration of monocytosis	Persistent (≥ 3 months)	
Blast/promonocyte threshold	20% blasts (including promonocytes) in PB and BM	
Clonality requirement	Cytogenetic or molecular evidence of clonality (acquired mutation or cytogenetic)	
Monocyte immunophenotyping	Increase in classical monocytes ($\geq 94\%$) supportive but not required	
Exclusions	Absence of <i>BCR::ABL1</i> , <i>PDGFRA/PDGFRB/FGFR1</i> rearrangements, <i>PCM1::JAK2</i> ; absence of AML-defining genetic abnormalities	
Cytopenia requirement	Not an explicit separate criterion	≥ 1 cytopenia (ineffective hematopoiesis)
Morphologic criteria	BM morphology consistent with CMML; dysplasia in ≥ 1 lineage	More stringent morphologic criteria (myeloid proliferation and monocytes) \rightarrow fewer cases qualifying as CMML compared to WHO
Practical impact	More oligomonocytosis cases as CMML than ICC.	More oligomonocytosis cases as MDS, MDS/MPN-NOS than WHO. 
Prognostic impact	NO DIFFERENCE	

Step 5: Classify the CMML subtype

COMPLEMENTARY

WHO 2022°
Blast-based

ICC 2022
WBC-based

CMML-1	CMML-2*	CMML-MD	CMML-MP*
PB blasts <5%	5-19%	WBC <13 x10 ⁹ /L	WBC ≥13 x10 ⁹ /L
BM blasts <10%	10-19% Auer rods+		

*higher risk of AML transformation 

Clonal monocytosis of undetermined significance (CMUS) – ICC 2022

= Clonal hematopoiesis *AND* monocytosis *AND* who do not meet the diagnostic criteria for CMML or any other myeloid neoplasm

Persistent monocytosis defined as monocytes $\geq 0.5 \times 10^9/L$ and $\geq 10\%$ of the WBC

Absence or presence of **cytopenia** (thresholds same as for MDS)*

Presence of at least one myeloid neoplasm associated mutation of appropriate allele frequency (ie, $\geq 2\%$)†

No significant dysplasia, increased blasts (including promonocytes) or **morphologic findings of CMML** on bone marrow examination‡

No criteria for a myeloid or other hematopoietic neoplasm are fulfilled

No reactive condition that would explain a monocytosis is detected

Clonal Cytopenia and Monocytosis of Undetermined Significance (CCMUS)

Dysplasia ≥ 1 lineage *AND/OR*
Hypercellularity, myeloid predominance +/- monocytes +/- monoblasts +/- blast equivalents



Mister M, 72 yo, 1 year later

Monocyte cytometry

Classical monocytes CD14+ CD16^{neg}: 97% (N 80-90)


Myelogram

- Hypercellular (80% cellularity)
- Monocytic proliferation
- Dysplasia in ≥ 1 lineages
- Blasts + promocytes : 6%

Diagnosis CMML-1, Proliferative subtype

Cytogenetics Normal karyotype (46, XY)

Molecular Studies TET2, SRSF2, ASXL1

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Molecular landscape

Gene	Frequency mutation	Prognostic impact
TET2*	60%	Favorable (if no adverse mutation)
SRSF2**	50%	Neutral
ASXL1*	40%	Adverse
NRAS***	15%	Adverse
RUNX1****	15%	Adverse
DNMT3A*	15%	Adverse
SETBP1	10%	Adverse

Co-mutation TET2+ SRSF2
Highly specific of CMML

- * Epigenetic regulators
- ** Spliceosome genes
- *** Signaling pathway
- **** Transcription factor

Step 6: Risk stratification

1. Such E. et al. Blood 2013 Apr 11;121(15):3005-15.
2. Elena C. et al. Blood 2016 Sep 8;128(10):1408-17.
3. Tefferi A. et al. Blood 2025 Aug 14;146(7):874-886.
4. Lanino L. et al. JCO 2026 Mar 27;JCO2502116.

Variable	CPSS ¹ (2013)	CPSS-Mol ² (2016)
WBC ≥13	✓	✓
Hb / TD	RBC-TD	RBC-TD
PLT		
Blasts	✓ BM	✓ BM
Cytogenetics	✓	✓
Molecular		✓ ASXL1, RUNX1, NRAS, SETBP1
Risk groups	4	4
mOS (mo)	72/31/18/12	144/59/31/18
1ary purpose 2ary	OS AML transformat°	OS AML transformat°
Best value	Limited resources	HSCT if no BLAST-Mol or iCPSS



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Variable	CPSS ¹ (2013)	CPSS-Mol ² (2016)	BLAST ³ (2025)	BLAST-Mol ³ (2025)
WBC ≥13	✓	✓	✓	✓
Hb / TD	RBC-TD	RBC-TD	Hb	Hb
PLT				
Blasts	✓ BM	✓ BM	✓ PB ≥2%	✓ PB ≥2%
Cytogenetics	✓	✓		✓
Molecular		✓ ASXL1, RUNX1, NRAS, SETBP1		✓ favorable: TET2, PHF6 Adverse: ASXL1, RUNX1, NRAS, SETBP1, DNMT3A, TP53, U2AF1, BCOR, PTPN11
Risk groups	4	4	3	3
mOS (mo)	72/31/18/12	144/59/31/18	63/28/13	N/A
1ary purpose 2ary	OS AML transformat°	OS AML transformat°	OS -	OS -
Best value	Limited resources	HSCT if no BLAST-Mol or iCPSS	Non transplant monitoring	HSCT

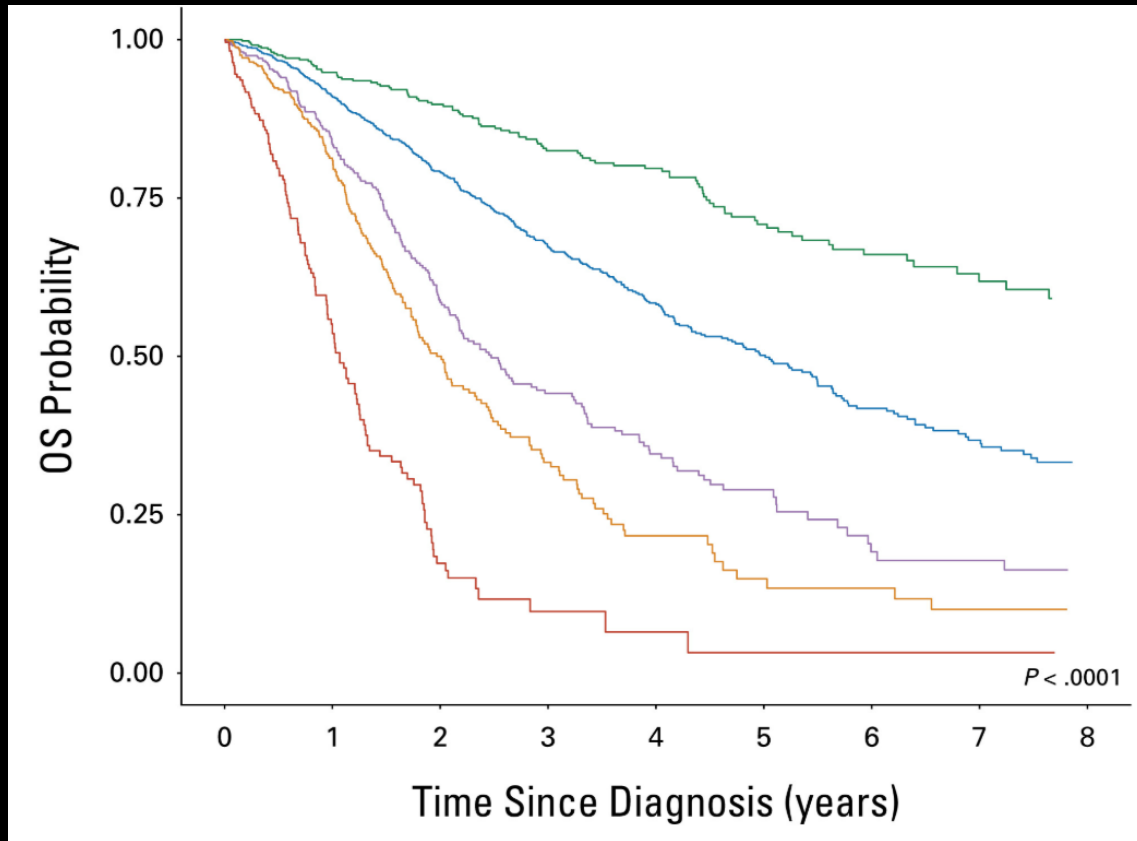


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Variable	CPSS ¹ (2013)	CPSS-Mol ² (2016)	BLAST ³ (2025)	BLAST-Mol ³ (2025)	iCPSS ⁴ (2026)
WBC ≥13	✓	✓	✓	✓	✓
Hb / TD	RBC-TD	RBC-TD	Hb	Hb	Hb
PLT					✓
Blasts	✓ BM	✓ BM	✓ PB ≥2%	✓ PB ≥2%	✓ BM
Cytogenetics	✓	✓		✓	✓
Molecular		✓ ASXL1, RUNX1, NRAS, SETBP1		✓ favorable: TET2, PHF6 Adverse: ASXL1, RUNX1, NRAS, SETBP1, DNMT3A, TP53, U2AF1, BCOR, PTPN11	✓ ASXL1, RUNX1, TET2, SETBP1, DNMT3A, TP53, EZH2, STAG2, U2AF1
Risk groups	4	4	3	3	5
mOS (mo)	72/31/18/12	144/59/31/18	63/28/13	N/A	N/A
1ary purpose 2ary	OS AML transformat°	OS AML transformat°	OS -	OS -	OS HSCT timing
Best value	Baseline, no NGS	HSCT if no BLAST-Mol or iCPSS	Blood → routine monitoring	HSCT	Transplant timing decisions

iCPSS (2026)



Risk	OS	HSCT timing
Very Low	~10y	No upfront HSCT
Low	~5y	Deferred HSCT
Intermediate	~3y	Individualized decision
High	~2y	Early HSCT
Very high	~1y	Urgent HSCT



Mister M, 72 yo, CMML-1, MP

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BM blasts		6%	<2%

CPSS Mol	Blast-Mol	iCPSS
Int-2 mOS 37 mo	High 12 mo	Intermediate ~ 30 mo

Cytogenetics Normal karyotype (46, XY)

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ONLINE CALCULATORS

CPSS-Mol

- https://qxmd.com/calculate/calculator_609/cmml-cpss-mol

BLAST and BLAST-MOL

- <https://sudheshk.com/projects/blast-model>

iCPSS

- <https://icpss-risk.com/>



Step 7: Determine transplant eligibility

- Age and performance status
- Comorbidity index (HCT-CI)
- Donor availability → early HLA typing
- Risk score (CPSS-Mol, BLAST-mol, iCPSS)
- Patient preference and goals of care



Mister M, 72 yo, CMML-1, MP

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Peripheral Blasts	0	4%	0%
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CPSS Mol	Blast-Mol	iCPSS
Int-2 mOS 37 mo	High 12 mo	Intermediate ~ 30 mo

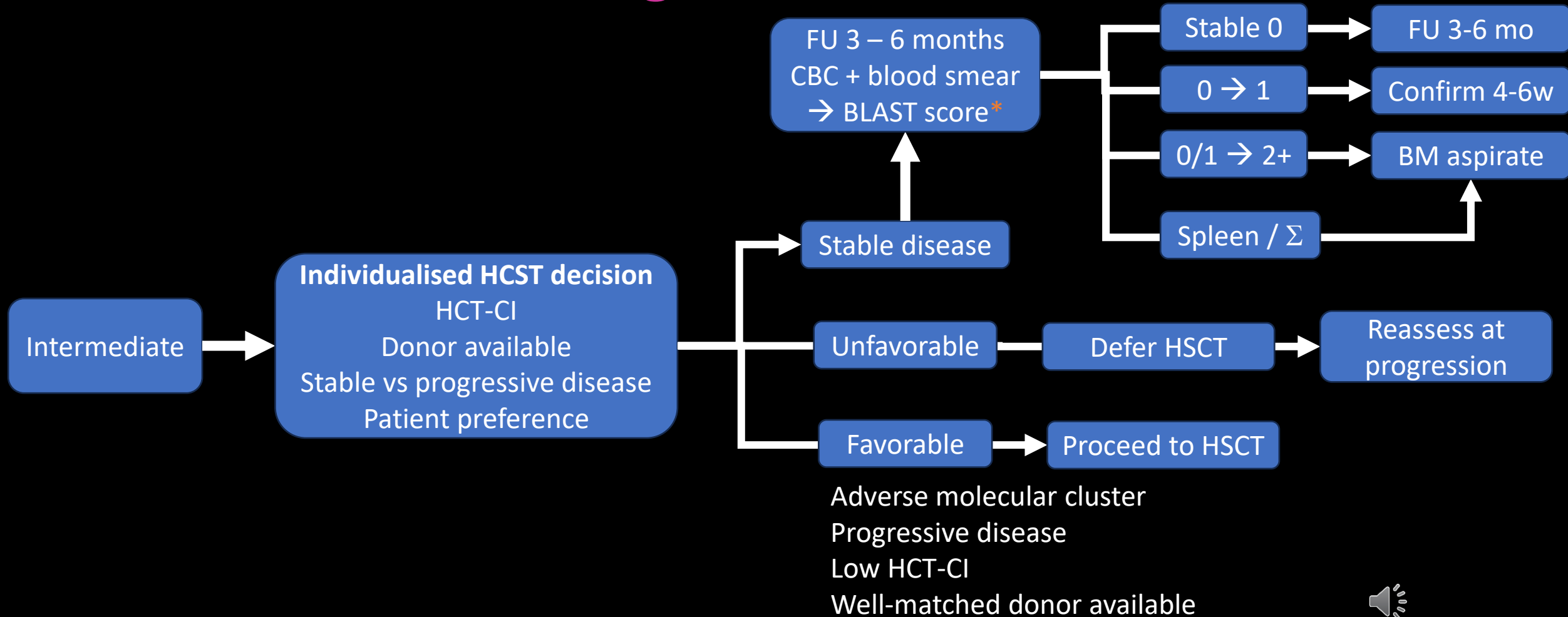
Cytogenetics Normal karyotype (46, XY)

Molecular Studies TET2, SRSF2, ASXL1

→ Transplant should be discussed early but not performed immediately.

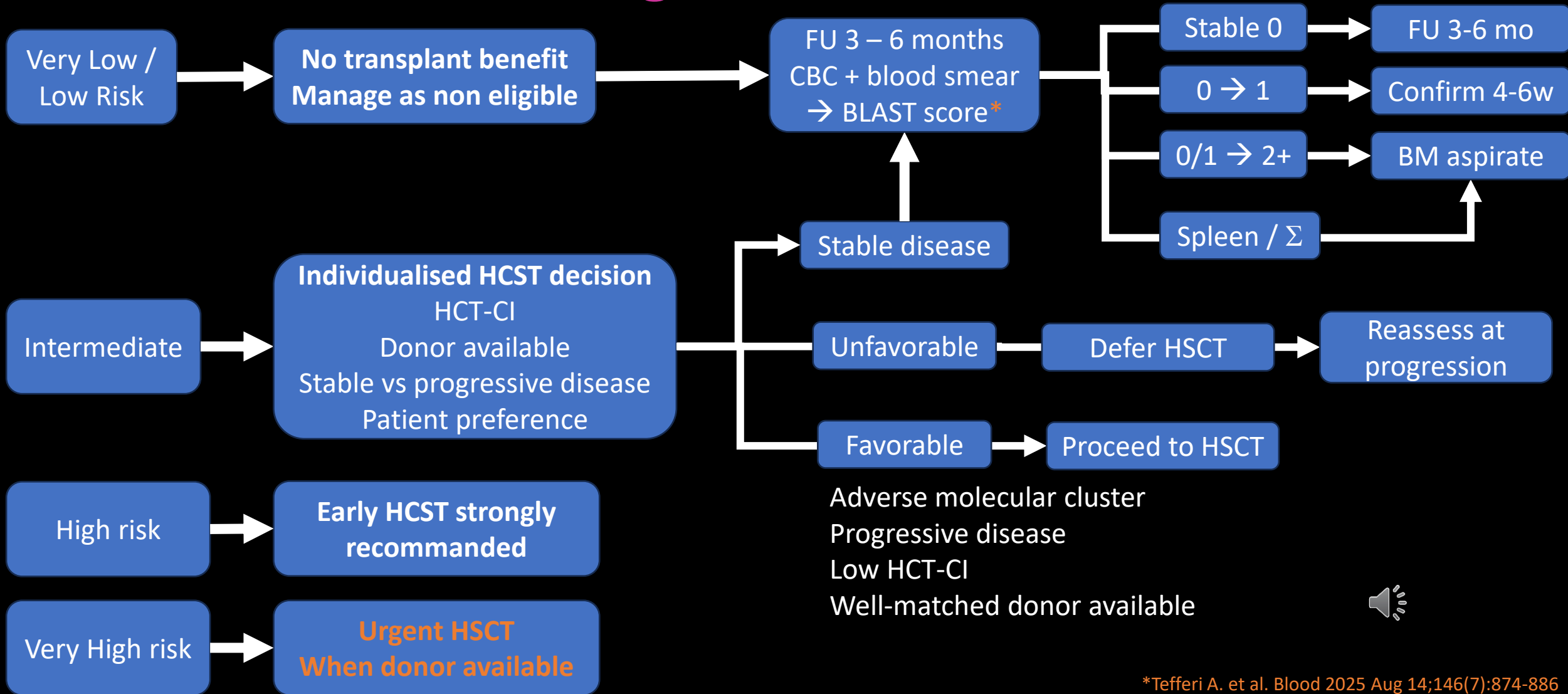
TRANSPLANT ELIGIBLE PATIENTS

Treatment according to iCPSS risk assessment



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Treatment according to iCPSS risk assessment

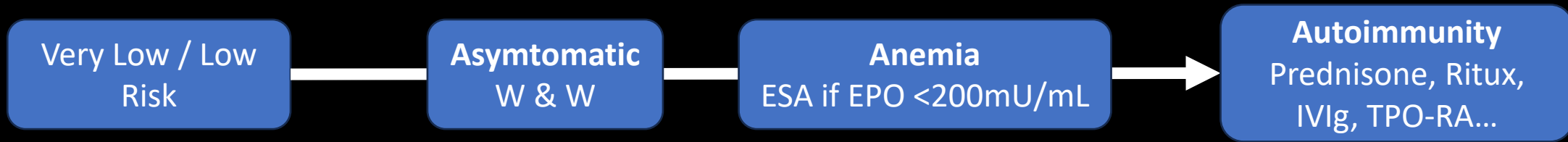


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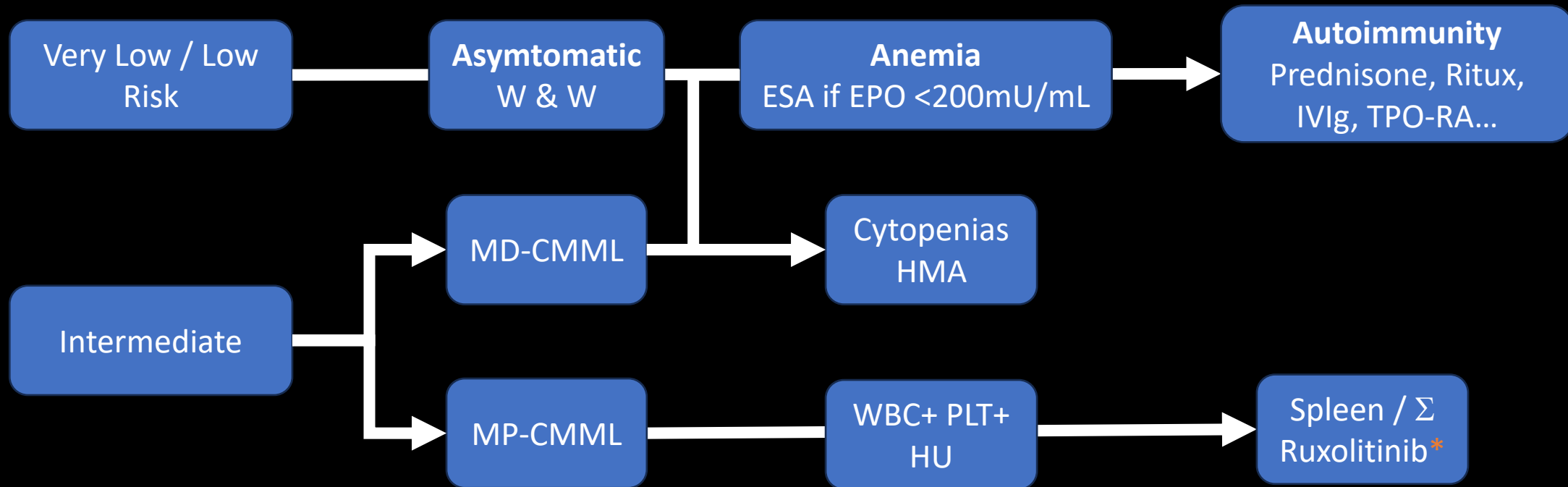
Step 8: Treatment of non transplant eligible patients according to iCPSS risk assessment

... in an ideal setting ...



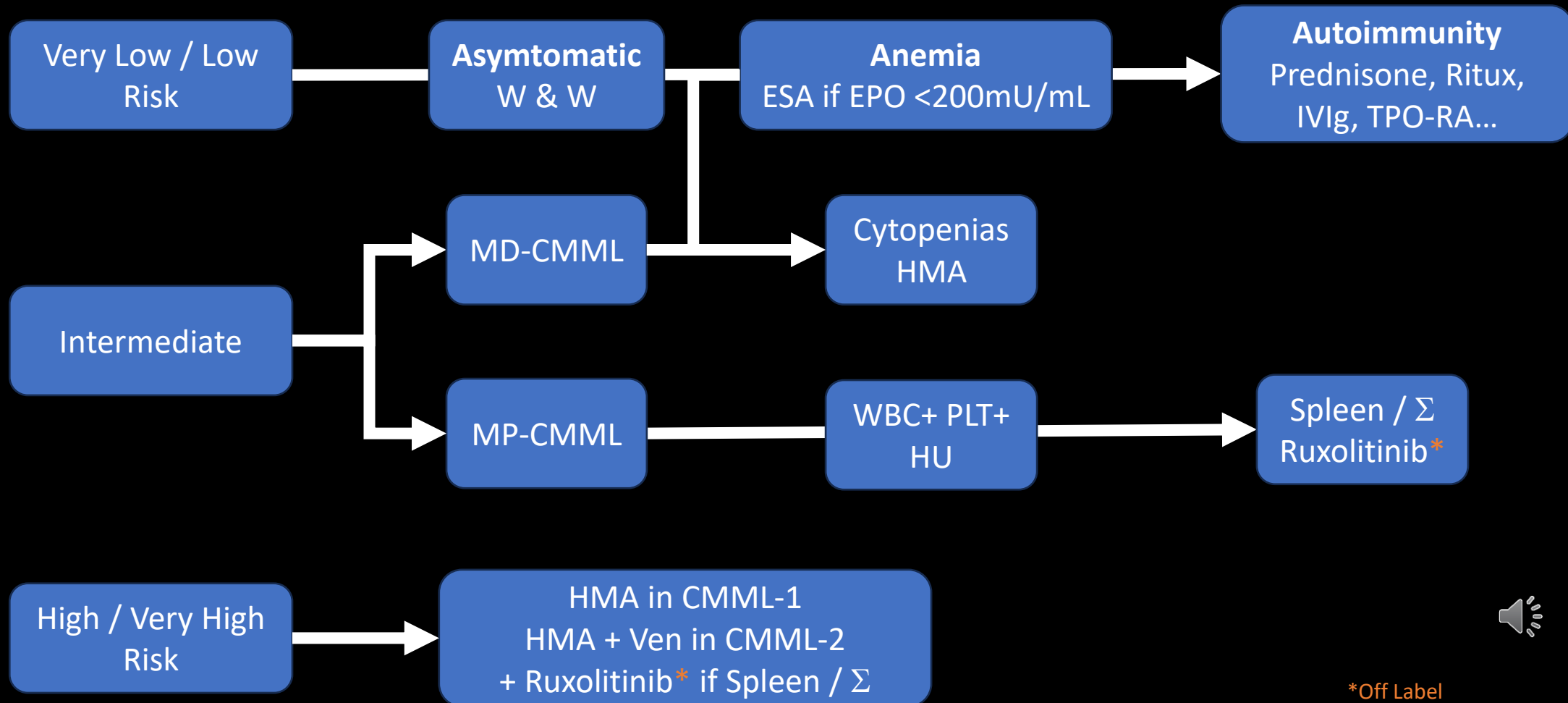
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... in an ideal setting ...



Step 8: Treatment of non transplant eligible patients according to iCPSS risk assessment

... in an ideal setting ...



Broader use of HMA across CMML subtypes

- **Multicenter retrospective**
- N 949 CMML
- First-line treatments
 - **HMA 43% (n=412)**
 - **Hydroxyurea 41% (n=391)**
 - **Intensive chemotherapy 9% (n=83)**

Subgroup	N (%)	HMA OS Benefit?	mOS: HMA vs. non-HMA	p-value
MD-CMML, blasts 10%	182 (21%)	No	No significant difference	NS
MD-CMML, blasts $\geq 10\%$	111 (13%)	Yes	23.8 vs. 9.4 months	0.0014
MP-CMML, blasts 10%	345 (40%)	Yes	24.7 vs. 13.1 months	0.0002
MP-CMML, blasts $\geq 10\%$	219 (26%)	Yes	15.5 vs. 6.6 months	0.0002

- mOS **HMA 20.7** vs mOS **HU 15,6mo** (p=0.0002)
- mOS **HMA 20.7** and mOS **intensive CT 14.0 mo** (p=0.0027)



Broader use of HMA across CMML subtypes?

- **Dacota trial: Randomized Phase 3**
- **1st line Decitabin vs HU**
- **Advanced MP-CMML: WBC >13 + at least 2**
 - BM blasts ≥5%,
 - Clonal cytogenetic abnormality,
 - Hemoglobin <10 g/dL,
 - Neutrophils >16 × 10⁹/L,
 - Platelets <100 × 10⁹/L,
 - Splenomegaly,
 - ECOG ≤2
- **Dissociation response rate / survival**
 - **Death without progression +++**
 - Treatment-related toxicity (infections...)

Endpoint	Decitabine (n=84)	Hydroxyurea (n=86)	p-value
Primary: Median EFS	12.1 months	10.3 months	0.27
Median OS	18.4 months	21.9 months	0.67
Overall Response Rate (ORR)	63% (53/84)	35% (30/86)	0.0004
Risk of AML transformation/progression	Reduced	Reference	0.005
Risk of death without progression	Increased	Reference	0.04
Median follow-up	17.5 months	17.5 months	—

Role of stringent cytoreduction?

- *Ancillary analysis of the DACOTA trial*
- **Monocytosis** $>1 \times 10^9/L$ or **Leucocytosis** $>10 \times 10^9/L$ after 6 cycles
→ **Increased death** x5.38 ($p = 0.0003$).
- **Independent of:**
 - Treatment arm (decitabine vs. hydroxyurea)
 - Baseline CPSS risk category
 - Persistence of BM blast excess
- *Persistent myeloproliferation after treatment*
→ *Worsens prognosis*
→ *Irrespective of treatment arm or bone marrow response.*
- **To do: Prospective validation** in independent cohorts



CMML: what really matters

- It's not one disease → it's **many diseases**
- Prognosis is not enough → we need **decision tools**
- Timing is everything → especially for **transplant**
- Treatment is not standardized → it must be **personalized**
- **Controlling proliferation** might matter more than response



Thank you

*Good judgment comes from experience.
Experience comes from bad judgment.*

Rita Mae Brown



