







UPDATE ON PLASMA CELL NEOPLASMS Falko Fend University Hospital Tübingen

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Research support from Stemline

Speaker honoraria from Stemline and EUSAPharma







RECORDATI

PLASMA CELL NEOPLASMS

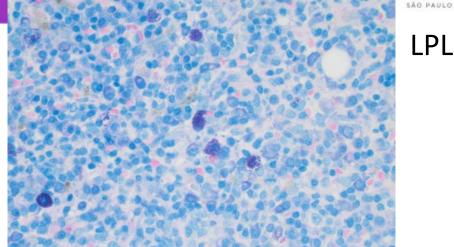


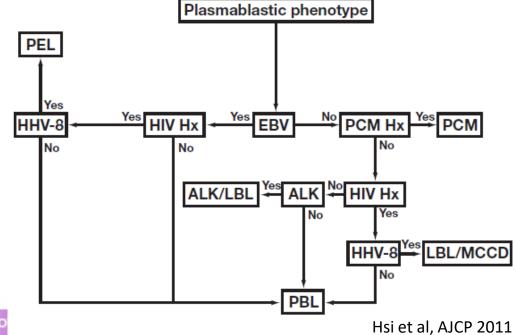
Plasma cells are the dominant and proliferating neoplastic population, +/- monoclonal immunoglobulin (M-protein)

Plasma cell tumors need to be differentiated from

- Small B cell neoplasms with plasma cell differentiation
- Terminally differentiated aggressive B-cell lymphomas with plasmablastic phenotype (PBL, PEL, ALK+ LBCL)

Multiple myeloma 10% of all hematological neoplasms, 1% of all cancers

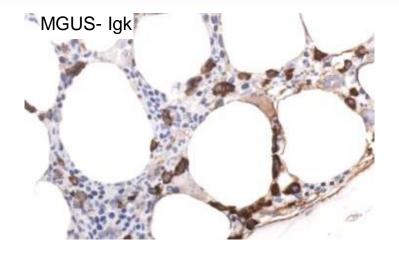




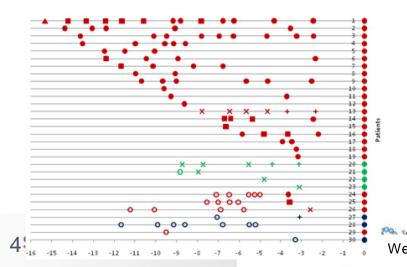


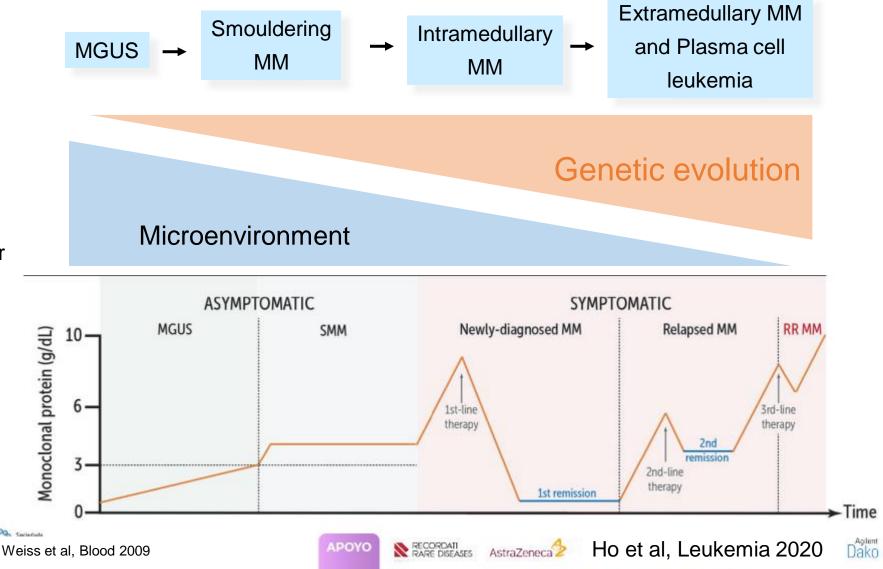
SYSTEMIC PLASMA CELL DISORDERS SHOW STEPWISE EVOLUTION





MGUS is (virtually) universal precursor to MM – progression rate 1%/year

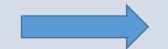






Evolution from clinically inapparent precursors through an indolent (smoldering) disease phase to manifest organ damage

Alternative occurrence as localized disease (plasmacytoma)



Integration of clinical, laboratory and imaging parameters for diagnosis

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International Myeloma Working Group (IMWG) consensus criteria form basis for classification plasma cell neoplasms in both ICC and WHO classifications







Clinical symptoms caused by end organ damage (CRAB) and biomarkers of malignancy (SLIM)

HyperCalcemia (>0.25 mmol/L (>1 mg/dL) above normal or >2.75 mmol/L (>11 mg/dL))

Renal insufficiency (creatinine clearance <40 mL or serum creatinine>177 µmol/L (>2 mg/dL))

Anemia (Hb >20g/l below normal or <100g/L)

Bone lesions (one or more lytic lesions on Rx, CT or PET-CT)

Sixty % BM plasma cells Serum free Light chain ratio >100

>1 focal lesion on MRI

*Rajkumar et al, Lancet Oncol 2014





IMWG CONSENSUS CRITERIA FOR PLASMA CELL NEOPLASMS

Non-IgM MGUS

Monoclonal serum protein <30 g/L <10% plasma cells in BM

No PCM-related end-organ damage

Smouldering (asymptomatic) myeloma

>10% and <60% plasma cells in BM and/or paraprotein >30 g/L 500mg/24h urine Absence of myeloma-defining events or amyloidosis

Symptomatic myeloma

>10% plasma cells in BM (or plasmacytoma) Monoclonal protein in serum or urine Myeloma-defining events End organ damage Biological markers of malignancy

*Rajkumar et al, Lancet Oncol 2014

Determination of PC count by cytology (not FCM!) and histology, higher value counts

3º CONGRESO LATINOAMERICAN HEMATOPATO

Incidence 3-4% >50y, >5% >70y Progression risk 1%/year (1.5% for IgA, 0.3% for LC)











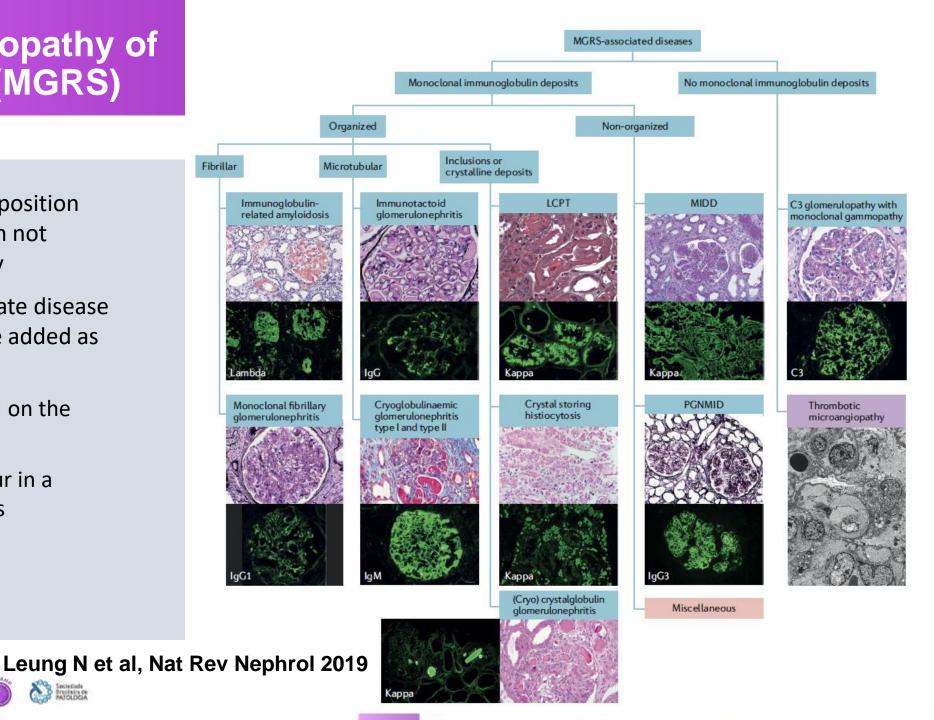
Monoclonal Gammopathy of renal significance (MGRS)

Renal damage caused by IG deposition due to clonal B/PC proliferation not fulfilling criteria for malignancy

MGRS not introduced as separate disease entities in 2022 ICC, but can be added as qualifier

- ICC nomenclature focused on the neoplastic process
- MGRS and MGCS can occur in a variety of B cell neoplasms

Extended concept MGClinicalS



IMWG CONSENSUS CRITERIA FOR PLASMA CELL NEOPLASMS

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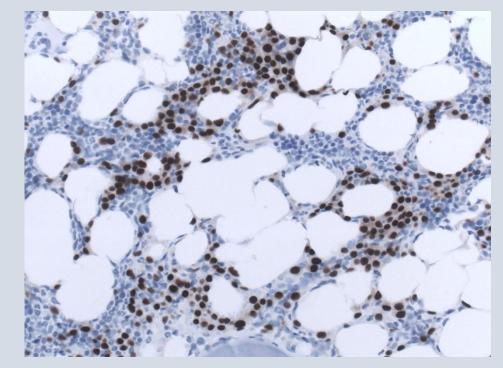
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*Rajkumar et al, Lancet Oncol 2014



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MUM1



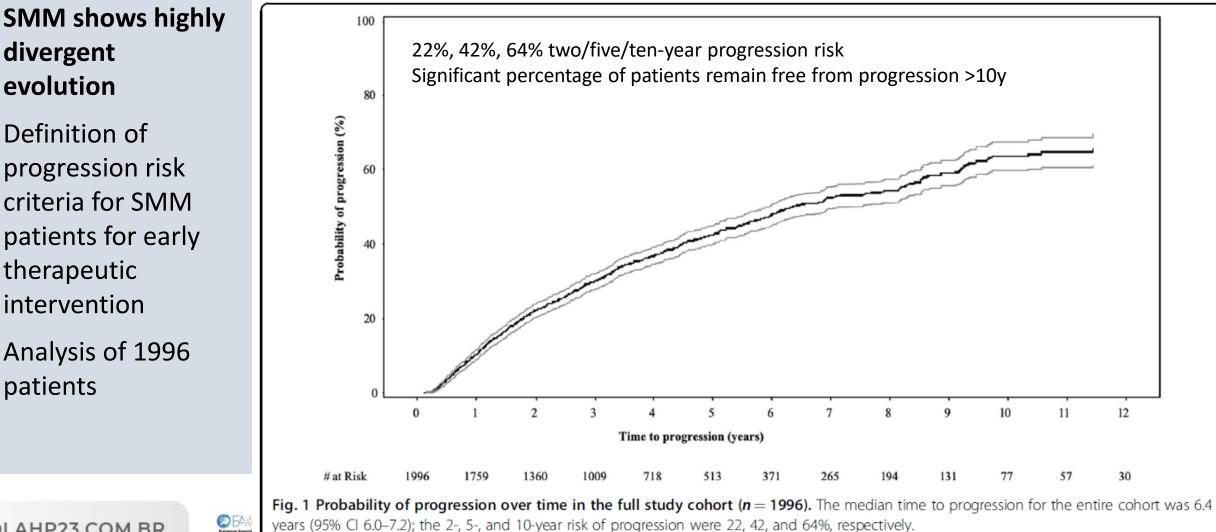








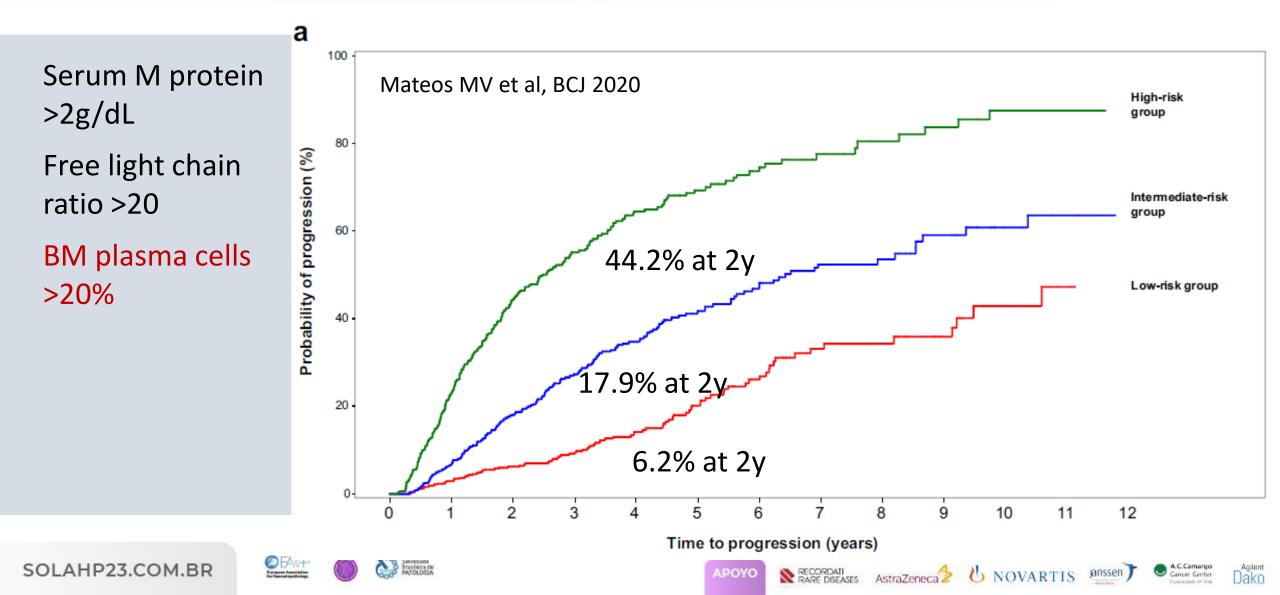
Mateos MV et al, BCJ 2020





3 PARAMETER MODEL (2/20/20) FOR DEFINING HIGH RISK SMM





IMWG CONSENSUS CRITERIA FOR PLASMA CELL NEOPLASMS

MGUS

Monoclonal serum protein <30 g/L <10% plasma cells in BM No PCM-related end-organ damage

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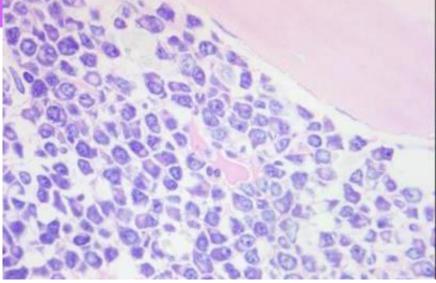


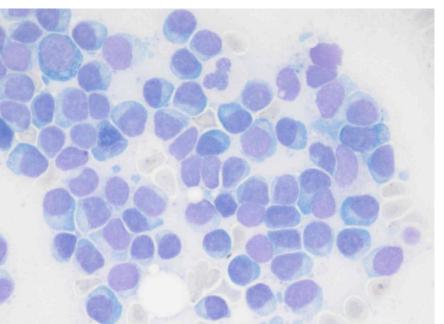
Jako

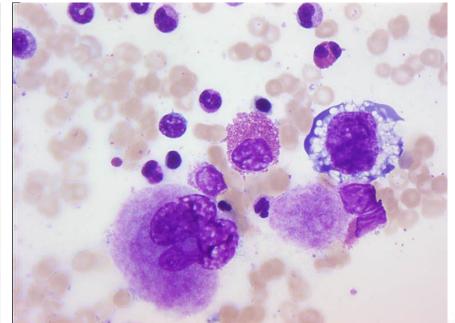
3º CONGRESO LATINOAMERICANO DE HEMATOPATO

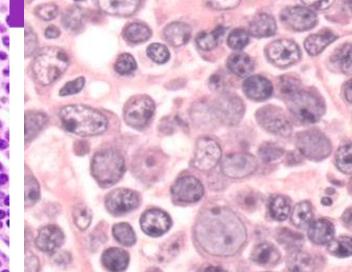
MM DIAGNOSIS IN PATHOLOGY – ALWAYS EASY?

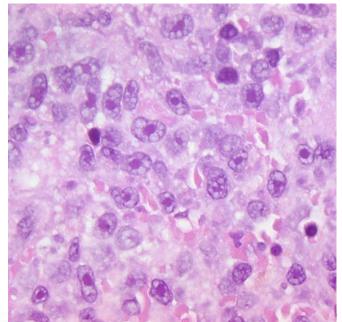


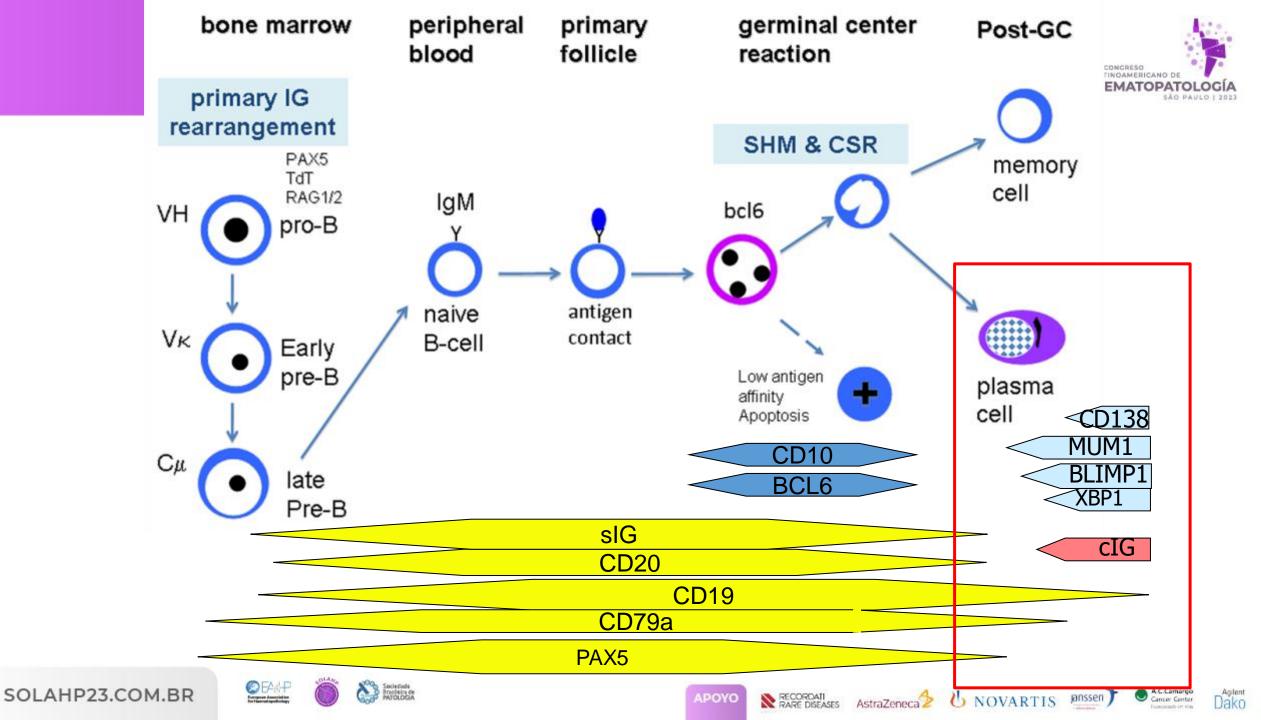












Relevant markers of plasma cell differentiation



CD138

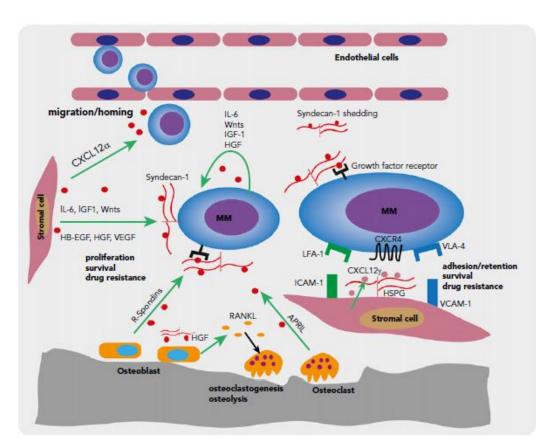
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CD138 (syndecan)

Transmembrane heparan sulfate proteoglycan (HSPG)

Provides alternative mode of interaction with microenvironment in the absence of BCR signalling



Ren Z et al, Blood 2021

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Sometimes variable intensity

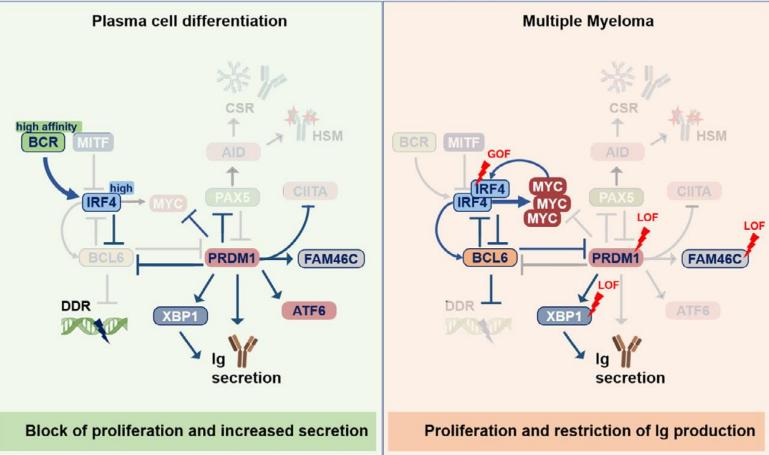
Expressed by epithelial cells

MUM1/IRF4



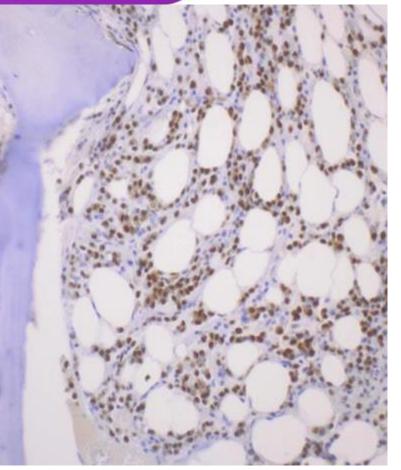
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Master regulator of PC development and key survival factor



Perrini et al, FEBS J 2021

Gain of function mutations in *IRF4* and loss of function mutations in XBP1 and PRDM1 shift from secretion to proliferation



Very reliable PC marker Expression in terminal diff. B cell neoplasms and ALCL

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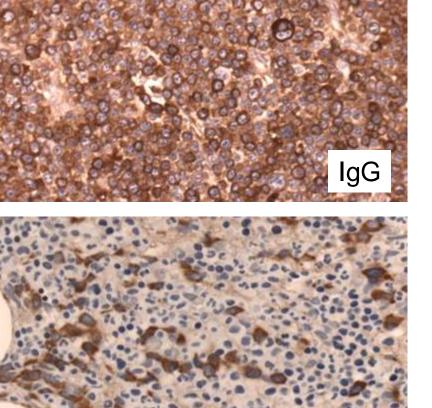
Immunoglobulin light and heavy chains



lambda

Monoclonal light chain expression by virtually all neoplastic plasma cells

- Expression levels often much lower than in normal PC, interpretation may be difficult – CISH as alternative
 IgG 50-60%, 15% IgA, 15-20% light chain only
 Rare MM express IgD, IgM or IgE
 - Biclonality (including both light chains) common in MGUS (5%)[§], but rare in established MM (0.91%)*
 - Likely suppression of second clone similar to immunoparesis (reduction in polyclonal IG)



§ Landgren et al 2014; *Campbell et al, BJC 2017

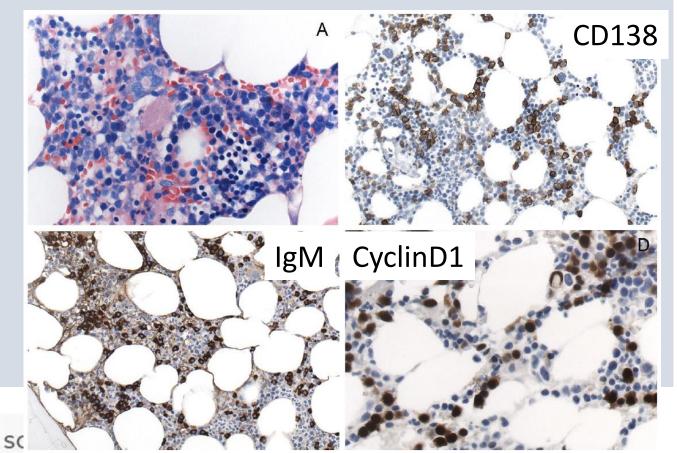


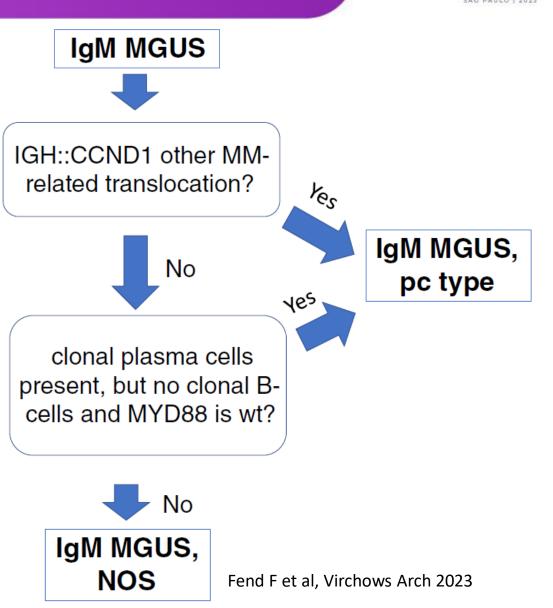
IGM MGUS AND IGM MM



Higher frequency of t(11;14)

Separation from LPL by lack of MYD88^{mut,} presence of clonal B cells and aberrant immunophenotype of PC



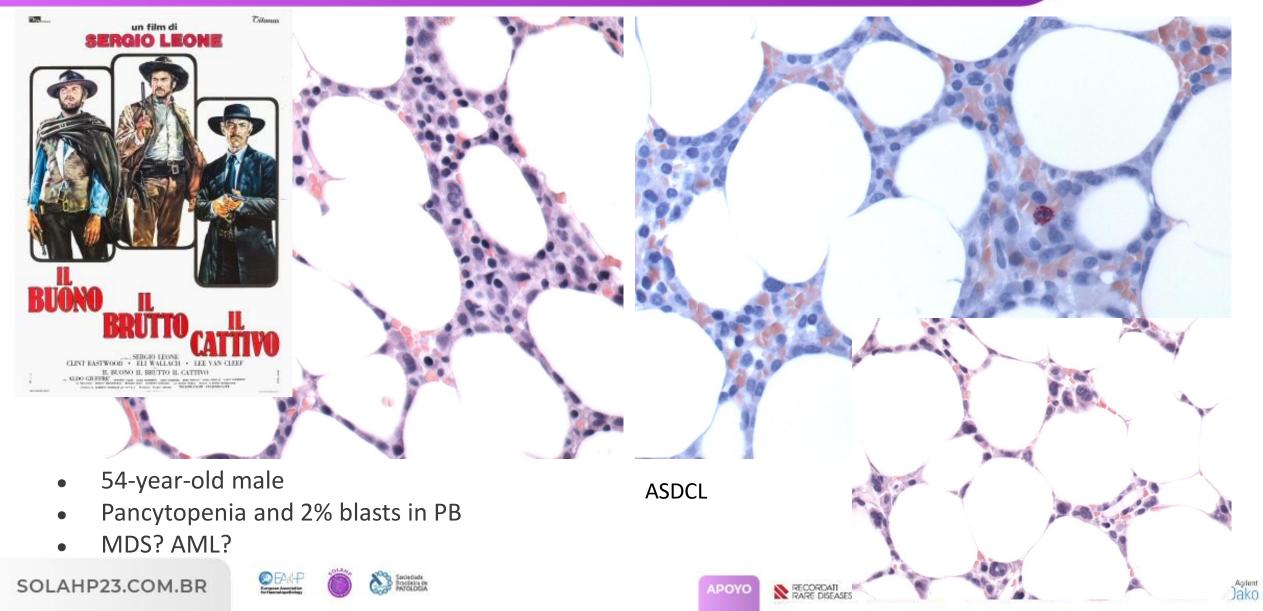


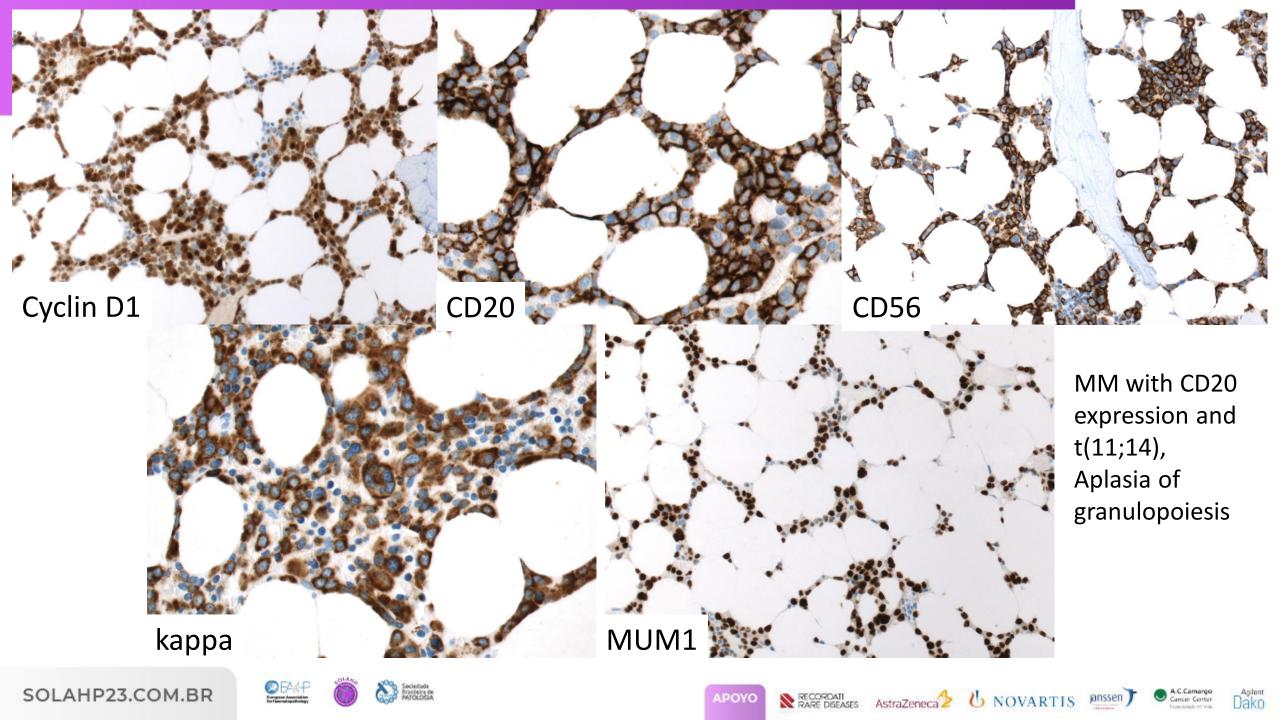


 $\geq 10\%$ CP in BM + SLiM biomarkers/CRAB criteria True non-secretory MM rare (<1-2%) M-protein (in IFX) Frequency decreased due to increased no , yes sensitivity of diagnostic tests (FLC) sFLC secretory MM Diagnosis of non-secretory MM requires yes no determination of serum free light chain ratio oligosecretory MM non-secretory MM Lower frequency of renal complications Ig detectable in the cytoplasm and paraneoplasia lno 15% 85% yes Charlinski et al, Adv Clin Exp Med 2022 non-producing MM true non-secretory MM Sociedade Brasileira de PATOLOGIA SOLAHP23.COM.BR RARE DISEASES AStraZeneca APOYO Dako

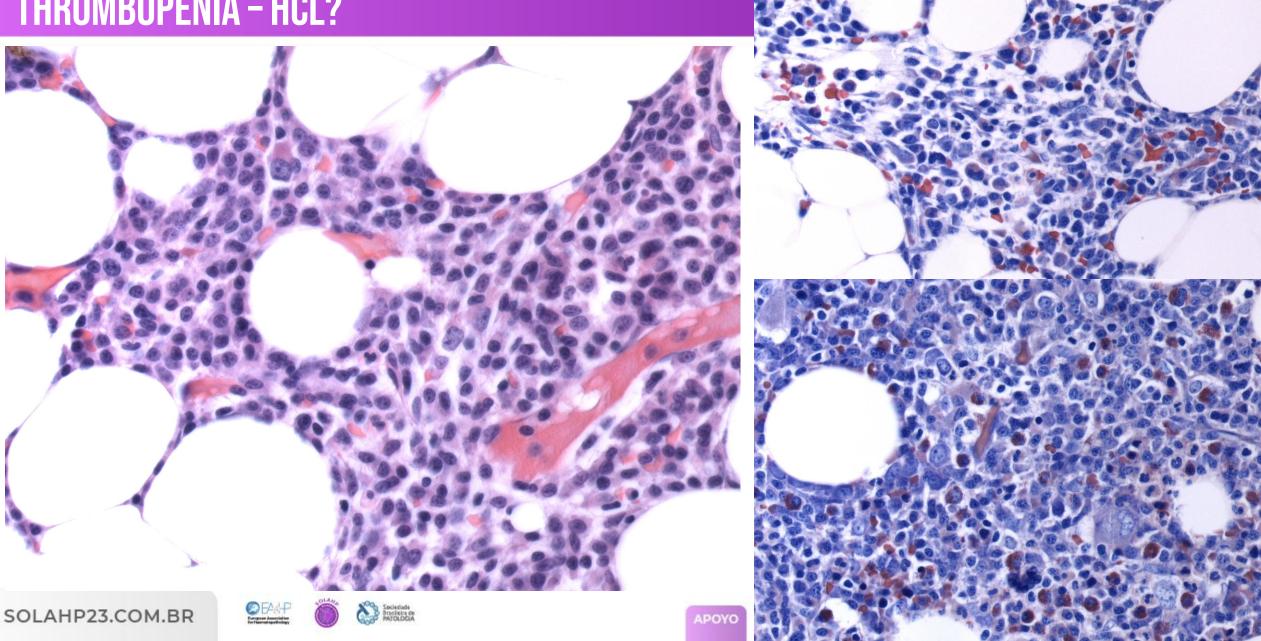
THE GOOD, THE BAD AND THE UGLY





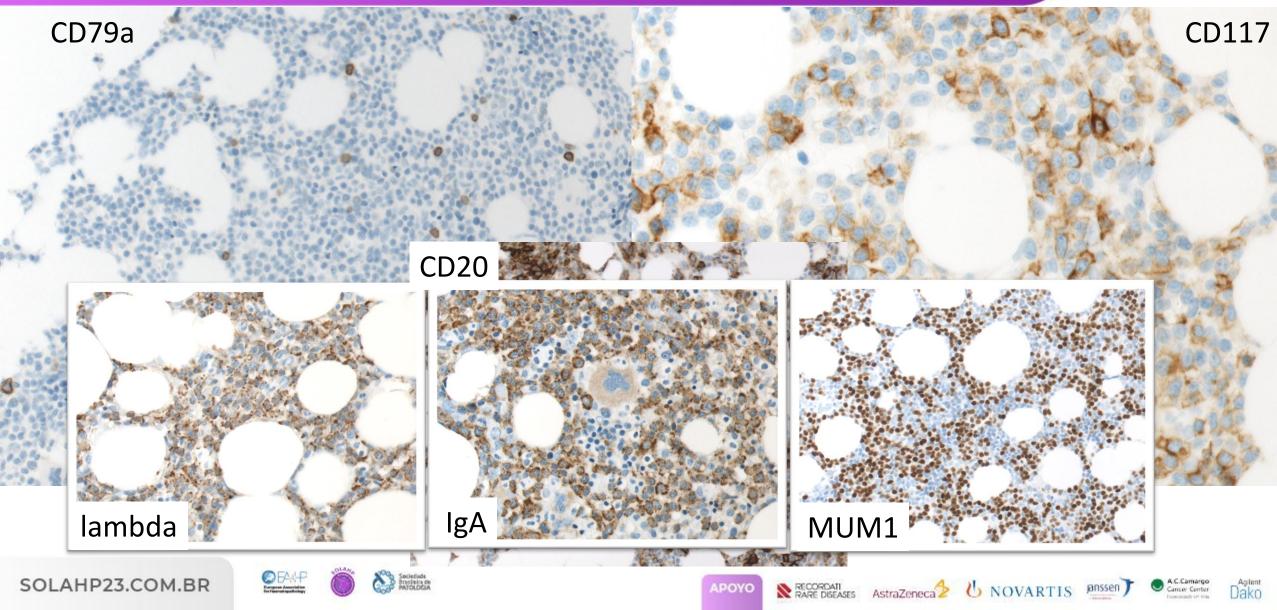


69-YEAR-OLD FEMALE, SPLENOMEGALY, THROMBOPENIA – HCL?



IGA PARAPROTEIN, DETECTION OF T(14;16) AND 13Q-





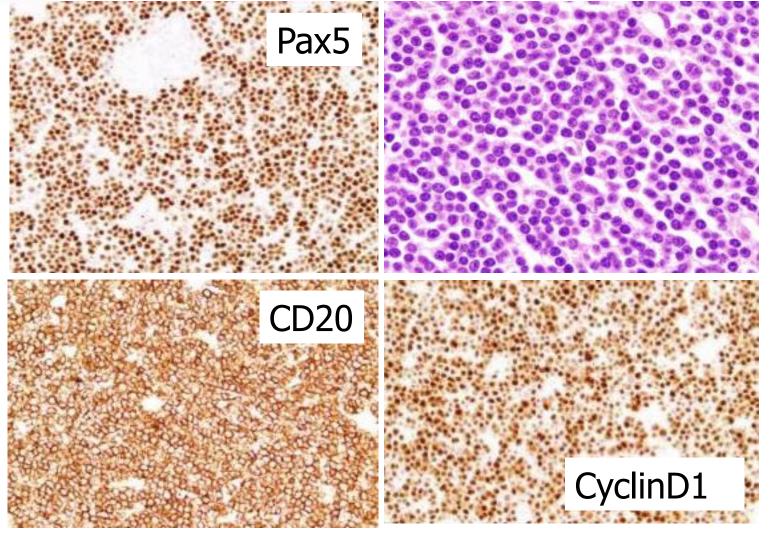
MM IMMUNOPHENOTYPE



Normal PC Negative: PAX-5, CD22, CD20 Positive: CD19, CD138, CD38, CD79a, cIG, CD45+/-, EMA

Malignant PC lack CD19 CD56 (75%), CD117 (3-30%), CD20 (20-30%), CyclinD1 (50%, 20% strong), PAX5 (rare), CD10, MYC

Partial preservation of B-cell program especially in t(11;14)+ PCM – sometimes in primary plasma cell leukemia





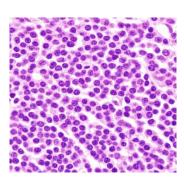
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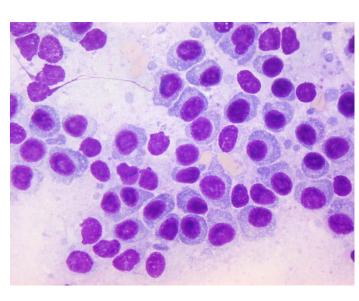
BORDERLANDS OF MM



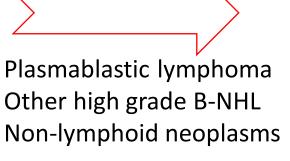


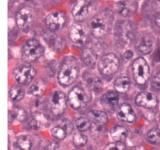
Lymphoplasmacytic morphology B-cell markers IgM MM

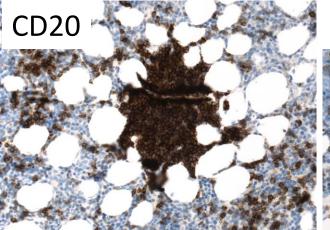
Lymphoplasmacytic lymphoma (non IgM) Mantle cell lymphoma Co-occurring B-NHL

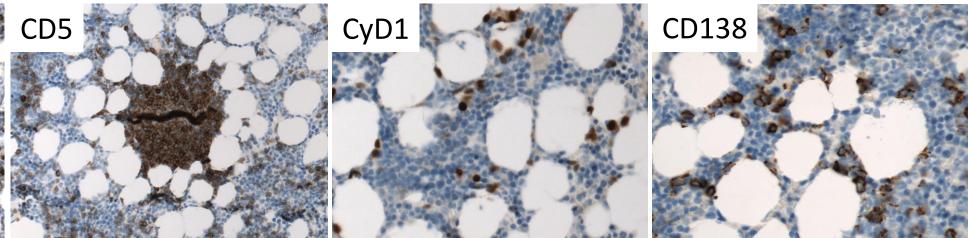


(Plasma-) blastic morphologyHigh proliferationLoss of PC markersEBV positivity





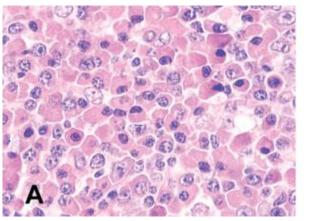




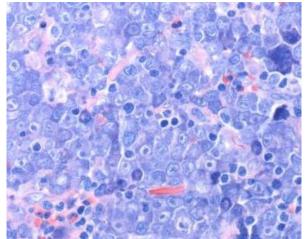
BORDERLANDS OF MM – EXTRAMEDULLARY PROLIFERATIONS OF PLASMACYTIC/PLASMABLASTIC MORPHOLOGY



	Plasmablastic MM	Plasmablastic lymphoma	Extraosseous plasmacytoma
Location	Any site, occ. leukemic	Extranodal, oral cavity	80% head & neck
Morphology	Plasmablastic/plas macytic	Plasmablastic large cell /immunobl.	Usually mature
M-protein	Most cases	Rare	30%, low level
Osteolytic lesions	Common	Rare	Occasionally (skull)
Bone marrow	Yes	May occur	No or at MGUS level
CD56	CD56+	-/+	-/(+)
Cyclin D1 / t(11;14)	15-20%	Neg.	Neg.
MYC rearrangement	common	common	rare
EBV	Rare	60-70%	rare
Genetics	MM-type tx. and trisomies, RAS, P53	MYC translocations, JAK/STAT3 pathway, TP53	Trisomies and IGH- translocations, no t(11;14)



Primary extraosseous plasmacytoma



Plasmablastic lymphoma AstraZeneca b NOVARTIS

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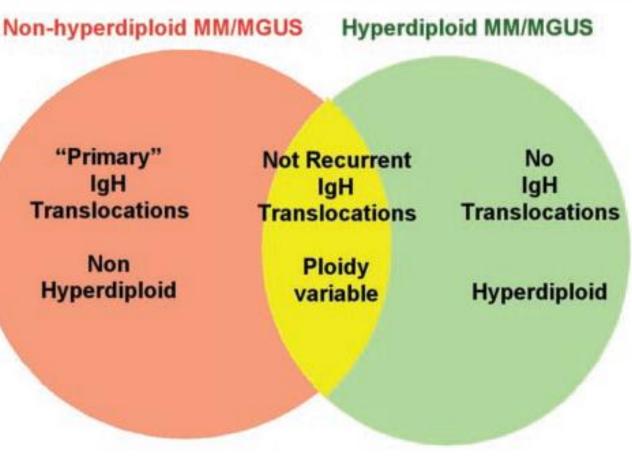
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GENETICS OF MULTIPLE MYELOMA



- MM is a genetically heterogeneous disease
 - 40-50 % show recurrent translocations
 - 45% show trisomies of uneven chromosomes (3,5,7,9,11,15,19,21)
- These primary alterations are already present in MGUS



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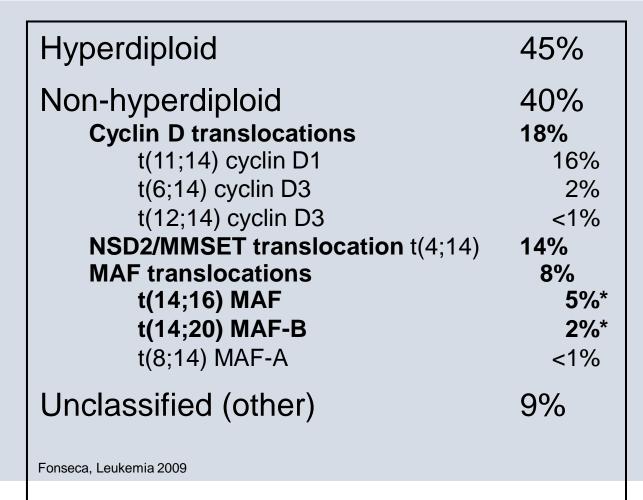
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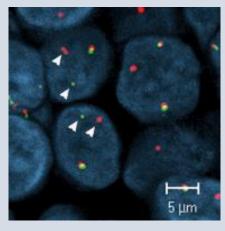
Fonseca et al, CCR 2004

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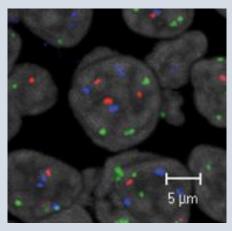
IMWG cytogenetic classification of MM





Break 14q32

Minimal Panel t(4;14) FGFR3/NSD2 t(14;16) und t(14;20) MAF Del 17p TP53



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Trisomy 9 and 15

Comprehensive Panel t(11;14) CCND1/cyclin D1 Del 13 Ploidy Chromosome 1 alterations

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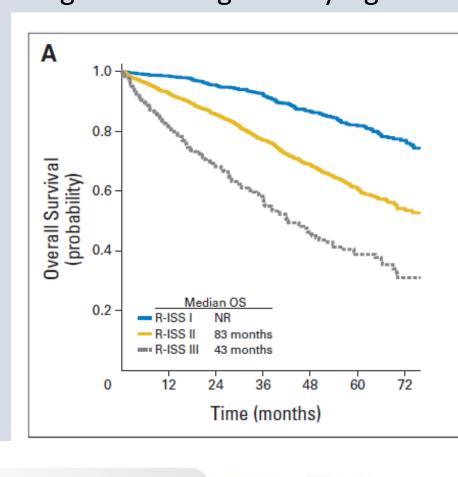




Revised International Staging System for Multiple Myeloma: A Report From International Myeloma Working Group

Sociedade Brasileira de PATOLOGIA

Integration of high risk cytogenetics provides improved stratification in MM and SMM



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Prognostic Easter	Criteria
Prognostic Factor	Criteria
ISS stage	
T	Serum β₂-microglobulin < 3.5 mg/L, serum albumin ≥ 3.5 g/dL
II	Not ISS stage I or III
III	Serum β_2 -microglobulin \geq 5.5 mg/L
CA by iFISH	
High risk	Presence of del(17p) and/or translocation t(4;14) and/or translocation t(14;16)
Standard risk	No high-risk CA
LDH	
Normal	Serum LDH < the upper limit of normal
High	Serum LDH > the upper limit of normal
A new model for risk stratification for MM	
R-ISS stage	
1	ISS stage I and standard-risk CA by iFISH and normal LDH
I	Not R-ISS stage I or III
Ш	ISS stage III and either high-risk CA by iFISH or high LDH

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High risk: del 17p (TP53) t(4;14) t(14;16) **PFS**: l: 55 mo II: 36 mo III: 24 mo

VOLUME 33 · NUMBER 26 · SEPTEMBER 10 2015

JOURNAL OF CLINICAL ONCOLOGY

Palumbo et al JCO 2015, Mateos et al, BCJ 2021

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Mutational Spectrum, Copy Number Changes, and Outcome: Results of a Sequencing Study of Patients With Newly Diagnosed Myeloma



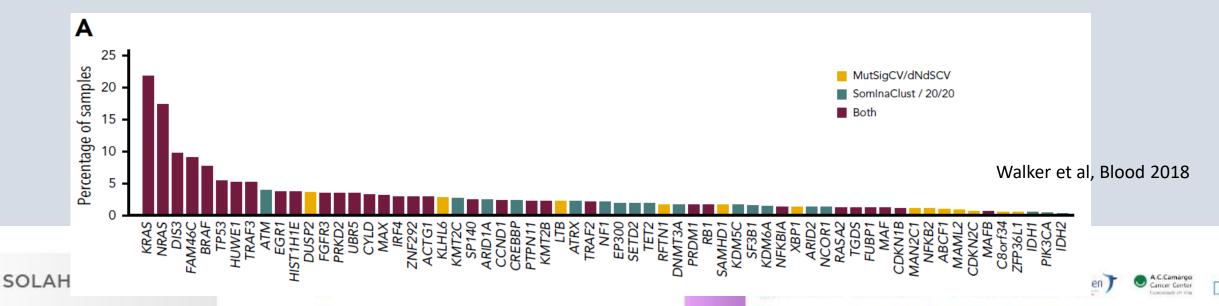
Walker et al, JCO 2017

WES in 463 patients with newly diagnosed MM showed 15 significantly mutated genes

- RAS pathway (43%)
- NFkB pathway (17%)

- Prognostically neutral
- G0/G1 cell cycle transition and epigenetic regulators

Mutations in CCND1, TP53, ATM associated with poor, IRF4 and EGR1 good prognosis



GENETIC EVOLUTION OF MULTIPLE MYELOMA

Initial Alterations establish the malignant clone (non-overlapping)

Rajan & Rajkumar, Boood Cancer J 2015

janssen

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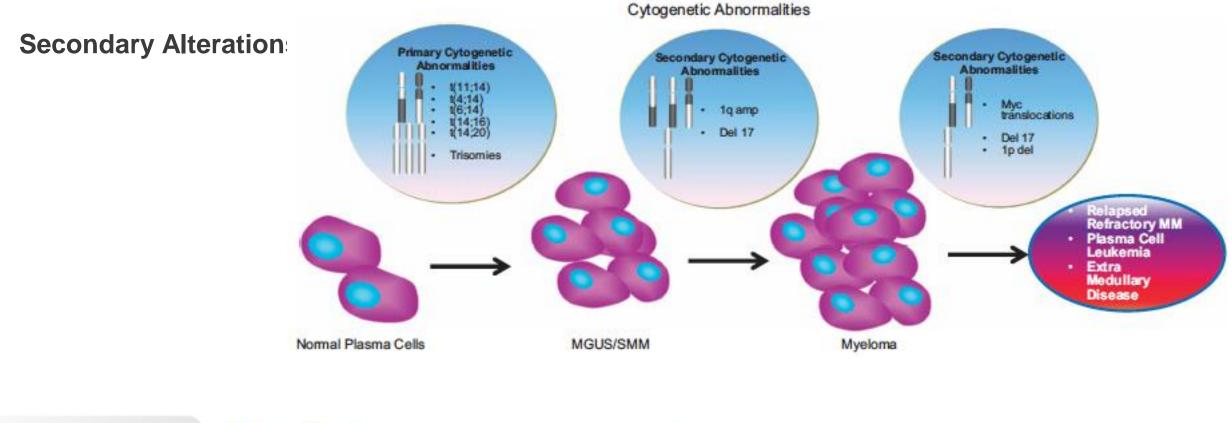
HEMATOPATOLOGÍA

SÃO PAULO | 2023

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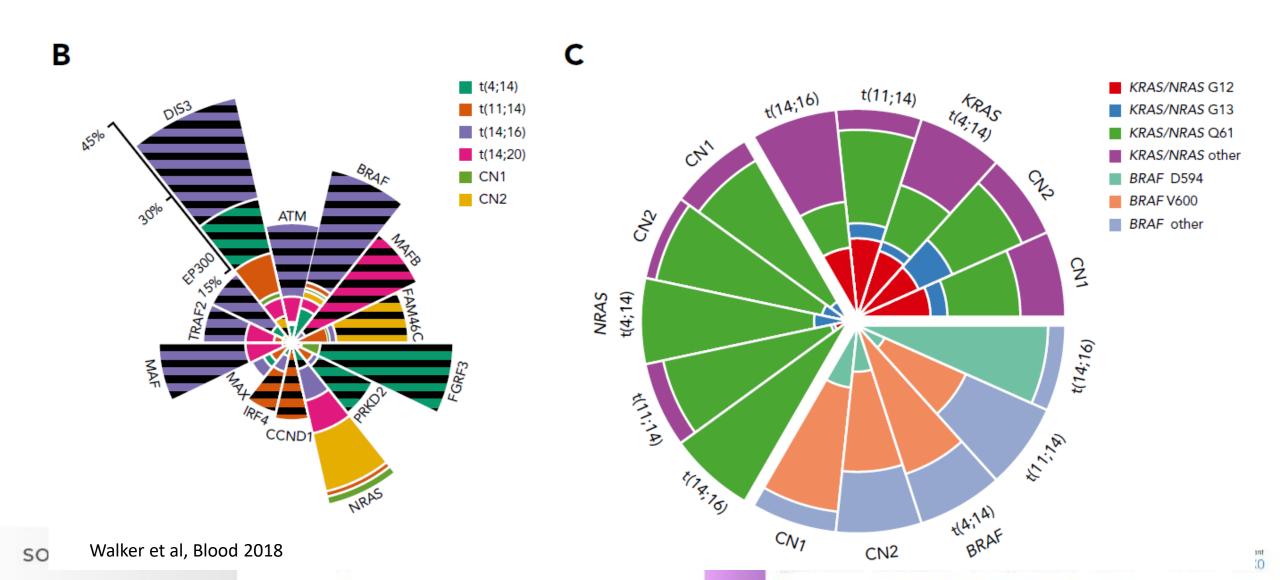
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STRONG ASSOCIATIONS OF PRIMARY AND SECONDARY GENETIC ALTERATIONS INDICATE ONCOGENIC DEPENDENCIES

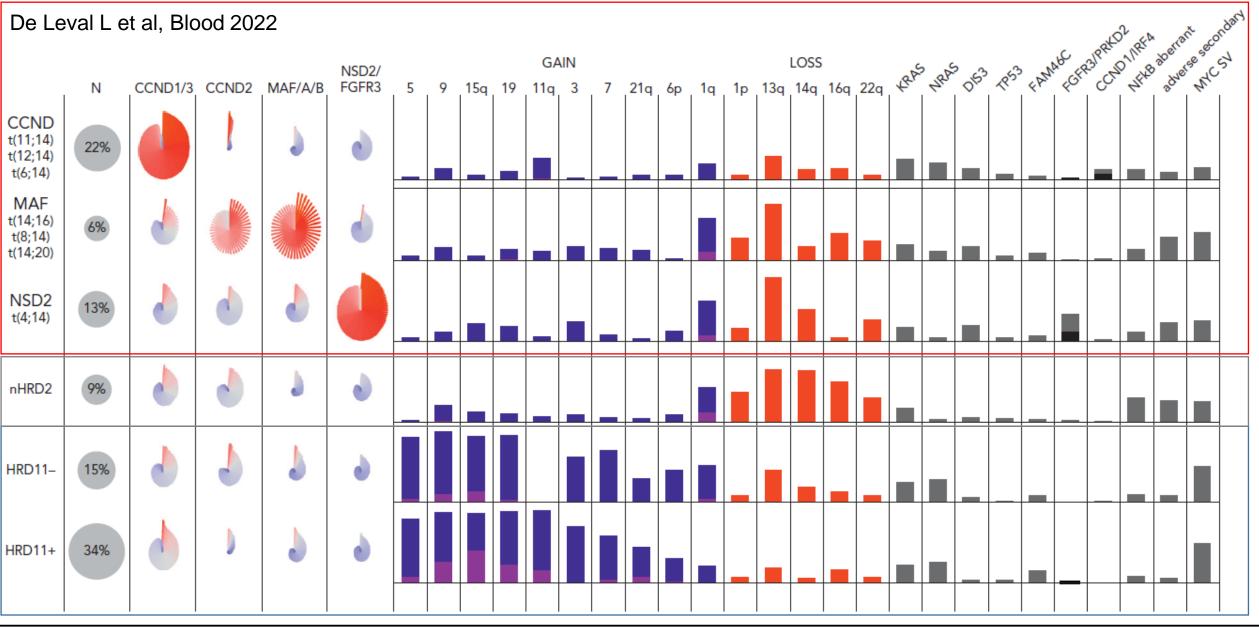




MOLECULAR CLASSIFICATION OF MM



De Leval L et al, Blood 2022



WORK-UP OF MM GENETICS



FISH (or NGS-) based detection of recurrent translocations required for genetic subtyping according to ICC and relevant for risk stratification

Prognostic impact of cytogenetics modified by therapy regimen

Entity	Genetic alteration: test	Diagnostic use	Clinical impact	Future assays
Multiple myeloma (MM) MM-NOS MM with recurrent genetic abnormality MM with CCND family translocation MM with MAF family translocation MM with <i>NSD2</i> translocation MM with hyperdiploidy	t(4;14) NSD2::IGH; t(14;16) IGH::MAF; t(11;14) CCND1::IGH;*,§ gain of odd numbered chromosomes: FISH on bone marrow plasma cells (CD138-positive selected sample strongly recommended)*	Diagnostic of the ICC subtypes of MM	t(11;14) predictive of response to venetoclax ¹³⁴	WGS for subtype assignment, risk stratification, and decision making MRD using HTS for decision making
	t(4;14) NSD2::IGH; t(14;16) IGH::MAF; amp(1q); del(1p), del(17p)*; TP53 mutations ⁴⁶⁴ For SMM: t(4;14) NSD2::IGH; t(14;16) IGH::MAF; 1q gain/ amplification; del(13) ¹⁴⁵ and MYC rearrangement ¹³⁹ : FISH and HTS	Risk stratification at diagnosis and relapse	The adverse prognosis of high-risk genetics is partially overcome by the addition of a proteasome inhibitor ¹³¹ and/or anti- CD38 MoAb ¹³² to first- line therapy	
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CHANGES IN CLASSIFICATION – ICC 2022

IgM MGUS separated into

- Plasma cell type with *MYD88* WT and MM-type cytogenetics (e.g. t(11;14)), precursor to IgM MM
- IgM MGUS NOS

Name change back to multiple myeloma Formal separation into cytogenetic groups Emphasis on the presence of minimal BM infiltration in SPB and EMP Name change to IG light chain amyloidosis

Introduction of localized AL amyloidosis as new category



Immunoglobulin M (IgM) monoclonal gammopathy of undetermined significance (MGUS)

☆ IgM MGUS, plasma cell type* IgM MGUS, not otherwise specified (NOS)*

Plasma cell neoplasms

- Non-IgM MGUS
- 🗡 Multiple myeloma (plasma cell myeloma)*

Multiple myeloma, NOS

Multiple myeloma with recurrent genetic abnormality Multiple myeloma with *CCND* family translocation Multiple myeloma with *MAF* family translocation Multiple myeloma with *NSD2* translocation Multiple myeloma with hyperdiploidy Solitary plasmacytoma of bone Extraosseous plasmacytoma Monoclonal Ig deposition diseases Monoclonal Ig deposition diseases ↓ Ig light chain amyloidosis (AL)* ↓ Localized AL amyloidosis*



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WHO CLASSIFICATION 5TH EDITION (PRE-VERSION)



	Plasma cell neoplasms and other diseases with paraproteins	
	Monoclonal gammopathies	
\star	Cold agglutinin disease	Not previously included
\star	IgM monoclonal gammopathy of undetermined significance	(Same)
	Non-IgM monoclonal gammopathy of undetermined significance	(Same)
\star	Monoclonal gammopathy of renal significance	Not previously included
	Diseases with monoclonal immunoglobulin deposition	
	Immunoglobulin-related (AL) amyloidosis	Primary amyloidosis
	Monoclonal immunoglobulin deposition disease	Light chain and heavy chain deposition disease
	Heavy chain diseases	
	Mu heavy chain disease	(Same)
	Gamma heavy chain disease	(Same)
	Alpha heavy chain disease	(Same)
	Plasma cell neoplasms	
	Plasmacytoma	(Same)
\star	Plasma cell myeloma	(Same)
*	Plasma cell neoplasms with associated paraneoplastic syndrome -POEMS syndrome	(Same) Except AESOP syndrome not previously included
SOLAHP	-TEMPI syndrome -AESOP syndrome	
	Algori Synaronic	



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ICC 2022

Minor adaptations Based on IMWG criteria Increased importance of genetics Relevance of minimal BM infiltration in solitary plasmacytomas

WHO 5th Edition

Minor adaptations Re-organisation of categories based on type of paraprotein and IG deposition Introduction of clinical syndromes

RARE DISEASES ASTRAZENECA

caused by paraprotein

Overall, differences seem minor between the two classifications





NOVEL BIOMARKERS FOR MM

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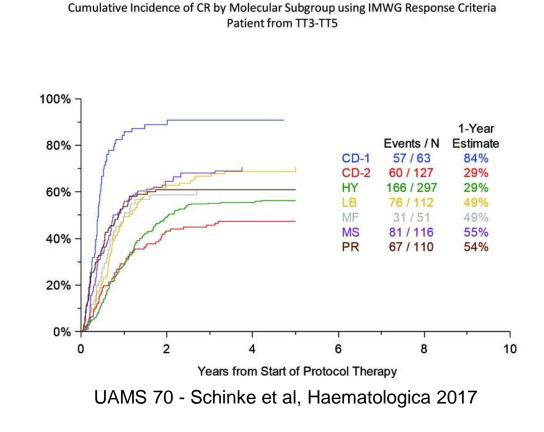
Gene expression analysis

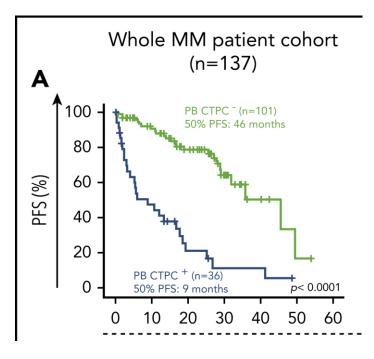
UAMS 70 signature

EMC 92 signature Mutational analysis Epigenetics ctDNA

Circulating tumor cells

Exploration of role of **microenvironment**





circulating PC with FCM post therapy Sanoja-Flores L et al, Blood 2020

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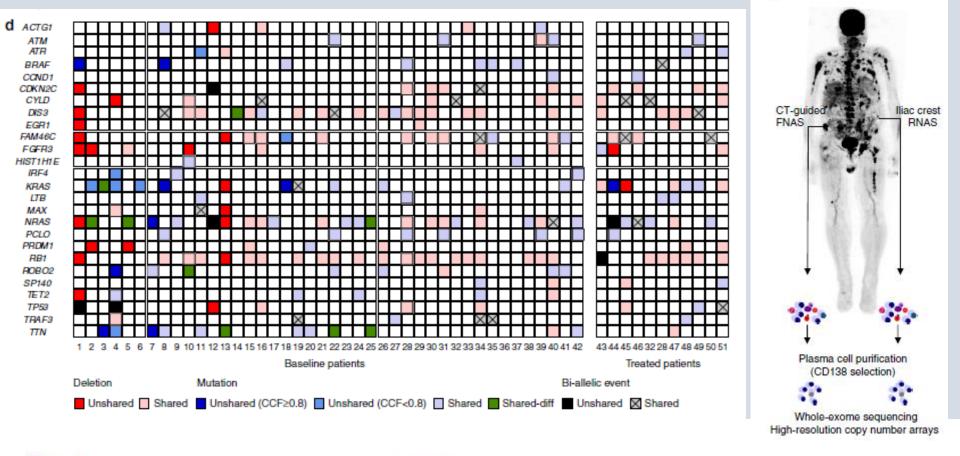


Spatial genomic heterogeneity in multiple myeloma revealed by multi-region sequencing

L. Rasche¹, S.S. Chavan¹, O.W. Stephens¹, P.H. Patel¹, R. Tytarenko¹, C. Ashby¹, M. Bauer ¹, C. Stein¹,
S. Deshpande¹, C. Wardell¹, T. Buzder¹, G. Molnar¹, M. Zangari¹, F. van Rhee¹, S. Thanendrarajan¹, C. Schinke¹,
J. Epstein¹, F.E. Davies¹, B.A. Walker ¹, T. Meissner², B. Barlogie¹, G.J. Morgan¹ & N. Weinhold¹

75% of patients show clonal heterogeneity, especially with large lesions (>2.5 cm)

Initial (founder) alterations detectable in all tumor cells



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MM AND THE MICROENVIRONMENT



Dako

Complex interaction with BM microenvironment through adhesion molecules, cytokines/chemokines

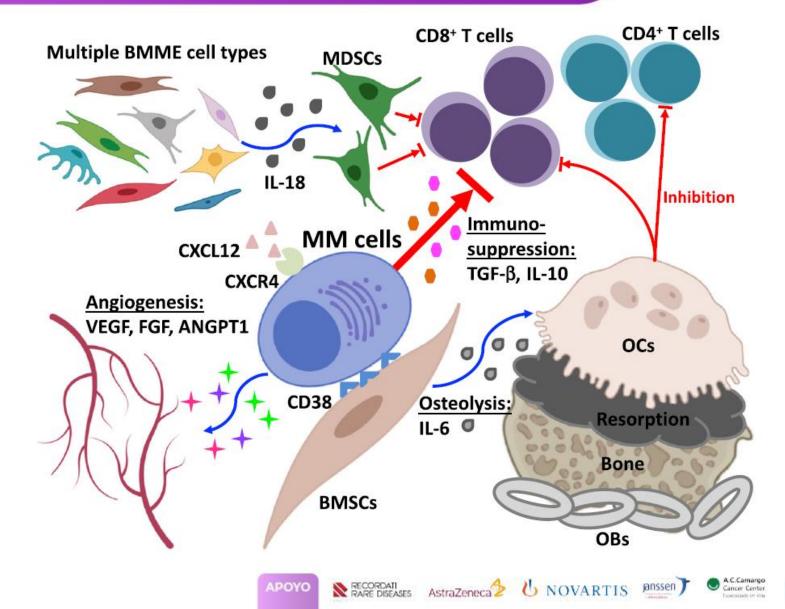
Induction of angiogenesis, Immunosuppression/evasion and bone disease

With progression increasing independence from BM ME development of extramedullary disease and PC leukemia

> Sociedade Brasileira de PATOLOCIA

Therapeutic targets

⁴¹SOLAHP23.COM.BR







- Diagnosis of plasma cell neoplasms requires integration of clinical, laboratory and radiological findings
- Precise quantification of plasma cells is essential
- Unusual morphological and phenotypical features provide pitfalls, which can be avoided by a limited number of immunostains
- Cytogenetic profile is an integral part of the current classification of plasma cell disorders





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SÃO PAULO | 2023





European Association for Haematopathology

REALIZACIÓN

APOYO





UNOVARTIS





