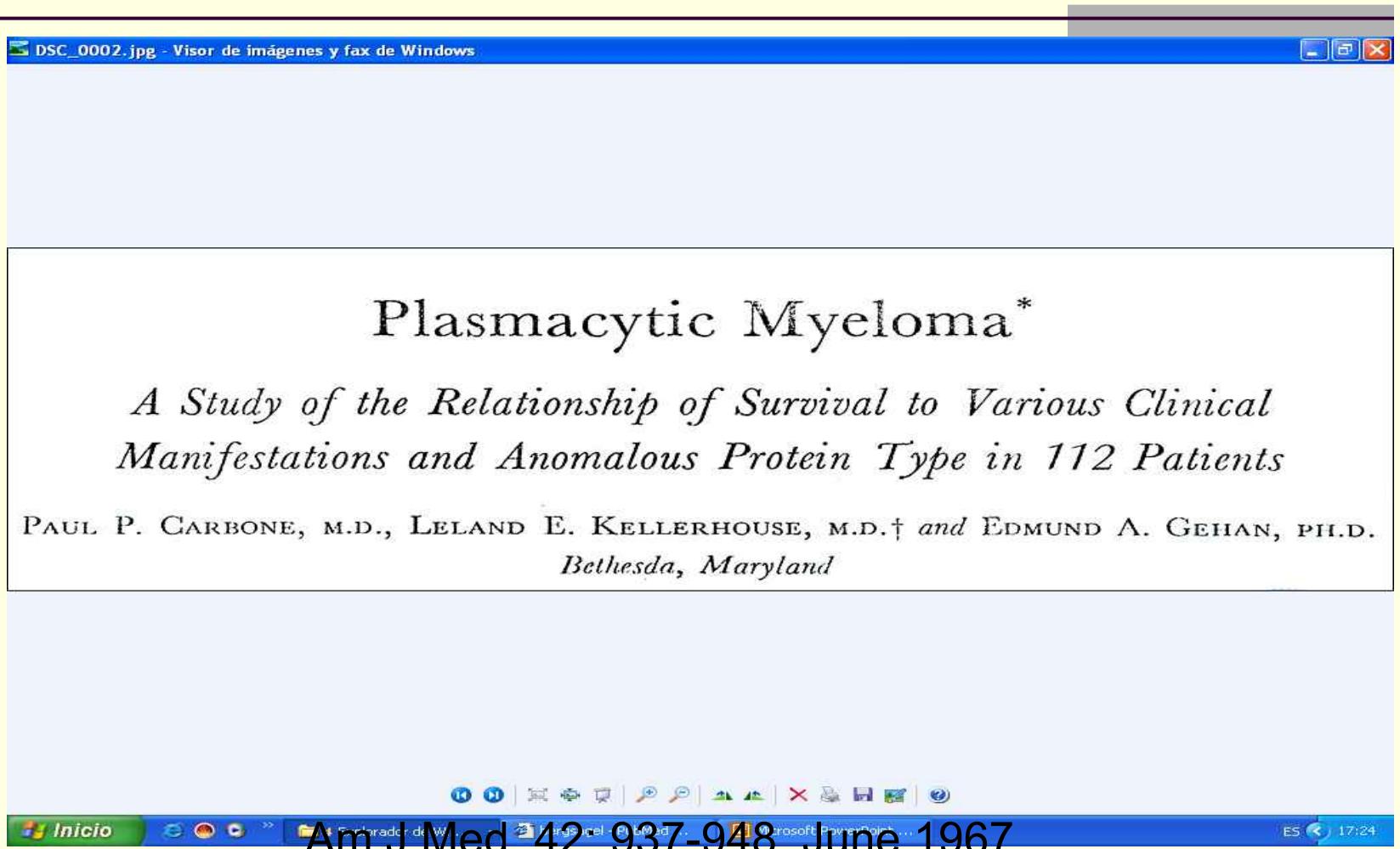


# **Which are the relevant prognostic factors in the current management of multiple myeloma with novel agents?**

**Joan Bladé**

**Athens Conference on Plasma Cell Dyscrasias**

**Athens, September 2009**



## Plasmacytic Myeloma\*

## *A Study of the Relationship of Survival to Various Clinical Manifestations and Anomalous Protein Type in 112 Patients*

PAUL P. CARBONE, M.D., LELAND E. KELLERHOUSE, M.D.<sup>†</sup> and EDMUND A. GEHAN, PH.D.  
*Bethesda, Maryland*

Am J Med, 42: 937-948, June 1967

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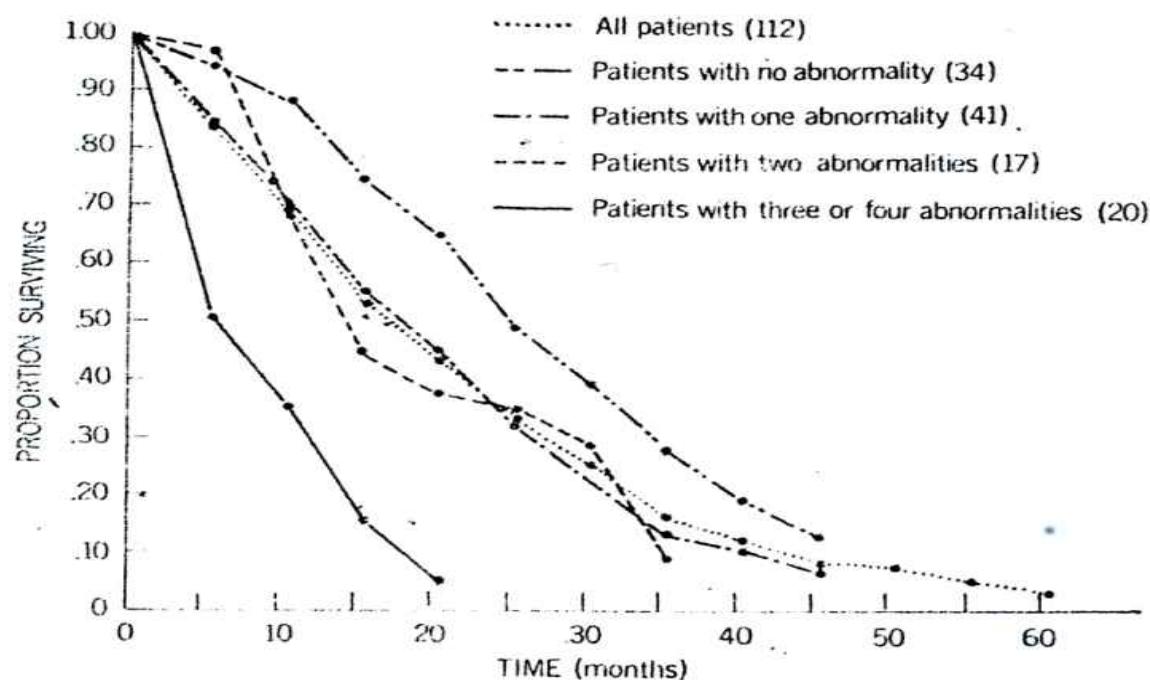


FIG. 2. Survival curves for patients with plasmacytic myeloma.

# Prognostic Factors in MM

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- Prognostic value...
- A new prognostic factor...
- A simple reliable marker...
- An easily available parameter...
- An independent prognostic factor...
- A new staging system...
- Proposal for a novel prognostic index...

# Prognostic Factors in MM

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- Clinical and laboratory features
- Staging systems
- Malignant clone: molecular genetic status
- Response to therapy
- Mechanisms of disease control/progression

# Prognostic Factors in MM Clinical and Laboratory Features

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## ■ Host characteristics

- Age
- PS

## ■ Tumor burden

- $\beta_2$ -microglobulin

## ■ Organ damage

- Renal function
- Hb

# Main Staging Systems in MM

Author, year	Parameters	Other
Durie and Salmon, 1975	Hb, Ca, M-protein, bone lesions	Renal function
Merlini et al, 1980	%PC, Cr, and Ca (IgG) Hb, Ca, M-protein (IgA)	
MRC, 1980	Hb, urea, PS	
Cavo et al, 1989	D & S, platelet count	
Greipp et al, 1988	$\beta$ 2-microglobulin, LI	
Bladé et al, 1989	Albumin, urea	
San Miguel et al, 1989	Hb, Cr, PS, PI	
San Miguel et al, 1995	S-phase, $\beta$ 2-microglobulin, age, PS	
<b>IMWG, 2005</b>	<b><math>\beta</math>2-microglobulin, albumin</b>	

MRC: Medical Research Council; IMWG: International Myeloma Working Group; Hb: haemoglobin; Ca: calcium; PC: plasma cells; Cr: creatinine; Ig: Immunoglobulin; PS: performance status; LI: labelling index; PI: paraprotein index.

# International Prognostic System (IPS)

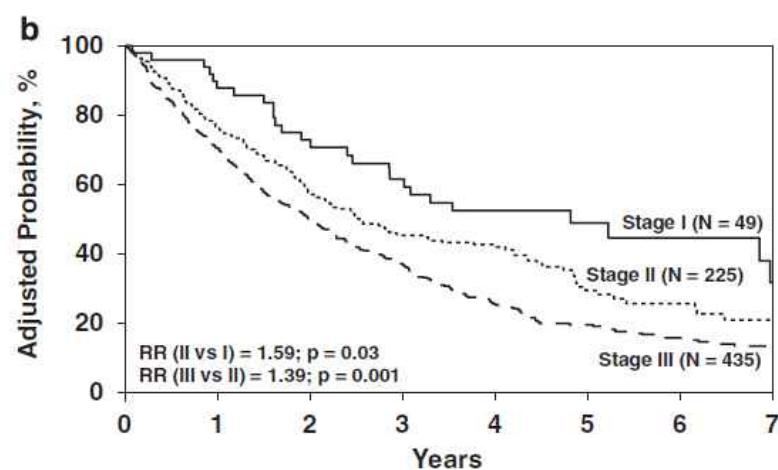
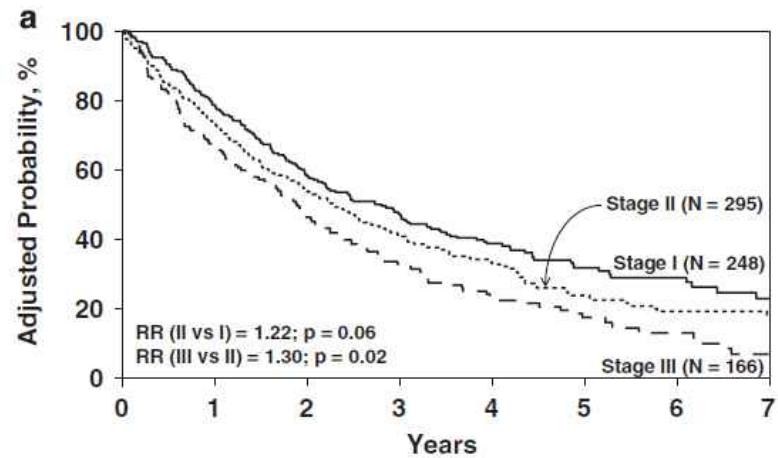
Stage		Overall Survival (months)
I	$\beta\text{-}2\text{M} < 3.5 \text{ mg/L}$ and albumin $\geq 3.5 \text{ g/dL}$	62
II	$\beta\text{-}2\text{M} < 3.5 \text{ mg/L}$ and albumin $< 3.5 \text{ g/dL}$ or $\beta\text{-}2\text{-m } 3.5 - 5.4 \text{ mg/L}$	44
III	$\beta\text{-}2\text{M} \geq 5.5 \text{ mg/L}$	29

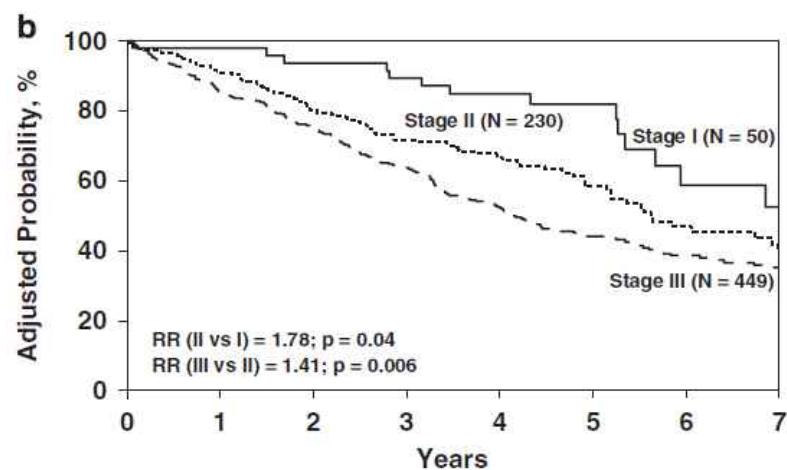
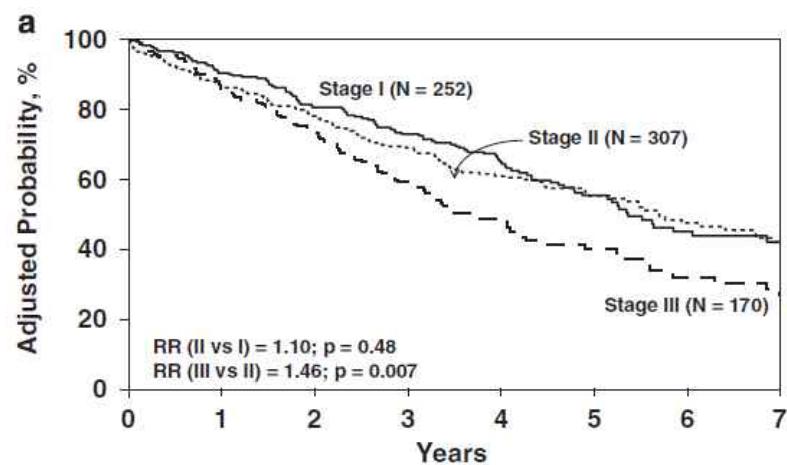
Greipp P et al. J Clin Oncol 2005; 23: 3412-20.

# Is the ISS Superior to the Durie & Salmon SS?

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- CIBMTR results in 729 patients who underwent up-front ASCT





# ISS versus Durie & Salmon SS

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- Durie & Salmon superior when adjusting with Brier Score
- Neither staging system strongly predictive of outcomes



Need for incorporation other prognostic markers (cytogenetics, GEP, imaging-MRI/PET)

# Cytogenetic Prognostic Subgroups in Multiple Myeloma

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- Good/average prognosis
  - Hyperdiploidy
  - t(11;14)(q32;q32): cyclin D1 upregulation
- Bad prognosis
  - Hypodiploidy
  - t(4;14)(p16.3;q32): FGFR3&MMSET upregulation
  - t(14;16)(q32;q23): c-MAF upregulation
  - Chromosome 1 abnormalities: 1q gains  
(overexpression CKS1B)
  - 17q deletions, 13q deletions

# 13q Deletion as Single Abnormality

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- No independent prognostic impact\*

\* Gutiérrez N et al. Leukemia 2007; 21: 541-9.

\* Avet-Loiseau H et al. Blood 2007; 109: 3489-95.

# Molecular Myeloma Subgroups Gene Expression Profiling

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- “Translocation/Cyclin D” classification\*:
  - 8 groups
  
- Recurrent translocations/hyperdiploidy\*\*:
  - 7 entities

\* Bergsagel PL et al. Blood 2005; 106: 296-303.

\*\* Zhan F et al. Blood 2006; 108: 2020-8.

# High-resolution Genomic Profiles\*

(aCGH/mRNA microarray/FISH/novel bioinformatics)

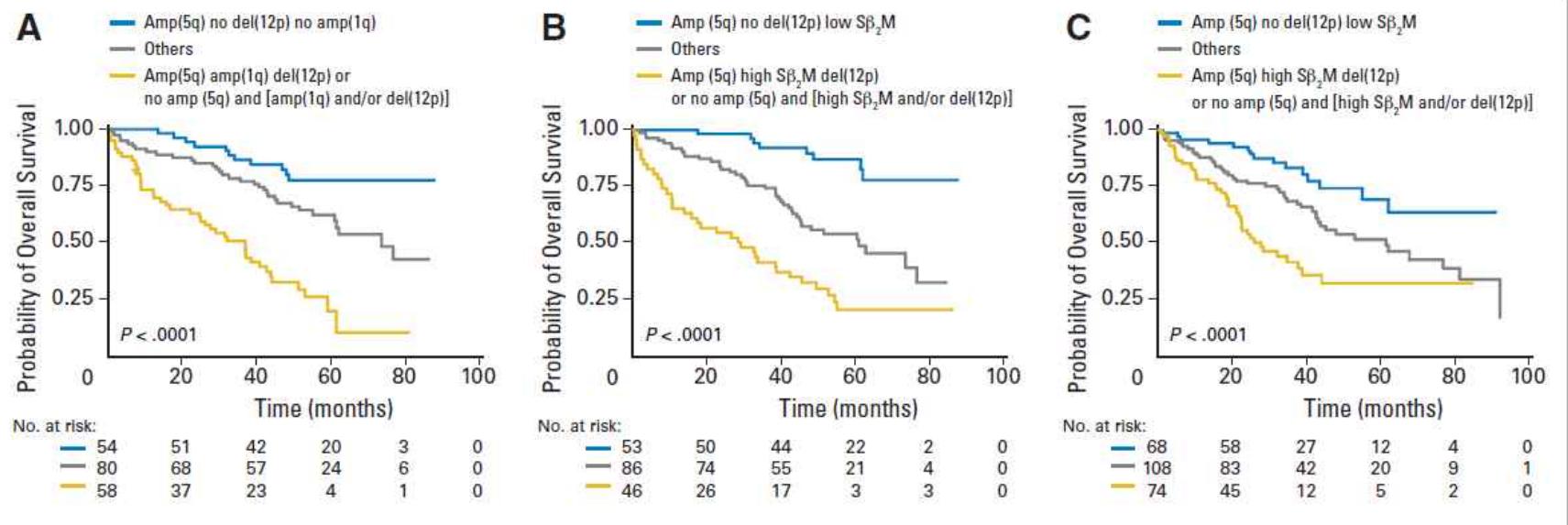
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- 4 different MM subtypes  
(recurrent DNA copy number changes), i.e.:
  - Hyperdiploid, 11q gains: good outcome
  - Hyperdiploid, 1q gains and/or 13 losses: poor outcome

\* Carrasco R et al. Cancer Cell 2006; 4: 313-25.

# High-resolution DNA copy number changes (SNP-based mapping array technology)

	HR	p
Del (12p13.31)	3.17	<0.0001
s $\beta$ 2M $\geq$ 5 mg/L	2.78	<0.0001
Amp (5q31.3)	0.37	0.0005



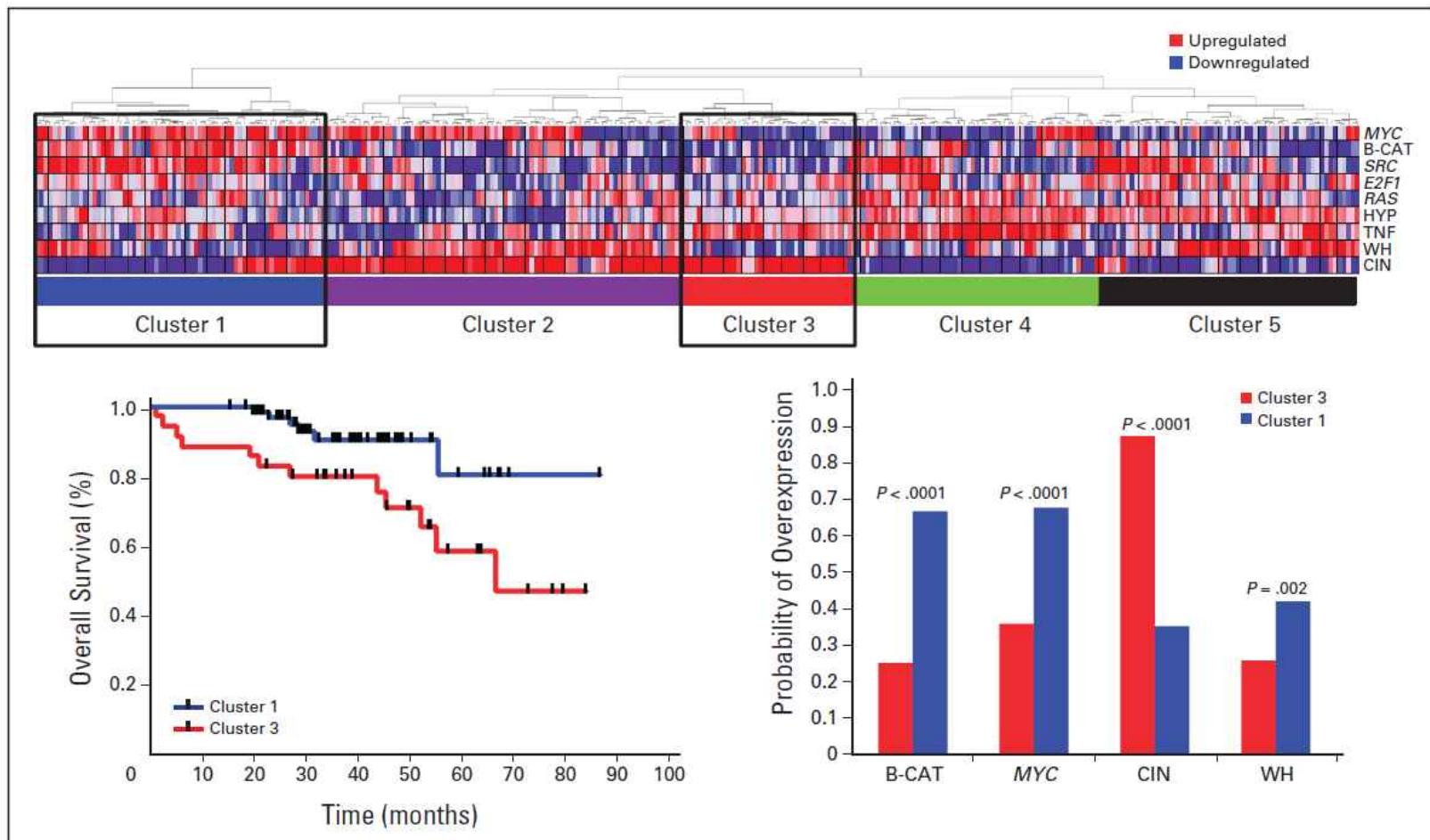
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## Gene Expression Profiles of Tumor Biology Provide a Novel Approach to Prognosis and May Guide the Selection of Therapeutic Targets in Multiple Myeloma

*Ariel Anguiano, Sascha A. Tuchman, Chaitanya Acharya, Kelly Salter, Cristina Gasparetto, Fenghuang Zhan, Madhav Dhodapkar, Joseph Nevins, Bart Barlogie, John D. Shaughnessy Jr, and Anil Potti*

- GEP of tumor biology / chemotherapy sensitivity can refine the ISS classification



# Response to Therapy as Prognostic Factor

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- Stabilization of disease
- Impact of CR
  - With primary therapy
  - After HDT/SCT

# Imaging Techniques with Prognostic Interest

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- MRI: number of focal lesions (FL)
  
- FDG-PET/CT:
  - FDG suppression (SUV-FL) prior ASCT
  - Metastatic spread (EMD)

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Walker R, et al. J Clin Oncol 2007; 25:1121-1128

Bartel TB, et al. Blood 2009 (prepublished online, May 14)

# Novel Drugs and New Molecular Targets

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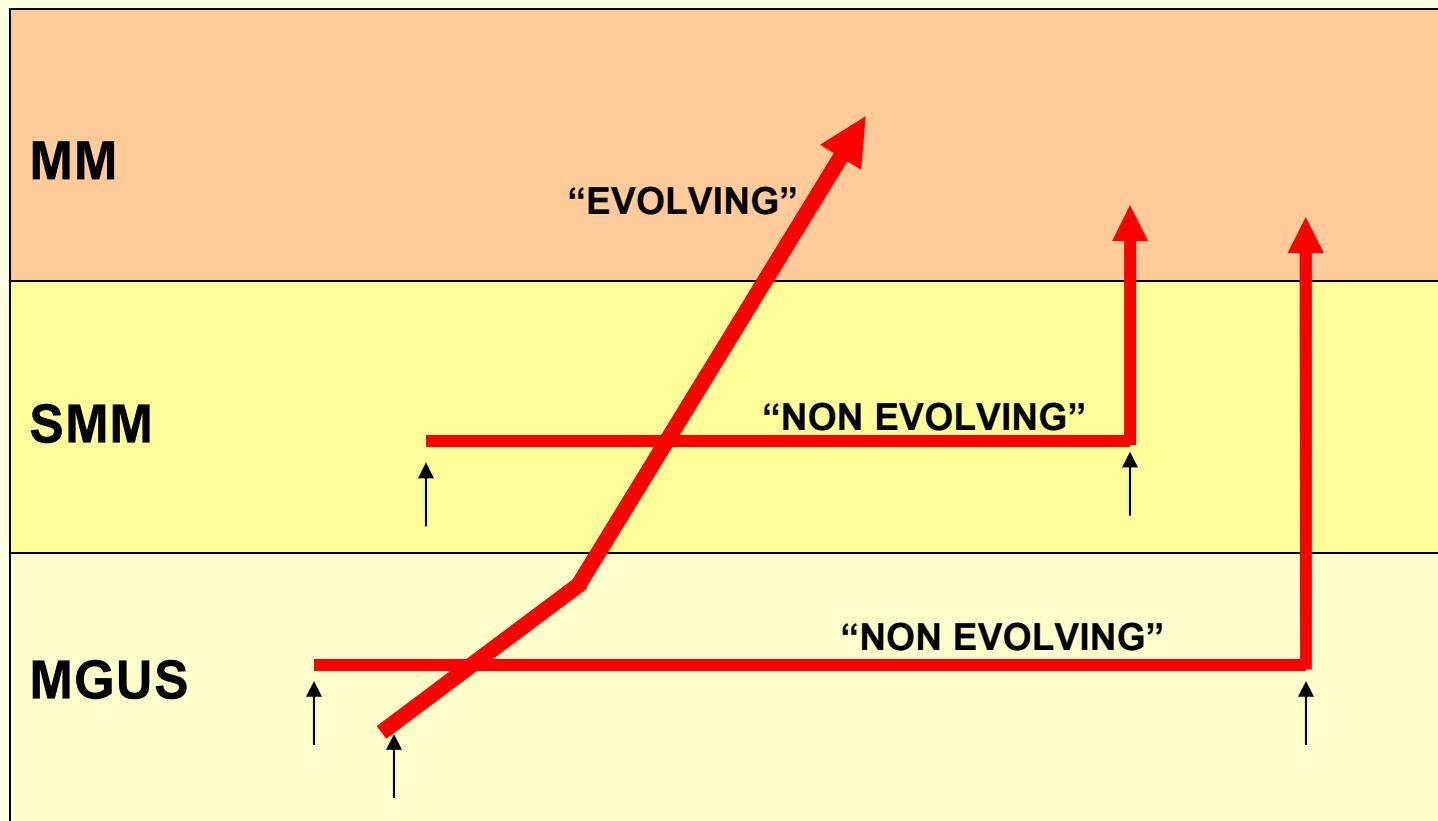
- Novel drugs can overcome drug resistance in poor cytogenetic subgroups
- New therapies should target specific molecular pathways

# Therapy Against Cancer “Myeloma” Stem Cell

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- Plasma cell killing
  - lowest possible tumor mass
- Meaning of CR achieved with novel therapies
- Different effect of old and new drugs on the **bulk of differentiated plasma cells** versus the **myeloma stem cell?**

# Monoclonal Gammopathies



Rosiñol et al. Br J Haematol 2003; 123: 631-36.

Rosiñol et al. Mayo Clin Proc 2007; 82: 428-34.

# Possible Impact of Influencing on Mechanisms of Disease Progression

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- Evolving MGUS  $\Rightarrow$  early/slowly evolving myeloma (escaping growth-restraining mechanisms)
- Avoid disease progression after decreasing tumor mass  $\Rightarrow$  MGUS state (growth-restraining influences)

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# **When Is a “Prognostic Factor” Really Prognostic?**

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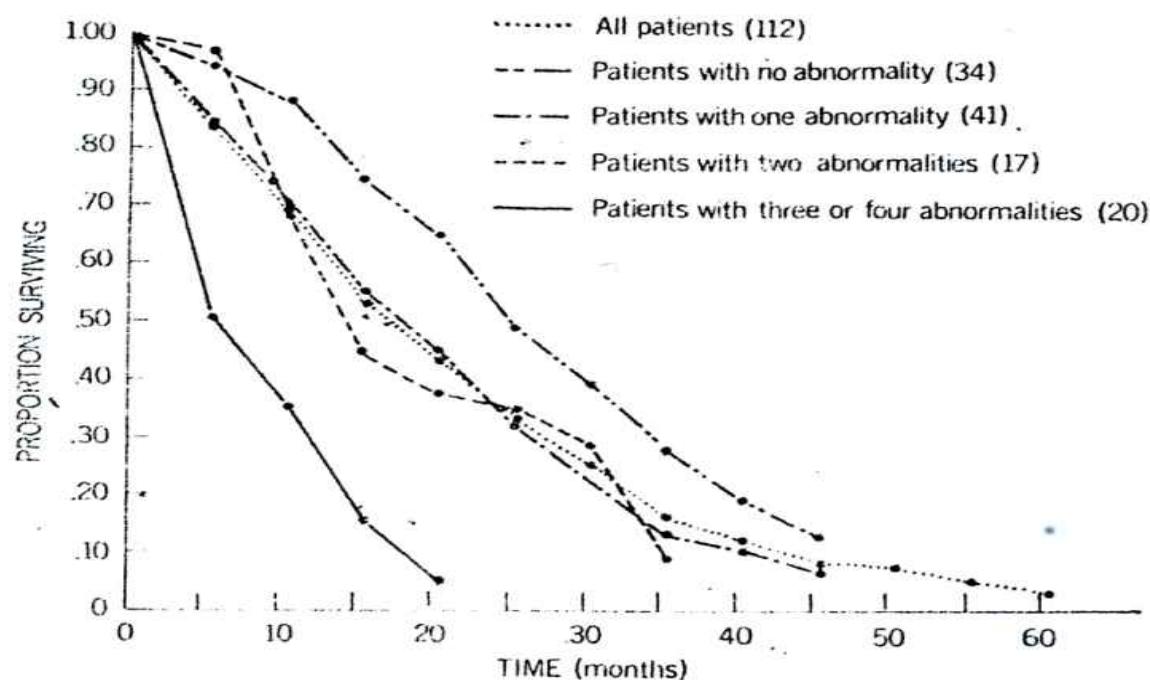
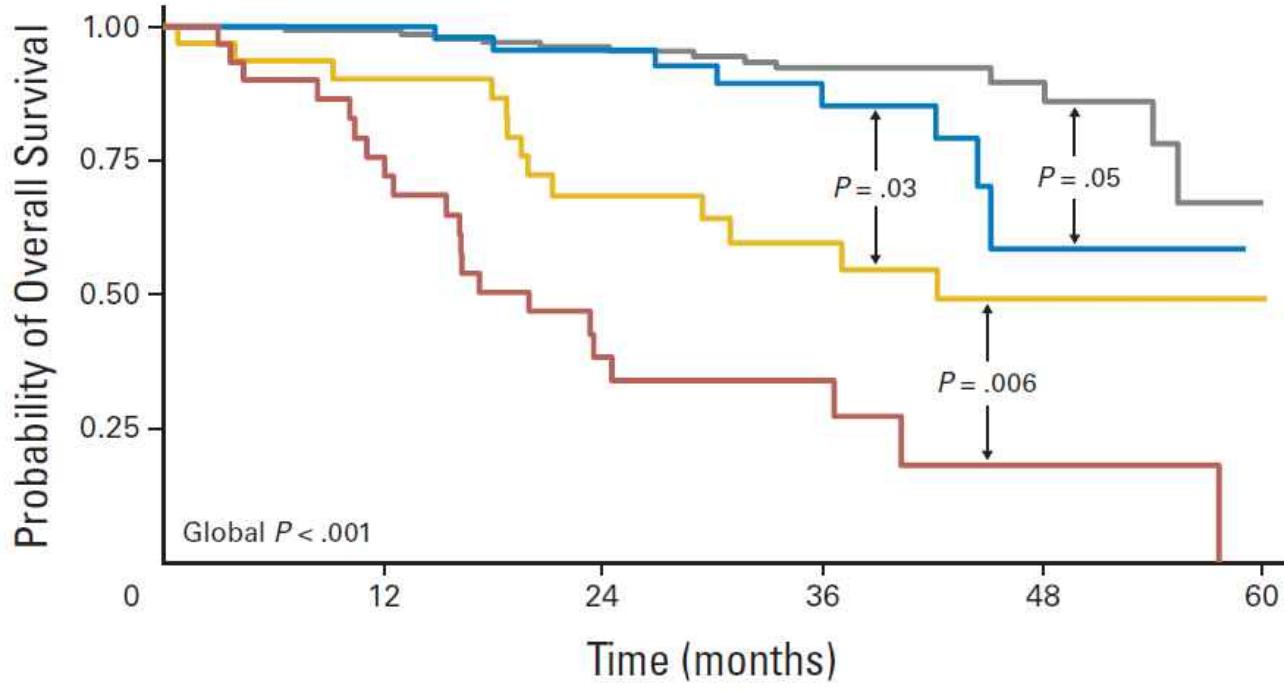


FIG. 2. Survival curves for patients with plasmacytic myeloma.



#### Patients at risk

15-gene low-risk + S $\beta$ 2M < 5.5 and not t(4;14)	140	129	116	71	26	1
15-gene low-risk + S $\beta$ 2M $\geq$ 5.5 and/or t(4;14)	48	45	36	19	2	0
15-gene high-risk + S $\beta$ 2M < 5.5 and not t(4;14)	31	27	18	12	7	1
15-gene high-risk + S $\beta$ 2M $\geq$ 5.5 and/or t(4;14)	31	21	9	5	1	0

