

Role of MRD in MM: different techniques for a crucial biomarker

Jaarbeurs – Media Plaza / Supernova
complex, Jaarbeursplein, Utrecht

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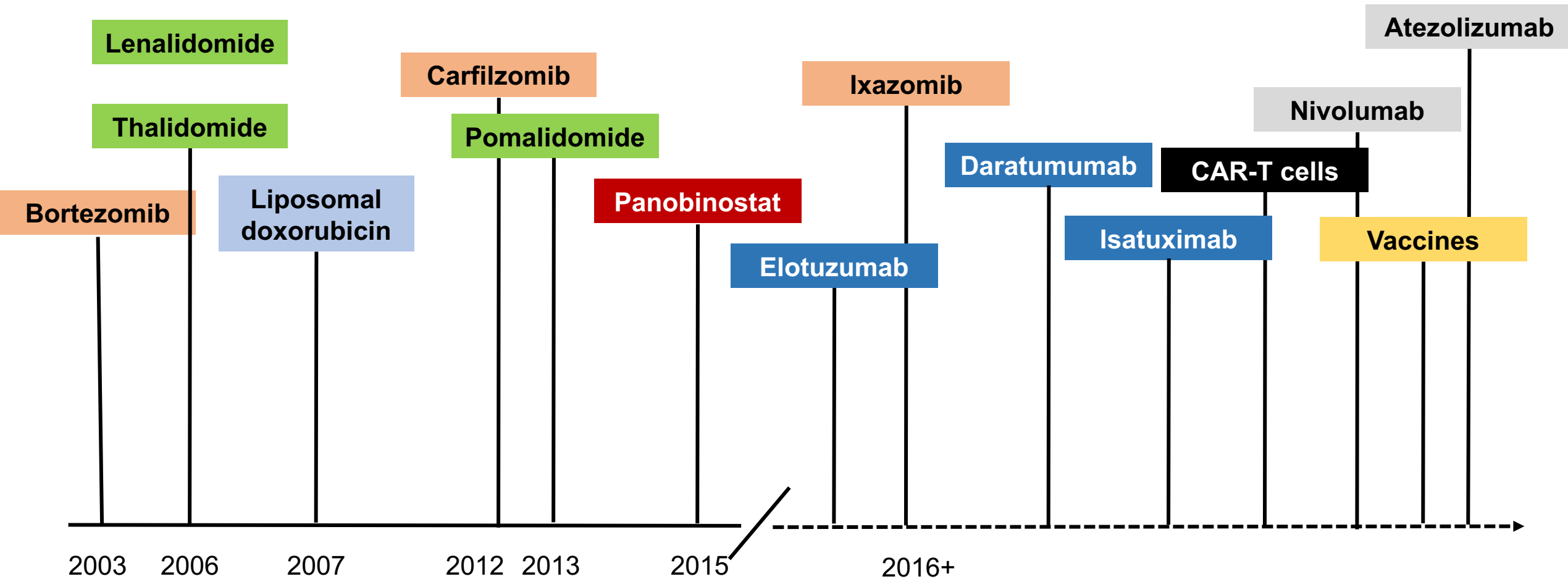
Disclosure commercial conflict of interest

	No, nothing to disclose
x	Yes, as specified below:

Company Name	Specification
Amgen	Research support, Scientific Advisory Board, Other
BMS	Research, Scientific Advisory Board, Speaker's bureau
Janssen	Research, Scientific Advisory Board, Other
Takeda	Research, Scientific Advisory Board, Other
The Binding Site	Scientific Advisory Board
Sanofi	Scientific Advisory Board
Pfizer	Research support

Novel Therapy

Novel Therapies and Immunotherapy



IMiD

HDAC inhibitor

Monoclonal antibody

Vaccines

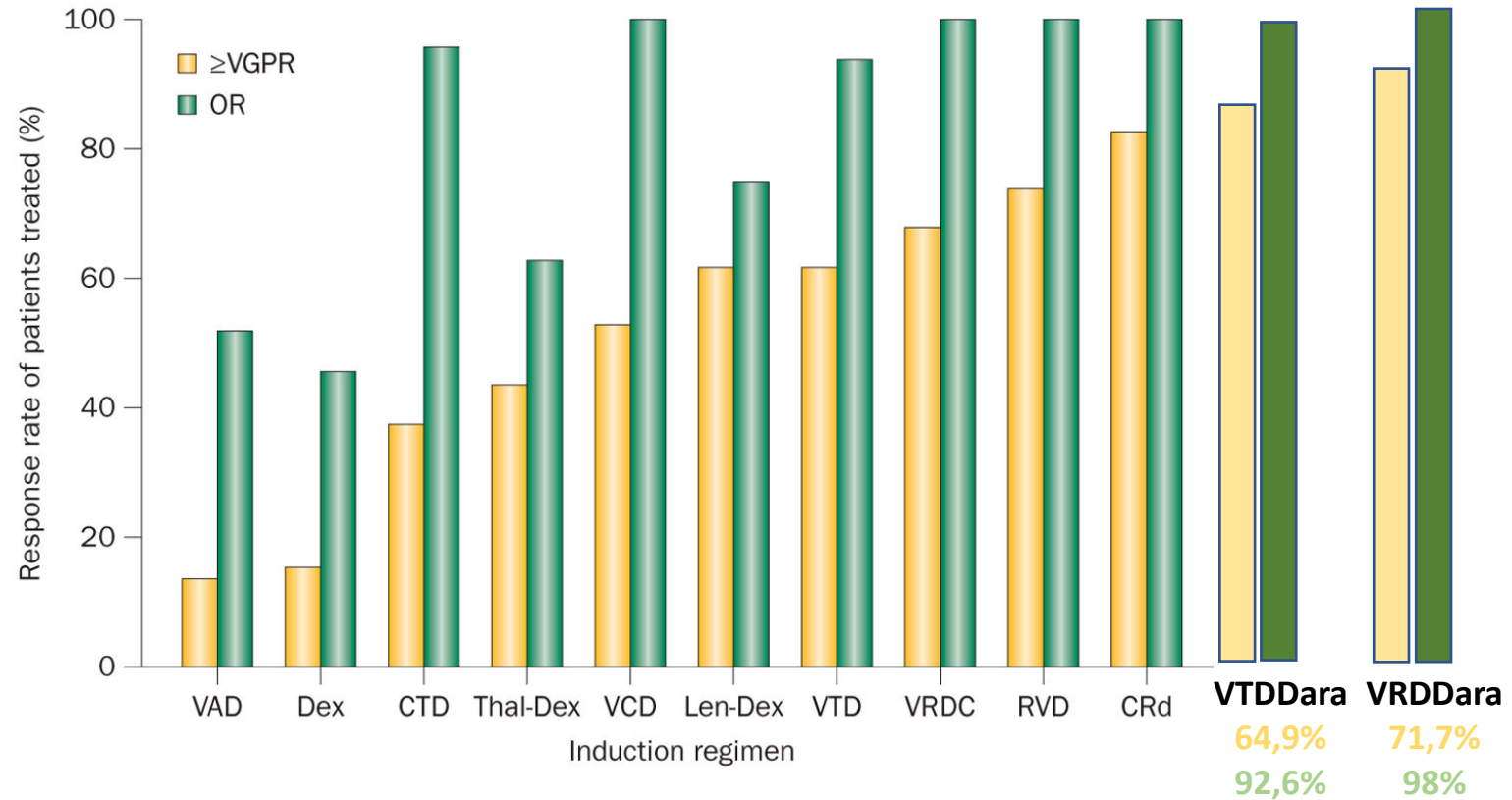
Proteasome inhibitor

Chemotherapy

Adoptive T-cell therapy

Checkpoint inhibitors

Treatment response with induction regimens in MM



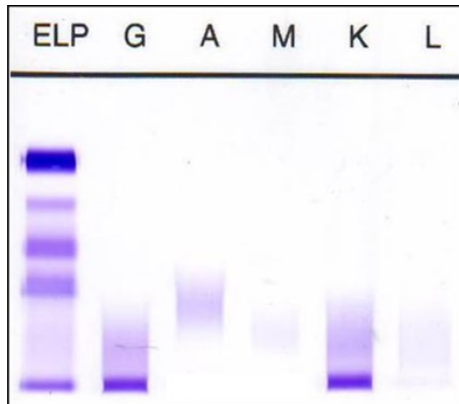
Nature Reviews | Clinical Oncology

Modified with permission from Springer Science+Business Media © Kumar, S. *Med. Oncol.* **27** (Suppl. 1), S14–S24 (2010)

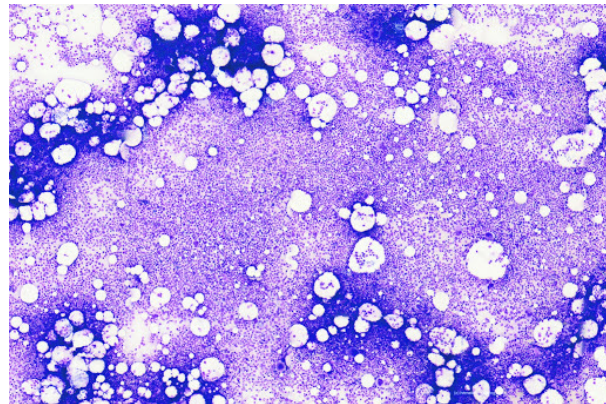
IMWG Standard Response Criteria

Standard IMWG response criteria

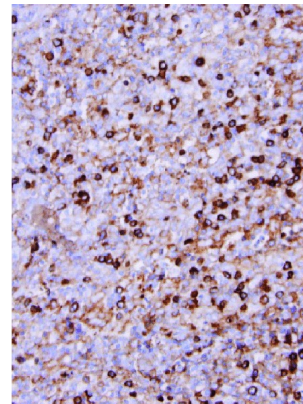
Stringent complete response	Complete response as defined below plus normal FLC ratio** and absence of clonal cells in bone marrow biopsy by immunohistochemistry (κ/λ ratio $\leq 4:1$ or $\geq 1:2$ for κ and λ patients, respectively, after counting ≥ 100 plasma cells)††
Complete response	Negative immunofixation on the serum and urine and disappearance of any soft tissue plasmacytomas and $<5\%$ plasma cells in bone marrow aspirates



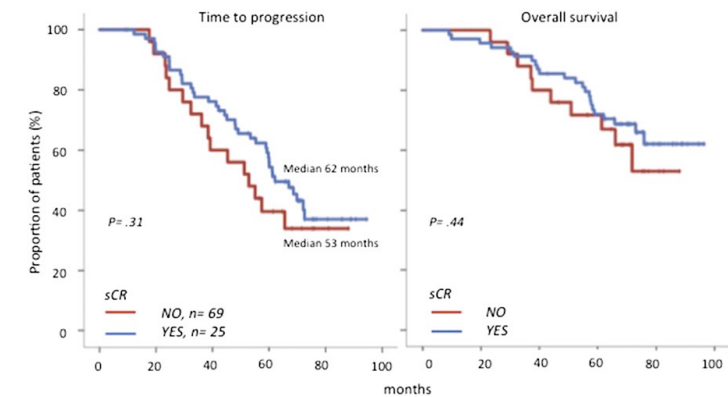
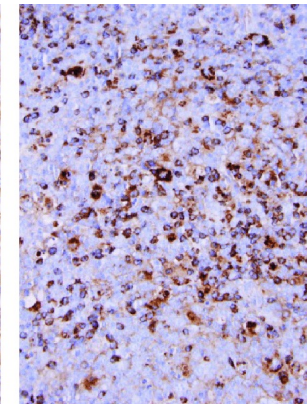
Serum Immunofixation



PC enumeration by morphology



IHC/IF: low sensitivity due to the recovery of NPCs that normalize K/L ratios



TTP and OS in patients in CR according to their status for the sCR criteria
Martínez-López J, et al; Blood 2015

Survival curves according to standard response

n = 1175

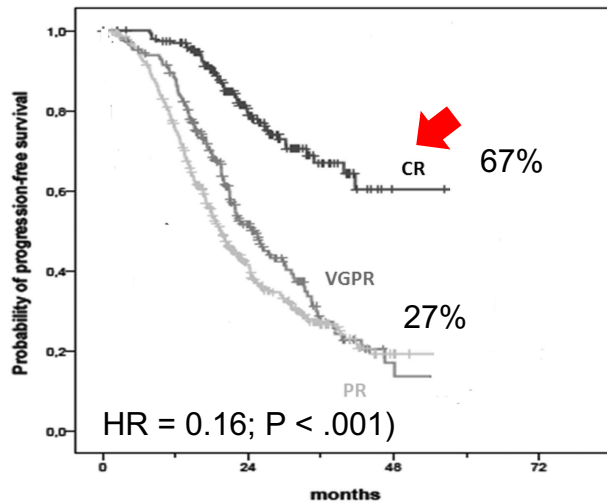
MP (n=332), MPT (n=332), MPBz (n=257), MPTBz (n=254)

Median FU 29 months

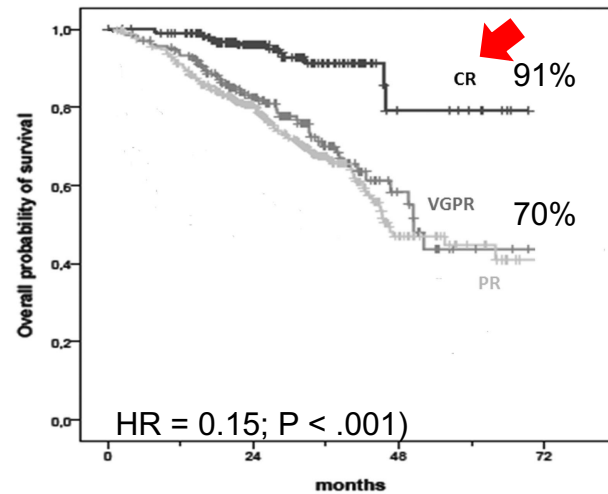
All patients

>75 yo

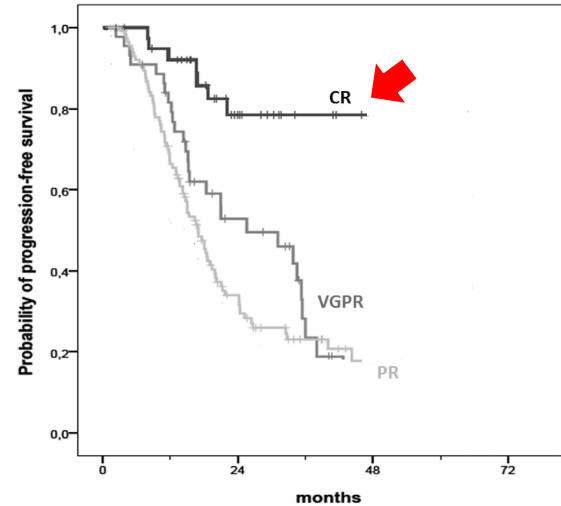
A 3-year PFS/OS



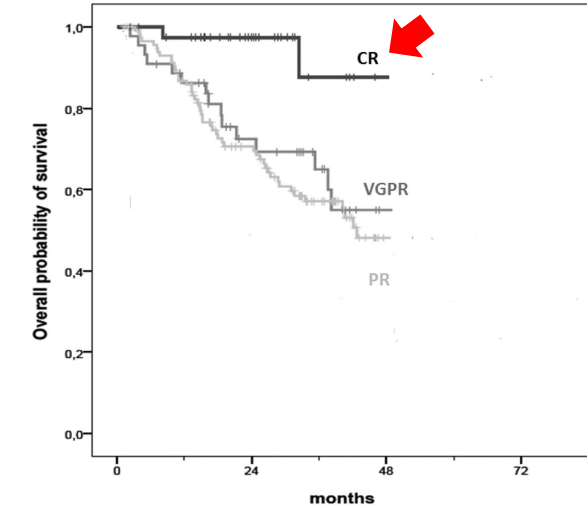
B



A

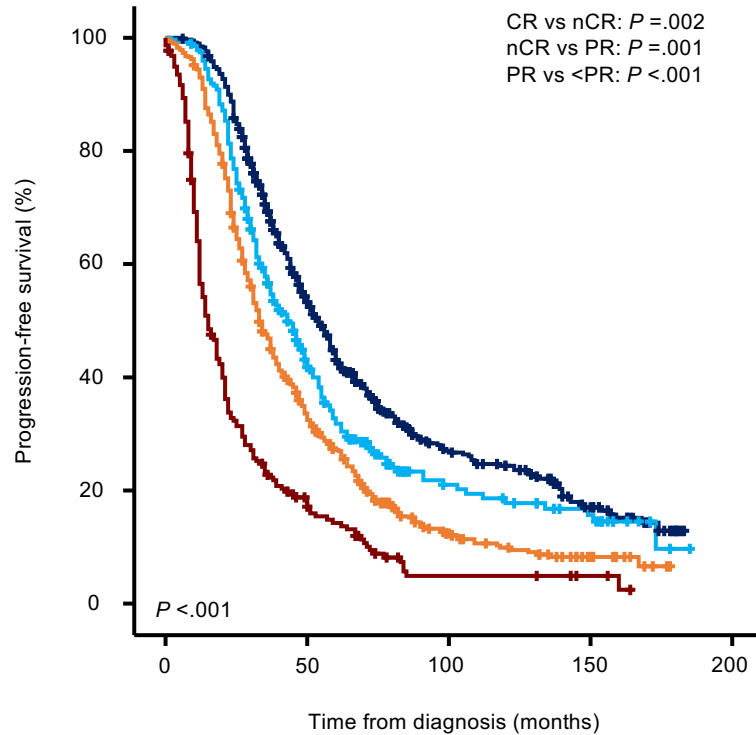


B

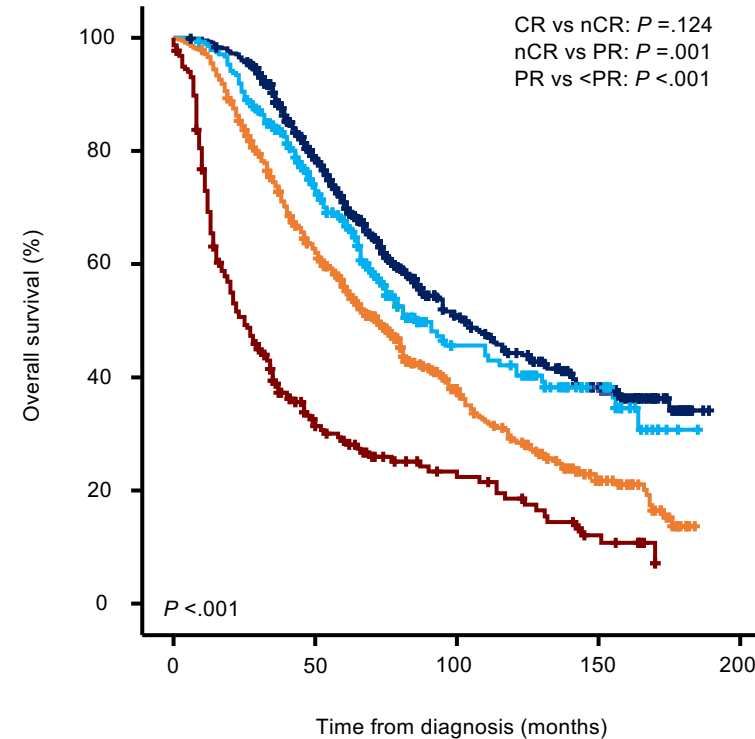


Direct relationship between the depth of response and prolonged PFS and OS

PFS and OS according to standard response



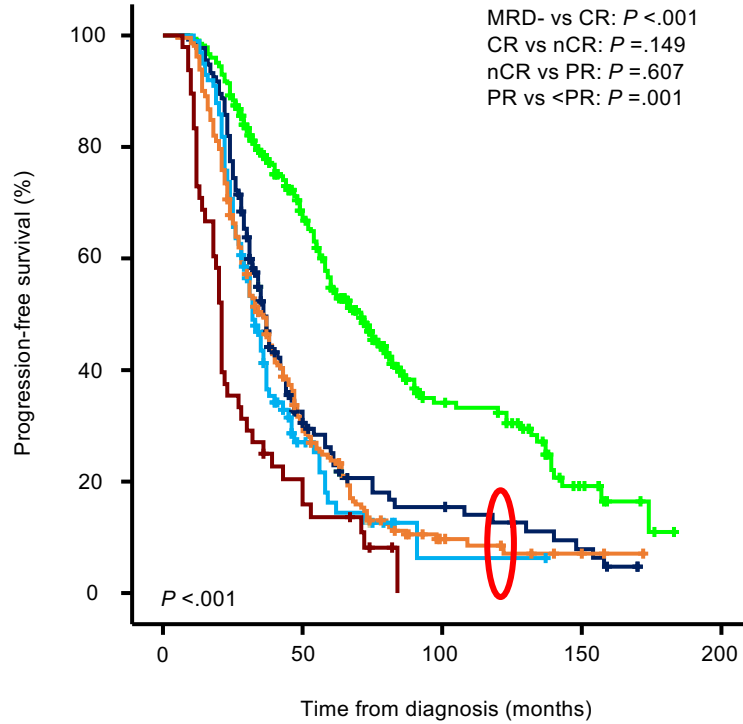
— CR (n=578) median PFS: 54 months
 — nCR (n=273) median PFS: 43 months
 — PR (n=553) median PFS: 33 months
 — <PR (217) median PFS: 15 months



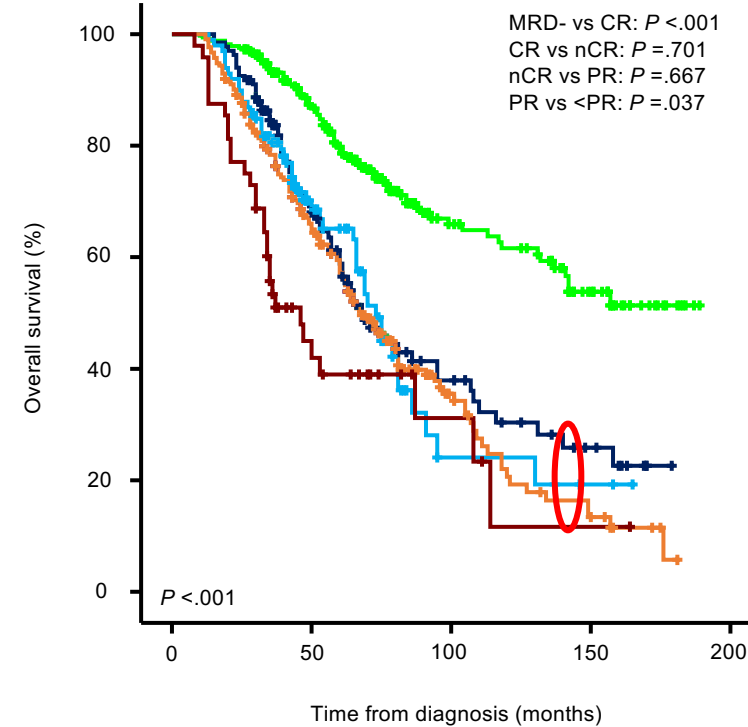
— CR (n=578) median OS: 103 months
 — nCR (n=273) median OS: 86 months
 — PR (n=553) median OS: 72 months
 — <PR (217) median OS: 25 months

Patients in CR have longer PFS and OS than those in VGPR, nCR, PR or <PR

The true value of CR depends on the MRD status



- MRD- (n=326) median PFS: 71 months
- CR (n=133) median PFS: 36 months
- nCR (n=99) median PFS: 32 months
- PR (n=211) median PFS: 35 months
- <PR (48) median PFS: 21 months

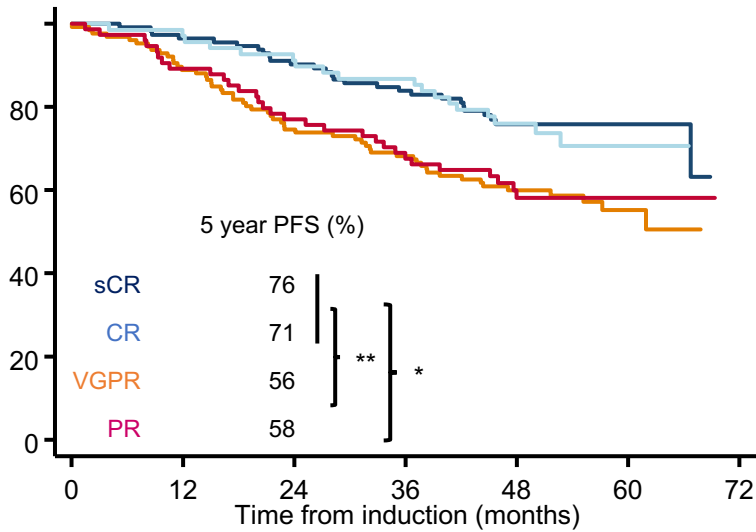


- MRD- (n=326) median OS: Not reached
- CR (n=133) median OS: 68 months
- nCR (n=99) median OS: 73 months
- PR (n=211) median OS: 68 months
- <PR (48) median OS: 46 months

Patients in CR with persistent MRD had the same outcome as patients in nCR/VGPR and even PR

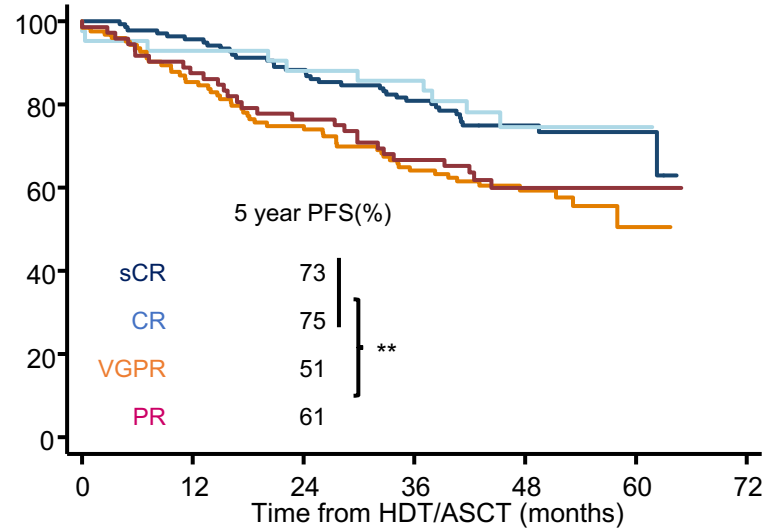
Prognostic value of standard response in patients with MM included in the GEM2012MENOS65 trial

Post-Induction



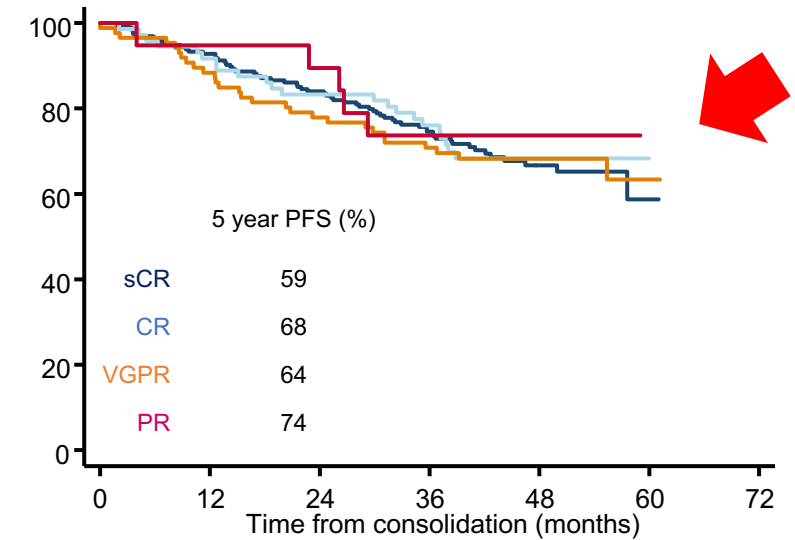
112	108	101	92	60	22	0
69	66	62	59	36	10	0
126	112	94	85	57	17	0
74	66	57	50	33	12	0

Post-ASCT



137	131	120	104	56	12	0
42	39	37	36	18	2	0
123	105	91	78	47	4	0
72	63	55	48	27	6	0

Post-Consolidation



194	180	163	135	55	3	0
72	66	59	51	23	0	0
86	76	66	60	30	3	0
19	18	17	14	6	0	0

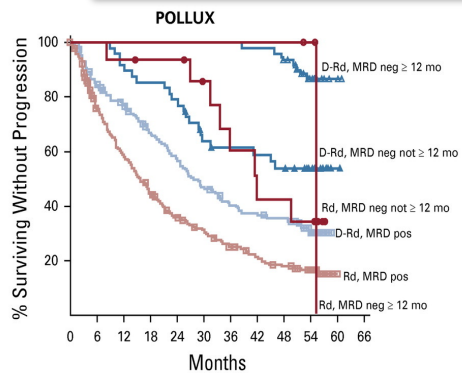
* $P < .05$; ** $P < .01$

IMWG Minimal Residual Disease criteria

IMWG MRD criteria (requires a complete response as defined below)

Sustained MRD-negative	MRD negativity in the marrow (NGF or NGS, or both) and by imaging as defined below, confirmed minimum of 1 year apart. Subsequent evaluations can be used to further specify the duration of negativity (eg, MRD-negative at 5 years)†
Flow MRD-negative	Absence of phenotypically aberrant clonal plasma cells by NGF‡ on bone marrow aspirates using the EuroFlow standard operation procedure for MRD detection in multiple myeloma (or validated equivalent method) with a minimum sensitivity of 1 in 10 ⁵ nucleated cells or higher
Sequencing MRD-negative	Absence of clonal plasma cells by NGS on bone marrow aspirate in which presence of a clone is defined as less than two identical sequencing reads obtained after DNA sequencing of bone marrow aspirates using the LymphoSIGHT platform (or validated equivalent method) with a minimum sensitivity of 1 in 10 ⁵ nucleated cells§ or higher
Imaging-positive MRD-negative	MRD negativity as defined by NGF or NGS plus disappearance of every area of increased tracer uptake found at baseline or a preceding PET/CT or decrease to less mediastinal blood pool SUV or decrease to less than that of surrounding normal tissue¶

Sustained



Flow



Sequencing



Imaging



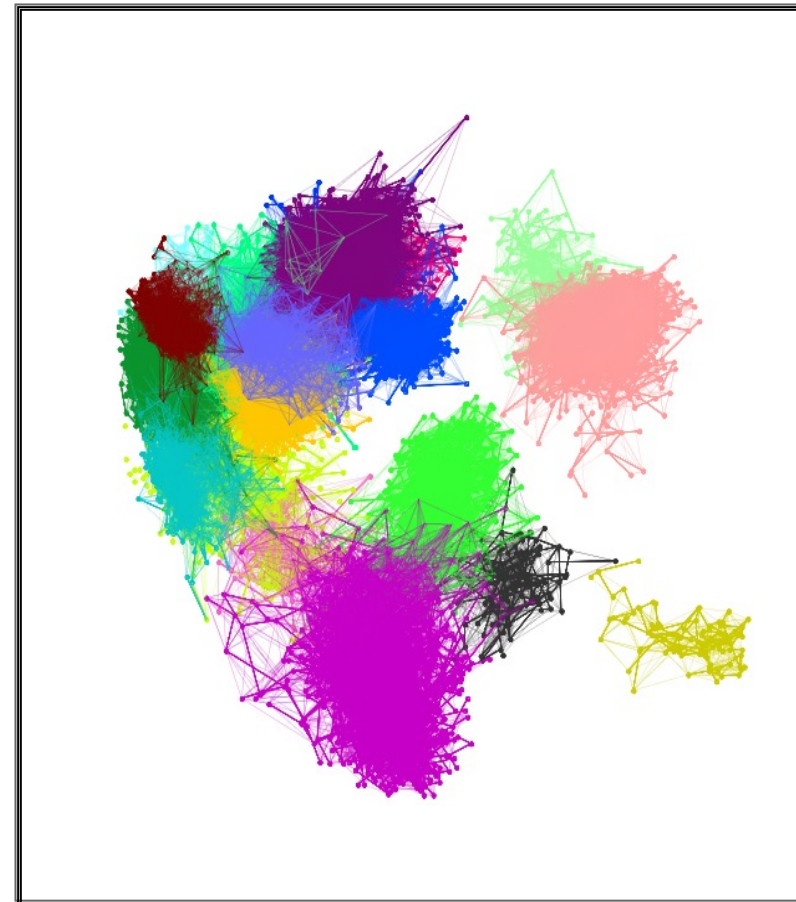
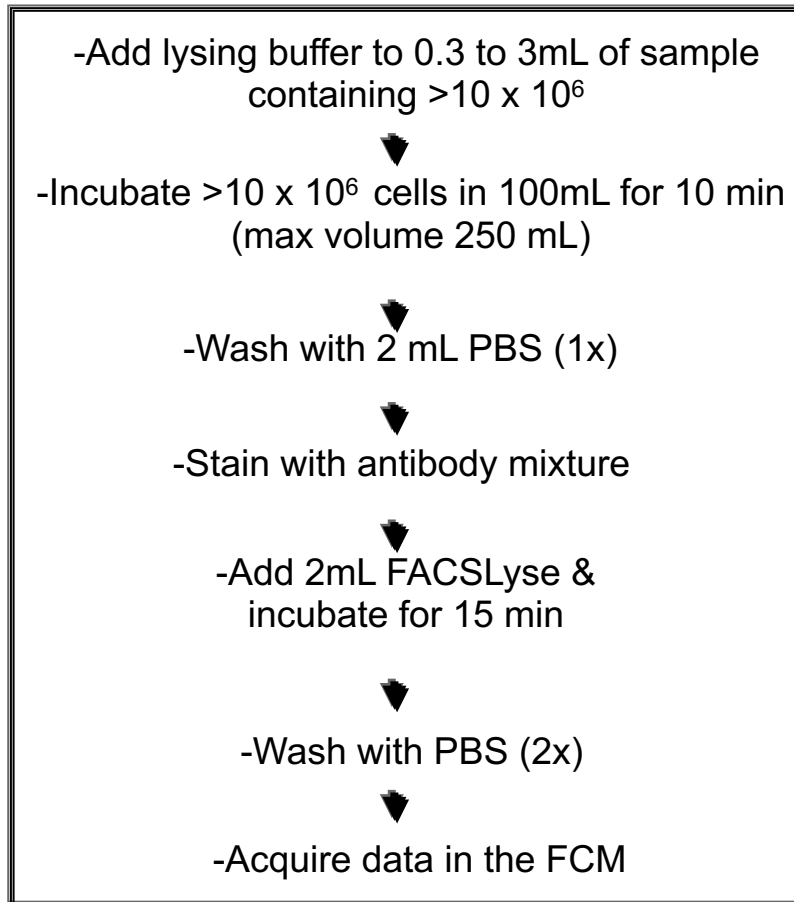
Next generation flow (NGF) cytometry

Optimal antibody panel

Tube	BV421	BV510	FITC	PE	PerCP Cy5.5	PECy7	APC	APCC750
1							CD117	CD81
2	CD138	CD27	CD38	CD56	CD45	CD19	cyKappa	cyLambda

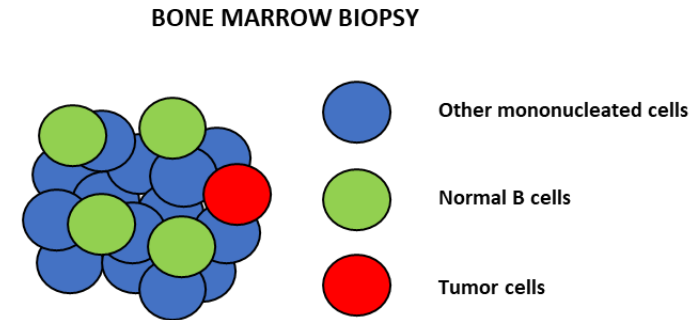
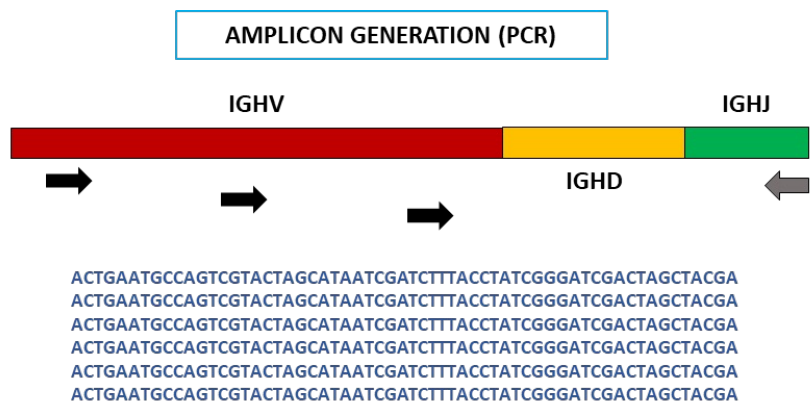
Merged files

Bulk lyse sample processing

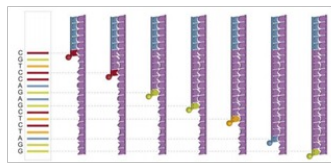


Novel analysis strategies

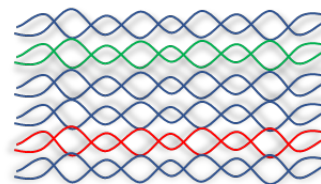
MRD detection in the BM by NGS



SEQUENCING (ILLUMINA MISEQ)



DNA ISOLATION AND QUANTIFICATION



Known quantity of spike in molecules were added (*MMWL1*)



BIOINFORMATICS

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TGGACTAGCATAAACTCGATCGATGATAATCGATCTAGCTATCGATCAATAAGCCTCGTAGCTTAGCTATAATCAGAACGATCAATAAGCCTCGTAGTGATAAA
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Normal B-cell reads (polyclonal)

Spike-in reads (monoclonal)

Tumor reads (monoclonal)

$$Tumor\ cells = \frac{tumor\ reads \times spike\ in\ reads}{spike\ in\ reads}$$

$$MRD\ level\ (\%) = \frac{tumor\ cells}{total\ cells} \times 100$$

Technical features of NGF and NGS for MRD detection

	NGF	NGS
Applicability (% cases)	99%	90%
Sensitivity	$2-4 \times 10^{-6}$	10^{-6}
Time to result	2-3 h	≥ 7 days
Number of cells required	2×10^7	$2-3 \times 10^6$
Need for fresh sample	Yes (within 24h)	No
Need for diagnostic sample	No	Yes
Quantitative	Yes	Yes
Intrinsic quality control for hemodilution	Yes	No
Cell characterization	Yes	No
Molecular characterization	No	Yes
Availability	Wide	Limited
Reproducibility among centers	High	Not reported
Harmonization	Yes	Not reported
Cost	+	++

Technical features of NGF and NGS for MRD detection

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Applicability (% cases)	99%	90%
Sensitivity	2-4 x 10⁻⁶	10⁻⁶
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Number of cells required	2 x 10 ⁷	2-3 x 10 ⁶
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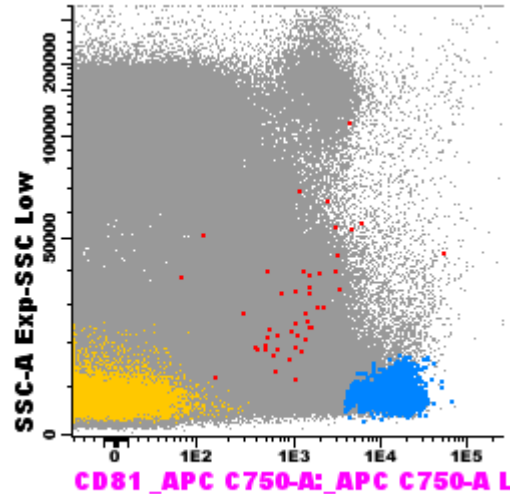
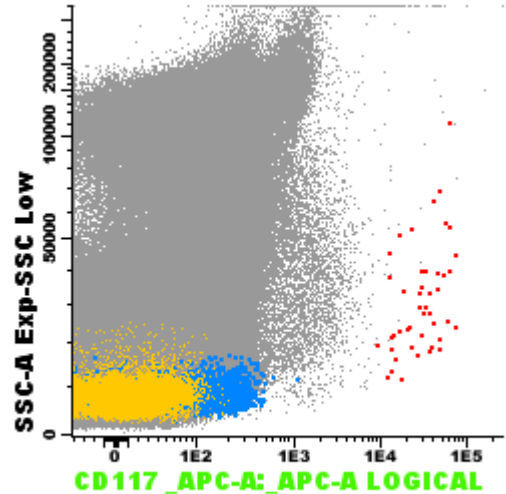
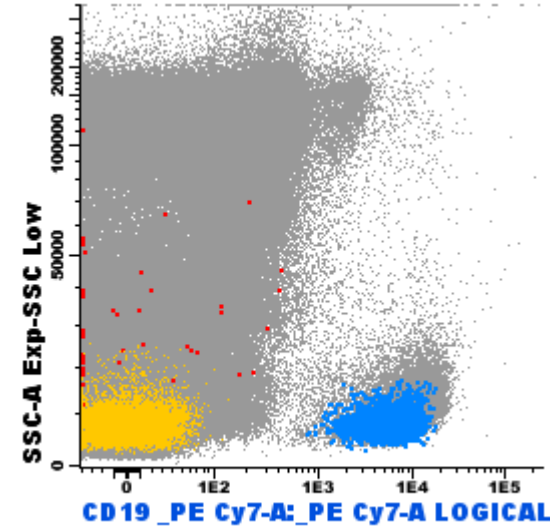
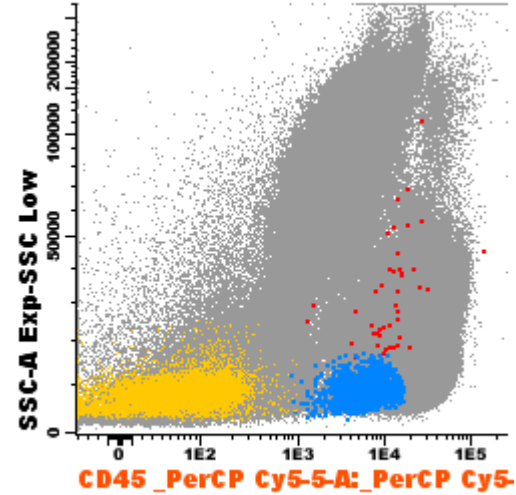
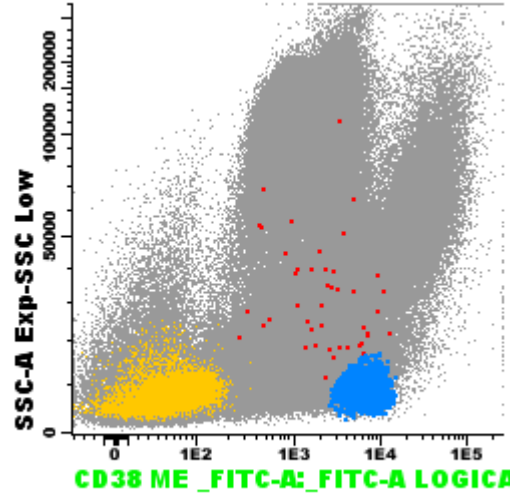
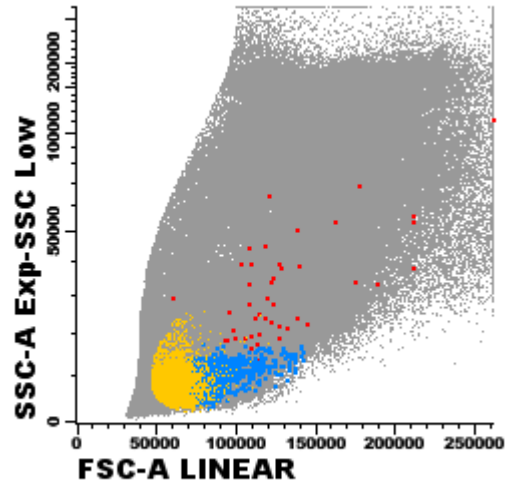
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NGF allows for a quality control check



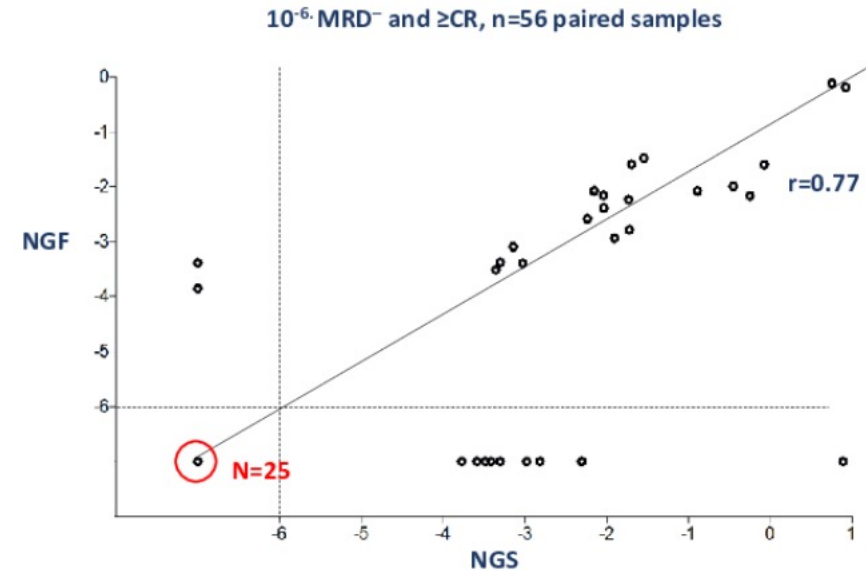
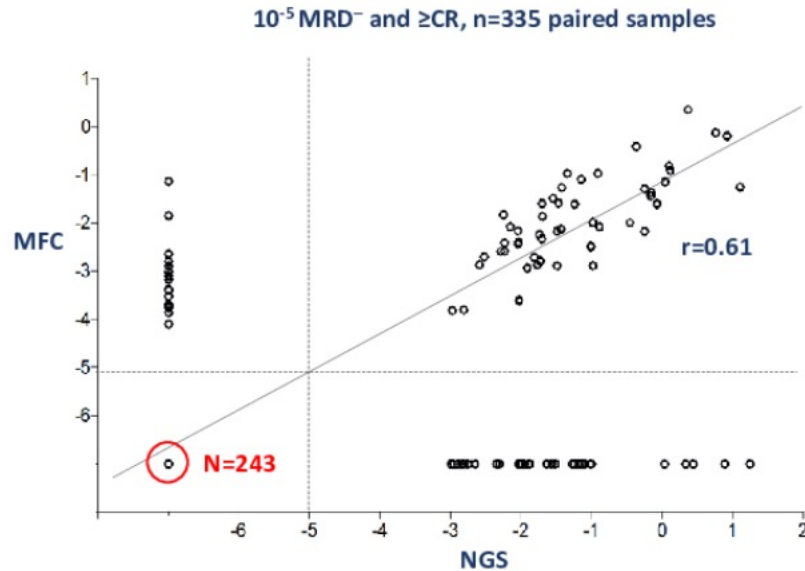
Mast cells (CD117bright, CD45dim)

Nucleated red blood cells (CD45-, CD38-, CD117-/+ , SSClo)

B-cell precursors (CD19+, CD45dim, CD38bright, CD81bright, CD27-)

Concordance between MRD results by the two techniques

FORTE trial Patients achieving VGPR or better pre-maintenance



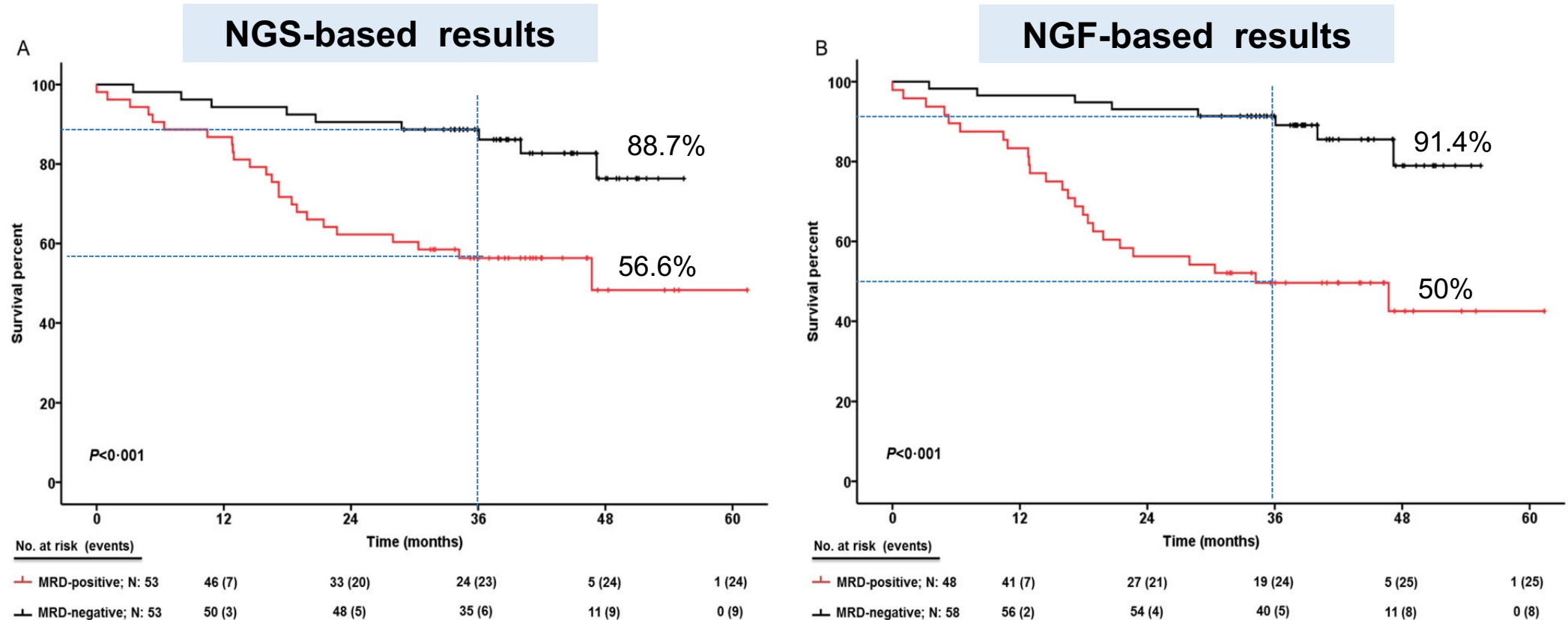
			NGS		
MRD ⁻ and ≥CR, n (%) 10 ⁻⁵	Flow cytometry	Total	Positive	Negative	Observed agreement
MRD status, n (%)	Positive	56	46 (82)	10 (18)	86%
	Negative	279	36 (13)	243 (87)	
			NGS		
MRD ⁻ and ≥CR, n (%) 10 ⁻⁶	NGF	Total	positive	Negative	Observed agreement
MRD status, n (%)	Positive	21	19 (90)	2 (10)	78%
	Negative	35	10 (28)	25 (72)	

MFC and NGS at 10⁻⁵ evaluable samples
(n: 335; r: 0.61)

MFC and NGS at 10⁻⁶ evaluable samples
(n: 56; r: 0.77).

Kaplan–Meier curves comparing PFS of MRD-positive and MRD-negative subsets

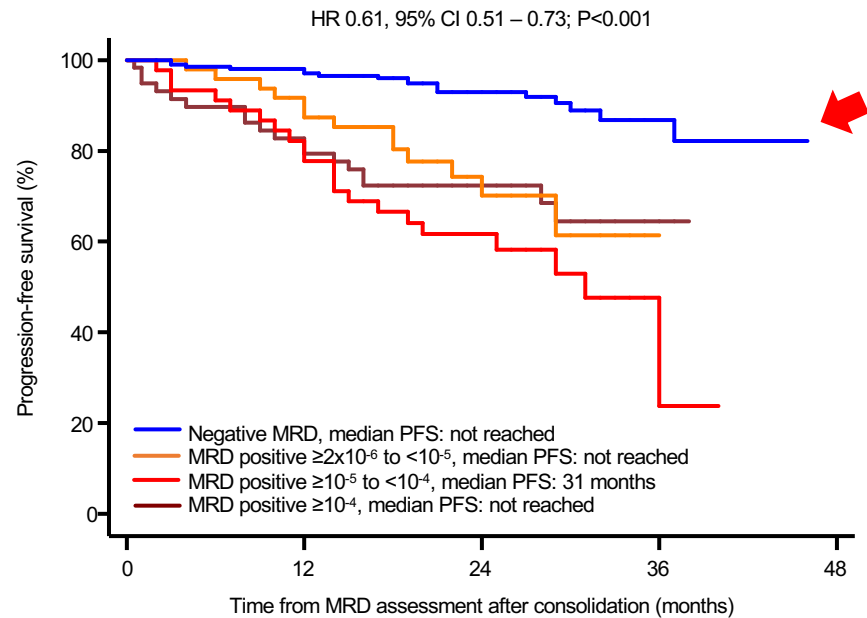
MRD in +100 post-ASCT
n = 106



Correlation between NGS and NGF was high ($R^2 = 0.905$)

The clinical impact of MRD is reproducible in... ... different centers and by different methods in the transplant setting

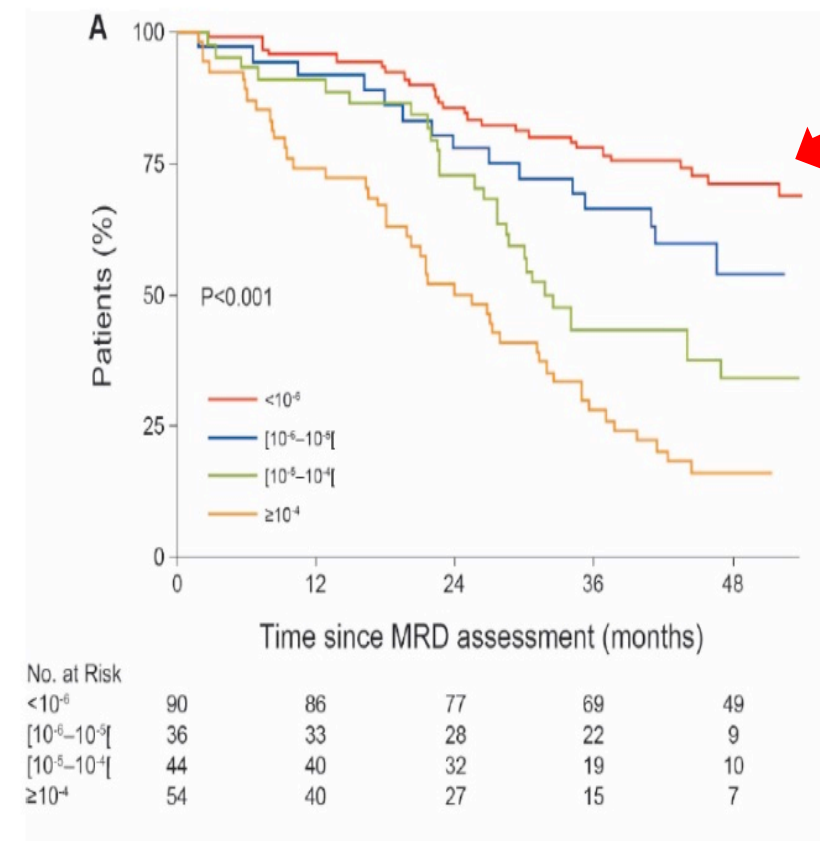
GEM2012: RVD->TASPE: NGF
Pre-maintenance, 2×10^{-6}



Number at risk	0	12	24	36	48
MRD positive $\ge 10^{-4}$	59	48	26	4	0
MRD positive $\ge 10^{-5}$ to $< 10^{-4}$	45	37	20	2	0
MRD positive $\ge 2 \times 10^{-6}$ to $< 10^{-5}$	48	43	18	1	0
Negative MRD	205	198	111	19	0

Paiva B, et al. JCO 2019

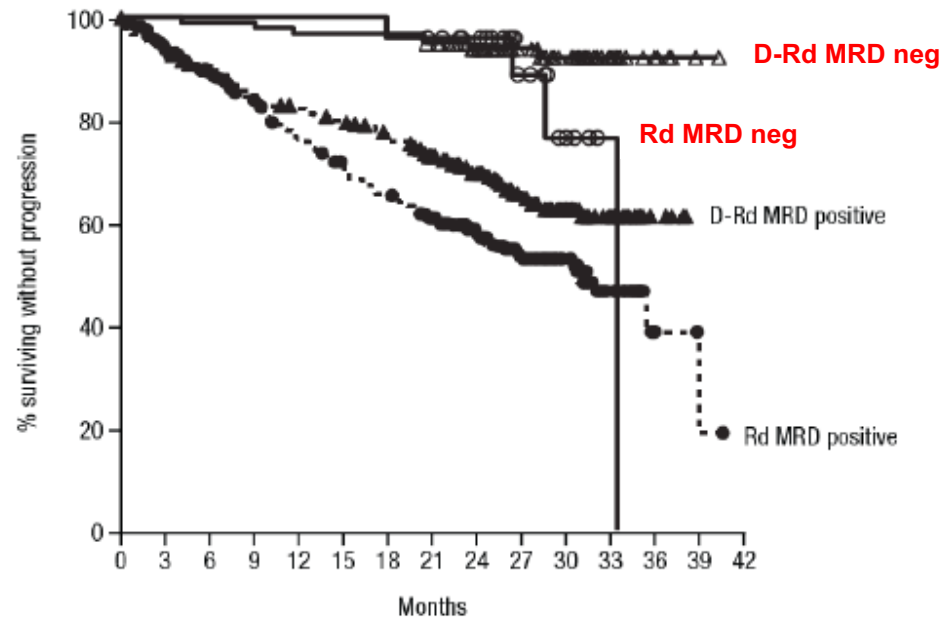
IFM2009: RVD->RVD/TASPE: NGS
Pre-maintenance, 1×10^{-6}



Perrot A, et al. Blood. 2018;132(23):2456-2464

... in the non-transplant setting

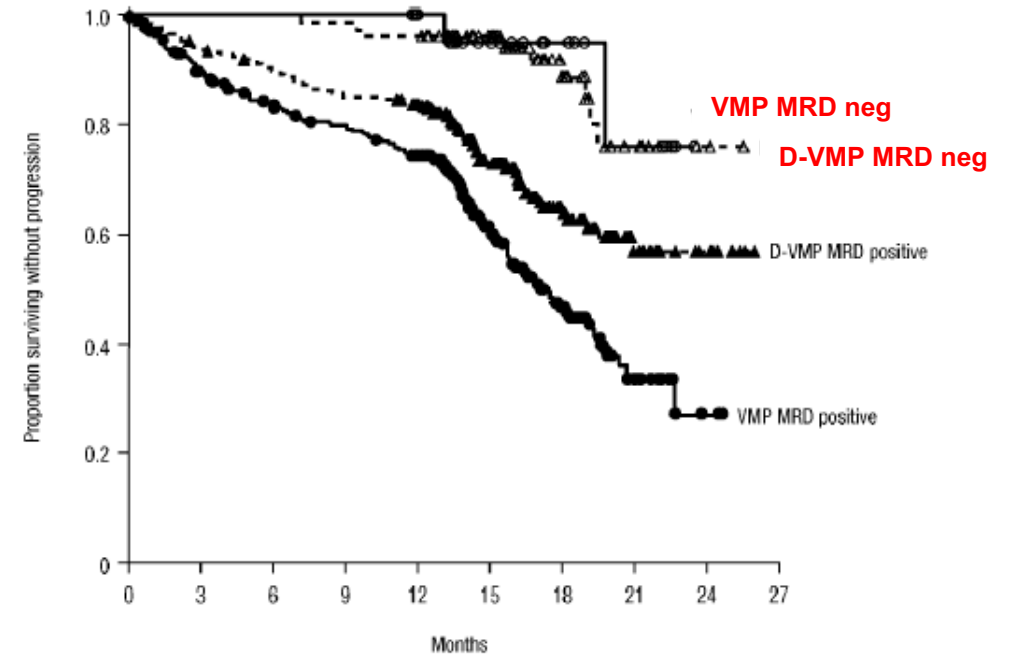
MAIA



No. at risk	0	3	6	9	12	15	18	21	24	27	30	33	36	39	42
Rd MRD negative	27	27	27	27	27	27	27	25	21	12	5	1	0	0	0
D-Rd MRD negative	89	89	88	88	86	86	86	84	70	55	33	12	5	1	0
Rd MRD positive	342	305	280	253	227	209	192	175	128	82	45	17	3	2	0
D-Rd MRD positive	279	258	247	232	223	214	204	187	133	91	53	23	6	0	0

n = 737

ALCYONE



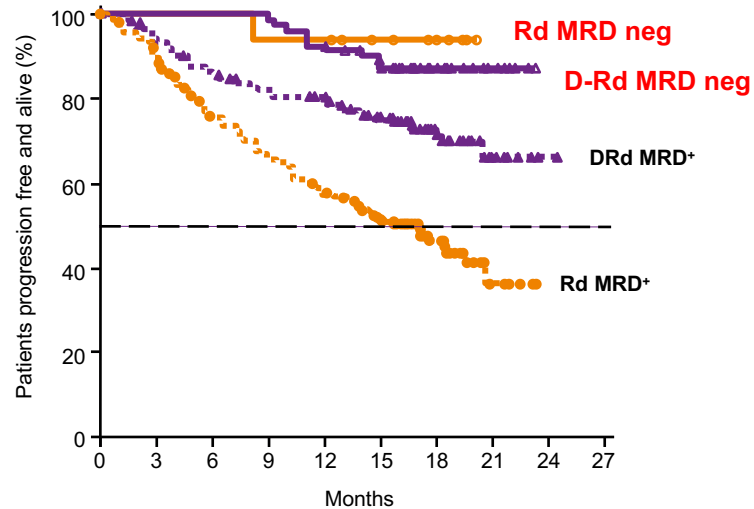
Patients at risk	0	3	6	9	12	15	18	21	24	27
VMP MRD negative	22	22	22	22	21	14	8	4	0	0
D-VMP MRD negative	78	78	78	77	75	58	31	14	2	0
VMP MRD positive	334	281	254	239	210	113	53	14	2	0
D-VMP MRD positive	272	244	234	221	210	121	62	21	8	0

n = 706

Adaptive Biotechnologies clonoSEQ NGS assay (version 2.0)

... and in relapsed or refractory patients

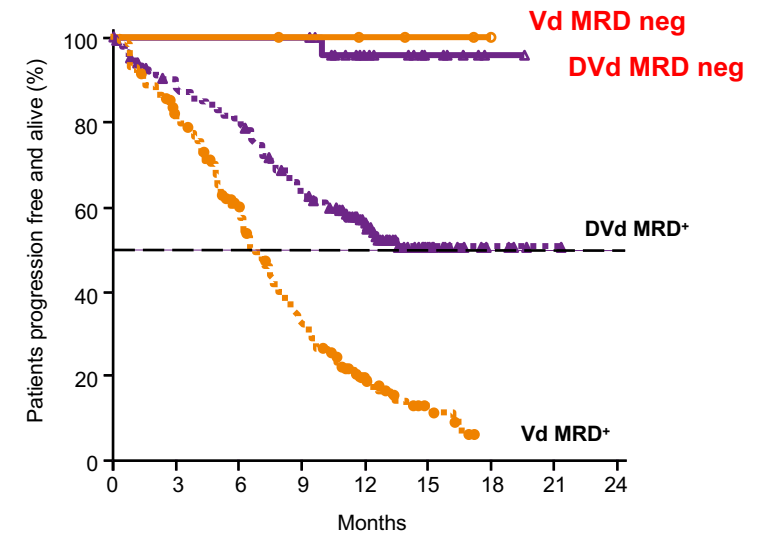
POLLUX



Patients at risk	0	3	6	9	12	15	18	21	24	27
Rd MRD negative	16	16	16	15	15	12	10	0	0	0
DRd MRD negative	71	71	71	70	66	57	28	6	0	0
Rd MRD positive	267	233	190	166	144	120	38	5	0	0
DRd MRD positive	215	195	178	167	161	137	54	9	1	0

n = 569

CASTOR

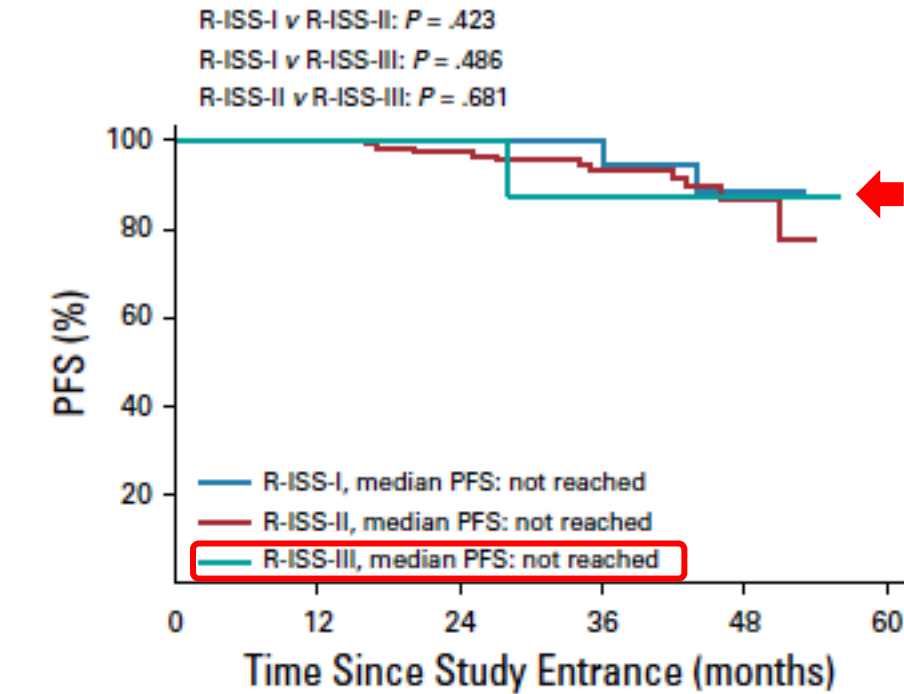


Patients at risk	0	3	6	9	12	15	18	21	24
Vd MRD negative	6	6	6	5	3	2	0	0	0
DVd MRD negative	26	26	26	26	15	7	1	0	0
Vd MRD positive	241	176	123	68	20	7	0	0	0
DVd MRD positive	225	189	172	134	76	26	4	1	0

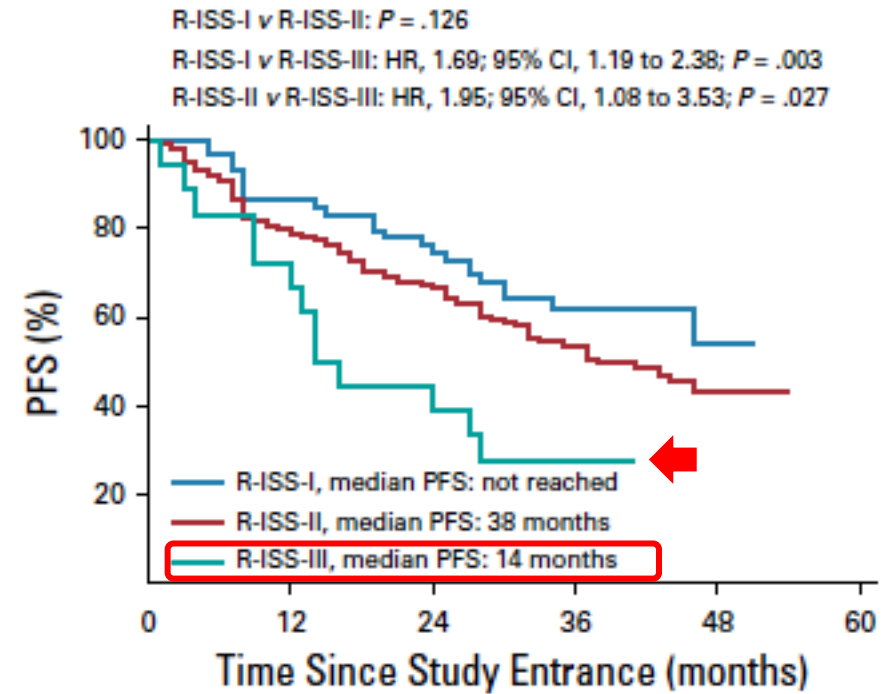
n = 498

Adaptive Biotechnologies clonoSEQ NGS assay

Achieving MRD^{neg} is clinically relevant in standard and high-risk disease



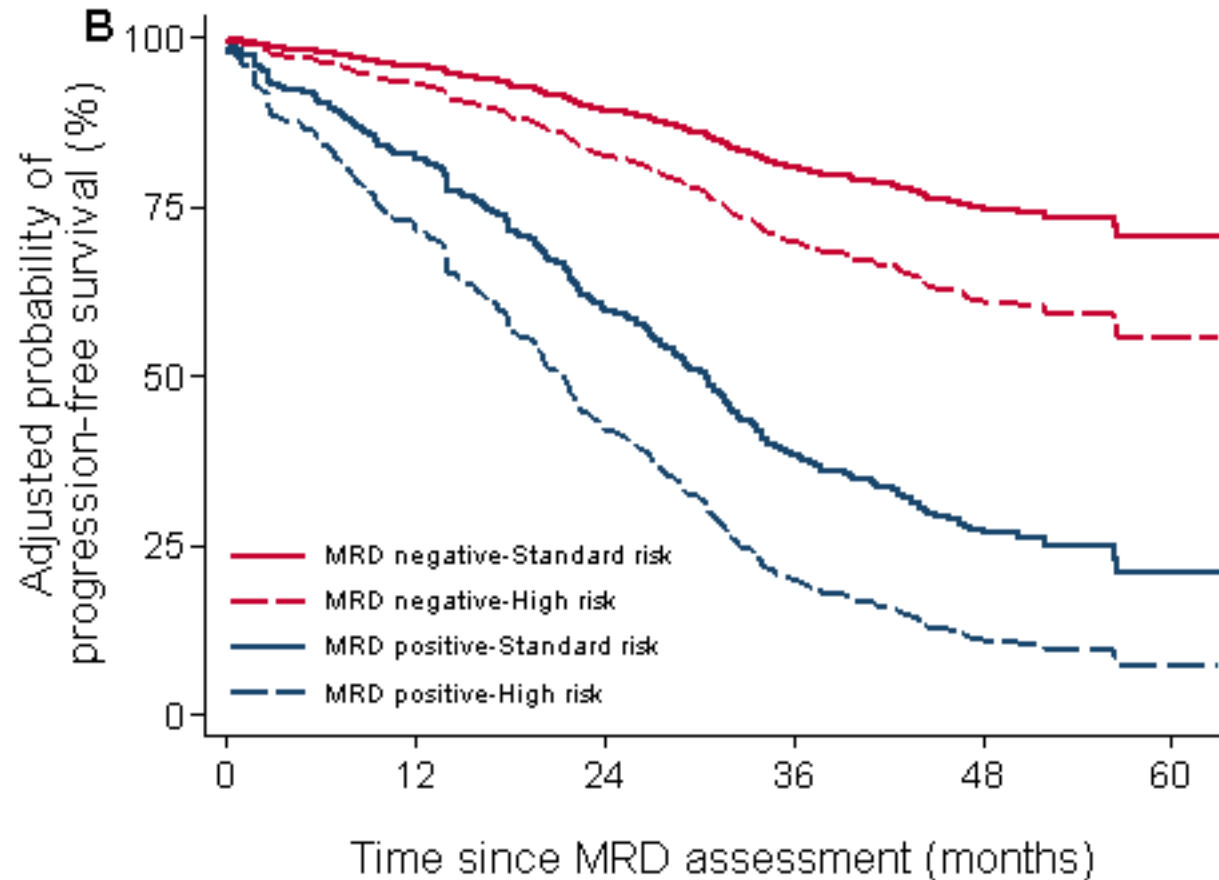
No. at risk	0	12	24	36	48	60
R-ISS-1	55	55	54	37	7	0
R-ISS-2	114	114	111	78	19	0
R-ISS-3	8	8	8	6	1	0



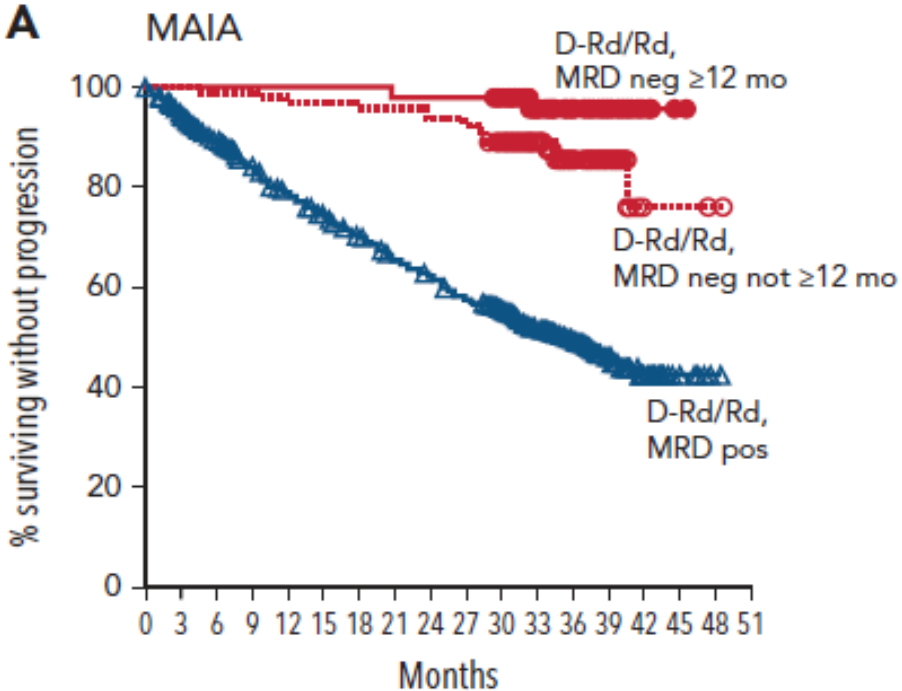
No. at risk	0	12	24	36	48	60
R-ISS-1	59	51	45	26	4	0
R-ISS-2	150	119	99	58	18	0
R-ISS-3	18	13	8	3	0	0

NGF, LoD 2×10^{-6}

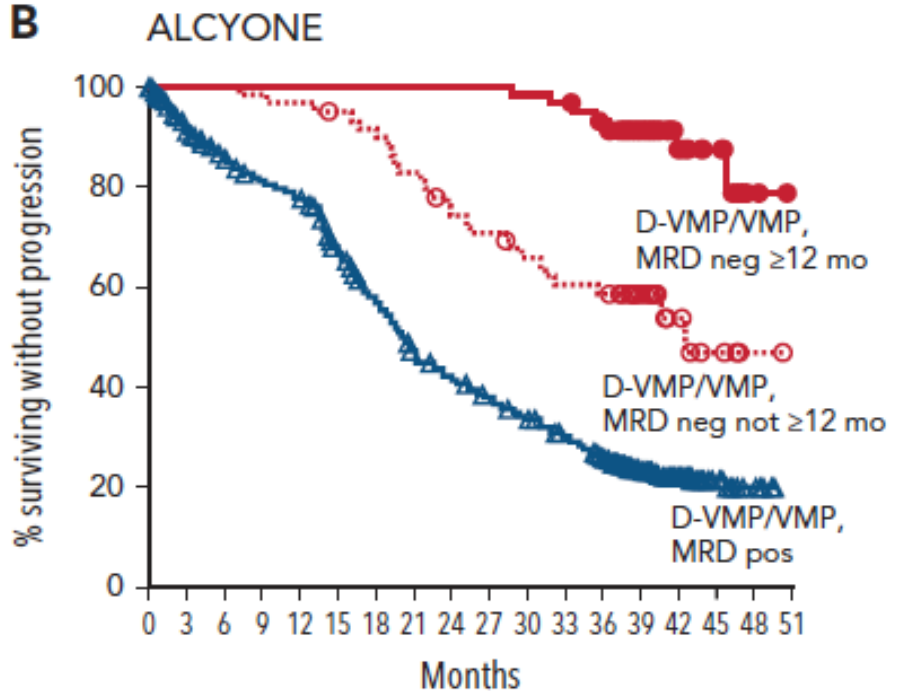
Achieving MRD^{neg} is clinically relevant in standard and high-risk disease



PFS based on sustained MRD negativity in MAIA and ALCYONE trials



No. at risk	0	3	6	9	12	15	18	21	24	27	30	33	36	39	42	45	48	51	
D-Rd/Rd, MRD neg ≥ 12 mo	49	49	49	49	49	49	49	48	48	48	48	47	37	27	13	6	2	0	0
D-Rd/Rd, MRD neg not ≥ 12 mo	91	91	90	90	88	88	88	87	85	84	75	56	41	21	2	2	1	0	0
D-Rd/Rd, MRD pos	597	540	503	461	426	399	372	345	327	301	272	194	127	69	26	5	1	0	0



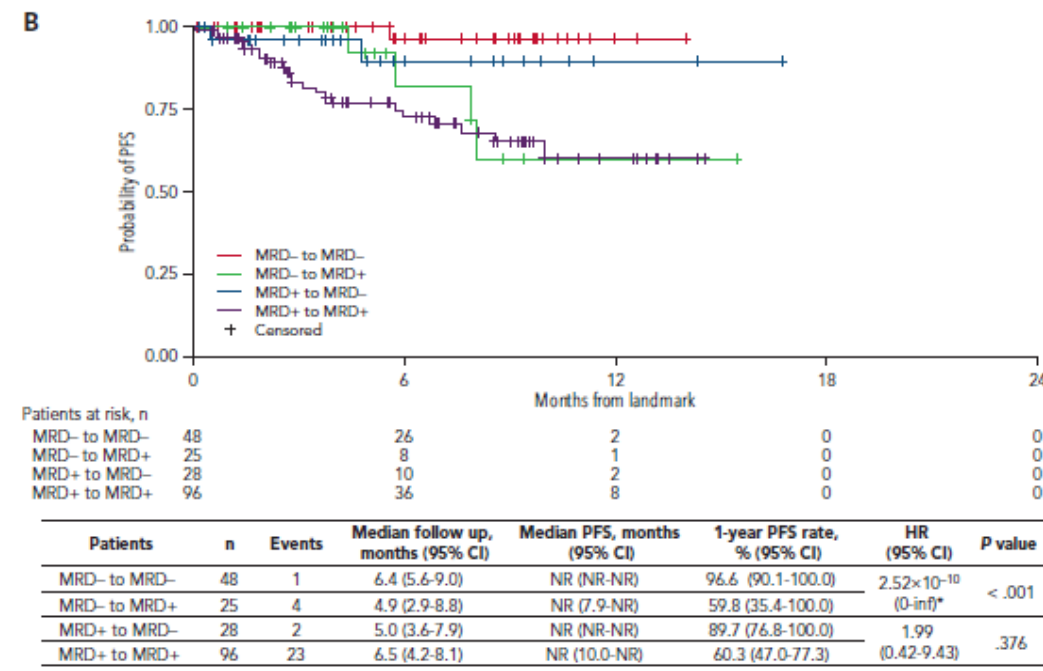
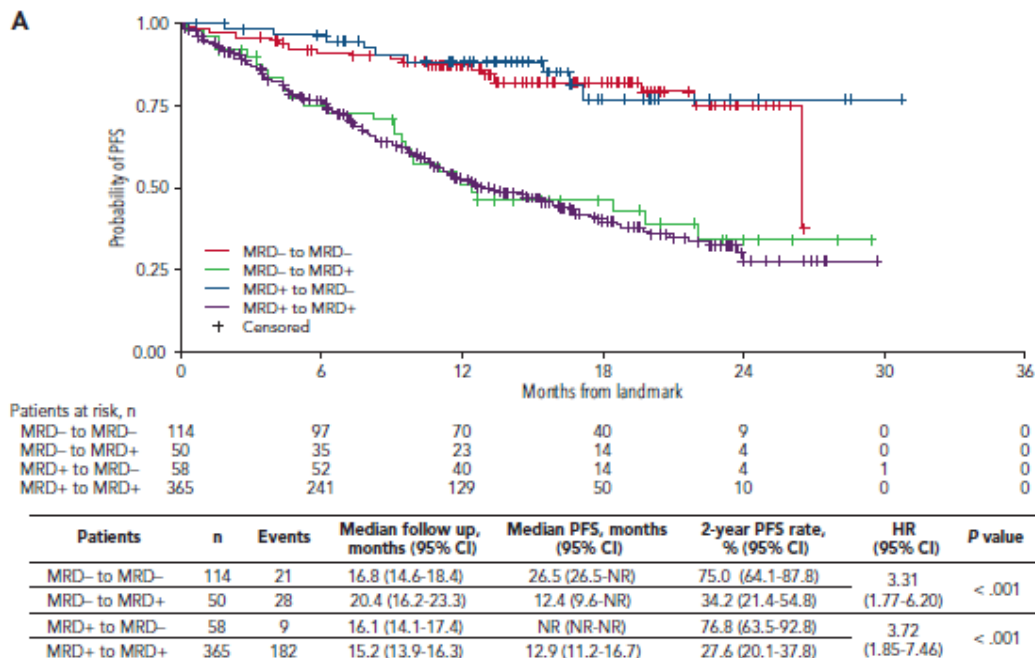
No. at risk	0	3	6	9	12	15	18	21	24	27	30	33	36	39	42	45	48	51	
D-VMP/VMP, MRD neg ≥ 12 mo	59	59	59	59	59	59	59	59	59	59	58	57	53	40	20	11	3	0	0
D-VMP/VMP, MRD neg not ≥ 12 mo	60	60	60	59	58	56	54	49	43	41	37	34	33	21	9	4	1	0	0
D-VMP/VMP, MRD pos	587	507	471	443	421	357	301	240	215	195	171	149	125	81	49	18	5	0	0

KM estimates of PFS by MRD negativity lasting ≥ 12 months among patients in the ITT populations.

Landmark analyses of PFS based on MRD kinetics from randomization in the TOURMALINE-MM3 and -MM4 trials

14 months

28 months

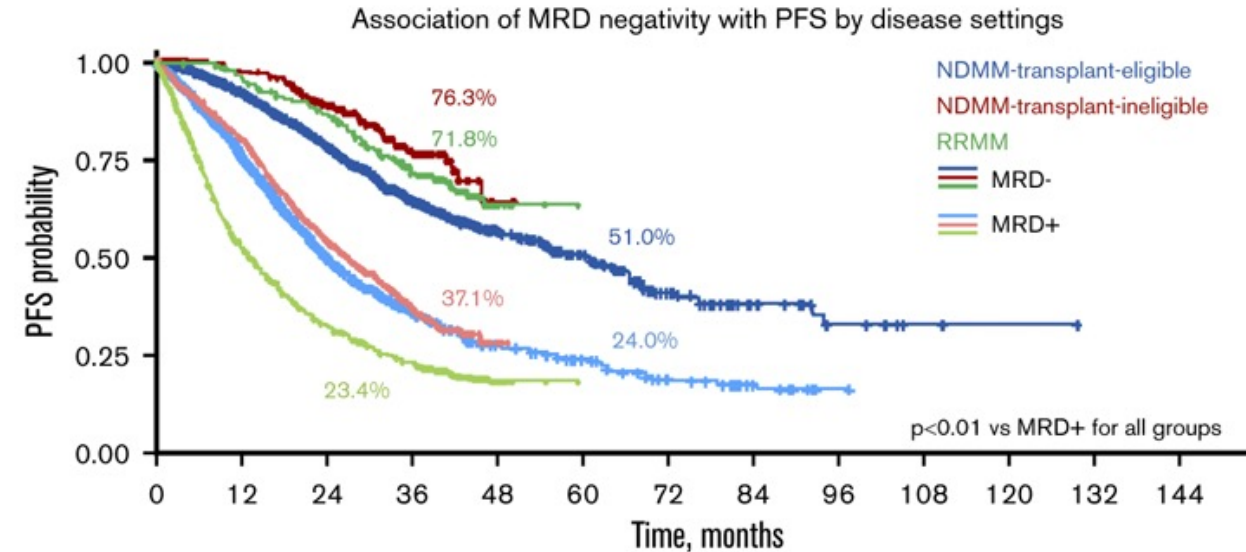


Conversion from **MRD- to MRD+** or from **MRD+ to MRD-** status during ixazomib or placebo maintenance modulates the risk of disease progression

Regardless of the treatment received

Paiva B, et al. Blood. 2023 Feb 9;141(6):579-591

Role of MRD negativity in long-term survival outcomes in patients with multiple myeloma



Number at risk

MRD-	1515	1055	589	332	164	95	47	22	10	3	1	0	0
MRD+	1180	719	317	153	72	50	30	13	2	0	0	0	0
MRD-	291	283	217	93	4	0							
MRD+	1328	983	516	133	5	0							
MRD-	164	155	135	97	10	0							
MRD+	960	456	269	179	11	0							

Clinical trials in which MRD guides treatment decisions

NCT	Official title	Country	Method
NCT02406144	Maintenance treatment with lenalidomide and dexamethasone versus lenalidomide, dexamethasone and ixazomib after autologous hematopoietic stem cell transplantation in patients With newly diagnosed symptomatic multiple myeloma-duration of maintenance guided by MRD status (GEM2014MAIN)	Spain	NGF
RADAR*	Risk adapted therapy directed according to response comparing treatment escalation and de-escalation strategies in newly diagnosed patients with multiple myeloma suitable for stem cell transplantation	UK	N/A
NCT03490344	Short course daratumumab in minimal residual disease (MRD) positive myeloma patients after induction therapy with/without consolidative high-dose chemotherapy/autologous stem cell support	USA	MFC
NCT03224507	Monoclonal antibody-based sequential therapy for deep remission in multiple myeloma (MASTER)	USA	NGS
NCT03742297*	Induction therapy with bortezomib-melphalan and prednisone (VMP) followed by lenalidomide and dexamethasone (Rd) versus carfilzomib, lenalidomide, and dexamethasone (KRd) plus/minus daratumumab, 18 cycles, followed by consolidation and maintenance therapy with lenalidomide and daratumumab: phase III, multicenter, randomized trial for elderly fit newly diagnosed multiple myeloma patients aged between 65 and 80 years	Spain	NGF
NCT03697655	Pre-emptive daratumumab therapy of minimal residual disease reappearance or biochemical relapse in multiple myeloma (PREDATOR)	Poland	N/A
NCT03710603	A phase 3 study comparing daratumumab, VELCADE (Bortezomib), lenalidomide, and dexamethasone (D-VRd) vs VELCADE, lenalidomide, and dexamethasone (VRd) in subjects with previously untreated multiple myeloma who are eligible for high-dose therapy (PERSEUS)	EMN	N/A
NCT03992170	A pilot study on the efficacy of daratumumab in multiple myeloma (MM) patients in >VGPR/MRD-positive by next-generation flow (DART4MM)	Italy	FC
NCT02969837	Open-label, single-arm, phase 2 study of initial treatment with elotuzumab, carfilzomib (Kyprolis), lenalidomide (Revlimid), and low-dose dexamethasone (E-KRd) in newly diagnosed, multiple myeloma requiring systemic chemotherapy	USA	NGS and MFC
NCT04071457	S1803, phase III study of daratumumab/rHuPH20 (NSC-810307) + lenalidomide or lenalidomide as postautologous stem cell transplant maintenance therapy in patients with multiple myeloma (MM) using minimal residual disease to direct therapy duration (DRAMMATIC study)	USA	NGS
NCT04096066	Carfilzomib-lenalidomide-dexamethasone (KRd) versus lenalidomide-dexamethasone (Rd) in newly diagnosed myeloma patients not eligible for autologous stem cell transplantation: a randomized phase III trial	Italy	N/A
NCT03376477	A randomized, double-blind, placebo-controlled phase II trial of an allogeneic myeloma GM-CSF vaccine with lenalidomide in multiple myeloma patients in complete or near complete	USA	NGS
NCT04108624	A multimodality approach to minimal residual disease detection to guide post-transplant maintenance therapy in multiple myeloma (MRD2STOP)	USA	NGS
NCT04221178	A single-arm, prospective study of maintenance therapy cessation for patients with multiple myeloma in sustained MRD-negative remissions	USA	NGF
NCT04140162	Phase 2 study with minimal residual disease (MRD) driven adaptive strategy in treatment for newly diagnosed multiple myeloma (MM) with upfront daratumumab-based therapy	USA	N/A

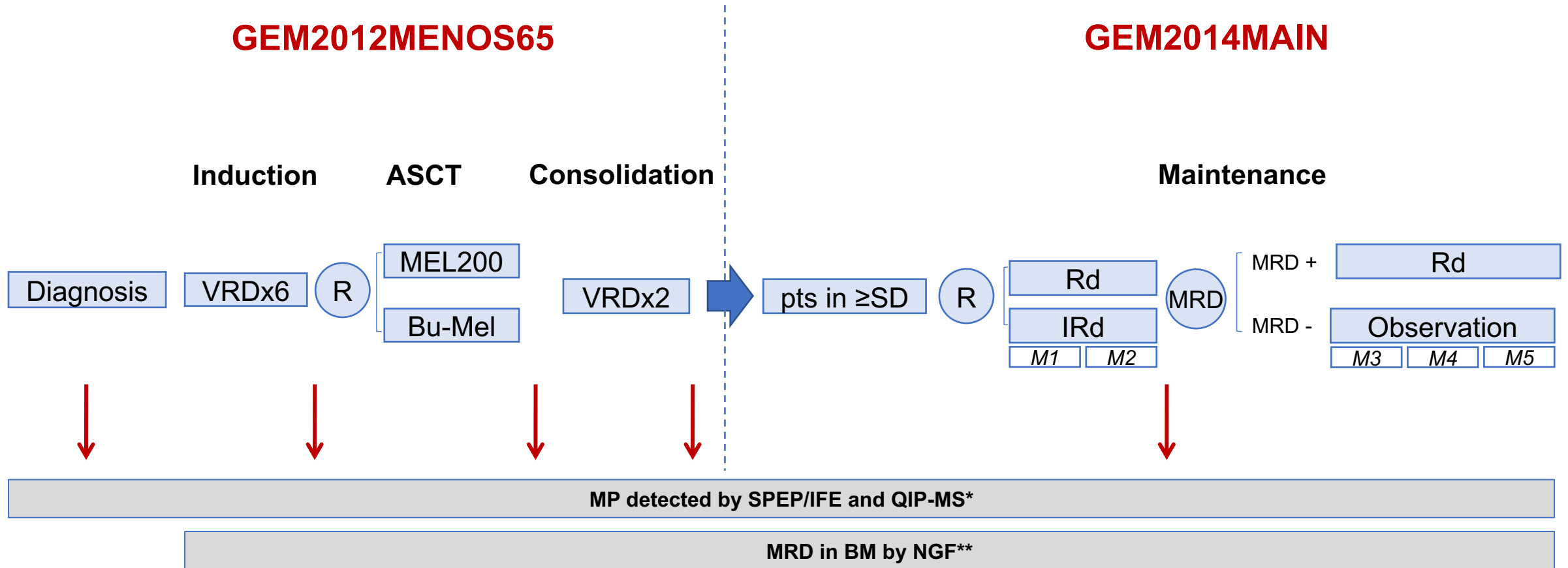
Ixazomib plus Lenalidomide/dexamethasone (IRd) versus lenalidomide/dexamethasone (Rd) maintenance after autologous stem cell transplant in patients with newly diagnosed multiple myeloma: results of the Spanish GEM2014MAIN trial

L. Rosiñol, A. Oriol, R. Rios, M^a J. Blanchard, I. Jarque, J. Bargay, M.T. Hernández, J. M. Moraleda, E. Carrillo, A. Sureda, J. Martínez-López, I. Krsnik, M.E. González, F. Casado, J.M. Martí, C. Encinas, F. de Arriba, L. Palomera, A. Sampol, Y. González-Montes, E. Cabezudo, M^a V. Mateos, J.F. San Miguel, J.J Lahuerta J. Bladé on behalf of the PETHEMA/GEM group.

ASH Annual Meeting, Atlanta, December 12, 2021



Monitoring of TE-NDMM patients: GEM trials

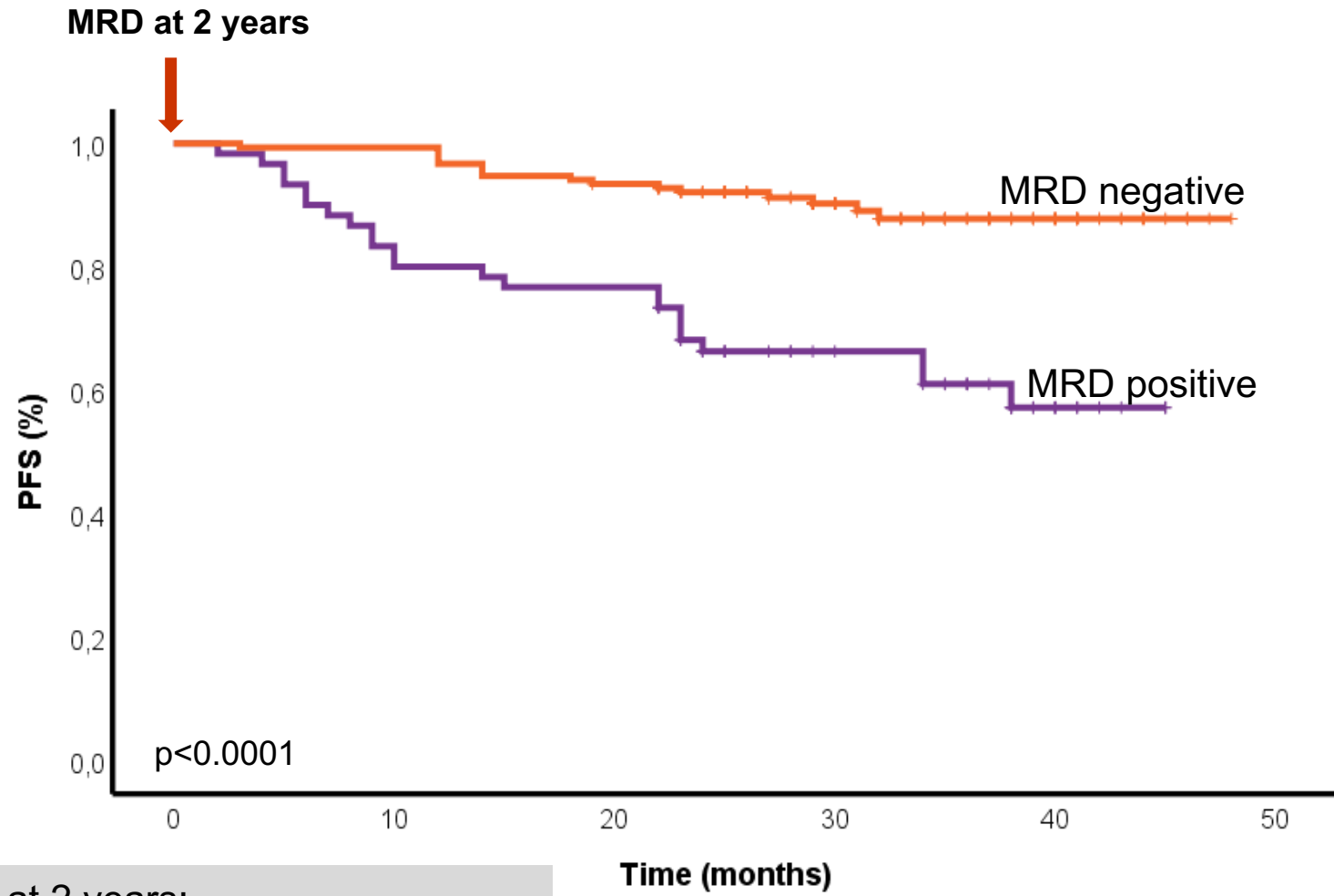


V: Bortezomib 1.3 mg/m² sc days 1, 4, 8, and 11, R: lenalidomide 25 mg po od days 1-21 and D: dexamethasone 40 mg po days 1-4 and 9-12 at 4-week intervals for 6 cycles; MEL200: melphalan 200 mg/m²; Bu-Mel: busulfan 9.6 mg/kg + melphalan 140 mg/m²

I: Ixazomib 4 mg po od days 1, 8, and 15; R: lenalidomide 15 mg po od days 1-21; d: dexamethasone 40 mg po od days 1-4 and 9-12, in 4 weeks cycles

*Performed using IgG/A/M and total $\kappa/\lambda \pm \kappa/\lambda$ free specific beads; ** following the Euroflow guidelines (sensitivity $\geq 10^{-5}$)

PFS from MRD at 2 years

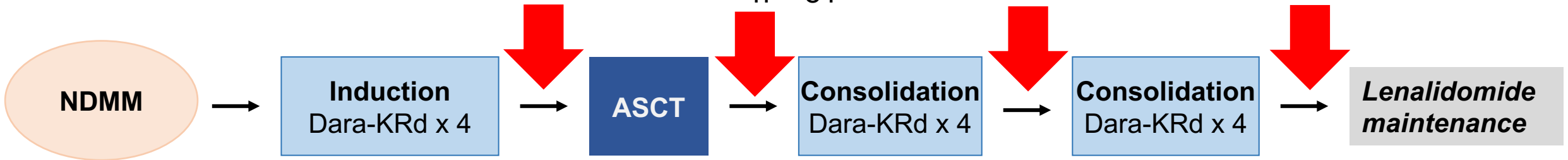


MRD at 2 years:

- negative: stop maintenance
- positive: Rd for 3 additional years

MASTER trial

Multicenter, single-arm phase II trial
n = 81

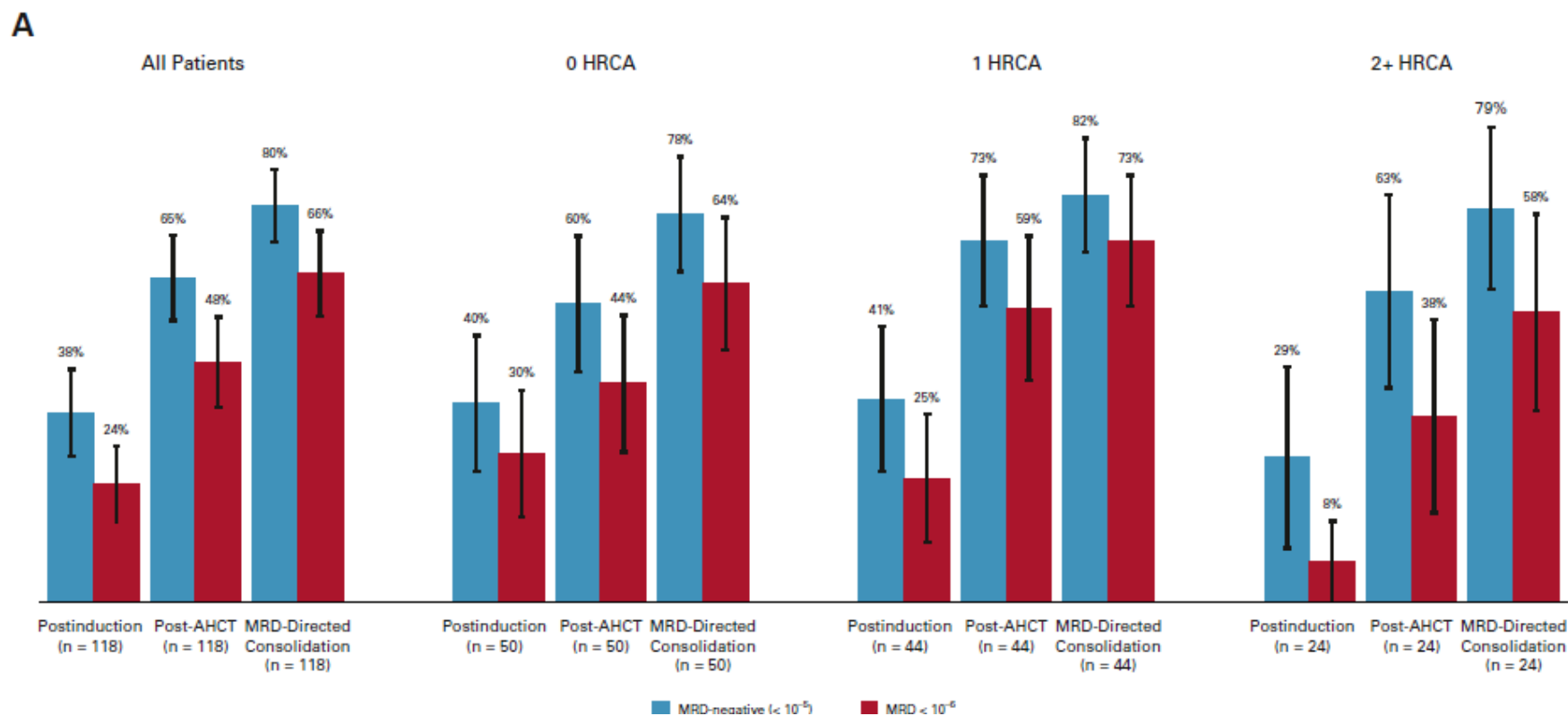


MRD assessment after each treatment phase; pts with confirmed (2nd) MRD-neg status ($<10^{-5}$) entered treatment-free observation phase with MRD assessment at 24 and 72 wks after EOT

- Primary endpoint: MRD-negative remission ($< 10^{-5}$) on NGS assay in pts receiving induction, AHCT, and response-adapted consolidation
- Secondary endpoints: safety, imaging frequency plus remission, MRD status post-AHCT, IMWG response, loss of MRD negativity in pts with no maintenance therapy
- Exploratory endpoint: MRD-negative rates on NGS assay (threshold $< 10^{-6}$)

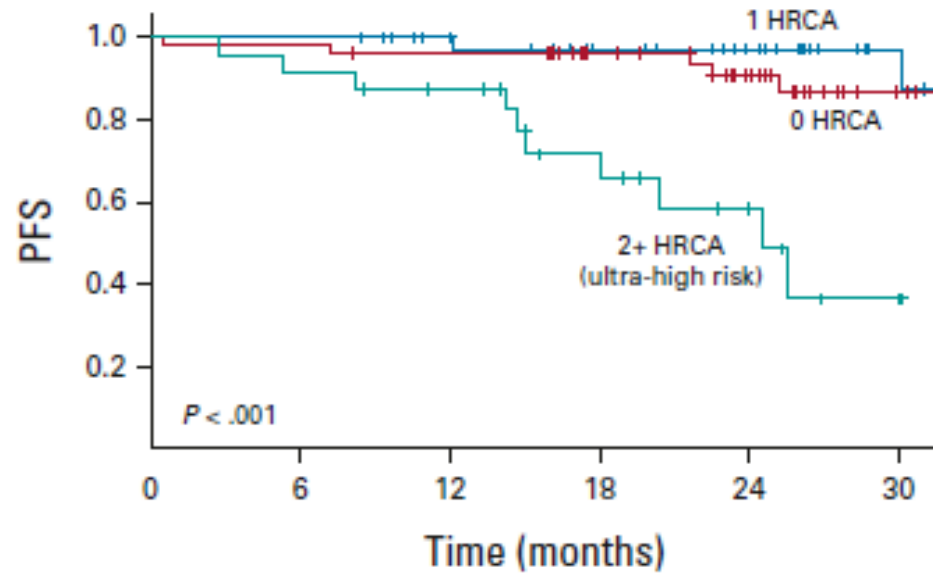
Dara-KRd dosing: daratumumab 16 mg/m² on Days 1,8,15,22 (Days 1,15 of Cycles 3-6; Day 1 Cycle > 6); carfilzomib 56 mg/m² Days 1,8,15; lenalidomide 25 mg Days 1-21; dexamethasone 40 mg PO Days 1,8,15,22. *1 VCD cycle permitted. †Planned recruitment N = 123.

Achievement of MRD negativity (MRD <math><10^{-5}</math> and MRD <math><10^{-6}</math>) according to phase of therapy and number of HRCA



PFS and OS for all participants according to the presence of HRCA in the MASTER trial

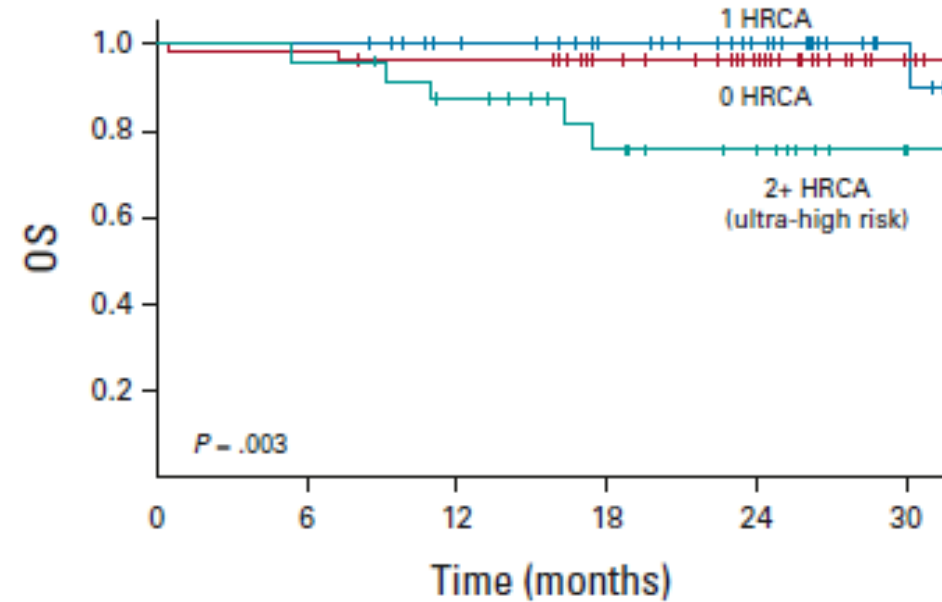
A



No. at risk:

	0	6	12	18	24	30
0 HRCA	50	49	46	36	27	10
1 HRCA	44	44	36	30	23	9
2+ HRCA	24	22	19	12	7	2

B

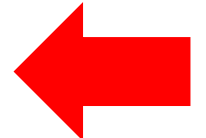


No. at risk:

	0	6	12	18	24	30
0 HRCA	50	49	46	36	29	11
1 HRCA	44	44	36	30	23	9
2+ HRCA	24	23	19	13	9	3

2016 IMWG Criteria for MRD in MM

IMWG MRD Negativity Criteria (Requires CR)	Sustained MRD negative	<ul style="list-style-type: none">MRD negative in marrow (NGF or NGS) and by imaging as defined below, confirmed ≥ 1 year apart. Subsequent evaluations can be used to further specify duration of negativity (eg, MRD negative at 5 years)
	Imaging MRD negative	<ul style="list-style-type: none">MRD negative as defined below (NGF or NGS) PLUS disappearance of all areas of increased tracer uptake observed at baseline or previous PETs/CTs or decrease to less mediastinal blood pool SUV or decrease to less than that of surrounding normal tissue
	Flow MRD negative	<ul style="list-style-type: none">Absence of phenotypically aberrant clonal PCs by NGF on bone marrow aspirates using the EuroFlow standard operating procedure for MRD detection in MM (or equivalent validated method) with minimum sensitivity of 1 in 10^5 nucleated cells
	Sequencing MRD negative	<ul style="list-style-type: none">Absence of clonal PCs by NGS on bone marrow aspirates where presence of a clone is defined as < 2 identical sequencing reads obtained after DNA sequencing of bone marrow aspirates using the LymphoSIGHT[®] platform (or equivalent validated method) with minimum sensitivity of 1 in 10^5 nucleated cells



CR, complete response; CT, computed tomography; IMWG, International Myeloma Working Group; MM, multiple myeloma; MRD, minimal residual disease; NGF, next-generation flow; NGS, next-generation sequencing; PC, plasma cell; PET, positron emission tomography; SUV, standardized uptake value. LymphoSIGHT[®] is a registered trademark of Sequentia, Inc.

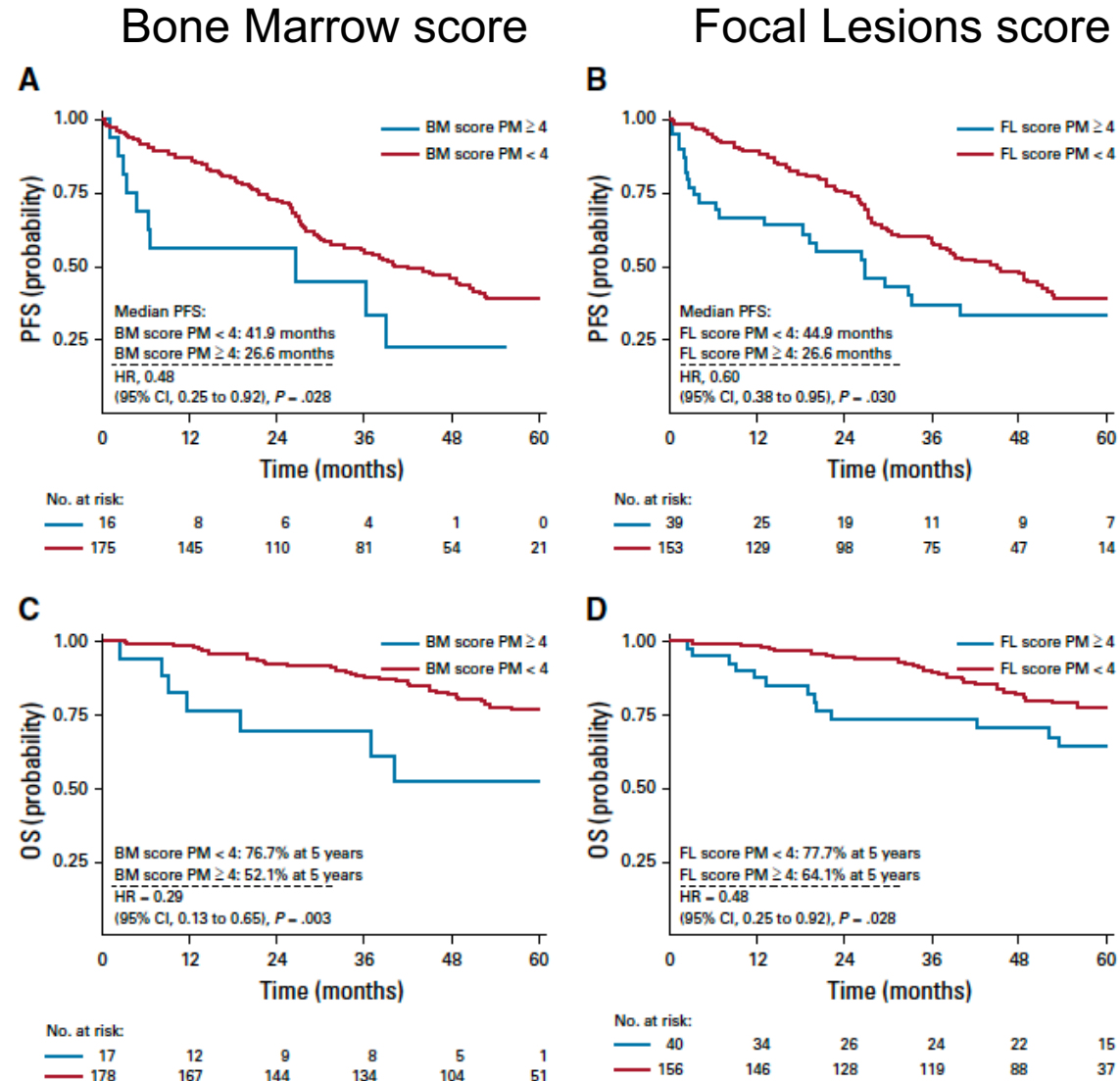
PET/CT

- 18-FDG is currently considered the gold standard to monitor treatment response
- Deauville scores proved to be applicable and representative of patients' outcomes
- 10-15% false negatives due to the lack of hexokinase (need for new tracers)
- Prognostic value
- Lower sensitivity than DW-MRI to detect both diffuse infiltration and focal lesions

PET/CT

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PFS and OS according to pre-maintenance PET/CT



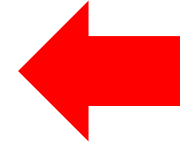
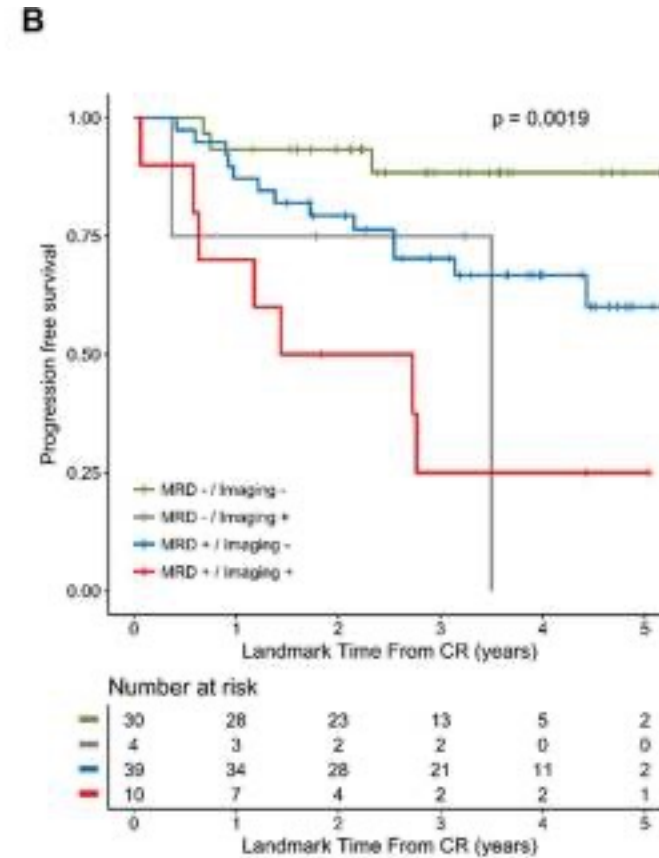
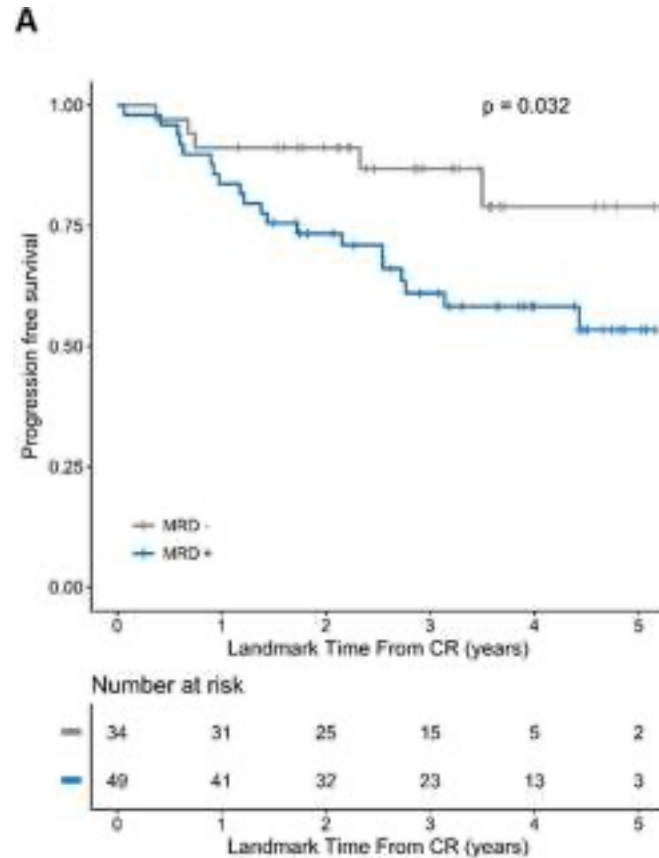
PET/CT

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- Deauville scores proved to be applicable and representative of patients' outcomes
- 10-15% false negatives due to the lack of hexokinase (need for new tracers)
- **Prognostic value**
- Lower sensitivity than DW-MRI to detect both diffuse infiltration and focal lesions

Complementarity between imaging and BM techniques in defining the prognosis of patients



PET/CT

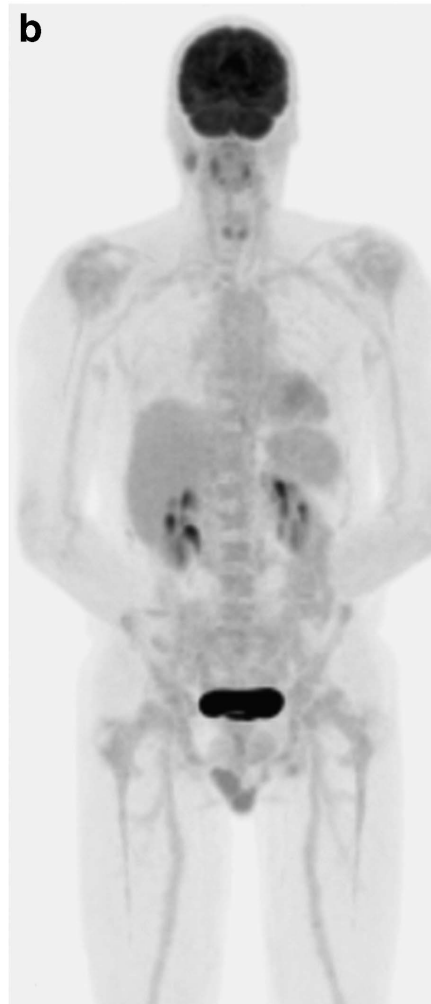
- 18-FDG is currently considered the gold standard to monitor treatment response
- Deauville scores proved to be applicable and representative of patients' outcomes
- 10-15% false negatives due to the lack of hexokinase (need for new tracers)
- Prognostic value
- **Lower sensitivity than DW-MRI to detect both diffuse infiltration and focal lesions**

DW MRI

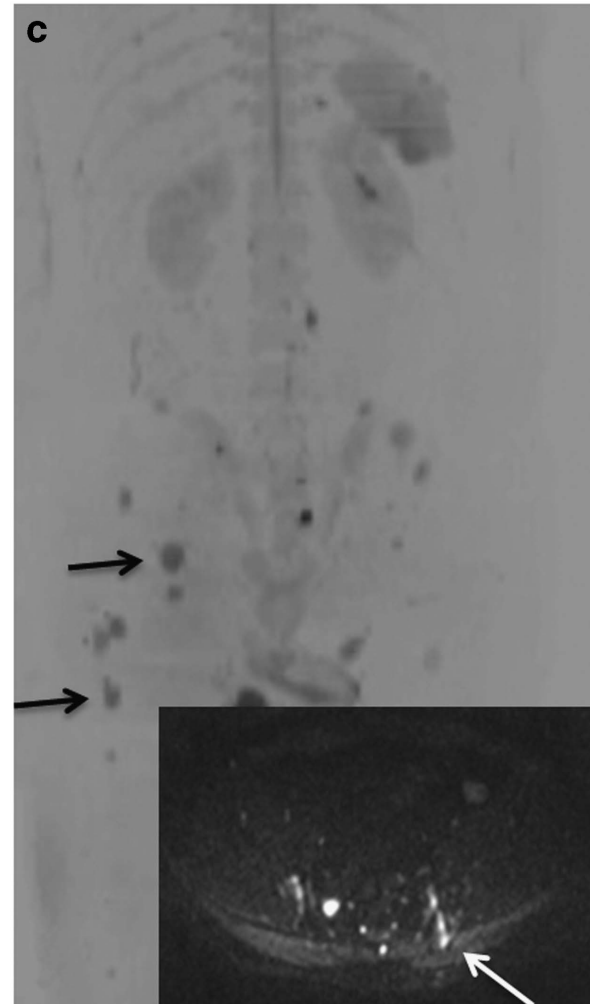
WB-DWI improves detection of diffuse BM infiltration compared with FDG PET-CT and detects trephine sampling error



WB-DWI



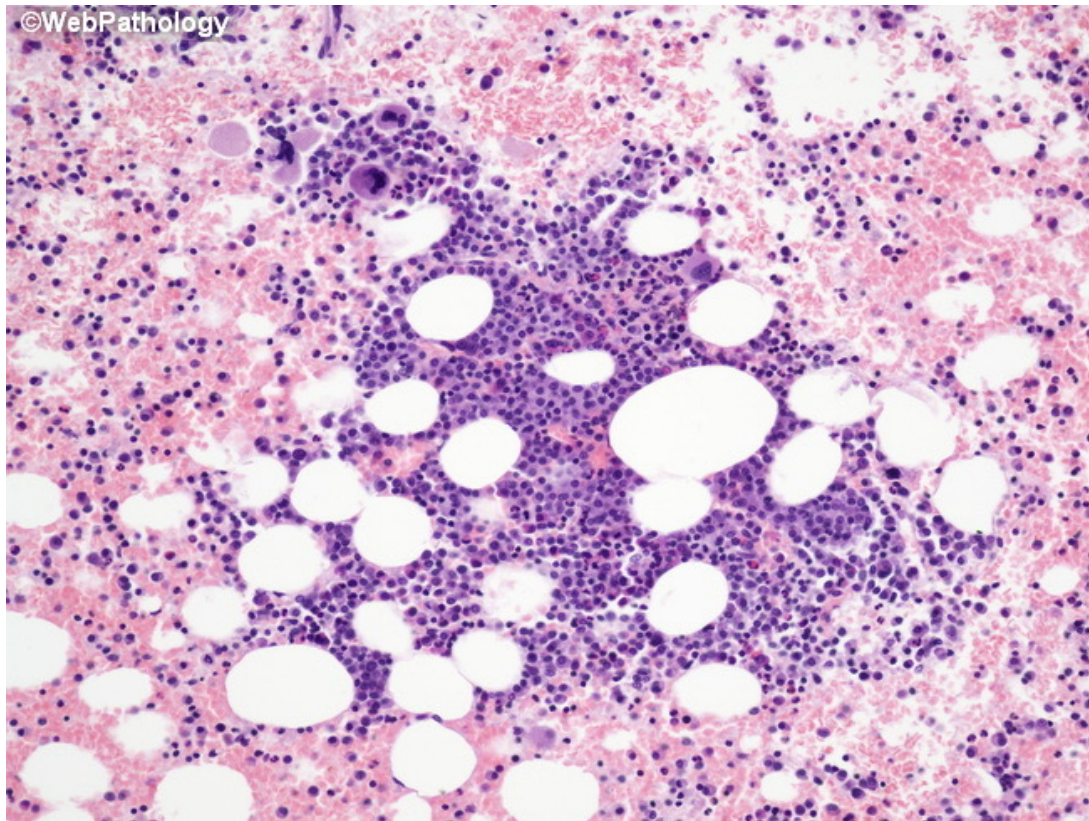
FDG PET-CT



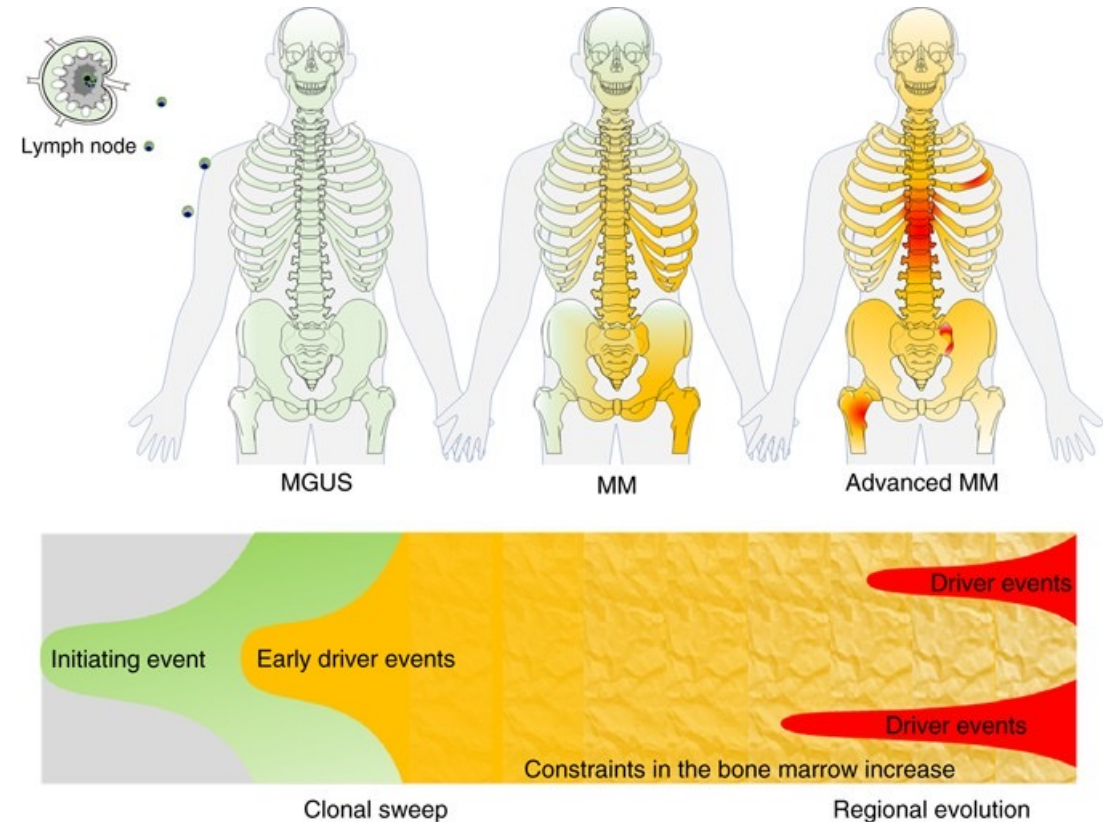
Pawlyn C, et al Leukemia. 2016 Jun;30(6):1446-8

Limitations of MRD assessment in MM

Patchy pattern of bone marrow infiltration

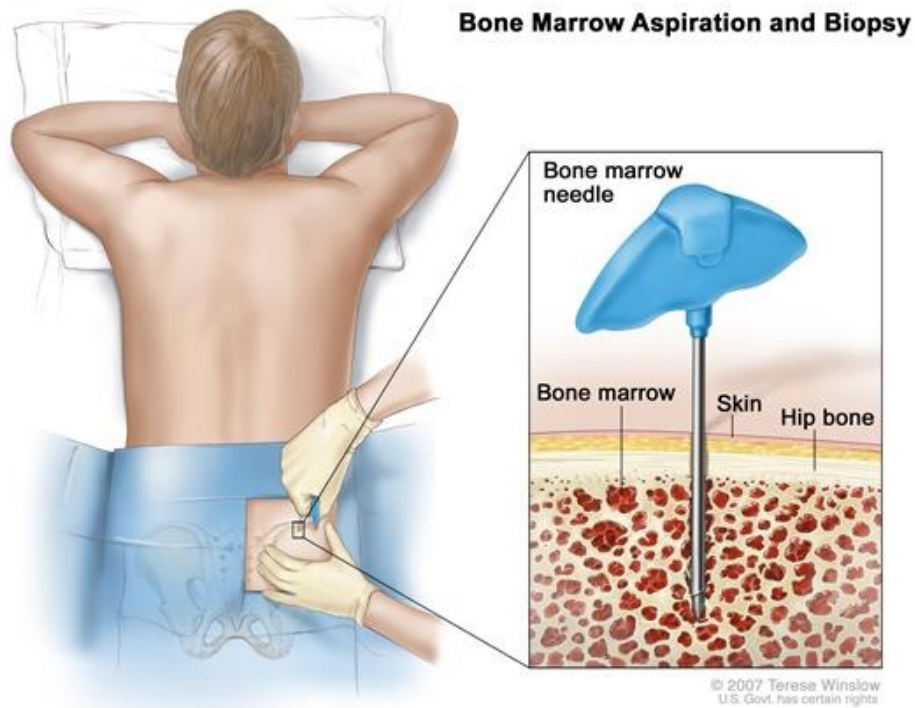


Spaciously molecular heterogeneity

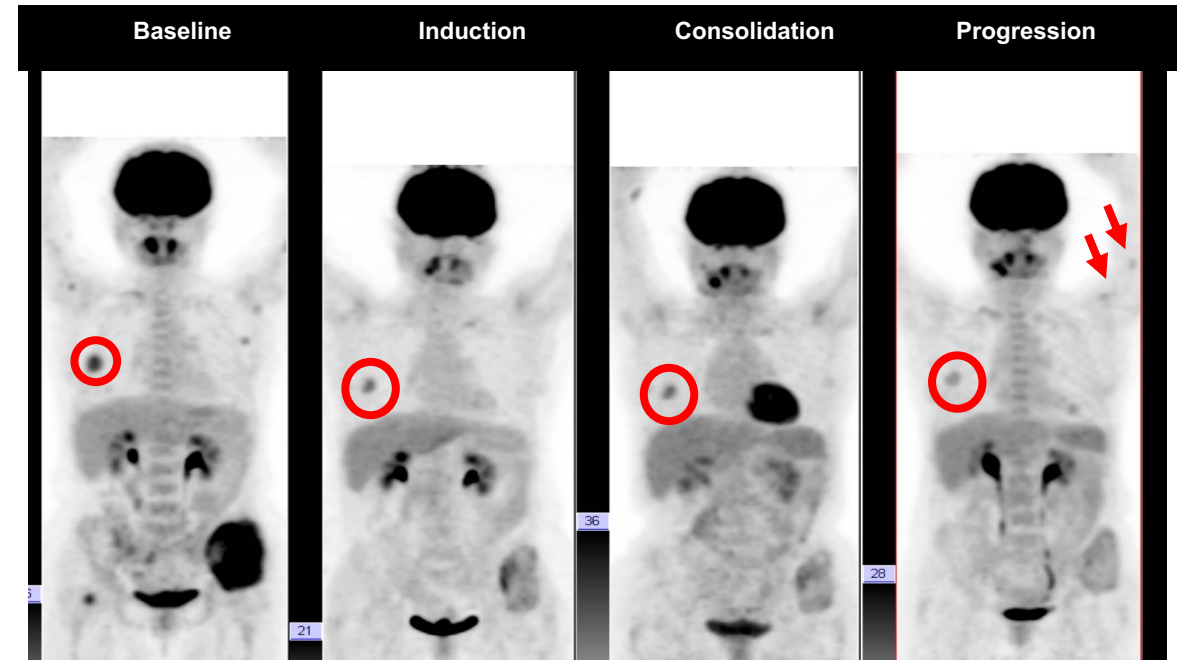


Limitations of MRD assessment in MM

BM aspiration is invasive and expensive and frequent sampling is impractical



Presence of extramedullary disease



Peripheral blood as alternative sample for MRD analysis in patients with MM

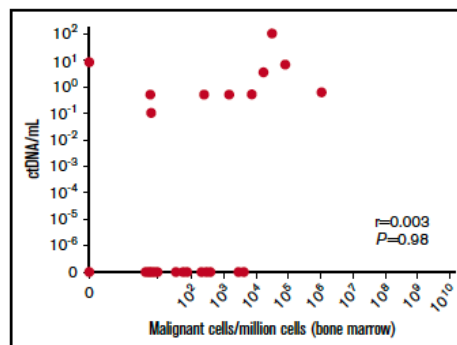
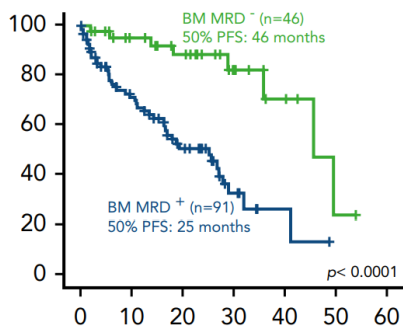
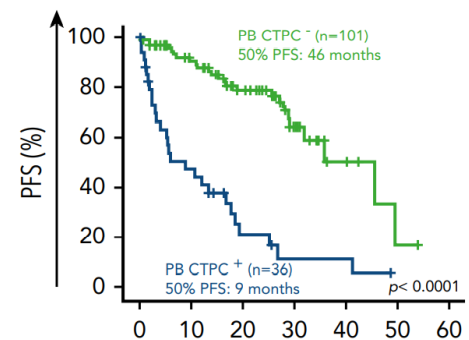
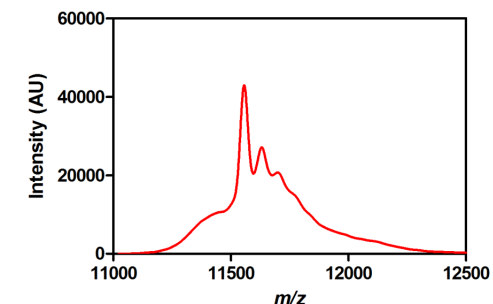
Flow cytometry



NGS



Mass spectrometry



Peripheral Blood Based MRD Approaches

cfDNA

CTPC

M-protein

Peripheral Blood Based MRD Approaches

cfDNA

CTPC

M-protein

MRD status in PB vs BM

CR/sCR MM cases (n=57)

	PB MRD			
BM MRD	Negative (-)	Positive (+)	Subtotal	<i>p</i> -value
Negative (-)	29/57 (51%)	0/57 (0%)	29/57 (51%)	
Positive (+)	19/57 (33%)	9/57 (16%)	28/57 (49%)	0.001
Subtotal	48/57 (84%)	9/57 (16%)	57/57 (100%)	

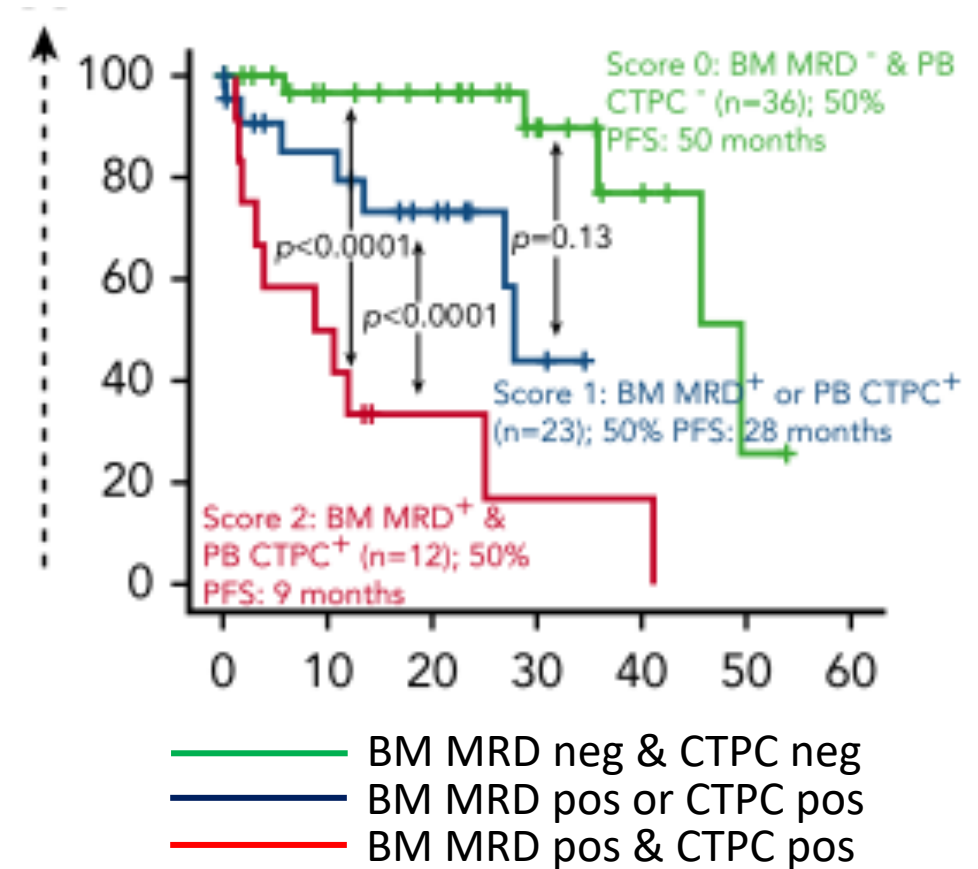
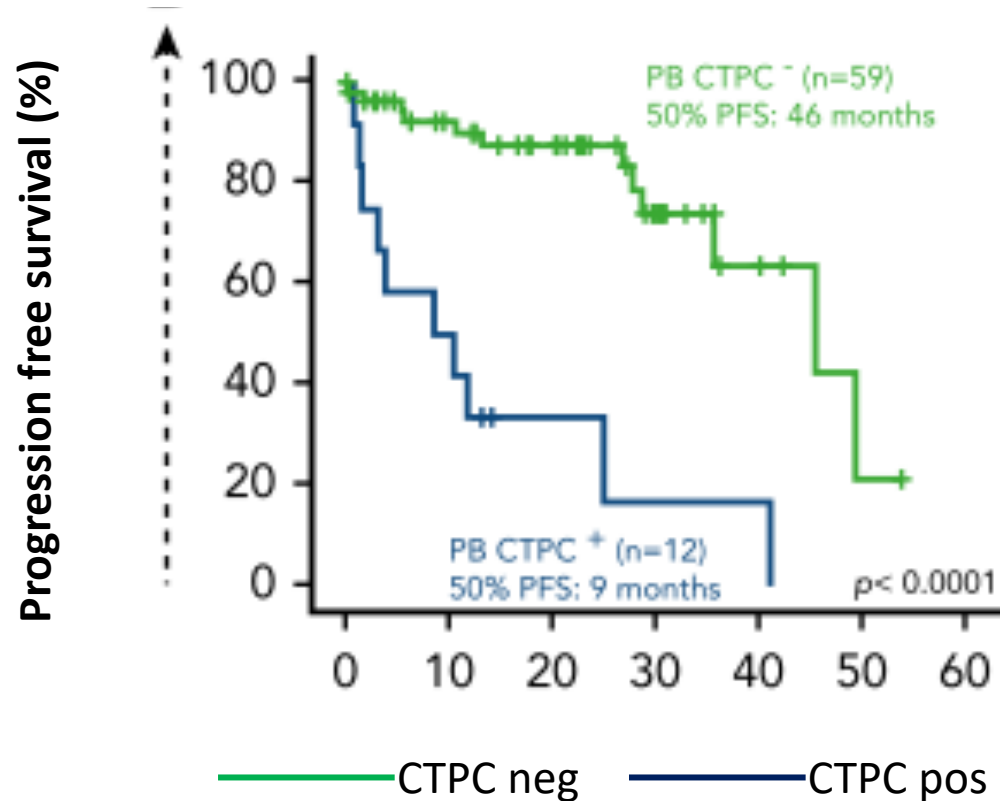
Relative and absolute cPC counts in PB and BM samples

Group	N. of cPC/mL of PB	% of cPC in PB	% of cPC in BM	<i>p</i> -value (% MRD in PB vs BM)
< CR (n=19)	307 cPC/mL (3-18,352)	0.007% (<0.0001%-0.6%)	0.3% (0.0005%-14.3%)	<0.0001
≥CR (n=9)	20 cPC/mL (5-457)	0.0002% (0.0001%-0.007%)	0.07% (0.0008%-1.6%)	0.008
All (n=28)	86 cPC/mL (3-18,352)	0.002% (<0.0001%-0.6%)	0.2% (0.0005%-14.3%)	<0.0001

<CR including VGPR, PR, SD and PD cases, ≥CR including sCR and CR cases

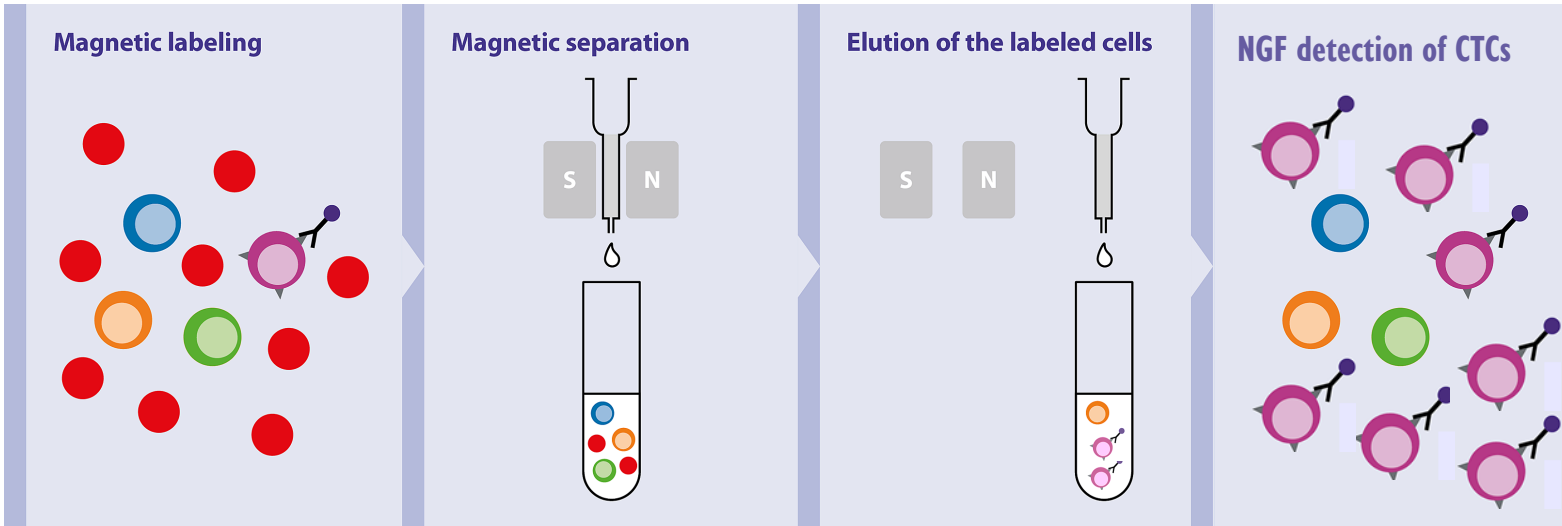
Prognostic impact of PB and BM MRD status by NGF

CR and sCR MM patients



BloodFlow

Immunomagnetic enrichment using MACS® MicroBeads prior NGF

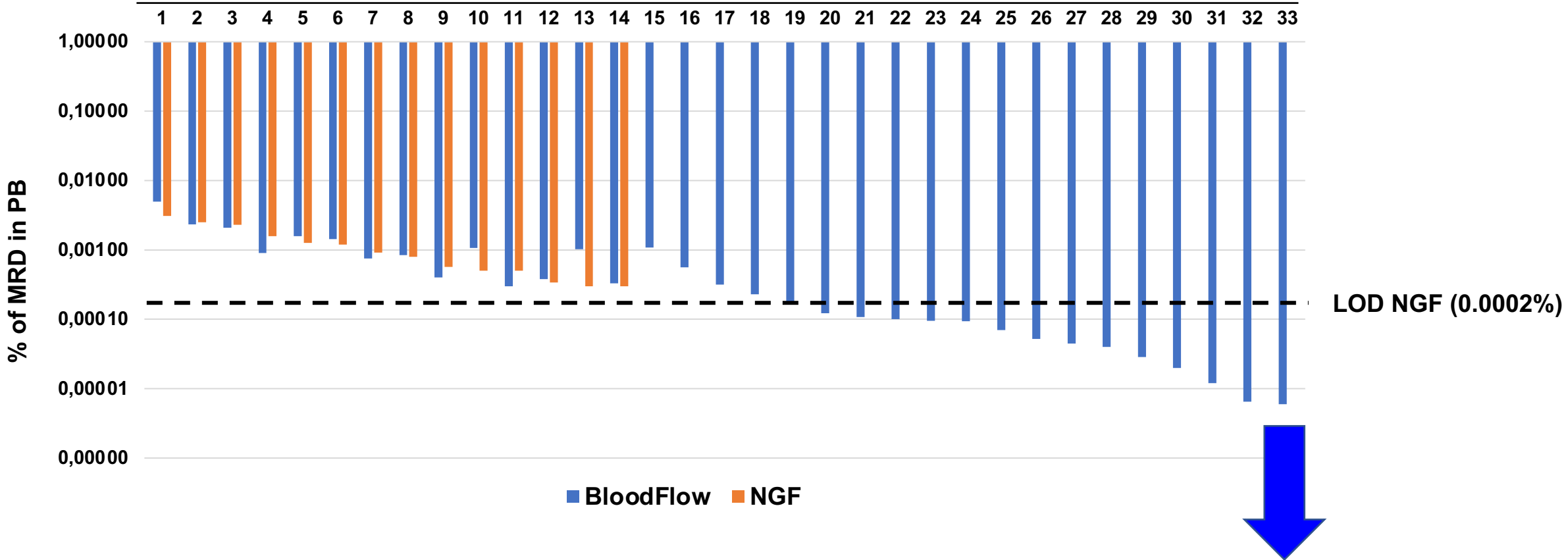


- A minimum sensitivity of 10^{-7} requires analyzing $\geq 2 \times 10^8$ cells (~50mL of PB)
- Large (~50mL) PB volumes were magnetically labeled and processed via MACS® columns, and ~100 μ L aliquots enriched with circulating PC were analyzed using EuroFlow NGF

Performance of BloodFlow vs NGF in PB (n = 353 samples)

BloodFlow detected MRD in 33/353 (9%); 19/33 (58%) were negative by NGF

Patients with positive MRD in PB using BloodFlow

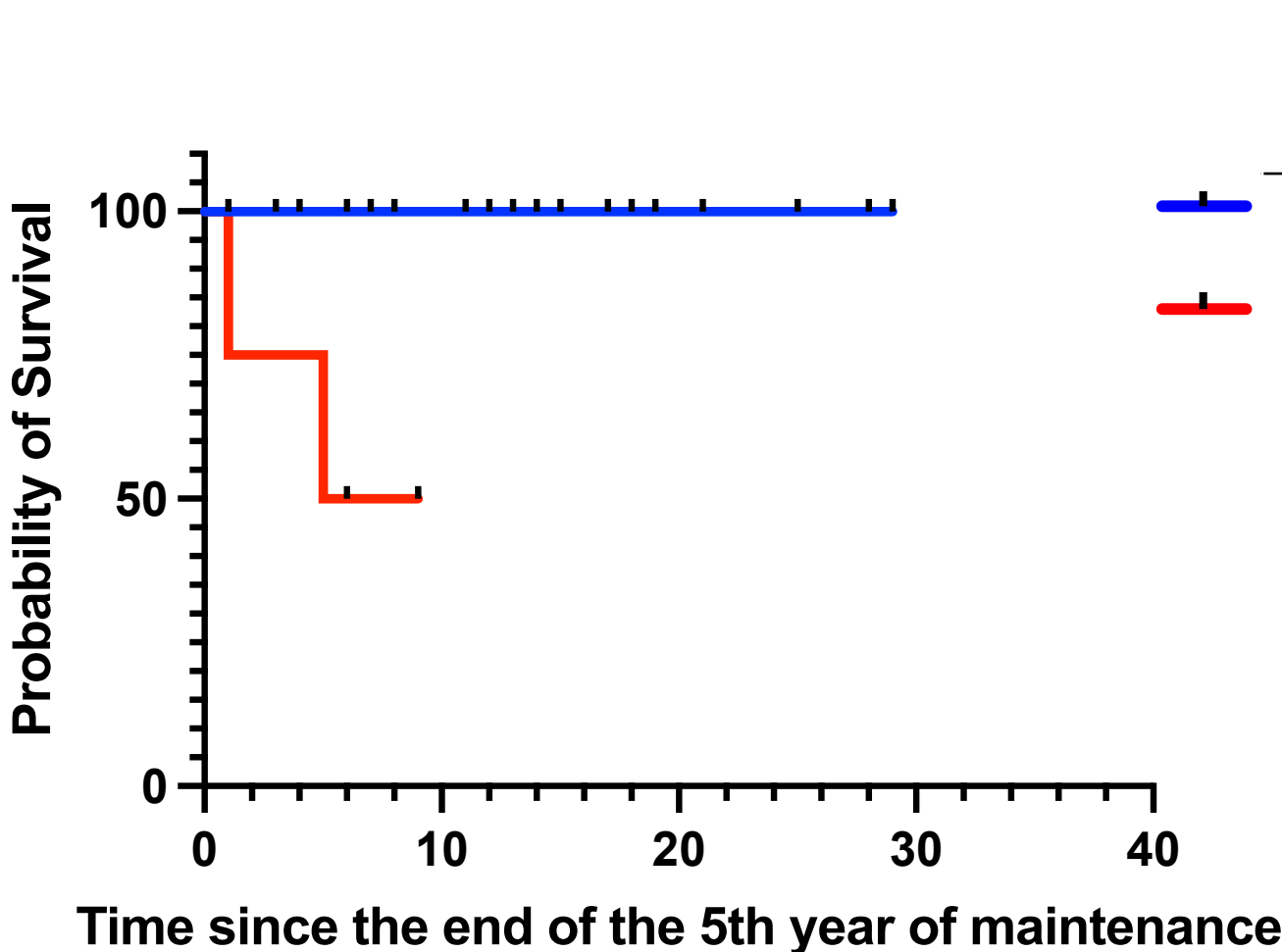


The lowest MRD level was 6x10⁻⁸

PB, peripheral blood; NGF, next-generation flow

Prognostic value of MRD assessment in PB using BloodFlow

GEM2014MAIN trial (n = 33)



MRD	N.	Median PFS	p value
Negative	29	NR	$p < .0001$
Positive	4	7 mo	

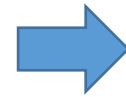
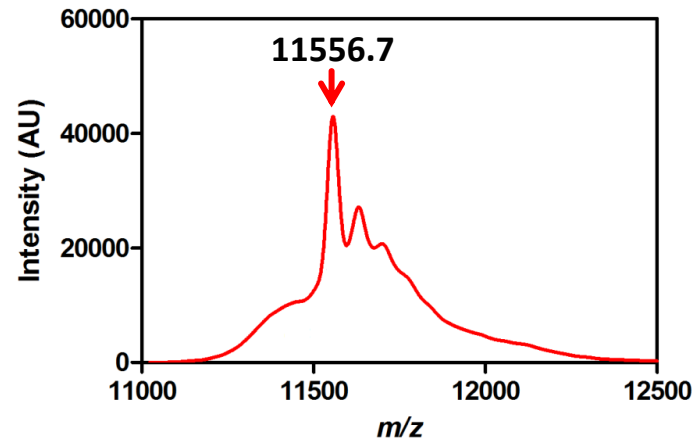
Peripheral Blood Based MRD Approaches

cfDNA

CTPC

M-protein

The innovative approach: identify M-protein molecular mass with high precision and accuracy

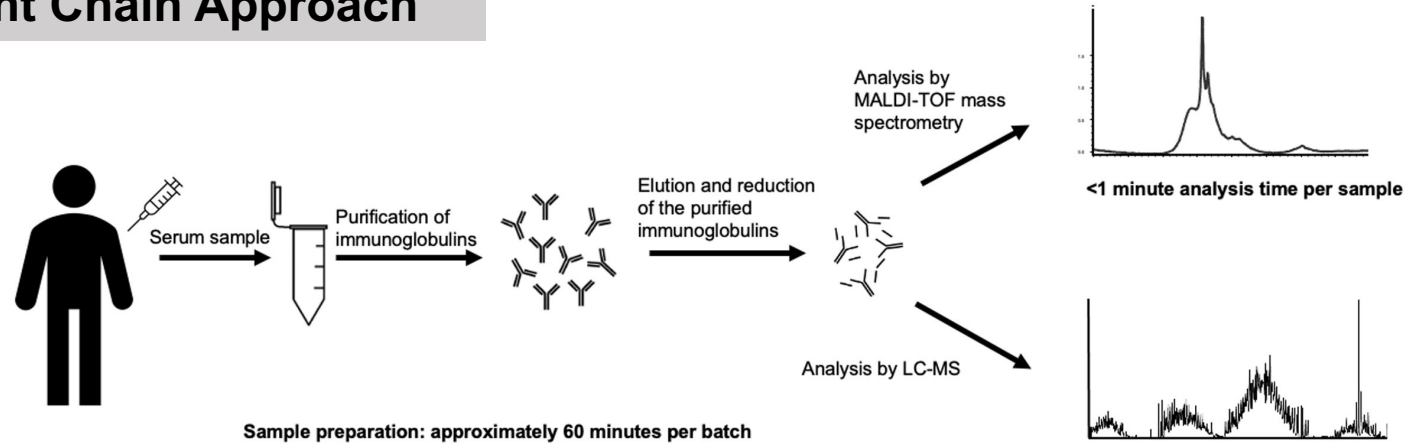


**Molecular mass defines clonality
/ intensity defines abundance**

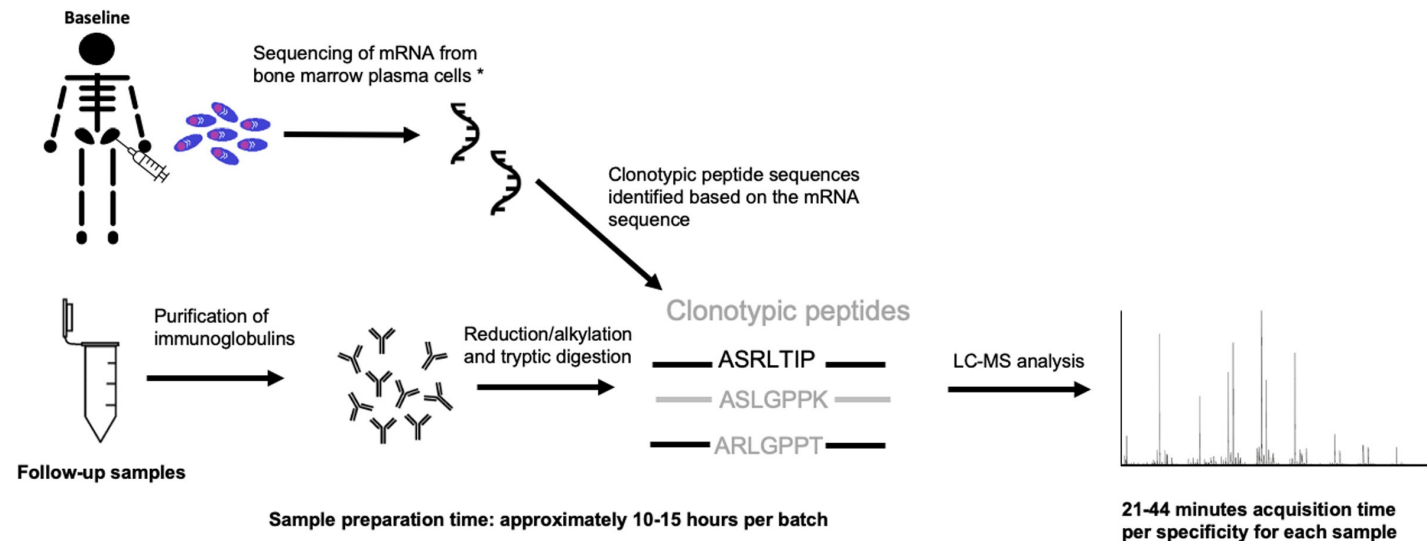
Each Ig has a specific amino acid sequence and therefore a specific molecular mass, constant over time, that can serve as a surrogate marker for the presence of clonal PC

MS methods for the identification on M-proteins in serum

Intact Light Chain Approach



Clonotypic Peptide Approach

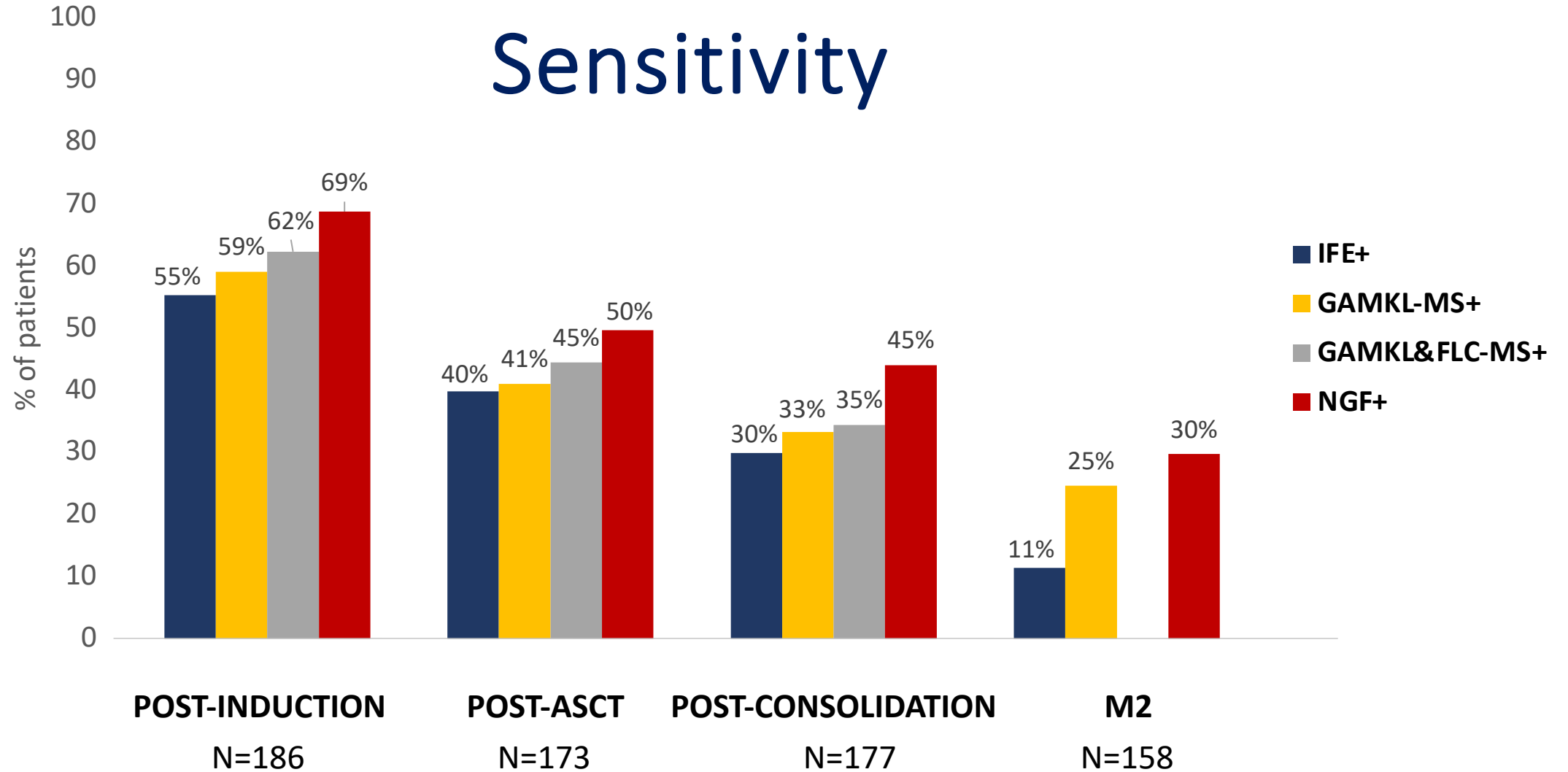


Increasing applicability
to routine clinical practice

Increasing Complexity, Sensitivity,
Time and Cost

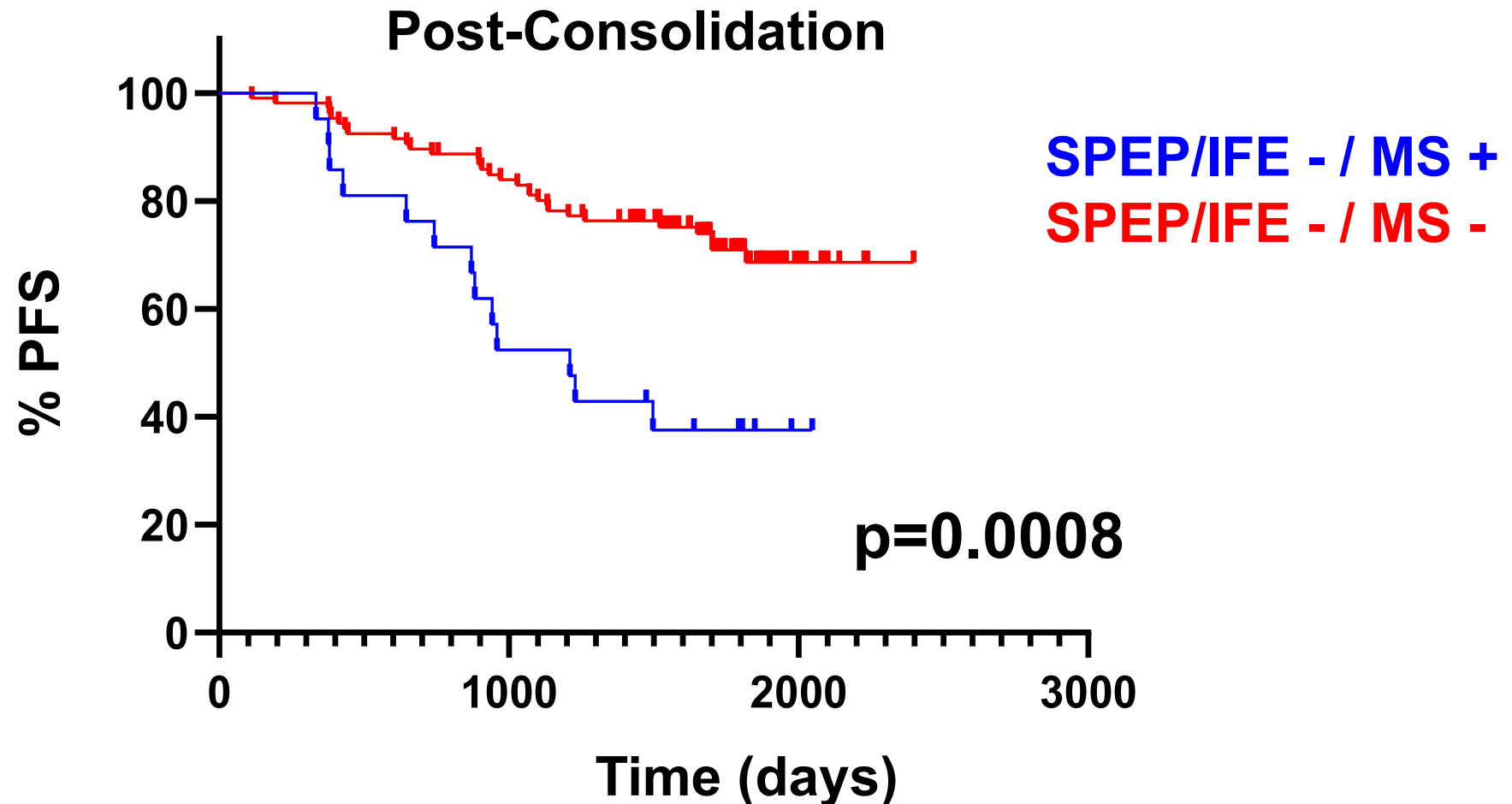
SPEP/IF vs BM-NGF vs PB-MS

Sensitivity



PFS according to the MS status in patients in \geq CR

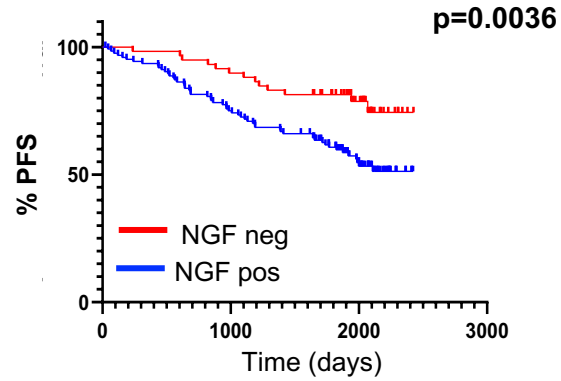
Among the 127 patients in CR post-consolidation, MS identified the presence of the MP in 21 of them (16.5%)



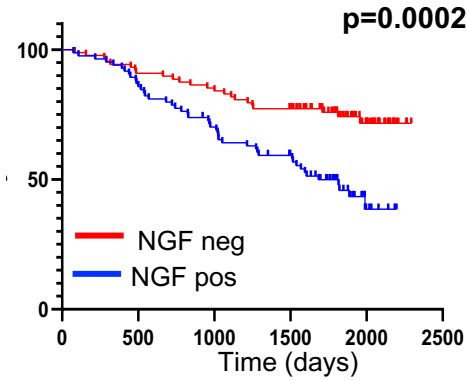
BM-NGF vs PB-MS: prognostic value

BM-NGF

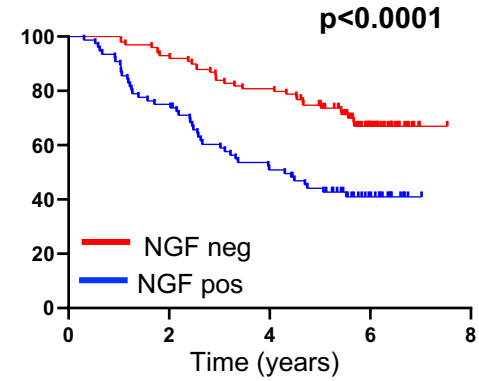
POST-INDUCTION



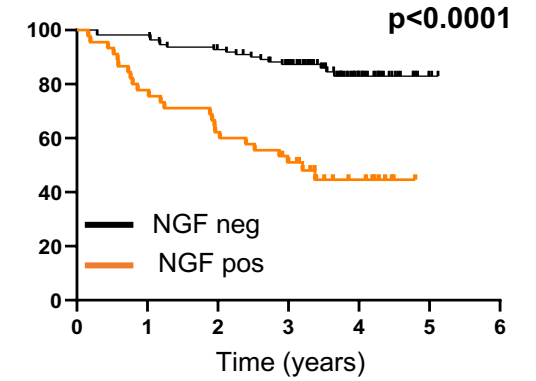
POST-ASCT



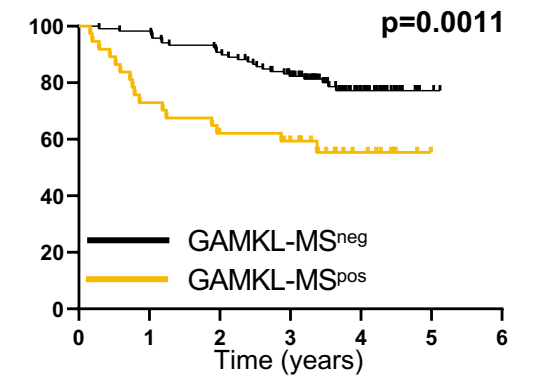
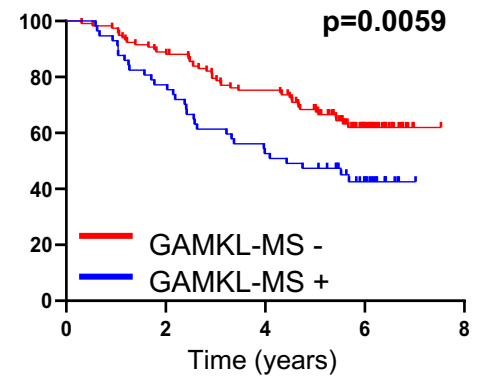
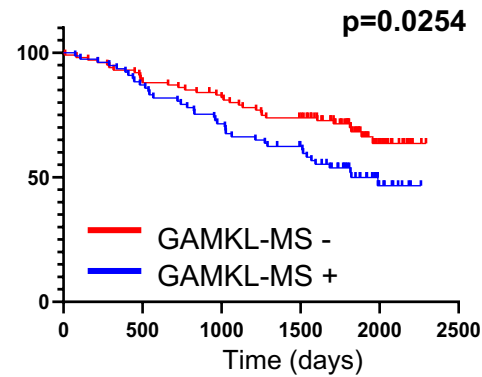
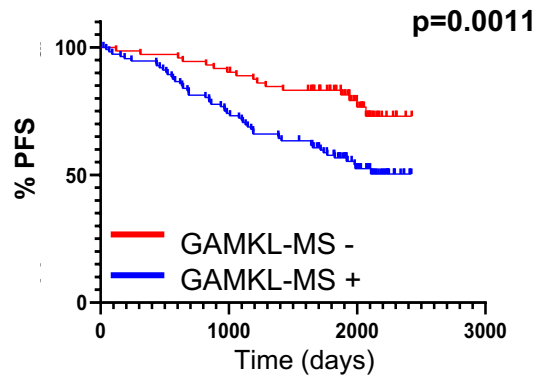
POST-CONSOLIDATION



M2



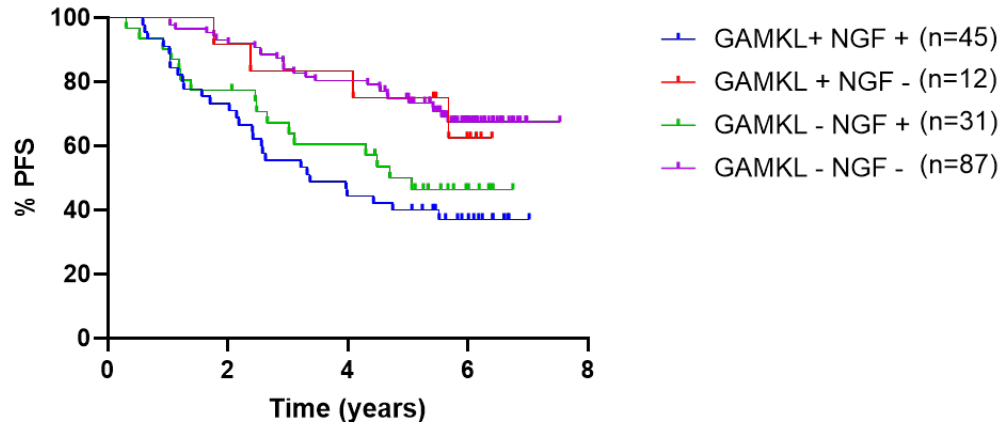
QIP-MS



BM-NGF & PB-MS

Combined results

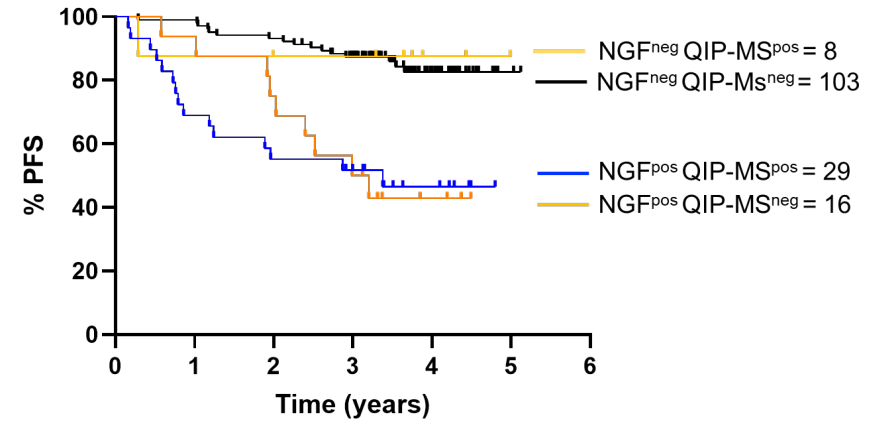
POST-CONSOLIDATION



Fisher's exact test

p-value	<0.0001
Positive Predictive Value	79% (95% CI, 0.67–0.88)
Negative Predictive Value	74% (95% CI, 0.65–0.81)

M2



Fisher's exact test

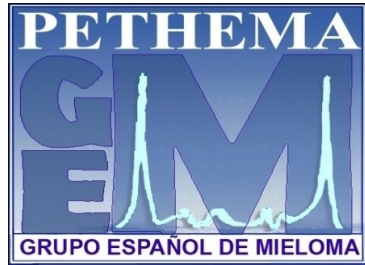
p-value	<0.0001
Positive Predictive Value	78% (95% CI, 0.63–0.89)
Negative Predictive Value	87% (95% CI, 0.79–0.92)

Conclusions

- An **extraordinary therapeutic progress** has been made in the last 20 years in the MM field and accordingly, **new techniques** to assess the presence of residual disease have been introduced
- Among them, the results obtained with **NGS and NGF** have broadly proven its value as a **prognostic factor**
- The use of **whole-body imaging techniques** to evaluate treatment response is **crucial** given the potential presence of patchy infiltration of the marrow and/or extramedullary disease

- However, to use MRD to make clinical decisions, we need the **results of randomized clinical trials** segregating patients to different treatment arms based on the MRD results
- New approaches to assess **MRD in PB** have shown very promising results, but more data are needed to define its definite role in MM

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**Thank you very much
for your attention!**

Role of MRD in MM: different techniques for a crucial biomarker

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