

2020



Progetto Ematologia Romagna

Mieloma Multiplo: lunghi sopravvissuti o guariti

Paola Tacchetti

Azienda Ospedaliero-Universitaria di Bologna; Istituto di Ematologia "Seràgnoli";
Dipartimento di Medicina Specialistica, Diagnostica e Sperimentale; Università degli Studi di Bologna

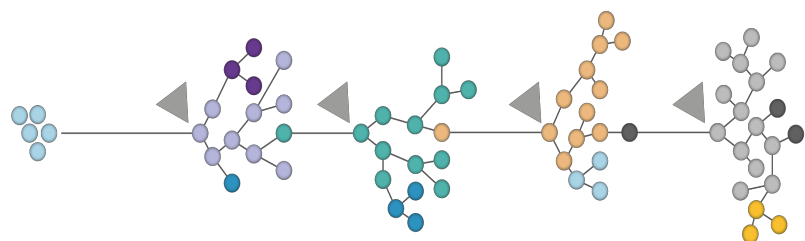
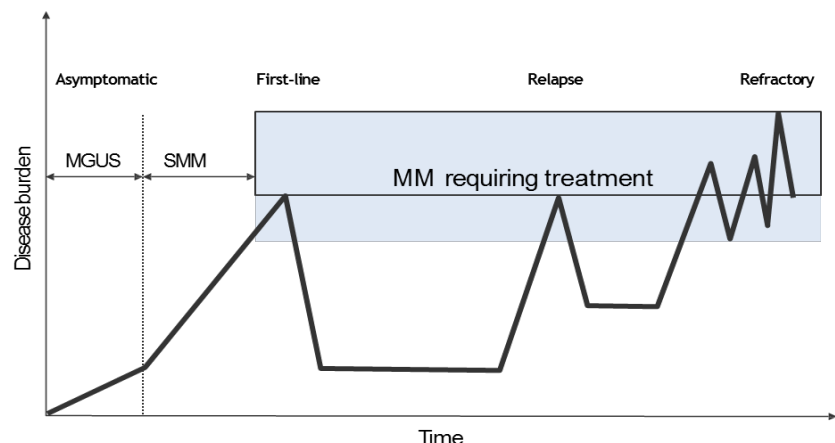


2020

Disclosures of Paola Tacchetti

Company name	Research support	Employee	Consultant	Stockholder	Speakers bureau	Advisory board	Other
Bristol-Myers Squibb/Celgene						x	honoraria
Janssen						x	honoraria
Amgen						x	honoraria
Takeda						x	honoraria
Abbvie							honoraria
Oncopeptides							honoraria

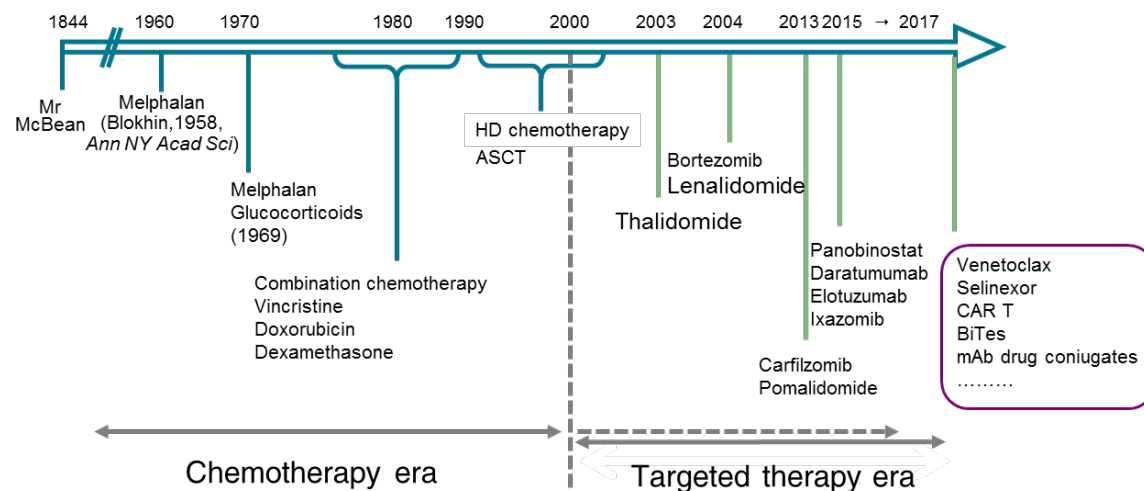
Treating myeloma – The changing landscape



Tumor cell diversity

Genetic lesions

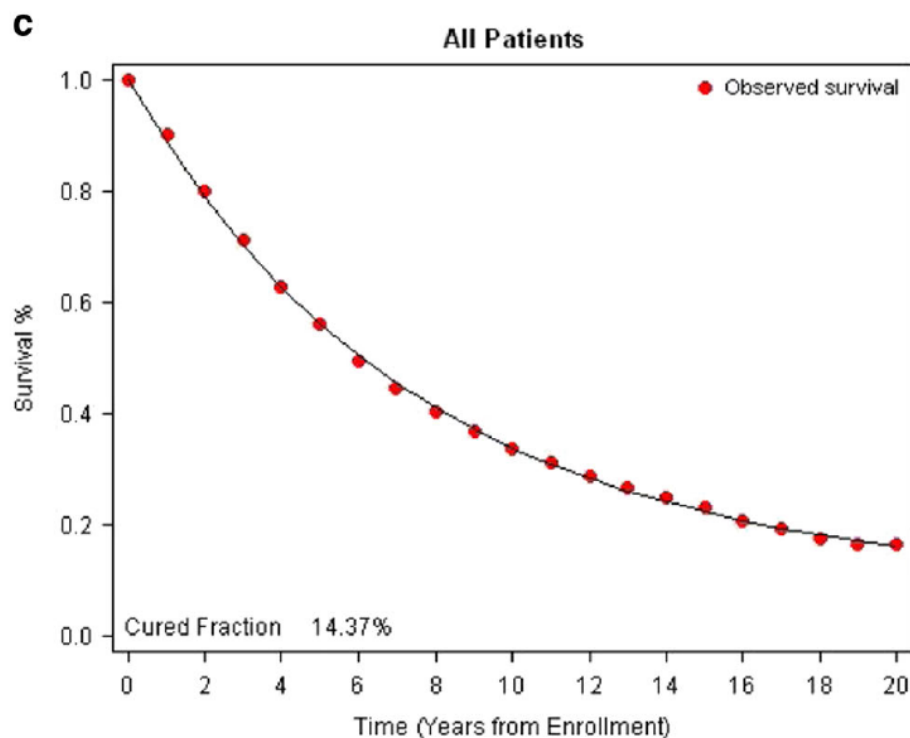
Image adapted from Morgan G et al. Nat Cancer Revs 2012;12:335-348



Is MM “curable”?

Clinical predictors of long-term survival in NDTE MM

An IMWG Research Project



7291 pts enrolled in clinical trials involving ASCT

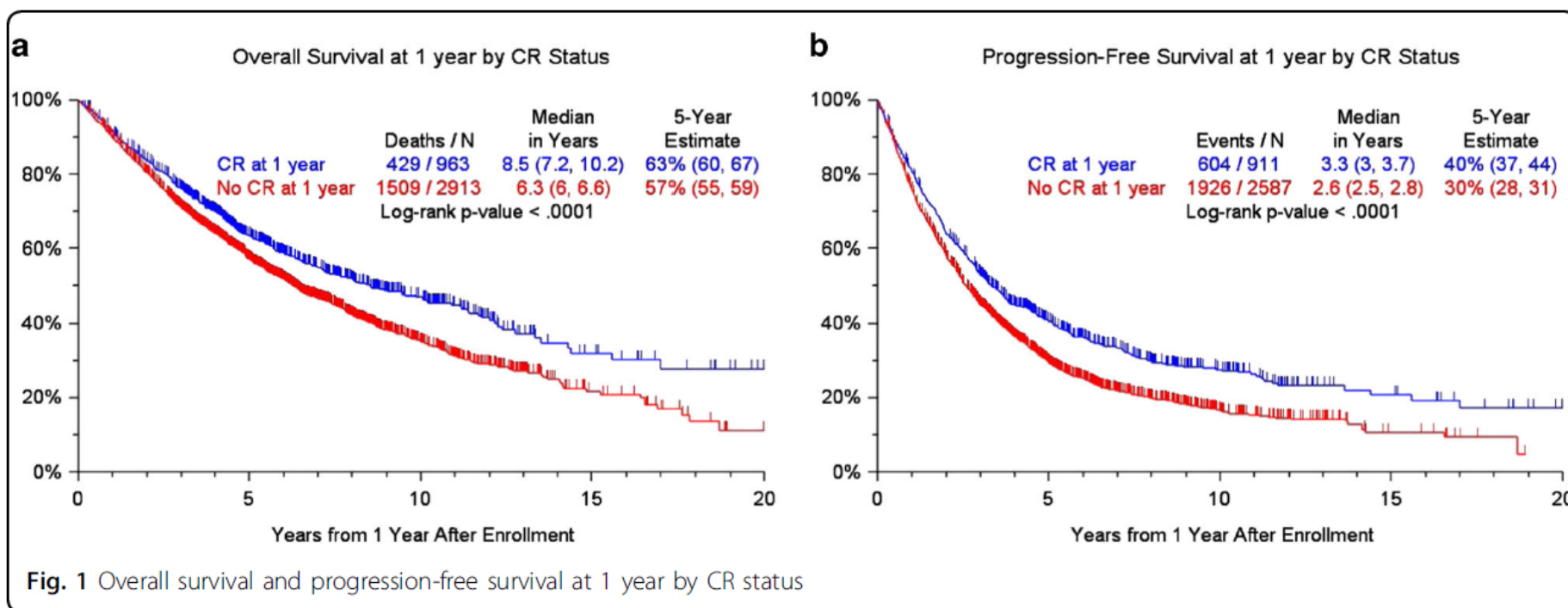
The **statistical cure fraction** for the whole group appears to be **14.3%**, which signifies the overall proportion of MM pts in this cohort who were able to achieve or exceed expected survival compared to matched general population.

Usmani SZ, et al. Blood Cancer Journal 2018;8:123



2020

IMWG Research Project: OS and PFS by CR status



Usmani SZ, et al. Blood Cancer Journal 2018;8:123

IMWG Research Project: Logistic regression, 10-yr survival vs 2-yr death

Variable	Factors differentiating 10-year survival vs. 2-year death				
	N	Survival less than 2 years	Survival more than 10 years	OR (95% CI)	P-value
Multivariate (stratified by country)					
Age at registration ≥ 65 yr	1230	98/144 (68%)	525/1086 (48%)	1.87 (1.26, 2.79)	0.002
IgA	1230	180/298 (60%)	443/932 (48%)	1.53 (1.15, 2.04)	0.004
Albumin < 3.5 g/dL	1230	294/474 (62%)	329/756 (44%)	1.36 (1.04, 1.78)	0.023
B2M ≥ 3.5 mg/L	1230	376/569 (66%)	247/661 (37%)	1.86 (1.41, 2.45)	< 0.001
Creatinine ≥ 2 mg/dL	1230	130/174 (75%)	493/1056 (47%)	1.77 (1.18, 2.65)	0.005
HGB < 10 g/dL	1230	288/429 (67%)	335/801 (42%)	1.55 (1.16, 2.06)	0.003
Platelet Count $< 150 \times 10^9/L$	1230	163/218 (75%)	460/1012 (45%)	2.26 (1.59, 3.22)	< 0.001

It is important to note that **over 90% of the pts** in the dataset were from the **pre-novel therapy induction era** and ~10% did received thalidomide as part of their upfront therapy (Total Therapy 2 thalidomide arm, GMMG-HD3 thalidomide arm and BO2002).

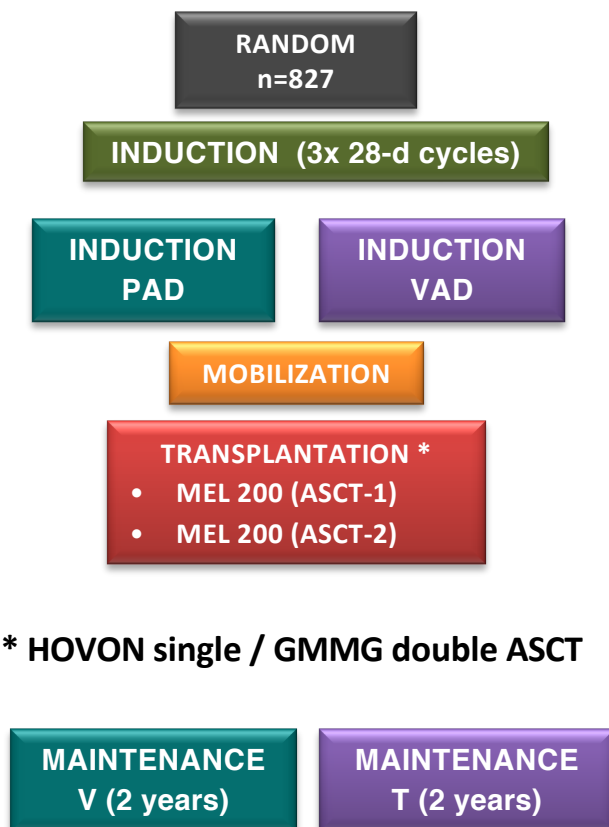
Usmani SZ, et al. Blood Cancer Journal 2018;8:123



2020

HOVON65MM/GMMG-HD41

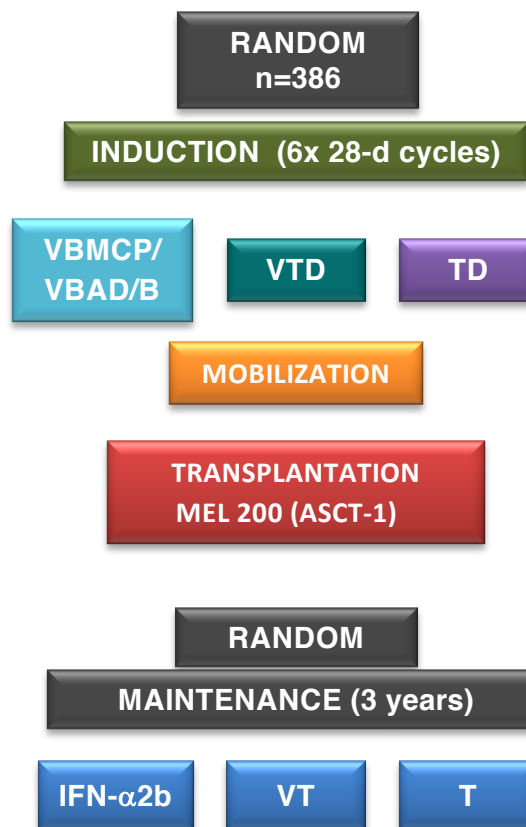
Sonneveld et al. JCO 2012;30:2946-55



* HOVON single / GMMG double ASCT

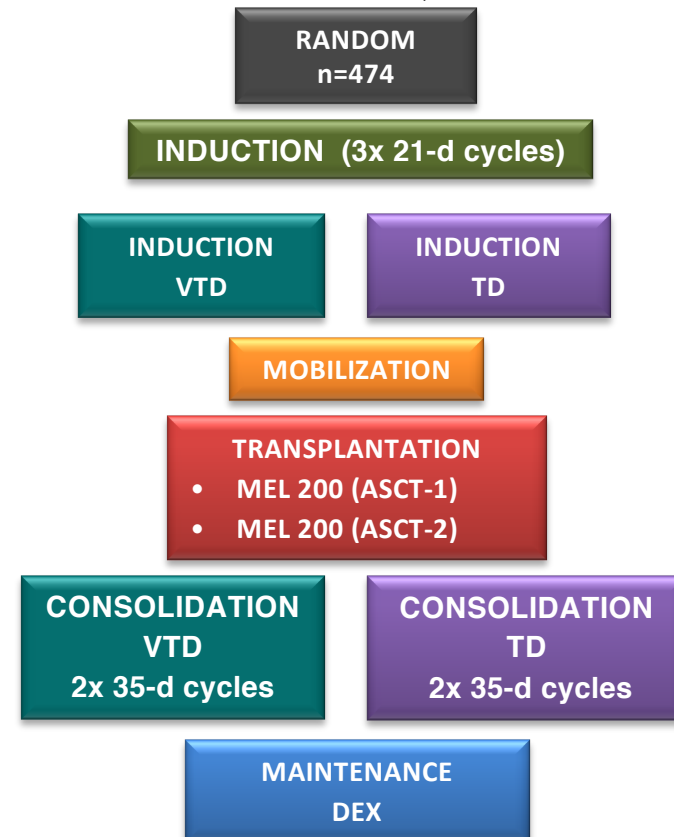
PETHEMA/GEM

Rosinol et al. Blood 2012;120:1589-96



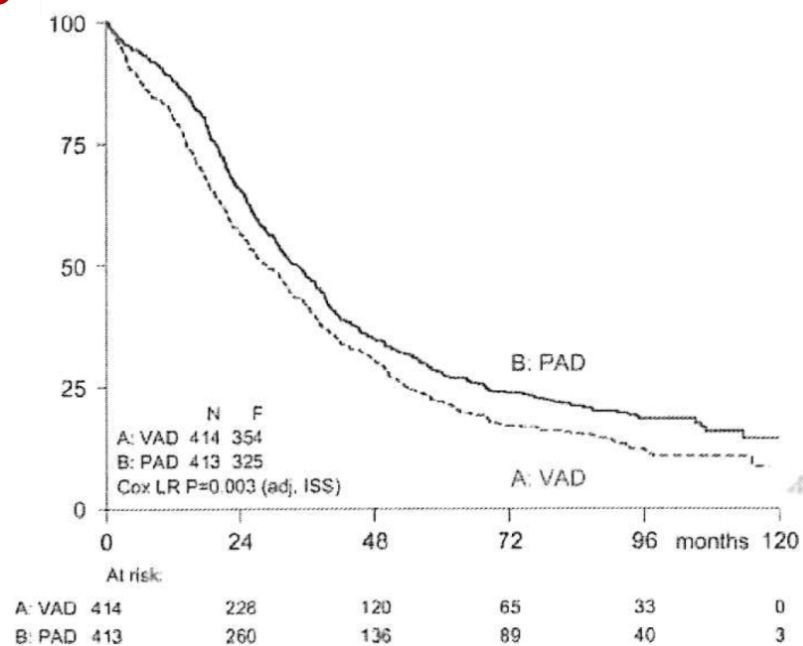
GIMEMA MMY-3006

Cavo M et al. Lancet 2010;376:2075-85

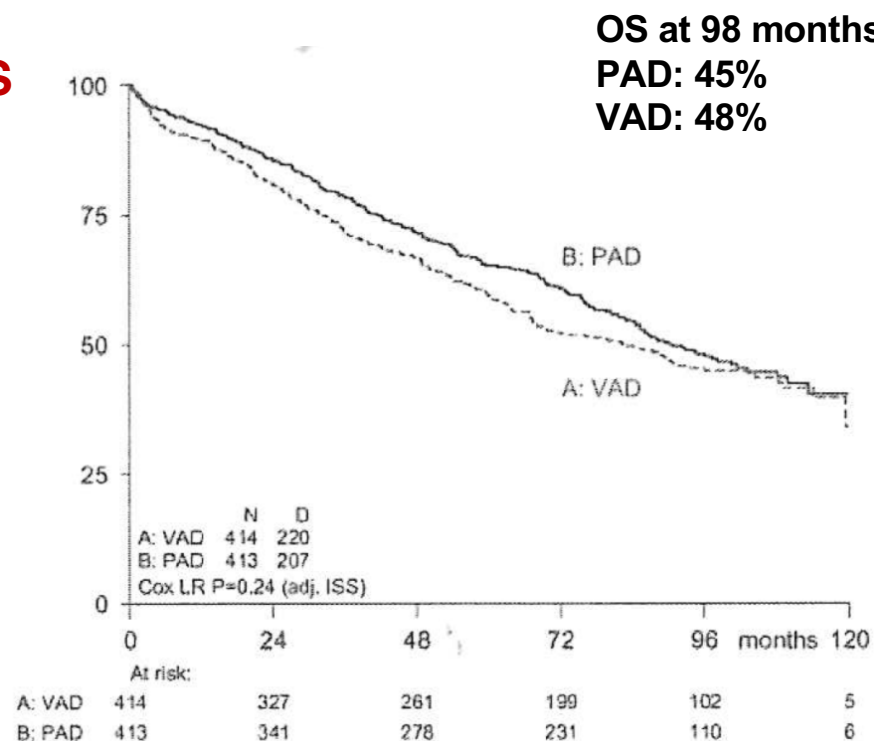


HOVON-65/GMMG-HD4 study: long-term analysis (98-mo follow-up)

PFS



OS

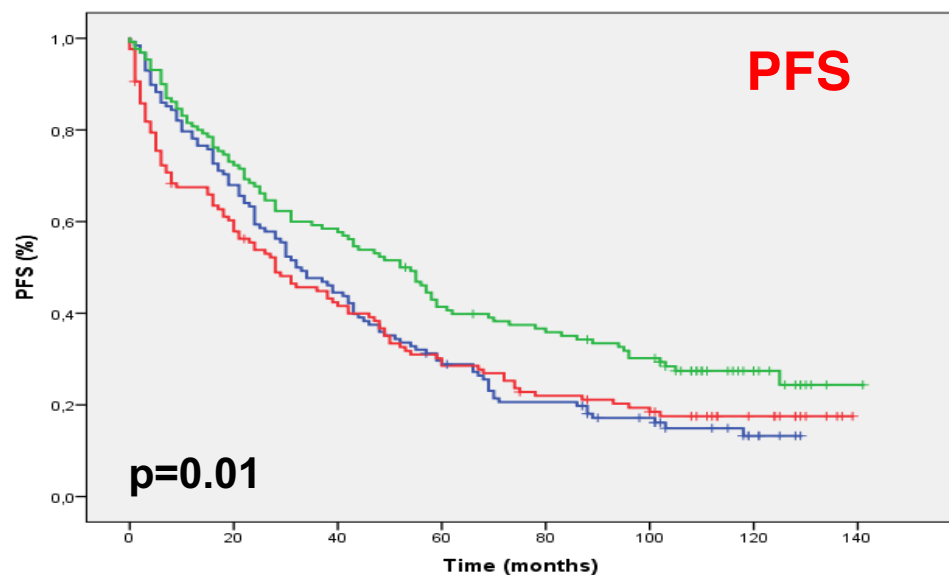


Goldschmidt H. et al., Leukemia 2018;32:383-390

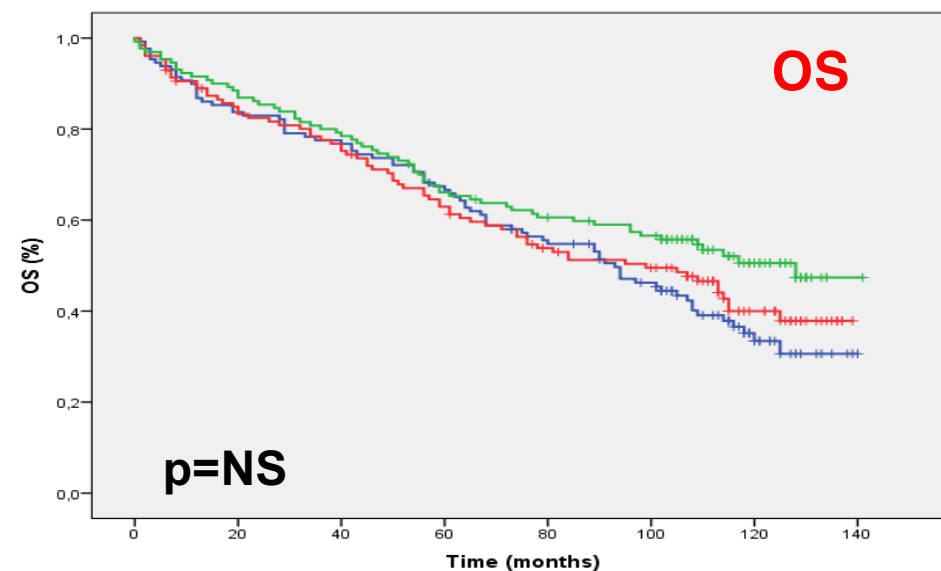


2020

PETHEMA/GEM study: long-term analysis (10-yr follow-up)



	Median PFS (mos)	10-yr PFS (%)
QT+V	32	13
TD	28	17
VTD	52	24



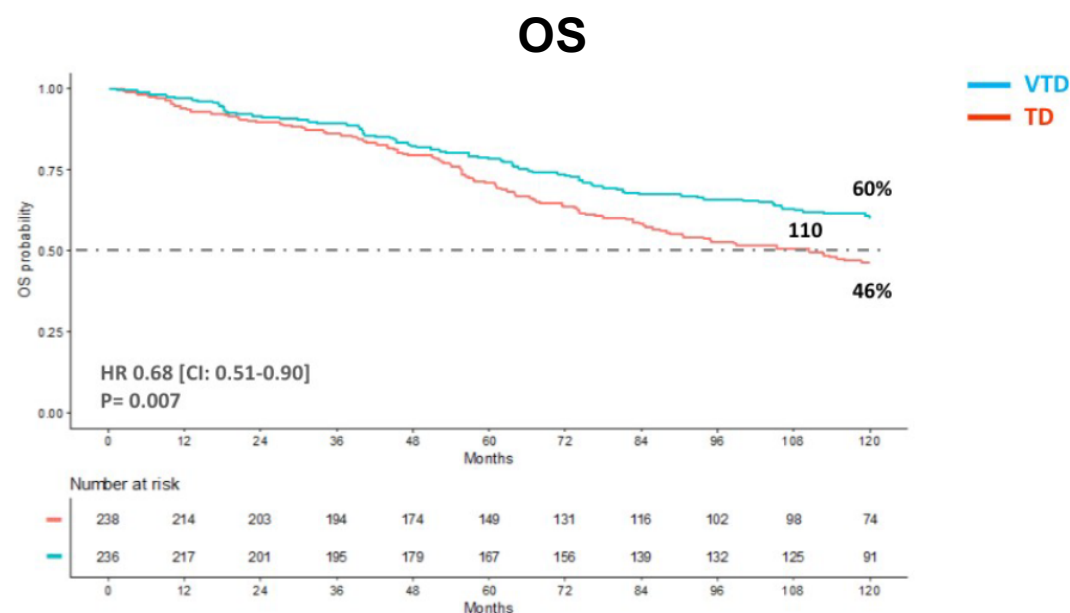
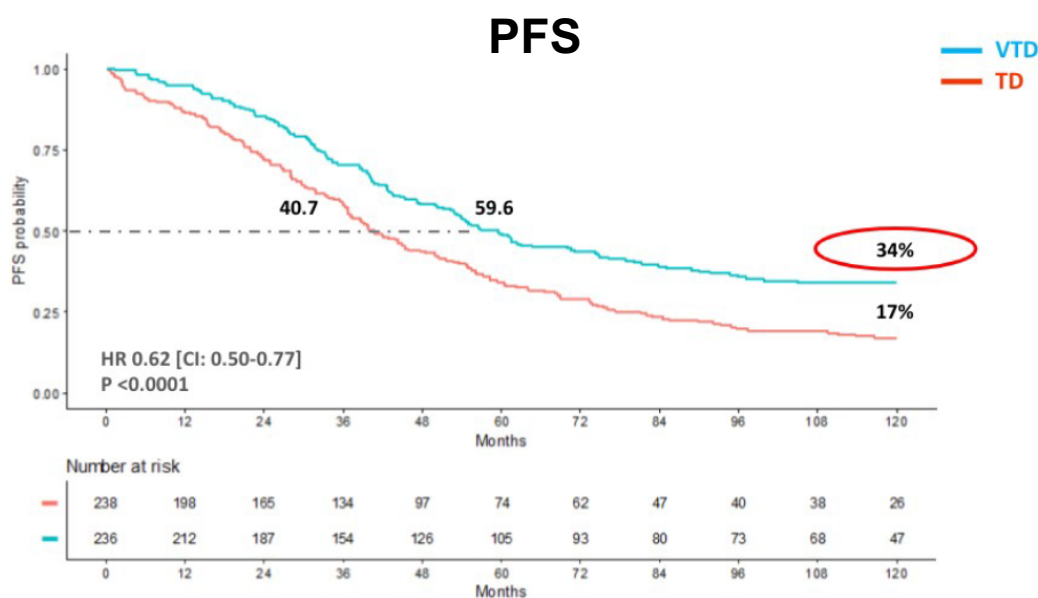
	Median OS (mos)	10-yr OS (%)
QT+V	93	33
TD	99	40
VTD	128	51

Rosinol L. et al., ASH 2018



2020

GIMEMA-MMY-3006 study: long-term analysis (10-yr follow-up)



HR, hazard ratio; PFS, progression-free survival; OS, overall survival; TD, thalidomide + dexamethasone; VTD, bortezomib + thalidomide + dexamethasone.

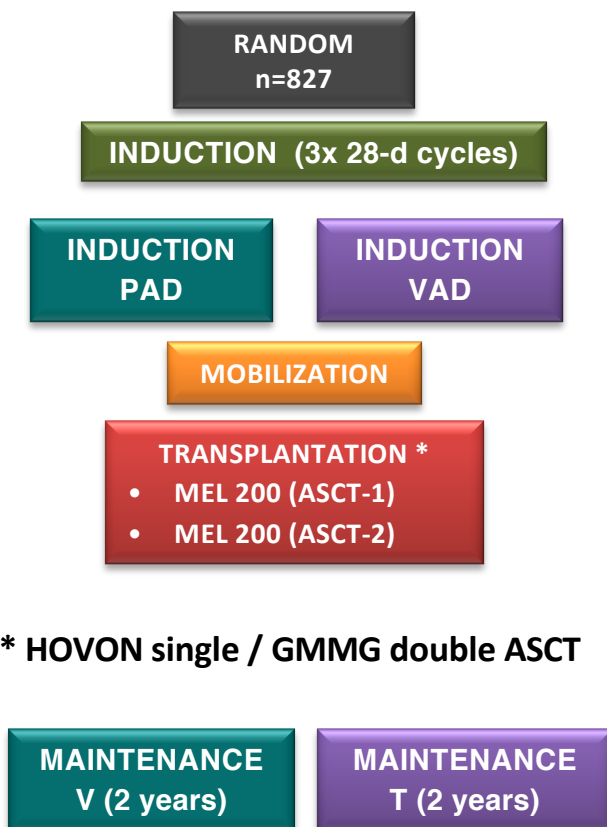
Tacchetti P. et al., *Lancet Haematology* 2020, accepted



2020

HOVON65MM/GMMG-HD41

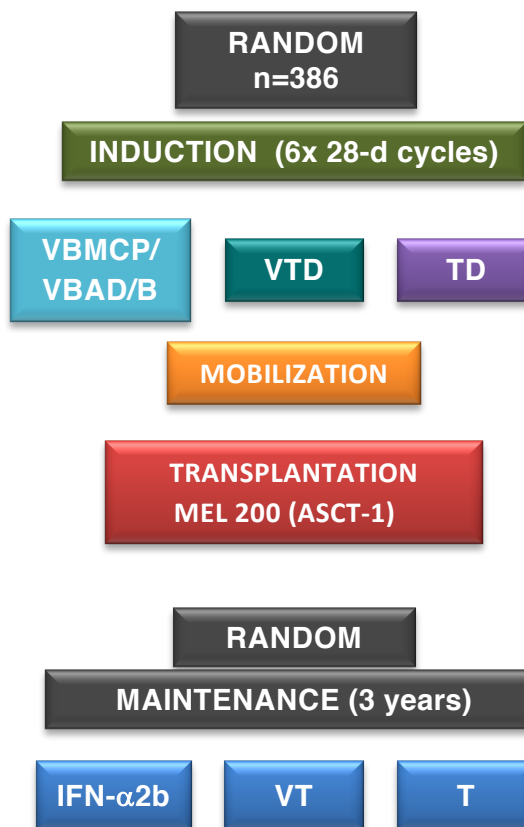
Sonneveld et al. JCO 2012;30:2946-55



* HOVON single / GMMG double ASCT

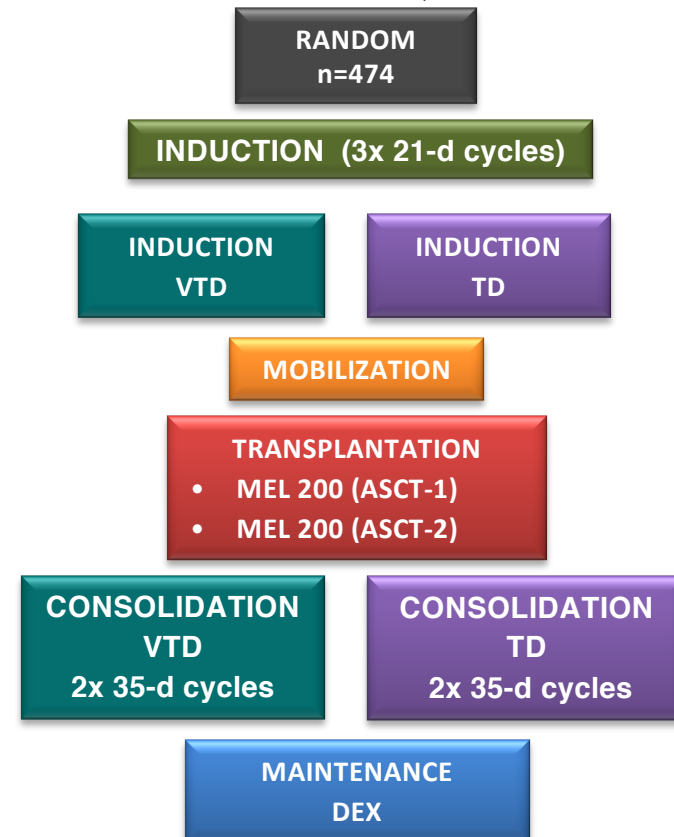
PETHEMA/GEM

Rosinol et al. Blood 2012;120:1589-96



GIMEMA MMY-3006

Cavo M et al. Lancet 2010;376:2075-85



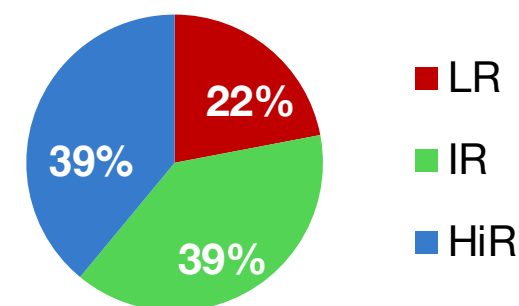
GIMEMA-MMY-3006 study: PROGNOSTIC SCORE

SPECIFIC MULTIVARIABLE REGRESSION ANALYSIS NOT INCLUDING THERAPY

Variables adversely affecting PFS	HR	95% CI	P-value
t(4;14) and/or del(17p) pos	1.857	1.452-2.375	<0.001
ISS stage II+III	1.384	1.099-2.743	0.006
Failure to achieve CR* (best response)	2.006	1.593-2.526	<0.001

*time-dependent variable

Risk group	Criteria
Low-risk (LR)	None of the three adverse variables
Intermediate-risk (IR)	1 adverse variable
High-risk (HiR)	2 or 3 adverse variables

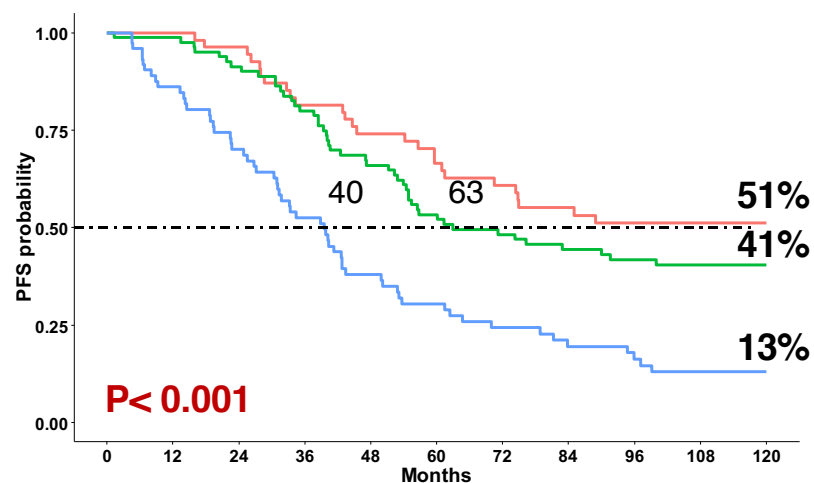


Tacchetti P. et al., Lancet Haematology 2020, accepted

GIMEMA-MMY-3006 study: PFS and OS by risk groups within the VTD arm

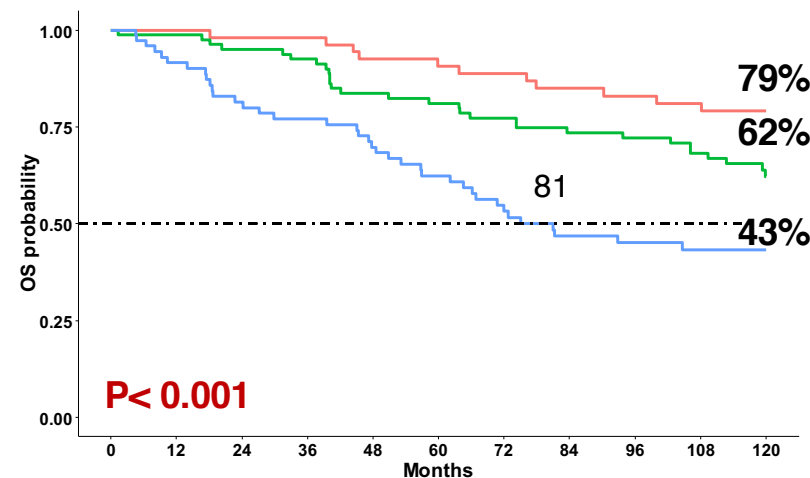
PFS

	HR (95% CI)	P
LR vs HiR	0.32 (0.20-0.51)	<0.001
IR vs HiR	0.44 (0.30-0.54)	<0.001



OS

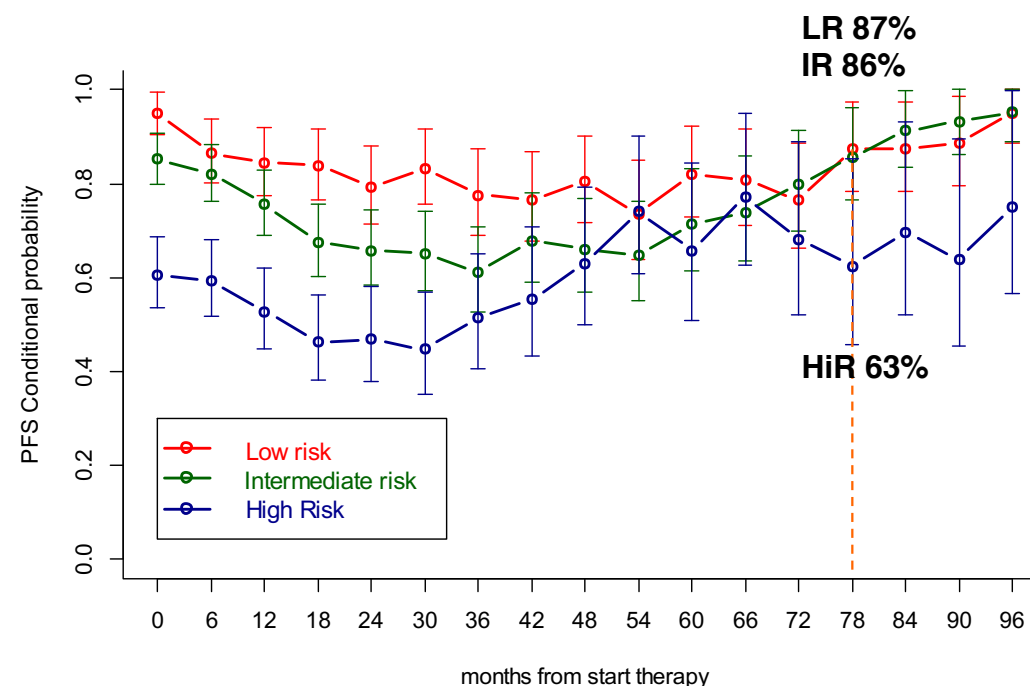
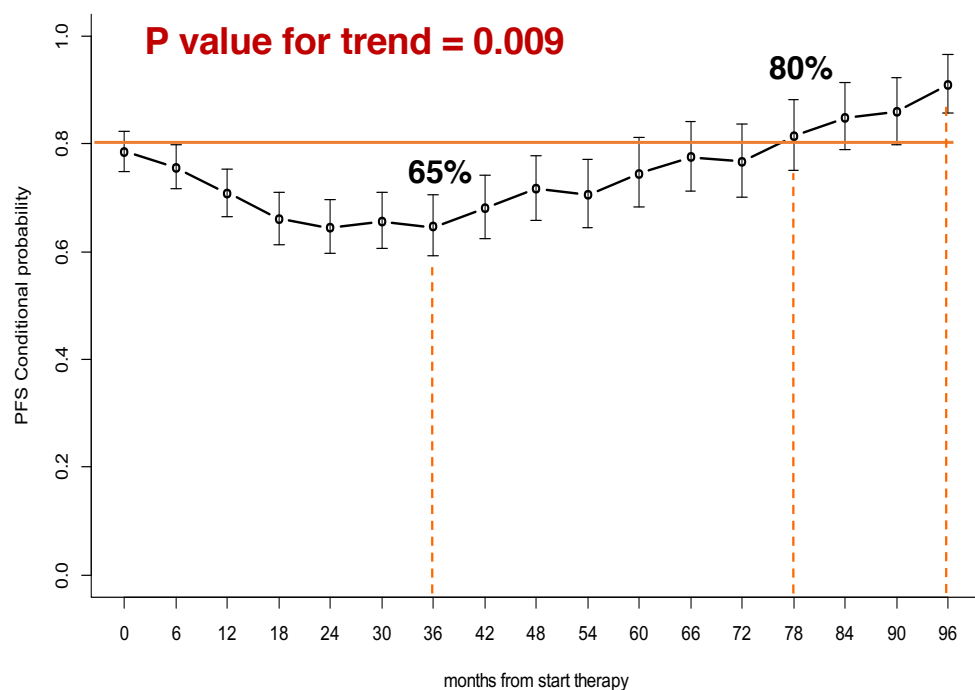
	HR (95% CI)	P
LR vs HiR	0.25 (0.13-0.50)	<0.001
IR vs HiR	0.50 (0.31-0.81)	<0.001



— LR
— IR
— HiR

Tacchetti P. et al., Lancet Haematology 2020, accepted

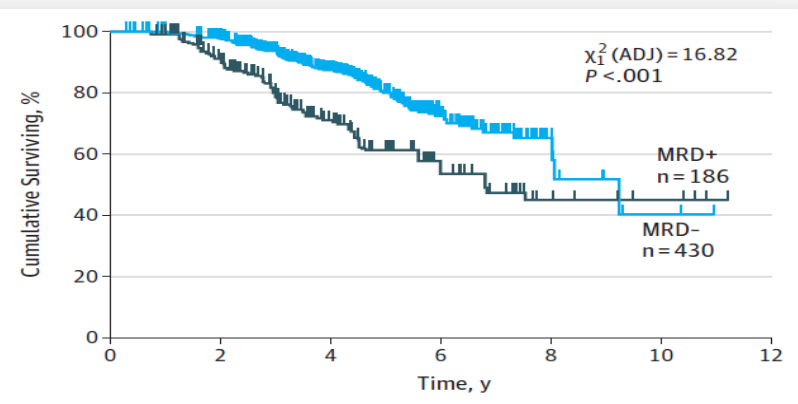
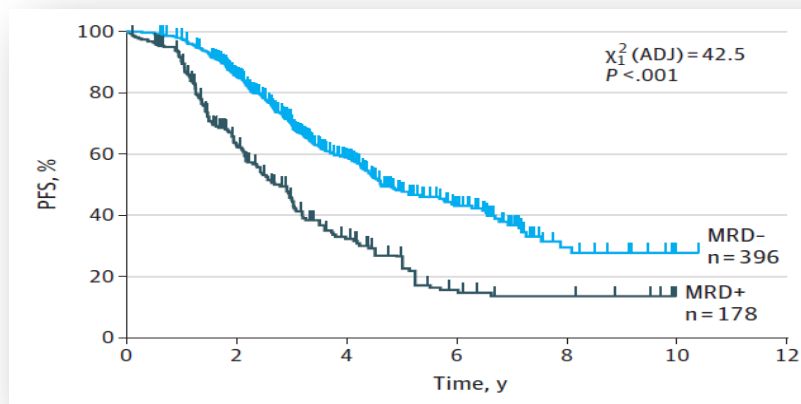
GIMEMA-MMY-3006 study: 2-YRS CONDITIONAL SURVIVAL ESTIMATE FOR PFS



Conditional survival estimate for PFS was calculated as the probability of surviving without progression a further 2 years given the years already survived

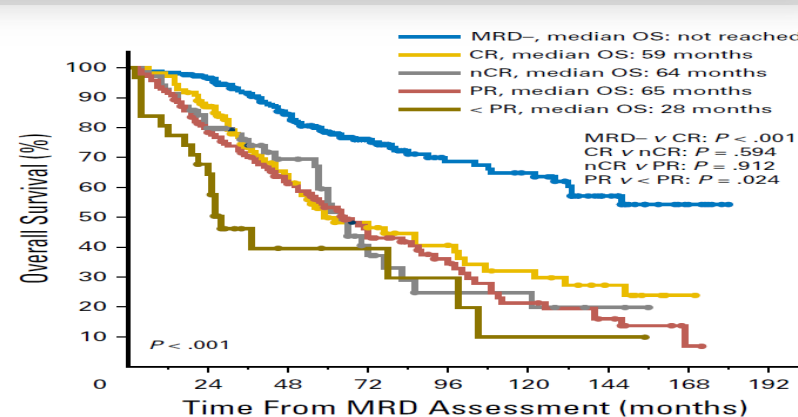
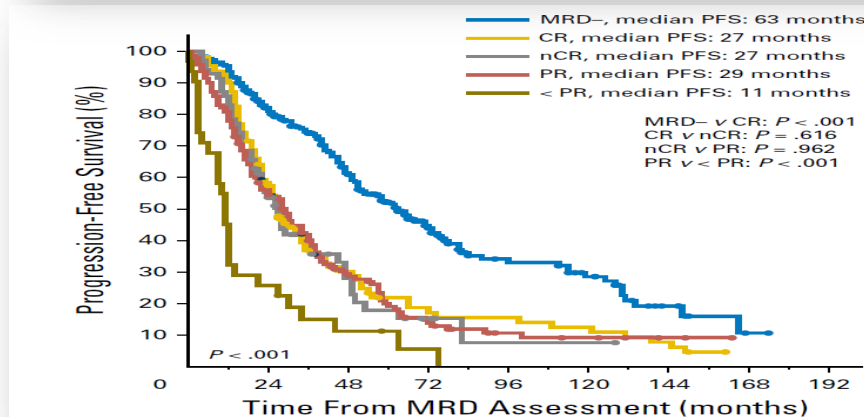
Tacchetti P. et al., Lancet Haematology 2020, accepted

Depth of response correlate with survival: MRD is the best biomarker to predict outcome



Meta-analysis of MRD studies (CR patients)

Munshi NC, et al. JAMA Oncol 2017;3(1):28-35



GEM2000 - GEM2005MENOS65 - GEM2010MAS65

Lahuerta JJ, et al. JCO 2017;35(25):2900-2910

PETHEMA/GEM study: Post-transplant negative MRD (1×10^{-4})*

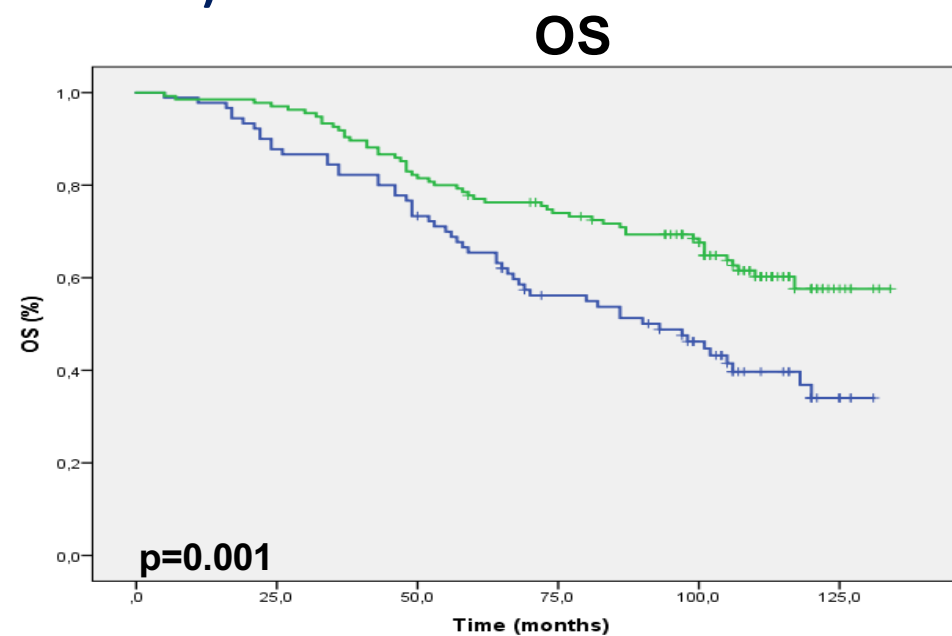
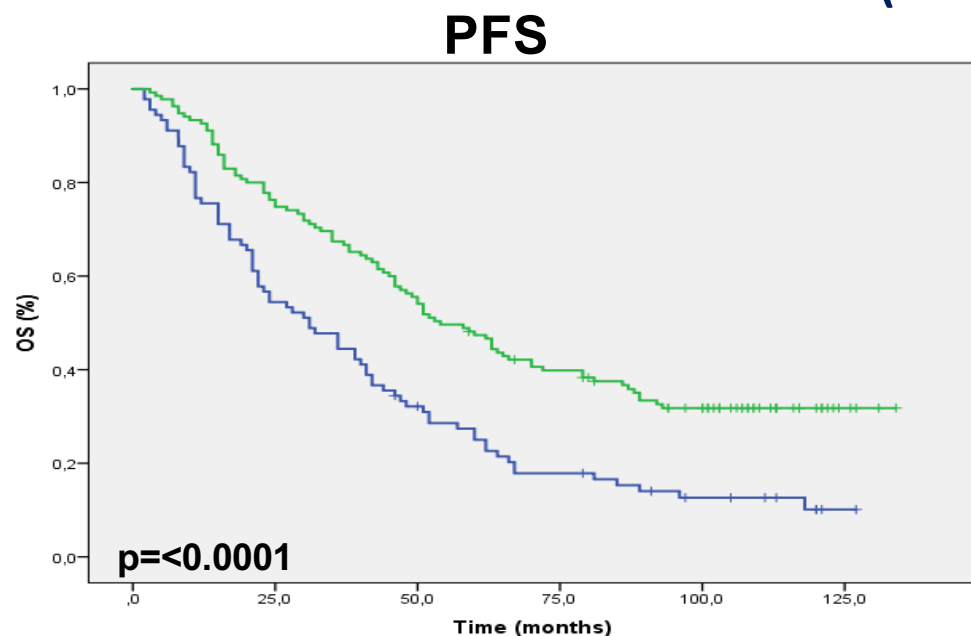
	Negative MRD
Overall series (n=226)	135 (60%)
CT+V (n=85)	48 (56%)
TD (n=50)	27 (46%)
VTD (n=83)**	60 (72%)

*MRD available in 226/ 284 (80%) transplanted patients

** VTD vs TD, $p=0.03$; VTD vs CT+V, $p=0.04$; TD vs CT+V, $p=0.9$

Rosinol L. et al., ASH 2018

PETHEMA/GEM study: PFS and OS according to post-transplant MRD (overall series)



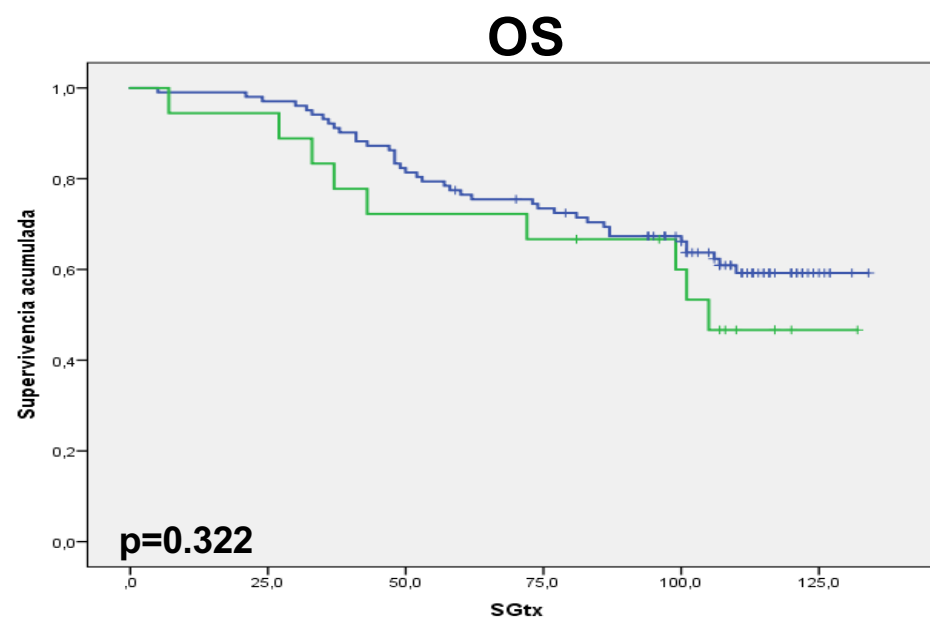
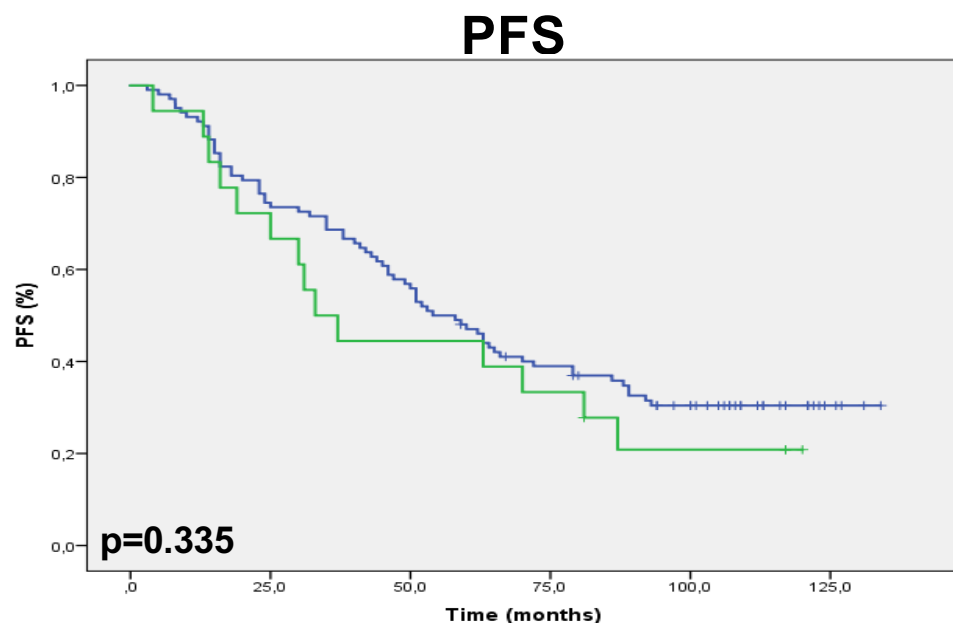
	Median PFS (mos)	Median OS (mos)
Negative MRD	55	NR
Positive MRD	31	93

Rosinol L. et al., ASH 2018



2020

PETHEMA/GEM study: Outcome of patients with **negative** post-transplant **MRD** according to cytogenetics



	Median PFS (mos)	Median OS (mos)
Standard- risk cytogenetics	54	NR
High-risk cytogenetics	33	105

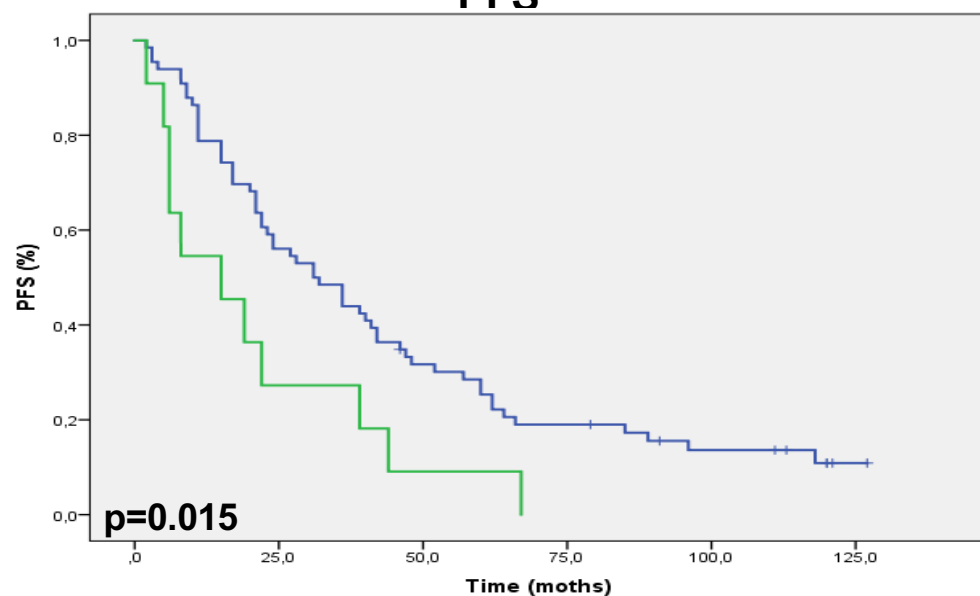
Rosinol L. et al., ASH 2018



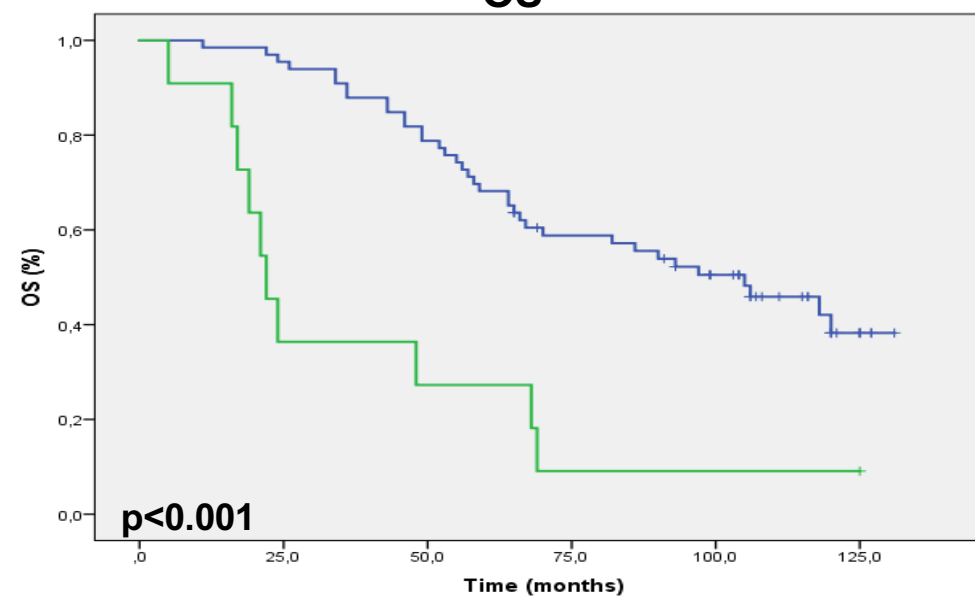
2020

PETHEMA/GEM study: Outcome of patients with **positive** post-transplant **MRD** according to cytogenetics

PFS



OS



	Median PFS (mos)	Median OS (mos)
Standard- risk cytogenetics	31	105
High-risk cytogenetics	15	22

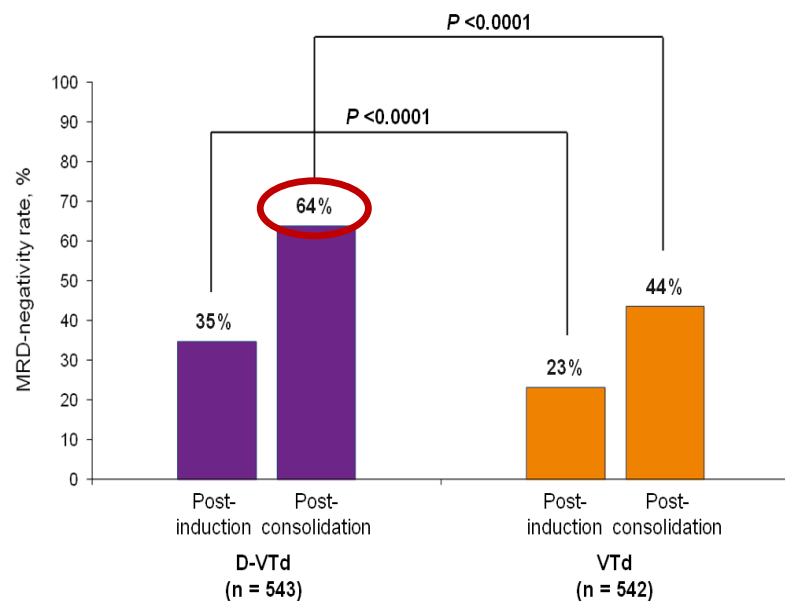
Rosinol L. et al., ASH 2018



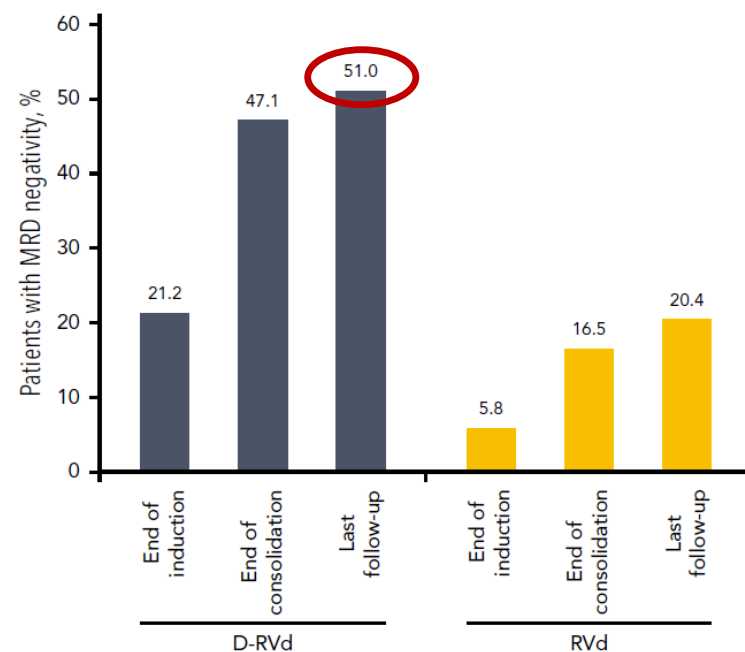
2020

Modern treatment strategies for NDTE: MRD data (10^{-5})

CASSIOPEIA ph3 trial



GRIFFIN ph2 trial



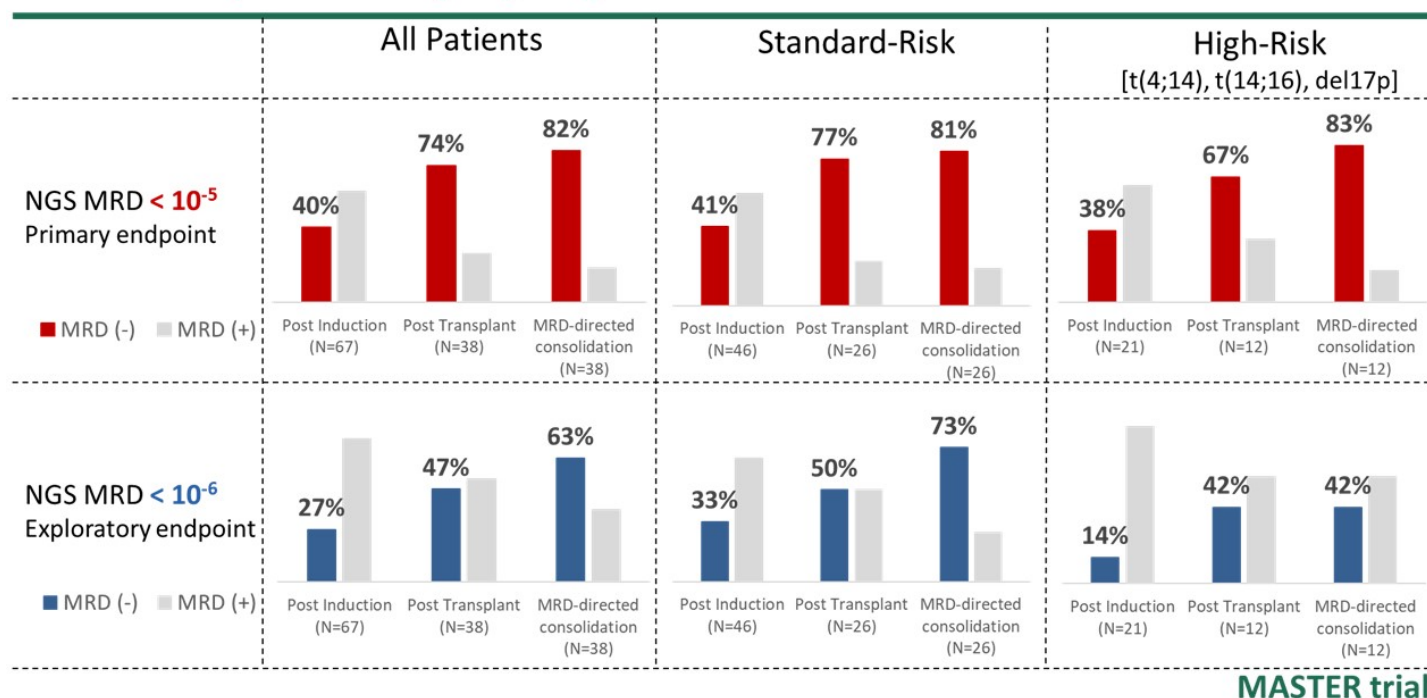
Moreau P, et al. *The Lancet* 2019;394:29-38; Avet-Loiseau H, et al. *IMW* 2019, Oral presentation; Voorhees PM et al., *Blood*. 2020;136(8):936-945



2020

DaraKRd MASTER Trial : Best MRD response by phase of therapy

MRD response by cytogenetic subset

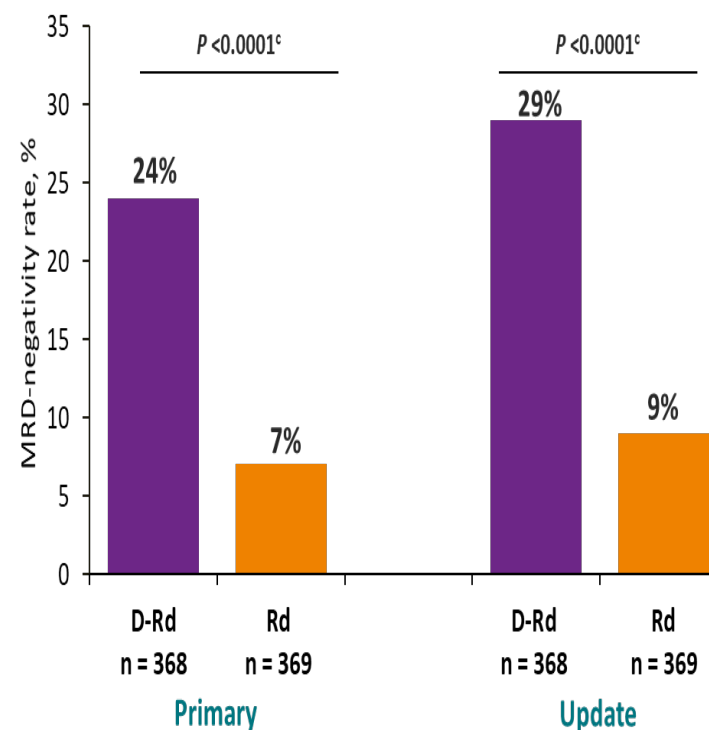
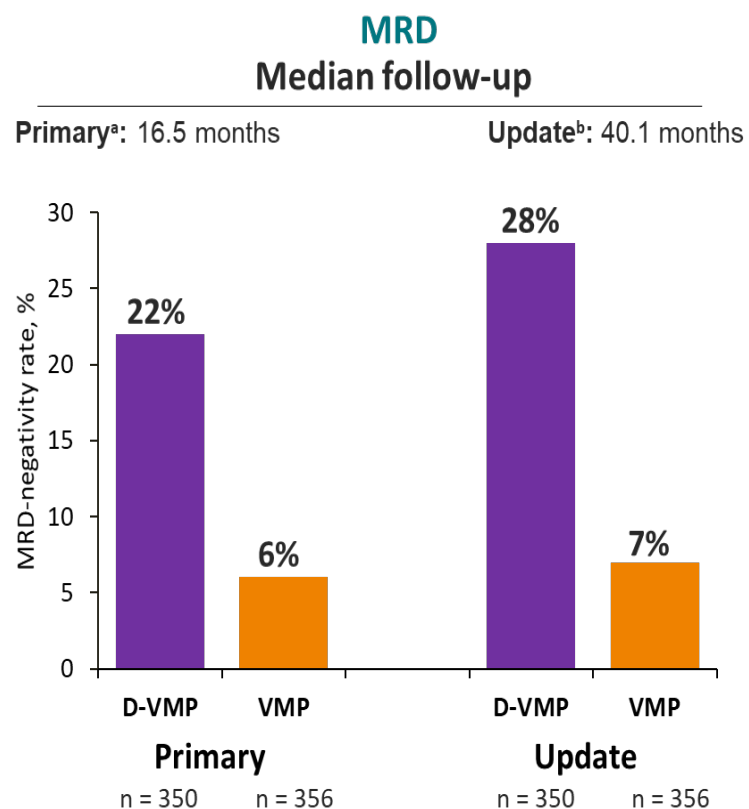


Costa LJ, et al., ASH 2019; oral presentation



2020

Modern treatment strategies for ND non TE: MRD data (10^{-5})



Mateos MV et al., Lancet 2020; Facon T et al. NEJM 2019

What do we know about long-term survivors

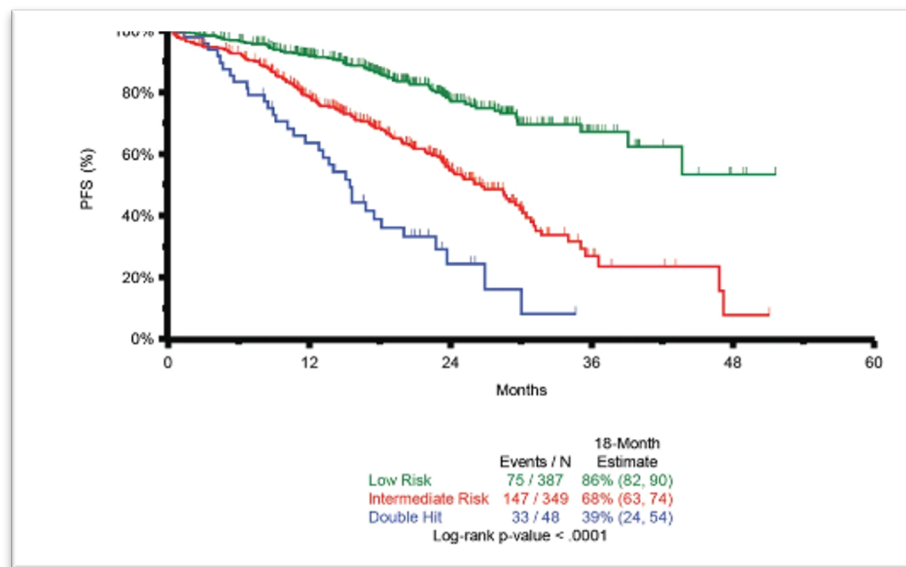
They Have

- » Low disease burden and minimal end-organ disease
- » Favorable disease biology
- » CR or better
- » Receive optimal up-front therapy (PI + IMiDs, ASCT, maintenance, mAB?)

They Do Not

- » High disease burden
- » Unfavorable disease biology (R-ISS3, HRCA, UltraHR, biallelic p53 del, EMD/PCL)
- » Suboptimal responses
- » Multiple co-morbidities

Unmet medical need: High-Risk and Ultra High-Risk Group: Double-Hit MM



Characteristic: Bi-allelic inactivation of TP53 or ISSIII + amplification of CKS1B

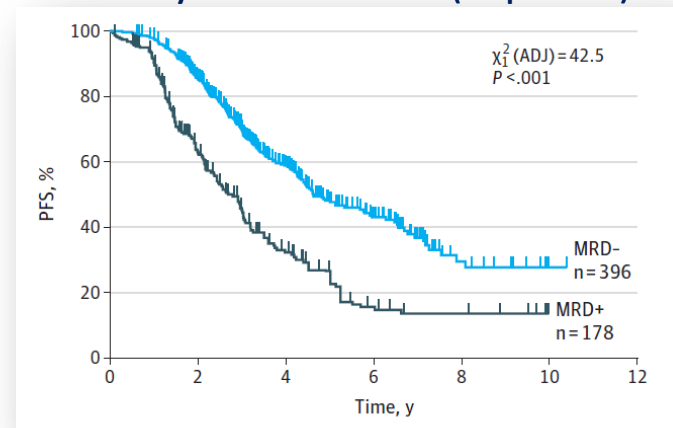
Median PFS = 15.4 mths; OS 20.7 mths

Walker BA, et al., *Leukemia* 2018;33(1):159-170

Depth of response correlate with survival MRD is the best biomarker to predict outcome

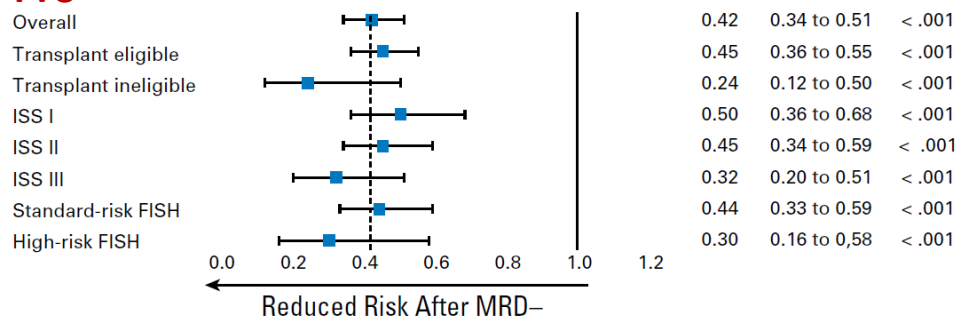
Munshi NC, et al. *JAMA Oncol* 2017;3(1):28-35

Meta-analysis of MRD studies (CR patients)



MRD negativity is a prognostic marker for PFS and OS across the spectrum of patients with MM

PFS

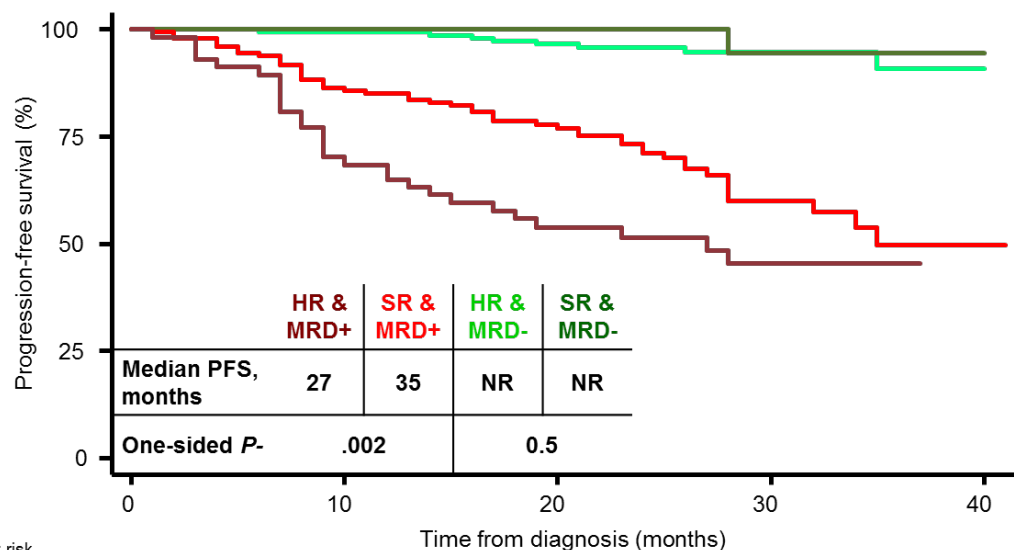


Lahuerta JJ, et al. *JCO* 2017;35(25):2900-2910

Can MRD-response modulate patients' risk at diagnosis?

Risk in dynamic: patients with adverse prognosis shift into a favorable one upon achieving deep responses to treatment

Progression-free survival according to FISH and NGF



The best pathway to overcome the poor prognosis of HRCA is through the achievement of MRD-negativity

Number at risk					
SR FISH & MRD-neg	0	151	122	59	1
HR FISH & MRD-neg	0	33	28	14	2
SR FISH & MRD-pos	300	126	93	31	3
HR FISH & MRD-pos	90	40	27	12	0

HR FISH: t(4;14), t(14;16) and/or del(17p)

Paiva B et al. ASH 2017

Conclusions

- There are some known **predictors of long-term survivors**: depth or response, disease biology, tumor burden
- **Not all long-term survivors** can be considered effectively **cured**
- The combination of an extended PFS time (ie 78 months), depth of response and absence of high risk features, can be associated with **survival curves potentially reaching a plateau**
- **MRD negativity is a strong predictor of survival**, showing a higher prognostic power than CR, patients with **adverse prognosis shift into a favorable one** upon achieving deep responses (sustained MRD negativity) to treatment
- Ongoing clinical trials will provide further insights into the role of **MRD disease-driven treatment strategies** for these patients in the near future