LYMPHOID NEOPLASIA

Prognostic implications of serial 18-fluoro-deoxyglucose emission tomography in multiple myeloma treated with total therapy 3

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Key Points

• Presence of more than 3 PET focal lesions after day 7 first cycle of induction chemotherapy can predict for inferior overall survival and progression free survival. Prognostic implications of 3 imaging tools, metastatic bone survey, magnetic resonance imaging, and positron emission tomography (PET), were evaluated in 2 consecutive Total Therapy 3 trials for newly diagnosed myeloma. Data including PET at baseline and on day 7 of induction as well as standard prognostic factors were available in 302 patients of whom 277 also had gene expression profiling (GEP)derived risk information. According to multivariate analysis, more than 3 focal lesions on day 7 imparted inferior overall survival and progression-free survival, overall and in the subset with GEP-risk data. GEP high-risk designation retained independent significance for all 3 end points examined. Thus, the presence of > 3 focal lesions on day 7

PET follow-up may be exploited toward early therapy change, especially for the 15% of patients with GEP-defined high-risk disease with a median overall survival expectation of 2 years. This trial was registered at www.clinicaltrials.gov as #NCT00081939 and # NCT00572169. (*Blood.* 2013;121(10):1819-1823)

Introduction

Magnetic resonance imaging (MRI) and fluoro-deoxy-glucose (FDG) positron emission tomography (PET) scanning are increasingly viewed as important state-of-the-art imaging tools in the initial workup of patients with multiple myeloma (MM).^{1,2} The prognostic implications of PET scanning performed at baseline and after induction and high-dose therapy interventions have recently been reported.³ Here we investigated the survival implications of the day 7 PET scanning of patients treated with Total Therapy 3A⁴ clinical trial and successor protocol Total Therapy 3B⁵ protocol.

Study design

Details of the TT3 protocols have been reported previously.^{4,5} The protocols and their modifications had been approved by the University of Arkansas Medical Sciences Institutional Review Board. Patients signed written informed consent in keeping with institutional, federal, and international guidelines (Helsinki Declaration). MM diagnostic and response criteria employed the European Bone Marrow Transplant criteria introduced by Blade et al.⁶

In addition to standard baseline variables, presence and type of metaphase cytogenetic abnormalities (CA, CA13) were considered.^{4,5} Imaging variables included radiograph-defined osteolytic lesions (OL), MRI-based focal lesions (FL), and diffuse hyperintense marrow involvement¹ as well as PET-FL and maximum FL standard uptake value (SUV) (SUVmax).³ PET studies were repeated

on day 7 of induction and before first transplant. Although MRI examinations were also performed before the second transplant, consolidation, and maintenance phases, this report considers only baseline and pretransplant data.

Survival distributions were estimated according to the Kaplan-Meier method.⁷ Cumulative incidence curves were estimated as described by Gooley.⁸ These estimates were compared using the log-rank test.⁹ As of March 9, 2012, median follow-up times are 6.8 years and 4.3 years for Total Therapy 3A and Total Therapy 3B, respectively. Univariate and multivariate regression analyses were performed using a stratified Cox regression model.¹⁰ Stepwise variable selection was used to select the multivariate models. Cut-points for imaging parameters were applied as previously reported, including multiple prognostic OL and FL number cutoffs (ie, >0 and >2 for metastatic bone survey (MBS)-OL, >0 and >7 for MRI-FL, and >0 and >3 for PET-FL).² An accounting of the patients included in the tables and figures can be found in supplemental Table 1.

Clinical end points included overall survival (OS), progressionfree survival (PFS), and complete response duration (CRD). OS events included death from any cause, whereas PFS events included disease relapse or progression. CRD was measured as the time from complete response onset to disease progression or death from any cause. Land-marking methods were employed in order to include post-induction PET variables in our analyses. In the absence of progressions, deaths, or censored follow-up before the day 14 landmark, baseline, and landmarked Cox models for OS

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Figure 1. Effects of imaging variables on OS, PFS, and CRD in Total Therapy 3A and Total Therapy 3B combined. (A) Baseline number of MBS-defined osteolytic lesions (OL). Superior OS and PFS were linked to the presence of no more than 2 OL. A trend was noted in case of complete response duration. (B) Baseline number of MRI-defined FL and diffuse hyperintense marrow. Trends were observed for superior OS and PFS in case of MRI-FL not exceeding 7 FL. (C) Baseline PET-defined number of FL. The presence of more than 3 PET-defined FL affected all 3 survival end points adversely. (D) Baseline PET-defined SUVmax. Higher SUVmax of PET-defined FL conferred inferior OS, PFS, and complete response (CR) duration. (E) Day 7 PET-FL. OS and PFS were inferior when more than 3 FL persisted on day 7 after starting protocol therapy. In case of CR duration, a strong trend in the same direction was noted. (F) Pretransplant MRI-FL. Trends were observed for both OS and PFS with the best survival at 3 years observed in those with 0 MRI FL, followed by 1 to 7 MRI FL and >7 MRI FL Similar to the baseline results, survival at 3 years for those with DHIM was more like patients with 1 to 7 or >7 MRI FL. (G) Pretransplant PET-FL. A graded negative impact was observed with increasing PET-defined FL remaining before first transplant.

and PFS are equivalent. Thus, only land-marked models for OS and PFS are included below. For CRD, additional land-marking is not required to include day-7 PET data since complete response onset was not observed for any patients within 14 days of initiation of therapy. Survival estimates for OS and PFS by pretransplant PET and MRI are landmarked at the first transplant date. CRD is not presented with pretransplant PET and MRI results, because this would exclude a number of posttransplant CRs.

Results and discussion

MBS-OL >2 adversely affected OS and PFS with a trend for CRD (OS: P < .0001, PFS: P = .0006, CRD: P = .15) (Figure 1A). OS differed among the 4 MRI baseline variables considered (P = .04) (Figure 1B). PFS tended to be superior in FL = 0 and FL 1-7 categories, whereas diffuse hyperintense marrow involvement and FL >7 baseline findings imparted borderline inferior outcomes

(P = .1). PET-FL impacted all 3 outcome variables in a consistent manner (OS: P < .0001, PFS: P = .0002, CRD: P = .01) (Figure 1C). Patients with 0 or 1-3 FL had equally favorable OS, PFS, and CRD, whereas those presenting with FL > 3 fared poorly. Similarly, PET-derived SUVmax of FL exceeding 3.9 conferred inferior OS, PFS, and CRD (OS: P = .03, PFS: P = .01, CRD: P = .02) (Figure 1D). A postinterventional PET scan follow-up examination on day 7 also provided prognostic value (Figure 1E). Presence of more than 3 FL was linked to inferior OS, PFS, and CRD with borderline significance (OS: P < .0001, PFS: P =.0003, CRD: P = .07). Day 7 SUVmax failed to affect outcomes significantly (data not shown). Consideration of pretransplant MRI FL suggested longer OS and PFS for cases without MRI FL. After 3 years, OS was >10% higher and PFS >12% higher for those without FL compared with all others (OS: P = .07, PFS: P = .19) (Figure 1F). PET-FL before transplant affected OS and PFS in a graded fashion (OS: P = .0002, PFS: P = .001) (Figure 1G).

Next we examined by multivariate Cox regression analysis which baseline variables in the context of day 7 PET-FL affected



Figure 1. (Continued).

Table 1. Multivariate analysis of baseline and day 7 PET findings on clinical outcomes landmarked from day 14 of induction therapy: OS, PFS, and CRD

Variable	Excluding GEP variables			Including GEP variables		
	n/N (%)	HR (95% CI)	P value	n/N (%)	HR (95% CI)	P value
os						
GEP70 high risk				52/277 (19%)	4.10 (2.57-6.52)	<.001
Baseline MBS OL >2	96/302 (32%)	1.82 (1.17-2.83)	.008	86/277 (31%)	1.81 (1.16-2.85)	.010
Day-7 PET FL >3	81/302 (27%)	1.86 (1.19-2.91)	.006	75/277 (27%)	1.67 (1.05-2.65)	.030
Age ≥65 years	78/302 (26%)	1.79 (1.16-2.76)	.008	72/277 (26%)	1.81 (1.16-2.83)	.009
Cytogenetic abnormalities	109/302 (36%)	2.15 (1.40-3.30)	<.001			
B2M >5.5 mg/L	73/302 (24%)	1.95 (1.25-3.03)	.003			
PFS						
GEP70 high risk				52/277 (19%)	3.23 (2.09-4.99)	<.001
Day 7 PET FL >3	81/302 (27%)	1.83 (1.23-2.71)	.003	75/277 (27%)	1.81 (1.21-2.70)	.004
B2M >5.5 mg/L	73/302 (24%)	1.82 (1.19-2.77)	.005	66/277 (24%)	1.60 (1.05-2.45)	.030
LDH ≥190 U/L	75/302 (25%)	1.69 (1.12-2.53)	.012			
CA13	57/302 (19%)	1.75 (1.13-2.71)	.012			
Albumin <3.5 g/dL	107/302 (35%)	1.60 (1.07-2.37)	.021			
CRD						
GEP-70 high risk				28/180 (16%)	3.61 (1.93-6.77)	<.001
B2M >5.5 mg/L	39/196 (20%)	3.77 (2.11-6.75)	<.001	36/180 (20%)	2.89 (1.56-5.37)	<.001
CA13	40/196 (20%)	1.98 (1.07-3.67)	.030			

Model selection and estimates based on the set of patients with complete data for the variables examined. *P* value from Wald χ -square test in Cox regression. Variables considered: age \geq 65 years, albumin <3.5 g/dL, B2M \geq 3.5 mg/L, B2M \geq 5.5 mg/L, creatinine \geq 2.0 mg/dL, C-reactive protein \geq 8 mg/L, hemoglobin <10 g/dL, LDH \geq 190 U/L, cytogenetic abnormalities, CA13, hypodiploid, CA 13/hypodiploid, GEP70 high risk, GEP Proliferation Index \geq 10, GEP Centrosome Index \geq 3, GEP Molecular Subgroup, baseline MBS OL (>0, >2), baseline MRI FL (>0, >7), baseline PET FL (>0, >3), baseline PET FL (>0, >3).

GEP, gene expression profiling; HR, hazard ratio; LDH, Lactate dehydrogenase



Figure 1. (Continued).

clinical outcomes applying a 14-day landmark from the beginning of induction therapy (Table 1). Data are presented also for the subgroup of 277 of 302 patients in whom GEP-based risk designation was available. Univariate data are summarized in supplemental Table 2. In the absence of GEP data, OS was dominantly affected by the presence of metaphase cytogenetic abnormalities (CA), high beta-2-microglobulin (B2M) (>5.5 mg/L), day 7 persistence of more than 3 FL, baseline MBS-OL >2, and older age (\geq 65 years). With knowledge of GEP data, GEP-defined highrisk status (70-gene model)⁵ was the dominant adverse feature imparting a 4.1-fold higher risk of death (95% CI: 2.57-6.52), whereas baseline MBS-OL, day-7 PET-FL, and older age all posed additional risk with hazard ratio values of 1.7 to 1.8. In the case of PFS, day 7 PET-FL persistence of >3 FL was the dominant adverse variable, followed by high B2M, CA subtype CA13, high lactate dehydrogenase (\geq 190 U/L) and low albumin (<3.5 g/dL). With access to GEP information, high risk dominated with a 3.23-fold higher risk of relapse or death (95% CI: 2.09-4.99), whereas day 7 PET-FL and B2M were the other independently adverse features. CRD, finally, was inferior in case of high B2M and in the presence of CA13. GEP-defined high-risk replaced CA13 in the subset of patients with GEP data. Day 7 PET data failed to enter the CRD model.

Collectively, our data confirm and extend observations by our group and others on the powerful prognostic implications of baseline and follow-up PET examinations.^{2,3} An important novel finding

relates to the superiority of day 7 PET-defined FL >3 over baseline findings, retaining such significance for OS and PFS in the presence of GEP data, displacing CA as a variable. B2M survived the multivariate models for all 3 end points examined and is only displaced from the OS model with knowledge of GEP risk. Although MBS-OL is a later event than the development of FL detected on MRI or PET scans, the presence of more than 2 OL retained independent adverse implications for OS also in the presence of GEP data, confirming the pioneering work of Durie and Salmon.¹¹ These findings strengthen the argument for validating such examinations in other trials as a basis for modifying therapy in the subset of patients with persisting FDG-avid FL >3. Such early corrective therapeutic measure should be of particular benefit in patients with GEP-defined high-risk MM.^{12,13}

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Authorship

Contributions: S.Z.U. and B.B. designed the study and wrote the manuscript; S.Z.U., B.B., A.M., A.H., J.C., N.C., N.P., and E.A. analyzed the data; B.B., E.A., F.v.R., S.W., and S.Z.U. contributed patients; and S.Z.U., A.M, S.W., J.C., A.H., N.P, T. Brown, T. Bartel, E.A., F.v.R., and B.B. reviewed and approved the manuscript.

Conflict-of-interest disclosure: S.Z.U. is a consultant to Celgene, Millennium, and Onyx; has received research funding from Onyx and Celgene and speaking honoraria from Celgene. B.B. has received research funding from Celgene, Millennium, and Onyx; has served as consultant to Celgene, Millennium, Onyx, and Genzyme; is a co-inventor on patents and patent applications related to use of gene expression profiling in cancer medicine. He has no financial interest in SignalGenetics.

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