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Br J Haematol. Author manuscript; available in PMC 2018 September 01.

Published in final edited form as:

Author manuscript

Br J Haematol. 2017 September ; 178(6): 888–895. doi:10.1111/bjh.14830.

## Elevation of serum lactate dehydrogenase in AL amyloidosis reflects tissue damage and is an adverse prognostic marker in patients not eligible for stem cell transplantation

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## Summary

The significance of serum lactate dehydrogenase (LDH) in light chain (AL) amyloidosis has not been previously explored. We studied 1019 newly diagnosed patients and correlated the elevation of LDH above the upper limit of normal (ULN) with disease characteristics and outcome. Four hundred and nine patients had an LDH above ULN, representing 40% of the study population. Patients with an elevated LDH were older, were less likely to be male and had more extensive organ involvement compared to patients with a normal LDH. Patients with high LDH had greater cardiac and renal dysfunction. Elevated LDH was an independent prognostic marker for overall survival and for death within 6 month of diagnosis, but this was restricted to patients not eligible for stem cell transplant. Serum LDH may act as a marker for organ damage and should be explored as a potential marker for tissue healing and organ recovery.

#### Authorship Contributions

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**Disclosure of Conflicts of Interest** 

Eli Muchtar: None; Angela Dispenzieri received research funding from Celgene, Millennium, Pfizer and Janssen, and travel grant from Pfizer; Martha Q. Lacy received research funding from Celgene; Francis K. Buadi: None; Prashant Kapoor received research funding from Takeda, Celgene and Amgen; Suzanne R. Hayman: None; Wilson Gonsalves: none; Rahma Warsame: None; Taxiarchis V. Kourelis: None; Stephen Russel: None; John A. Lust: None; Yi Lin: None; Ronald S. Go: None; Steven Zeldenrust: None; David Dingli received research funding from Karyopharm Therapeutics, Amgen and Millenium Pharmaceuticals; Nelson Leung: None; S. Vincent Rajkumar: None; Robert A. Kyle: None; Shaji Kumar received consultancy from Celgene, Millennium, Onyx, Janssen and BMS and research funding from Celgene, Millennium, Novartis, Onyx AbbVie, Janssen and BMS; Morie A. Gertz received consultancy from Millenium and honoraria from Celgene, Millenium, Onyx, Novartis, Smith Kline, Prothena, Ionis and Amgen.

E.M. designed the study, collected and analysed the data, wrote the first draft and approved the final version of the manuscript; A.D., M.Q.L., F.K.B., P.K., S.R.H., W.G., R.W., T.V.K., S.R., J.A.L., Y.L., R.S.G., S.Z., D.D., N.L., S.V.R., S.K.K. performed patient management, participated in final data analysis, revised the manuscript critically and approved of the final version of the manuscript. R.C. analysed data, provided critical review of the manuscript and approved the last version of the manuscript; R.A.K. Performed patients' follow-up, participated in final data analysis, revised the manuscript critically and approved of the final version of the manuscript; M.A.G. performed patient management, designed the study, analysed the data, wrote the first draft and approved the final version of the manuscript.

## Keywords

LDH; tissue; damage; prognosis; early death

## Introduction

Lactate dehydrogenase (LDH) is a cytoplasmic enzyme found in all tissues. Elevation of serum LDH is associated with various causes of cell damage,(Glick 1969) such as myocardial infarction,(Danese and Montagnana 2016) acute necrotizing pancreatitis, (Waldemar, *et al* 1995) acute ischaemic renal tubular necrosis(Green, *et al* 2016) and haemolysis(Tabbara 1992). Elevated serum LDH is associated with various malignancies, potentially as an indicator of tumour metabolic activity, proliferation and invasiveness. (Gallo, *et al* 2015) Serum LDH is elevated in both solid and haematological cancers (Petrelli, *et al* 2015, Walenta and Mueller-Klieser 2004, Wulaningsih, *et al* 2015) and is used for prognostic purposes in several types of cancer.(Miles, *et al* 2012, Palumbo, *et al* 2015, Petrelli, *et al* 2015, Winter 2007)

Immunoglobulin light chain (AL) amyloidosis is a unique syndrome due to a monoclonal expansion of plasma cells, in which nearly half of patients reach the threshold of multiple myeloma, with more than 10% clonal plasma cells in the bone marrow.(Kourelis, *et al* 2013) However, amyloidosis is associated with misfolded protein deposition and tissue damage. We hypothesized that elevation of serum LDH in AL amyloidosis may be a marker for tissue damage rather than a measure of tumour mass in this low proliferative clonal process.

## Methods

One thousand and nineteen patients (n=1019) with biopsy-proven systemic AL amyloidosis who were seen in our institution within 90 days of diagnosis between 1 January 2000 and 31 August 2015 and had serum LDH measurement before treatment were included in this study. Patients were excluded if they had been previously treated for AL amyloidosis or multiple myeloma (MM), had AL amyloidosis due to a lymphoproliferative disorder or had incidental amyloid deposits without demonstrable visceral involvement.

Data were extracted from prospectively maintained databases and patient records were reviewed for data accuracy and completeness. The study was approved by the Institutional Review Board (IRB). All patients gave written informed consent for medical record review according to the IRB and Minnesota state law.

Serum LDH was measured using a commercially available reagent (Roche diagnostics, Indianapolis, IN, USA). Patients with serum LDH activity greater than the upper limit of normal (ULN) were designated as "high serum LDH" group, while those with serum LDH below ULN were categorized in the "normal serum LDH" group.

Assessment of organ involvement, haematological response and organ response were determined by consensus criteria.(Comenzo, *et al* 2012, Gertz, *et al* 2005) Bone marrow plasma cells (BMPCs) were considered as the highest estimate of the bone marrow

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aspiration and biopsy. Immunoparesis was assessed using a quantitative method, where significant immunoparesis was defined when the average relative difference of the uninvolved immunoglobulins from their respective lower limit of normal was below zero, as previously described.(Muchtar, *et al* 2017a) Treatment was categorized at autologous stem cell transplant (ASCT) and non-ASCT regimens. Current eligibility for ASCT in our centre has been published.(Muchtar, *et al* 2016a)

The Pearson  $\chi^2$  test and the Kruskal-Wallis test were used to ascertain differences between nominal and continuous variables, respectively. Survival analysis was performed using the Kaplan-Meier method. Cox proportional hazards regression models were built to identify independent prognostic factors for survival. One model included the whole study cohort and another model was restricted to patients who received standard therapy without ASCT. In addition, a binary logistic regression model was built to identify independent predictors of death within the first 6 months of diagnosis. All variables in the univariate analysis were entered into the multivariate analysis, regardless of significance reached in the univariate analysis. P values less than 0.05 were considered significant. Statistical analysis was performed using JMP software (SAS, Cary, NC).

## Results

Of the 1019 patients included, 610 (60%) had serum LDH within the normal range and 409 patients (40%) had serum LDH level above the ULN.

Baseline characteristic of the entire cohort and by serum LDH groups can be viewed in Table I. The two serum LDH groups differed by age (older population in the high serum LDH group, median 63 vs 61 years; P=0.01) and by sex (44% females in high LDH group vs 34% in the normal LDH group; P=0.001). Patients with an elevated LDH were more likely to present with a systolic blood pressure below 100 mmHg (26% vs 20%; P=0.02). Of note, patients presented with a lower frequency of extensive organ involvement (>2 organs) in the most recent period (2008–2015) compared to the earlier period (2000–2007) (22% vs 31%; P=0.002) as well as a lower proportion of cardiac involvement (66% vs 77%; P<0.001) and hepatic involvement (14% vs 26%; P<0.001), but not renal involvement (64% vs 64%; P=0.99). There was no difference in the proportion of high LDH between periods (42% vs 40%; P=0.57).

#### Correlation between LDH and organ involvement pattern

Patients in the high serum LDH group were more likely to have extensive organ involvement (>2 organs) compared to patients in the normal serum LDH group (32% vs 25%; *P*=0.008). Also, elevated serum LDH patients were more likely to have cardiac (83% vs 66%; *P*<0.001), renal (73% vs 57%; *P*<0.001) and hepatic involvement (25% vs 19%; *P*=0.01) compared to their counterparts. The high LDH group had a lower proportion of nerve involvement (14% vs 22%; *P*<0.001). In an attempt to better characterize the source of LDH elevation, we analysed the frequency of LDH elevation among patients with single-organ involvement. Three hundred and forty-six patients (n=346) had predominantly single organ involvement. Among patients with cardiac involvement alone (n=150), 36% had a LDH above the ULN. In comparison, 29% of patients with only renal involvement (n=150) had an

elevated LDH (P=0.17). Numbers of patients were lower in other organs (n=12 for nerve and n=6 for liver involvement as a single organ), with 17% of patients in each of these organ groups having an elevated LDH.

#### Correlation of serum LDH and laboratory studies

In an attempt to characterize the source of serum LDH, we correlated laboratory findings with the two study groups. The results are in given in Table II. Parameters associated with the plasma cell compartment (BMPCs, difference between involved and uninvolved light chains [dFLC], labelling index and immunoparesis state) did not demonstrate a correlation with LDH. A correlation with haemoglobin and bilirubin was not found, making haemolysis the cause of LDH elevation unlikely. Given that organ involvement distribution was unbalanced between the LDH groups, we analysed organ function in an organ-stratified fashion. Among patients with cardiac involvement, the cardiac biomarkers N-terminal pro btype natriuretic peptide (NT-proBNP) and troponin T were significantly higher among patients with high LDH than those with LDH below the ULN (median NT-proBNP 6164 vs 2385 pg/ml; median troponin T 0.08 vs 0.02 ng/ml; P<0.001 for both). Among patients with renal involvement, the high LDH group had a lower estimated glomerular filtration rate (median 53 vs 66 ml/min/1.73 m<sup>2</sup>; P<0.001) and a higher proteinuria level (median 5.3 vs 4.1 g/24 h collection; *P*=0.008) compared with their counterparts. Among patients with hepatic amyloidosis, we found no association between LDH groups and serum alkaline phosphatase or total bilirubin levels.

#### Correlation to response to first line treatment

LDH elevation did not correlate with the haematological response to first line treatment. Partial response or better was similar between groups (82% in the normal LDH group and 80% in the high LDH group; P=0.64). There was also no association with depth of response (very good partial response or better 59% vs 61%, respectively; P=0.58). Correspondingly, the LDH groups were not associated with depth of response whether stem cell transplantation was used as primary therapy (P=0.48) or not (P=0.73). Similarly, no difference in organ response rate was seen between the LDH groups (51% vs 49%; P=0.56).

## **Survival analysis**

Patients with an elevated LDH had a shorter OS compared to patients with LDH within the normal range (median 18 vs 59 months; P<0.001) (Figure 1). This survival difference was seen among non-ASCT patients (median OS 9 vs 14 months; P=0.002), but not among patients who underwent ASCT as their primary therapy (median 125 vs 135 months; P=0.36). A survival difference was also seen in subgroup analysis by number of involved organs. In patients with single organ involvement, the high serum LDH group had a shorter OS compared to those with normal LDH (median 47 vs 102 months; P<0.001). Similarly, among patients with 2 or more involved organs, survival was shorter in the high LDH group (median 14 vs 31 months; P<0.001). In a multivariate analysis, high serum LDH did not demonstrate an independent impact on OS (P=0.08) (Table III, model A). Since an OS difference was restricted to the non-ASCT cohort, we performed a multivariate analysis limited to non-ASCT patients. In this patient group, high serum LDH was associated with an independent effect on OS with a hazard ratio of 1.3 (95% confidence interval [CI] 1.02–1.7;

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*P*=0.03) (Table III, model B). When multivariate analysis of model B was performed using the variables in their continuous form (when applicable), LDH retained its independent association with survival (hazard ratio 1.6 per one unit change; 95% CI 1.2–2.2; P=0.003).

## Effect on early death ( 6 months of diagnosis)

Finally, the impact of LDH on the risk of early death (within 6 month of diagnosis) was assessed. Patients with elevated LDH had a greater likelihood of early death compared to patients with normal LDH (32% vs 20%; P<0.001). Sub-analysis by the primary treatment showed this difference to be seen among the non-ASCT cohort (43% vs 32%; P=0.005), but not among ASCT patients (4% vs 7%; P=0.34). A binary logistic regression analysis was performed to identify independent predictors of early death in the whole study cohort and in the non-ASCT cohort. The results of this analysis can be viewed in Table IV. Serum LDH above ULN was associated with an independent increase in early death (odds ratio [OR] 1.9, 95% CI 1.1–3.1; P=0.01). Other predictors for early death in this model were dFLC 18 mg/dl (OR=1.7); systolic blood pressure (SBP) <100 mmHg (OR=1.7); NT-proBNP 1800 pg/ml (OR=5.1) and troponin T 0.025 ng/ml (OR=2.1).

## Discussion

AL amyloidosis can be viewed as a two-compartment disease. The first compartment is the clonal plasma cell disorder that resides in the bone marrow and produces the amyloidogenic protein depositing in this disease. The biological characteristics of the plasma cells, such as their percentage in the bone marrow,(Kourelis, *et al* 2013) their proliferation rate(Gertz, *et al* 1989, Kumar, *et al* 2012) and their cytogenetic content(Warsame, *et al* 2015) have been shown to be prognostic. Cytogenetics also correlates with response to various regimens. (Bochtler, *et al* 2015, Muchtar, *et al* 2016b) The second disease compartment in AL amyloidosis is the organ dysfunction caused by the amyloid deposits. These deposits accumulate in organs and tissues throughout the body, leading to damage at both the cellular level and tissue architecture, which lead to organ dysfunction.

No data is available on LDH elevation in AL amyloidosis. In MM, serum LDH is elevated in approximately 12% of patients, is prognostic and used to define high-risk disease. (Gkotzamanidou, *et al* 2011, Palumbo, *et al* 2015) This study demonstrated that elevation of LDH above normal occurs in 40% of newly diagnosed AL amyloidosis patients, higher than observed in MM. This prevalence is unexpected in light of the fact that AL amyloidosis is a plasma cell disorder, often with a low clonal cell burden (median 7–10%). The source of LDH does not appear to be from the plasma cell compartment. Our analysis did not show any correlation between serum LDH and various measures of the clonal plasma cell burden, including the percentage of bone marrow plasma cells, the light chain level, the proliferation rate and the presence of significant immunoparesis. In contrast, we have demonstrated that high levels of serum LDH are associated with more extensive disease (>2 involved organs) as well as higher levels of dysfunction of the heart and kidneys. High LDH was also more likely in patients with kappa light chain. While light chain restriction is a feature of the plasma cells, kappa light chain has a tissue tropism for liver and thyroid gland involvement.

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(Huang, *et al* 2015, Muchtar, *et al* 2017b) Serum LDH was not associated with response to therapy. This also suggests the plasma cell compartment is not the source of serum LDH.

The exact source of the elevated LDH in AL amyloidosis cannot be determined from our data and may be widespread given the abundance of LDH in various body tissues. In patients with single organ involvement, a similar rate of elevated LDH was seen between those with heart involvement and those with renal involvement. Therefore, our data do not suggest any predilection of either heart or kidneys. The high LDH group was more likely to have SBP below 100 mmHg, which may contribute to tissue hypoxemia and cellular damage, with no organ specificity. Amyloid deposits, soluble amyloid aggregates and the precursor amyloidogenic proteins, the free light chains, have been implicated as the mediators for tissue injury.(Merlini, *et al* 2011) Assessment of LDH level in other forms of amyloidosis will enable a better understanding of tissue injury by amyloidosis subtype.

We have shown that LDH was an independent prognostic marker (overall survival and early death alike), but only in the non-ASCT cohort. Non-ASCT patients tend to be more affected from the extent of organ deposition rather than the plasma cell compartment. The non-ASCT patient cohort is usually older, has a poorer performance status and more affected organs and the degree of organ dysfunction is greater.(Muchtar, *et al* 2017c) Therefore, organ repair may not occur once a haematological response is achieved. It is not uncommon to have a deep haematological response, without impact on organ damage. Therefore, LDH may act as a potential marker for tissue damage and amyloid burden and is more likely to have an impact on patients with advanced organ dysfunction.

This study has several limitations. It is a retrospective single-centre study, which impacts the generalizability of its findings. However, given the rarity of AL amyloidosis, this study provides the first dataset on LDH in AL amyloidosis. We lacked data for several variables in 10–30% of patients, owing to the study period extending back to 2000, a time in which cardiac biomarkers and dFLC were not routinely performed. Disease presentation and management have changed considerably over the years, and this requires caution when interpreting the data. Nevertheless, the multivariate analysis included variables associated with severity of disease as well as type of treatment employed. Finally, our data interpretation and conclusions are based on inference rather than causality. However, we believe that our analysis justifies the conclusions.

In summary, we have demonstrated that serum LDH is elevated in 40% of newly diagnosed AL amyloidosis. The source of elevation is probably tissue damage caused by the amyloid deposits, with no clear organ source. Elevated serum LDH above the normal reference values is an independent prognostic marker in the non-ASCT population, predicting both overall survival and early death. The clinical significance of elevated LDH should be further investigated as a marker for tissue healing and organ recovery. Measuring LDH in non-ASCT candidates is an easy way to assess for a poor prognosis. ASCT may be able to overcome the poor prognosis associated with LDH due to the higher associated haematological responses seen with stem cell transplantation.

## Acknowledgments

The study was supported by the Jabbs foundation (Birmingham, UK), the Henry J. Predolin Foundation (USA) and the Specialized Programs of Research Excellence (SPORE), grant CA186781 from the National Cancer Institute, National Institutes of Health, USA.

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#### Figure 1.

Overall survival Kaplan-Meier curves by serum LDH groups. A. Whole cohort. B. Non-ASCT cohort C. ASCT cohort.

ASCT, autologous stem cell transplant; CI, confidence interval; LDH, Lactate dehydrogenase;

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## Table I

Baseline characteristics of the entire cohort and serum LDH groups

Characteristic	Entire cohort (n=1019)	Normal serum LDH (n=610)	High serum LDH (n=409)	Р
Age, years; Median (IQR)	62 (55-69)	61 (54–68)	(54–68) 63 (55–70)	
Male sex, N (%)	633 (62%)	403 (66%)	230 (56%)	0.001
Systolic blood pressure <100mmHg	226 (22%)	121 (20%)	105 (26%)	0.02
Organ involved, N (%)				
>2 organs	283 (28%)	151 (25%)	132 (32%)	0.008
Cardiac	744 (73%)	405 (66%)	339 (83%)	<0.001
Renal	645 (63%)	348 (57%)	297 (73%)	<0.001
Hepatic	217 (21%)	114 (19%)	103 (25%)	0.01
Nerve	194 (19%)	137 (22%)	57 (14%)	<0.001
GI	168 (16%)	108 (18%)	60 (15%)	0.19
Lambda restricted, N (%)	755 (74%)	480 (79%)	275 (67%)	<0.001
2004 Mayo AL amyloidosis stage, % (n=729)				
I / II / III	23 / 37 / 40	32/44/24	11 / 29 / 60	<0.001
2012 Mayo AL amyloidosis stage, % (n=787)				
I / II / III / IV	25 / 24 / 22 / 29	30 / 31 / 21 / 18	17 / 15 / 25 / 43	<0.001
First line treatment, N (%)				
ASCT	406 (40%)	289 (47%)	118 (29%)	
MDex	272 (27%)	120 (20%)	152 (37%)	
MP	104 (10%)	69 (11%)	35 (9%)	
Bortezomib-based	59 (6%)	31 (5%)	28 (7%)	<0.001
IMiD-based	45 (4%)	30 (5%)	15 (4%)	
Dexamethasone alone	32 (3%)	20 (3%)	12 (3%)	
No treatment	71 (7%)	18 (3%)	6(1%)	
Unknown	24 (2%)	30 (5%)	41 (10%)	
Other	5 (<1%)	3 (<1%)	2 (<1%)	

Abbreviations: ASCT, Autologous stem cell transplantation, GI, Gastrointestinal; IMiD, Immunomodulatory drug; IQR, interquartile range; LDH, Lactate dehydrogenase; MDex, melphalan-dexamethasone; MP, melphalan-prednisone

## Table II

Laboratory evaluation in the whole cohort and by serum LDH groups

	Entire cohort (n=1019)	Normal serum LDH (n=610)	High serum LDH (n=409)	Р
Plasma cell compartment evaluation				
BMPCs, median (IQR)	9 (5–15)	8 (5–14)	9 (5–15)	0.52
dFLC, mg/dl, median (IQR) (n=888)	23 (9–57)	22 (8–57)	24 (10-60)	0.26
Labelling index (n=668)	0 (0-0.4)	0 (0–0.3)	0(0-0.4)	0.42
Significant immunoparesis, N (%) (n=928)	335 (36%)	212 (38%)	123 (33%)	0.11
Organ function evaluation				
Haemoglobin, g/l median (IQR)	131 (119–144)	132 (119–144)	131 (118–146)	0.94
Serum albumin, g/l, median (IQR)	29 (23–33)	30 (24–34)	26 (19–32)	<0.001
Thyroid stimulating hormone, u/ml, median (IQR)	2.8 (1.8-4.5)	2.6 (1.7-4.2)	3.1 (2.1–4.7)	<0.001
Cardiac amyloidosis population				
NT-proBNP, pg/ml, median (IQR)	3766 (1333–8934)	2385 (999–5128)	6164 (2166–12691)	<0.001
Troponin T, ng/ml, median (IQR)	0.04 (0.01-0.1)	0.02 (<0.01-0.05)	0.08 (0.03-0.15)	<0.001
Renal amyloidosis population				
eGFR, ml/min/1.73 m <sup>2</sup> , median (IQR)	60 (41–78)	66 (49–81)	53 (30–70)	<0.001
Proteinuria, g/24 h, median (IQR)	4.8 (1.8-8.2)	4.1 (1.8–7.4)	5.3 (2.0–9.4)	0.008
Hepatic amyloidosis population				
Alkaline phosphatase, xUNL, median (IQR)	2.4 (1.7–4.5)	2.4 (1.6–4.2)	2.6 (1.7–4.8)	0.14
Total bilirubin, µmol/l, median (IQR)	13.7 (8.6–25.7)	13.7 (8.6–23.9)	13.7 (8.6–29.1)	0.71

Abbreviations: BMPCs, bone marrow plasma cells; dFLC, difference between involved and uninvolved light chains; eGFR, estimated glomerular filtration rate; IQR, interquartile range; LDH, Lactate dehydrogenase; NT-proBNP, N-terminal pro b-type natriuretic peptide; ULN, upper limit of normal;.

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Table III

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	Mod	el A (Whole	cohort)	Model B	(non-ASCT	cohort)
Prognostic factor	HR	95% CI	Р	HR	95% CI	Р
Age ≻65 years	1.4	1.1 - 1.7	0.003	1.4	1.1–1.8	0.002
Male sex	1.3	1.1–1.7	0.006	1.2	0.95-1.6	0.11
>2 involved organs	1.1	0.8–1.3	0.66	0.9	0.7–1.1	0.31
dFLC 18 mg/dl	1.3	1.03-1.6	0.02	1.2	0.95–1.6	0.11
BMPCs 10%	1.1	0.9–1.3	0.46	1.0	0.8–1.3	0.98
Serum LDH above ULN	1.2	0.97–1.5	0.08	1.3	1.02–1.7	0.03
eGFR <30 ml/min/1.73 m <sup>2</sup>	0.8	0.6–1.1	0.25	0.8	0.5–1.04	0.08
Proteinuria >2 g/24 h	0.8	0.6-0.98	0.03	0.8	0.6–1.01	0.06
Troponin T 0.025 ng/ml	1.4	1.1 - 1.9	0.008	1.6	1.2–2.2	0.003
NT-proBNP 1800 pg/ml	1.9	1.4–2.5	<0.001	1.9	1.4–2.6	<0.001
Systolic blood pressure <100 mmHg	1.5	1.2–1.9	<0.001	1.4	1.1–1.8	0.01
ASCT as primary therapy (vs non-ASCT regimen)	0.4	0.3-0.5	<0.001	Z	Vot included	
Treatment category MDex				Reference		
<b>Bortezomib-based</b>		Not includ	po	0.7	0.5 - 1.1	0.12
IMiD-based			10	1.3	0.8 - 1.9	0.27
MP/Dex alone				1.4	0.9–2.1	0.14
No treatment				5.4	3.6-7.9	<0.001

Br J Haematol. Author manuscript; available in PMC 2018 September 01.

type natriuretic peptide; SBP, systolic blood pressure; ULN, upper limit of normal.

#### Table IV

Univariate and multivariate analysis for predictor of early death ( 6 months) in the non-ASCT cohort

	Univariate a	nalysis	Multivariate analysis	
	OR (95% CI)	Р	OR (95% CI)	Р
Age 65 years	0.8 (0.6–1.1)	0.11	0.9 (0.6–1.4)	0.65
>2 organs involved	1.9 (1.4–2.7)	<0.001	1.4 (0.9–2.3)	0.14
SBP <100 mmHg	3.1 (2.1-4.5)	<0.001	1.6 (0.9–2.7)	0.08
dFLC 18 mg/dl	2.3 (1.5–3.5)	<0.001	1.7 (1.0–2.9)	0.05
NT-proBNP 1800 pg/ml	8.5 (4.4–18.6)	<0.001	5.8 (2.6–14.9)	<0.001
Troponin T 0.025 ng/ml	5.3 (3.3-8.8)	<0.001	2.0 (1.03-4.0)	0.03
LDH above ULN	1.6 (1.2–2.2)	0.005	1.8 (1.1–3.0)	0.02

Abbreviations: ASCT, autologous stem cell transplant; CI, confidence interval; dFLC, difference between involved and uninvolved light chains; eGFR, estimated glomerular filtration rate; LDH, Lactate dehydrogenase; NT-proBNP, N-terminal pro b-type natriuretic peptide; OR, odd ratio; SBP, Systolic blood pressure; ULN, upper limit of normal.