

SUPPLEMENTARY INFORMATION

**Predictors of early morbidity and mortality in newly diagnosed multiple myeloma:
data from five randomized, controlled, phase III trials in 3700 patients**

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SUPPLEMENTARY TABLES

Table S1. Characteristics and treatments in the trials included in the multi-cohort analysis.

	GMMG-HD4/HOVON-65	GMMG-MM5	GMMG-HD6	EMN02/HOVON-95	GMMG-HD7
Recruitment period	05/2005–05/2008	07/2010–11/2013	06/2015–09/2017	02/2011–04/2014	10/2018–09/2020
Induction regimen	Arm B: PAD bortezomib i.v. 1.3 mg/m ² on days 1, 4, 8, 11 / doxorubicin i.v. 9 mg/m ² on days 1–4 / dexamethasone p.o. 40 mg on days 1–4, 9–12, 17–20; repeated on day 29 (3x)	Arms A1+B1: PAd bortezomib i.v./s.c. 1.3 mg/m ² on days 1, 4, 8, 11 / doxorubicin i.v. 9 mg/m ² on days 1–4 / dexamethasone p.o. 20 mg on days 1–4, 9–12, 17–20; repeated on day 29 (3x)	Arms A1+A2: RVd lenalidomide p.o. 25 mg on days 1–14 / bortezomib s.c. 1.3 mg/m ² on days 1, 4, 8, 11 / dexamethasone 20 mg on days 1–2, 4–5, 8–9, 11–12; repeat on day 22 (4x)	VCD: bortezomib i.v. 1.3 mg/m ² on days 1, 4, 8, 11 / cyclophosphamide i.v. 500 mg/m ² on days 1 and 8 / dexamethasone p.o. 40 mg on days 1–2, 4–5, 8–9, 11–12; repeated on day 22 (3x or 4x)	Arm IA: RVd lenalidomide p.o. 25 mg on days 1–14 and 22–35 / bortezomib s.c. 1.3 mg/m ² on days 1, 4, 8, 11, 22, 25, 29, 32 / dexamethasone 20 mg on days 1–2, 4–5, 8–9, 11–12, 15, 22–23, 25–26, 29–30, 32–33; repeat on day 43 (3x)

Anti-bacterial prophylaxis^a	Recommended	Mandatory	Mandatory	Recommended	Recommended
N of patients per induction treatment	192/218 (PAD only)	596 (PAd: 296; VCD: 300)	545 (RVd: 272; elotuzumab-RVd: 273)	1491 (VCD only)	658 (RVd: 328; isatuximab-RVd: 330)
CTCAE version	3.0	4.0	5.0	4.0	5.0
Trial registry	EudraCT No. 2004-000944-26	EudraCT No. 2010-019173-16	NCT 02495922	NCT 01208766	NCT 03617731

CTCAE National Cancer Institute Common Terminology Criteria for Adverse Events, *EMN* European Myeloma Network, *GMMG* German-speaking Myeloma Multicenter Group, *HOVON* Dutch–Belgian Cooperative Trial Group for Hematology Oncology Foundation.

^aRecommendations according to study protocols – HD4/HO-65: cotrimoxazol; MM5: cotrimoxazol or ciprofloxacin; HD6: cotrimoxazol or ciprofloxacin; EMN02/HO-95: not specified/according to local protocols; HD7: cotrimoxazol.

Table S2. Patient demographics and baseline characteristics in the trials included in the training cohort.

n (%)		GMMG-HD4	GMMG-MM5	GMMG-HD6	All
Evaluable	Yes	192 (100)	596 (100)	545 (100)	1333 (100)
Completed IT	Yes	180 (93.8)	567 (95.1)	514 (94.3)	1261 (94.6)
Median age	Years (range)	55 (31–65)	58 (32–70)	58 (32–70)	58 (31–70)
Age > 60 years	No	140 (72.9)	347 (58.2)	312 (57.2)	799 (59.9)
	Yes	52 (27.1)	249 (41.8)	233 (42.8)	534 (40.1)
	Missing	-	-	-	-
Sex	Female	121 (63.0)	348 (58.4)	309 (56.7)	778 (58.4)
	Male	71 (37.0)	248 (41.6)	236 (43.3)	555 (41.6)
	Missing	-	-	-	-
WHO performance status	0–1	177 (92.2)	533 (90.5)	496 (91.0)	1206 (91.0)
	> 1	15 (7.8)	56 (9.5)	49 (9.0)	120 (9.0)
	Missing	-	7	-	7
Body mass index	≤ 30 kg/m ²	162 (84.4)	488 (81.9)	438 (81.0)	1088 (81.9)
	> 30 kg/m ²	30 (15.6)	108 (18.1)	103 (19.0)	241 (18.1)
	Missing	-	-	4	4
White blood cell count	≥ 4.0/nL	169 (88.0)	487 (81.7)	436 (80.0)	1092 (81.9)
	< 4.0/nL	23 (12.0)	109 (18.3)	109 (20.0)	241 (18.1)
	Missing	-	-	-	-
Hemoglobin	≥ 10.0 g/dL	121 (63.0)	394 (66.1)	385 (70.6)	900 (67.5)
	< 10.0 g/dL	71 (37.0)	202 (33.9)	160 (29.4)	433 (32.5)
	Missing	-	-	-	-
Platelets	≥ 150/nL	172 (89.6)	520 (87.2)	489 (89.7)	1181 (88.6)
	< 150/nL	20 (10.4)	76 (12.8)	56 (10.3)	152 (11.4)
	Missing	-	-	-	-
Creatinine	≤ 2.0 mg/dL	173 (90.1)	530 (88.9)	512 (93.9)	1215 (91.2)
	> 2.0 mg/dL	19 (9.9)	66 (11.1)	33 (6.1)	118 (8.8)
	Missing	-	-	-	-
Calcium	≤ 2.75 mmol/L	176 (92.6)	554 (93.0)	525 (96.3)	1255 (94.3)
	> 2.75 mmol/L	14 (7.4)	42 (7.0)	20 (3.7)	76 (5.7)
	Missing	2	-	-	2
C-reactive protein	≤ 5.0 mg/L	120 (65.2)	340 (57.7)	351 (65.0)	811 (61.8)

	> 5.0 mg/L	64 (34.8)	249 (42.3)	189 (35.0)	502 (38.2)
	Missing	8	7	5	20
Lactate dehydrogenase	Normal	151 (83.4)	489 (82.3)	451 (83.2)	1091 (82.8)
	> ULN	30 (16.6)	105 (17.7)	91 (16.8)	226 (17.2)
	Missing	11	2	3	16
International Staging System	I/II	141 (78.3)	433 (72.7)	423 (77.6)	997 (75.5)
	III	39 (21.7)	163 (27.4)	122 (22.4)	324 (24.5)
	Missing	12	-	-	12
Cytogenetics	Standard risk	126 (75.5)	387 (73.3)	322 (72.0)	835 (73.1)
	High risk	41 (24.6)	141 (26.7)	125 (28.0)	307 (26.9)
	Missing	25	68	98	191
Anti-bacterial prophylaxis	No	21 (35.6)	67 (11.2)	9 (1.6)	97 (8.1)
	Yes	38 (64.4)	529 (88.8)	536 (98.3)	1103 (91.9)
	Missing	133	-	-	133

GMMG German-speaking Myeloma Multicenter Group, *IT* induction therapy, *ULN* upper limit of normal, *WHO* World Health Organization.

Table S3. Study endpoints in the training cohort.

n (%)	GMMG-HD4	GMMG-MM5	GMMG-HD6	All
Evaluable	192 (100)	596 (100)	545 (100)	1333 (100)
Severe infection (grade ≥ 3)	52 (27.1)	60 (10.1)	46 (8.4)	158 (11.8)
Death	4 (2.1)	8 (1.3)	12 (2.2)	24 (1.8)
Severe infection/death	53 (27.6)	64 (10.7)	50 (9.2)	167 (12.5)
Cause of death	(n = 4)	(n = 8)	(n = 12)	(n = 24)
Infection	2 (50.0)	4 (50.0)	9 (75.0)	15 (62.5)
Multiple myeloma	1 (25.0)	2 (25.0)	2 (16.7)	5 (20.8)
Thromboembolic event	-	2 (25.0)	1 (8.3)	3 (12.5)
Unknown	1 (25.0)	-	-	1 (4.2)

GMMG German-speaking Myeloma Multicenter Group.

Table S4. Odds ratio and 95% confidence interval estimates comparing risk score stages for endpoints severe infection, death, and severe infection/death in the training cohort.

	Odds ratio	95% Confidence interval	p
Severe infection			
High vs low risk	3.40	1.99–5.79	< 0.001
High vs intermediate risk	2.25	1.39–3.66	< 0.001
Intermediate vs low risk	1.51	0.91–2.50	0.06
Death			
High vs low risk	15.67	2.68–91.56	< 0.001
High vs intermediate risk	4.70	1.58–13.99	< 0.001
Intermediate vs low risk	3.33	0.51–21.57	0.13
Severe infection/death			
High vs low risk	3.63	2.15–6.13	< 0.001
High vs intermediate risk	2.34	1.46–3.76	< 0.001
Intermediate vs low risk	1.55	0.94–2.55	0.04

Comparison of risk groups for endpoints severe infection, death, or combined endpoint of severe infection/death.

Table S5. Distribution of baseline and risk factors in the validation cohorts.

n (%)		HOVON-65	EMN02/HOVON-95	GMMG-HD7
Evaluable	Yes	218 (100)	1491 (100)	658 (100)
Completed IT	Yes	197 (90.3)	1340 (89.9)	605 (92.0)
Age > 60 years	No	157 (72.0)	970 (65.1)	372 (56.5)
	Yes	61 (28.0)	521 (34.9)	286 (43.5)
	Missing	-	-	-
WHO performance status	0–1	184 (84.8)	1270 (85.2)	592 (90.2)
	> 1	33 (15.2)	221 (14.8)	62 (9.8)
	Missing	1	-	2
Platelets	≥ 150/nL	189 (87.1)	1263 (85.9)	592 (90.0)
	< 150/nL	28 (12.9)	208 (14.1)	66 (10.0)
	Missing	1	20	-
International Staging System	I/II	153 (78.9)	1163 (78.0)	516 (78.2)
	III	41 (21.1)	328 (22.0)	142 (21.6)
	Missing	24	-	-
Anti-bacterial prophylaxis	No	27 (25.7)	361 (24.5)	16 (2.4)
	Yes	78 (74.3)	1112 (75.5)	642 (97.6)
	Missing	113	18	-

EMN European Myeloma Network, *GMMG* German-speaking Myeloma Multicenter Group, *HOVON* Dutch–Belgian Cooperative Trial Group for Hematology Oncology Foundation, *IT* induction therapy, *WHO* World Health Organization.

Table S6. Study endpoints in the validation cohorts.

n (%)	HOVON-65	EMN02/HOVON-95	GMMG-HD7
Evaluable	218	1491	658
Severe infection (grade ≥ 3)	56 (25.7)	101 (6.8)	77 (11.7)
Death	13 (6.0)	25 (1.7)	12 (1.8)
Severe infection/death	61 (28.0)	118 (7.9)	81 (12.3)

EMN European Myeloma Network, *GMMG* German-speaking Myeloma Multicenter Group, *HOVON* Dutch–Belgian Cooperative Trial Group for Hematology Oncology Foundation.

Table S7. Odds ratio and 95% confidence interval estimates comparing risk score stages for endpoints severe infection, death, and severe infection/death in the HO65 trial.

	Odds ratio	95% Confidence interval	p
Severe infection			
High vs low risk	3.12	1.15–8.45	0.007
High vs intermediate risk	2.23	0.80–6.20	0.07
Intermediate vs low risk	1.40	0.56–3.53	0.39
Death			
High vs low risk	6.67	0.89–49.86	0.03
High vs intermediate risk	1.49	0.33–6.70	0.54
Intermediate vs low risk	4.47	0.64–31.38	0.07
Severe infection/death			
High vs low risk	3.24	1.21–8.65	0.005
High vs intermediate risk	1.96	0.72–5.31	0.11
Intermediate vs low risk	1.65	0.68–4.02	0.19

Comparison of risk groups for endpoints severe infection, death, or combined endpoint of severe infection/death.

Table S8. Odds ratio and 95% confidence interval estimates comparing risk score stages for endpoints severe infection, death, and severe infection/death in the EMN02/HO95 trial.

	Odds ratio	95% Confidence interval	p
Severe infection			
High vs low risk	2.36	1.25–4.46	0.002
High vs intermediate risk	1.48	0.83–2.66	0.11
Intermediate vs low risk	1.59	0.87–2.91	0.07
Death			
High vs low risk	5.11	1.05–24.97	0.02
High vs intermediate risk	1.17	0.40–3.43	0.73
Intermediate vs low risk	4.37	0.96–19.79	0.02
Severe infection/death			
High vs low risk	2.56	1.41–4.66	< 0.001
High vs intermediate risk	1.50	0.87–2.59	0.08
Intermediate vs low risk	1.70	0.97–3.01	0.03

Comparison of risk groups for endpoints severe infection, death, or combined endpoint of severe infection/death.

Table S9. Odds ratio and 95% confidence interval estimates comparing risk score stages for endpoints severe infection, death, and severe infection/death in the HD7 trial.

	Odds ratio	95% Confidence interval	p
Severe infection			
High vs low risk	3.38	1.62–7.04	< 0.001
High vs intermediate risk	2.47	1.25–4.90	0.002
Intermediate vs low risk	1.37	0.66–2.82	0.31
Death			
High vs low risk	5.08	0.71–36.52	0.05
High vs intermediate risk	2.56	0.52–12.54	0.17
Intermediate vs low risk	1.98	0.26–15.19	0.43
Severe infection/death			
High vs low risk	3.68	1.78–7.62	< 0.001
High vs intermediate risk	2.49	1.28–4.84	0.001
Intermediate vs low risk	1.48	0.73–3.02	0.20

Comparison of risk groups for endpoints severe infection, death, or combined endpoint of severe infection/death.

SUPPLEMENTARY FIGURE

Fig. S1. Logistic regression analyses of factors influencing risk of severe infection, death, or combined endpoint of severe infection/death during induction therapy for each trial included in the training cohort.

A Forest plots on factors associated with risk of severe infection. **B** Forest plots on factors influencing risk of death. **C** Forest plots on factors influencing risk of severe infection and/or death. Analyses are presented for each trial in the training cohort (HD4, MM5, and HD6) and overall. All logistic regression analyses including the complete training cohort accounted for trial effects. *P* values from univariable analysis were adjusted for multiple testing.

BMI body mass index, *CI* confidence interval, */SS* International Staging System, *LDH* lactate dehydrogenase, *ULN* upper limit of normal, *WHO* World Health Organization.

Fig. 1A

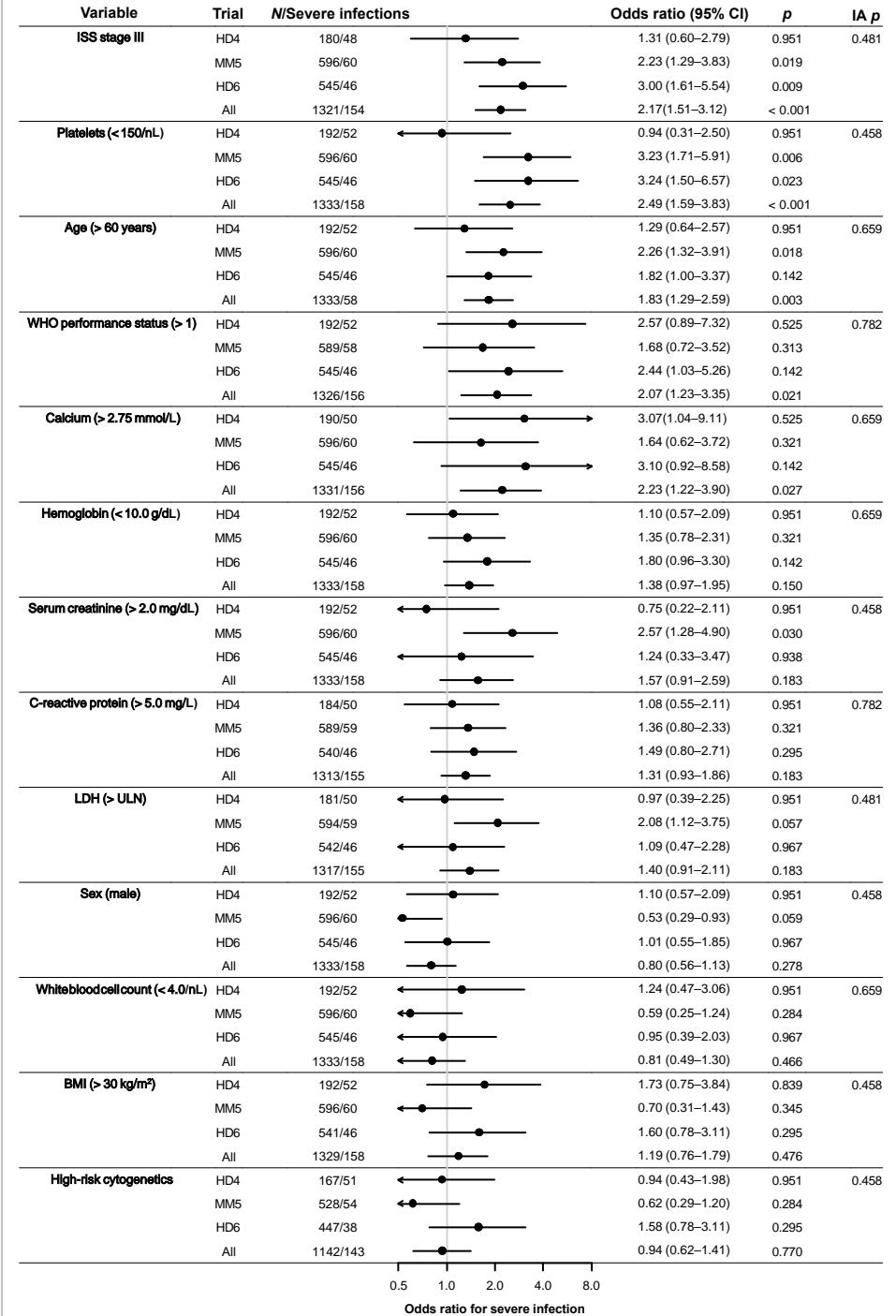


Fig. 1B

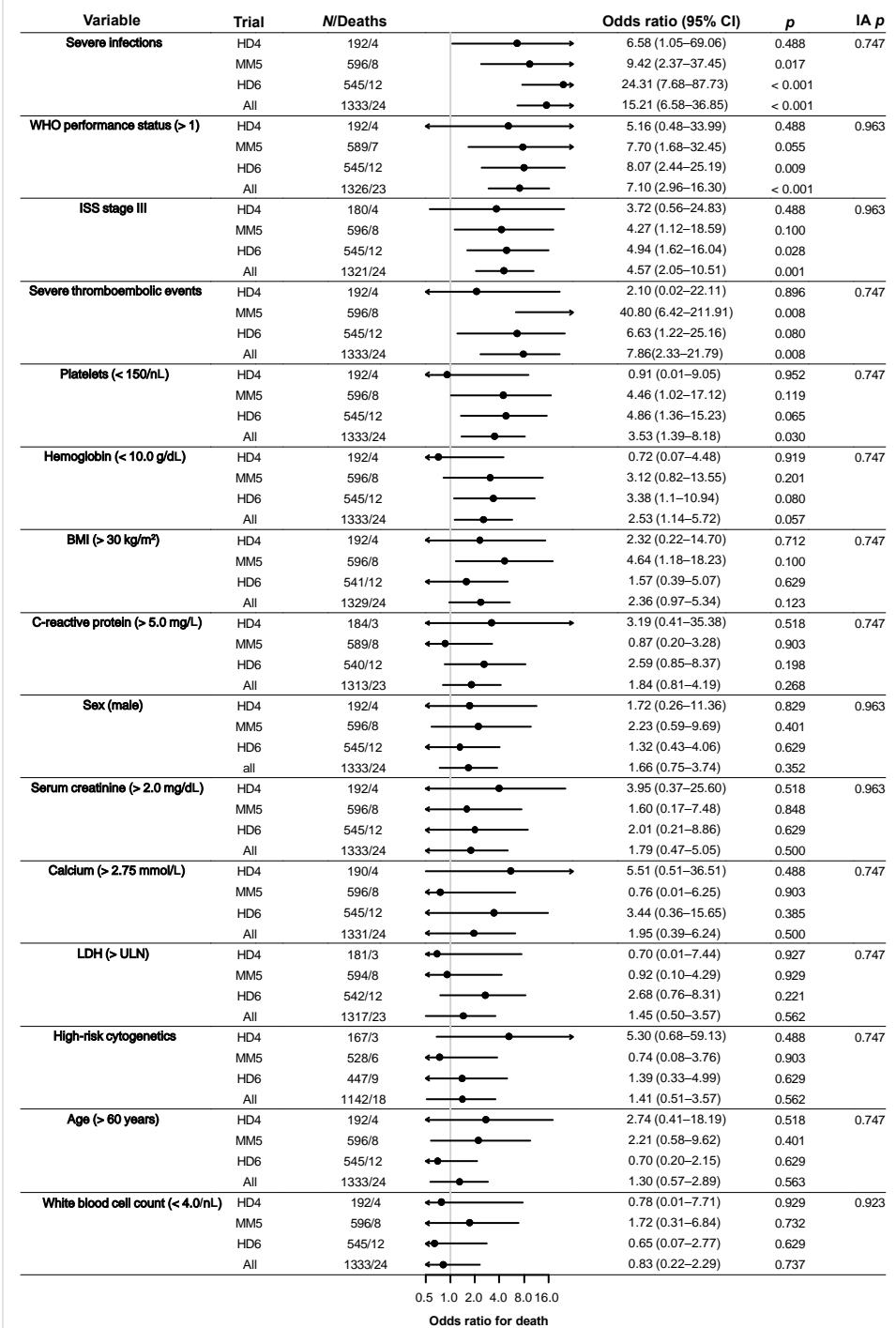


Fig. 1C

