The Geriatric Prognostic Index: a clinical prediction model for survival of older diffuse large B-cell lymphoma patients treated with standard immunochemotherapy

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Supplemental Methods

Study design and patients

The Cancer Registry of Norway (CRN) has an estimated 98.8% completeness on diagnosis and receives data on patients' vital status from the Norwegian Population Registry, and prospectively reported clinical and treatment features from treating physicians.¹ In the current study, patients with primary central nervous system lymphoma or prior lymphoproliferative disease were excluded, while a concurrent diagnosis of indolent lymphoma was allowed.

The division into training- and test set was based on geography/hospitals and independent investigators collecting the data. This was done to create a robust, external validation design that allowed for non-random variation between the training- and test set.²

In this study, an exception from informed consent was granted by the Norwegian Regional Health Research Ethics Committee (REK 2017/1861) based on the high age of the patients and the potential benefit for future patients.

Definition of full-dose and attenuated R-CHOP

The cutoff for attenuated R-CHOP was set at an initial "intended" dosage of 80% or lower, in line with cutoffs used in previous studies on older DLBCL patients.³⁻⁷ Doxorubicin was used to define R-CHOP dosage as it is considered a key component of the regimen and dose reduction of cyclophosphamide is seldom done without a concurrent reduction in doxorubicin. The initial dosage was used to define treatment intensity, rather than the mean dose per course or accumulated dose as dose reductions during therapy may have many causes like toxicity and disease progression, especially in older patients.

In the training set, four patients had received R-COP and etoposide (R-CEOP) in the first treatment cycle. Here, treatment intensity was defined by the initial dosage of etoposide and cyclophosphamide.

Candidate predictors

Data on candidate predictors were retrieved from the CRN and through review of clinical records of all patients to quality-check data, collect missing data and obtain information not routinely reported to the CRN.

Geriatric Nutritional Risk Index (GNRI) is an adaption to elderly of the Nutritional Risk Index and consists of albumin and estimated weight loss calculated from current and ideal weight, and has been validated in older DLBCL patients.^{8,9} Heart disease included heart failure, coronary artery disease, cardiac arrhythmia, operated valve disease or an implanted pacemaker. Heart failure was defined as a diagnosis of heart failure or clinical or radiological signs of heart failure (ejection fraction <50% measured with echocardiography or multigated acquisition (MUGA) scan) documented in clinical records at the time of diagnosis. Coronary artery disease included prior myocardial infarction, percutaneous coronary intervention, bypass surgery and angina pectoris.

Candidate predictors that were collected as part of the study, but were not included in model development due to a high fraction of missing values were "double hit" status detected by fluorescence in situ hybridization (FISH) for MYC, BCL2 and BCL6, double-protein expression of MYC and BCL2, and EBV positivity. The reason for the high fraction of missing values for double-hit status was that FISH for MYC, BCL2 and BCL6 was not routinely performed for patients in the time period for the training cohort (2006-2016), especially not for older patients who would not be candidates for further treatment intensification.

Outcome variables

In the training set, time of death was retrieved from the CRN, while data on progression, relapse and causes of death were retrieved from clinical records. Date of diagnosis was registered as the day the diagnosis was confirmed by the pathologist. Date of progression or relapse was retrieved from clinical records by the investigators and registered as the date when there was a biopsy-confirmed relapse or progression, radiological findings or a strong clinical suspicion of progression or relapse, whichever came first.

Cause of death was registered retrospectively by the investigators and divided into the following categories: lymphoma, treatment-related toxicity, other non-lymphoma related cause and unknown cause. Death from treatment-related toxicity included deaths occurring during or shortly after treatment where the death was considered likely to have been caused by acute treatment toxicity, or later deaths that were likely a result of long term toxicity.

Statistical methods and model development

Patient characteristics were compared using the χ^2 test or Fisher's exact test when indicated for categorical variables, and Mann-Whitney U test for continuous variables when comparing two groups and Kruskal-Wallis test when comparing three groups.

In the training set, the few missing values were assumed to be missing at random (MAR) and imputed using multivariate imputation by chained equations (MICE) creating 21 imputed datasets.¹⁰ To prevent bias in the results, the outcome was not included in the imputation model. For simplicity, the 21 multiply imputed datasets were combined into one dataset during the model-selection process by using the mean of the imputed continuous variables and the most frequent imputed category for categorical variables for each patient with missing values. For the external test set, no imputations of missing values were performed.

For the multivariable Cox models, model selection with stepwise backward elimination was performed using the stepAIC function in the MASS package in R. Candidate predictors showing a high degree of correlation (Spearman correlation >0.90) were examined in separate models (lymphocyte/monocyte ratio (log) and monocyte/lymphocyte ratio). For the final model, dichotomization of continuous variables (age, albumin and LDH) were examined. For age, a cutoff at 80 years decreased model performance and the continuous definition was kept. For albumin, two categorical cutoffs were examined: a cutoff at the median (38 g/L) and a cutoff at 36 g/L as this cutoff had been used in other studies on the same patient population.¹¹⁻¹⁴ LDH was examined as a continuous predictor and categorized as in NCCN-IPI.

For the final Cox model, the model without imputed data was used for estimation of regression coefficients and calculation of the Geriatric prognostic index (GPI), as there were only a few missing observations (n=10) for the variables included in the final model.

The proportional-hazards assumption for the final model was checked using Schoenfeld residuals and showed violation for stage (p=0.009), while the global test for the model showed no significant violation of the proportional-hazards assumption (p=0.11).¹⁵

Division into risk groups

When creating 3 risk groups from the Geriatric prognostic index (GPI), three different divisions were examined: cutoffs at the 33th and 66th percentile, at the 25th and 75th percentile and at the 30th and 80th percentile. Further exploration of cutoffs, including an optimal cutoff, was not examined to avoid overfitting of the model to our data. The cutoffs at the 30th and 80th percentile were chosen as unequal group sizes enabled identification of a high-risk group with a more extreme prognoses and a relatively large low-risk group with a very favorable prognosis.

Model performance

Model performance was assessed with discrimination and calibration. Discrimination measures the models' ability to distinguish between patients who experience the outcome or not, and was quantified using Harrell's C-index. Calibration measures the agreement between observed and predicted survival, and was assessed with calibration slope and calibration plots.

Internal validation

Internal validation was performed by applying the final model to 200 bootstrap resamples of the training set with the validate function in the rms package in R. The bootstrap resampling technique simulates the process of repeatedly sampling from an underlying patient population to assess the likely model overfit and correct for the resulting optimism in model performance.² For simplicity, optimism was estimated for the final model, not including all modelling steps, and thus the optimism-corrected performance might still include some optimism.

In the training set, this resulted in an optimism-corrected C-index of 0.752, and an optimismcorrected calibration slope of 0.89. A calibration slope <1 indicated some anticipated optimism in the model. When assessing calibration in a calibration plot comparing predicted and observed 2-year overall survival (OS), the model showed fairly good concordance between observed and predicted survival (Figure S2).

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Supplemental Tables

Table S1. Univariate Cox regression analyses for the association between candidate predictors and 2-year overall survival in the training set.

Candidate predictors	n (%)	HR (95% CI)	р
Age, years, continuous	365	1.04 (1.00-1.08)	0.06
Age group	365		
70-79 years	273 (75%)	1	
≥80 years	92 (25%)	1.41 (0.97-2.04)	0.07
ADL	365		
Independent	327 (90%)	1	
Dependent	38 (10%)	3.06 (1.98-4.75)	5.5e-07
ССІ	365		
0-1	254 (70%)	1	
≥2	111 (30%)	2.41 (1.70-3.41)	6.5e-07
Polypharmacy	365		
<5 regular medications	249 (68%)	1	
≥5 regular medications	116 (32%)	1.33 (0.93-1.91)	0.12
BMI, continuous	361	0.98 (0.94-1.02)	0.31
GNRI	360		
Absent	159 (44%)	1	
Low	96 (27%)	1.98 (1.20-3.27)	0.008
Moderate/severe	105 (29%)	5.05 (3.25-7.84)	5.3e-13
Albumin g/L, continuous	360	0.91 (0.89-0.93)	6.5e-14
Albumin	360		
≥36 g/L	230 (64%)	1	
<36 g/L	130 (36%)	3.93 (2.74-5.63)	9.3e-14
ECOG PS	362		
ECOG 0-1	244 (67%)	1	
ECOG ≥2	118 (33%)	3.28 (2.31-4.66)	3.2e-11
Stage	365		
1/11	161 (44%)	1	
III/IV	204 (56%)	2.40 (1.63-3.54)	9.9e-06
Extranodal sites	365		
Extranodal sites 0-1	280 (77%)	1	
Extranodal sites >1	85 (23%)	1.34 (0.91-1.97)	0.137
Bone marrow, liver or lung infiltration*	360		
No	293 (81%)	1	
Yes	67 (19%)	1.89 (1.28-2.80)	0.00135
Sex	365		
Female	178 (49%)	1	
Male	187 (51%)	1.44 (1.01-2.05)	0.0412

Bulky disease (≥7 cm)	352		
No	223 (63%)	1	
Yes	129 (37%)	1.64 (1.16-2.33)	0.00543
B-symptoms	361		
No	227 (63%)	1	
Yes	134 (37%)	2.17 (1.53-3.08)	1.3e-05
Heart failure	365		
No	338 (93%)	1	
Yes	27 (7%)	1.52 (0.86-2.69)	0.154
Hypertension	365		
No	178 (49%)	1	
Yes	187 (51%)	1.26 (0.89-1.78)	0.194
Coronary artery disease	365		
No	281 (77%)	1	
Yes	84 (23%)	1.13 (0.76-1.69)	0.548
Heart disease**	365		
No	238 (65%)	1	
Yes	127 (35%)	1.14 (0.80-1.63)	0.473
Cell-of-origin (IHC)	273		
GCB	156 (57%)	1	
Non-GCB	117 (43%)	1.00 (0.65-1.53)	0.995
Ki67 (IHC), continuous	339	1.00 (0.99-1.01)	0.626
Ki67 (IHC), median	339		
≤80%	119 (35%)	1	
>80%	220 (65%)	0.88 (0.60-1.28)	0.495
BCL2 positive (cutoff 1%) (IHC)	319		
No	44 (14%)	1	
Yes	275 (86%)	1.13 (0.63-2.02)	0.678
CD5 positive (IHC)	304		
No	280 (92%)	1	
Yes	24 (8%)	1.54 (0.80-2.96)	0.197
LDH, U/L, continuous (log transformed)	361	1.89 (1.56-2.29)	5.2e-11
LDH	361		
Not elevated	179 (50%)	1	
Elevated	182 (50%)	2.25 (1.55-3.26)	1.9e-05
LDH	361		
Not elevated	179 (50%)	1	
Elevated x 1-3 x ULN	151 (42%)	1.86 (1.25-2.76)	0.00208
Elevated >3 x ULN	31 (8%)	5.11 (3.06-8.52)	4.04e-10
Hb, g/dL, continuous	365	0.83 (0.77-0.91)	2.4e-05
Lymphocytes, x10^9/L (log transformed)	359	0.52 (0.40-0.67)	8.9e-07
Monocytes, x10^9/L, continuous	356	1.62 (0.98-2.69)	0.061
Neutrophils, x10^9/L (log transformed)	360	1.90 (1.35-2.67)	0.000226

LMR, continuous (log transformed)	356	0.51 (0.40-0.66)	2.2e-07
MLR, continuous	356	2.00 (1.56-2.56)	3.2e-08
NLR, continuous (log transformed)	359	1.71 (1.43-2.04)	3.2e-09
eGFR, mL/min/1,73m ² , continuous	363	0.99 (0.98-1.00)	0.0036
CRP, mg/L, continuous (log transformed)	348	1.45 (1.28-1.64)	2.3e-09
ALAT, U/L, continuous (log transformed)	353	1.08 (0.82-1.43)	0.581

Abbreviations: ADL, Activities of daily living; ALAT, alanine aminotransferase; BMI, body mass index; CCI, Charlson comorbidity index; CI, confidence interval; CRP, C-reactive protein; ECOG, Eastern Cooperative Oncology Group; ; eGFR, estimated glomerular filtration rate; GCB, germinal center B-cell like; GNRI, Geriatric Nutritional Risk Index; Hb, hemoglobin; HR, hazard ratio; IHC, immunohistochemistry; LDH, lactate dehydrogenase; LMR, lymphocyte/monocyte ratio; MLR, monocytes/lymphocyte ratio; NLR, neutrophile/lymphocytes ratio; PS, performance status; ULN, upper limit of normal. *Polypharmacy: \geq 5 regular medications vs <5 regular medications. **See Figure S1 for details on Cox univariate analyses for spesific extranodal sites. ***Includes heart failure, coronary artery disease, cardiac arrhythmia, operated valve disease or an implanted pacemaker. Further details are provided in the supplemental text. **Table S2**. Results from model selection in the training set using Cox multivariable models for 2year overall survival (OS) and stepwise backward elimination with Akaike's information criterion (AIC) as stopping criterion. **(A)** Prognostic model with missing variables imputed with multivariate imputation by chained equations. **(B)** Prognostic model in the training set without imputed data.

	Cox multivariable model with imputed data (n=365)					
	β	SE	HR (95% CI)	Р		
Years >70 years	0.04117	0.02053	1.04 (1.00-1.09)	0.045		
ADL dependent	0.71470	0.24282	2.04 (1.27-3.29)	0.003		
CCI ≥2	0.70756	0.18842	2.03 (1.40-2.94)	< 0.001		
GNRI						
Low	0.35118	0.25815	1.42 (0.86-2.36)	0.174		
Moderate/Severe	0.92099	0.24916	2.51 (1.54-4.09)	< 0.001		
ECOG ≥2	0.36940	0.20784	1.45 (0.96-2.17)	0.076		
Stage III/IV	0.48950	0.21429	1.63 (1.07-2.48)	0.022		
Male	0.44087	0.18916	1.55 (1.07-2.25)	0.020		
LDH (log)	0.21997	0.12758	1.25 (0.97-1.60)	0.085		
NLR (log)	0.22009	0.10008	1.25 (1.02-1.52)	0.028		

A)

B)

-	Cox multivariable me	odel without im	puted data (n=349)	
	β	SE	HR (95% CI)	Р
Years >70 years	0.03821	0.02133	1.04 (1.00-1.08)	0.073
ADL dependent	0.69544	0.24798	2.00(1.23-3.26)	0.005
CCI ≥2	0.68431	0.19548	1.98 (1.35-2.91)	<0.001
GNRI				
Low	0.29455	0.26819	1.34 (0.79-2.27)	0.272
Moderate/Severe	0.86783	0.25823	2.38 (1.44-3.95)	<0.001
ECOG ≥2	0.46088	0.21774	1.59 (1.03-2.43)	0.034
Stage III/IV	0.43911	0.22171	1.55 (1.01-2.40)	0.048
Male	0.42021	0.19507	1.52 (1.04-2.23)	0.031
LDH (log)	0.20115	0.13318	1.22 (0.94-1.59)	0.131
NLR (log)	0.24016	0.10307	1.27 (1.04-1.56)	0.020

Abbreviations: ADL, Activities of daily living; CCI, Charlson comorbidity index; ECOG, Eastern Cooperative Oncology Group; GNRI, Geriatric Nutritional Risk Index; LDH, lactate dehydrogenase; NLR, neutrophil/lymphocyte ratio. β indicates the regression coefficient; SE, standard error; HR, hazard ratio; CI, confidence interval. **Table S3.** Overall survival (OS) and hazard ratio (HR) for the Geriatric prognostic index (GPI) groups in the training set when **(A)** restricted to patients receiving full-dose R-CHOP (initial R-CHOP dosage >80%) (n=231, number of events =59), and **(B)** restricted to patients receiving attenuated R-CHOP (initial R-CHOP dosage ≤80%) (n=124, number of events =63).

A)				
GPI risk group	n (%)	2-year OS (95% CI)	HR (95% CI)	Р
Low-risk	92 (40)	95 % (90-99)	1	
Intermediate risk	112 (48)	69 % (61-78)	6.72 (2.63-17.1)	<0.001
High-risk	27 (12)	30 % (17-53)	21.7 (8.09-58.5)	<0.001
High-risk vs intermediate risk			3.24 (1.85-5.68)	<0.001
В)				
GPI risk group	n (%)	2-year OS (95% CI)	HR (95% CI)	Р
Low-risk	16 (13)	88 % (73-100)	1	
Intermediate risk	64 (52)	58 % (47-71)	3.93 (0.93-16.5)	0.062
High-risk	44 (35)	18 % (10-34)	10.49 (2.51-43.8)	0.001
High-risk vs intermediate risk			2.67 (1.60-4.45)	<0.001

Survival estimated from Kaplan-Meier curves. HR estimated from Cox regression for the three GPI risk groups in the training set. Abbreviations: HR, hazard ratio; CI, confidence interval.

Characteristics	Low risk	Intermediate risk	High risk	p
	n=57 (%)	n=71 (%)	n=46 (%)	
Age, years, median (IQR)	76 (72-79)	78 (75-80)	78 (74-81)	0.031
Age ≥80 years	14 (25)	20 (28)	18 (39)	0.253
Stage III-IV	14 (25)	38 (54)	40 (87)	<0.001
ECOG ≥2	1 (2)	26 (37)	37 (80)	<0.001
ADL dependent	2 (4)	7 (10)	13 (28)	<0.001
CCI ≥2	0 (0)	14 (20)	22 (48)	<0.001
IPI				<0.001
Low (1)	29 (51)	10 (14)	1 (2)	
Low-intermediate (2)	18 (32)	32 (45)	5 (11)	
High-intermediate (3)	8 (14)	18 (25)	8 (17)	
High (4-5)	2 (4)	11 (16)	32 (70)	
R-IPI				
Good (1-2)	47	42	6	
Poor (3-5)	10 (18)	29 (41)	40 (87)	<0.001
NCCN-IPI				<0.001
Low-intermediate (2-3)	38 (67)	10 (14)	2 (4)	
High-intermediate (4-5)	19 (33)	48 (68)	15 (33)	
High (6-8)	0 (0.0)	13 (18)	29 (63)	
Frailty status*				<0.001
Fit	48 (91)	29 (45)	0 (0.0)	
Unfit	5 (9)	28 (44)	29 (64)	
Frail	0 (0.0)	7 (11)	16 (36)	
Missing	4	7	1	
Treatment intensity**				
R-CHOP >80%	50 (88)	47 (68)	15 (36)	
R-CHOP ≤80%	7 (12)	22 (32)	27 (64)	
Missing	0	2	4	<0.001

Table S4. Patient characteristics for the Geriatric prognostic index (GPI) groups in the test set.

Abbreviations: ADL, Activities of daily living; CCI, Charlson comorbidity index; ECOG Eastern Cooperative Oncology Group; IPI, International Prognostic Index; IQR, interquartile range; NCCN, the National Comprehensive Cancer Network; R-CHOP, rituximab, cyclophosphamide, doxorubicin, vincristine and prednisone; R-IPI, revised IPI. *Frailty status assessed with our previously published frailty calculator, Isaksen et al, Blood Advances 2021, https://wide.shinyapps.io/app-frailty/. **Treatment intensity defined by the initial dosage of R-CHOP. Further details are provided in the supplemental text.

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set.				
Table S5. Patient characteristic	cs for the Geriatric pr	rognostic index (GPI) groups in the training	

Characteristics	Low risk	Intermediate risk	High risk	р
	n=108 (%)	n=176 (%)	n=71 (%)	
Age, years, median (IQR)	75 [72, 78]	77 [74, 80]	77 [73, 81]	<0.001
Age ≥80 years	15 (14)	47 (27)	24 (34)	0.005
Stage III-IV (%)	27 (25)	111 (63)	60 (85)	<0.001
ECOG ≥2	3 (3)	54 (31)	59 (83)	<0.001
ADL dependent	0 (0)	10 (6)	27 (38)	<0.001
CCI ≥2	2 (2)	66 (38)	39 (55)	<0.001
IPI (%)				<0.001
Low (1)	61 (56)	28 (16)	2 (3)	
Low-intermediate (2)	27 (25)	45 (26)	0 (0)	
High-intermediate (3)	18 (17)	62 (35)	19 (27)	
High (4-5)	2 (2)	41 (23)	50 (70)	
R-IPI (%)				<0.001
Good (1-2)	88 (82)	73 (41)	2 (3)	
Poor (3-5)	20 (18)	103 (59)	69 (97)	
NCCN-IPI (%)				<0.001
Low-intermediate (2-3)	71 (66)	29 (16)	1 (1)	
High-intermediate (4-5)	35 (32)	112 (64)	17 (24)	
High (6-8)	2 (2)	35 (20)	53 (75)	
Frailty status (%)*				<0.001
Fit	103 (95)	62 (35)	1 (1)	
Unfit	4 (4)	105 (60)	36 (51)	
Frail	1 (1)	9 (5)	34 (48)	
Treatment intensity**				
R-CHOP >80%	92 (85)	112 (64)	27 (38)	
R-CHOP ≤80%	16 (15)	64 (36)	44 (62)	
COD (at 2 years follow-up)				<0.001
Censored (alive)	101 (94)	114 (65)	18 (25)	
Lymphoma	3 (3)	30 (17)	30 (42)	
Treatment-related toxicity	4 (4)	20 (11)	16 (23)	
Other cause	0 (0)	11 (6)	5 (7)	
Unknown	0 (0)	1 (1)	2 (3)	

Abbreviations: ADL, Activities of daily living; CCI, Charlson comorbidity index; COD, cause of death; ECOG Eastern Cooperative Oncology Group; IPI, International Prognostic Index; IQR, interquartile range; NCCN, the National Comprehensive Cancer Network; R-CHOP, rituximab, cyclophosphamide, doxorubicin, vincristine and prednisone; R-IPI, revised IPI. *Frailty status assessed with our previously published frailty calculator, Isaksen et al, Blood Advances 2021, https://wide.shinyapps.io/app-frailty/. **Treatment intensity defined by the initial dosage of R-CHOP. Further details are provided in the supplemental text.

Supplemental Figures

Figure S1. Forest plot showing results from univariate Cox regression analyses for association between specific extranodal sites and 2-year overall survival (OS) in the training set. HR, hazard ratio; CI, confidence interval.

Outcome	Extranodal site		HR	95% CI	р	n
2-year OS	Bone marrow (DLBCL/indolent clone)	·	2.07	(1.34–3.18)	<0.001	360
ſ	Bone marrow (DLBCL)	│ └ ─ ♦ ──i	2.35	(1.41–3.92)	0.001	360
1	Liver	│	2.18	(1.20–3.95)	0.010	365
ſ	Lung	└─ ◆──1	1.68	(1.00–2.84)	0.051	365
	Testis		0.48	(0.15–1.52)	0.215	365
ſ	Bone marrow (indolent clone)		1.48	(0.75–2.91)	0.260	360
ſ	Colorectal	⊢	1.50	(0.66–3.40)	0.333	365
ſ	CNS	▲ 	0.39	(0.05–2.76)	0.343	365
ſ	GI tractus	↓ ↓ ↓	1.18	(0.81–1.72)	0.398	365
ſ	Mammae	▶ ► ► ► ► ► ► ► ► ► ► ► ► ► ► ► ► ► ► ►	0.43	(0.06–3.10)	0.404	365
;	Small intestine	▶ •	0.88	(0.45–1.74)	0.721	365
ſ	Oral cavity/pharynx/esophagus	↓ ↓	0.80	(0.11–5.73)	0.825	365
ſ	Pancreas	▶ ► ► ► ► ► ► ► ► ► ► ► ► ► ► ► ► ► ► ►	0.87	(0.22–3.52)	0.846	365
1	Kidney	⊢	0.94	(0.38–2.30)	0.891	364
(Stomach		1.03	(0.59–1.80)	0.915	365
	Adrenal gland	• • • • • • • • • • • • • • • • • • •	0.95	(0.30–2.99)	0.931	364

Figure S2. Internal calibration. Calibration plot of the final prognostic model from 200 bootstrap resamples of the training set (n=355). The plot shows comparison of observed and predicted 2-year overall survival (OS) from 200 bootstrap resamples of the training set. The grey line represents perfect predictions where predicted 2-year OS is equal to observed 2-year OS estimated with the Kaplan-Meier method. The black points show mean predicted 2-year OS with corresponding 95% confidence interval for groups of patients with similar prognosis (about 70 cases per group). Crosses represent optimism-corrected 2-year OS. Histogram on top shows number of patients.



Figure S3. Progression-free survival of the Geriatric prognostic index (GPI) groups in the training set.



Figure S4. Overall survival of the Geriatric prognostic index (GPI) groups in the training set for patients treated with **(A)** full-dose R-CHOP (initial dosage >80%) and **(B)** attenuated R-CHOP (initial dosage ≤80%)



Figure S5. Overall survival of the Geriatric prognostic index (GPI) groups in the training set for patients aged **(A)** 70-79 years and **(B)** \ge 80 years.



Figure S6. Calibration of survival probabilities for the Geriatric prognostic index (GPI) groups in the test set.



The figure shows observed versus predicted survival for the GPI risk groups in the test set. The colored lines represent observed survival estimated with the Kaplan-Meier method and the smooth lines represent predicted mean survival for the GPI risk groups.