

## A laboratory-based scoring system predicts early treatment in Rai 0 chronic lymphocytic leukemia

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## Supplemental Materials for

### A Laboratory-Based Scoring System Predicts Early Treatment in Rai 0 Chronic Lymphocytic Leukemia

#### Supplemental information:

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**Supplemental Table 1.** Clinical and laboratory features of CLL patients.

<b>Parameter</b>	<b>n/total cases (%)</b>				
	<b>Training</b>	<b>Gemelli</b>	<b>Cardiff</b>	<b>Mayo</b>	<b>O-CLL</b>
Age $\geq$ 65, years	254/478 (53.1)	92/144 (63.9)	225/395 (60.0)	241/540 (44.6)	97/322 (30.1)
WBC count $>32 \times 10^3$ cell/ $\mu$ L	141/478 (29.5)	5/144 (3.5)	110/395 (27.8)	92/540 (17.0)	29/322 (9.0)
IGHV unmutated	94/478 (19.7)	35/144 (25.3)	83/395 (21.0)	178/540 (33.0)	97/322 (30.1)
FISH category					
del17p	30/478 (6.3)	3/144 (2.1)	17/395 (4.3)	18/540 (3.3)	6/322 (1.9)
del11q	36/478 (7.5)	6/144 (4.2)	32/395 (8.1)	38/540 (7.0)	18/322 (5.6)
tri12	67/478 (14.0)	14/144 (9.7)	39/395 (9.9)	59/540 (10.9)	38/322 (11.8)
Beta-2-microglobulin $>3.5$ mg/L	36/418 (8.6)	21/144 (14.6)	92/395 (23.3)	70/540 (13.0)	1/239 (0.4)
Follow-up median, months (95%CI)	62 (57-68)	91 (83-103)	94 (83-104)	77 (68-88)	89 (85-95)

FISH categories were as reported by Dohner et al (Dohner H, N Engl J Med. 2000;343:1910-6). Abbreviations: WBC, white blood cell.

**Supplemental Table 2.** Treatment free survival in CLL cases identified as high-risk according to the CRO model, stratified by score (N=309).

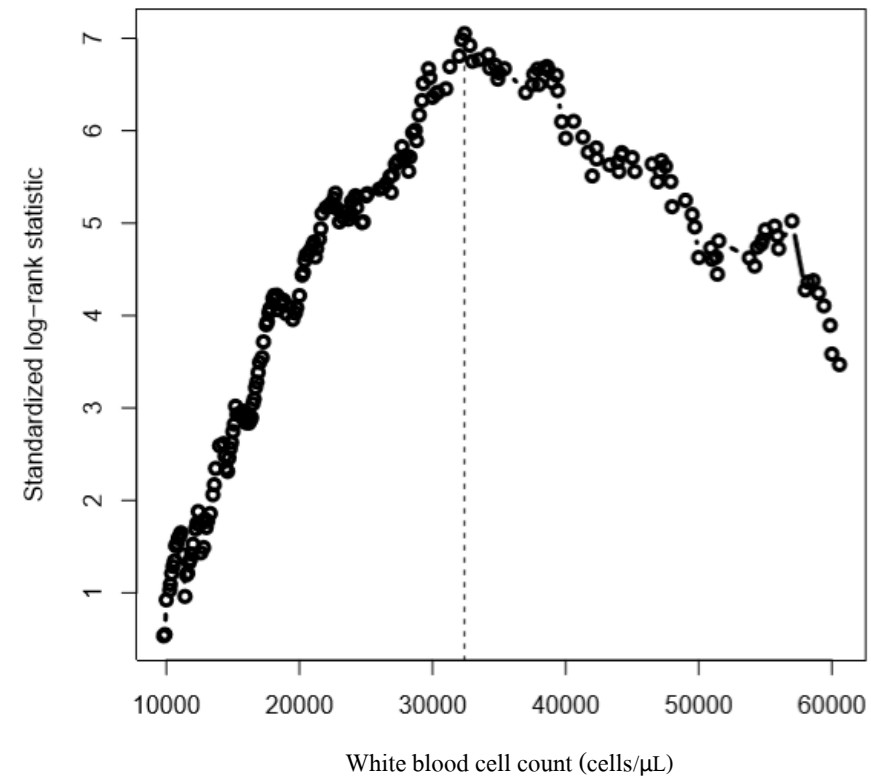
<b>CRO score</b>	<b>Median, months</b>	<b>95% CI, months</b>
3	57.6	45.9-68.0
4	44.0	36.0-64.0
5	28.0	14.7-29.1
Overall	51.6	42.6-60.0

CLL cases were taken from the training and validation cohorts. Abbreviations: CI, confidence interval.

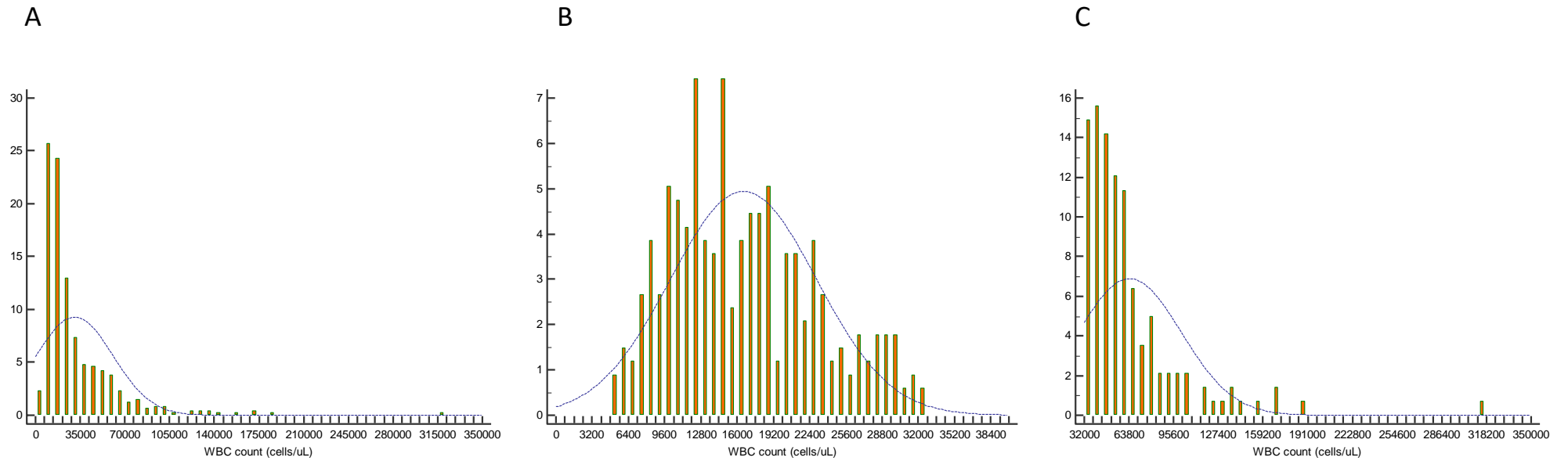
**Supplemental Table 3:** Subgroup univariable and reduced multivariable analysis involving 304/478 cases from the training cohort with available TP53 and NOTCH1 gene mutational status.

Factor	Univariable Analysis			Reduced Multivariable Analysis		
	P	HR	95% CI	P	HR	95% CI
WBC>32K cells/ $\mu$ L	<0.0001	2.69	1.80-4.00	0.0009	2.02	1.33-3.06
Dohner category						
del11q	<0.0001	4.25	2.41-7.49	0.0234	1.95	1.09-3.47
tri12	0.0001	2.75	1.67-4.53	0.0130	1.94	1.15-3.28
UM IGHV	<0.0001	3.98	2.64-5.98	<0.0001	2.57	1.64-4.04
TP53 disrupted	<0.0001	3.12	1.97-4.94	0.0086	2.01	1.94-3.39
Del17p	<0.0001	4.00	2.12-7.55			
TP53 mutated	<0.0001	3.33	2.07-5.38			
CD49d+	0.0022	1.85	1.25-2.75			
NOTCH1 mutated	0.0003	2.25	1.45-3.48			
Age >65 years	0.0818	1.42	0.96-2.12			
Male	0.9251	1.02	0.69-1.51			

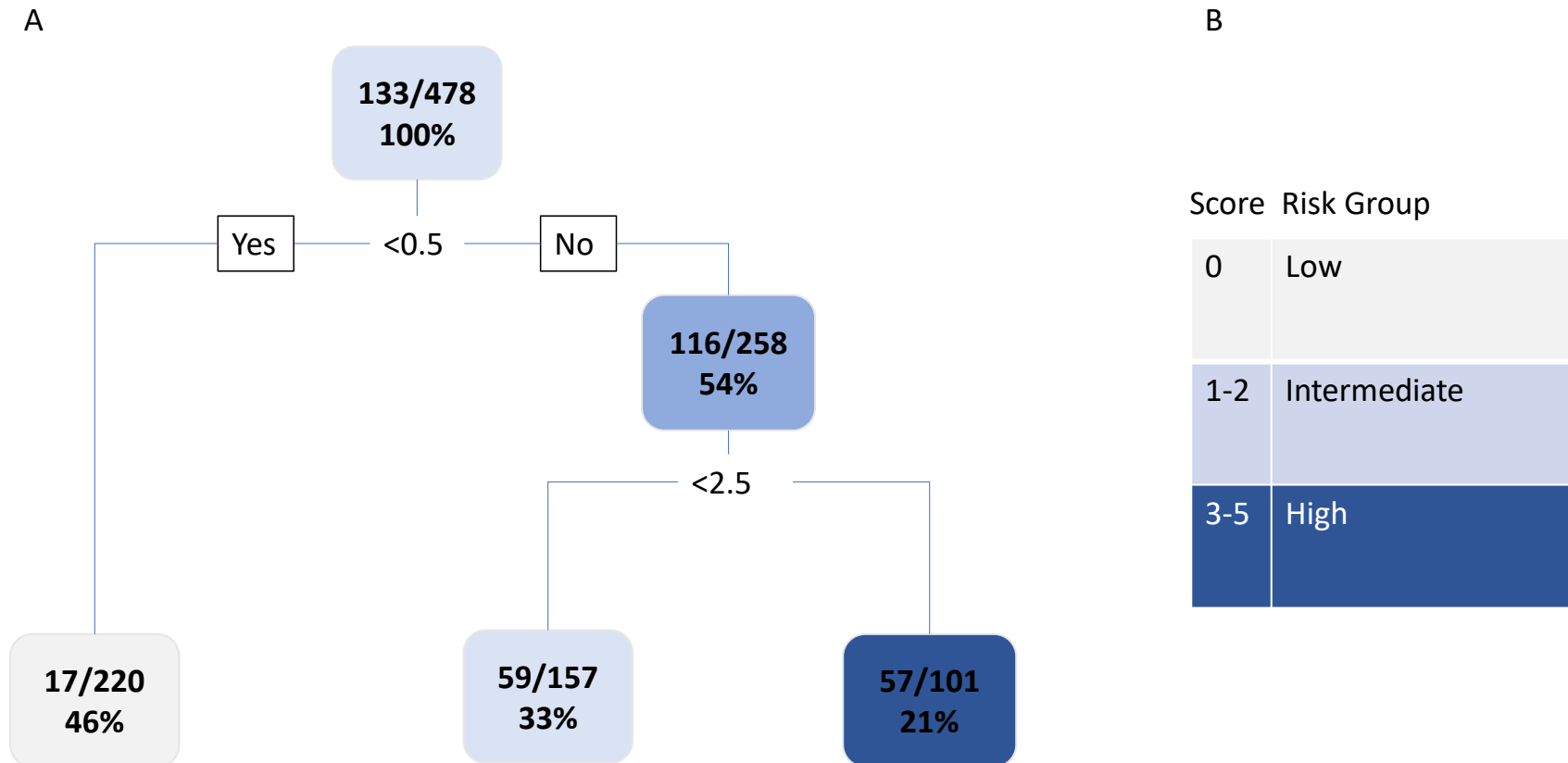
FISH categories were as reported by Dohner et al (ref. 7). Abbreviations: HR, hazard ratio; CI, confidence interval; WBC, white blood cell; UM, unmutated.



**Supplementary Figure 1. Establishing a WBC count cutoff.** Maximally selected log-rank statistic establishing a WBC count >32K cells/μL as our threshold.

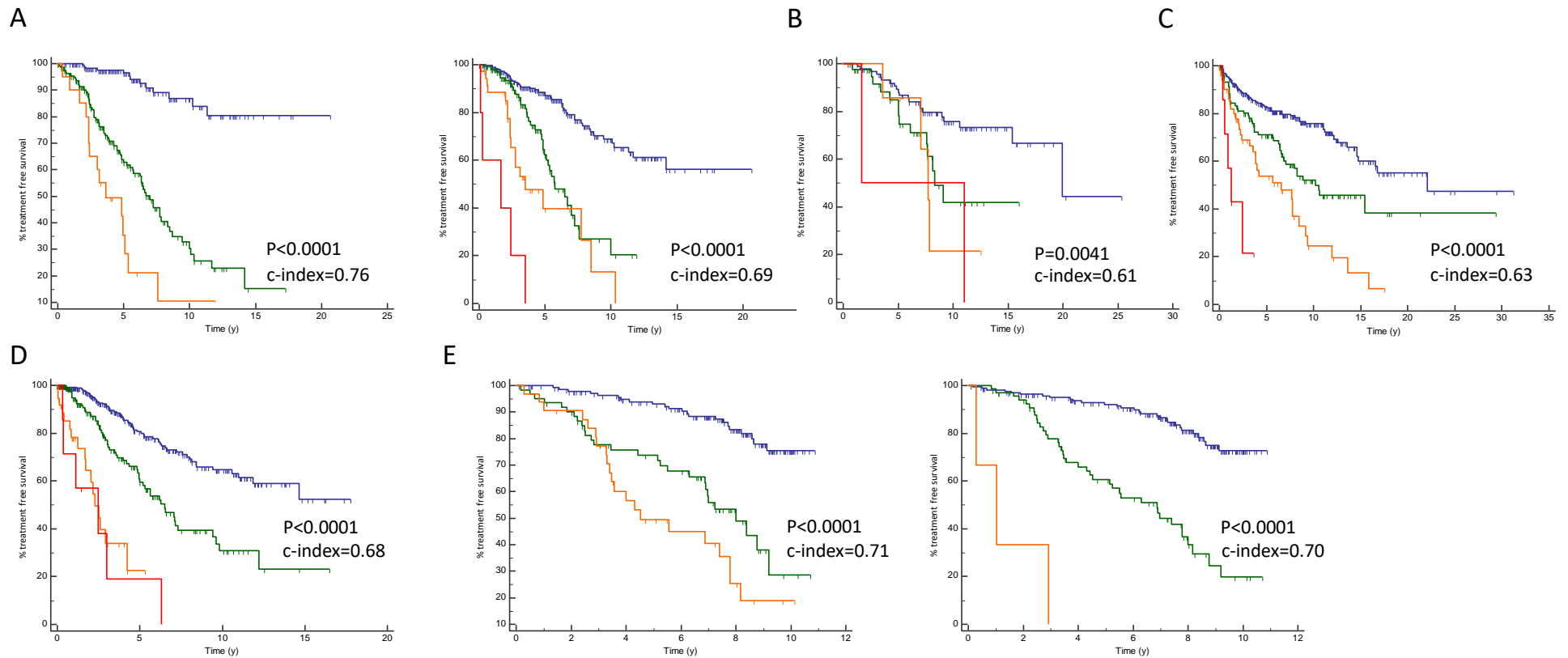


**Supplementary Figure 2. Distribution plots demonstrating WBC count frequencies in the training cohort.** Histogram of WBC count frequencies in (A) all 478 cases comprising the training cohort (B) cases with WBC count  $\leq 32 \times 10^3$  and (C) cases with WBC count  $> 32 \times 10^3$  cell/ $\mu$ L.



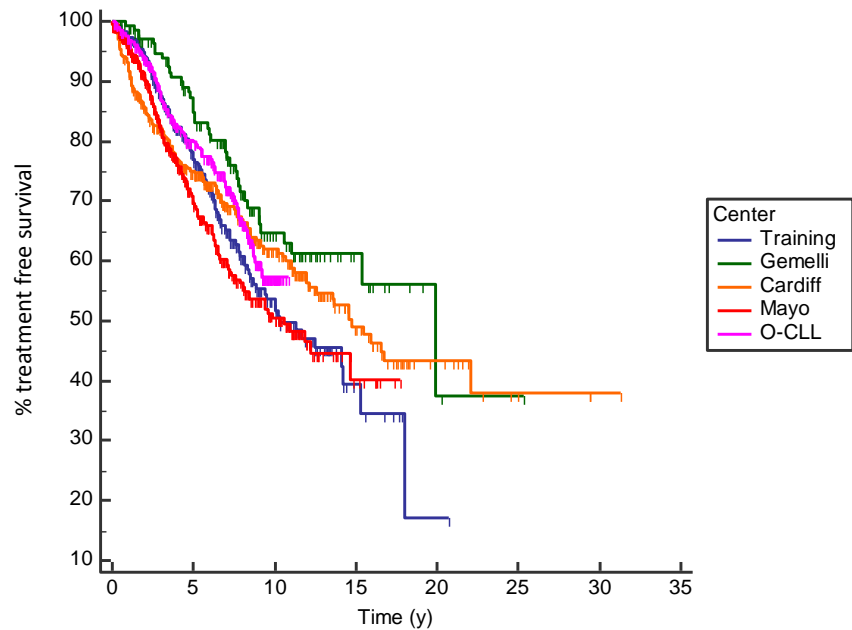
**Supplementary Figure 3. Risk group determination.** A) Recursive partitioning demonstrating 2 cut-offs at  $<0.5$  and  $<2.5$  (within nodes, # of events/cases and % of total cases reported). B) Risk group categorization into low (0) intermediate (1-2) and high (3-5).





**Supplementary Figure 4. Comparison to CLL-IPI.** Kaplan-Meier curves demonstrating treatment free survival (%; Y-axis) in: (A) the training cohort (left panel, CRO score; right panel, CLL-IPI); (B) the Gemelli validation cohort (CLL-IPI; the corresponding CRO score is reported in Figure 3A); (C) the Cardiff cohort (CLL-IPI; the corresponding CRO score is reported in Figure 3B); (D) the Mayo cohort (CLL-IPI) the corresponding CRO score is reported in Figure 3C); (E) the O-CLL cohort (left panel, CRO score; right panel, CLL-IPI). Blue: low-risk category, green: moderate-risk category, orange: high-risk category, red: very high-risk category (CLL-IPI only); x axis: time in years.

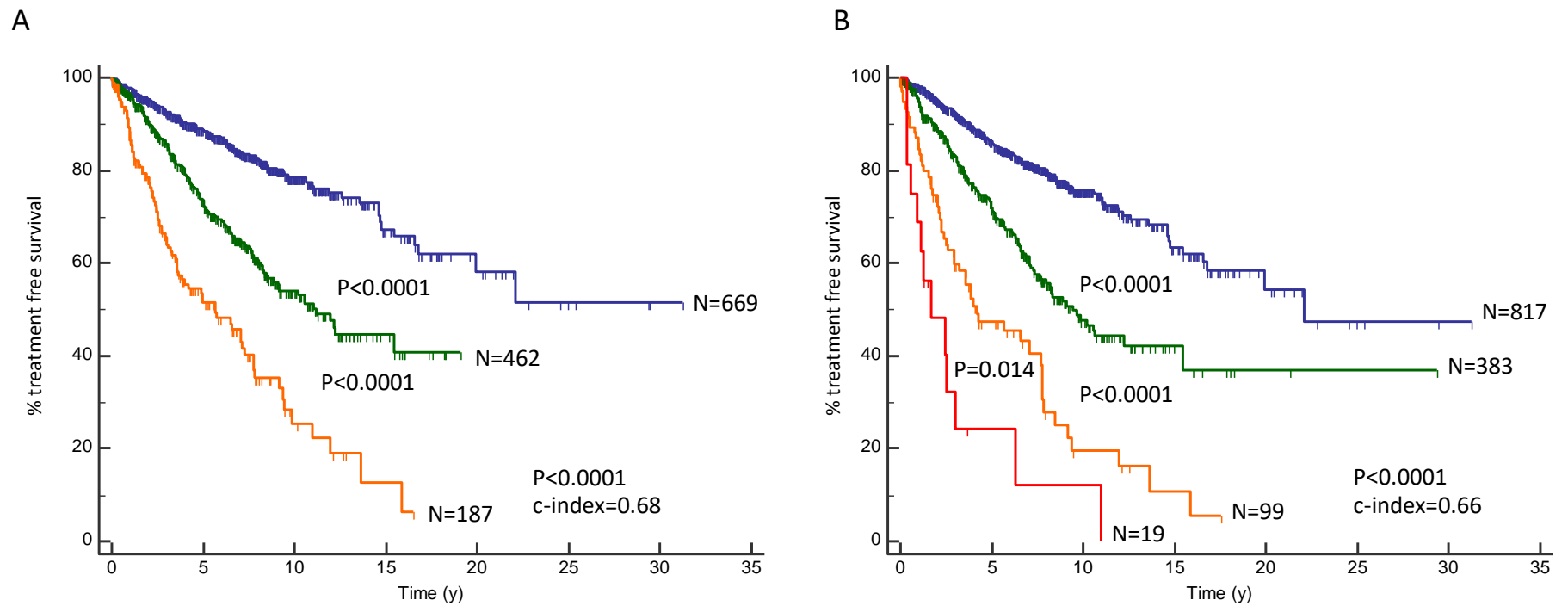
A



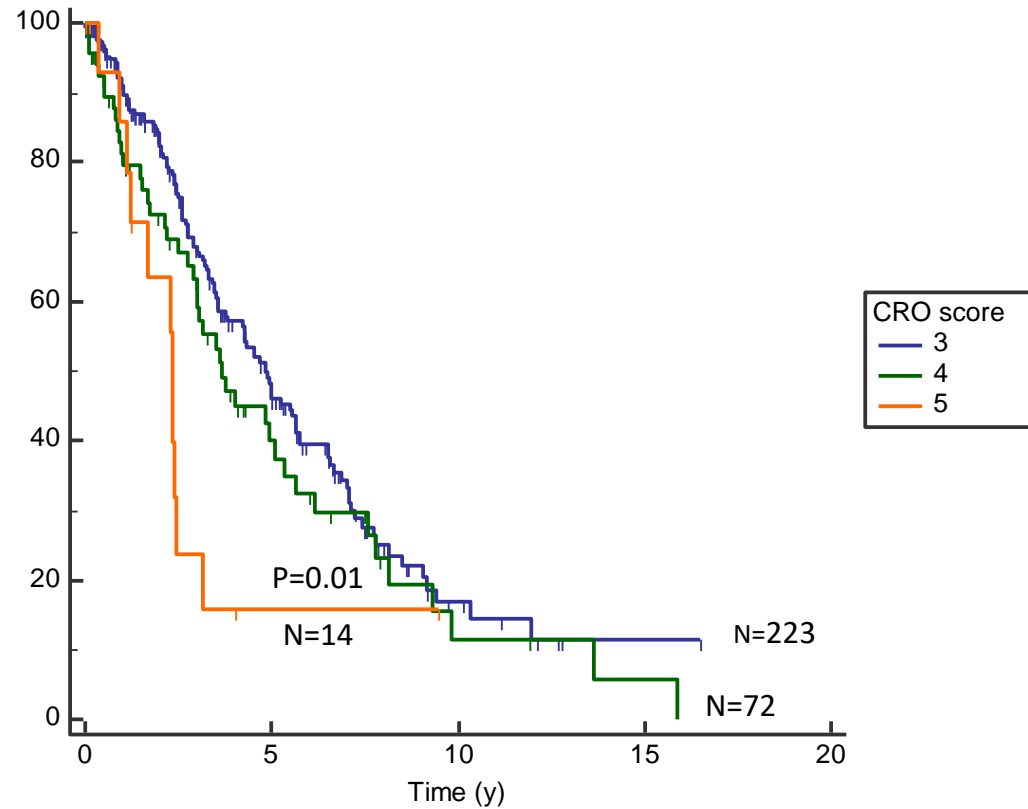
B

	Mayo	Training	O-CLL	Cardiff	Gemelli
Mayo	--	--	--	--	--
Training	0.15	--	--	--	--
O-CLL	0.01	0.25	--	--	--
Cardiff	0.12	0.55	0.66	--	--
Gemelli	0.0007	0.0047	0.15	0.06	--

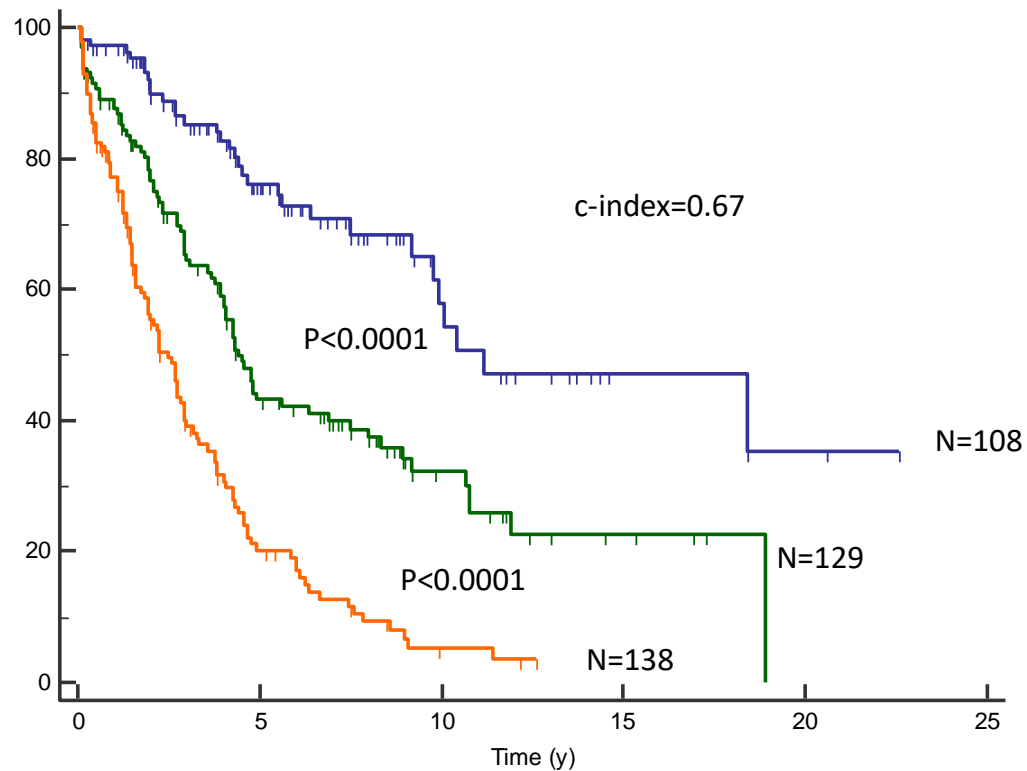
**Supplementary Figure 5. Overall treatment free survival in all cohorts.** A) Kaplan-Meier curves demonstrating treatment free survival (%; Y-axis) for the training and 4 validation cohorts. B) Statistical significance (P-value) comparing median treatment free survivals between all cohorts.



**Supplementary Figure 6. Comparison of CRO score and CLL-IPI in the composite cohort.** Kaplan-Meier curves demonstrating treatment free survival (%; Y-axis) in the composite validation cohort determined by the CRO score (A) and CLL-IPI (B). Blue: low-risk category, green: moderate-risk category, orange: high-risk category, red: very high-risk category (CLL-IPI only).



**Supplementary Figure 7. Treatment free survival in all high-risk cases determined according to the CRO score.** Kaplan-Meier curve demonstrating treatment free survival (%; Y-axis) stratified by CRO score in all high risk cases (N=309) taken from the training and validation cohorts.



**Supplementary Figure 8. Application of the CRO score in Rai 1 patients.** Kaplan-Meier survival curves demonstrating treatment free survival (%; Y-axis) using the CRO score in Rai I patients applied to a consecutive series of Italian multicenter patients referred to our center for immunocytogenetic analyses between 2006 and 2017. Blue: low-risk category, green: moderate-risk category, orange: high-risk category.