



IOR
Un istituto
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Sviluppo di Prognostic Scoring Systems

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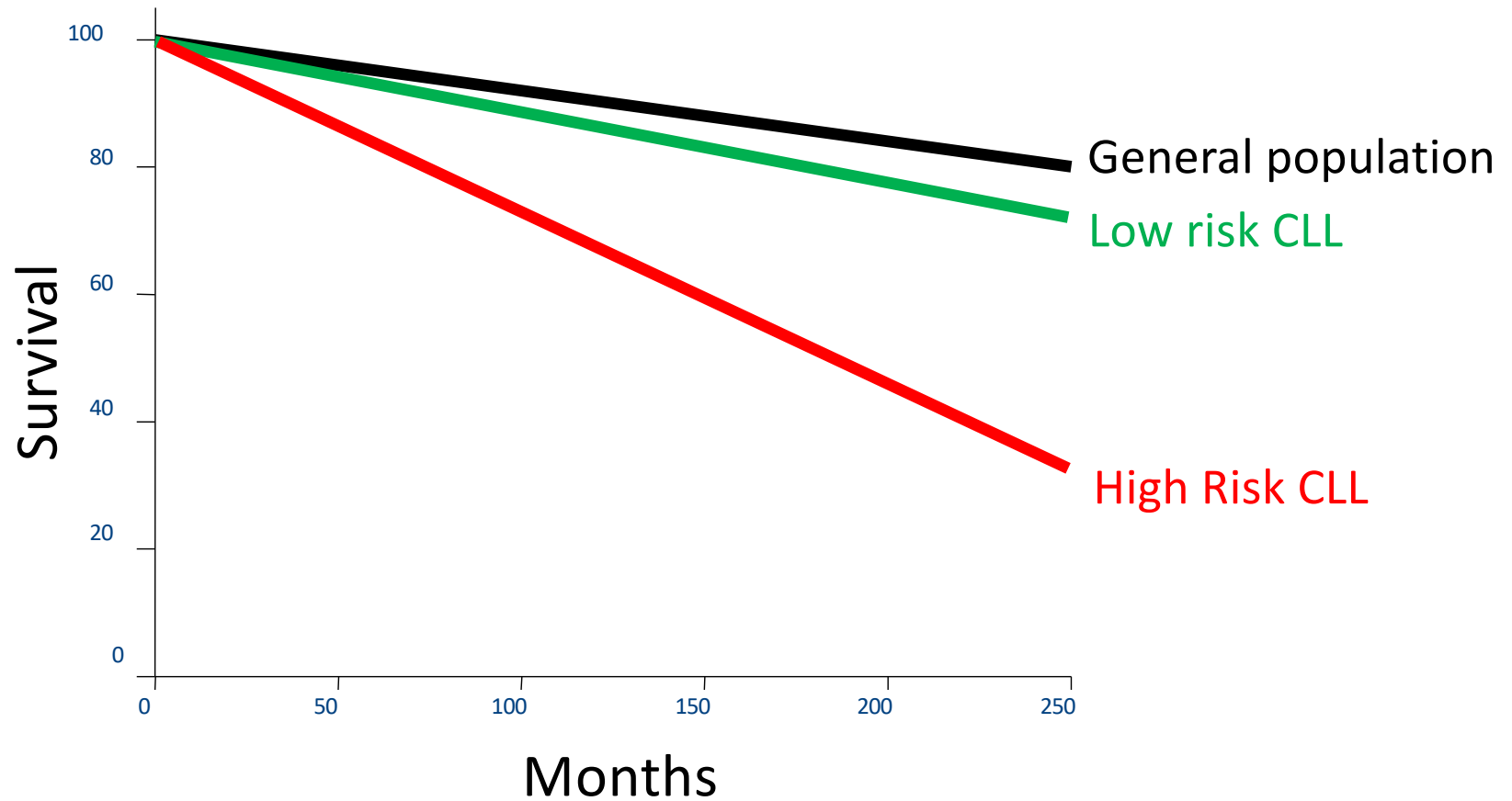
IOR - Institute of Oncology Research

USI – Università' della Svizzera Italiana

Bellinzona - Switzerland

Overall survival

CLL: Homogeneous phenotype but heterogeneous clinical course

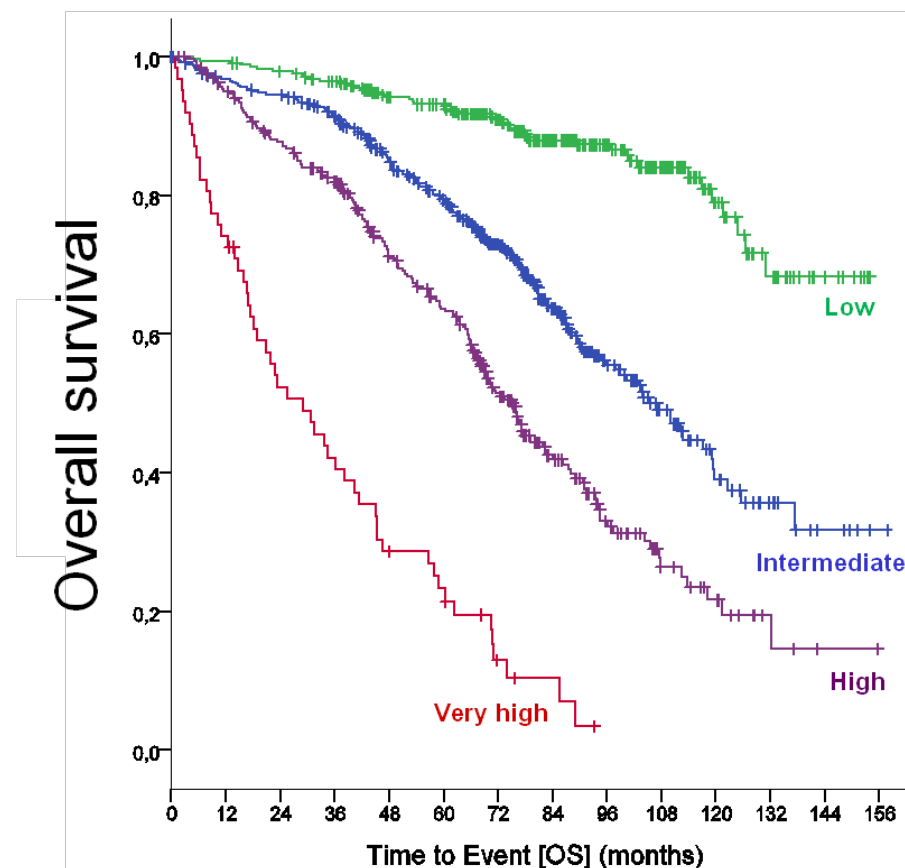


Prognostic scoring system of survival for CLL treated with chemoimmunotherapy



Variable	Adverse factor	Coeff.	HR	Grading
TP53 (17p)	deleted and/or mutated	1.442	4.2	4
IGHV status	Unmutated	0.941	2.6	2
B2M, mg/L	> 3.5	0.665	2.0	2
Clinical stage	Binet B/C <u>or</u> Rai I-IV	0.499	1.6	1
Age	> 65 years	0.555	1.7	1
Prognostic Score				0 – 10

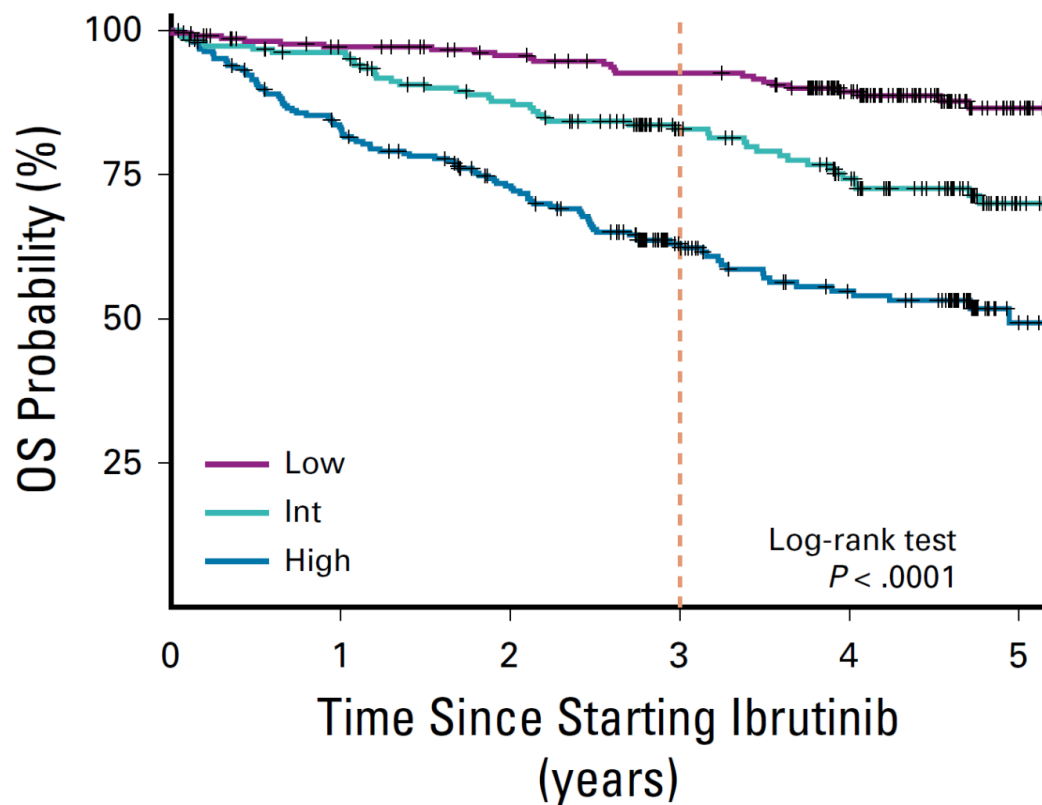
Risk group	Score	Patients N (%)	5-year OS, %
Low	0 – 1	340 (29)	93.2
Intermediate	2 – 3	464 (39)	79.4
High	4 – 6	326 (27)	63.6
Very High	7 – 10	62 (5)	23.3



Prognostic scoring system of survival for CLL treated with ibrutinib

Variable	Points
TP53 aberration	1
Prior treatment	1
B2M ≥ 5 mg/l	1
LDH ≥ 250 U/l	1

Risk group	Score
Low risk	0-1
Intermediate risk	2
High risk	3-4

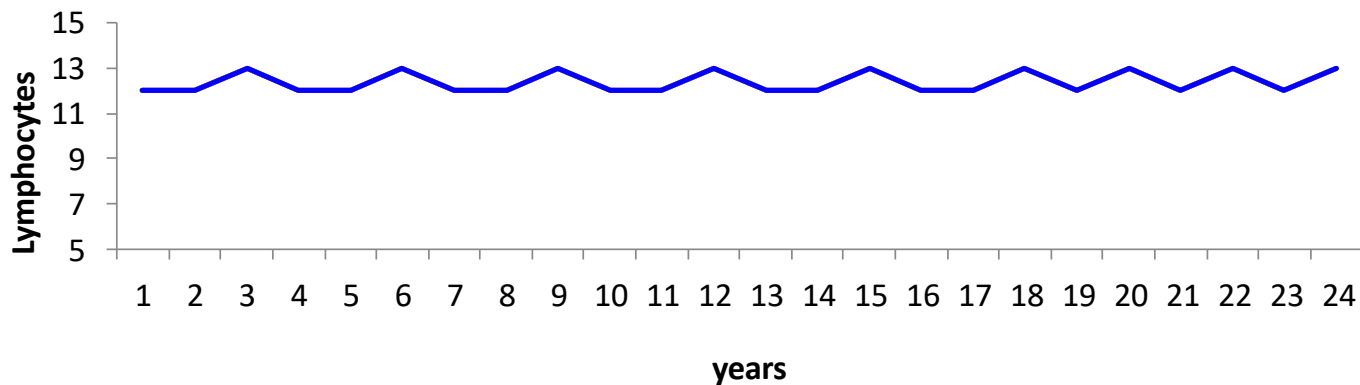


Time to first treatment

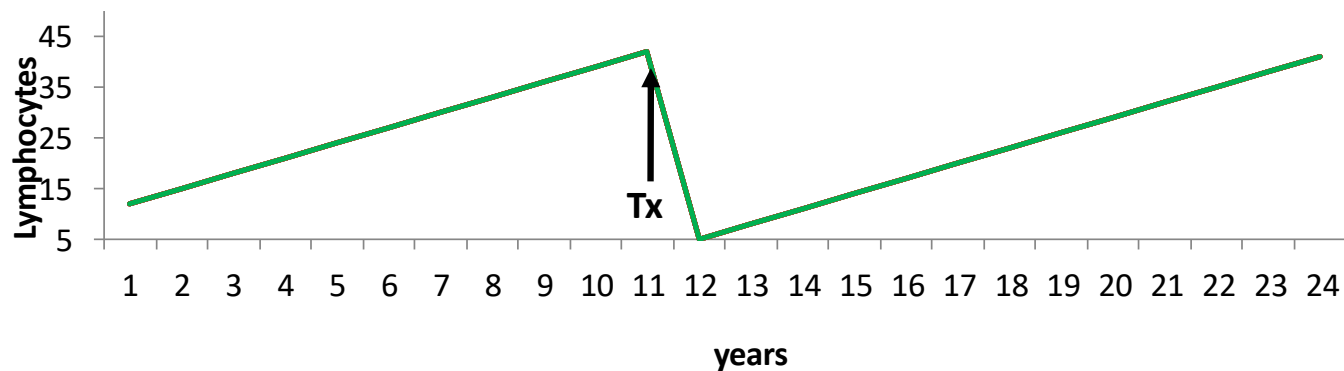
Binet A CLL: Homogeneous phenotype but heterogeneous clinical course



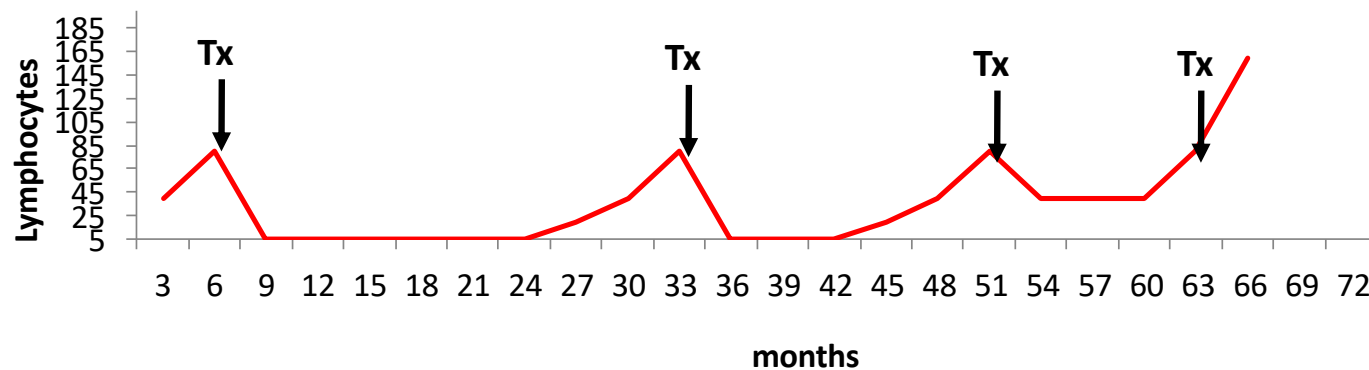
Highly stable
1/3



Slowly progressive
1/3



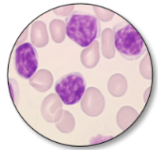
Rapidly progressive
1/3



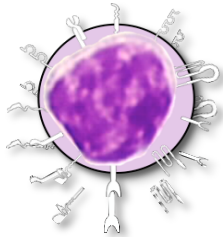
Biomarker: variable that associates with disease outcome



Host Factors: **Age**, **sex**, etc

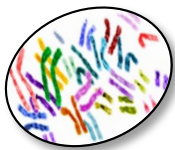


Disease Markers: **Stage**, lymphocyte count, **LDT**, etc

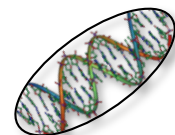


Ag expression: CD38, Zap70, **CD49d**, etc

Serology: **β 2M**, TK, LDH, sCD23, etc

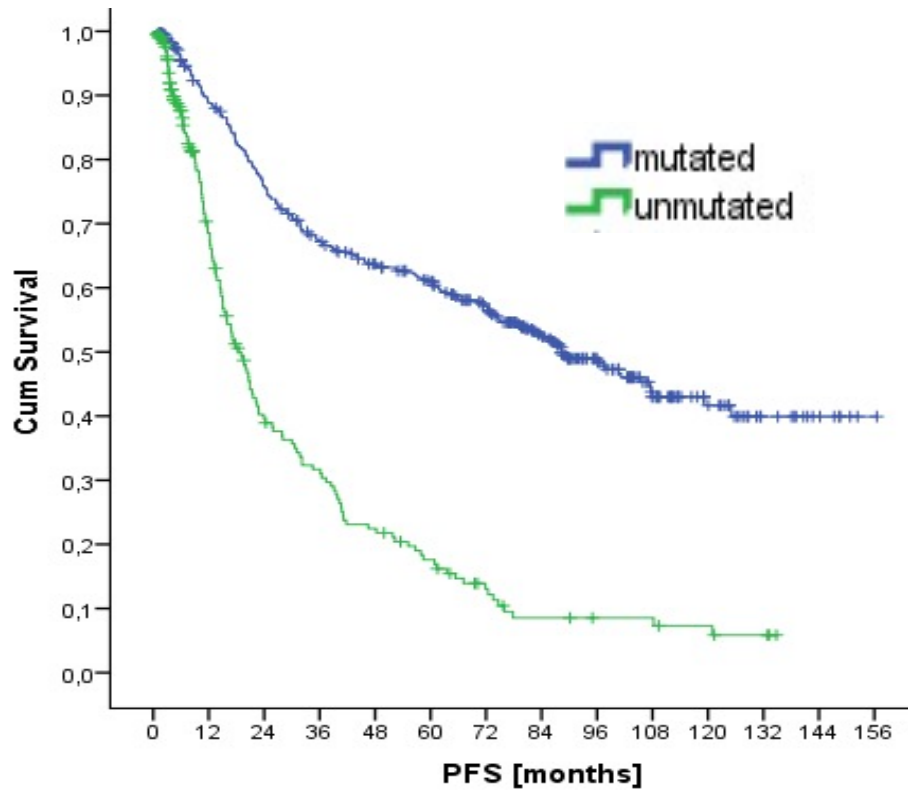


Genetics: **del17p**, **TP53 mutation**, del11q22, del13q14, trisomy 12, NOTCH1 mutation, SFRB1 mutation, etc

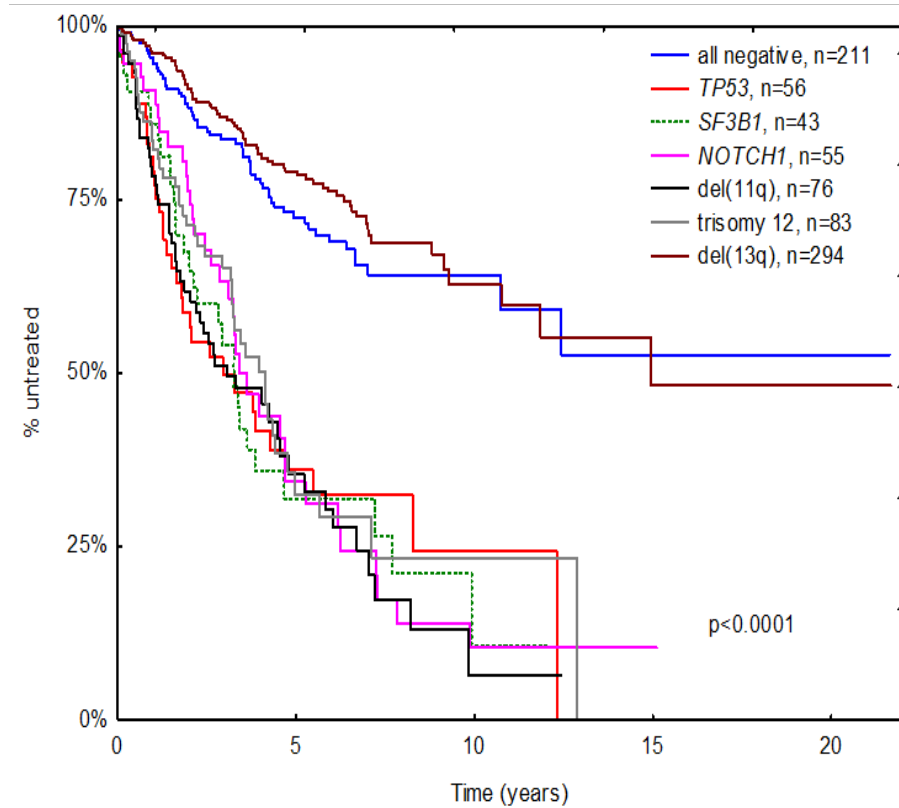


Biology Markers: **IGVH-sequence**, BCR-structure

Patient counseling on risk of progression: genetic-based models



GCLLSG CLL1 study



Baliakas et al, Leukemia 2014

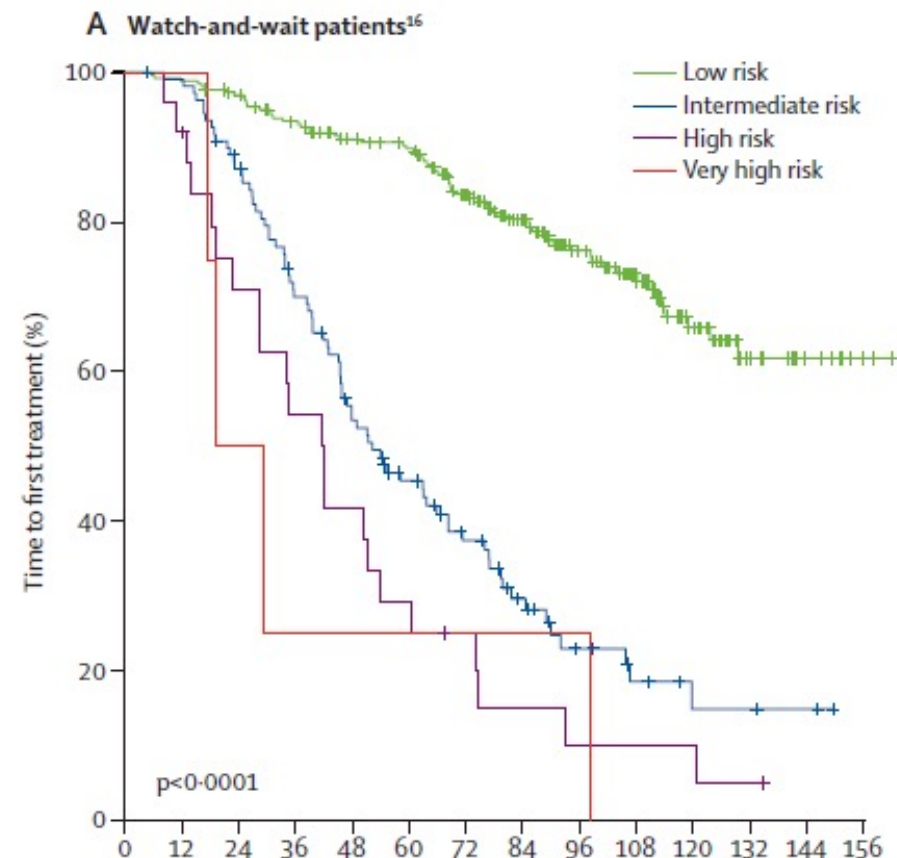
Patients with *TP53* disruption and *IGHV* mutated status show indolent clinical course: a study on 1,327 CLL

Table: Time to treatment (TTT) and overall survival (OS) according to *TP53* disruption and *IGHV* mutational status.

	n	TTT (years)	<i>TP53</i> wt/ <i>IGHV</i> -M	<i>TP53</i> disruption/ <i>IGHV</i> -M	<i>TP53</i> wt/ <i>IGHV</i> -U
		5-year OS			
<i>TP53</i>wt/<i>IGHV</i>-M	669	14			
	674	91%			
<i>TP53</i> disruption/<i>IGHV</i>-M	33	10	<u>n.s.</u>		
	35	76%	0.009		
<i>TP53</i>wt/<i>IGHV</i>-U	370	4	<0.001	<0.001	
	381	81%	<0.001	<u>n.s.</u>	
<i>TP53</i> disruption/<i>IGHV</i>-U	51	2	<0.001	<0.001	<u>n.s.</u>
	53	57%	<0.001	0.006	<0.001

CLL-IPI:

- Based on clinical trial CLL
- Developed for OS as endpoint
- Secodarily assessed in W&W CLL
- Multiple biomarkers (Age, clinical stage, B2M, IGHV, TP53)
- Complex scoring requiring a calculator



Multicenter, international, retrospective, observational study
(NCT03436524)

Study inclusion criteria:

- flow cytometry confirmed diagnosis of CLL after 1996^{1,2}
- early stage at diagnosis as defined by blood cell count and physical examination^{1,2}
- active surveillance as initial management after diagnosis

1. Cheson BD et al. Blood. 1996

2. Hallek M et al. Blood. 2008

Characteristics of the study cohorts

Variable	Training				Validation						
	Clinical trial series				Institutional series						Population based series
	UEP N=333	CLL1 N=547	CLL7 N=339	O-CLL-1 N=312	MDACC N=1225	Mayo Clinic* N=881	Barcelona N=355	Brno N=269	SU N=223	Southampton N=226	SCAN N=223
	%	%	%	%	%	%	%	%	%	%	%
Age >65	62	28	29	26	30	43	47	42	12	53	43
Male gender	53	61	63	61	59	67	57	62	56	59	59
Palpable lymph nodes	20	22	34	20	44	48	23	44	51	41	42
Lymphocytes >15x10 ⁹ /l	19	51	42	44	50	40	26	74	34	34	57
B2M >3.5 mg/l	10	8	2	1	10	10	11	12	4	21	7
Del 13q	50			46	45		50	58	44	67	41
Trisomy 12	18	9	8	9	16	18	16	11	10	11	7
Del 11q	5			7	9		9	15	10	9	5
Del 17p	6	3	2	3	6	5	4	7	4	4	2
Unmutated IGHV	28	29	22	34	39	45	39	50	31	31	25

Hb, hemoglobin; B2M, beta-2-microglobulin; FISH, fluorescence in situ hybridization; IGHV, immunoglobulin heavy variable gene

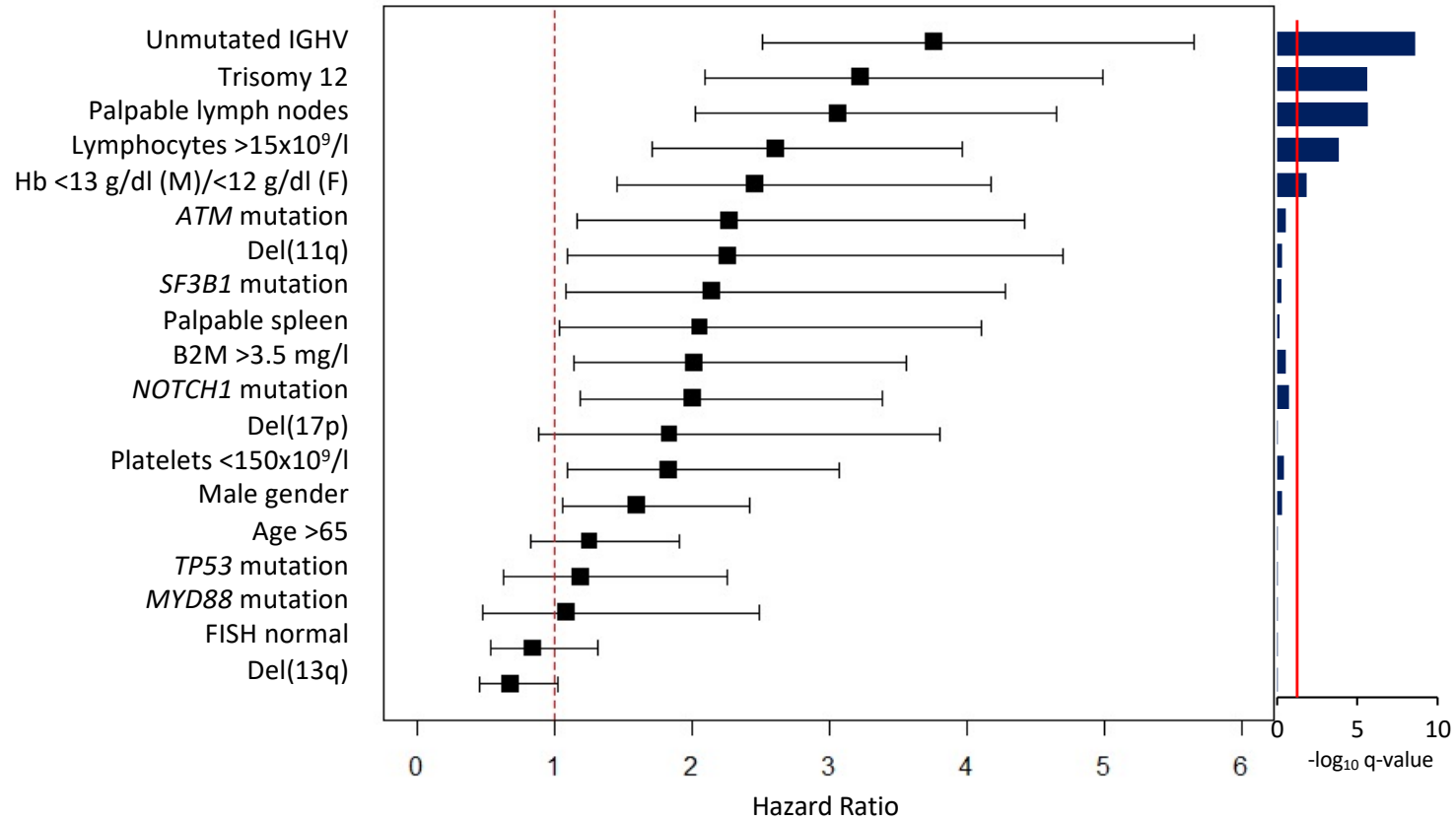
*early stage CLL according to Rai system (0-II)

N=4933

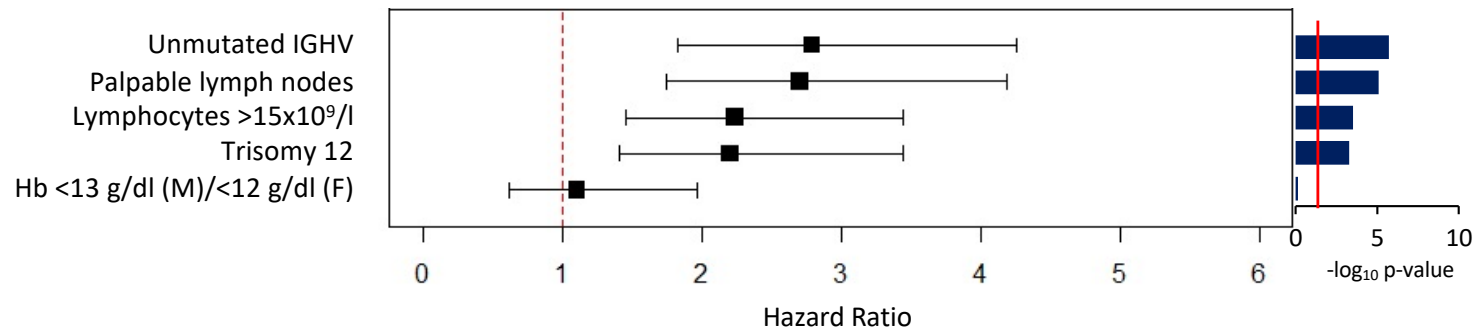
Univariate and multivariate associations and initial model for Time to First Treatment (TTFT)



Univariate



Multivariate

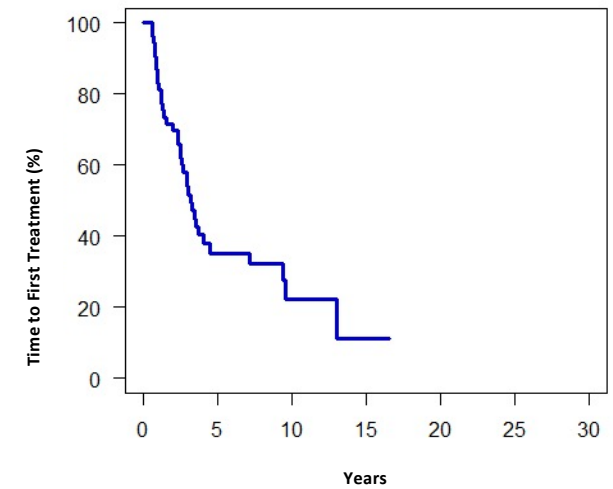
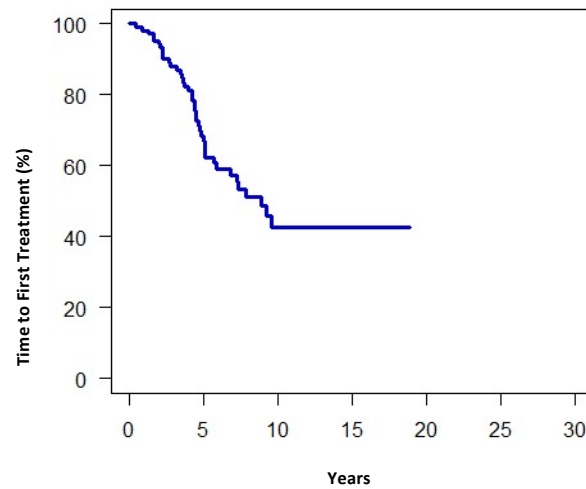
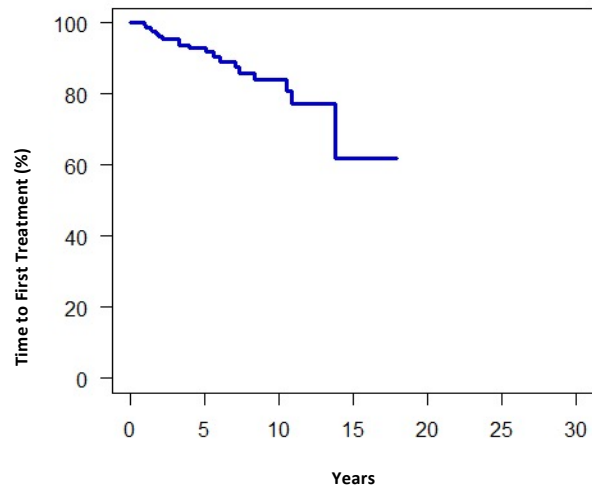
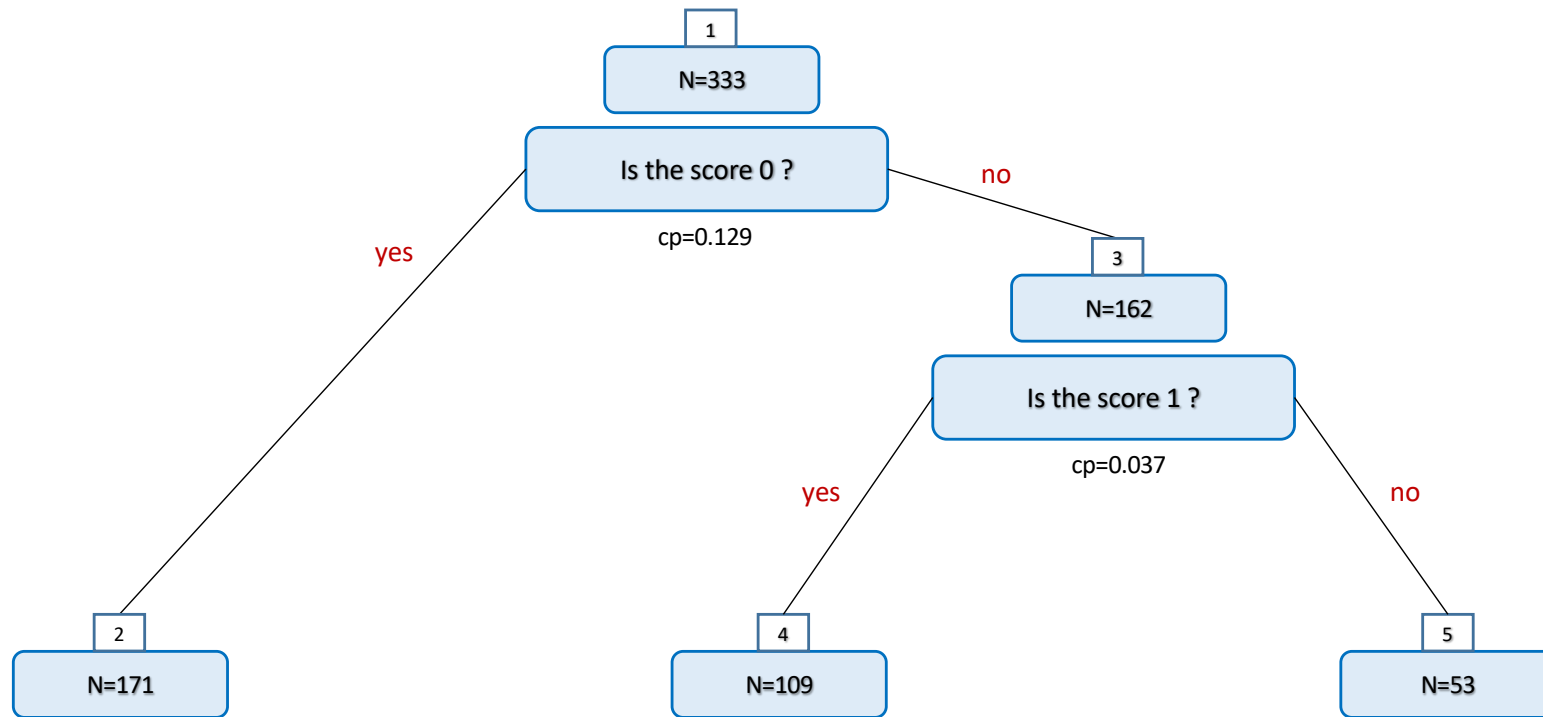


Characteristics consistently associated with TTFT



Variable	Percentage of selection in the final Cox model across the 10 Binet A study cohorts
Unmutated IGHV	100%
Lymphocytes >15x10⁹/l	90%
Palpable lymph nodes	90%
Trisomy 12	40%

Risk group stratification and IPS-E generation

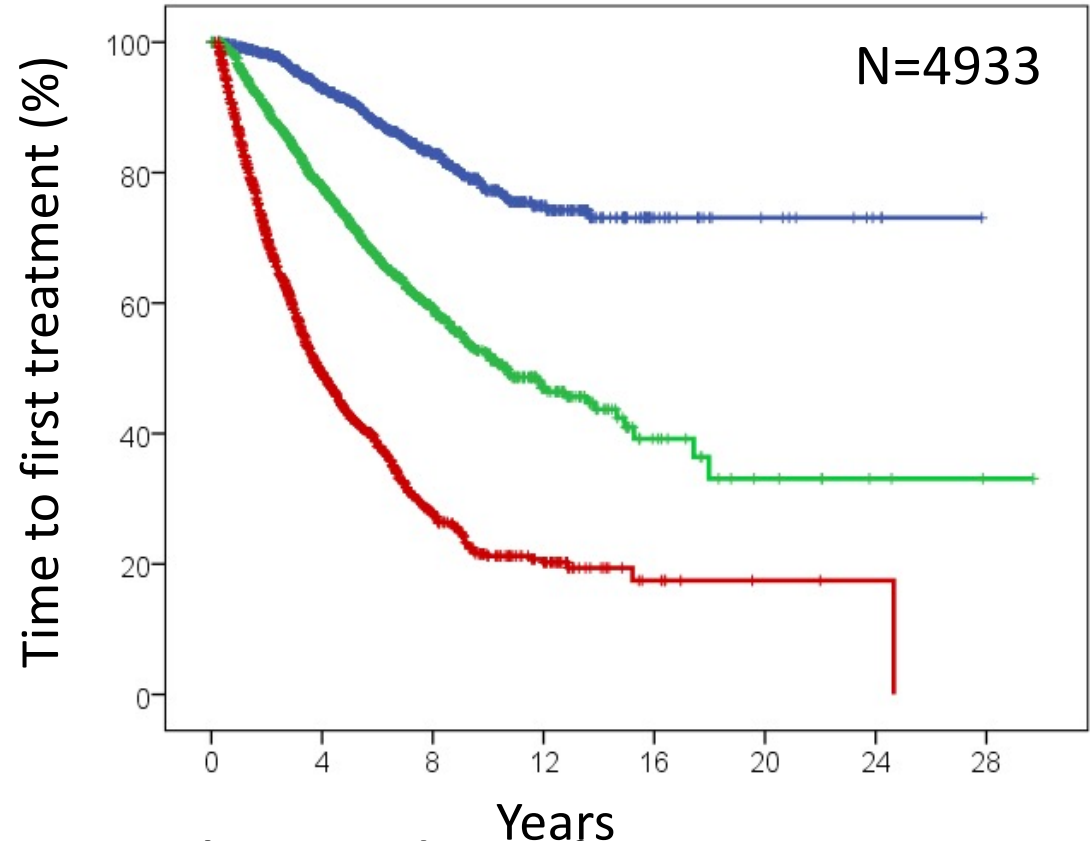


Risk group stratification and IPS-E generation



Variable	Points
IGHV unmutated	1
Lymphocytes $>15 \times 10^9/L$	1
Nodal involvement	1

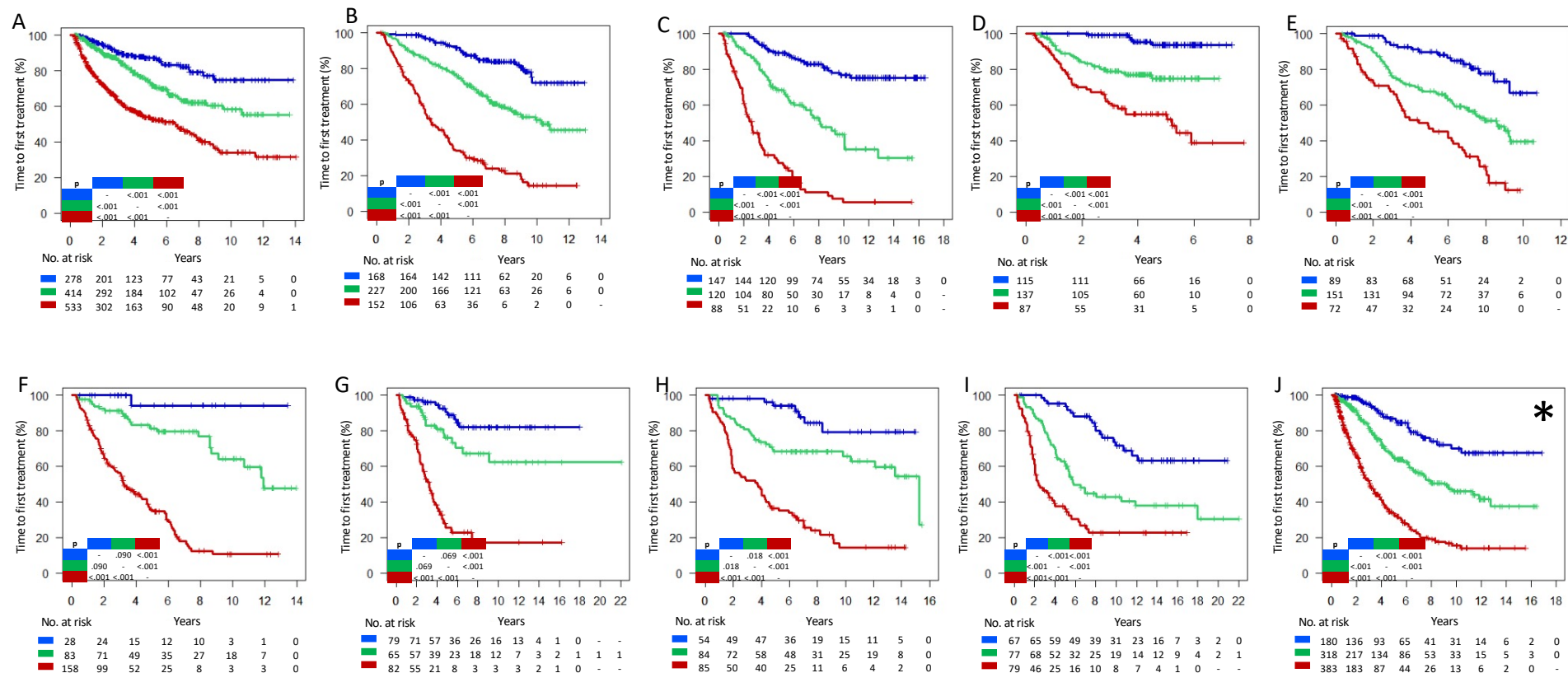
Risk group	Score
Low risk	0
Intermediate risk	1
High risk	2-3














Cumulative incidence of treatment

	1 year	5 years
Low risk	<1%	8%
Intermediate risk	3%	28%
High risk	14%	61%

IPS-E in the validation cohorts



*early stage CLL according to Rai system (0-II)
Risk group stratification maintains also after adjusting for death as a competing risk

Cohort	N	Weight (%)		IPS-E c-index
Barcelona	355	8.8		0.75
Southampton	226	5.6		0.75
UEP	333	8.2		0.74
CLL7	339	8.4		0.73
CLL1	547	13.5		0.71
SU	223	5.5		0.71
SCAN	223	5.5		0.71
Brno	269	6.6		0.69
MDACC	1225	30.2		0.66
0-CLL1	312	7.7		0.66
Total	4052	100		0.70



1 year

- high-risk: 14.1%
- intermediate-risk: 2.1%
- low-risk: <0.1%

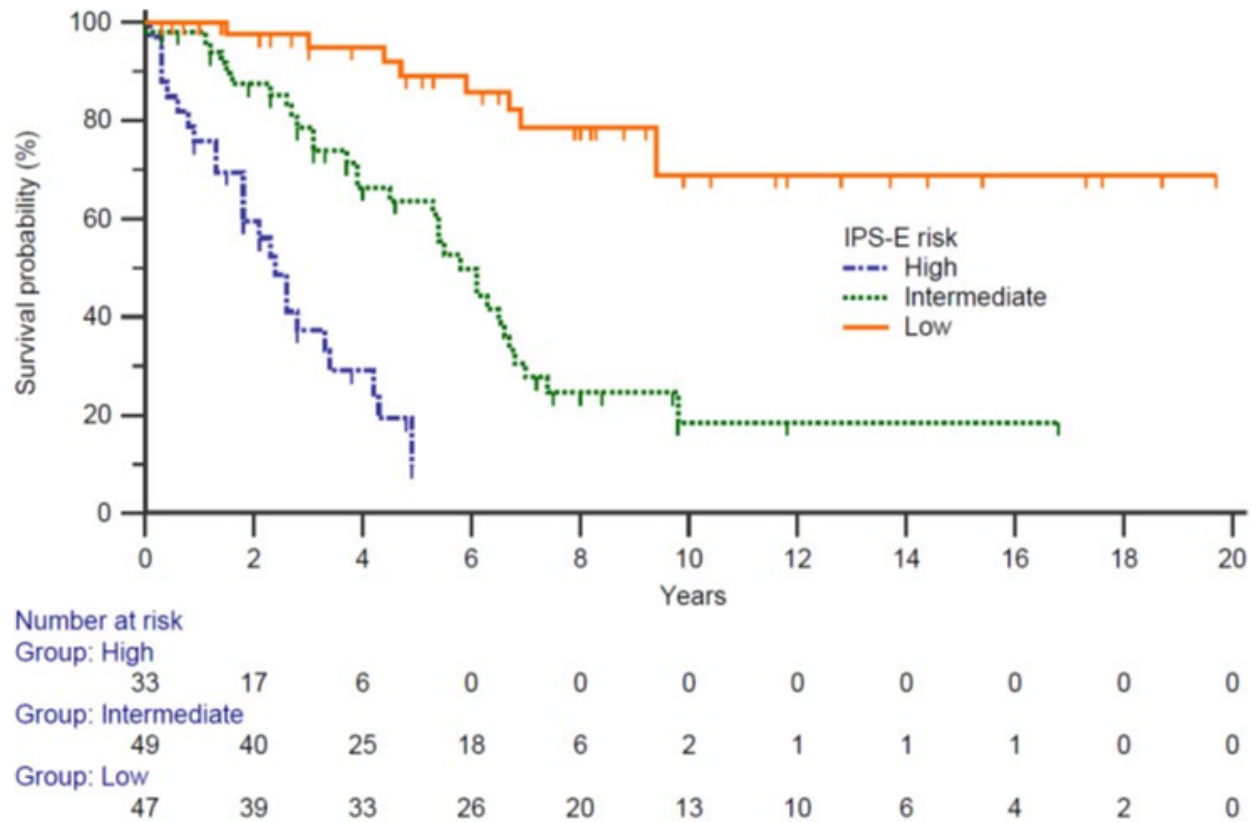
5 years

- high-risk: 61.2%
- intermediate-risk: 28.4%
- low-risk: 8.4%

External validation of IPS-E (I)

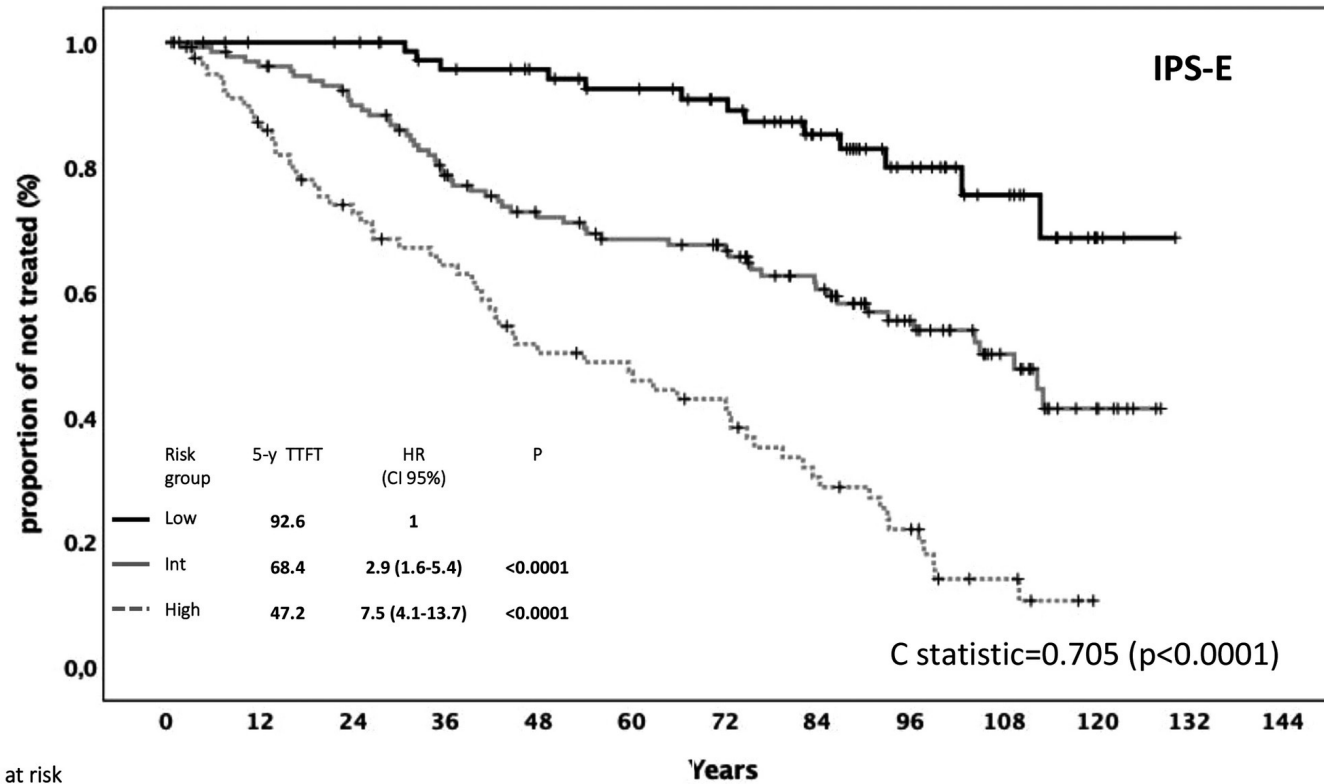


N=130 Binet A CLL



5-year TTFT of 8%, 28%, and 61%

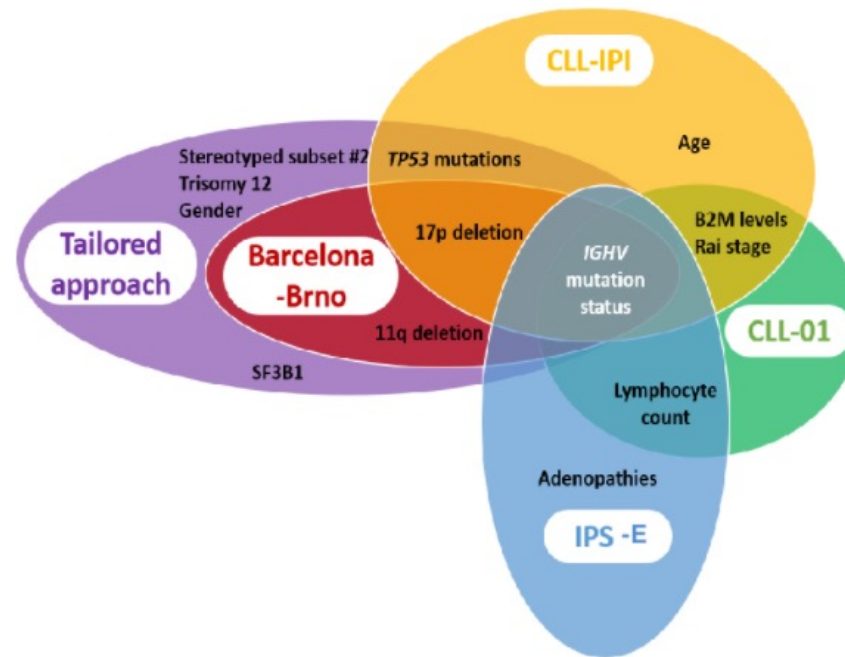
External validation of IPS-E (III)



No. of pts at risk

Low risk	77	74	72	65	61	57	50	38	24	114	5	0	0
Int risk	135	125	114	95	83	75	70	56	36	21	7	0	0
High risk	80	67	54	44	35	31	27	18	12	5	0	0	0

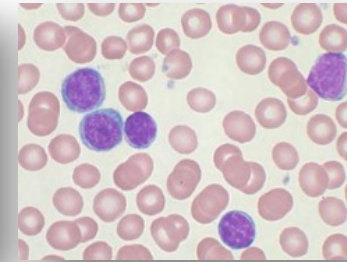
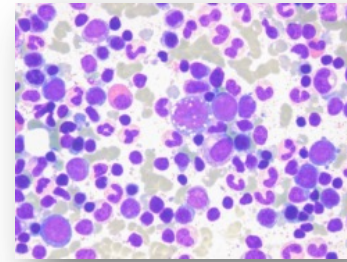
External validation of IPS-E (II)



Score risk	Low			Intermediate			High			P	C-index
	N	%	Median TTFT	N	%	Median TTFT	N	%	Median TTFT		
CLL-IPI	265	62	188	133	31	52	30	7	27	<0.001	0.67
Barcelona-Brno	283	66	188	127	30	53	18	4	31	<0.001	0.67
IPS-A	185	43	NR	181	42	88	62	15	31	<0.001	0.72
CLL-01	231	54	188	160	37	61	37	9	31	<0.001	0.69
Tailored-M	265	92	188	24	8	60				<0.001	0.61
Tailored-UM	46	33	60	75	47	44	18	13	30	<0.001	0.58

Upfront definition of the risk of treatment requirement can benefit:

- **patients**, who can be informed about the likely course of their disease
- **physicians**, who can allocate medical resources according to patients' risk
- **researchers**, who can design risk-adapted clinical trials



• Lymphoma

• Leukemia



Laboratory of experimental Hematology IOR

- Post Doc**
 - Alessio Bruscaffin PhD
 - Deborah Piffaretti PhD
 - Simone Bocchetta PhD
- Lab Technician**
 - Gabriela Forestieri BSc
 - Katia Pini BSc
- Research Fellow**
 - Marco Marangon BSc
 - Adalgisa Condoluci MD, PhD student
 - Cristina Piroso MD, PhD student
- Bioinformatician**
 - Joyce Marques De Almeida MD, PhD student
 - Lodovico Terzi Di Bergamo MSc, PhD student
 - Matin Salehi MSc, PhD student



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