

HIGHLIGHTS IN EMATOLOGIA

TREVISO
7-8 NOVEMBRE 2025



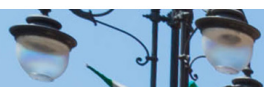
LINFOMA DI HODGKIN CHE COSA C'È DI NUOVO IN TERMINI PROGNOSTICI?

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Disclosures: ALESSANDRO BROCCOLI

Company name	Research support	Consultant	Stockholder	Speakers bureau	Advisory board	Other
Sandoz	X				X	X
Gilead				X	X	X
Merck	X					X
Janssen	X			X		X
Takeda		X		X	X	X
Kyowa Kirin	X			X	X	X
Incyte						X
Recordati						X
Astra Zeneca				X	X	
Roche				X		X
GlaxoSmithKline		X		X	X	
BeOne	X			X	X	X
Eli Lilly						X
SOBI				X		
SERB Pharma		X			X	

Prognostication for early- and advanced-stage Hodgkin lymphoma

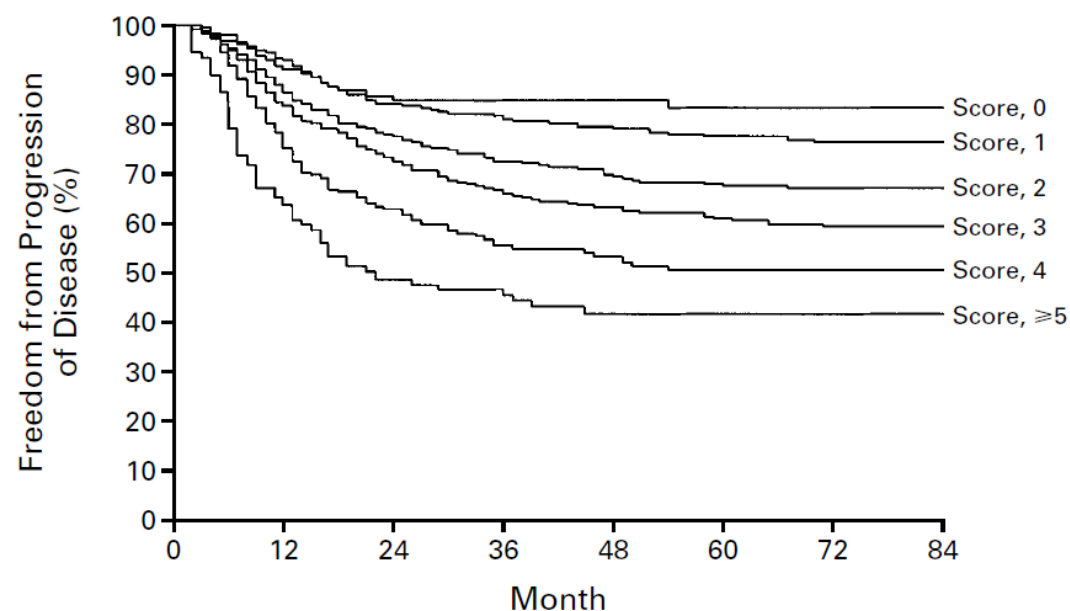
- Prognostic models based on pre-treatment factors can help identify patients with early- and advanced-stage classic Hodgkin lymphoma who are at increased risk of relapse (or death).
- In early-stage disease, EORTC and GHSG classifications have been followed for the past 40-50 years.
- Original EORTC early-stage prognostic factors date back to early '70s, when staging laparotomy and mantle radiation (alone) were the predominant ways of treatment.
- In advanced-stage disease, the International Prognostic Score (IPS7) had been a standard index in classic Hodgkin lymphoma for 25 years.
- Performance of IPS7 decreased when analyzed in patients treated in a more recent era, and is almost surpassed when regimens containing newer drugs are applied frontline (poor calibration).
- Updated analyses with modern clinical trial and registry (real-world data) is desired.

A PROGNOSTIC SCORE FOR ADVANCED HODGKIN'S DISEASE

DIRK HASENCLEVER, PH.D., AND VOLKER DIEHL, M.D.,
FOR THE INTERNATIONAL PROGNOSTIC FACTORS PROJECT ON ADVANCED HODGKIN'S DISEASE*

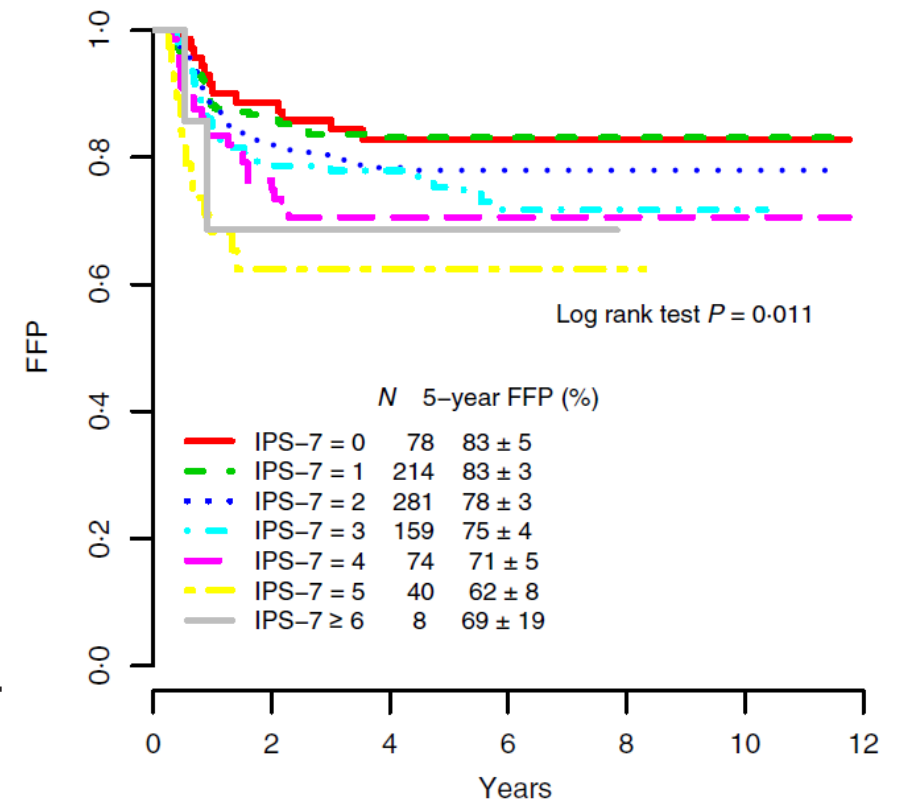
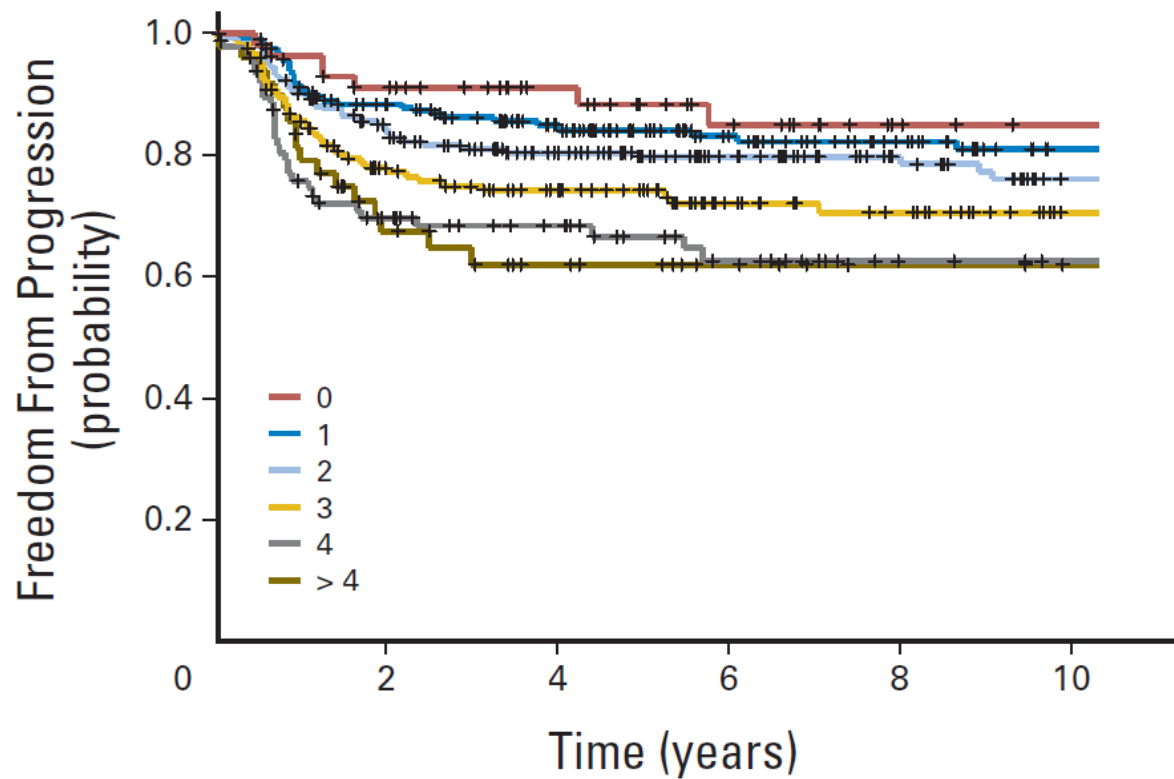
THE FINAL COX REGRESSION MODEL.

FACTOR	LOG HAZARD RATIO	P VALUE	RELATIVE RISK
Serum albumin, <4 g/dl	0.40±0.10	<0.001	1.49
Hemoglobin, <10.5 g/dl	0.30±0.11	0.006	1.35
Male sex	0.30±0.09	0.001	1.35
Stage IV disease	0.23±0.09	0.011	1.26
Age, ≥45 yr	0.33±0.10	0.001	1.39
White-cell count, ≥15,000/mm ³	0.34±0.11	0.001	1.41
Lymphocyte count, <600/mm ³ or <8% of white-cell count	0.31±0.10	0.002	1.38



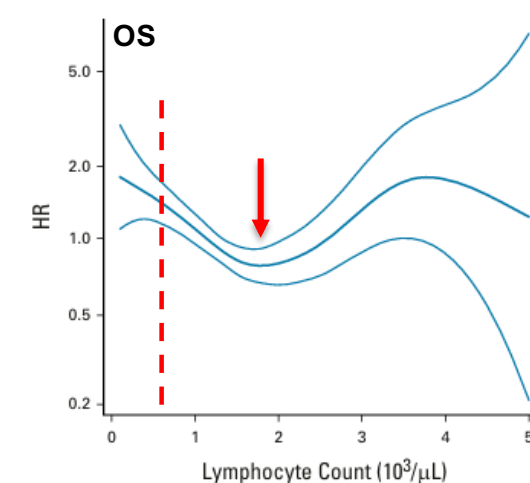
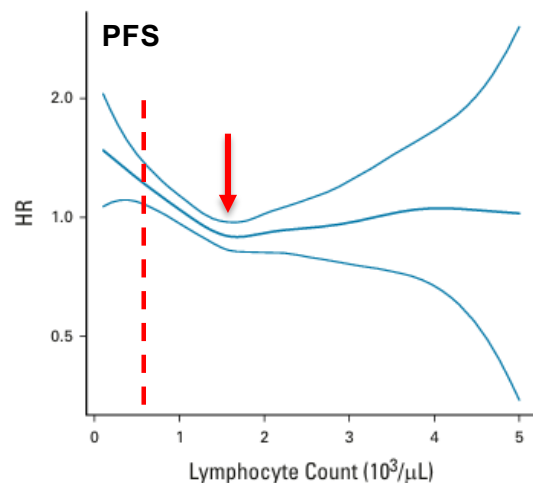
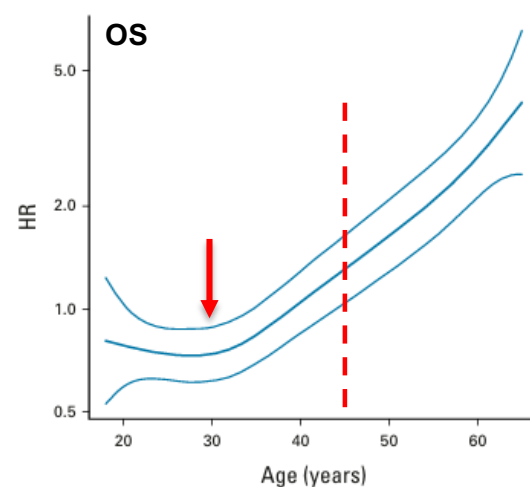
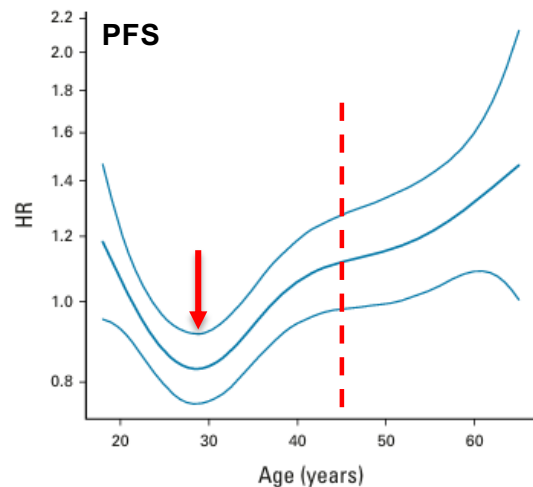
Treatments started **before January 1992**. More than 75 percent of the patients were treated with standard doxorubicin-containing regimens; 20 percent received MOPP or a similar regimen. Sixty percent of the patients received no radiotherapy. Thirty-three percent received full or selected involved field irradiation; 2 percent underwent more extensive irradiation with a mantle or inverted-Y field, and 5 percent underwent subtotal or total nodal irradiation.

Performance of IPS7 in modern era



The Advanced-Stage Hodgkin Lymphoma International Prognostic Index: Development and Validation of a Clinical Prediction Model From the HoLISTIC Consortium

- **Why did IPS7 have to be refined?** Due to improvements in diagnostic accuracy, use of PET in more accurate stage definition, optimization of drug dosing and treatment dose intensity, treatment adaptation according to early PET results, improved post-relapse salvage treatment modalities.
- **How was it performed?**
 - A population of adult patients with newly diagnosed classic Hodgkin lymphoma aged 18-65 years and with advanced stage (IIB, III, IV).
 - Model developed on 4,022 patients from 8 clinical trials enrolling advanced-stage patients between 1996 and 2014; validated on 1,431 patients from 4 cancer registries (between 1996 and 2019).
- Continuous variables (e.g. age, laboratory parameters) were no more considered as dichotomized.
- **Novel non-linear relationships between age and absolute lymphocyte counts have been identified, along with unique prognostic variables significant for PFS versus OS.**
- An online calculator to assist clinicians and patients in estimating individualized prognosis.



The Advanced-Stage Hodgkin Lymphoma International Prognostic Index: Development and Validation of a Clinical Prediction Model From the HoLISTIC Consortium

	5-year PFS HR (95% CI)	5-year OS HR (95% CI)
Age (years)		
Linear effect in 18 to 30 years	0.97 (0.95, 1.00)	0.98 (0.94, 1.02)
Linear effect in >30 years	1.02 (1.01, 1.02)	1.05 (1.04, 1.07)
Female	--	0.78 (0.61, 1.00)
Stage		
Stage IIB		
Stage III	1.23 (1.03, 1.48)	
Stage IV	1.53 (1.27, 1.83)	1.33 (1.04, 1.70)
Bulk	--	1.37 (1.05, 1.78)
Hemoglobin (g/dL)	--	0.88 (0.81, 0.96)
Albumin (g/dL)	0.74 (0.66, 0.82)	0.67 (0.53, 0.84)
Lymphocyte count ($10^3/\text{mm}^3$)		
Linear effect in .1 to 2	0.75 (0.65, 0.87)	0.61 (0.46, 0.80)
Linear effect in 2 to 5	1.21 (0.96, 1.52)	1.49 (0.99, 2.22)

The Advanced-Stage Hodgkin Lymphoma
International Prognostic Index: Development and
Validation of a Clinical Prediction Model From
the HoLISTIC Consortium

ADVANCED STAGE HODGKIN LYMPHOMA INTERNATIONAL PROGNOSTIC INDEX (A-HIPI)

This model predicts 5-year progression-free survival and overall survival in adults (18 to 65 years of age) with newly diagnosed advanced stage Hodgkin Lymphoma (stage 2B, 3 and 4)

Age (years)

22

Values from 18 to 65 allowed.

Gender:

Male

Stage:

4

Any Bulk:

Yes

Lymphocyte count:

2.4

10⁹/μL

Values from 0.1 to 5 are allowed.

Hemoglobin:

11

g/dL

Values from 8 to 16.5 are allowed.

Albumin:

4.5

g/dL

Values from 1 to 6 are allowed.

CALCULATE

Progression Free Survival at 5 years

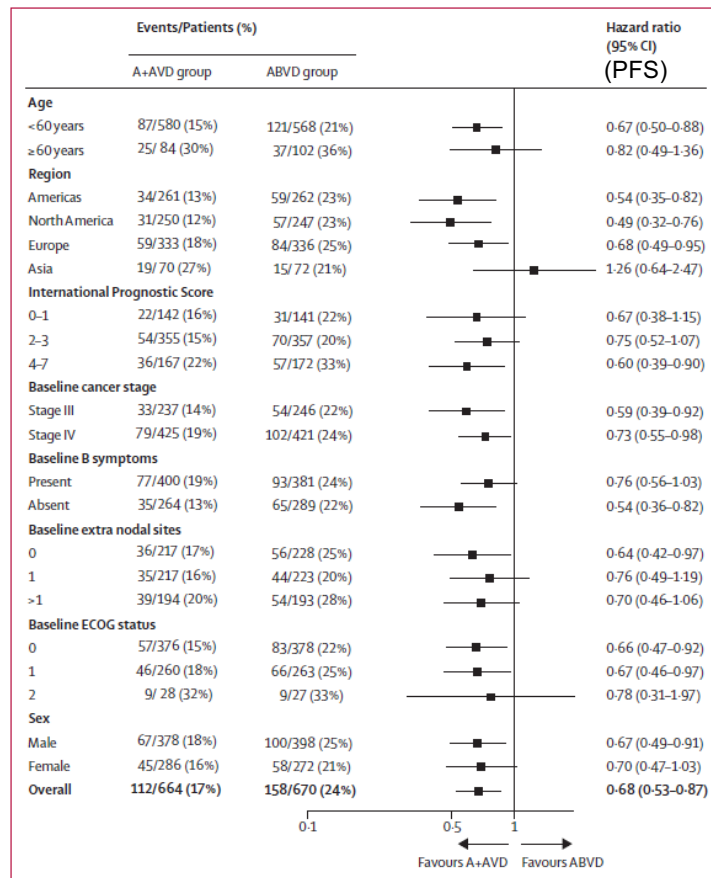
80.19%

Overall Survival at 5 years

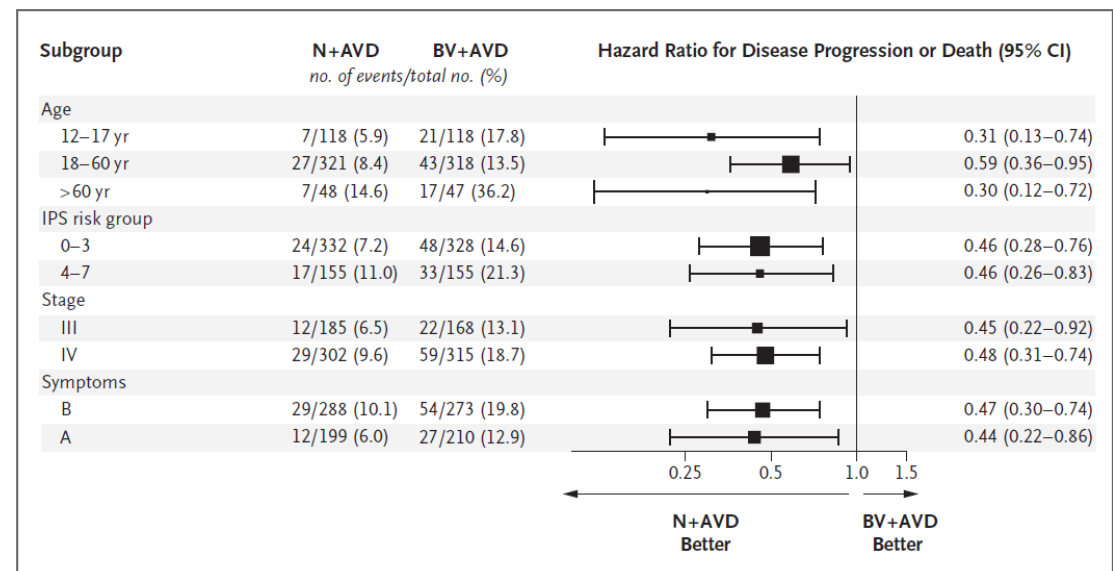
94.49%

Significance of IPS7 in contemporary randomized clinical trials with new drugs (1)

A+AVD vs ABVD

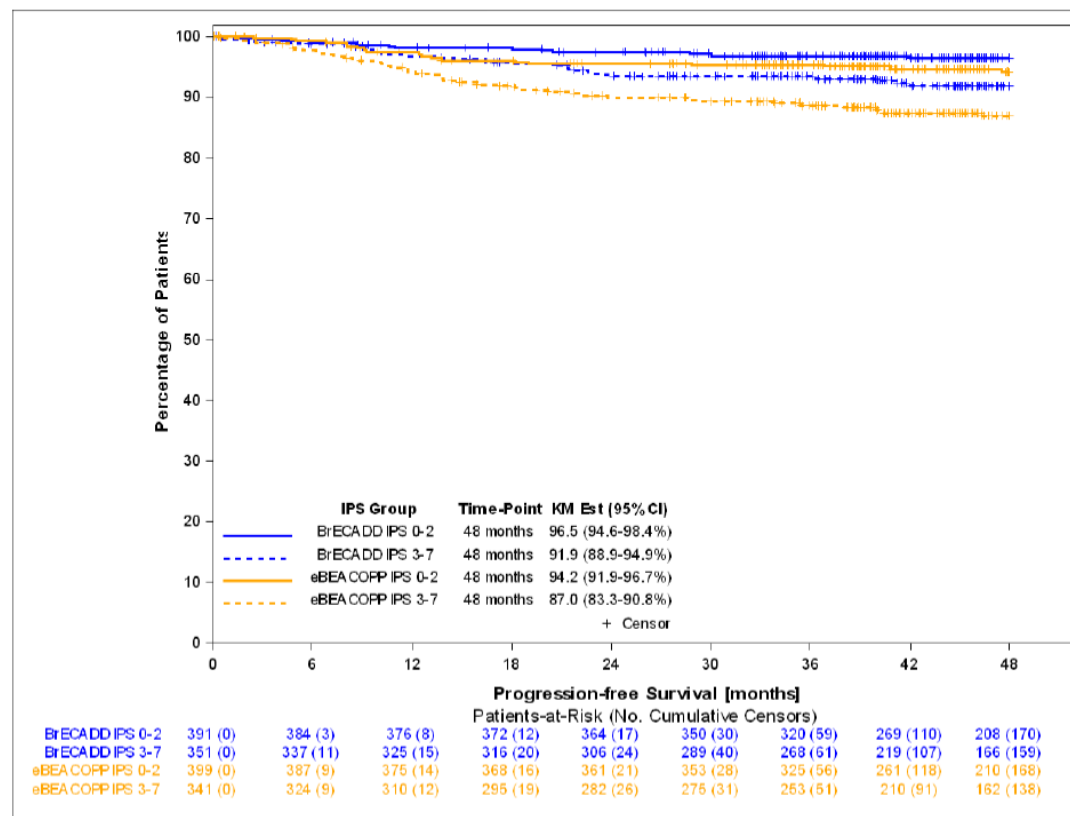


N+AVD vs A+AVD



Significance of IPS7 in contemporary randomized clinical trials with new drugs (2)

BrECADD vs eBEACOPP



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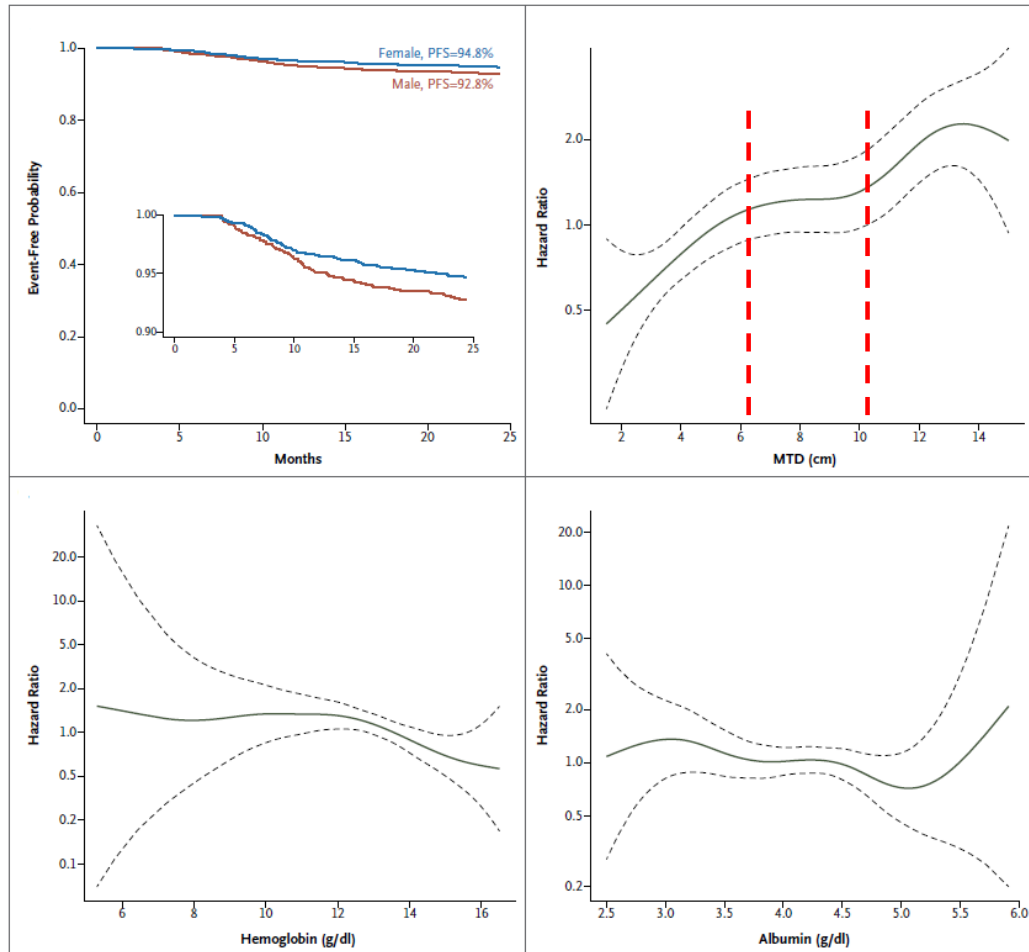
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Shaping treatment according to risk factors and stage in Hodgkin lymphoma

Category	Features	Approach
Early-stage favorable	Stage I-II with no risk factors	ABVD x 2 + 20 Gy IFRT (non PET-adapted) ⁴ ABVD x 2 + 20 Gy IFRT (PET-adapted) ⁵ ABVD x 3 + 30 Gy IFRT (PET-adapted) ⁶
Early-stage unfavorable	Stage I-II with risk factors ¹⁻³ • Age ≥ 50 years (*) • Large mediastinal mass • B symptoms • ≥ 3 involved lymph node areas • Elevated ESR • Extranodal involvement	ABVD x 4 + 30 Gy IFRT (PET-adapted) ⁶ ABVD x 6 (PET-adapted) ⁶ ABVD x 2 → AVD x 4 (PET-adapted) ⁷
Advanced stage	Stage II B with risk factors Stage III-IV	ABVD x 2 → AVD x 4 (PET2 negative) ⁷ ABVD x 2 → BEACOPP (PET2 positive) ^{8,9} A-AVD x 6 (non PET-guided) ¹⁰ N-AVD, BrECADD (non PET-guided) ^{11,12}

(*) EORTC only

1. Tubiana M. *Cancer Res*, 1971; 31: 1801-1810 — 2. Tubiana M. *Natl Cancer Inst Monogr*, 1973; 36: 513-530 — 3. Tubiana M. *Int J Radiat Oncol Biol Phys*, 1985; 11: 23-30
4. Engert A. *N Engl J Med*, 2010; 363: 640-652 — 5. Fuchs M. *J Clin Oncol*, 2019; 37: 2835-2845 — 6. Federico M. *J Clin Oncol*, 2024; 42: 19-25
7. Johnson P. *N Engl J Med*, 2016; 374: 2419-2429 — 8. Press OW. *J Clin Oncol*, 2016; 34: 2020-2027 — 9. Gallamini A. *J Clin Oncol*, 2018; 36: 454-462
10. Ansell SM. *N Engl J Med*, 2022; 387: 310-320 — 11. Herrera AF. *N Engl J Med*, 2024; 391: 1379-1389 — 12. Borchmann P. *Lancet*, 2024; 404: 341-352



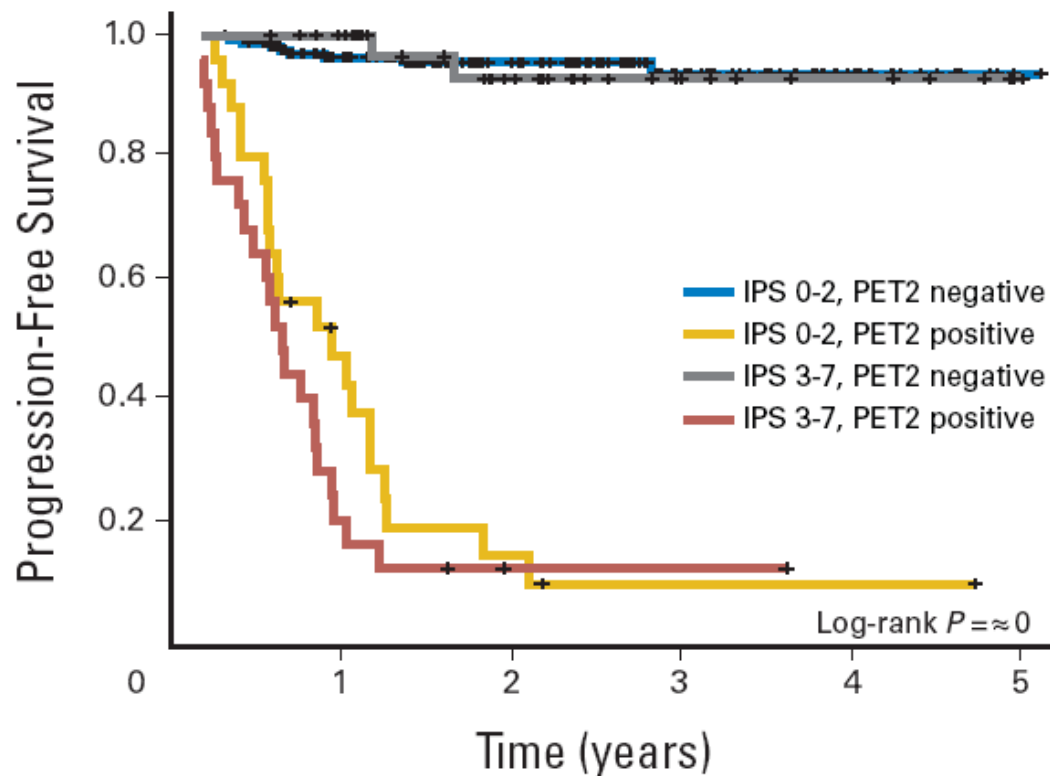
An Individualized Prediction Model for Early-Stage Classic Hodgkin's Lymphoma

PFS-related variables

	Development (N=3000)	Validation 1 (N=1488)	Validation 2 (N=872)
Age (years), mean (SD)	34 (12)	34 (12)	38 (14)
Female sex	51%	53%	46%
Stage			
Stage I	23%	14%	25%
Stage IIA	55%	60%	46%
Stage IIB	22%	26%	30%
MTD, mean (SD)	6.5 (3.5)	6.5 (3.3)	5.3 (2.8)
Hemoglobin (g/dL), mean (SD)	13.0 (1.6)	13.1 (1.6)	13.4 (1.8)
Albumin (g/dL), mean (SD)	4.2 (0.5)	4.0 (0.5)	4.1 (0.5)
2-year PFS (KM)	93.7%	90.3%	91.6%

3,000 patients from 4 seminal early-stage trials with accrual from 1996 to 2011; two model validation cohorts with 2,360 patients from 5 cancer registries, diagnosed between 1996 and 2019 (treated with curative intent and outside clinical trials).

Predictive role of PET2 in Hodgkin lymphoma and its superiority to IPS7



	All	95% confidence level
All	260	---
True positive	33	---
True negative	203	---
False positive	12	---
False negative	12	---
Positive predictive value	0.73	0.68-0.79
Negative predictive value	0.94	0.92-0.97
Sensitivity	0.73	0.68-0.79
Specificity	0.94	0.92-0.97
Accuracy	0.91	0.87-0.94

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PET2-adapted treatment design in 4 clinical trials

Study	Pts.	Clinical data	PET (+2) positive	Intensification (adherence)	Final PET neg	PFS	Toxicity (gr. 3-4)
Johnson ¹ (2016)	1135	Age: 32 yrs Bulky: 33% B sympt: 61% Stage III-IV: 59%	16% (Deauville)	BEACOPP (95%)	74%	68% (3 yrs) [85% A(B)VD] $\Delta = 17\%$	Hematol: 72% Infections: 37%
Press ² (2016)	331	Age: 32 yrs Bulky: 18% B sympt: 62% Stage III-IV: 100%	18% (Deauville)	BEACOPP (92%)	55%	64% (2 yrs) [82% ABVD] $\Delta = 18\%$	Hematol: NR Infections: 42%
Zinzani ³ (2016)	512	Age: 33 yrs Bulky: 35% B sympt: 64% Stage III-IV: 81%	20% (Juweid)	HDT HSCT (79%)	72%	74% (2 yrs) [81% ABVD] $\Delta = 7\%$	Hematol: 72% Infections: 16%
Gallamini ⁴ (2018)	780	age: 31 yrs Bulky: 58% B sympt: 81% Stage III-IV: 64%	19% (Deauville)	BEACOPP (99%)	54%	60% (3 yrs) [87% ABVD] $\Delta = 27\%$	Hematol: 76% Infections: 10%

1. Johnson P. *N Engl J Med*, 2016; 374: 2419-2429 — 2. Press OW. *J Clin Oncol*, 2016; 34: 2020-2027

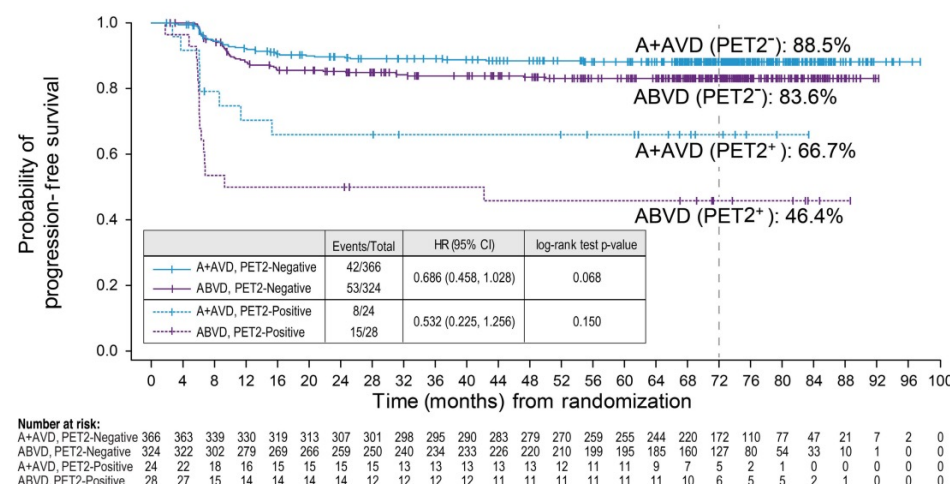
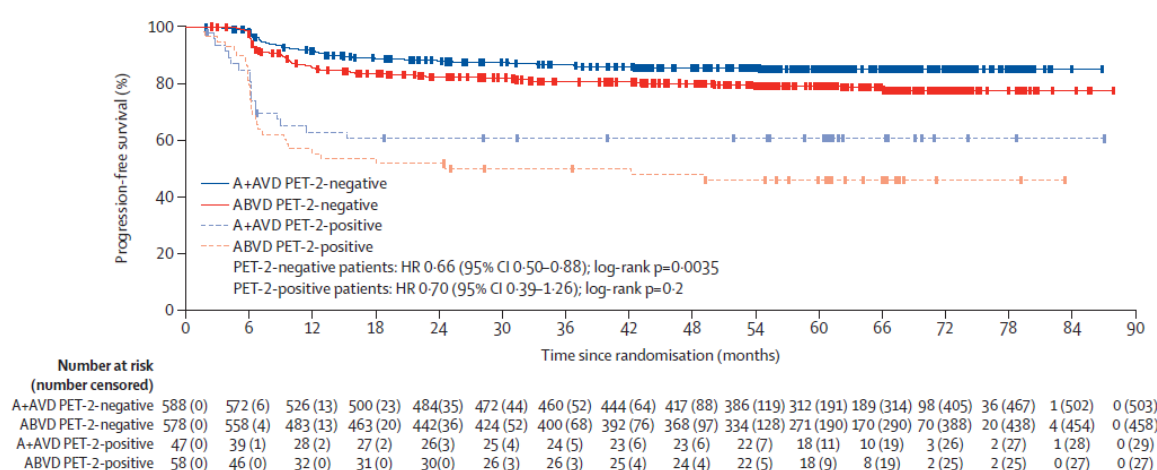
3. Zinzani PL. *J Clin Oncol*, 2016; 34: 1376-1385 — 4. Gallamini A. *J Clin Oncol*, 2018; 36: 454-462

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Is PET2 still informative in frontline approaches with new drugs?

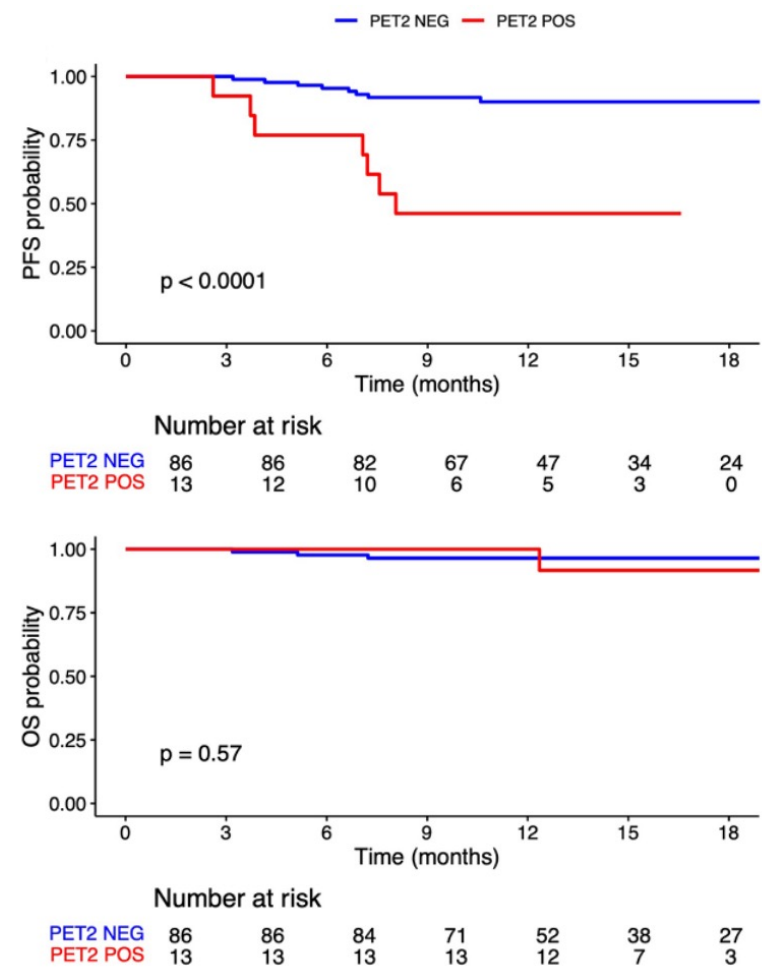
ECHELON-1 was not a PET2-adapted trial. Patients with positive PET2 (Deauville 4 and 5) continued on treatment according to randomization. Patients with Deauville 5 at PET2 could be switched to another off-protocol treatment (e.g. escalation).



All patients	Deaths	Progression events	Death + Progression events	Age 18-39	Death + Progression events
A-AVD PET2 negative	9/588 (1.5%)	76/588 (12.9%)	85/588 (14.5%)	A-AVD PET2 negative	42/366 (11.5%)
ABVD PET2 negative	26/578 (4.5%)	94/578 (16.3%)	120/578 (20.8%)	ABVD PET2 negative	53/324 (16.4%)
A-AVD PET2 positive	0/47 (0)	18/47 (38.3%)	18/47 (38.3%)	A-AVD PET2 positive	8/24 (33.3%)
ABVD PET2 positive	3/58 (5.2%)	28/58 (48.3%)	31/58 (53.4%)	ABVD PET2 positive	15/28 (53.6%)

Straus DJ. *Lancet Haematol*, 2021; 8: e410-e421 — Crosswell HE. *Haematologica*, 2024; 109: 982-987

Interim-PET predicts progression-free survival in stage IV Hodgkin lymphoma treated with upfront brentuximab vedotin-AVD

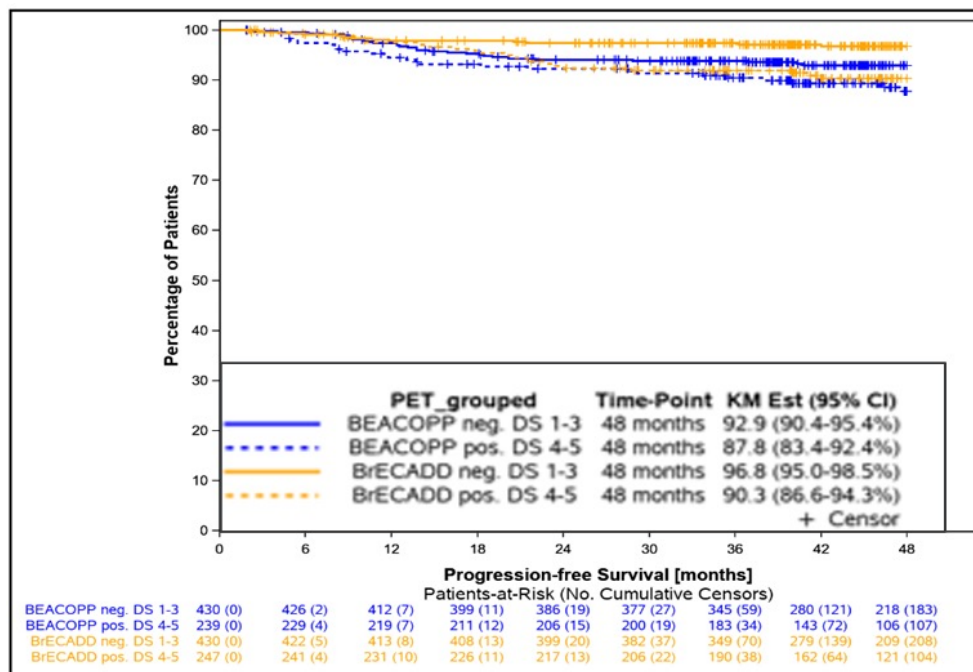


Variable	Comparison	HR	95% CI	p-value	
Age	>65 vs <=65 years	1.56	0.37-6.49	0.542	
B symptoms	Yes vs No	1.64	0.34-7.9	0.535	
Extranodal involvement	>1 vs 0-1	1.86	0.59-5.91	0.293	
Bulky	Yes vs No	1.39	0.4-4.88	0.607	
Hasenclever score	>=4 vs <4	1.81	0.55-6.02	0.331	
PET2 DS	Pos vs Neg	4.64	1.47-14.62	0.009 **	
Treatment delay for neutropenia	Yes vs No	3.23	0.95-11	0.061	

Is PET2 still informative in frontline approaches with new drugs?

HD21 (BrECADD vs eBEACOPP) is a PET2-adapted trial in which PET2-negative patients receive 2+2 cycles of either BrECADD or eBEACOPP according to randomization, whereas PET2-positive patients receive 2+4 cycles of therapy.

BrECADD vs eBEACOPP

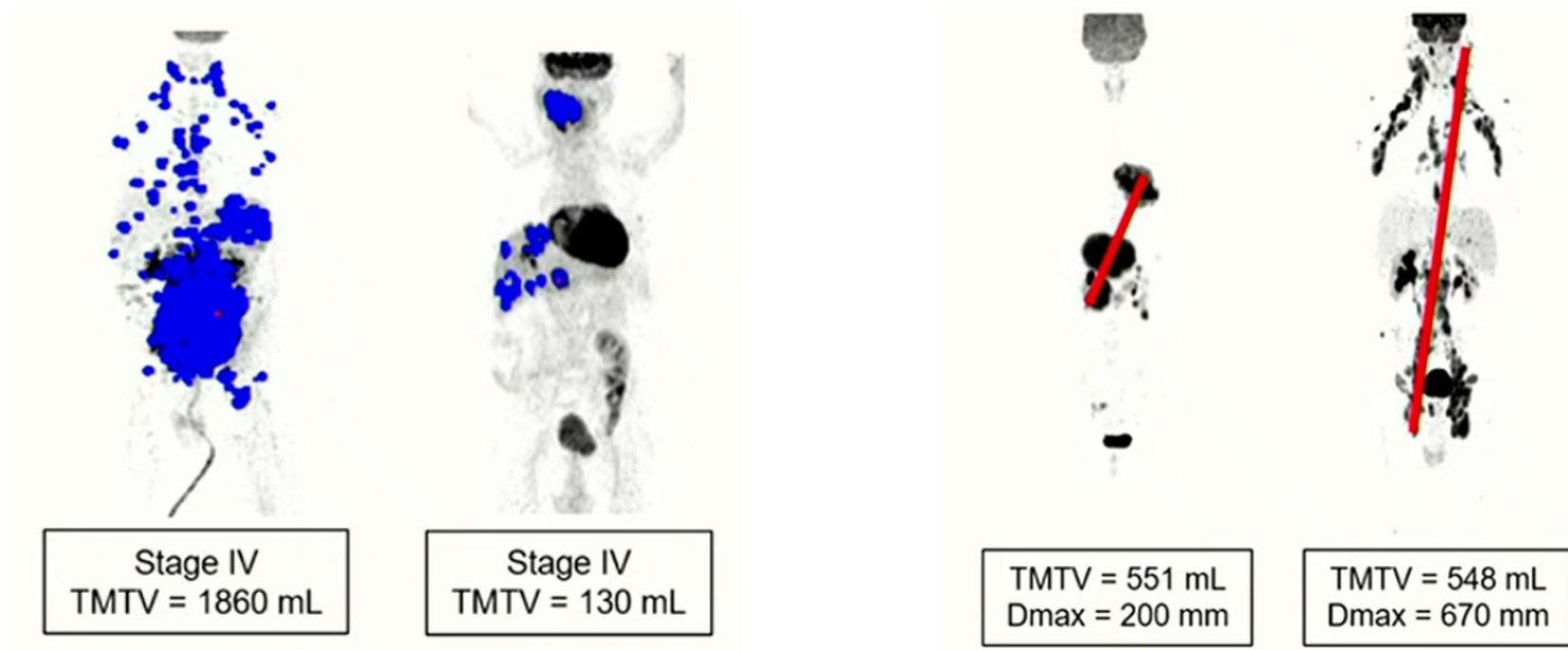


	BV-AVD	PET Adapted BrECADD	Nivo-AVD
Median f/u	73 months	48 months	25 months
PFS	83%	94.3%	92%
iPET neg	85%	96.8%	NA
iPET pos	60.6%	90.3%	NA
# cycles	6	4-6	6
OS	93.4%	98.6%	99%

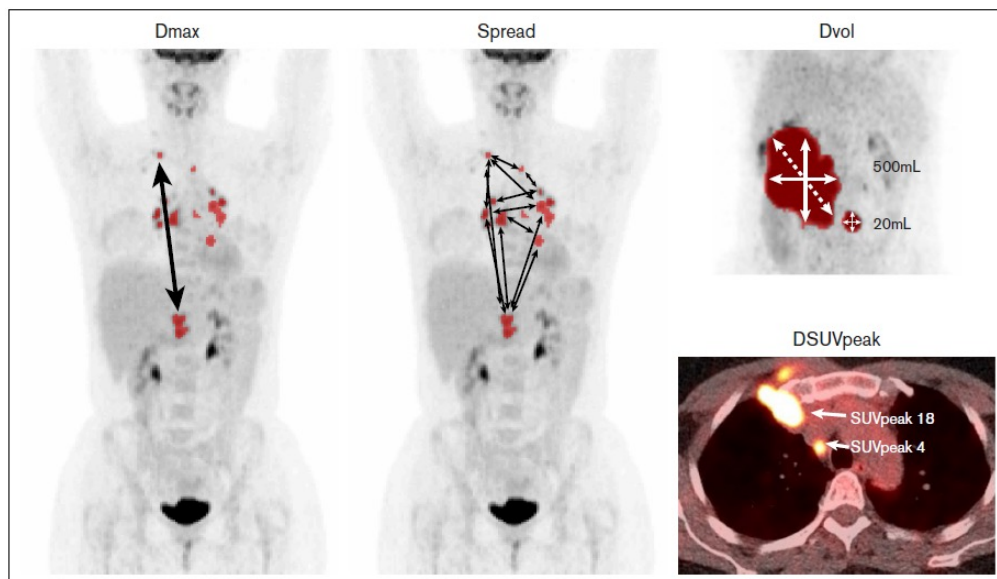
Prognostic parameters upon baseline PET scan

Baseline PET features could provide additional prognostic parameters in various lymphoma subtypes:

- tumor burden quantification: Total Metabolic Tumor Volume (TMTV);
- tumor dissemination: Distance maximum (Dmax) as the largest distance between two lesions.



A radiomics glossary



Variable	Definition
MTV	Metabolic tumor volume; The FDG-avid tumor volume
TLG	Total lesion glycolysis; MTV multiplied by SUVmean
SUVmean	The mean SUV value of the VOI
SUVmax	The SUV of the voxel with the highest SUV within the VOI
SUVpeak	The SUV of the 3mL with the highest SUV within the VOI (global peak)
TLRSUVmean	Tumor to liver ratio of the lesional SUVmean and the liver SUVmean
TLRSUVpeak	Tumor to liver ratio of the lesional SUVpeak and the liver SUVmean
Number of lesions	The number of separated lesion selections within the VOI
Dmax	The maximum distance between two lesions
DmaxBulk	The maximum distance between the largest lesion and any other lesion
Spread	The sum of the distance between all lesions
SpreadBulk	The sum of the distance between the largest lesion and all other lesions
Dvol	The difference in volume between the largest and the smallest lesion
VolSpread	The sum of the differences in volume between all lesions
VolSpreadBulk	The sum of the differences in volume between the largest lesion and all other lesions
DSUVmax	The difference in SUVmax between the lesion with the highest SUVmax and the lesion with the lowest SUVmax
DSUVmaxSum	The sum of the differences in SUVmax of all lesions
DSUVmaxBulk	The differences in SUVmax between the largest lesion and all other lesions
DSUVmaxSumBulk	The sum of the differences in SUVmax between the largest lesion and all other lesions
DSUVmaxSumHot	The sum of the differences in SUVmax between the lesion with the highest SUVmax and all other lesions
DSUVpeak	The difference in SUVpeak between the lesion with the highest SUVpeakmax and the lesion with the lowest SUVpeak
DSUVpeakSum	The sum of the differences in SUVpeak of all lesions
DSUVpeakBulk	The differences in SUVpeak between the largest lesion and all other lesions
DSUVpeakSumBulk	The sum of the differences in SUVpeak between the largest lesion and all other lesions
DSUVpeakSumHot	The sum of the differences in SUVpeak between the lesion with the highest SUVpeak and all other lesions

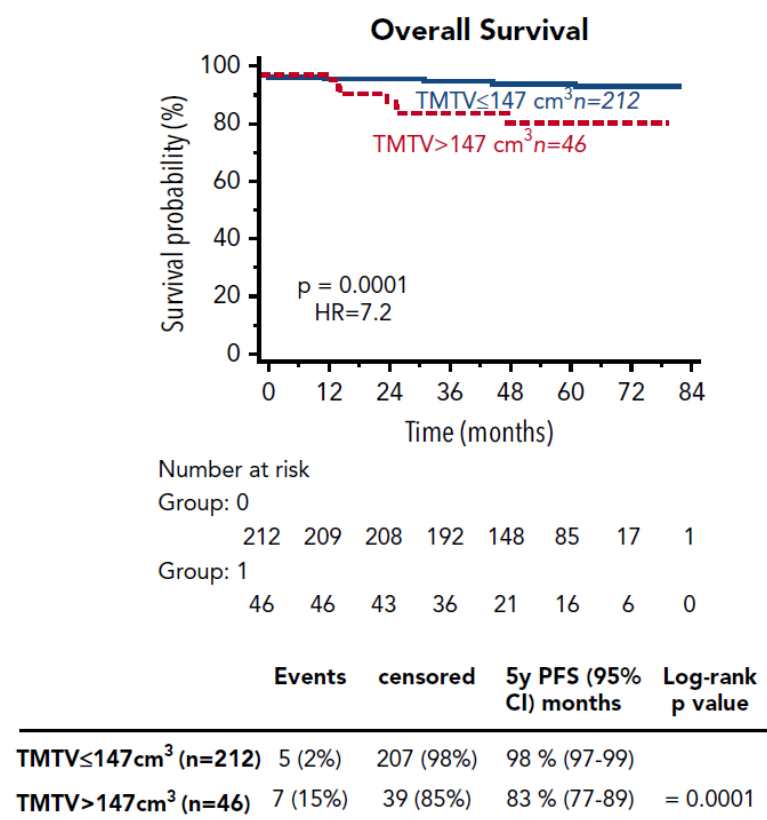
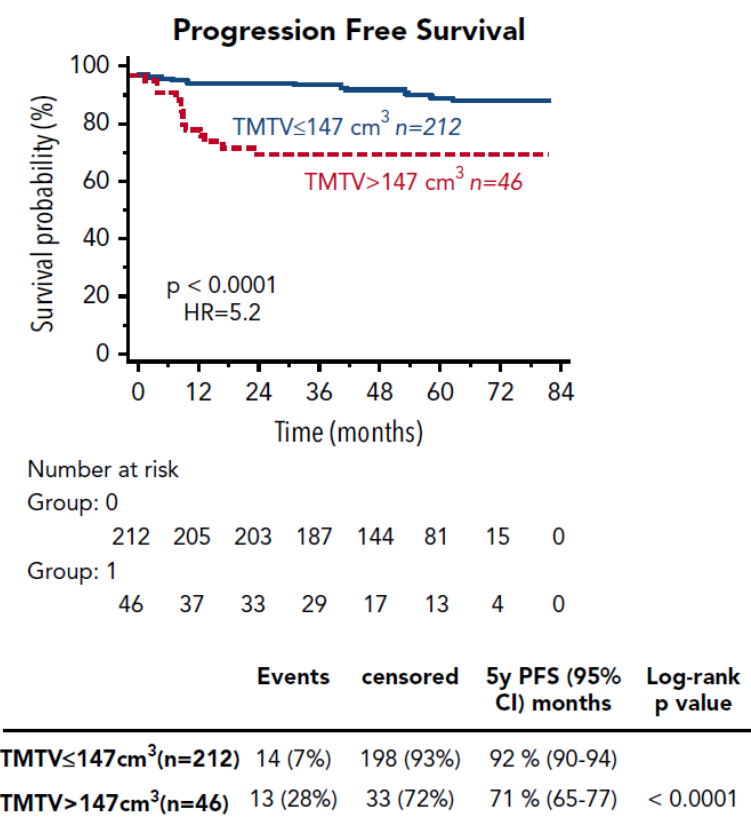
Prognostic value of baseline metabolic tumor volume in early-stage Hodgkin lymphoma in the standard arm of the H10 trial

Characteristics	Total population (N = 258)	Low TB TMTV ≤147 cm ³ (n = 212)	High TB TMTV >147 cm ³ (n = 46)	P
Median age (range), y	31 (15-71)	32 (15-71)	27 (17-63)	.009
Age ≥50 y (%)	34 (13)	31 (15)	3 (7)	.22
Male, n (%)	129 (50)	104 (49)	25 (54)	.62
Nodular sclerosis histology n (%)	207 (80)*	168 (79)	37 (80)	.68
Ann Arbor stage II, n (%)	198 (77)	157 (74)	41 (89)	.03
B symptoms, n (%)	85 (33)	60 (28)	25 (54)	.001
Median ESR (interquartile range), mm/h	26	23 (12-50)	49 (26-72)	.0001
≥4 Involved sites, n (%)	28 (11)	17 (8)	11 (24)	.004
Bulk mediastinum (M/T ≥ 0.35), n (%)	62 (24)	34 (16)	29 (63)	<.0001
U EORTC, n (%)	157 (61)	117 (55)	40 (87)	.0001
U GSHG, n (%)	177 (69)	133 (63)	44 (96)	<.0001
U NCCN, n (%)	164 (64)	121 (57)	43 (93)	<.0001
Positive iPET2 (DS 4-5), n (%)	21 (8)	13 (6)	8 (17)	.028

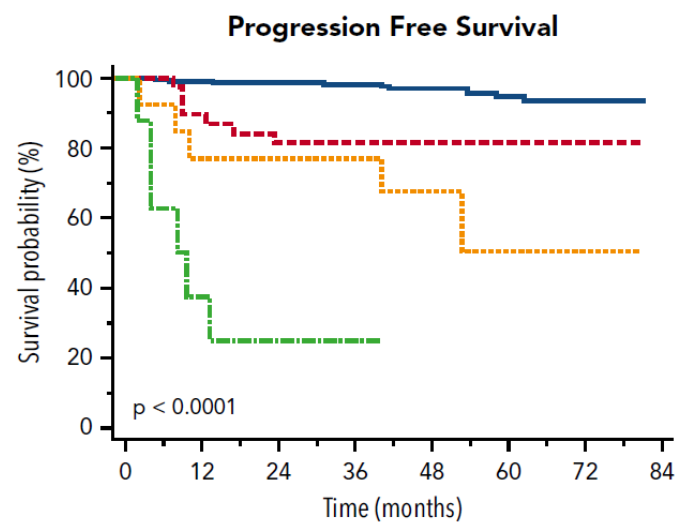
M/T, mass/thoracic ratio.

*Data not available for 2 patients.

Prognostic value of baseline metabolic tumor volume in early-stage Hodgkin lymphoma in the standard arm of the H10 trial

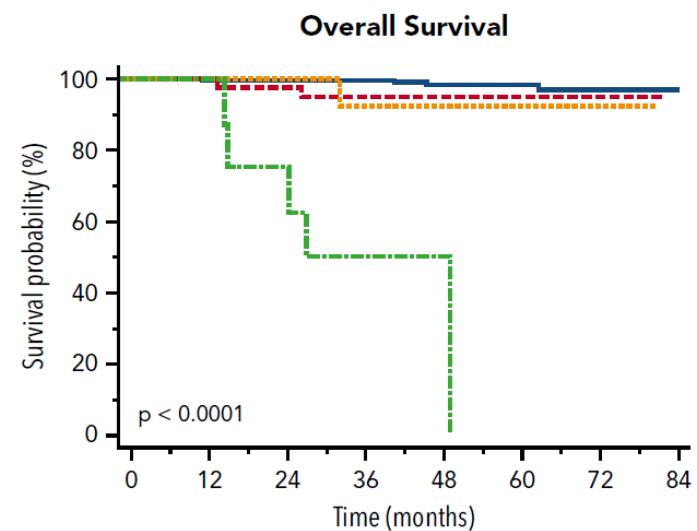


Prognostic value of baseline metabolic tumor volume in early-stage Hodgkin lymphoma in the standard arm of the H10 trial



Number at risk

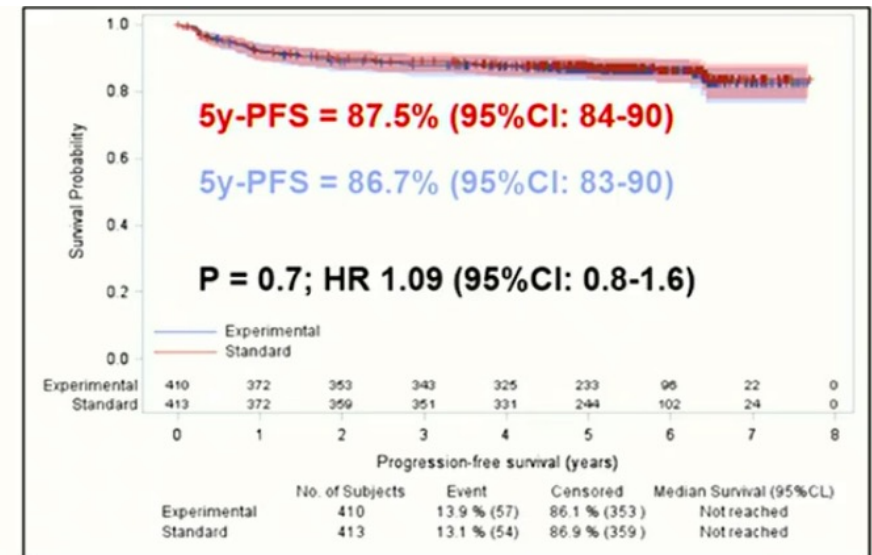
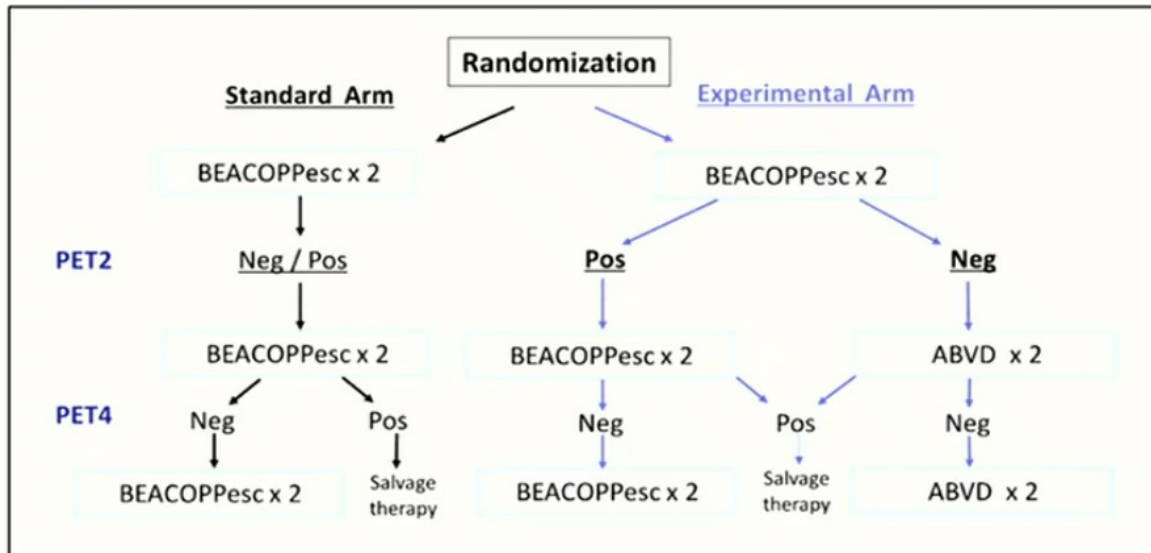
TMTV $\leq 147\text{cm}^3$ +iPET2 negative	199	195	193	178	137	79	14	0
TMTV $> 147\text{cm}^3$ +iPET2 negative	38	34	31	27	17	13	4	0
TMTV $\leq 147\text{cm}^3$ +iPET2 positive	13	10	10	9	7	2	1	0
TMTV $> 147\text{cm}^3$ +iPET2 positive	8	3	2	2	0	0	0	0



Number at risk

TMTV $\leq 147\text{cm}^3$ +iPET2 negative	199	196	195	181	139	82	16	1
TMTV $> 147\text{cm}^3$ +iPET2 negative	38	38	37	32	20	16	6	0
TMTV $\leq 147\text{cm}^3$ +iPET2 positive	13	13	13	11	9	3	1	0
TMTV $> 147\text{cm}^3$ +iPET2 positive	8	8	6	4	1	0	0	0

PROGNOSTIC VALUE OF BASELINE TOTAL METABOLIC TUMOR VOLUME (TMTV) AND TUMOR SPREAD (Dmax) IN ADVANCED HODGKIN LYMPHOMA: ANCILLARY STUDY OF THE AHL2011 TRIAL



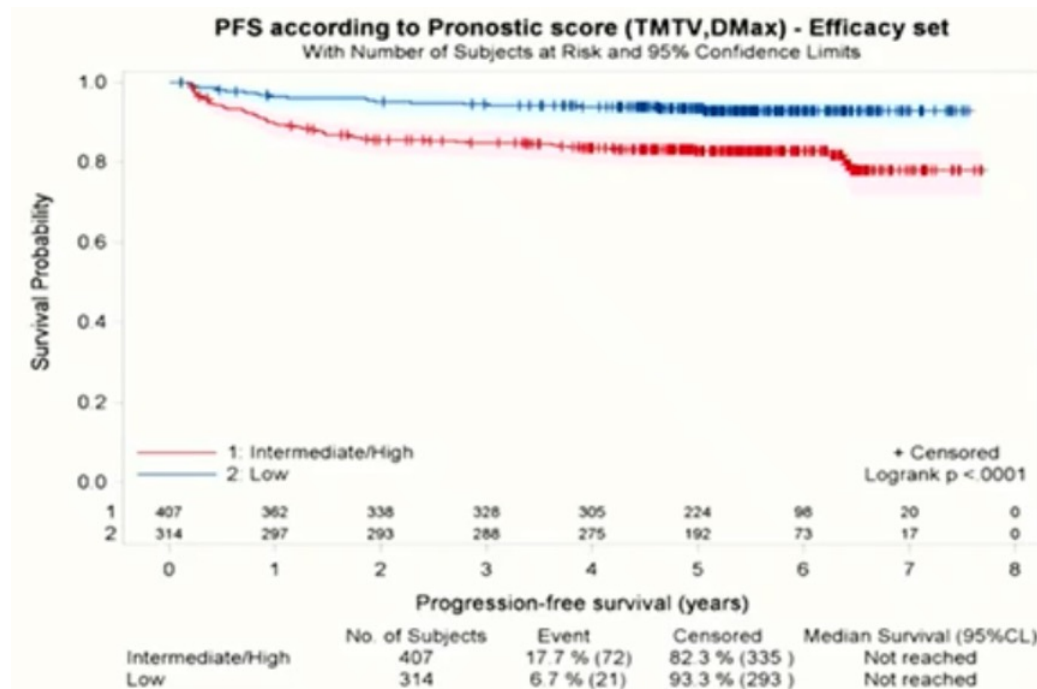
**PROGNOSTIC VALUE OF BASELINE TOTAL
METABOLIC TUMOR VOLUME (TMTV) AND TUMOR
SPREAD (D_{max}) IN ADVANCED HODGKIN LYMPHOMA:
ANCILLARY STUDY OF THE AHL2011 TRIAL**

	Median (IQR)	Cutoff	% patients (out of the 721 pts)
TMTV (mL)	210 (123-390)	220	48% with high TMTV (≥ 220)
D _{max} (mm ⁻¹)	220 (160-335)	325	26% with high D _{max} (≥ 325)

Two subgroups of patients are identifiable:

- patients with low TMTV and low Dmax;
- patients with either high TMTV and/or high Dmax.

PROGNOSTIC VALUE OF BASELINE TOTAL METABOLIC TUMOR VOLUME (TMTV) AND TUMOR SPREAD (Dmax) IN ADVANCED HODGKIN LYMPHOMA: ANCILLARY STUDY OF THE AHL2011 TRIAL



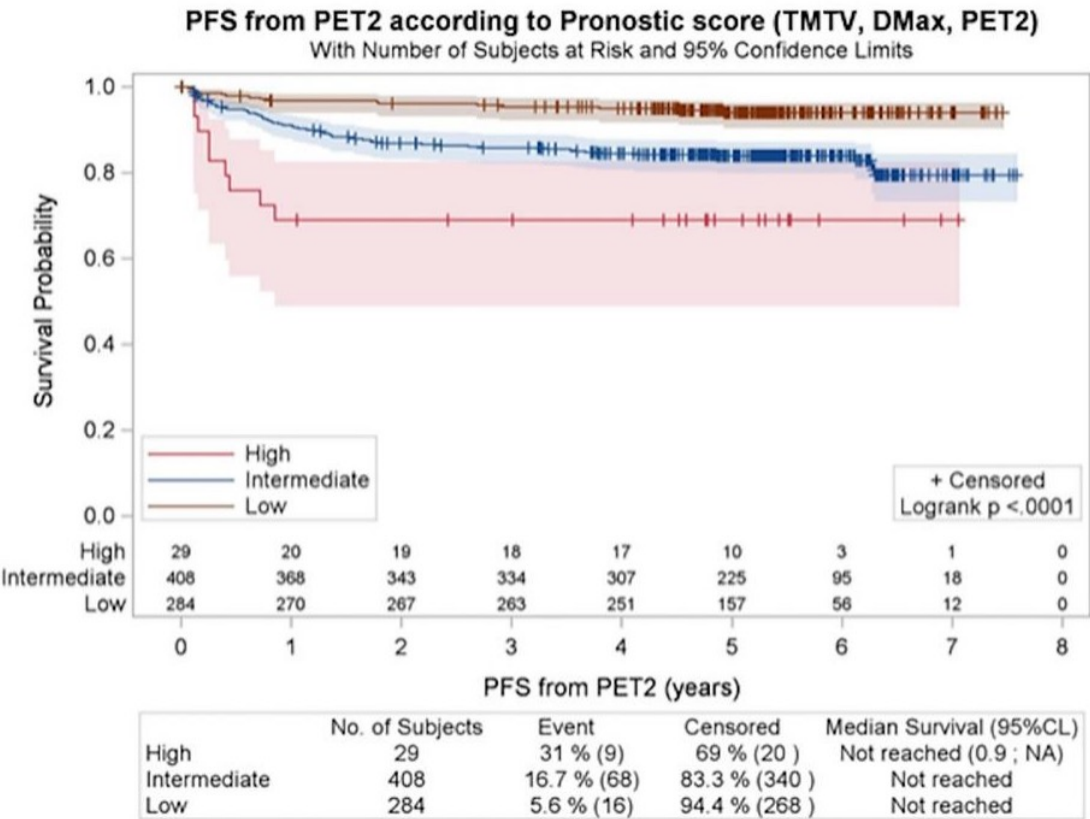
**5-year PFS: 92% vs 83.4%,
HR=2.2 [95%CI: 1.4-3.6],
p<0.001**

**PROGNOSTIC VALUE OF BASELINE TOTAL
METABOLIC TUMOR VOLUME (TMTV) AND TUMOR
SPREAD (Dmax) IN ADVANCED HODGKIN LYMPHOMA:
ANCILLARY STUDY OF THE AHL2011 TRIAL**

	TMTV		Dmax	
	Low TMTV ($<220\text{mL}$)	High TMTV ($\geq 220\text{mL}$)	Low Dmax ($<325\text{mm}^{-1}$)	High Dmax ($\geq 325\text{mm}^{-1}$)
PET2 positive*	10% (n=37)	16% (n=57)	11% (n=58)	19% (n=36)
PET2 negative*	90% (n=337)	84% (n=290)	89% (n=473)	81% (n=154)
P-value (Fisher exact test)	0.011		0.008	

*In AHL trial, PET2 evaluations were centrally reviewed and interpreted using modified Deauville score (partial response if residual uptake > 140% liver background)

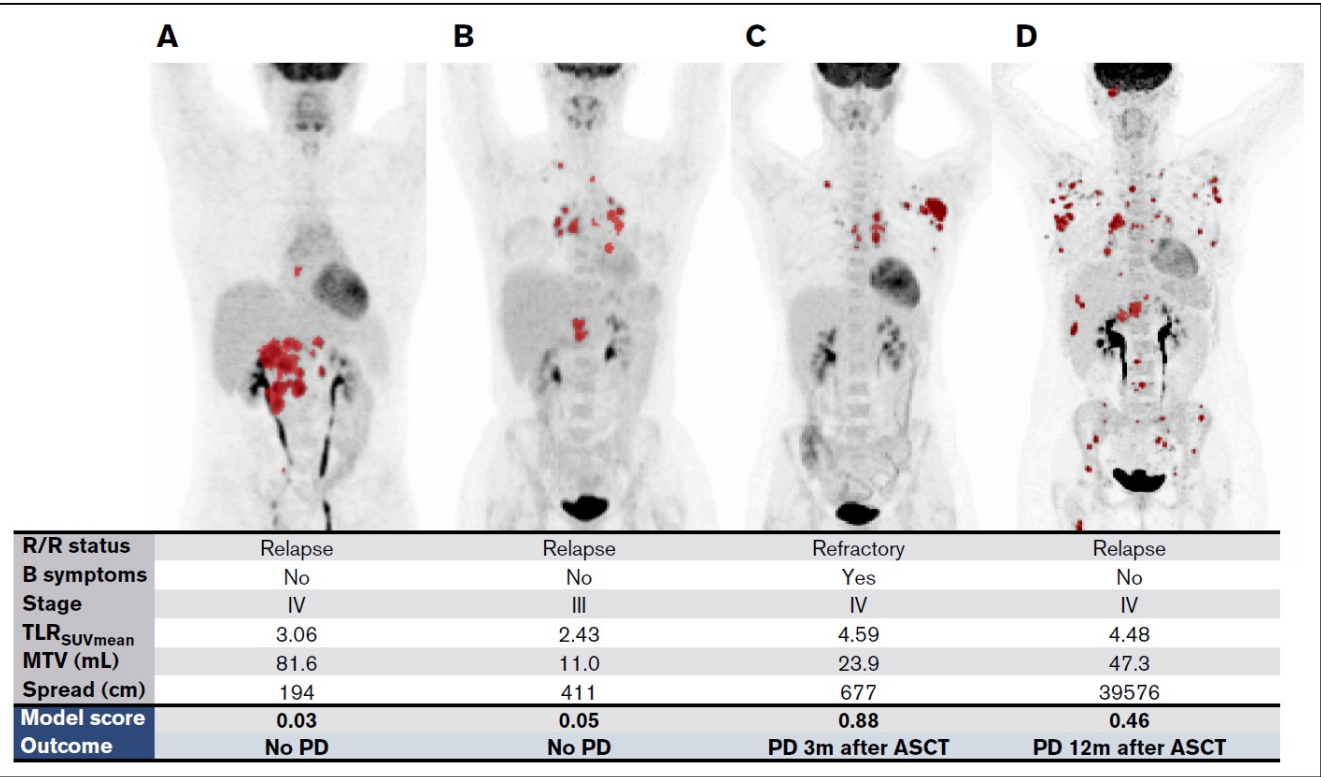
PROGNOSTIC VALUE OF BASELINE TOTAL METABOLIC TUMOR VOLUME (TMTV) AND TUMOR SPREAD (Dmax) IN ADVANCED HODGKIN LYMPHOMA: ANCILLARY STUDY OF THE AHL2011 TRIAL



	No of pts (%)	5-year PFS	PFS event rate
0 risk factor	284 (39%)	94%	6%
1 or 2 risk factors	408 (57%)	84%	17%
3 risk factors	29 (4%)	69%	31%

Prognostic model using ¹⁸F-FDG PET radiomics predicts progression-free survival in relapsed/refractory Hodgkin lymphoma

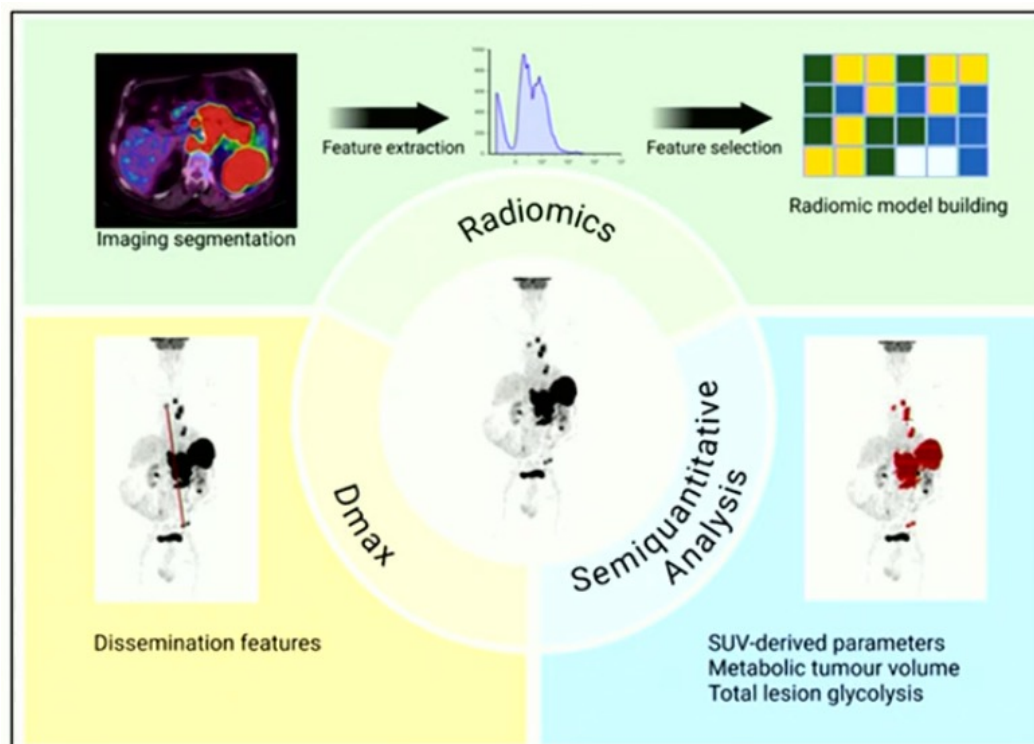
Study	Training		Validation		Total		P value
	(n = 113)		(n = 69)		(n = 182)		
	No.	%	No.	%	No.	%	
BV-DHAP	58	51	0	0	58	32	<.001
BV-ICE	55	49	0	0	55	30	
ICE-GVD	0	0	69	100	69	38	
Female sex	61	54	32	46	93	51	.319
Median age, (range)	30 (13-65)		34 (18-66)		31 (13-66)		.175
Primary refractory	55	50	25	37	80	45	.062
Ann Arbor stage							.002
I	10	9	1	1	11	6	.042
II	46	41	43	62	89	49	.004
III	19	17	2	3	21	12	.004
IV	38	34	23	33	61	34	.970
Extranodal disease	44	39	25	36	69	38	.715
B symptoms	28	25	7	10	35	20	.011



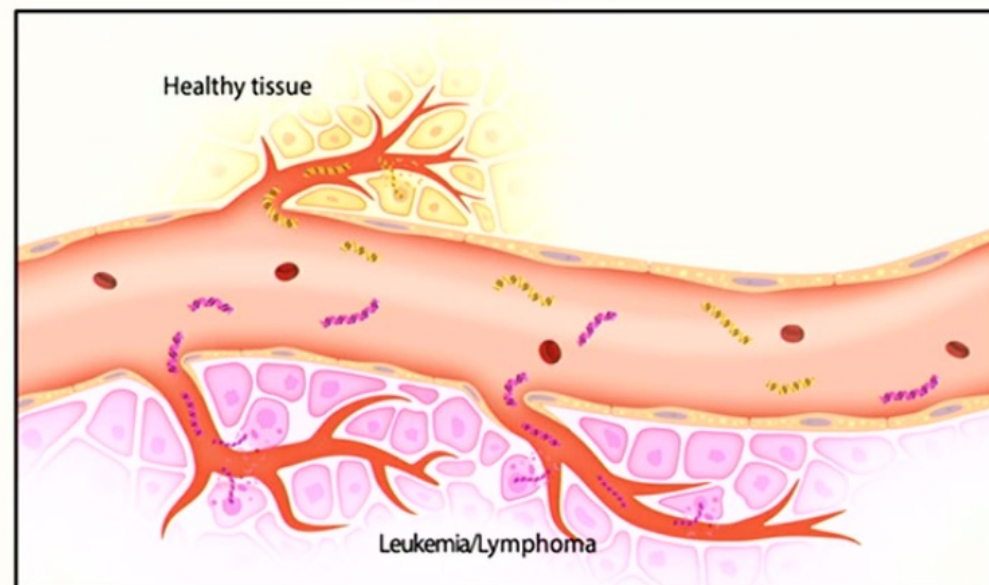
$$-2.472 - [2.478 * (\text{Relapsed}=1, \text{refractory}=0)] + [1.010 * (\text{B symptoms} = 1, \text{no B symptoms} = 0)] - [0.384 * \log(\text{MTV in uL})] + [0.413 * \log(\text{Spread})] + [2.409 * \log(\text{SUVmean}/\text{liverSUVmean})].$$

Ann Arbor stage: not part of the final model as outperformed by spread.
TLR: tumor to liver ratio.

PET parameters



Circulating tumor DNA (ctDNA)



Prognostication for early- and advanced-stage Hodgkin lymphoma... What's new?

- Prognostic models built upon continuous variables better stratify patients according to risk in both advanced- and early-stage disease.
- These models are complex, but can be easily applied at bedside by using tools available online.
- Models based on binary cutoffs are nowadays outdated and reduce their performance in the era of new drugs (e.g. brentuximab vedotin, checkpoint inhibitors) used frontline.
- *Interim* PET scan maintains its value if PET-adapted strategies are adopted (e.g. stage IIB, III), although its predictive value may be questioned when new drugs are applied frontline.
- New PET parameters (*radiomics*) evaluated at baseline may offer new prognostic interpretations and emerge as novel predictive factors, both in treatment naïve and relapsed/refractory patients.