



Third international workshop on
interim-PET in lymphoma

PET AND BIOMARKERS

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FDG-PET studies

Evaluate the prognostic role of an early interim fluorodeoxyglucose-PET scan in advanced Hodgkin's Lymphoma

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JOURNAL OF CLINICAL ONCOLOGY

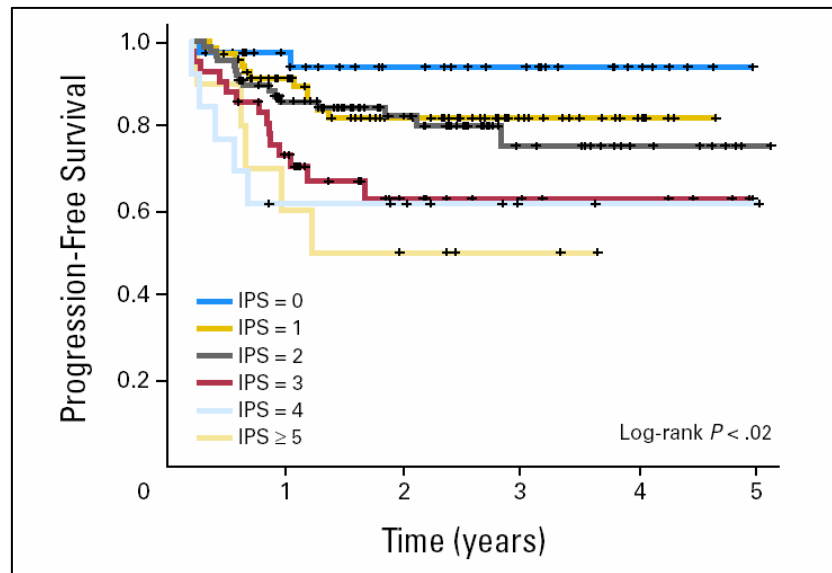
ORIGINAL REPORT

Early Interim 2-[¹⁸F]Fluoro-2-Deoxy-D-Glucose Positron Emission Tomography Is Prognostically Superior to International Prognostic Score in Advanced-Stage Hodgkin's Lymphoma: A Report From a Joint Italian-Danish Study

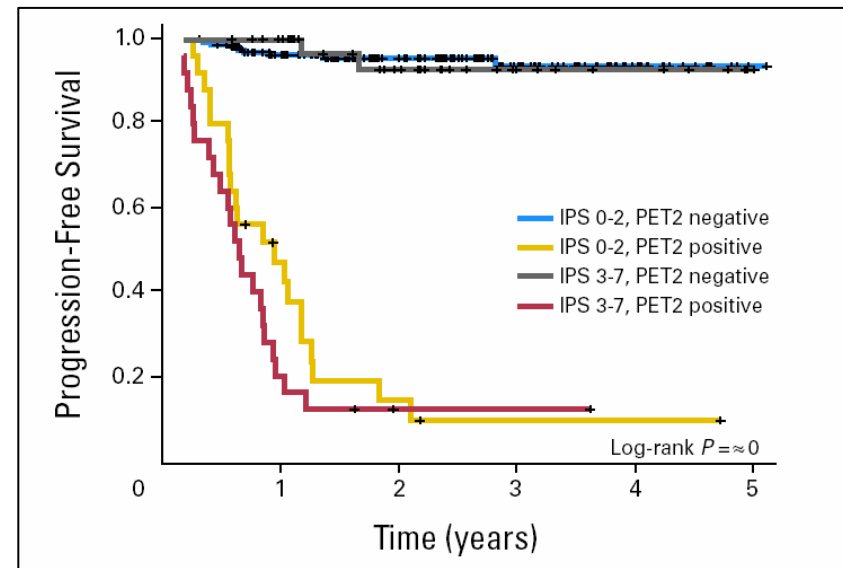
Andrea Gallamini, Martin Hutchings, Luigi Rigacci, Lena Specht, Francesco Merli, Mads Hansen, Caterina Patti, Annika Loft, Francesco Di Raimondo, Francesco D'Amore, Alberto Biggi, Umberto Vitolo, Caterina Stelitano, Rosario Sancetta, Livio Trentin, Stefano Luminari, Emilio Iannitto, Simonetta Viviani, Ivana Pierri, and Alessandro Levis

Prognostic biologic factors in Hodgkin's lymphoma

Progression-Free Survival



According to International Prognostic Score



According to PET-2 results for patients with a low or a high IPS

Prognostic biologic factors in Hodgkin's lymphoma

PubMed

interim PET and Hodgkin lymphoma

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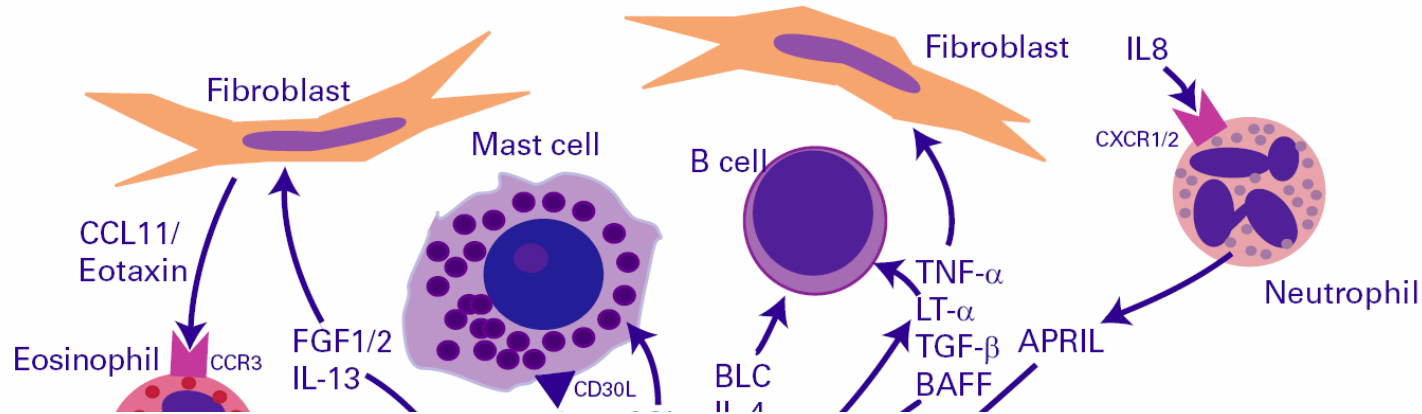
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1. [The role of FDG-**PET** in early and late therapy assessment of patients with advanced **Hodgkin Lymphoma** treated with BEACOPP.](#)
Markova J, Kahraman D, Kobe C, Skopalova M, Mocikova H, Klaskova K, Dedekova K, Eich H, B LI B, Dietlein M, Kozak T.
Leuk **Lymphoma**. 2011 Jul 7. [Epub ahead of print]
PMID: 21740300 [PubMed - as supplied by publisher]
2. [Epratuzumab with rituximab, cyclophosphamide, doxorubicin, vincristine, and prednisone chemotherapy \(ER-CHOP\) in patients with previously untreated diffuse large B-cell **lymphoma**.](#)
Micallef IN, Maurer MJ, Wiseman GA, Nikcevic DA, Kurtin PJ, Cannon MW, Perez DG, Soori GS, Link BK, Habermann TM, Witzig TE.
Blood. 2011 Jun 14. [Epub ahead of print]
PMID: 21673350 [PubMed - as supplied by publisher]
3. [Report of satellite workshop on **interim-PET** in **Hodgkin lymphoma**: 8th International Symposium on **Hodgkin Lymphoma**, Cologne, 23 October 2010.](#)
Gallamini A, O'Doherty M.
Leuk **Lymphoma**. 2011 Apr;52(4):583-6. Review. No abstract available.
PMID: 21438829 [PubMed - indexed for MEDLINE]
4. [Radioguided lymph node biopsy of a chemoresistant lymph node detected on **interim** FDG **PET-CT** in **Hodgkin lymphoma**.](#)
Györke T, Kollár A, Bottlik G, Szepesi A, Bodó I, Masszi T, Bérczi V, Garai I.
Int J Hematol. 2011 Apr;93(4):545-50. Epub 2011 Mar 1.
PMID: 21360009 [PubMed - indexed for MEDLINE]

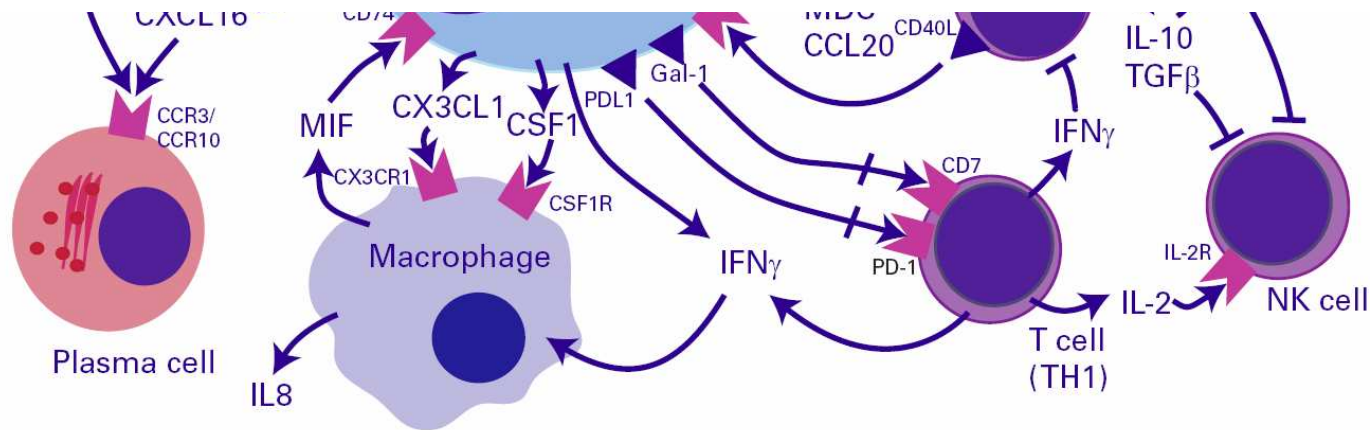
Problems: costs and availability as well as *ad interim* test

“If at time of diagnosis we could identify patients who are destined to have a poor response to treatment, most patients could be spared a combination of therapies or radiotherapy with its attendant long-term toxic effects”. De Vita
NEJM 2010

CHL: cross-talk between HRSCs and microenvironment



Biomarkers referred to neoplastic cells and microenvironmental components



Immunoistochemical studies

- BCL2
- CD20
- p53
- EBV
- TOP2A
- HGAL
- IRF4
- HLA class II
- FOXP3
- Tia1/GyB

BLOOD, 15 JANUARY 2003 • VOLUME 101, NUMBER 2

Hodgkin and Reed-Sternberg cells harbor alterations in the major tumor suppressor pathways and cell-cycle checkpoints: analyses using tissue microarrays

Juan F. García, Francisca I. Camacho, Manuel Morente, Máximo Fraga, Carlos Montalbán, Tomás Álvaro, Carmen Bellas, Ángel Castaño, Ana Díez, Teresa Flores, Carmen Martín, Miguel A. Martínez, Ana I. Sáez, Lydia Sánchez, and Miguel A. Piris, for the Spanish

Vol. 11, 1467–1473, February 15, 2005 Clinical Cancer Research

Outcome in Hodgkin's Lymphoma Can Be Predicted from the Presence of Accompanying Cytotoxic and Regulatory T Cells

Tomás Álvaro,¹ Marylène Lejeune,¹ M^a Teresa Salvadó,¹ Ramón Bosch,¹ Juan F. García,² Joaquín Jaén,¹ Alison H. Banham,⁵ Giovanna Roncador,³ Carlos Montalbán,⁴ and Miguel A. Piris²

VOLUME 22 • NUMBER 9 • MAY 1 2004

JOURNAL OF CLINICAL ONCOLOGY ORIGINAL REPORT

Influence of Biologic Markers on the Outcome of Hodgkin's Lymphoma: A Study by the Spanish Hodgkin's Lymphoma Study Group

Carlos Montalbán, Juan F. García, Víctor Abaira, Leocricia González-Camacho, Manuel M. Morente, Jose L. Bello, Eulogio Conde, Miguel A. Cruz, Ramón García-Sanz, José García-Laraña, Carlos Grande, María Llanos, Rafael Martínez, Eduardo Flores, Miguel Muñoz, Concepción Poveda, Concepción Pavia,

BLOOD, 1 OCTOBER 2005 • VOLUME 106, NUMBER 7

Impact of tumor Epstein-Barr virus status on presenting features and outcome in age-defined subgroups of patients with classic Hodgkin lymphoma: a population-based study

Ruth F. Jarrett, Gail L. Stark, Jo White, Brian Angus, Freda E. Alexander, Andrew S. Krajewski, June Freeland, G. Malcolm Taylor, and Penelope R. A. Taylor, for the Sc

Prognostic Significance of Cell Proliferation and Apoptosis-Regulating Proteins in Epstein-Barr Virus Positive and Negative Pediatric Hodgkin Lymphoma

SAFIYE AKTAŞ, M.D., Ph.D.,¹ AYDANUR KARGI, M.D.,² NUR OLGUN, M.D.,² GULDEN DINİZ, M.D.,¹ AYŞE ERBAY, M.D.,¹ and CANAN VERGIN, M.D.¹

haematologica | 2008; 93(2) | 193 |

Correlation of high numbers of intratumoral FOXP3⁺ regulatory T cells with improved survival in germinal center-like diffuse large B-cell lymphoma, follicular lymphoma and classical Hodgkin's lymphoma

Alexandar Tzankov,¹ Cecile Meier,¹ Petra Hirschmann,¹ Philip Went,¹ Stefano A. Pileri,² and Stephan Dirnhofer¹

CLINICAL OBSERVATIONS, INTERVENTIONS, AND THERAPEUTIC TRIALS

BCL-2 expression in Hodgkin and Reed-Sternberg cells of classical Hodgkin disease predicts a poorer prognosis in patients treated with ABVD or equivalent regimens

George Z. Rassidakis, L. Jeffrey Medeiros, Theodoros P. Vassilikopoulos, Simonetta Viviani, Valeria Bonfante, Gianpaolo Nadali, Marco Herling, Maria K. Angelopoulou, Roberto Giardini, Marco Chilosi, Christos Kittas, Timothy J. McDonnell, Gianni Bonadonna, Alessandro M. Gianni, Giovanni Pizzolo, Gerassimos A. Pangalis, Fernando Cabanillas, and Andreas H. Sarris

VOLUME 23 • NUMBER 16 • JUNE 1 2005

JOURNAL OF CLINICAL ONCOLOGY ORIGINAL REPORT

Expression of bcl-2 in Classical Hodgkin's Lymphoma: An Independent Predictor of Poor Outcome

Stephen J. Sup, Carlos A. Alemany, Brad Pohlman, Paul Elson, Serena Malhi, Snehal Thakkar, Roxanne Steinle, and Eric D. Hsi

Prognostic biologic factors in Hodgkin's lymphoma

Steidl C, Connors JM, Gascoyne RD. JCO 2011, 29:1812-26

Table 2. Biomarkers With Outcome Correlations Described in the Literature

	Gene Expression Profiling		
	Main Gene Components	Outcome Correlation	Reference No.
Angiogenic signature	<i>ADH1B, CD93, SRPX, PLA2G2A, GPR126</i>	Adverse (primary treatment failure)	132
Adipocyte signature	<i>GLUL, MGST1, COL1A2, FABP4</i>	Adverse (primary treatment failure)	132
Fibroblast function/extracellular matrix remodeling	Adverse: <i>MMP2, MMP3, TIMP1, COL1A1, COL4A1, COL4A2, COL5A1, COL18A1, COL16A1, MFAP2, THBS1/2, FN1, EDNRA, ITGB5, LAMA4</i> ; favorable: <i>TIMP4, SPON1, LAMB1, TACR1, CCL26</i>	Discordant: adverse/favorable (primary treatment outcome)	142,148
B-cell signature	<i>BCL11A, BANK1, STAP1, BLNK, FCER2, CD24, CCL21</i>	Favorable (primary treatment outcome)	132,140
Cytotoxic T-cell signature	<i>CD3D, CD8B1, CTSL, CD26, SH2D1A, IFI16, RGS13, CR2, ELL3, CCDC23, PPM1L, TRA@, PIK3CA</i>	Adverse (primary treatment outcome)	131,132,142
Plasmacytoid dendritic cells	<i>ITM2A, SRPX, CTSB, APP</i>	Adverse (primary treatment outcome)	132
Macrophage signature	<i>ALDH1A1, LYZ, STAT1, ITGA4, CCL13, MS4A4A, CCL23, VCAN, HSP90AB3P, HSP90AB1, CTSB, CFL1, JMJD6, MAPK7, IKBKG, RAB7A, RXRA, MAPK13</i>	Adverse (primary treatment outcome)	131,132,142

131. Sánchez-Espiridión B, Sánchez-Aguilera A, Montalbán C, et al: A TaqMan low-density array to predict outcome in advanced Hodgkin's lymphoma using paraffin-embedded samples. Clin Cancer Res 15:1367-1375, 2009

132. Steidl C, Lee T, Shah SP, et al: Tumor-associated macrophages and survival in classic Hodgkin's lymphoma. N Engl J Med 362:875-885, 2010

140. Chetaille B, Bertucci F, Finetti P, et al: Molecular profiling of classical Hodgkin lymphoma tissues uncovers variations in the tumor microenvironment and correlations with EBV infection and outcome. Blood 113:2765-3775, 2009

142. Sánchez-Aguilera A, Montalbán C, de la Cueva P, et al: Tumor microenvironment and mitotic checkpoint are key factors in the outcome of classic Hodgkin lymphoma. Blood 108:662-668, 2006

148. Devillard E, Bertucci F, Tremat P, et al: Gene expression profiling defines molecular subtypes of classical Hodgkin's disease. Oncogene 21:3095-3102, 2002

Macrophages predict treatment outcome in Hodgkin's lymphoma

Christian Steidl, Pedro Farinha, and Randy D. Gascoyne

Departments of Pathology and Experimental Therapeutics, British Columbia Cancer Agency and the BC Cancer Research Centre, BC Cancer Agency, University of British Columbia, Vancouver, BC, Canada

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Haematologica 2010, 96:186-9

Markers used	Method	#	Outcome correlation	Reference
PNA	Histochemistry	43	Adverse (refractory disease, early relapse)	Ree <i>et al.</i> , Cancer 1985 ¹⁰
STAT1, ALDH1A1	GE, IHC	235	Adverse (disease-specific survival)	Sanchez-Aguilera <i>et al.</i> , Blood 2006 ¹¹
LYZ, STAT1, ALDH1A1	GE, IHC	194	Adverse (refractory disease, early relapse)	Sanchez-Espiridion <i>et al.</i> , Clinical Cancer Research 2009 ¹³
CD68	IHC	166	Adverse (progression-free survival, disease-specific survival)	Steidl <i>et al.</i> , NEJM 2010 ⁹
LYZ, STAT1	GE	262	Favorable (failure-free survival)	Sanchez-Espiridion <i>et al.</i> , Blood 2010 ¹²
CD68, CD163	IHC	288	Adverse (event-free survival, overall survival)	Kamper <i>et al.</i> , Haematologica 2011 ⁸
CD68	IHC	59	Adverse (refractory disease)	Benedicte <i>et al.</i> , Blood 2010 [abstr.] ³⁴
CD68 (also in combination with FOXP3)	IHC	122	Adverse (freedom from treatment failure, overall survival)	Greaves <i>et al.</i> , Blood 2010 [abstr.] ²⁵
CD68	IHC	144	Adverse (event-free survival, disease-specific survival)	Yoon <i>et al.</i> , Blood 2010 [abstr.] ³⁵
CD68	IHC	105	Adverse (overall survival)	Tzankov <i>et al.</i> [personal communication]
CD68	IHC	45	Adverse (progression-free survival)	Hohaus & Larocca [personal communication]
CD68	IHC	153	Adverse (overall survival, progression-free survival)	Farinha <i>et al.</i> [abstr.] ³⁶

PNA: peanut agglutinin, GE: gene expression (mRNA), IHC: immunohistochemistry.

Macrophages predict treatment outcome in Hodgkin's lymphoma

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E-mail: rgascoyn@bccancer.bc.ca doi:10.3324/haematol.2010.033316

Haematologica 2010, 96:186-9

Antibody and scoring system

Further study is needed to determine the optimal antigen (e.g. CD68 versus CD163), anti-CD68 antibody clone (e.g. KP1 versus PGM1) and scoring thresholds (e.g. manual versus computer-assisted) for detecting HL-associated macrophages.

Biomarker combination

The preliminary results of an immunohistochemistry study combining two markers, CD68 and FOXP3 (a marker for regulatory T cells), were presented at the ASH 2010 meeting. The authors showed that a combined FOXP3/CD68 immunohistochemistry score was an improvement over the predictive value of the individual markers alone and that this score was applicable to both limited and advanced-stage disease. The value of this com-

EBV infection

In a re-analysis of our data we were able to confirm a relationship between increased tumor-associated macrophages and EBV positivity; however, virtually all of our cases were of the nodular sclerosis subtype and EBV alone was not associated with treatment outcome (*unpublished observations, 2010*). EBV infection of HRS cells has been reported in up to 60% of patients and is more frequent in mixed cellularity subtype, although varying with geographical location, age, gender, clinical stage and histological subtype.¹⁴ The impact of EBV infection on outcome remains controversial, but appears to be dependent on age.

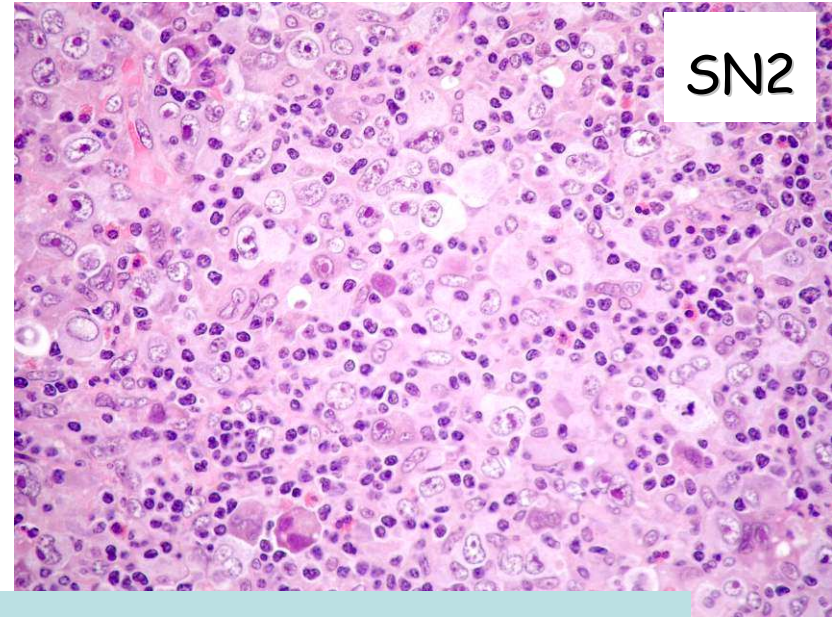
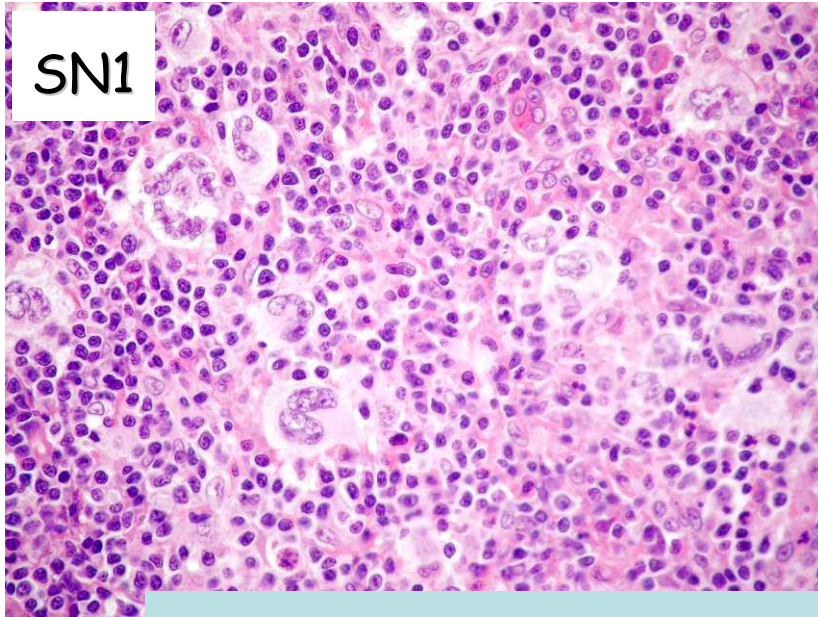
Bologna study

- Biopsy samples from cHL patients at diagnosis enrolled by 13 Italian and 3 Danish haematological centres
- Construction of TMAs to collect cases of interest in the same block and optimization of immunohistochemical procedures
- Ab tested:
 - 11 proteins encoded by genes shown as prognostically relevant by DNA-microarray studies (STAT1, PCNA, SAP, TOP2A, RRM2, CDC2, MAD2L1, ALDH1A1, CD68, CD163, and BCL11a)
 - 9 markers previously reported to have prognostic value in conventional studies (CD20, EBER, Bcl-2, p53, PD1, FOXP3, TIA1, Granzyme B, and Perforin)
- The molecules were assessed in both neoplastic (HRSC) and micro-environmental cell (MEC) components
- Evaluation of the prognostic impact of such markers on Hodgkin's lymphoma outcome
- Comparison with the predictive value of ad interim PET
- Construction of a predictive model

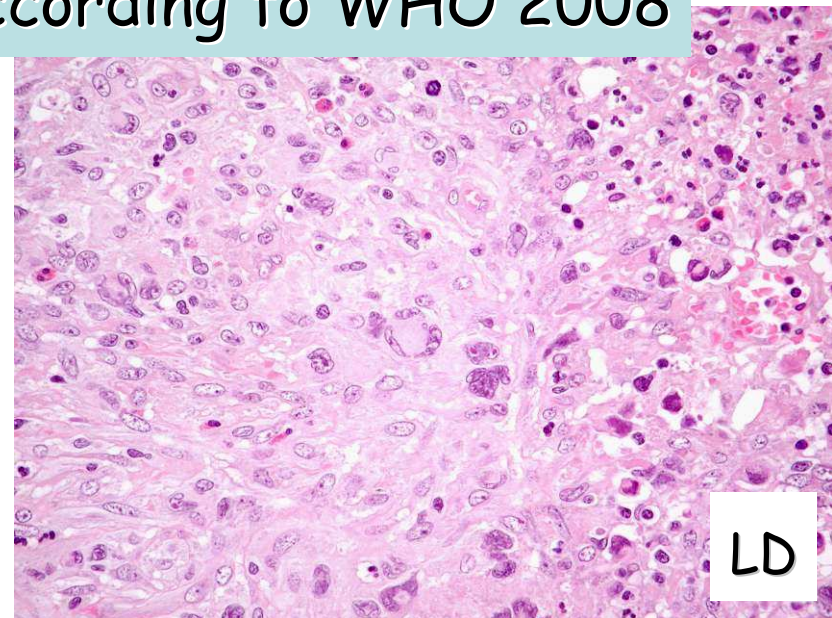
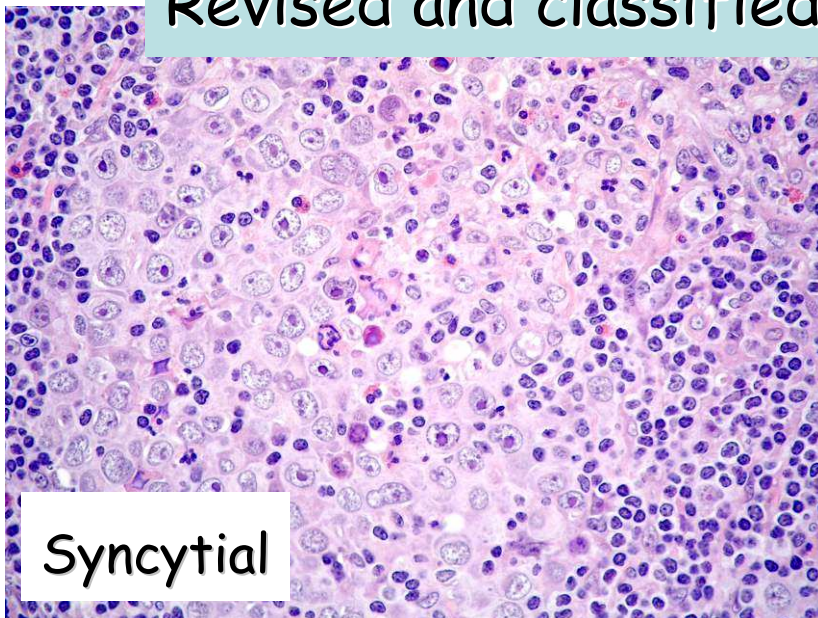
Inclusion criteria

- Diagnosis of **cHL**
- **HIV negative** status
- **Biopsy sample at diagnosis** available
- **Clinical and follow-up (FU) data** were always available
- Treatment with courses of **ABVD with or without radiotherapy**
- FDG-PET appraisal of the treatment response performed after two courses of chemotherapy (**PET-2**) available

209 cases enrolled



Revised and classified according to WHO 2008



Patients' characteristics

Age, years		Ann Arbor Stage	
mean	36	I	9 (4,3%)
median	32	II	118 (56,5%)
range	14-80	III	50 (23,9%)
Sex		IV	32 (15,3)
male	103 (49,3%)	Constitutional symptoms	
female	106 (50,7%)	A	110 (52,6%)
Follow-up, months		B	99 (47,4)
mean	50,26	Bulky disease	
median	52,30	Yes	42 (20.1%)
range	3-93	No	167 (79,9%)
Histologic subtype		First-line treatment	
CHL, nos	19 (9,1%)	ABVD	209 (100%)
NS-nos	10 (4,8%)	RT	52 (24,9)
NS-cellular phase	9 (4,3%)	PET after 2 cycles	
NS-1	93 (44,5%)	Negative	171 (81,9%)
NS-2	37 (17,7%)	Positive	38 (18,1%)
NS-syncytial variant	8 (3,8%)	Clinical outcome	
MC	29 (13,9%)	Failure	49 (23.4%)
LR	1 (0,5%)	<i>progression</i>	31
LD	3 (1,4%)	<i>relapse</i>	18

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- The molecules were assessed in both neoplastic (HRSC) and microenvironmental cell (MEC) components
- Evaluation of the prognostic impact of such markers on Hodgkin's lymphoma outcome
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Scoring system for HRSC markers

score 0 : 0%(+)

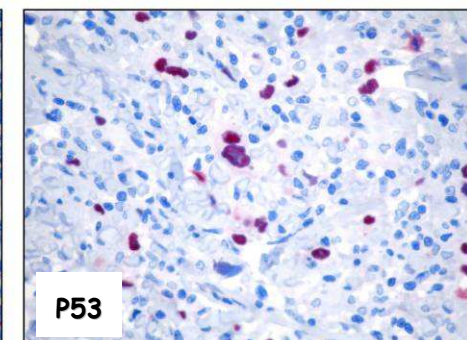
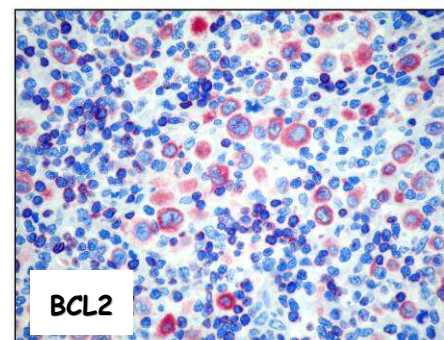
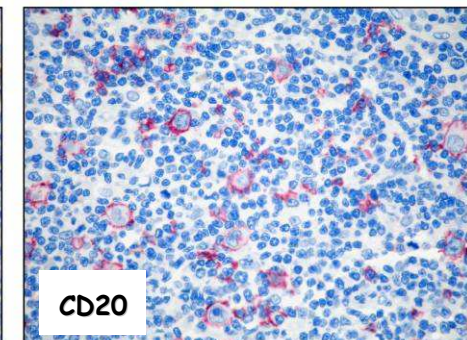
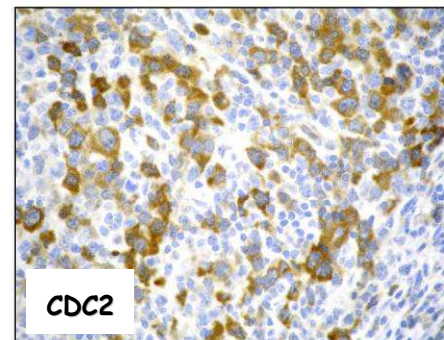
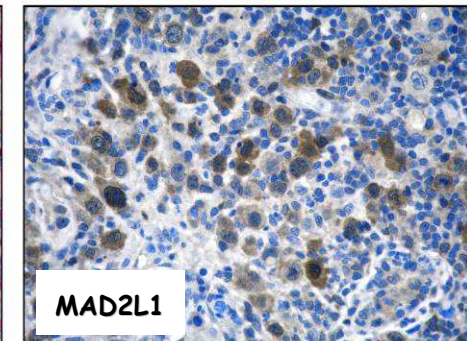
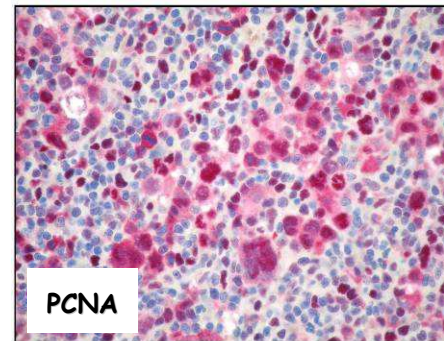
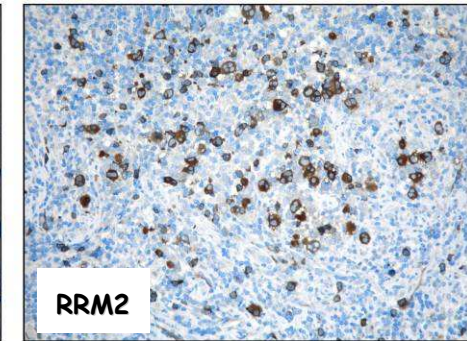
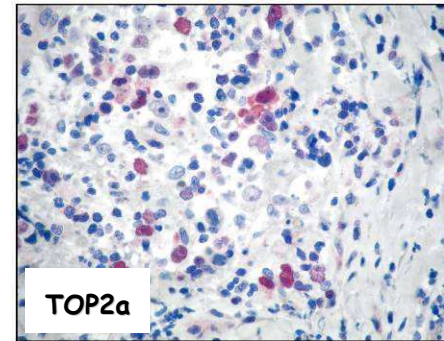
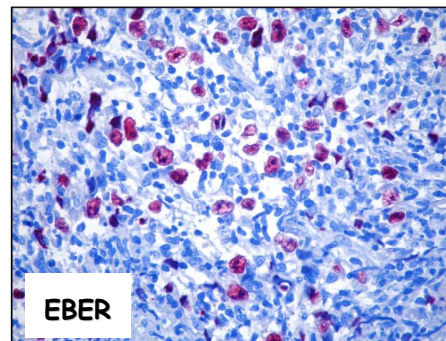
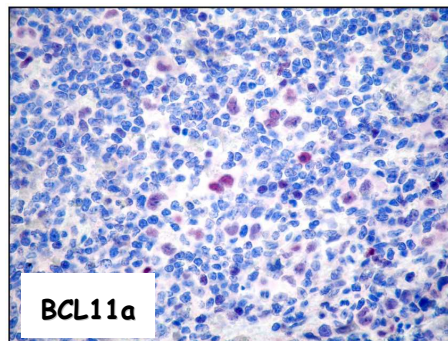
score 1 : 1-9%(+)

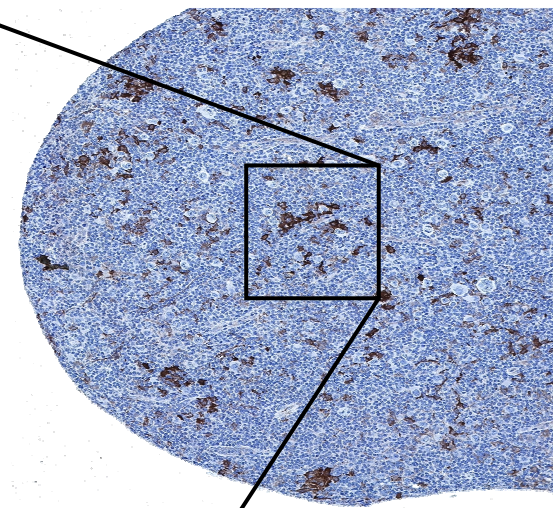
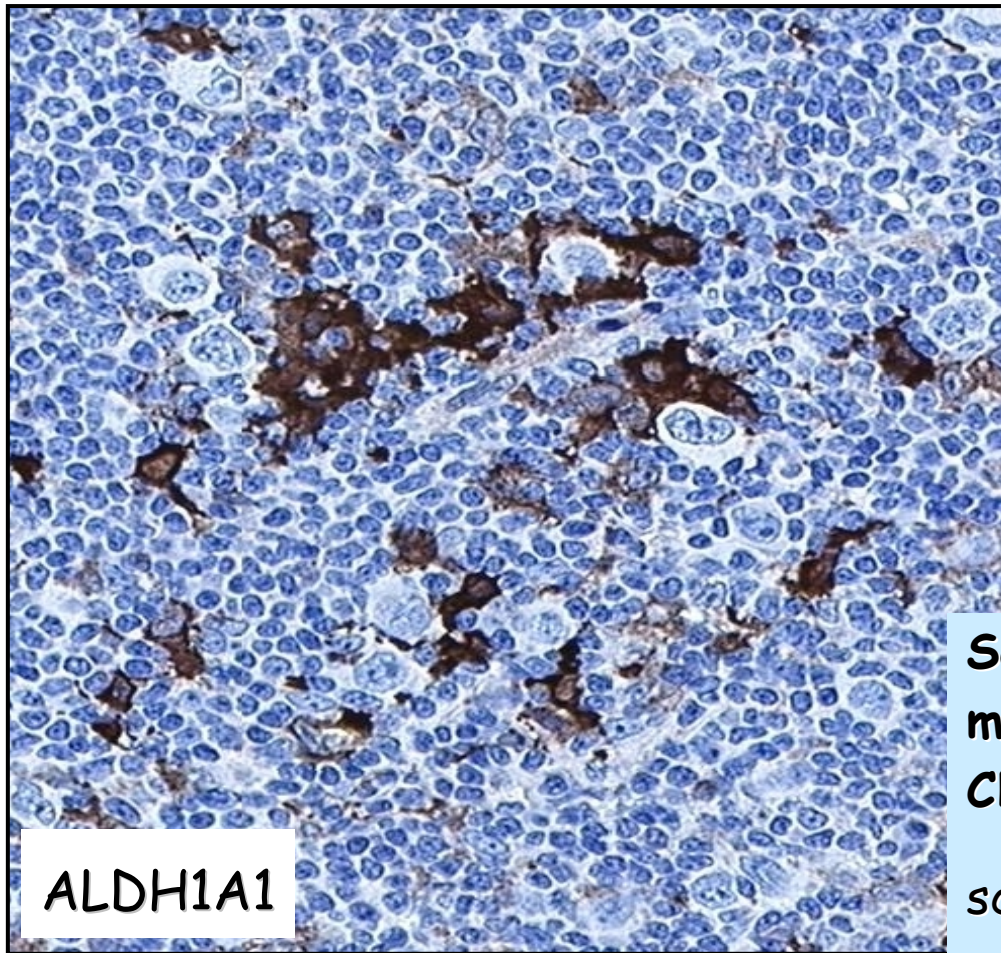
score 2: 10-24%(+)

score 3: 25-49%(+)

score 4: 50-74%(+)

score 5: >75%(+)





Scoring system for macrophage markers (ALDH1A1, CD68/PGM1, CD68/KP1, CD163)

score 0 : 0%(+)

score 1 : 1-4%(+)

score 2: 5-24%(+)

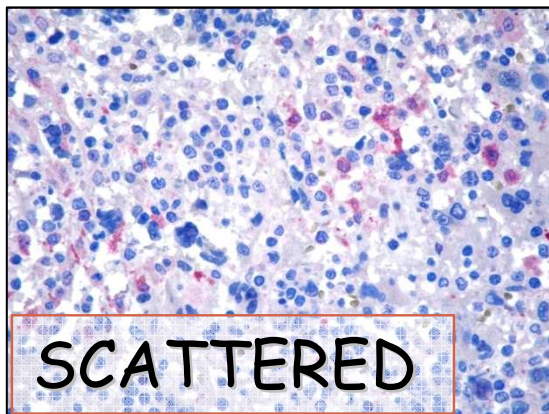
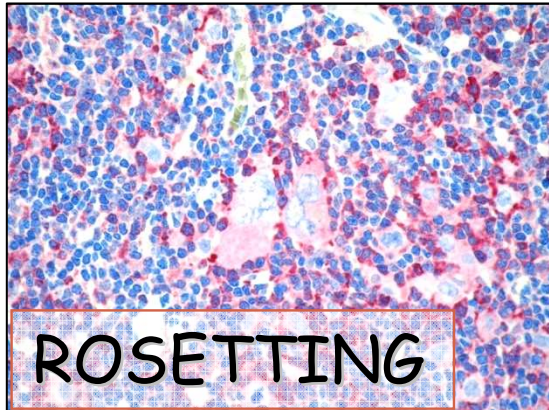
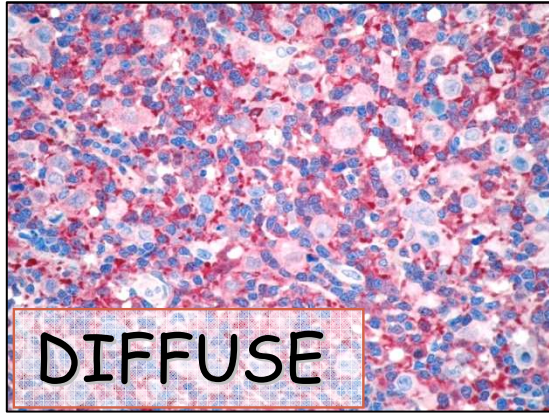
score 3: 25-49%(+)

score 4: 50-74%(+)

score 5: >75%(+)

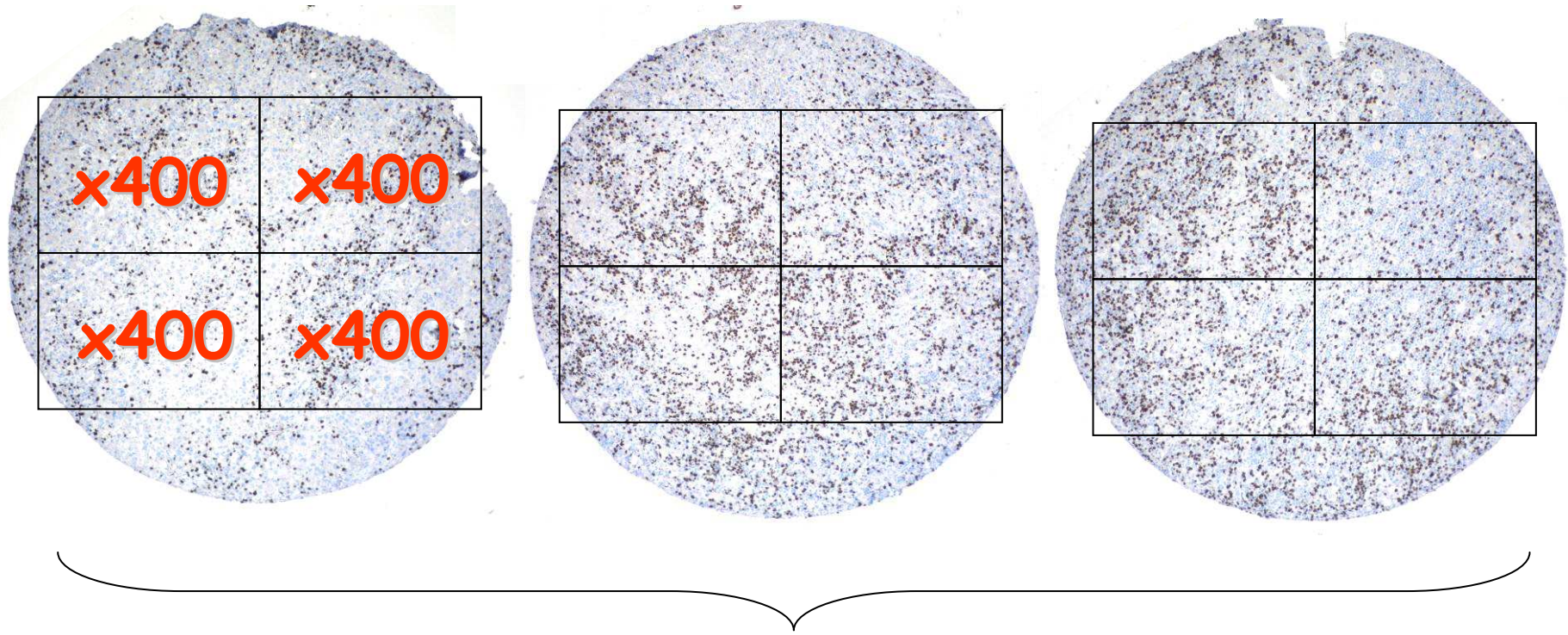
Prognostic indicators in Hodgkin's Lymphoma

STAT1, SAP, PD1 microenvironment expression patterns



- **Diffuse** : diffuse pattern of staining in MC cells between and surrounding neoplastic cells
- **Rosetting** : expressed only in cells forming rosettes around neoplastic cells
- **Scattered** : few cells positive in the microenvironment

FOXP3 and Cytotoxic markers (Tia1, GyB, Perforin) evaluation

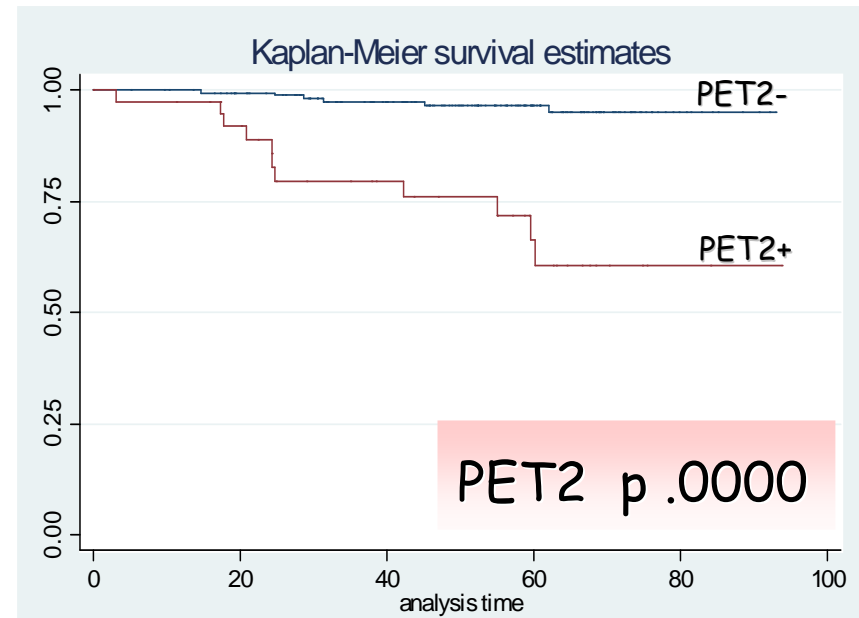
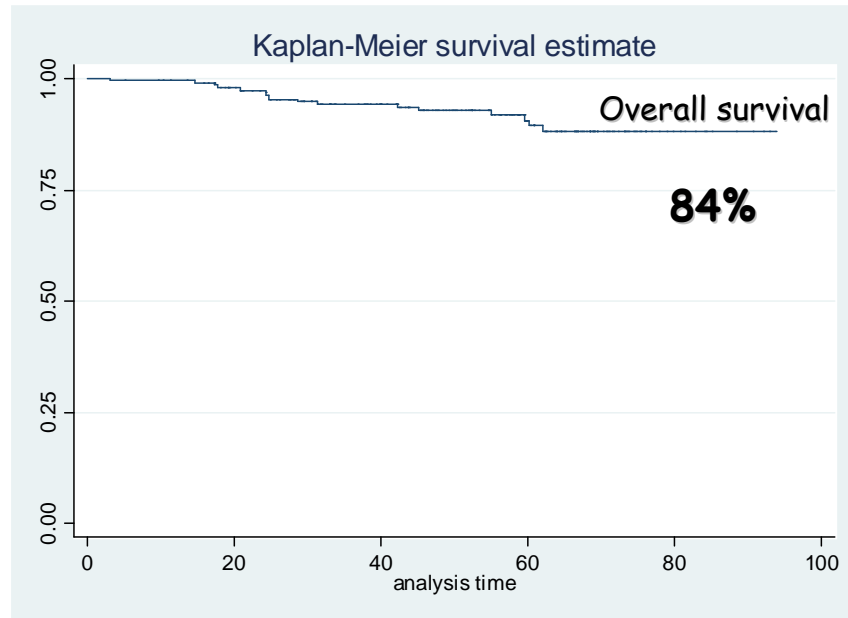


mean value
calculated on evaluable cores

Data processing

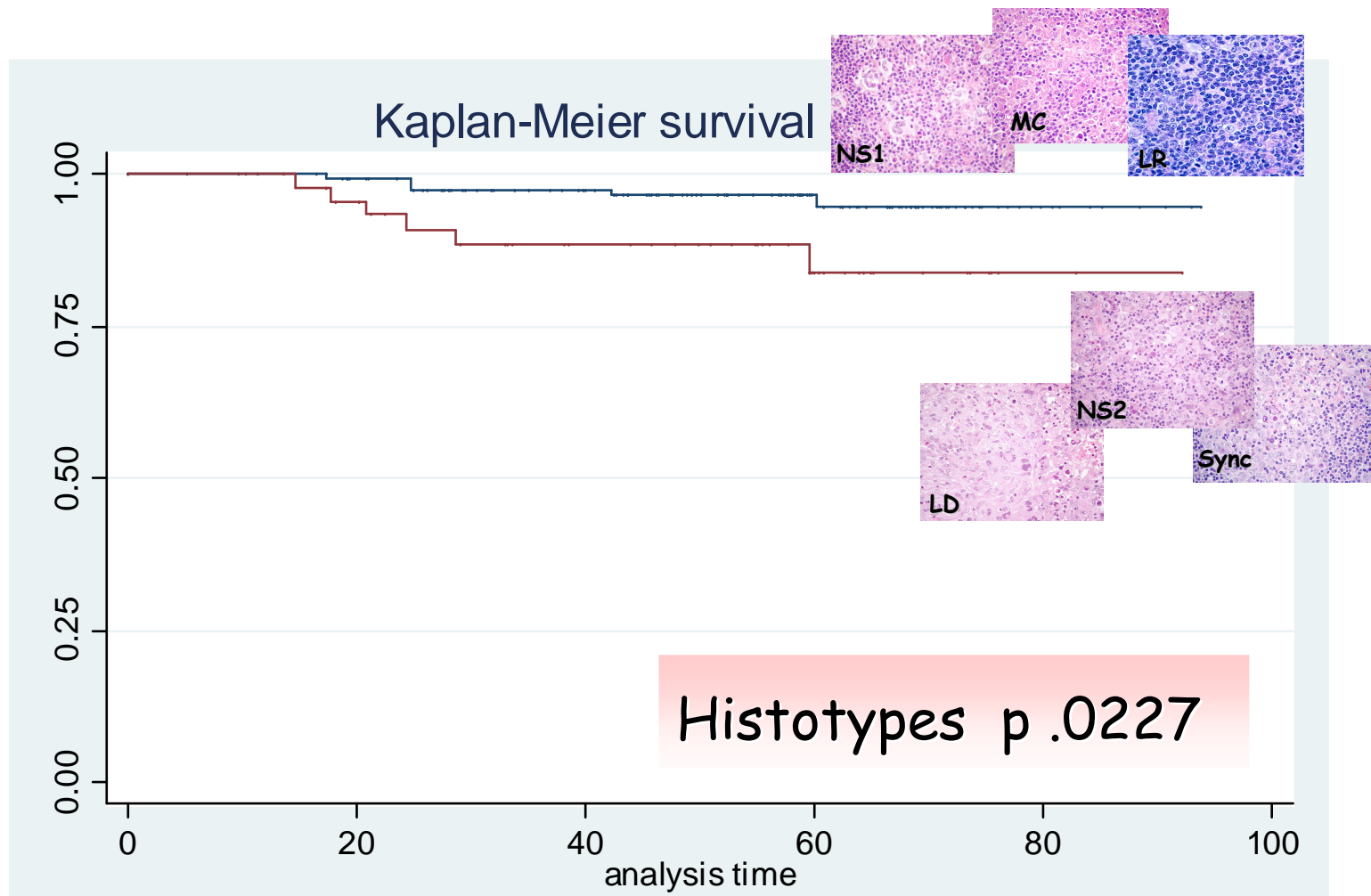
- Every result evaluated to find correlation with patients' outcome: percentage and intensity of expression, nuclear or cytoplasmic localization, both in tumour cells and microenvironment
- Every cut-off assessed
- Every pattern tested

Overall survival

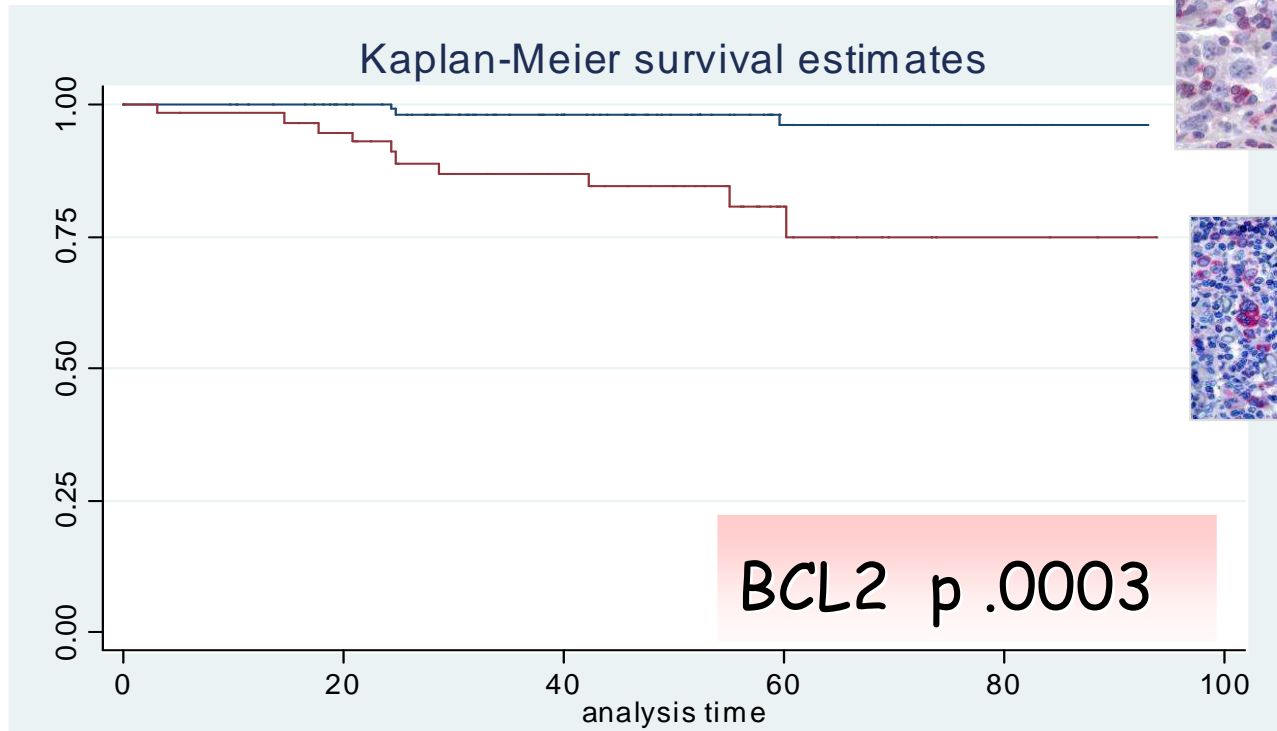


Median follow-up: 62.3 months

Overall survival



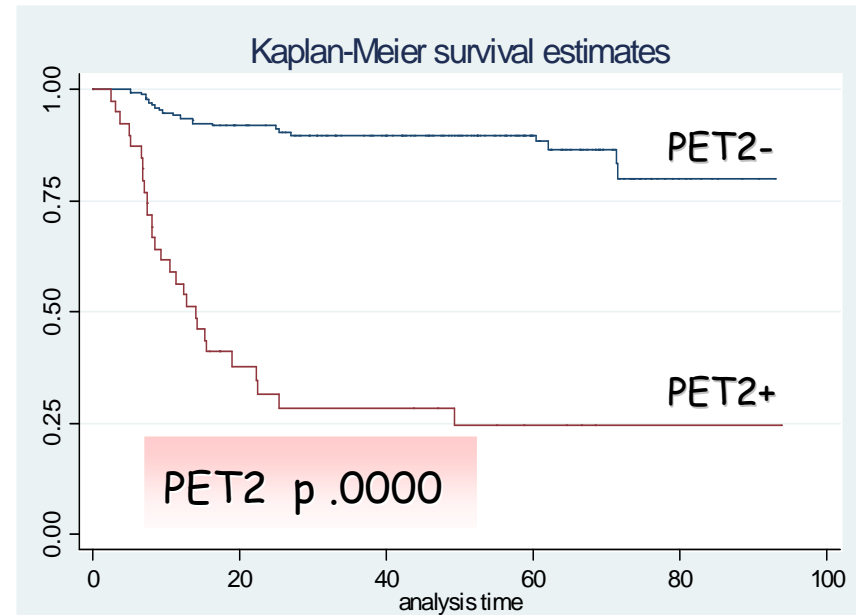
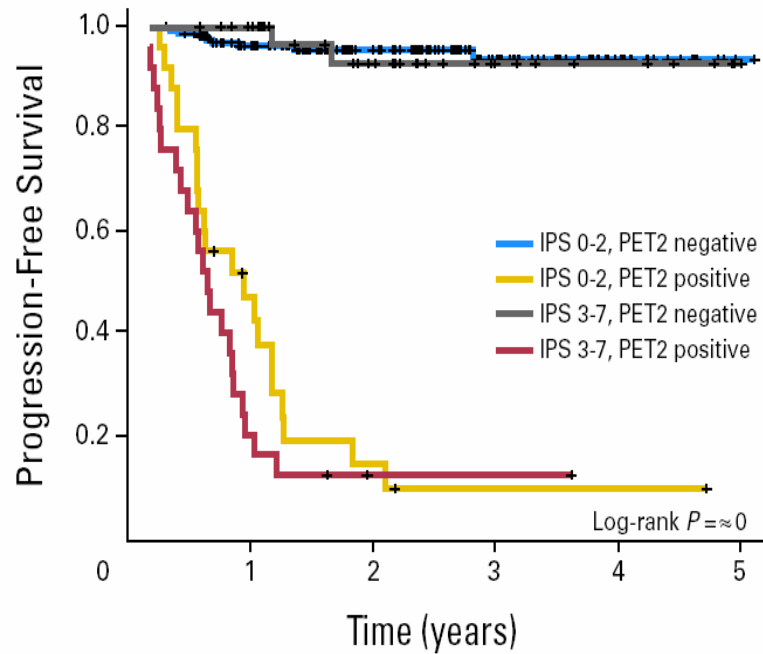
Overall survival



variable	cut-off	n	Hazard ratio of event risk	95% C.I.	P
BCL2	≥ 50%	33,5%	7.63	(2.09-27.80)	.0003

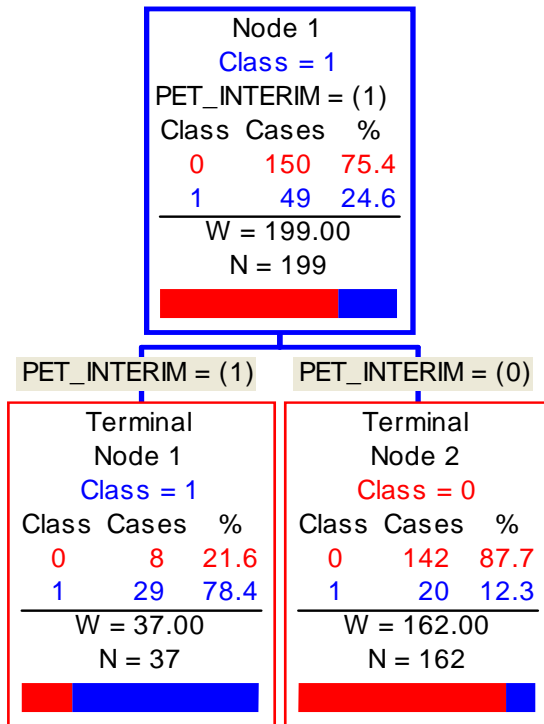
variable	cut-off	OS- 93 months	95% C.I.
BCL2	< 50%	96%	(87.2-98.8)
BCL2	≥ 50%	75%	(55.4-86.9)

Progression Free Survival

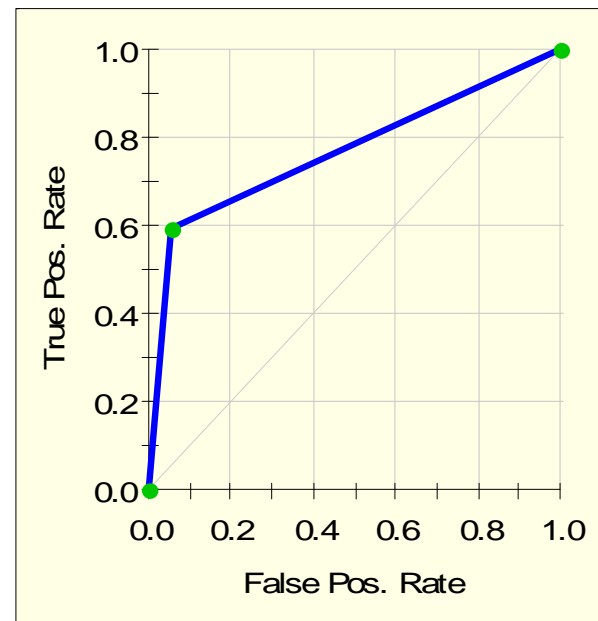


Misclassification for Learn and **Test** Data

Class	N Cases	N Mis-Classed	Pct Error
0	150	8	5.33
1	49	20	40.82
Tot	199	28	14.07

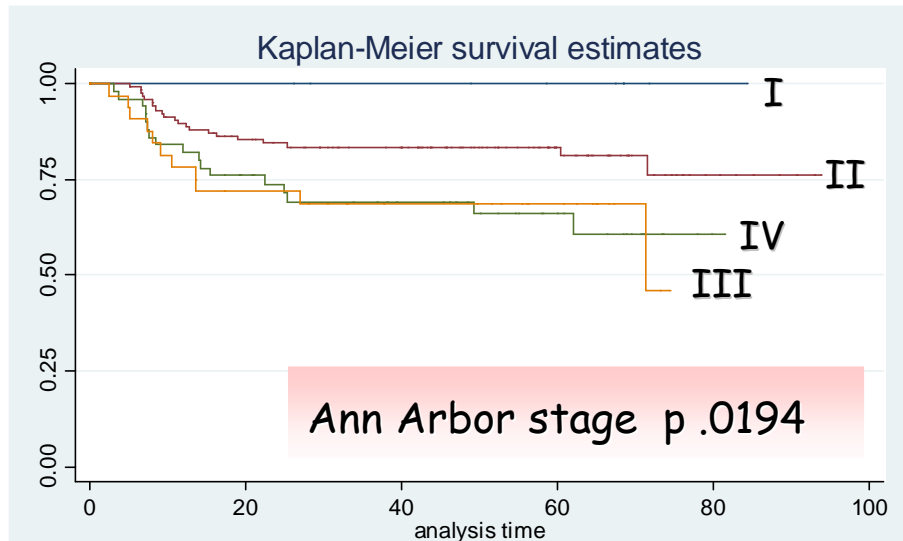


ROC Integral: 0.770

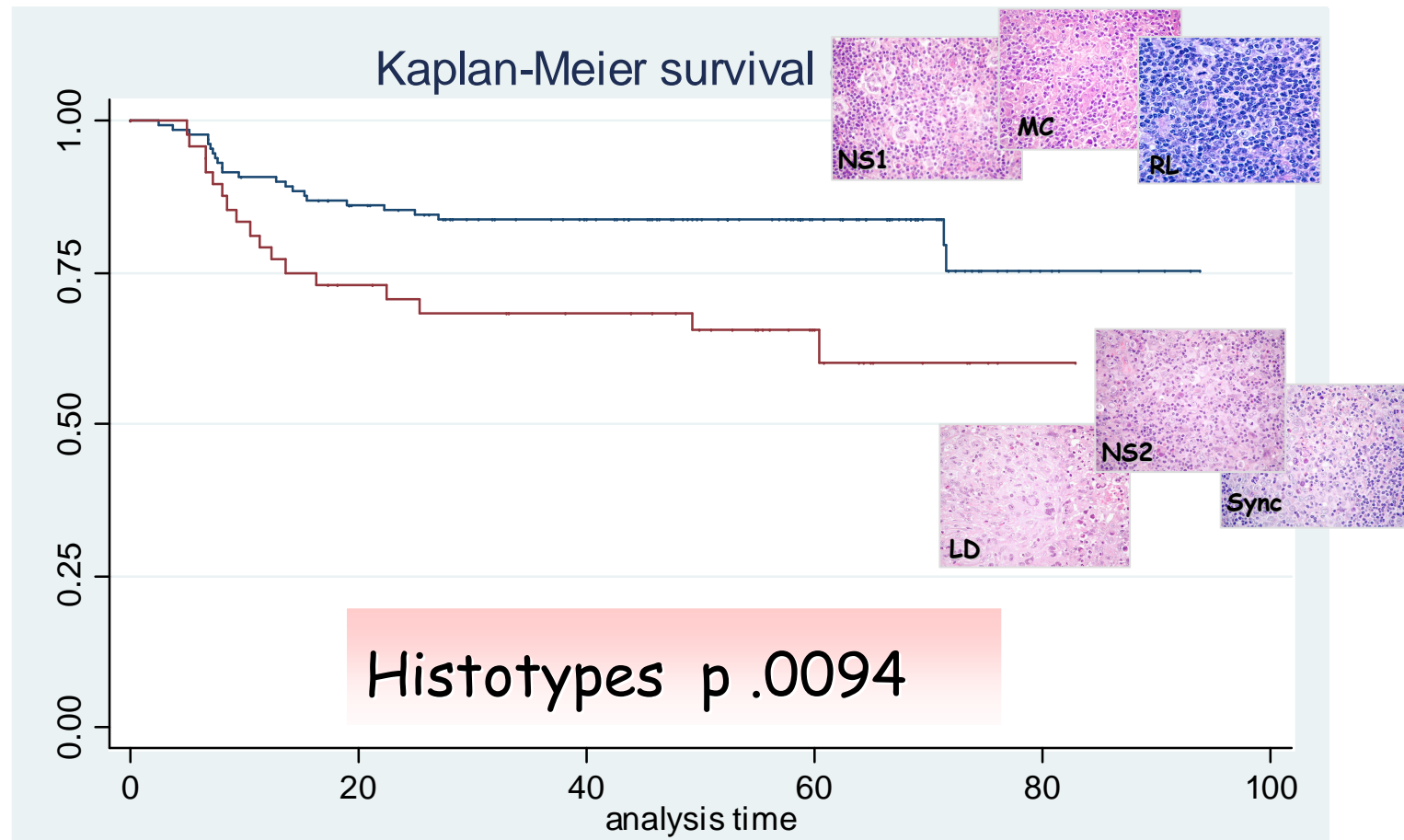


Progression Free Survival

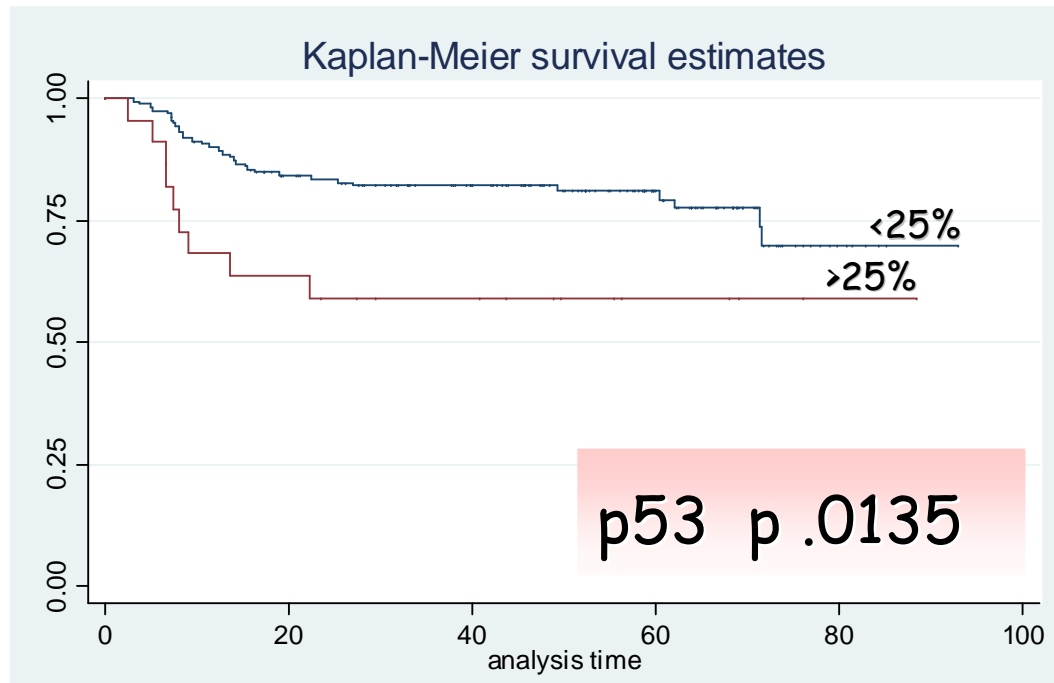
variable	Hazard ratio of event risk	95% C.I.	P
Stage I	0		
Stage II	4.03	(1.94-8.36)	.000
Stage III	8.31	(3.89-17.80)	.000
Stage IV	8.59	(4.93-11.17)	.000



Progression Free Survival

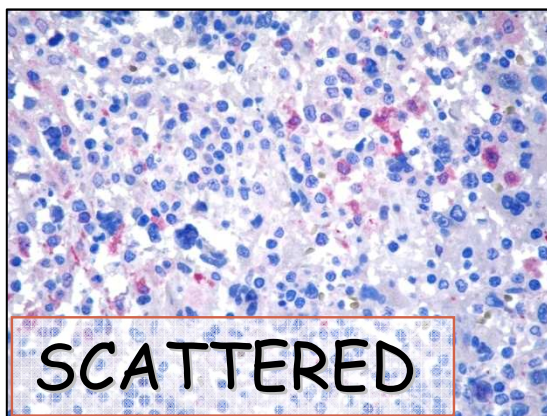
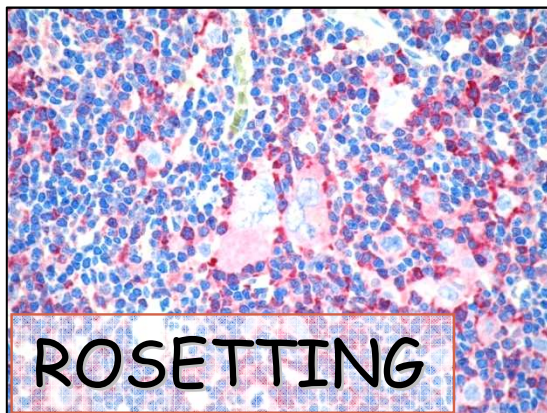
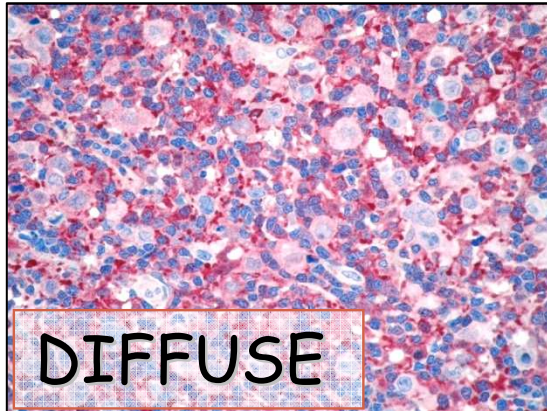


Progression Free Survival



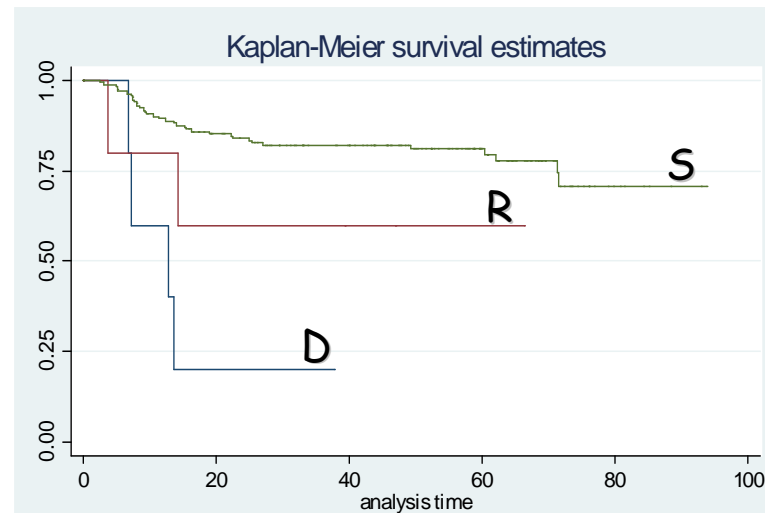
variable	cut-off	PFS- 93 months	95% C.I.
p53	< 25%	69.3%	(55.3-79.6)
p53	≥ 25%	61.1%	(35.3-79.1)

PD1 and PFS

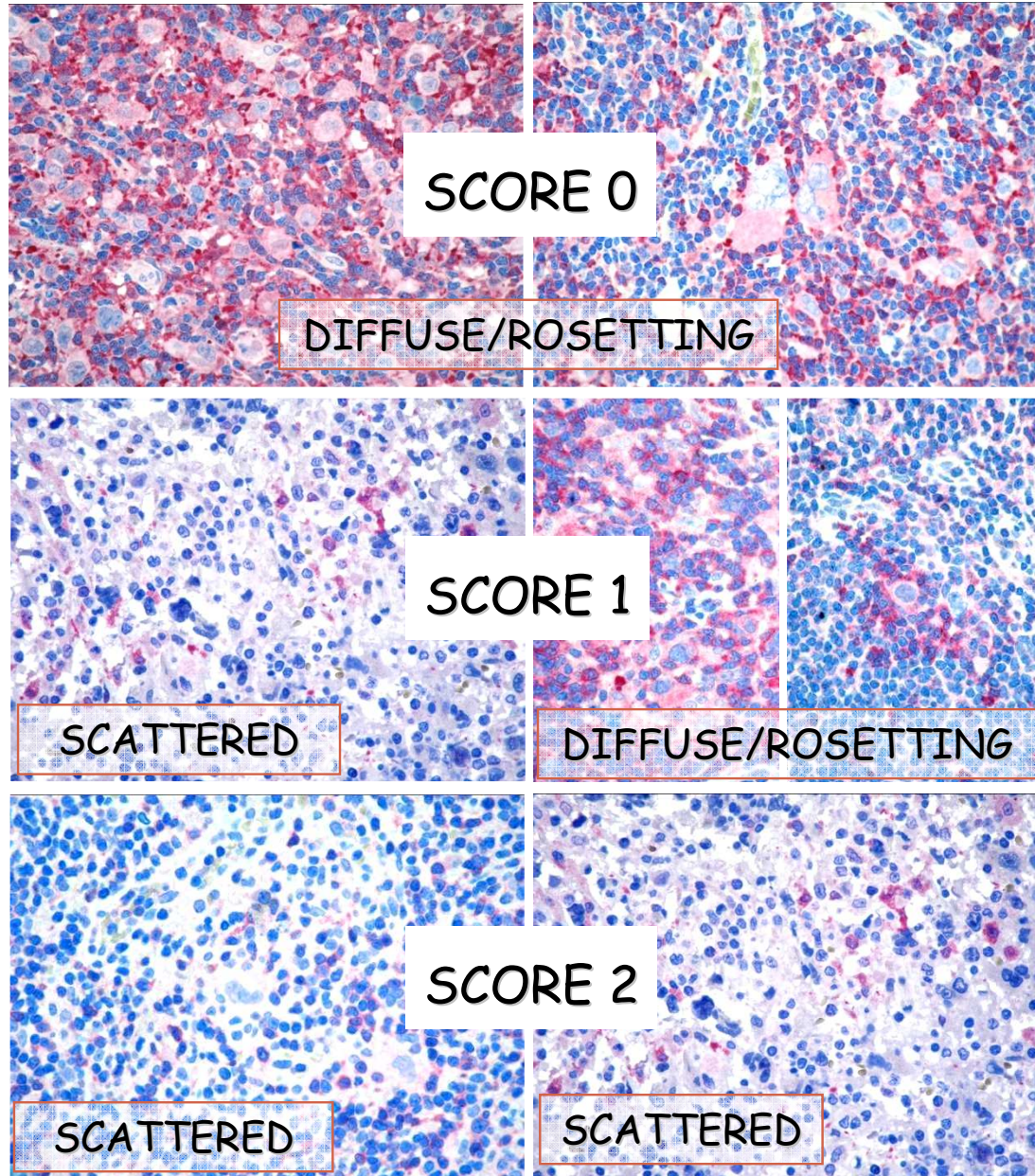


Membranous staining

- PD1 is involved in regulation of TCR-signaling
- Expressed by follicular helper T-cells
- In our series, the expression of PD1 by lymphocytes of microenvironment is related to adverse outcome (p.0000)

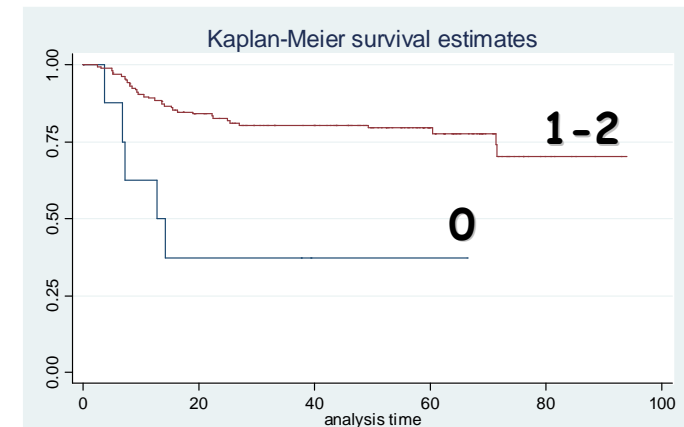


PD1/SAP uni-variate analysis PFS



Combined expression of FTH markers in microenvironment is associated with worse prognosis (p .0018):

Score 0 = worse prognosis



Prognostic biologic factors in Hodgkin's lymphoma

Multivariate analysis

Cox's regression model:

Overall Survival

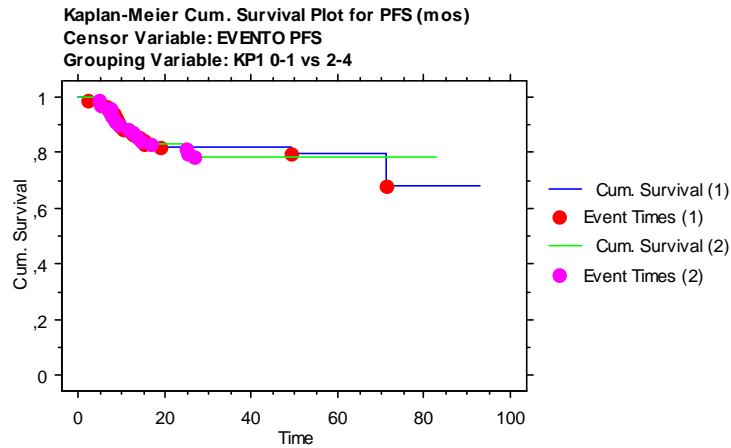
variable	Hazard ratio of event risk	95% C.I.	P
BCL2	1.51	(1.06-2.15)	.021
PET2	11.5	(3.0-43.5)	.000

Progression Free Survival

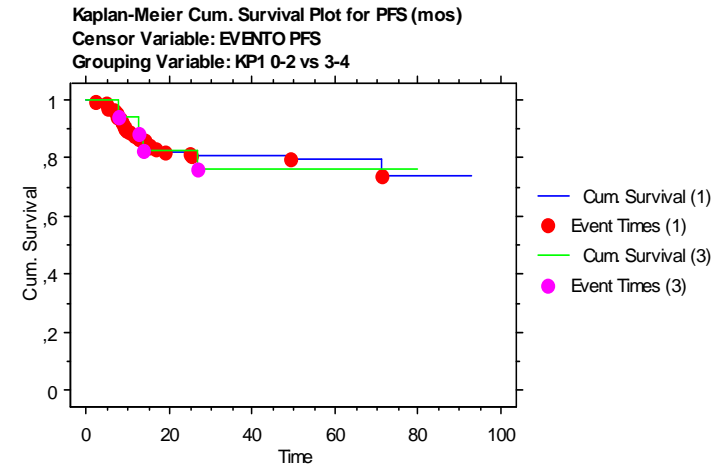
variable	Hazard ratio of event risk	95% C.I.	P
Stage	2.16	(1.46-3.09)	.000
P53	3.64	(1.55-8.51)	.003
PET2	14.97	(7.53-29.78)	.000

Prognostic biologic factors in Hodgkin's lymphoma

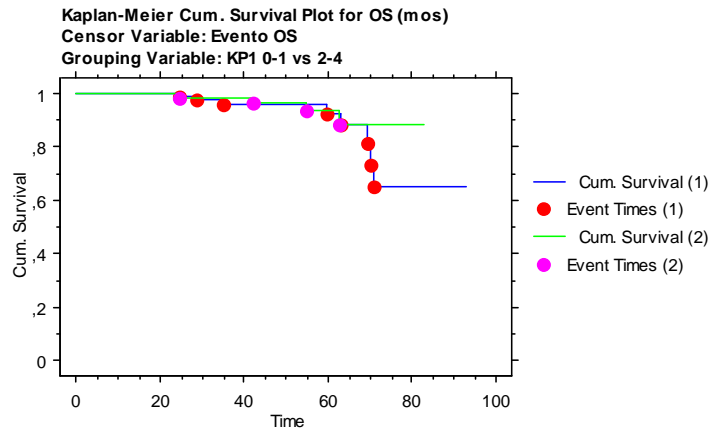
CD68/KP1



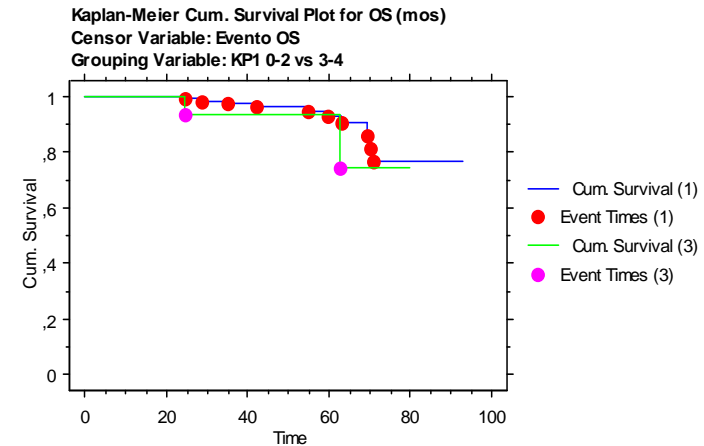
PFS, Cut off 5%, $p=0.95$



PFS, Cut off 25%, $p=0.81$



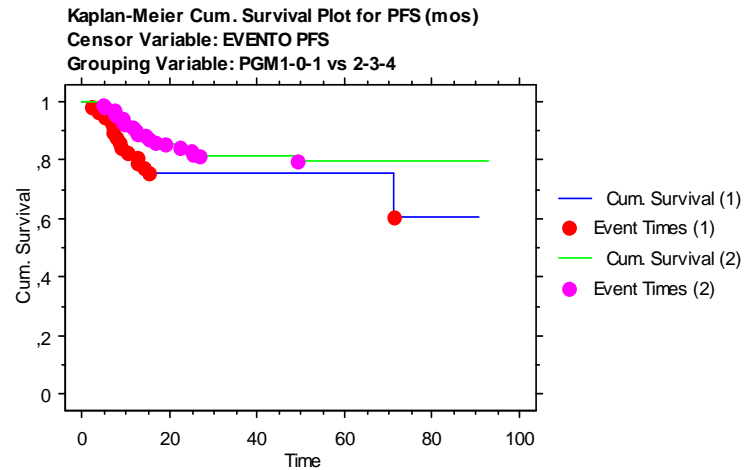
OS, Cut off 5%, $p=0.36$



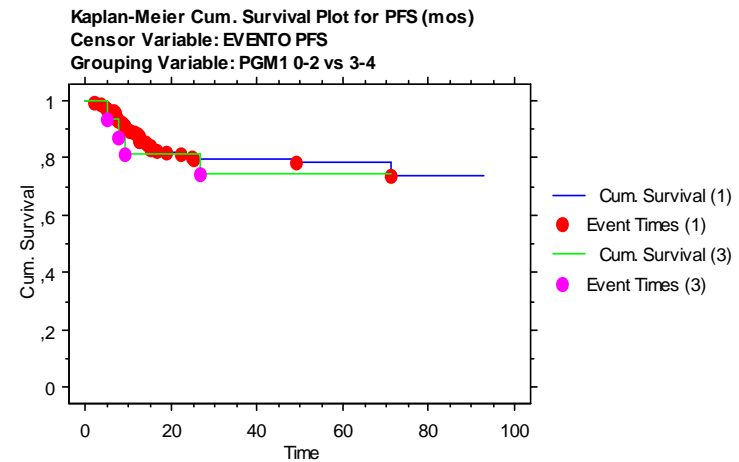
OS, Cut off 25%, $p=0.5$

Prognostic biologic factors in Hodgkin's lymphoma

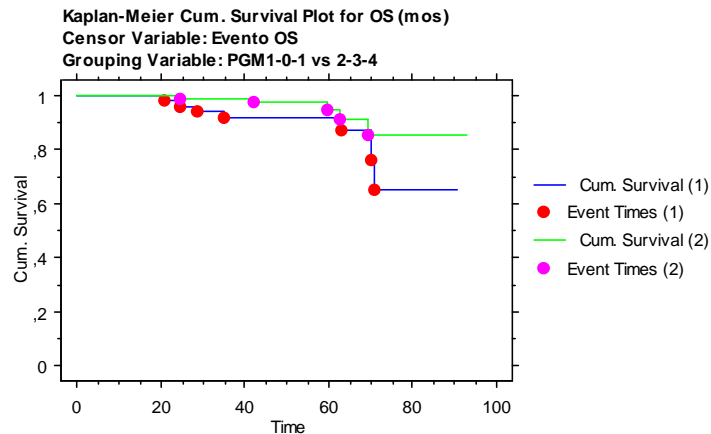
CD68/PGM1



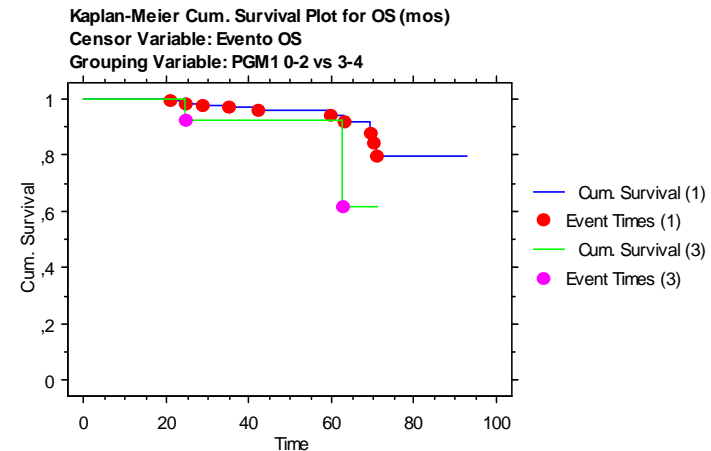
PFS, Cut off 5%, p=0.26



PFS, Cut off 25%, p=0.67



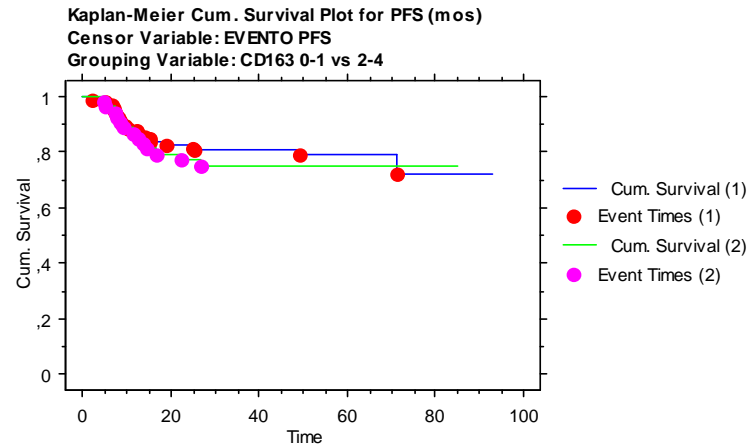
OS, Cut off 5%, p=0.1



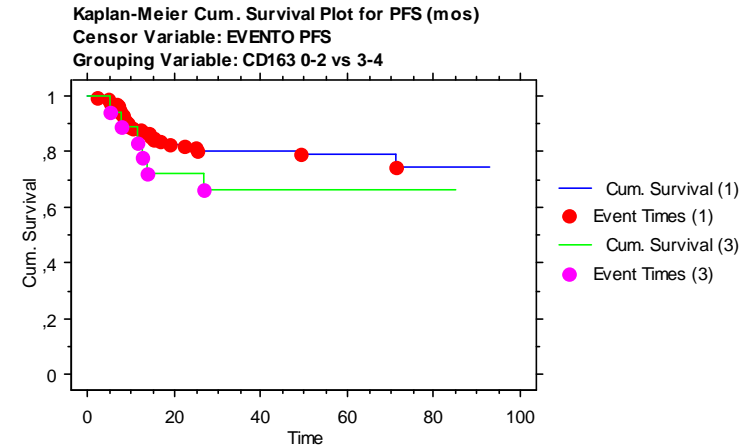
OS, Cut off 25%, p=0.2

Prognostic biologic factors in Hodgkin's lymphoma

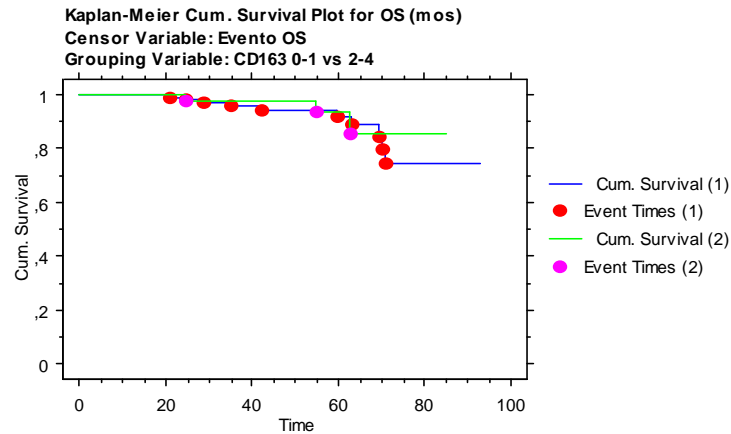
CD163



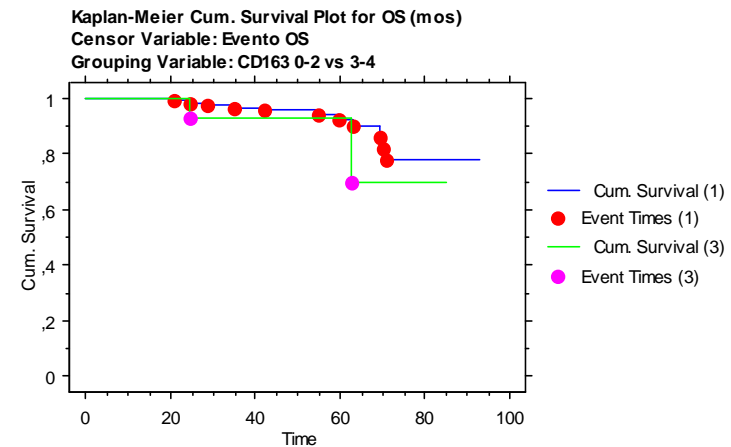
PFS, Cut off 5%, $p=0.64$



PFS, Cut off 25%, $p=0.2$



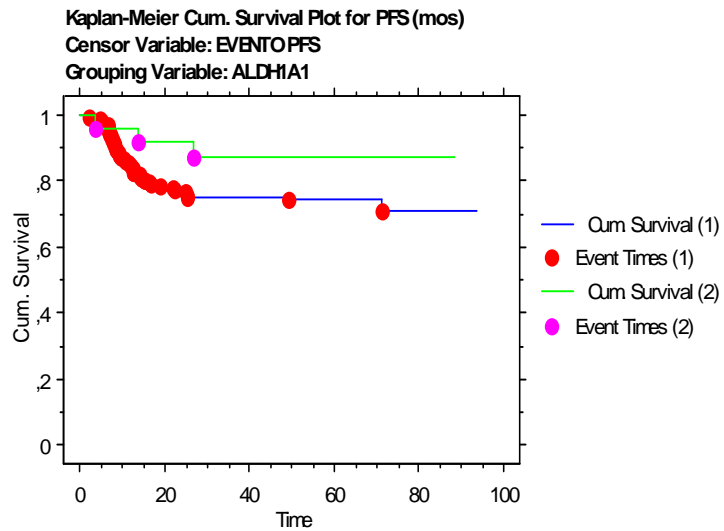
OS, Cut off 5%, $p=0.36$



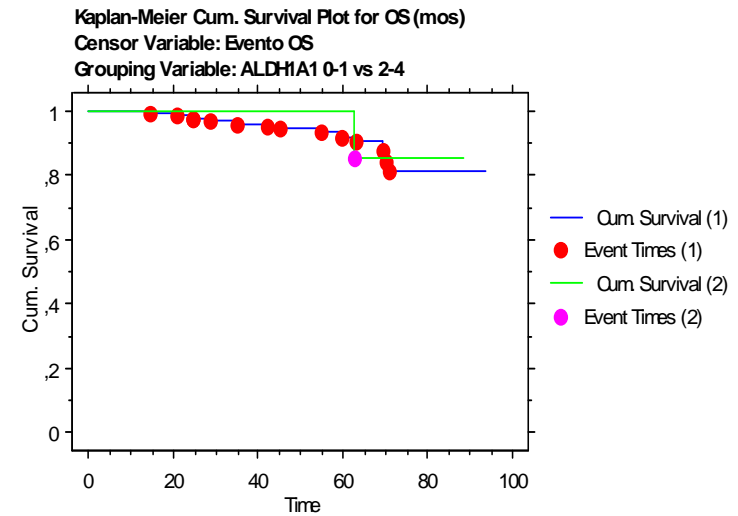
OS, Cut off 25%, $p=0.34$

Prognostic biologic factors in Hodgkin's lymphoma

ALDH1A1



PFS, Cut off 5%, $p=0.1$



OS, Cut off 5%, $p=0.5$

Prognostic biologic factors in Hodgkin's lymphoma

Lack of association of tumor-associated macrophages with clinical outcome in patients with classical Hodgkin's lymphoma

D. Azambuja¹, Y. Natkunam², I. Biasoli³, I. S. Lossos⁴, M. W. Anderson², J. C. Morais³ and N. Spector^{3,*}

Abstract

Background: A recent study demonstrated that an increased number of CD68+ macrophages were correlated with primary treatment failure, shortened progression-free survival (PFS) and disease-specific survival (DSS) in patients with classical Hodgkin's lymphoma (cHL).

Patients and methods: The aim of the present study was to verify the relationship between the number of CD68+ and CD163+ macrophages with clinical outcomes in a cohort of 265 well-characterized patients with cHL treated uniformly with the standard doxorubicin, bleomycin, vinblastine and dacarbazine chemotherapy regimen. Two pairs of hematopathologists carried out independent pathological evaluations of tissue microarray slides.

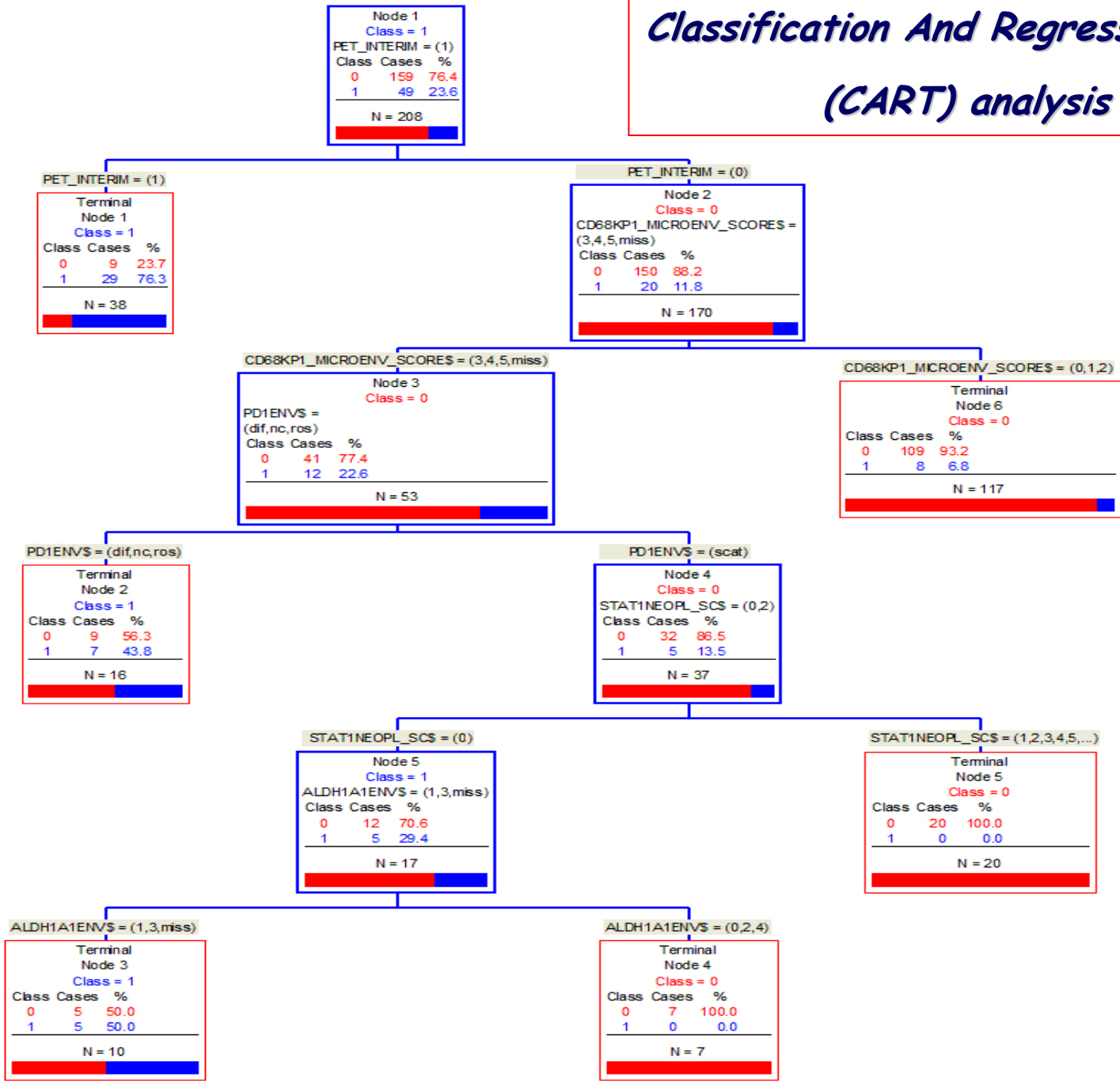
Results: There were no associations between clinical characteristics and the expression of CD68 or CD163. However, higher levels of CD68 and CD163 expression were correlated with the presence of Epstein-Barr virus-positive Hodgkin tumor cells ($P = 0.01$ and 0.037 , respectively). The expression of CD68 or CD163 was not associated with either the PFS or the DSS.

Conclusion: CD68 and CD163 expression require further evaluation before their use can be recommended for prognostic stratification of patients with cHL.

Comments on macrophages

- Steidl et al. (NEJM) found that the macrophage content correlates with DFS in a cohort of patients with a median follow-up of 16.4 years.
- In two other papers, Kamper et al (Haematologica) and Tzankov et al (Pathobiology) also observed that the amount of macrophages correlates with OS by using however different counting systems; the two series spanned over 17 and 26 years, respectively.
- It is likely that the relatively short follow-up (median 62.3 months) due to the selection criteria used (i.e. cases with PET-2 available), has limited the statistical power of the analysis.

Classification And Regression Tree (CART) analysis



Macrophages predict treatment outcome in Hodgkin's lymphoma

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Haematologica 2010, 96:186-9

Antibody and scoring system

Further study is needed to determine the optimal antigen (e.g. CD68 versus CD163), anti-CD68 antibody clone (e.g. KP1 versus PGM1) and scoring thresholds (e.g. manual versus computer-assisted) for detecting HL-associated macrophages.

Biomarker combination

The preliminary results of an immunohistochemistry study combining two markers, CD68 and FOXP3 (a marker for regulatory T cells), were presented at the ASH 2010 meeting. The authors showed that a combined FOXP3/CD68 immunohistochemistry score was an improvement over the predictive value of the individual markers alone and that this score was applicable to both limited and advanced-stage disease. The value of this com-

EBV infection

In a re-analysis of our data we were able to confirm a relationship between increased tumor-associated macrophages and EBV positivity; however, virtually all of our cases were of the nodular sclerosis subtype and EBV alone was not associated with treatment outcome (*unpublished observations, 2010*). EBV infection of HRS cells has been reported in up to 60% of patients and is more frequent in mixed cellularity subtype, although varying with geographical location, age, gender, clinical stage and histological subtype.¹⁴ The impact of EBV infection on outcome remains controversial, but appears to be dependent on age.

Conclusions

- **PET2 still maintains the highest predictive value** but remains an *ad interim* parameter that doesn't avoid the risk of induced chemo-resistance produced by two cycles of ABVD.
- Several promising up-front prognostic markers are proposed by the present study, including p53 and Bcl2 that have a bit been neglected during the last few years.
- The impact of microenvironment including macrophages, is certainly relevant; however, some further work (e.g. standardization of cut-off values and markers) seems needed.
- **CART analysis allowed the retrieval of most patients misclassified by interim PET as negative and may therefore represent an interesting operational tool .**



C Agostinelli, PP Piccaluga, E Sabattini, F Bacci, C Sagramoso, M Rossi, S Righi, A Gazzola, T Sista, M Piccioli, MR Sapienza, C Mannu, F Sandri, P Artioli, G De Biase, G Da Pozzo, C Tigrini and I Barese

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