

blood

MYC/BCL2 protein co-expression contributes to the inferior survival of activated B-cell subtype of diffuse large B-cell lymphoma and demonstrates high-risk gene expression signatures: a report from The International DLBCL Rituximab-CHOP Consortium Program Study.

Hu S, Xu-Monette ZY, Tzankov A, Green T, Wu L, Balasubramanyam A, Liu WM, Visco C, Li Y, Miranda RN, Montes-Moreno S, Dybkær K, Chiu A, Orazi A, Zu Y, Bhagat G, Richards KL, Hsi ED, Choi WW, Zhao X, van Krieken JH, Huang Q, Huh J, Ai W, Ponzoni M, Ferreri AJ, Zhou F, Slack GW, Gascoyne RD, Tu M, Variakojis D, Chen W, Go RS, Piris MA, Møller MB, Medeiros LJ, Young KH.

Blood. 2013 May 16;121(20):4021-31

**ΒΙΒΛΙΟΓΡΑΦΙΚΗ ΕΝΗΜΕΡΩΣΗ ΓΝΑ ΕΥΑΓΓΕΛΙΣΜΟΣ
Θ. ΚΑΝΕΛΛΟΠΟΥΛΟΥ**



ΕΙΣΑΓΩΓΗ

Ταξινόμηση WHO 2008 - DLBCL

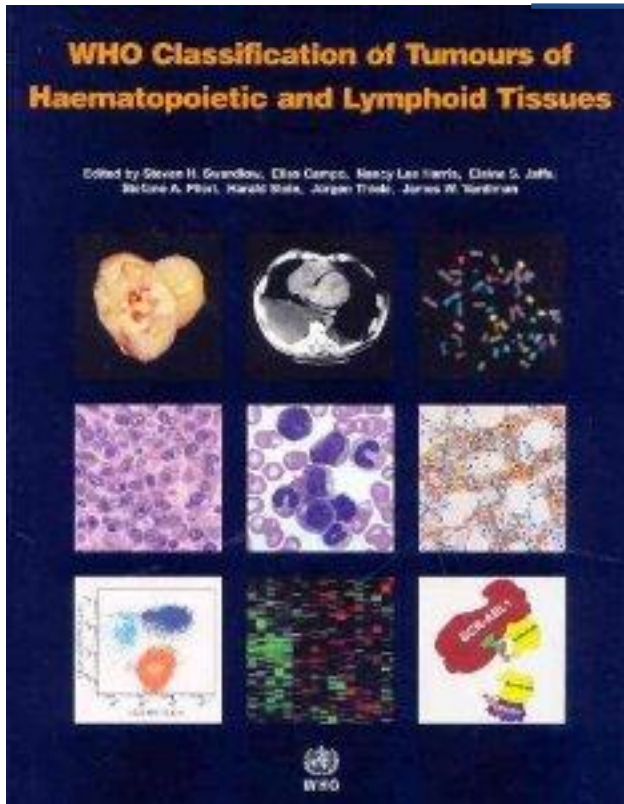


Table 10.14 Diffuse large B-cell lymphoma: variants, subgroups and subtypes/entities.

Diffuse large B-cell lymphoma, not otherwise specified (NOS)

Common morphologic variants

- Centroblastic
- Immunoblastic
- Anaplastic

Rare morphologic variants

Molecular subgroups

- Germinal centre B-cell-like (GCB)
- Activated B-cell-like (ABC)

Immunohistochemical subgroups

- CD5-positive DLBCL
- Germinal centre B-cell-like (GCB)
- Non-germinal centre B-cell-like (non-GCB)

Diffuse large B-cell lymphoma subtypes

- T-cell/histiocyte-rich large B-cell lymphoma
- Primary DLBCL of the CNS
- Primary cutaneous DLBCL, leg type
- EBV positive DLBCL of the elderly

Other lymphomas of large B cells

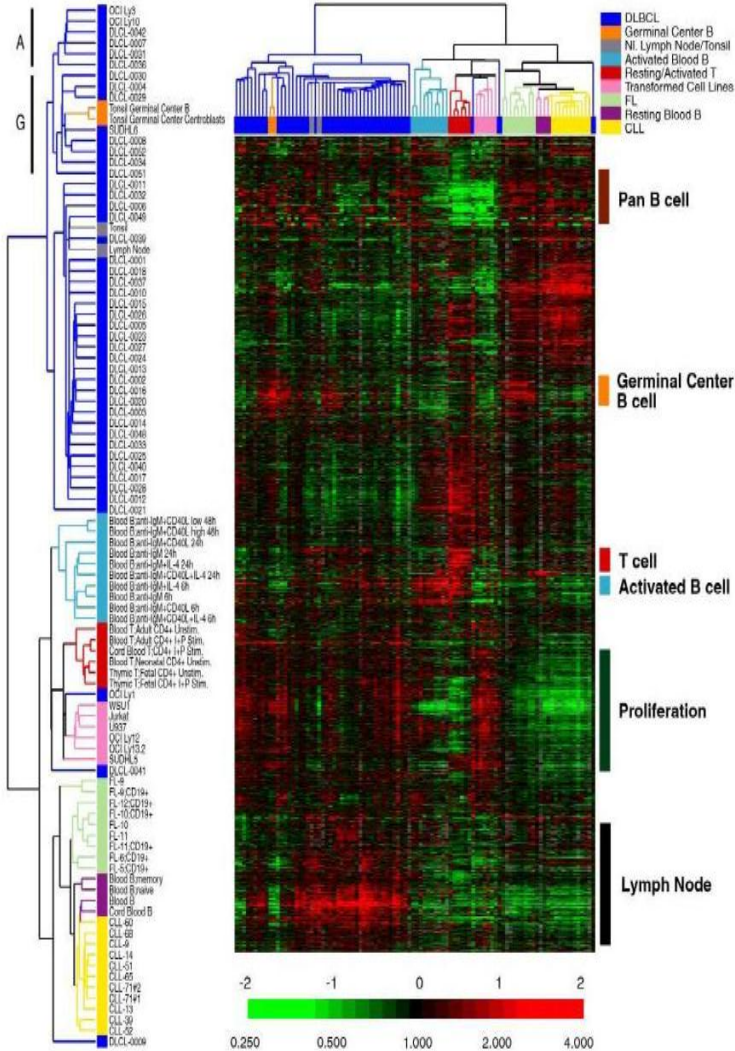
- Primary mediastinal (thymic) large B-cell lymphoma
- Intravascular large B-cell lymphoma
- DLBCL associated with chronic inflammation
- Lymphomatoid granulomatosis
- ALK-positive LBCL
- Plasmablastic lymphoma
- Large B-cell lymphoma arising in HHV8-associated multicentric Castlemann disease
- Primary effusion lymphoma

Borderline cases

- B-cell lymphoma, unclassifiable, with features intermediate between diffuse large B-cell lymphoma and Burkitt lymphoma
- B-cell lymphoma, unclassifiable, with features intermediate between diffuse large B-cell lymphoma and classical Hodgkin lymphoma

Gene-expression profiles

DLBCL – NOS



- Τύπος βλαστικού κέντρου (GCB)

- *CD10, BCL6*

- Τύπος ενεργοποιημένου β-λεμφοκυττάρου (ABC)

- *MYC, BCL2, MUM1, CD44, FLIP, cyclinD2*

- Ενεργοποίηση NF-κB

- Χειρότερη πρόγνωση

Gene-expression profiles

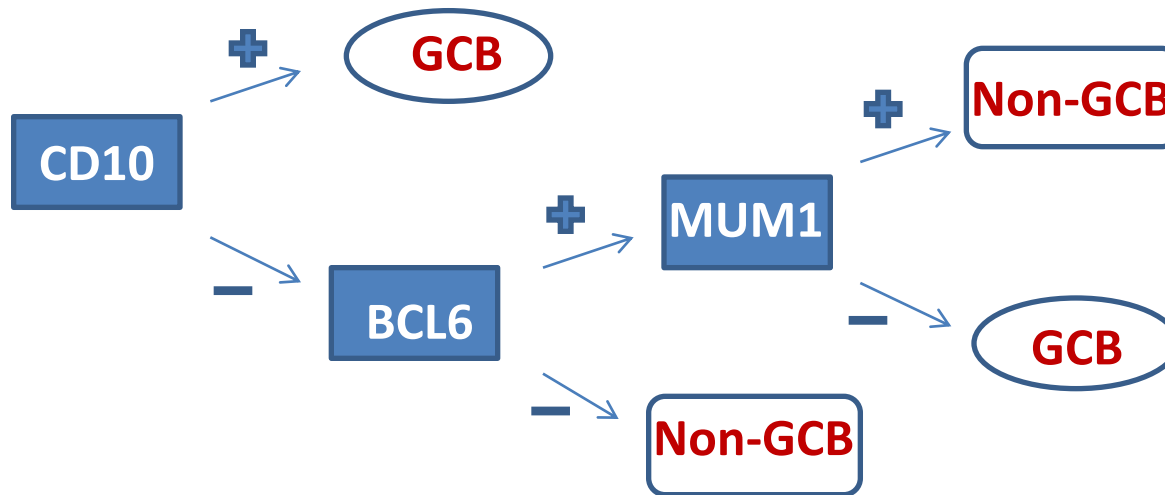
- **Περιορισμοί ...**

- Η τεχνολογία GEP δεν είναι διαθέσιμη στην καθημερινή κλινική πράξη
- Αναπαραγωγιμότητα;;;
- Παραμένει άγνωστο ποια γονιδιακά προϊόντα συμβάλλουν στη χειρότερη πρόγνωση των ασθενών με ABC-DLBCL

Εναλλακτικές λύσεις ;;;

Ανοσοϊστοχημεία

- Αλγόριθμος του Hans



Hans CP, et al. Confirmation of the molecular classification of diffuse large B-cell lymphoma by immunohistochemistry using a tissue microarray. Blood. 2004 Jan 1;103(1):275-82

- Συμφωνία με GEP 80%

“Double-Hit”

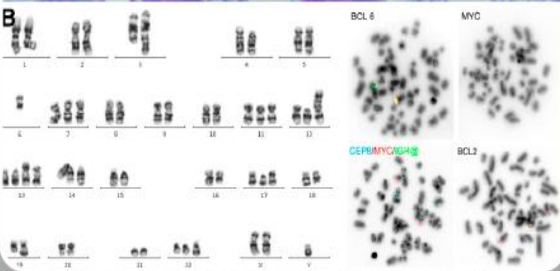
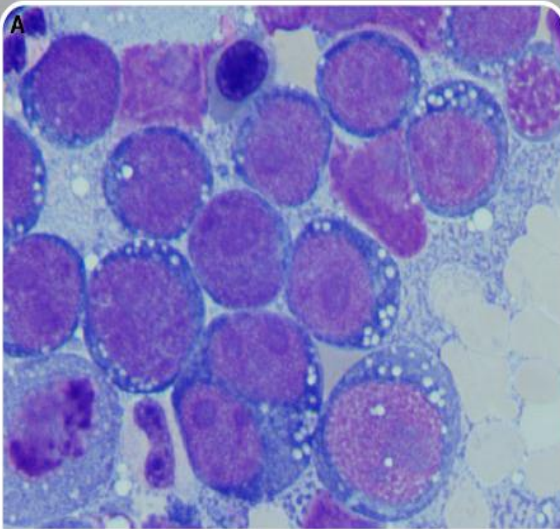
- Λεμφώματα με επαναλαμβανόμενες μεταλλαγές που ενεργοποιούν πολλαπλά ογκογονίδια

- Επιθετική κλινική πορεία και αντοχή στη θεραπεία

- ***BCL2+/MYC+*** , ***BCL6+/MYC+***, ***BCL3+/MYC+***

- Συνήθως **GCB φαινότυπος**

- **CD10+**, **BCL6+**, **MUM1/IRF4-**



• Rudolf Benz, Kantonsspital Münsterlingen; and Joelle Tchinda, Children's Hospital Zürich

Νεότερα βιβλιογραφικά δεδομένα

- Ασθενείς με DLBCL με συνέκφραση BCL2/MYC με ή χωρίς αναδιατάξεις MYC ή BCL2 έχουν χειρότερη πρόγνωση

- *Johnson NA, et al. Concurrent expression of MYC and BCL2 in diffuse large B-cell lymphoma treated with rituximab plus cyclophosphamide, doxorubicin, vincristine, and prednisone. J Clin Oncol. 2012 Oct 1;30(28):3452-9*
- *Green TM, et al. Immunohistochemical double-hit score is a strong predictor of outcome in patients with diffuse large B-cell lymphoma treated with rituximab plus cyclophosphamide, doxorubicin, vincristine, and prednisone. J Clin Oncol. 2012 Oct 1;30(28):3460-7*

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Blood. 2013 Feb 28. [Epub ahead of print]

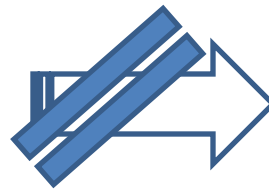


ΥΛΙΚΑ & ΜΕΘΟΔΟΙ

Ασθενείς

International DLBCL R-CHOP Consortium Program Study

893 ασθενείς
De novo DLBCL
MYC+/BCL2+
(ανοσοϊστοχημικά)
R-CHOP



Χαμηλής κακοήθειας B-NHL
HIV
DLBCL δέρματος/ΚΝΣ
EBV (+) DLBCL

466 training set

234 validation set #1

193 validation set #2

GEP σε 451 ασθενείς

9% αταξινόμητα

ΑΠΟΤΕΛΕΣΜΑΤΑ

Χαρακτηριστικά ασθενών

	N (%)	Overall		DP	Non-DP	p-value
		OS (p-value)	PFS (p-value)	N (%)	N (%)	
Patients	466 (100%)			157 (100%)	309 (100%)	
Gender						
Male	272 (58%)	0.7477	0.4730	90 (57%)	182 (59%)	0.7445
Female	194 (42%)			67 (43%)	127 (41%)	
Age						
<60	194 (42%)	0.0004	0.0016	49 (31%)	145 (47%)	0.0011
>60	272 (58%)			108 (69%)	164 (53%)	
B symptoms*						
Absence	276 (68%)	0.0015	0.0014	88 (62%)	188 (72%)	0.0541
Presence	127 (32%)			53 (38%)	74 (28%)	
ECOG performance status*						
<2	350 (88%)	<0.0001	<0.0001	111 (83%)	239 (90%)	0.0453
≥2	50 (12%)			23 (17%)	27 (10%)	
Stage*						
I-II	219 (49%)	<0.0001	<0.0001	50 (33%)	169 (57%)	<0.0001
III-IV	228 (51%)			100 (67%)	128 (43%)	
Extranodal Sites*						
<2	346 (78%)	<0.0001	<0.0001	106 (72%)	240 (82%)	0.0160
≥2	96 (22%)			42 (28%)	54 (18%)	
LDH*						
Normal	168 (40%)	0.0003	<0.0001	51 (36%)	117 (42%)	0.2908
Elevated	252 (60%)			89 (64%)	163 (58%)	
IPI risk group*						
0-2	263 (64%)	<0.0001	<0.0001	70 (51%)	193 (70%)	0.0001
3-5	148 (36%)			67 (49%)	81 (30%)	
Tumor size (cm)*						
<7.5	253 (77%)	0.0100	0.0172	81 (73%)	172 (79%)	0.2587
≥7.5	77 (23%)			30 (27%)	47 (21%)	
Treatment response						
CR	354 (76%)	<0.0001	<0.0001	103 (66%)	251 (84%)	<0.0001
Others	112 (24%)			54 (34%)	48 (16%)	
COO Classification						
GCB	241 (52%)	0.0080	0.0075	53 (34%)	188 (61%)	<0.0001
ABC	225 (48%)			104 (66%)	121 (39%)	
KI-67*						
<70%	158 (34%)	0.2998	0.3434	41 (26%)	117 (38%)	0.0086
≥70%	304 (66%)			116 (74%)	188 (62%)	
TP53 mutations						
Absence	357 (77%)	0.0005	0.0004	117 (75%)	240 (78%)	0.4480
Presence	109 (23%)			40 (25%)	69 (22%)	



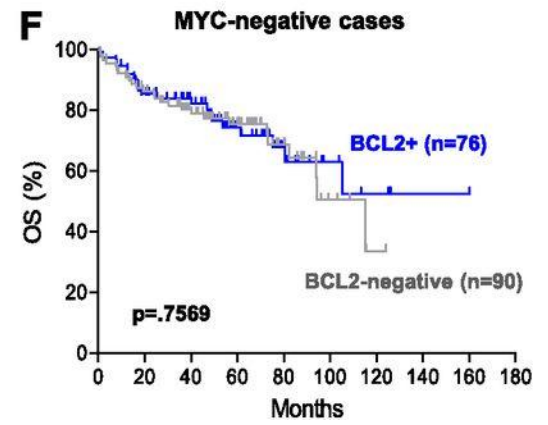
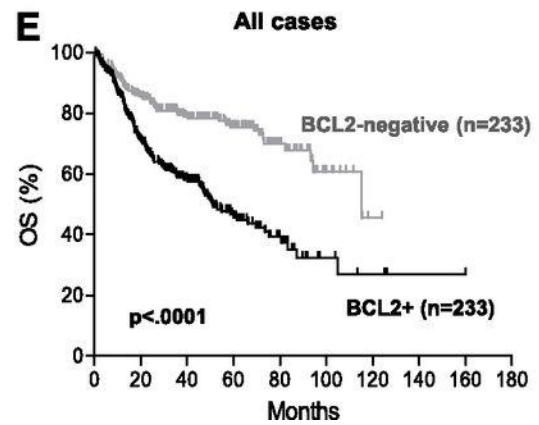
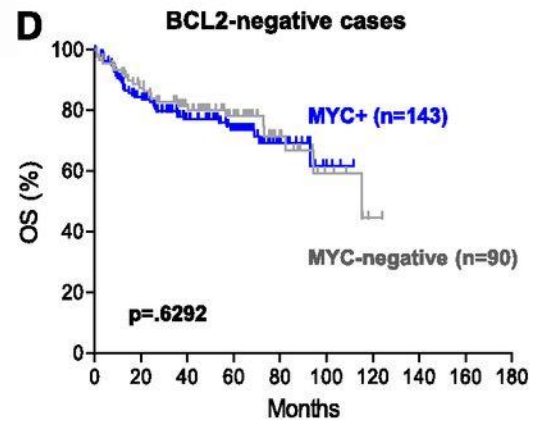
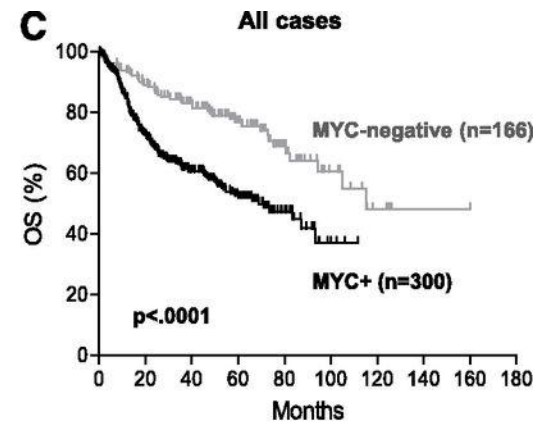
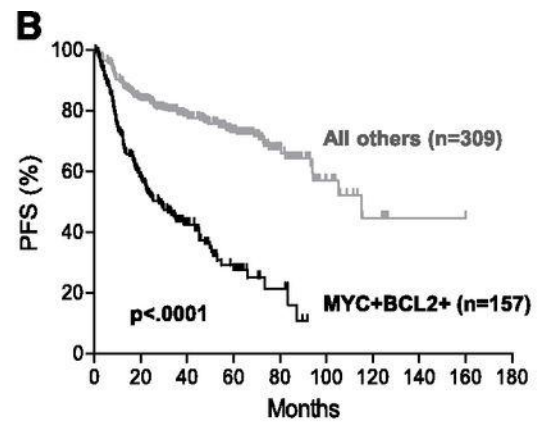
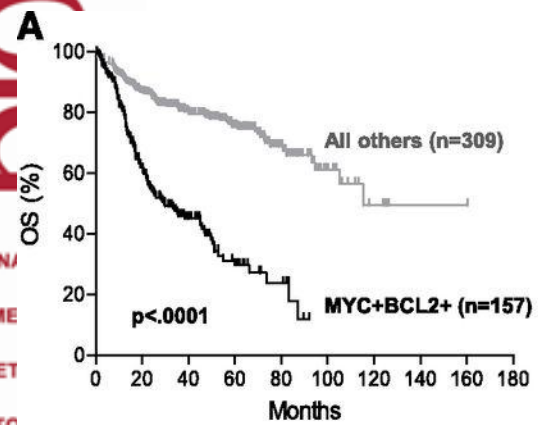
Χαρακτηριστικά ασθενών

Table 2. Multivariate analysis of clinicopathological parameters in DLBCLs treated with R-CHOP

	OS			PFS		
	HR	95% CI	<i>P</i>	HR	95% CI	<i>P</i>
B symptoms	1.47	1.04-2.09	.0310	1.45	1.03-2.03	.0314
Tumor size, ≥ 7.5 cm	1.22	0.87-1.71	.2467	1.21	0.86-1.69	.2708
IPI risk, >2	2.38	1.67-3.38	$<.0001$	2.22	1.59-3.11	$<.0001$
COO classification, ABC	1.17	0.79-1.72	.4329	1.18	0.82-1.71	.3750
<i>TP53</i> mutation	1.72	1.17-2.52	.0057	1.63	1.12-2.37	.0105
MYC/BCL2 coexpression	2.52	1.73-3.67	$<.0001$	2.45	1.71-3.51	$<.0001$

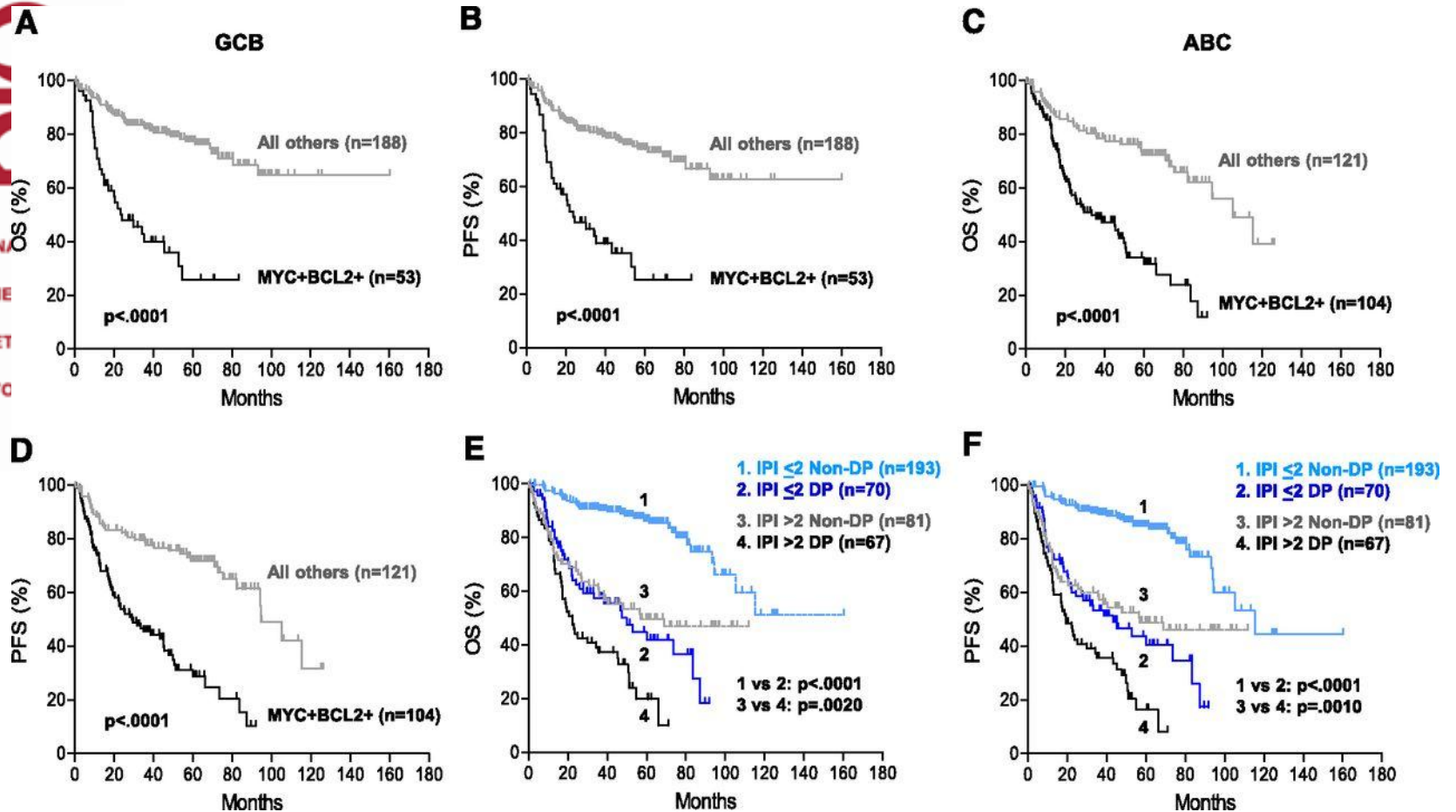
CI, confidence interval; HR, hazard ratio.

Prognostic impact of MYC/BCL2 coexpression in DLBCL. (A-B) OS (A) and PFS (B) of patients with DLBCL with MYC/BCL2 coexpression (MYC+BCL2+) in the training set.



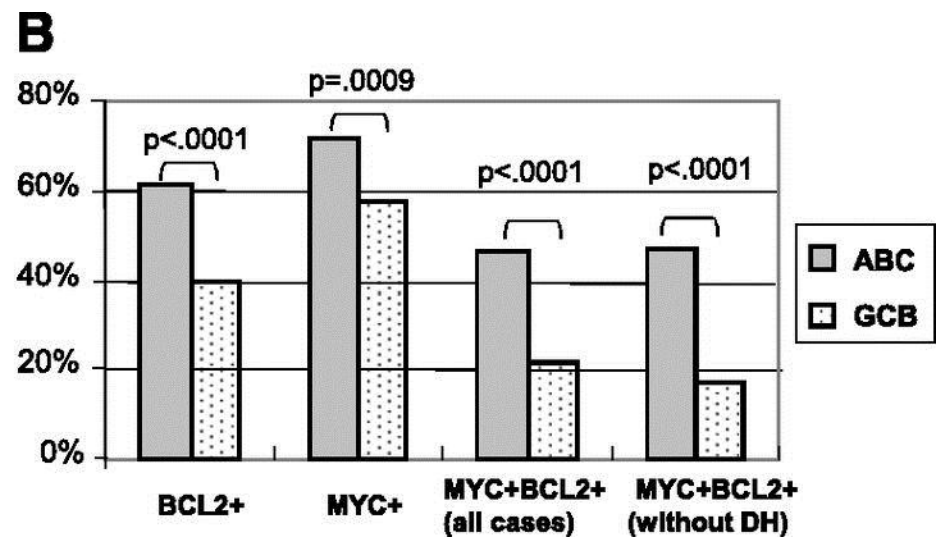
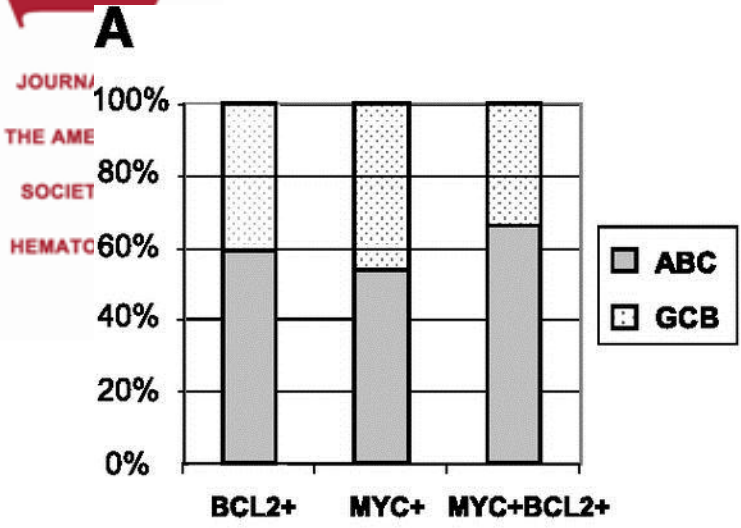
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Prognostic impact of MYC/BCL2 coexpression in DLBCL risk-stratified according to clinicopathologic parameters.



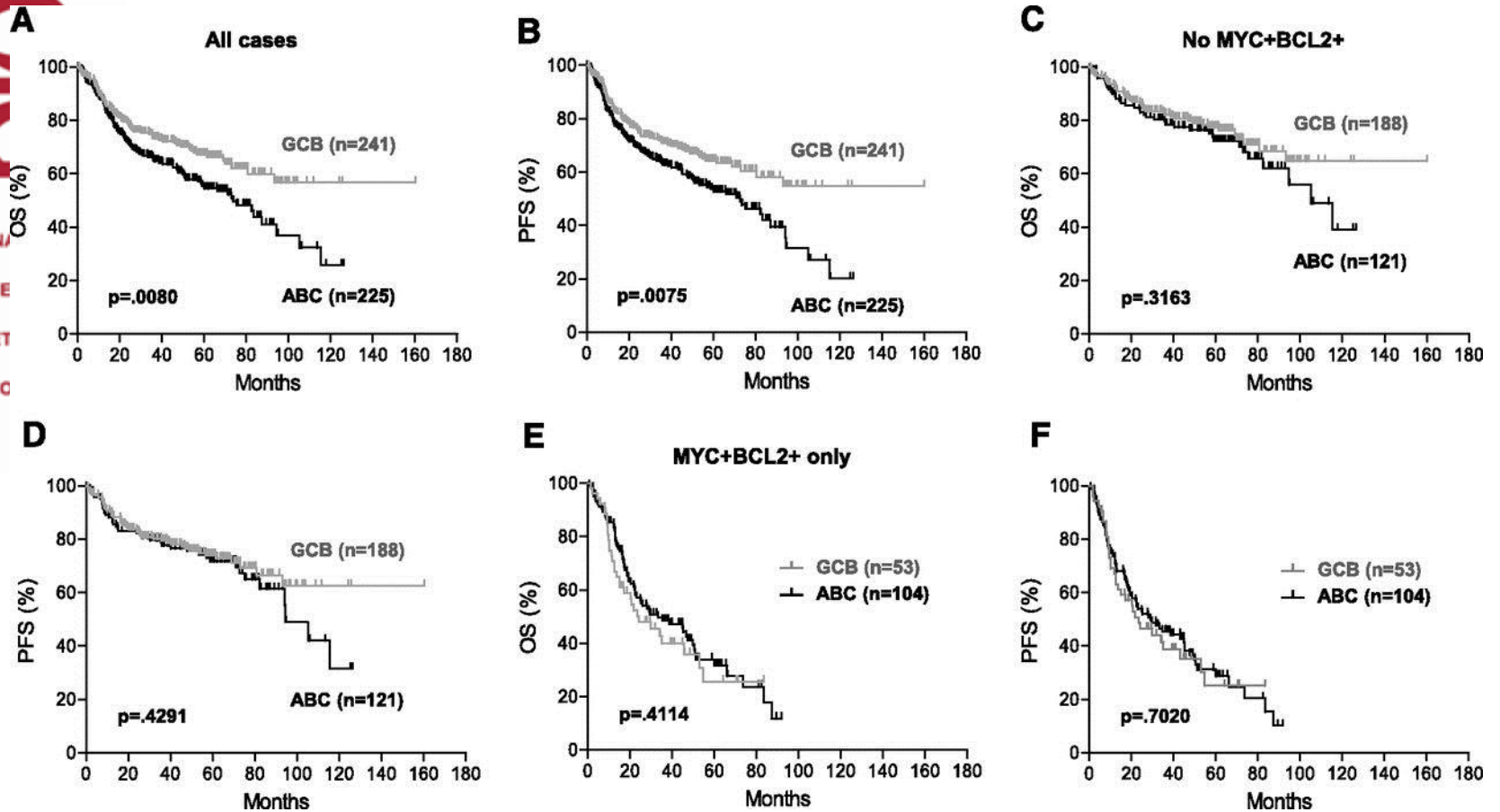
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Frequency of BCL2 and MYC expression in COO subtypes of DLBCL. (A) Relative frequency of the ABC vs GCB subtype in DLBCL positive for BCL2 expression, MYC expression, or MYC/BCL2 coexpression in the training set.



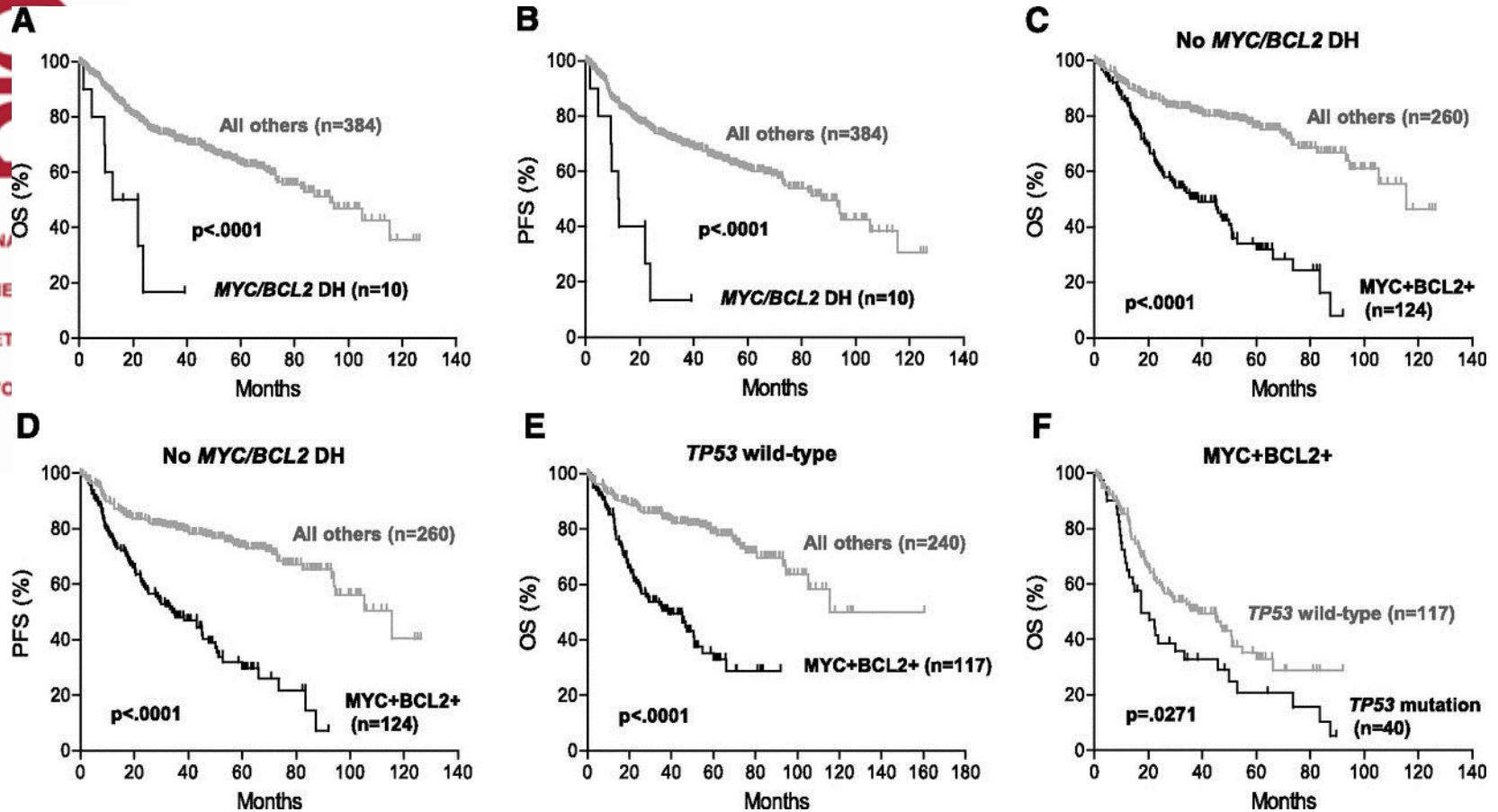
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MYC/BCL2 coexpression contributes to the inferior prognosis of ABC-DLBCL.



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Prognostic impact of MYC/BCL2 coexpression in DLBCL is independent of MYC/BCL2 corearrangement and TP53 mutation status.

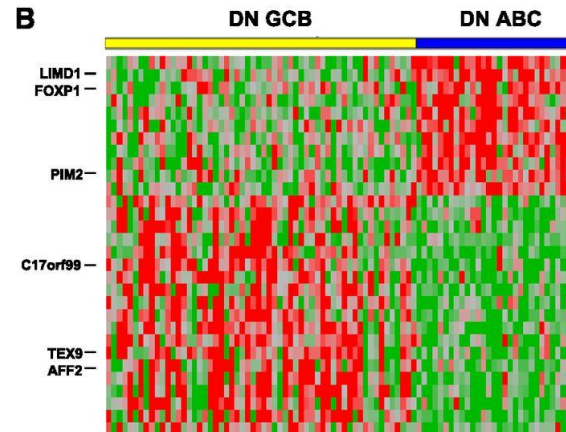
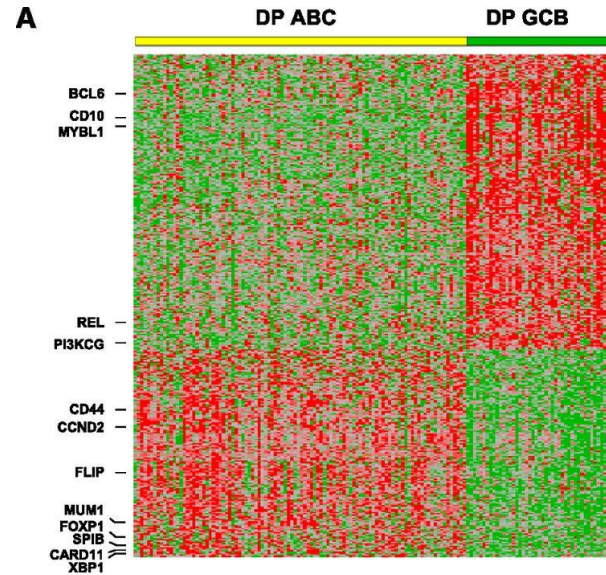


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MYC/BCL2 coexpression contributes to the different gene expression profiles between GCB and ABC subtypes of DLBCL.

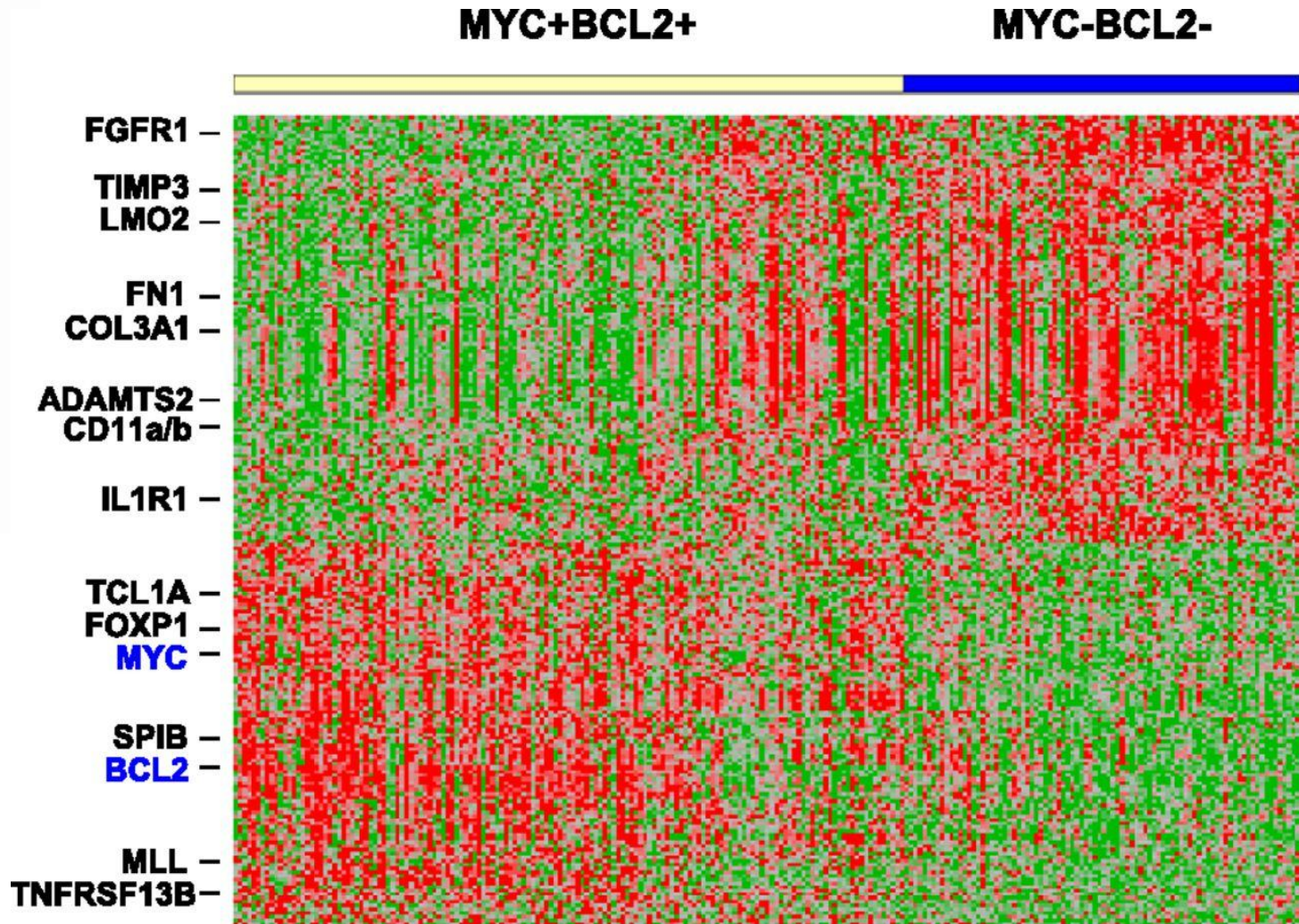
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Gene expression signature of DLBCL with MYC/BCL2 coexpression.



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Table 3. Differentially expressed genes in MYC⁺BCL2⁺ de novo DLBCL

Gene functional categories	No. of genes	Representative genes
Downregulated genes		
ECM, ECM production and remodeling	33	<i>COL3A1, VCAN, TNS1, FN1, THBS2, TIMP3, SPARC, SULF1, SPINK2, MMP2, ADAM12, FGFR1, FAP</i>
Cell adhesion and cytoskeletal organization	21	<i>CD11A/CD11B, CD58, THY1, RFTN1, ANTXR1, RHOB, MICAL2</i>
Cell growth regulation	16	<i>LM02, TRAF1, CDK14, SGK1, RGS1, NBL, PDE4D</i>
Others, including unknown	18	<i>PSAP, LYZ, LOC115110, ZNF662</i>
Upregulated genes		
Cell proliferation	20	<i>MYC, BCL2, TCL1A, MLL, FOXP1, SPIB, TCF4, TNFRSF13B, PMDAIP1, GAB1, PLOR3G</i>
Cell metabolism	5	<i>DCTPP1, CYB5R2, HK2, TMEM97, CYB5R2</i>
Miscellaneous cell functions	13	<i>PPIL1, PIGW, FUT8, SPINK5</i>
Unknown	27	<i>KIAA0664, C9orf91, ZNF107</i>

ΣΥΖΗΤΗΣΗ

Συζήτηση

- Ασθενείς με DLBCL και συνέκφραση MYC/BCL2 έχουν κακή πρόγνωση με 5ετή επιβίωση <30% ανεξαρτήτως αν πρόκειται για GCB ή ABC φαινότυπο
- 30% των ασθενών με DLBCL έχουν συνέκφραση MYC/BCL2 ενώ μόνο το 3% αντιπροσωπεύει DH λέμφωμα που αναδεικνύει ότι η επιθετικότητα των παραπάνω λεμφωμάτων είναι πέρα από το γενετικό επίπεδο
- Ασθενείς με MYC-/BCL2- έχουν μη στατιστικά σημαντική διαφορά στην επιβίωση ανεξαρτήτως GCB ή ABC φαινοτύπου
- Η συνέκφραση MYC/BCL2 παρατηρείται πιο συχνά στον ABC φαινότυπο που πιθανόν να ευθύνεται για την χειρότερη πρόγνωση αυτής της κατηγορίας ασθενών

LYMPHOID NEOPLASIA

CME Article

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Disclosures

The authors, Associate Editor A. Keith Stewart, and CME questions author Charles P. Vega, Associate Professor and Residency Director, Department of Family Medicine, University of California-Irvine, declare no competing financial interests.

Learning objectives

Upon completion of this activity, participants will be able to:

1. Assess genetic abnormalities associated with diffuse large B-cell lymphoma (DLBCL).
2. Analyze the prevalence and survival impact of MYC and BCL2 co-expression in the current study.
3. Distinguish the relationship between MYC/BCL2 co-expression and other negative prognostic variables in the current study.
4. Evaluate the relative effect of MYC/BCL2 co-expression on survival in the context of DLBCL subtypes.

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