

Webinars Cutaneous Lymphoma

EuroBloodNet  Topic on Focus

Primary cutaneous Diffuse Large B-cell Lymphoma, Leg Type

Prof. Anne Pham-Ledard

CHU Bordeaux, Dermatology Department, France

INSERM U 1053 UMR BaRITON; Cutaneous lymphoma oncogenesis team;

ERN-EuroBloodNet subnetwork cutaneous lymphoma

Bordeaux– France, November 2020



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of the European Union



université
de BORDEAUX



European
Reference
Network
for rare or low prevalence
complex diseases

Network
Hematological
Diseases (ERN EuroBloodNet)



Investigator : Celgene, Roche, BMS, MSD
Travel, Accomodation, Expenses: BMS, MSD.



First line therapy = Rituximab + polychemotherapy

In refractory / relapsing cases ... Which therapy?

- Radiation ?
- Second line chemotherapy?
- Lenalidomide?
- Other targeted therapies ?



Clinical case : chemoresistance with second line therapy

82 years old women
PCDLBCL Leg type (knee)



Cutaneous relapse (leg)
3 months later



Diffuse finger tumor infiltration
while receiving chemo



R-CHOPx6 → CR

Rituximab + Gemcitabine + Oxaliplatine

- Poor response
- Impaired quality of life, chemo side effects in elderly



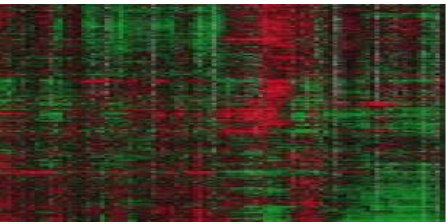
It is mandatory to :

Understand genomic landscape
Oncogenesis mechanisms

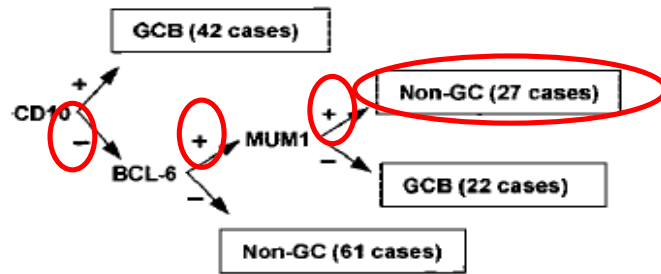
→ **To develop new therapies**



Leg type Lymphoma exhibit same features as ABC DLBCL



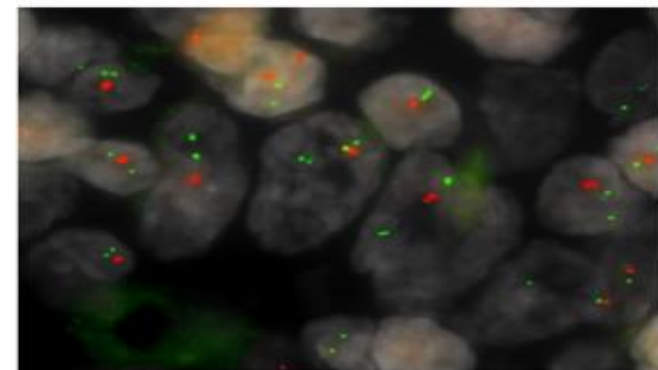
Transcriptomic profile



Han's algorithm, immunophenotype

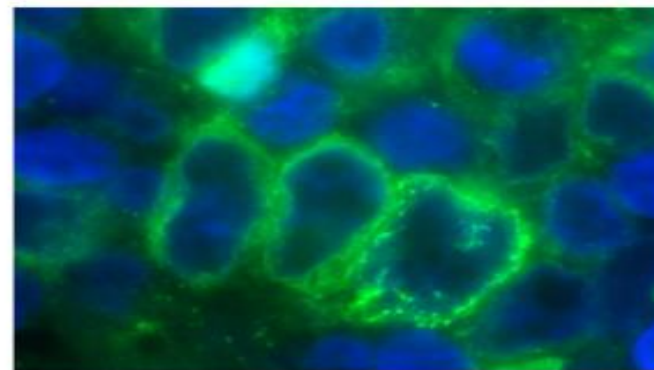
- **Transcriptomic and phenotypic profile** consistent with ABC DLBCL

Hoefnagel et al, Blood, 2005
Menguyet al, Histopathology 2019



Cmu deletion (switch)

D



IgM expression

Pham-Ledard et al, J Derm Sci, 2017,

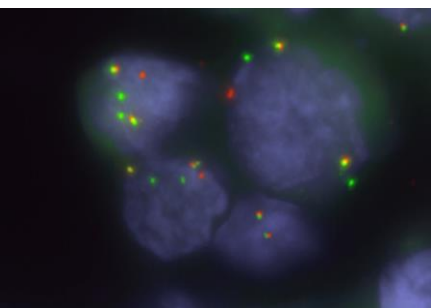
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- **IgH Gene analysis:** post germinal-center profile

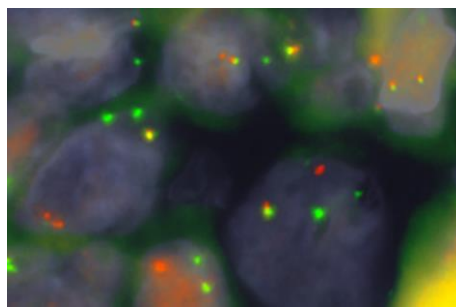
- **Oncogene alterations**

- **CDKN2A/ p16 deletion**
- **MYC (10-30%), BCL6 (25%)**
- No BCL2 (0%)

- = as DLBCL, ABC (non GC) subtype



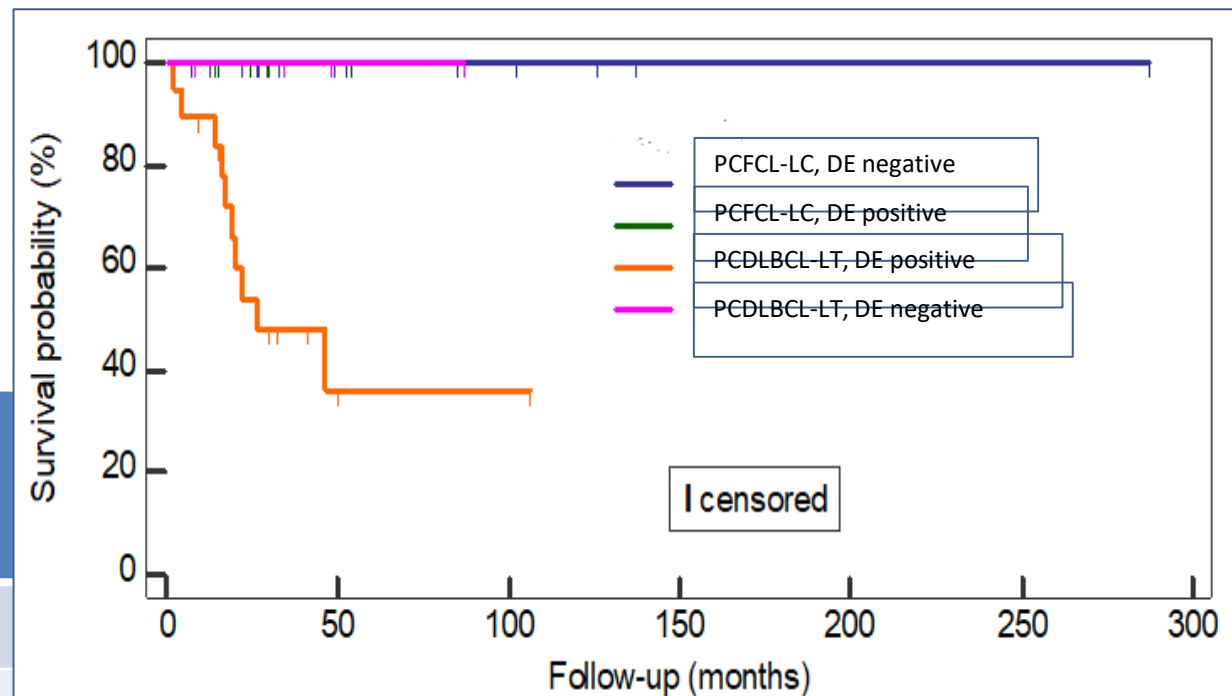
FISH *BCL6*



FISH *MYC*

Controversies about MYC/BCL2 Hit / Dual Expression

| | <i>BCL2 split</i> |
|--------------------------|-------------------|
| Pham-Ledard et al., 2013 | 1/23 (4%) |
| Lucioni et al., 2016 | 1/20 (5%) |
| Menguy et al., 2018 | 0/23 |
| Schrader et al., 2018 | |



Poor Specific survival in PCDLBCL leg type with BCL2-MYC expression

14/44 (32%) 2/44 (4%) 20/45 (44%)

MYC split (in this study)

Associated with **reduced specific survival and disease free survival** but not for overall survival

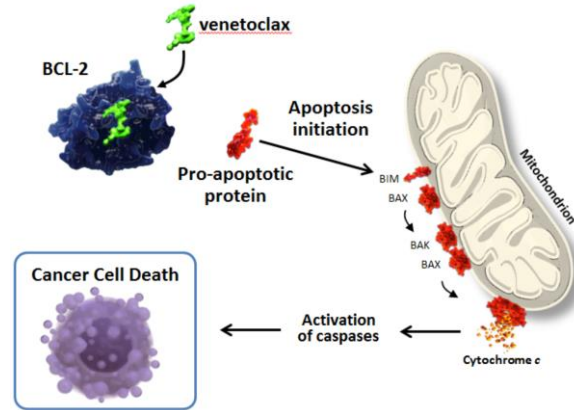
DH = Very rare event

**DE=Frequent
Prognosis impact?**

Successful Treatment of Primary Cutaneous Diffuse Large B-Cell Lymphoma Leg Type With Single-Agent Venetoclax

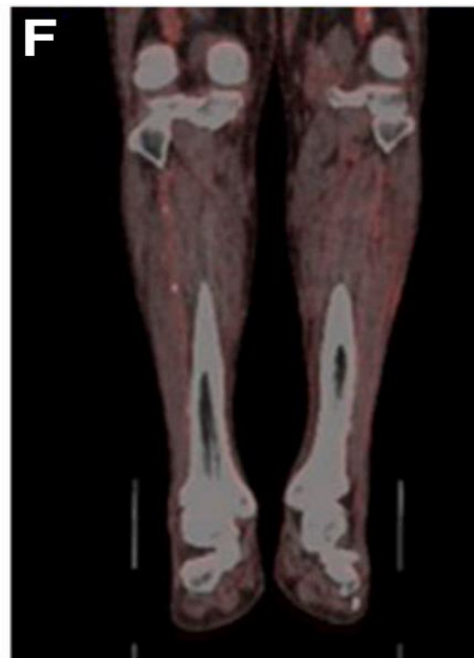
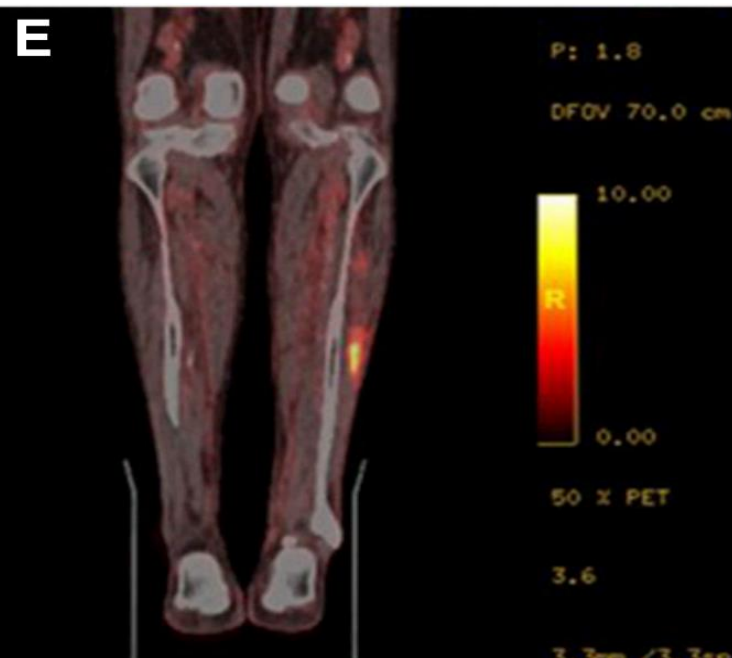
Harriet S. Walter, MBChB, PhD^{1,2}; Christopher S. Trethewey, MSc¹; Matthew J. Ahearne, MBChB, MD^{1,2}; Ross Jackson¹; Sandrine Jayne, PhD¹; Simon D. Wagner, MD, PhD^{1,2}; Gerald Saldanha, MBChB^{1,2}; and Martin J.S. Dyer, MD, DPhil^{1,2}

BCL2 Inhibitor Venetoclax



- Active in DLBCL with BCL2 alterations (ampli/DH)
- In PCDLBCL leg type: no *BCL2* mutation/ampli/hit
- **BCL2 strong expression**

75 years old man, Refractory to poly chemo
MYD88 WT, *BCL2* WT



ORIGINAL ARTICLE

See related commentary on pg 1831

Identification of Somatic Mutations in Primary Cutaneous Diffuse Large B-Cell Lymphoma, Leg Type by Massive Parallel Sequencing



Sylvain Mareschal¹, Anne Pham-Ledard^{2,3}, Pierre Julien Viailly¹, Sydney Dubois¹, Philippe Bertrand¹, Catherine Maingonnat¹, Maxime Fontanilles¹, Elodie Bohers¹, Philippe Ruminy¹, Isabelle Tournier⁴, Philippe Courville⁵, Bernard Lenormand⁵, Anne Bénédicte Duval⁵, Emilie Andrieu⁵, Laurence Verneuil⁶, Beatrice Vergier^{2,3}, Hervé Tilly¹, Pascal Joly⁴, Thierry Frebourg⁴, Marie Beylot-Barry^{2,3}, Jean-Philippe Merlio^{2,3} and Fabrice Jardin¹

Journal of Investigative Dermatology (2018) 138, 2365–2376; doi:10.1016/j.jid.2018.04.038



ORIGINAL ARTICLE

Genomic Analyses Identify Recurrent Alterations in Immune Evasion Genes in Diffuse Large B-Cell Lymphoma, Leg Type

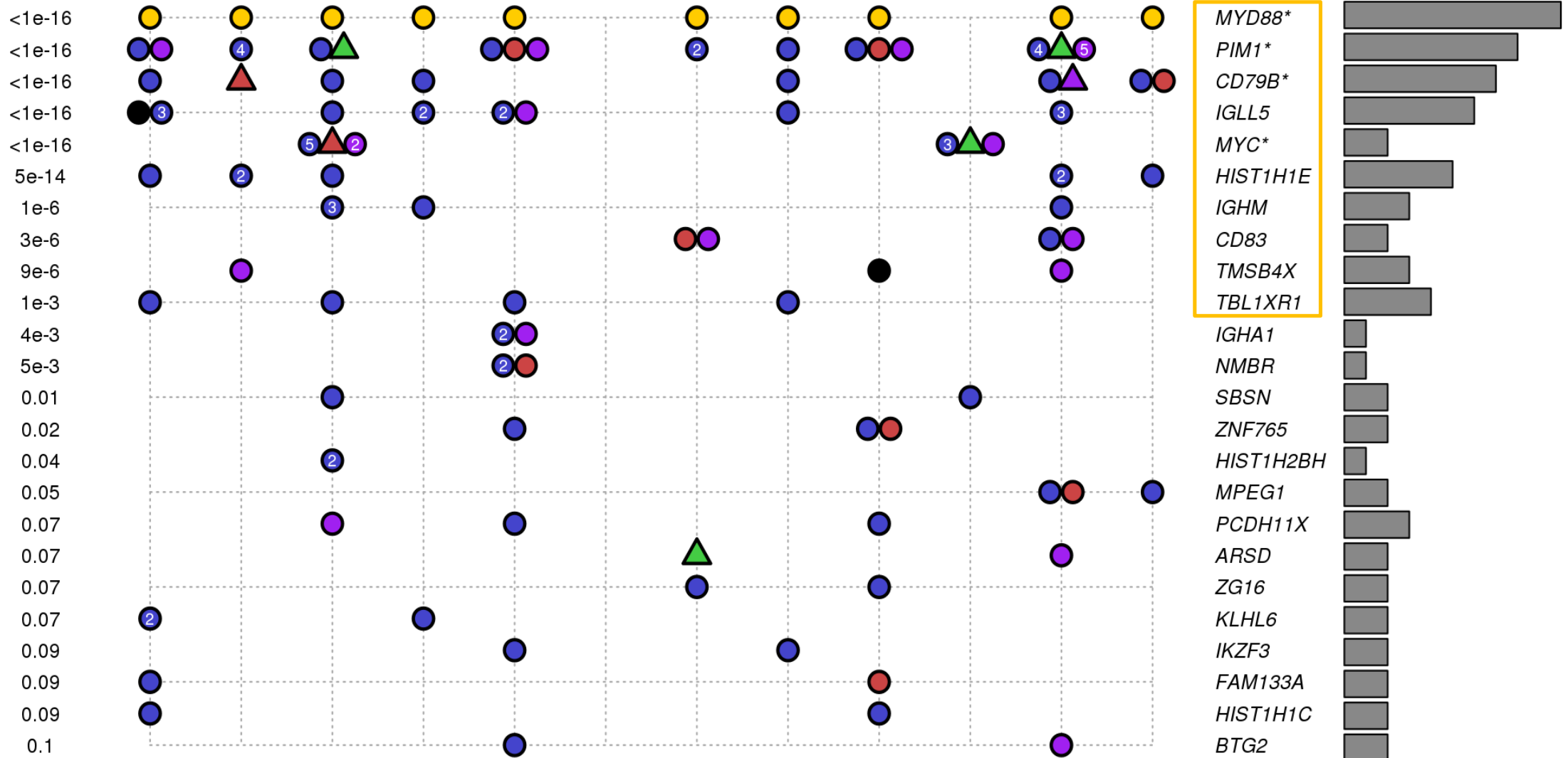


Xiaolong Alan Zhou^{1,12}, Abner Louissaint Jr.^{2,3,12}, Alexander Wenzel⁴, Jingyi Yang^{4,5}, Maria Estela Martinez-Escala¹, Andrea P. Moy^{2,3}, Elizabeth A. Morgan⁶, Christian N. Paxton⁷, Bo Hong⁸, Erica F. Andersen⁸, Joan Guitart¹, Amir Behdad⁹, Lorenzo Cerroni¹⁰, David M. Weinstock^{3,11} and Jaehyuk Choi^{1,4,5}

Webinars
Cutaneous Lymphoma

EuroBloodNet  Topic on Focus

NGS profile: confirms MYD88 frequency and shows recurrent genes mutations



Mareschal, Pham-Ledard et al, J Invest Dermatol 2018



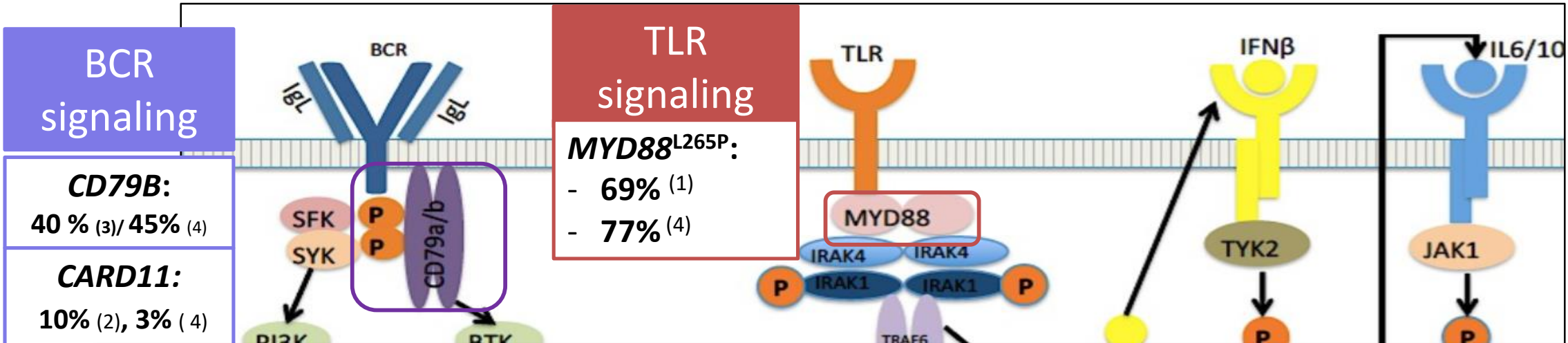
| | PCLBCL-LT n = 20 | PCNSL n = 25 | ABC n = 81 | GCB n = 83 | PMBL n = 18 | FDR |
|----------|---------------------|-----------------|---------------|---------------|----------------|---------|
| STAT6 | 0% | 0% | 0% | 14% | 72% | 1.6e-56 |
| MYD88 | 75% | 80% | 28% | 10% | 0% | 1.9e-47 |
| XPO1 | 0% | 0% | 1% | 1% | 39% | 5.3e-32 |
| CIITA | 5% | 0% | 12% | 10% | 56% | 8.6e-30 |
| PIM1 | 70% | 32% | 33% | 8% | 0% | 8.6e-30 |
| SOCS1 | 15% | 4% | 6% | 16% | 56% | 5.0e-23 |
| TNFAIP3 | 5% | 24% | 15% | 11% | 61% | 2.2e-22 |
| CD58 | 5% | 0% | 6% | 10% | 39% | 3.3e-18 |
| B2M | 5% | 12% | 9% | 18% | 50% | 4.3e-17 |
| CD79B | 40% | 32% | 25% | 2% | 0% | 3.0e-16 |
| MFHAS1 | 0% | 0% | 1% | 10% | 28% | 3.7e-16 |
| BCL2 | 0% | 4% | 1% | 24% | 0% | 1.5e-15 |
| GNA13 | 10% | 16% | 9% | 12% | 50% | 2.3e-15 |
| TNFRSF14 | 0% | 0% | 2% | 17% | 0% | 5.8e-12 |
| ITPKB | 5% | 8% | 9% | 16% | 39% | 1.4e-11 |
| EZH2 | 0% | 4% | 0% | 18% | 6% | 4.6e-08 |
| KMT2D | 15% | 28% | 41% | 46% | 17% | 3.2e-07 |
| CREBBP | 15% | 8% | 6% | 31% | 11% | 2.6e-06 |
| TP53 | 0% | 8% | 19% | 16% | 11% | 2.6e-04 |
| NOTCH2 | 5% | 0% | 2% | 10% | 0% | 4.0e-04 |
| MYC | 20% | 4% | 5% | 10% | 6% | 5.1e-04 |
| PRDM1 | 5% | 8% | 16% | 6% | 0% | 5.1e-04 |
| CDKN2A | 0% | 8% | 2% | 1% | 0% | 5.1e-04 |
| MEF2B | 10% | 4% | 12% | 23% | 11% | 1.8e-03 |
| CARD11 | 5% | 8% | 14% | 7% | 0% | 3.7e-03 |
| BRAF | 0% | 4% | 0% | 0% | 0% | 3.7e-03 |
| EP300 | 5% | 4% | 15% | 14% | 17% | 6.2e-03 |
| FOXO1 | 5% | 0% | 4% | 12% | 6% | 6.5e-03 |
| CD79A | 5% | 0% | 0% | 2% | 0% | 8.9e-03 |
| ID3 | 0% | 0% | 5% | 2% | 6% | 1.7e-02 |
| NOTCH1 | 5% | 0% | 7% | 1% | 6% | 3.4e-02 |
| TCF3 | 0% | 4% | 1% | 2% | 0% | 9.3e-02 |
| IRF4 | 15% | 12% | 14% | 5% | 11% | 2.0e-01 |
| CDKN2B | 0% | 0% | 0% | 1% | 0% | 4.1e-01 |



**Mutations in PCDLBCL-leg
 type: closer pattern to CNS
 (Central Nervous System)
 than other DLBCL**

- High **MYD88**
- PIM1**
- CD79B**
- **TBL1XR1: 30%**
- CREBBP**
- MYC**
- IRF4**

Genetic landscape of PCDLBCL Leg -Type



Whether the genetic profile may predict complete durable response or relapsing-refractory disease?

(1) Pham-Ledard et al. *J Invest Dermatol.* 2012; (2) Koens L et al. *J Invest Dermatol.* 2014 ;

(2) (3) Mareschal J et al. *J Invest Dermatol.* 2012 (4) Zhou et al. *J Invest Dermatol.* 2019

Targeted NGS of PCDLBCL leg type : RR versus good response

Biopsy at diagnosis
n=32

Refractory/
Relapse
n= 18

Responders
n= 14

Targeted NGS

Lymphopanel designed for DLBCL targeting 36 genes

NOTCH
NOTCH1,
NOTCH2

**Epigenetic
Regulation**
ARID1A,
EZH2, EP300,
MEF2B,
CREBBP

MAP Kinase
BRAF

**Apoptosis/
Cell cycle**
CCND3,
XPO1, MYC,
CDKN2A/B,
FOXO1, TP53,
GNA13, BCL2

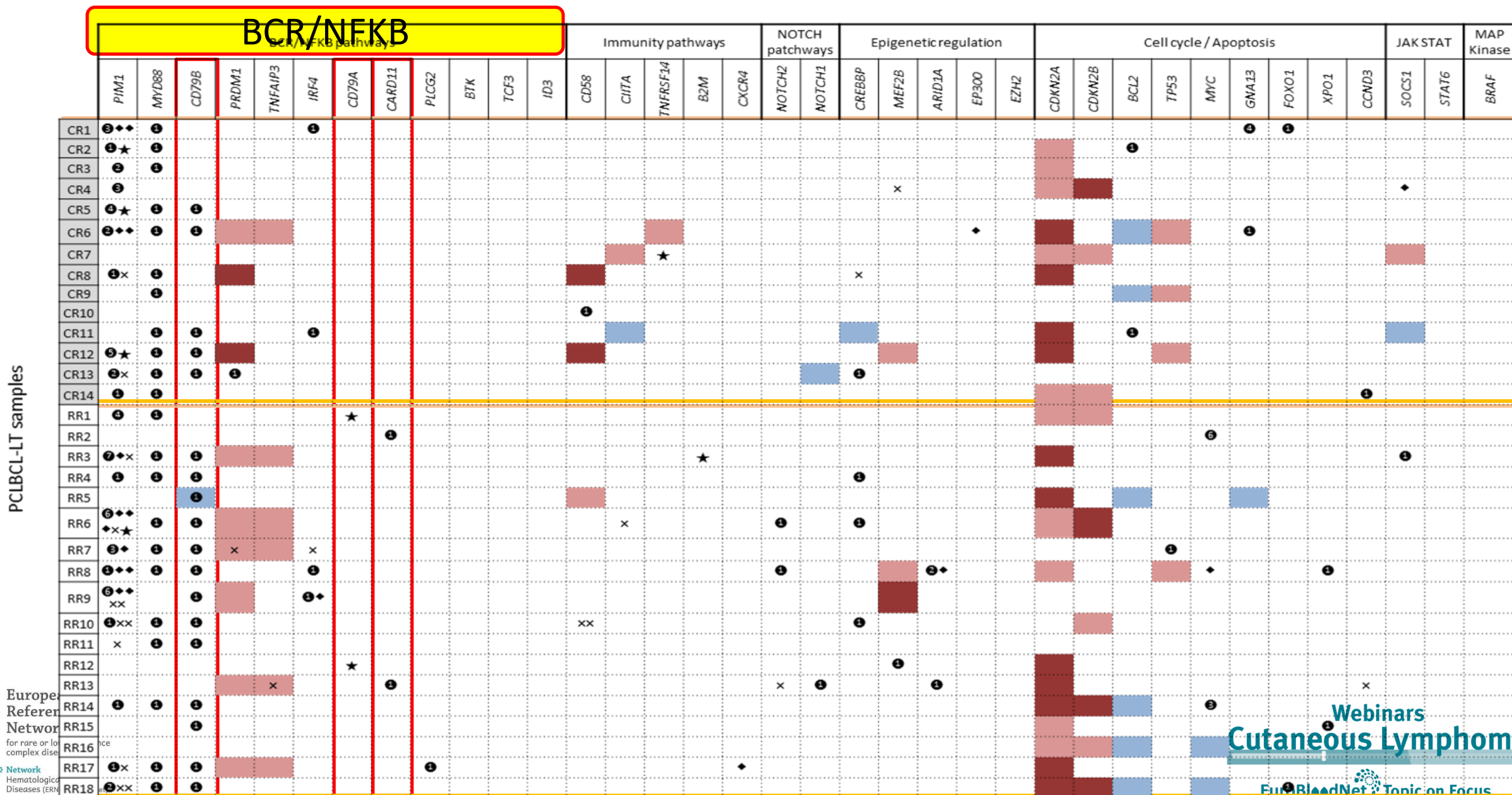
NF-kB
TNFAIP3,
MYD88, PIM1,
CARD11,
IRF4, PRDM1

BCR
BTK,
CD79A/B,
TCF3, ID3

JAK STAT
SOCS1,
STAT6

Immunity
CIITA, CXCR4,
B2M, PLCG2,
TNFRSF14,
CD58

More BCR signaling mutations (CD79A/B / CARD11) in the RR subgroup

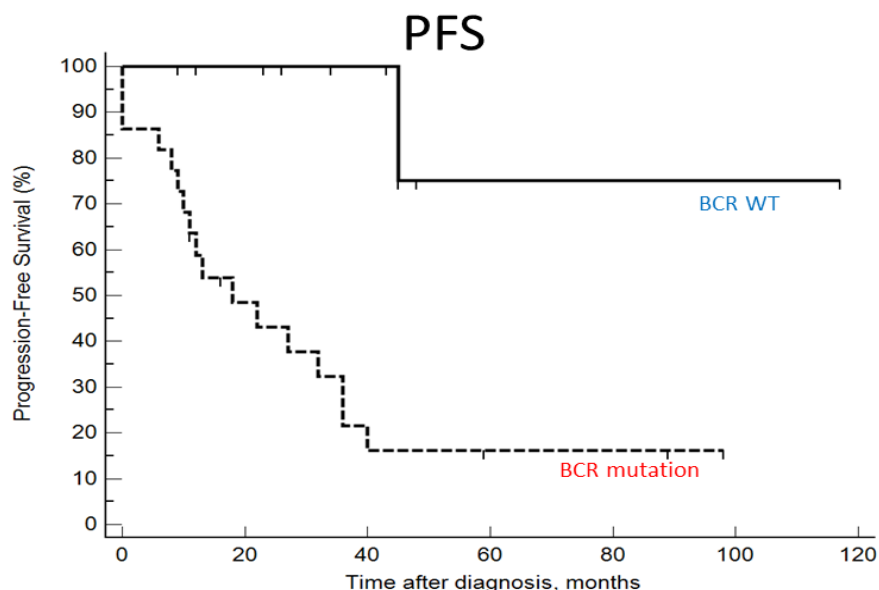


Mutations of the B-Cell Receptor Pathway Confer Chemoresistance in Primary Cutaneous Diffuse Large B-Cell Lymphoma Leg Type

Océane Ducharme^{1,2}, Marie Beylot-Barry^{1,2}, Anne Pham-Ledard^{1,2}, Elodie Bohers³, Pierre-Julien Viailly³, Thomas Bandres⁴, Nicolas Faur⁴, Eric Frison⁵, Béatrice Vergier^{2,6}, Fabrice Jardin³, Jean-Philippe Merlio^{2,4} and Audrey Gros^{2,4}

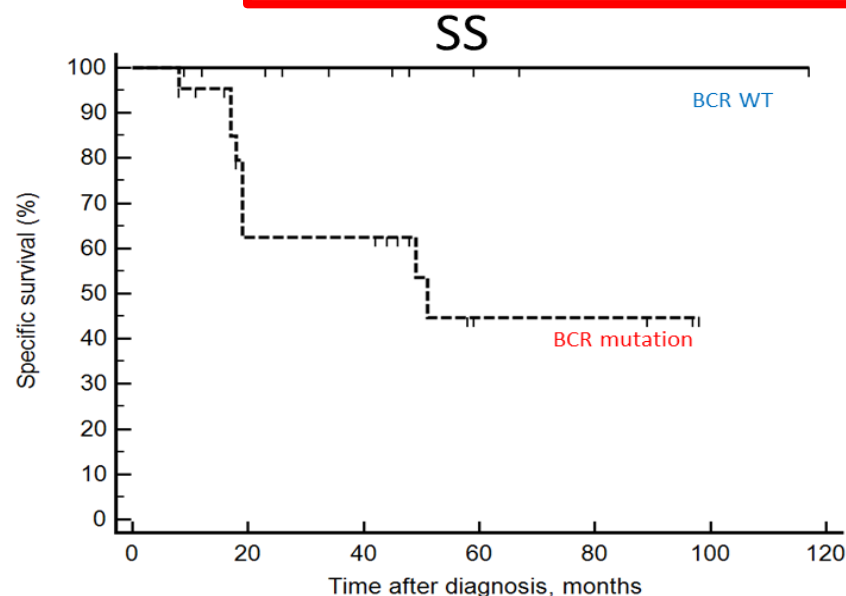


Patients with BCR mutation displayed a reduced PFS and specific survival



| Number at risk | 0 | 20 | 40 | 60 | 80 | 100 | 120 |
|----------------|----|----|----|----|----|-----|-----|
| Group: 0 | 10 | 8 | 5 | 1 | 1 | 1 | 0 |
| Group: 1 | 19 | 9 | 3 | 2 | 2 | 0 | 0 |

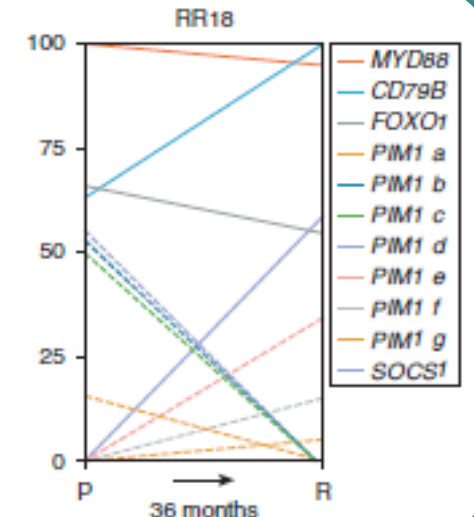
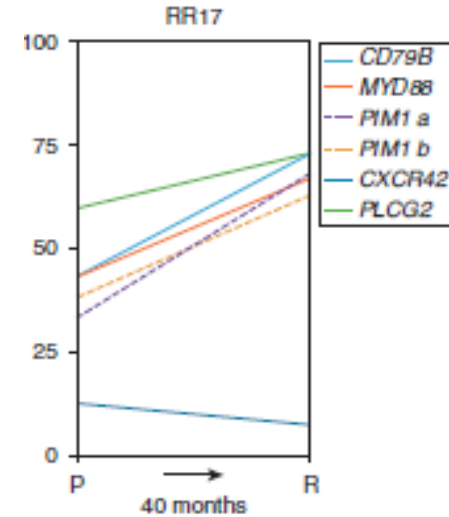
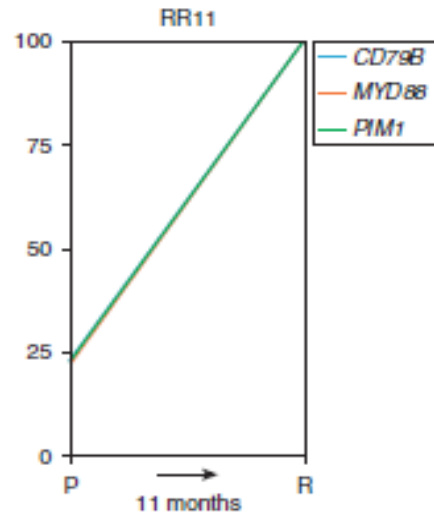
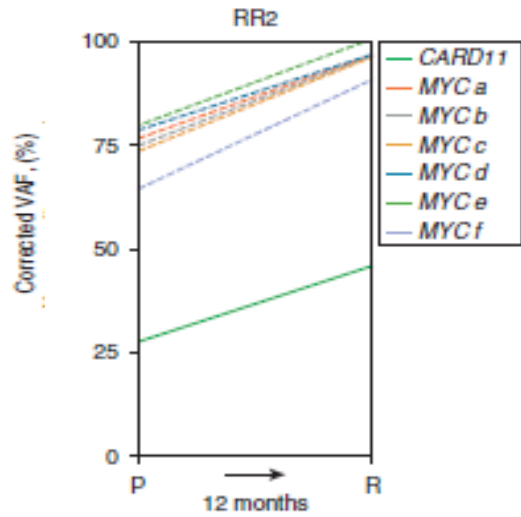
Median 18 months, Log-rank, $P=0.002$
not reached in the wild-type group



| Number at risk | 0 | 20 | 40 | 60 | 80 | 100 | 120 |
|----------------|----|----|----|----|----|-----|-----|
| Group: 0 | 10 | 8 | 5 | 2 | 1 | 1 | 0 |
| Group: 1 | 22 | 11 | 11 | 3 | 3 | 0 | 0 |

Median 51 months, Log-rank, $P=0.03$
not reached in the wild-type group

Clonal selection between diagnosis and relapse: BCR and *MYD88* mutations

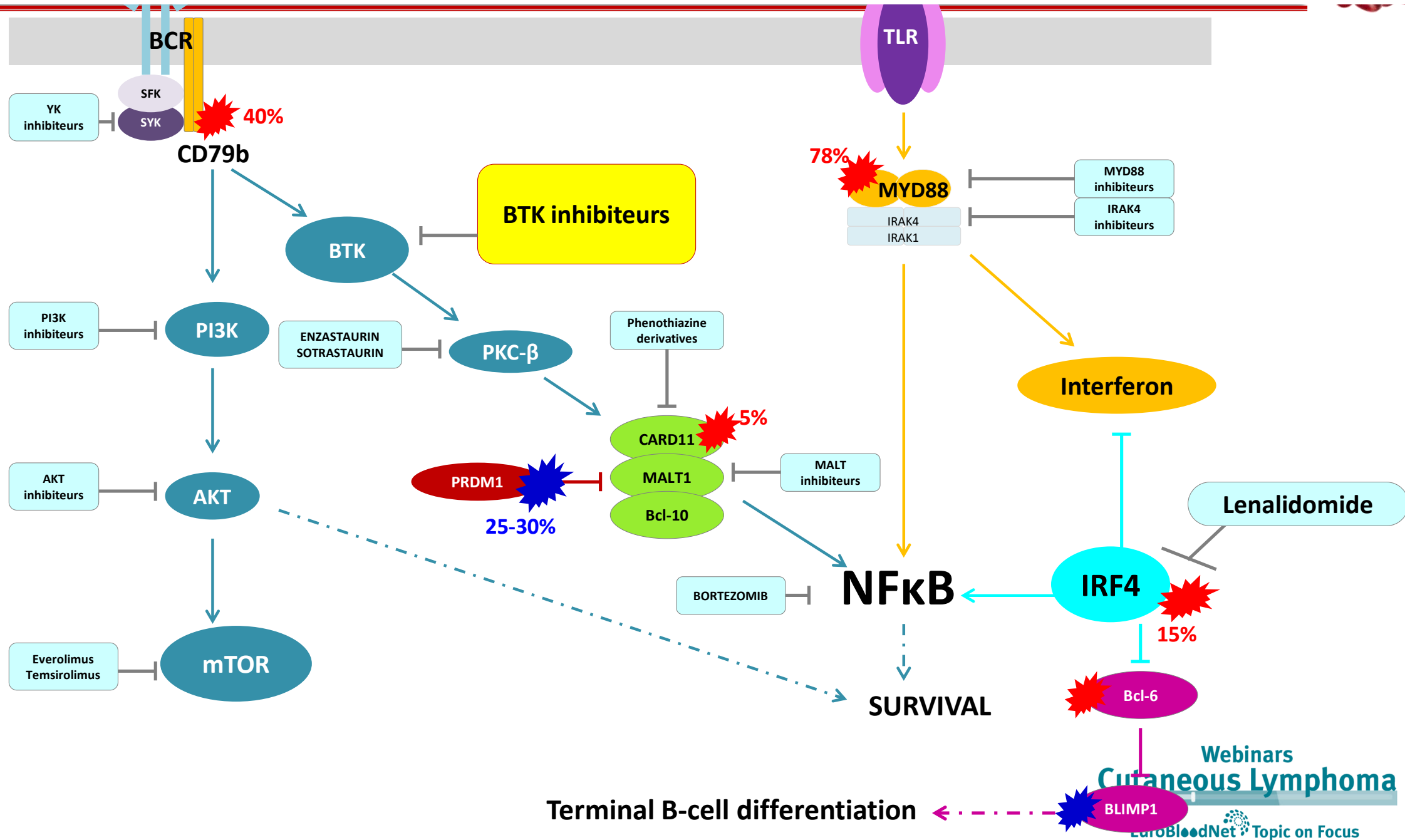


VAF enrichment of primary mutations affecting

- *CD79B*
- *CARD11*
- *MYD88*

➔ **pivotal role in disease oncogenesis**

BCR targeting in these patients?



Excellent outcome of immunomodulation or Bruton's tyrosine kinase inhibition in highly refractory primary cutaneous diffuse large B-cell lymphoma, leg type

Eva Gupta,¹ Joseph Accurso,² Jason Sluzevich,³ David M. Menke,⁴ Han W. Tun¹

¹Division of Hematology and Oncology,
²Department of Diagnostic Radiology,
³Department of Dermatology,
⁴Department of Pathology, Mayo Clinic, Jacksonville, FL, USA

Case Report

Molecular Mechanisms of Disease Progression in Primary Cutaneous Diffuse Large B-Cell Lymphoma, Leg Type during Ibrutinib Therapy

Lucy C. Fox, Costas K. Yannakou, Georgina Ryland, Stephen Lade, Michael Dickinson, Belinda A. Campbell and Henry Miles Prince *

Addictive response of primary cutaneous diffuse large B cell lymphoma leg type to low-dose ibrutinib. Pang A, Au-Yeung R, Leung RYY, Kwong YL. Ann Hematol. 2019

| Genomic Alteration | Pre-Ibrutinib (Skin) | Post-Ibrutinib (Node) |
|---------------------|---|--|
| Variants | MYD88 c.794T > C;p.L265P CD79B c.586T > C;p.Y196H | MYD88 c.794T > C;p.L265P CD79B c.586T > C;p.Y196H CARD11 c.367G > T;p.G123C CARD11 c.644A > T;p.K215M NFKBIE c.1379G > C;p.G460A |
| Copy-number changes | Gain chr7, chr8, chr9, chr10, 11q, chr12, 19q, chrX Amplification 18q (BCL2/TNFRSF11A) Del 4q Homozygous deletion 9p containing CDKN2A Second homozygous deletion 9p containing PTRPD | Gain 8q, chr9, 11q, chr12, 16p Amplification 18q (BCL2/TNFRSF11A) Del 2q, 4p, 4q, 6p, 7q, 8p, 8q, 16p, 16q, 17p, 17q, 19p, Homozygous deletion 9p containing CDKN2A Second homozygous deletion 9p containing PTRPD |
| Translocation | Not detected | t(14;16)(q32.33;q24.1) IgH-IRF8 |

BTK Inhibitor

July 2019
65 years woman
PCDLBCL Leg type



R-CHOPx6 → CR

Feb 2020
3 months later
Cutaneous Relapse



LENALIDOMIDE + Rituximab

June 2020
Lenalidomide refractory



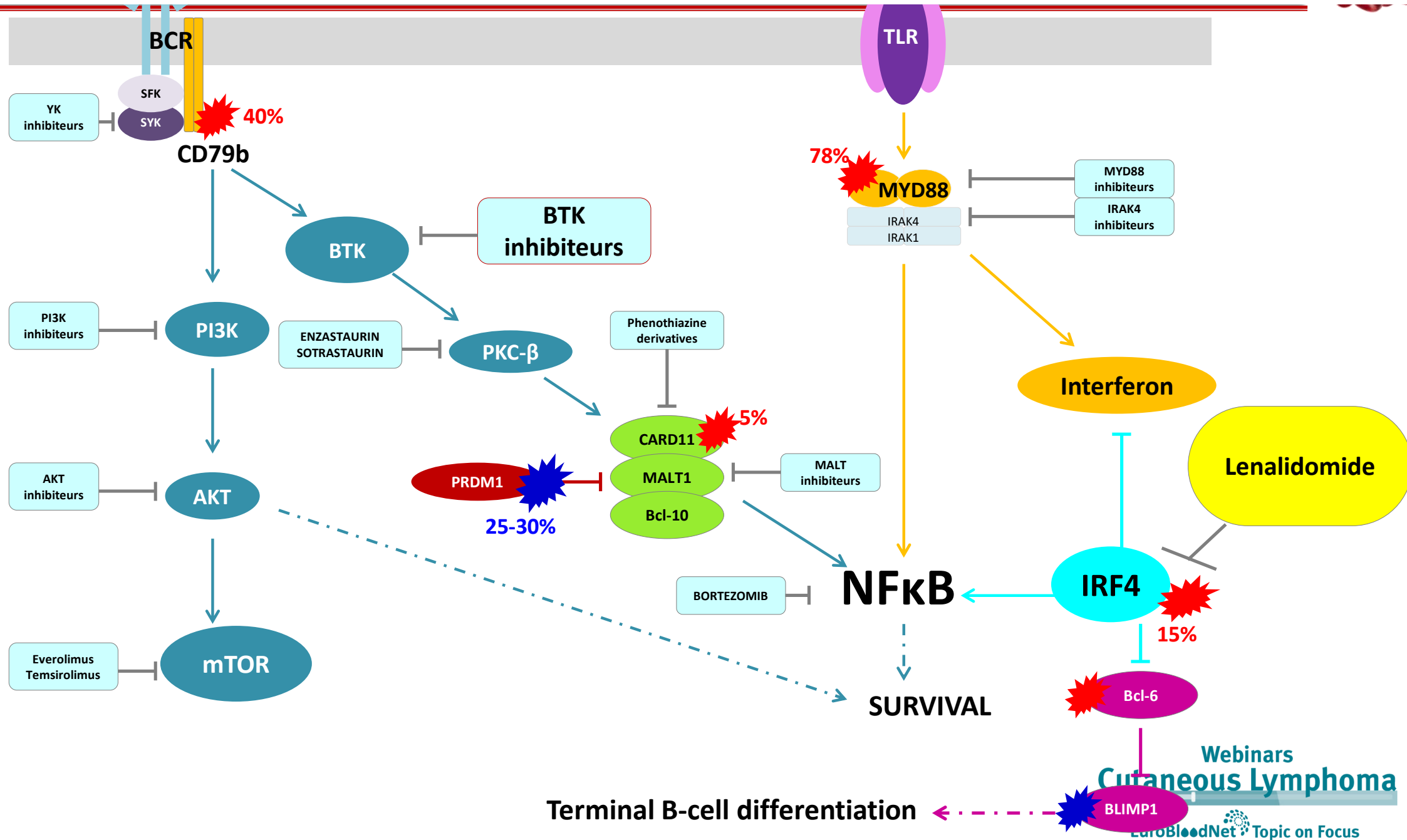
IBRUTINIB

Oct 2020
Complete Remission



Mutations: MYD88, CD79B, PIM1

Lenalidomide in PCDLBCL Leg type?



A Single-Arm Phase II Trial of Lenalidomide in Relapsing or Refractory Primary Cutaneous Large B-Cell Lymphoma, Leg Type



Marie Beylot-Barry^{1,2,19}, Diane Mermin^{1,19}, Aline Maillard³, Reda Bouabdallah⁴, Nathalie Bonnet⁵, Anne-Bénédicte Duval-Modeste⁶, Laurent Mortier⁷, Saskia Ingen-Housz-Oro⁸, Caroline Ram-Wolff⁹, Stéphane Barette¹⁰, Stéphane Dalle¹¹, Eve Maubec^{12,20}, Gaelle Quereux¹³, Isabelle Templier¹⁴, Martine Bagot⁹, Florent Grange¹⁵, Pascal Joly⁶, Béatrice Vergier^{2,16}, Pierre-Julien Vially¹⁷, Audrey Gros^{2,18}, Anne Pham-Ledard^{1,2}, Eric Frison³ and Jean-Philippe Merlio^{2,18}



Lenalidomide

Clinical case

Feb 2019



Diffuse cutaneous relapse



Low dose
Lenalidomide
10 mg/day
10 days / month

**October 2020:
minimal disease**



Aug 2018
85 years man
PCDLBCL Leg type



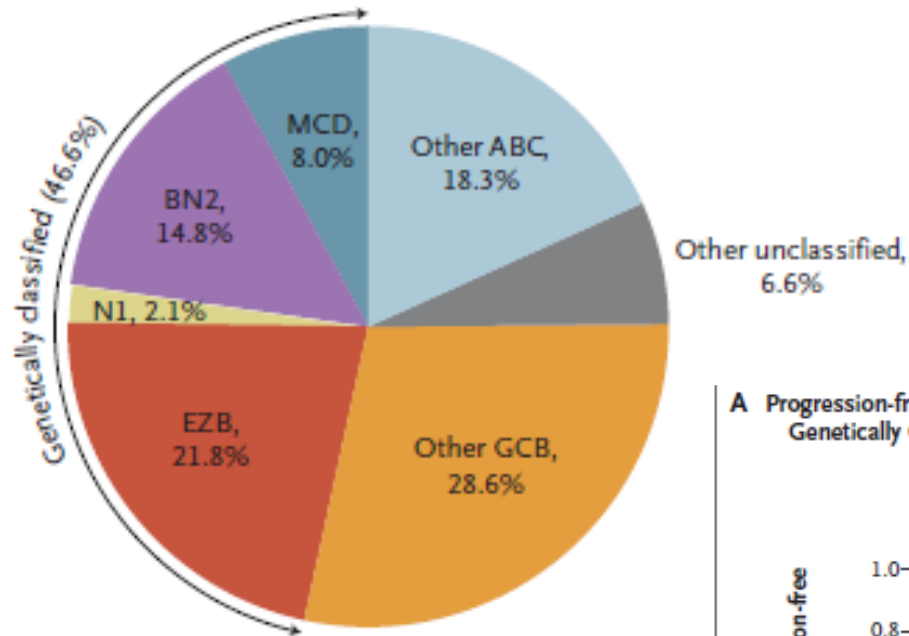
R-CHOPx6 → CR

NGS: mutation **MYD88**
PIM1
GNA13

LENALIDOMIDE + Rituximab



240 DLBCL at diagnosis, prior to R-CHOP therapy



Schmitz et al, NEJM 2018

Exome and RNAseq classification

MCD: *MYD88*^{L265P} and *CD79B* mutated

N1: *NOTCH1* mutation

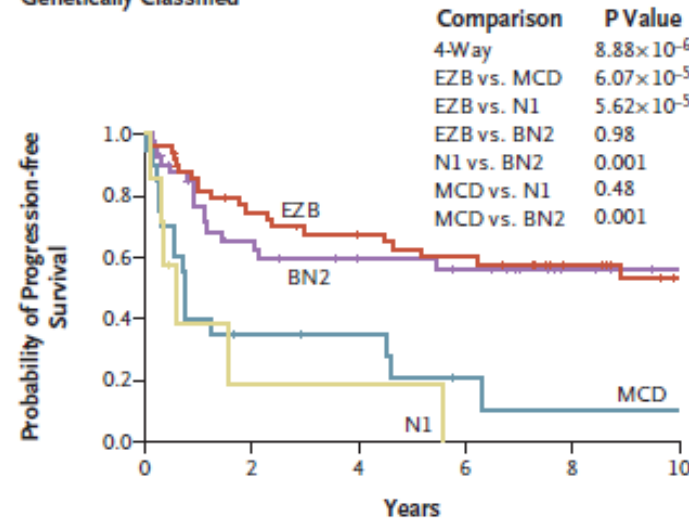
BN2: *BCL6* fusion and *NOTCH2* mutation

EZB: *EZH2* mutation and *BCL2* rearrangement

ABC

GC

A Progression-free Survival among Patients Whose Tumors Were Genetically Classified



No. at Risk

| | 0 | 2 | 4 | 6 | 8 | 10 |
|-----|----|----|----|----|----|----|
| MCD | 20 | 6 | 5 | 2 | 1 | 1 |
| BN2 | 41 | 22 | 17 | 15 | 8 | 5 |
| N1 | 7 | 1 | 1 | 0 | 0 | 0 |
| EZB | 49 | 32 | 27 | 24 | 17 | 11 |

As primary CNS DLBCL
PCDLBCL leg type belong
to « MCD » subtype

MYD88 + CD79b

9p24.1 locus (PDL1/PDL2) in PCDLBCL leg type?

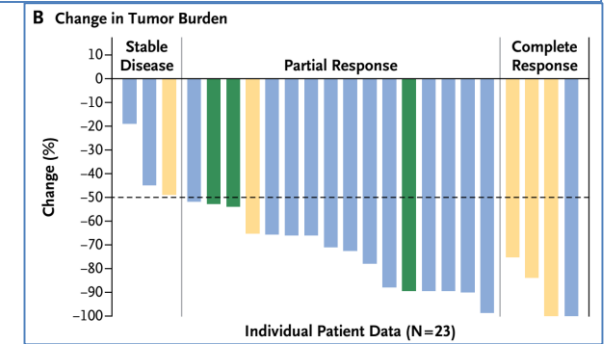


- Hodgkin lymphoma :

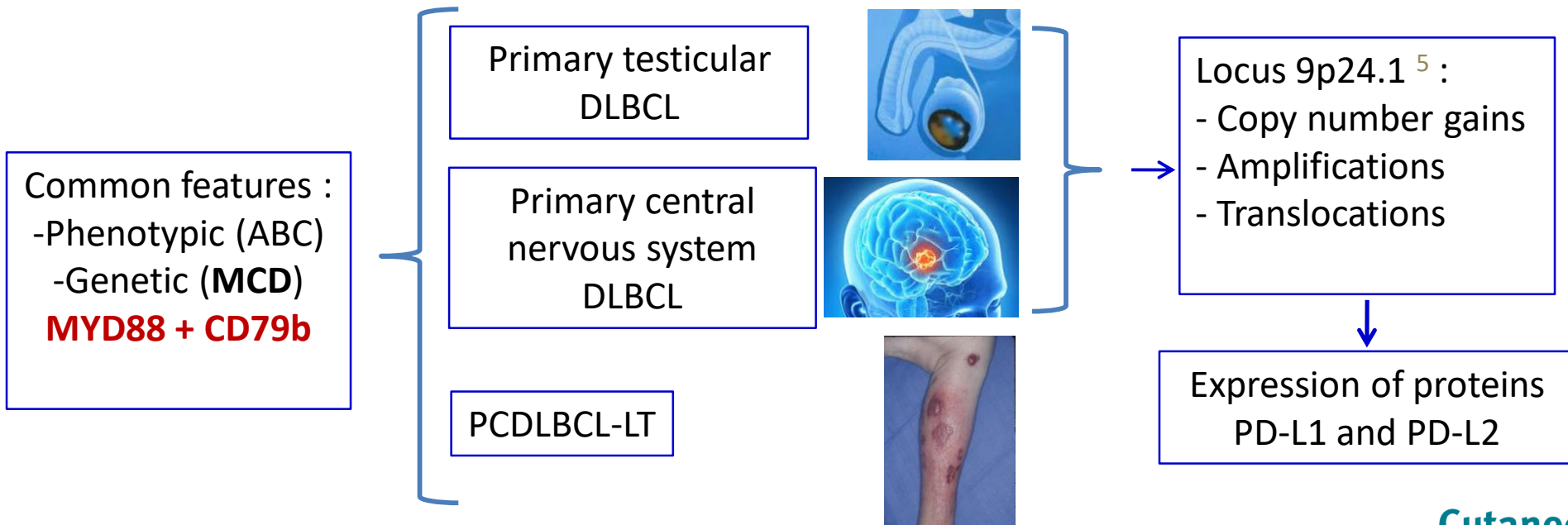
Amplification of 9p24.1 locus encoding for *PD-L1* and *PD-L2*

associated with protein expression → Efficacy of PD-1 inhibitors.⁴

PD-1 Blockade with Nivolumab in Relapsed or Refractory Hodgkin's Lymphoma



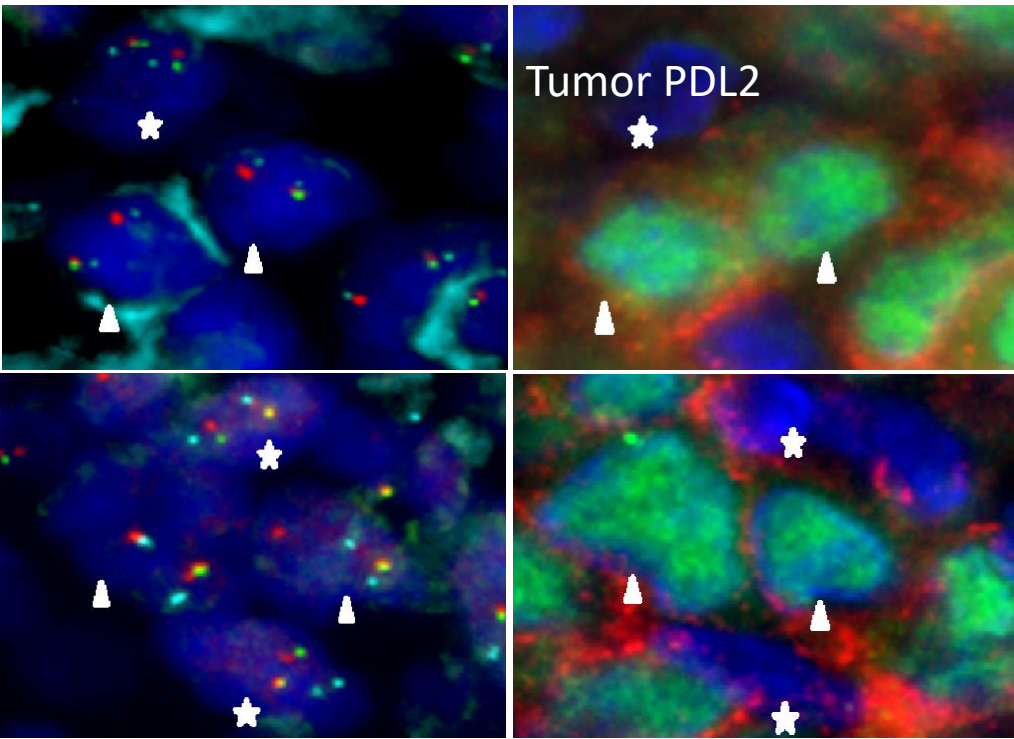
- Primary testicular and primary central nervous system DLBCLs :



4. Ansell, S. M. *et al.* PD-1 Blockade with Nivolumab in Relapsed or Refractory Hodgkin's Lymphoma. *N. Engl. J. Med.* **372** (2015) 25

5. Chapuy, B. *et al.* Targetable genetic features of primary testicular and primary central nervous system lymphomas. *Blood* **127**(2016).

Rares PDL1/PDL2 rearrangements but PDL1 expression by TAM

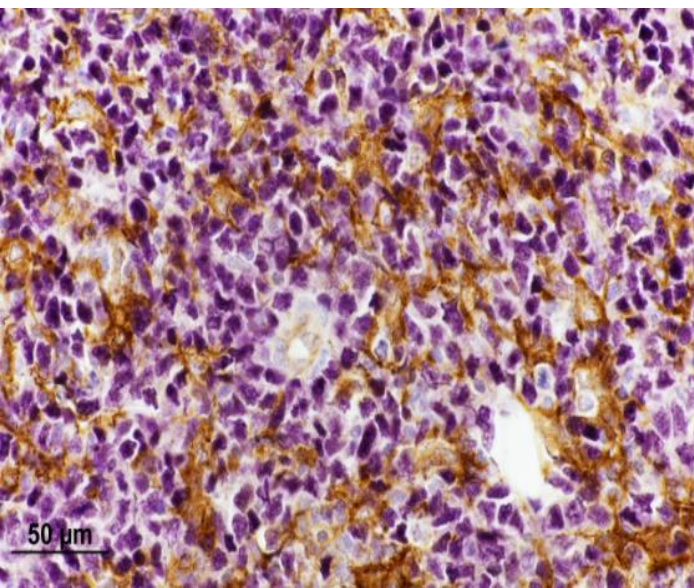


PD-L1 and PD-L2 are Differentially Expressed by Macrophages or Tumor Cells in Primary Cutaneous Diffuse Large B-Cell Lymphoma, Leg Type

Sarah Menguy, MD,† Martina Prochazkova-Carlotti, PhD,* Marie Beylot-Barry, MD, PhD,*‡
Frédéric Saltel, PhD,§ Béatrice Vergier, MD, PhD,*†
Jean-Philippe Merlio, MD, PhD,*|| and Anne Pham-Ledard, MD, PhD*‡*

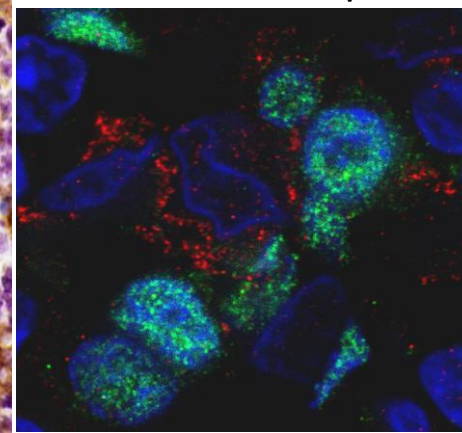
Am J Surg Pathol 2018

- **PDL1/2 rearrangements: 2/27 (7,4%)**
- → expression of PDL2 in these 2 only cases (tumor)
- **Microenvironnement expression of PDL1: 27/27** (TAM tumor associated macrophages)

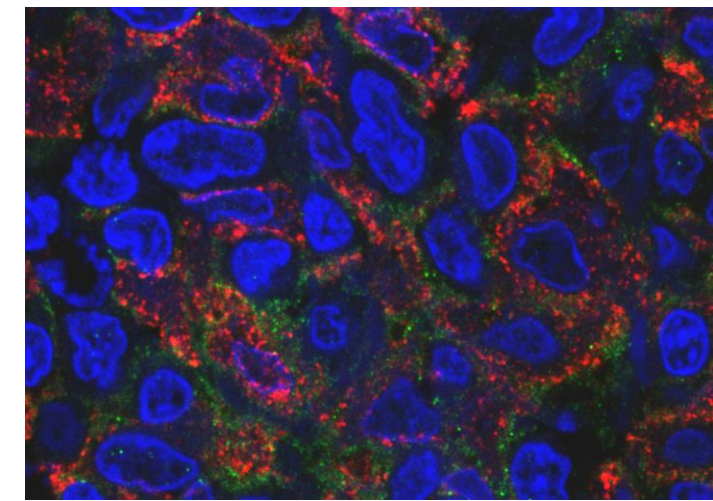


PDL2/PAX5

PDL1/PAX5



PDL1/CD163

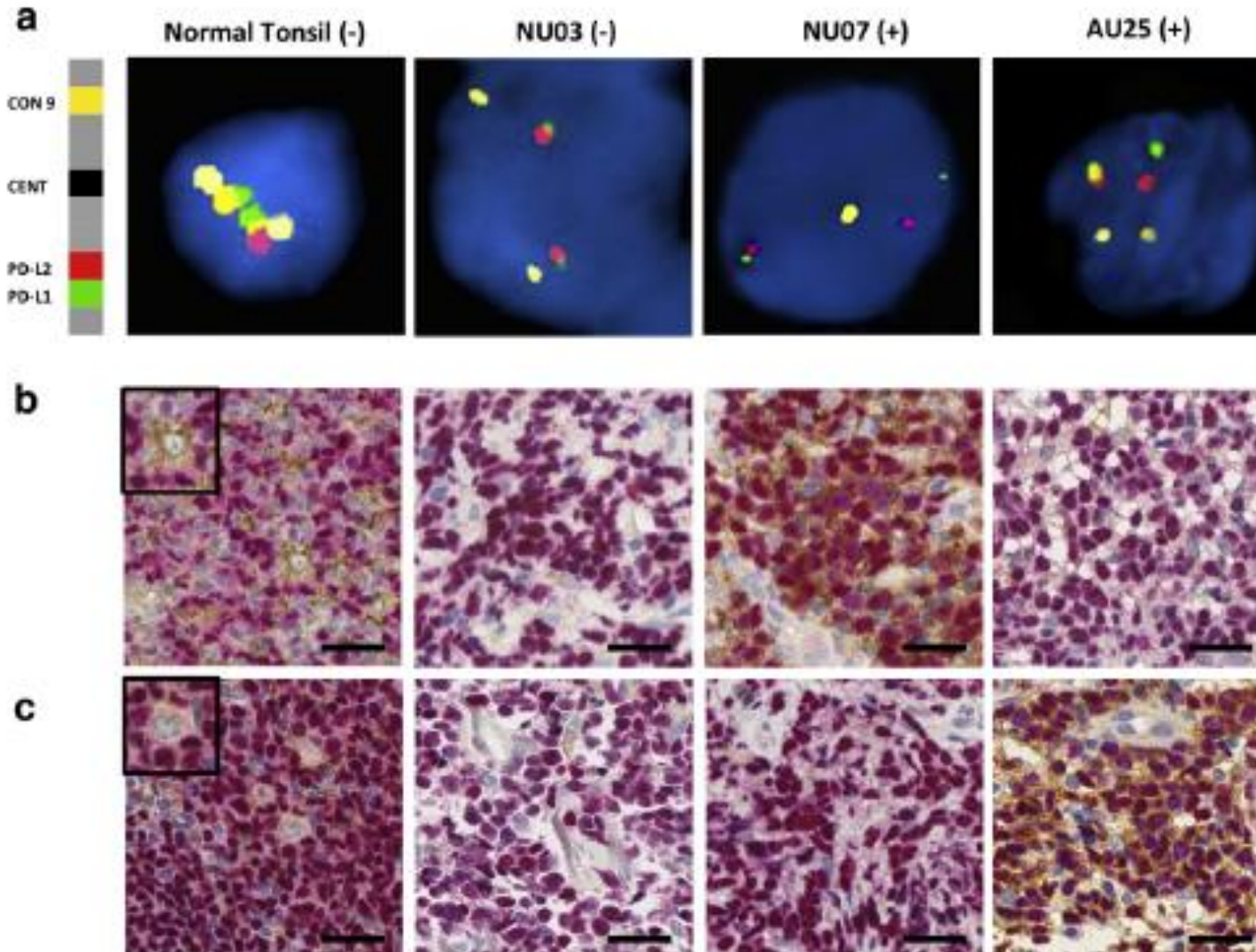


Some *PDL1/PDL2* rearrangements in this paper

Genomic Analyses Identify Recurrent Alterations in Immune Evasion Genes in Diffuse Large B-Cell Lymphoma, Leg Type

Xiaolong Alan Zhou^{1,12}, Abner Louissaint Jr.^{2,3,12}, Alexander Wenzel⁴, Jingyi Yang^{4,5}, Maria Estela Martinez-Escala¹, Andrea P. Moy^{2,3}, Elizabeth A. Morgan⁶, Christian N. Paxton⁷, Bo Hong⁸, Erica F. Andersen⁸, Joan Guitart¹, Amir Behdad⁹, Lorenzo Cerroni¹⁰, David M. Weinstock^{3,11} and Jaehyuk Choi^{1,4,5}

Zhou XA et al, *J Invest Dermatol* 2019



- 4/10 cases with *PDL1/ PDL2* break
- 50% with tumor (n=2) expression of *PDL1/2*

for rare or low prevalence complex diseases

Network
Hematological
Diseases (ERN EuroBloodNet)

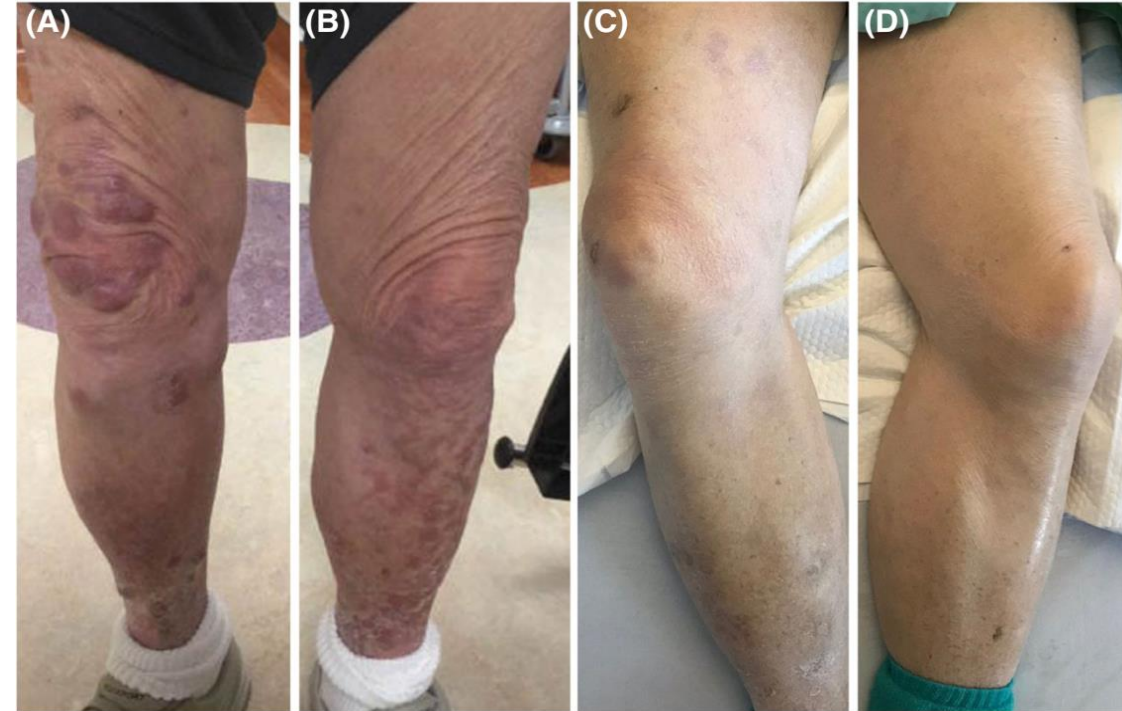
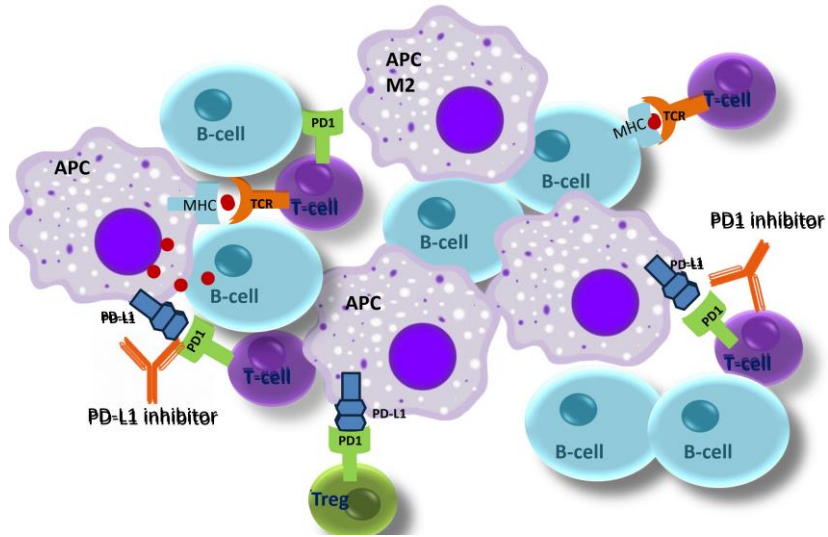
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Cutaneous Lymphoma

EuroBloodNet Topic on Focus

Checkpoint inhibitors in PCDLBCL Leg Type?

Rituximab, lenalidomide and pembrolizumab in refractory primary cutaneous diffuse large B-cell lymphoma, leg type

Di Ramondo et al, Br J Haematol, 2019

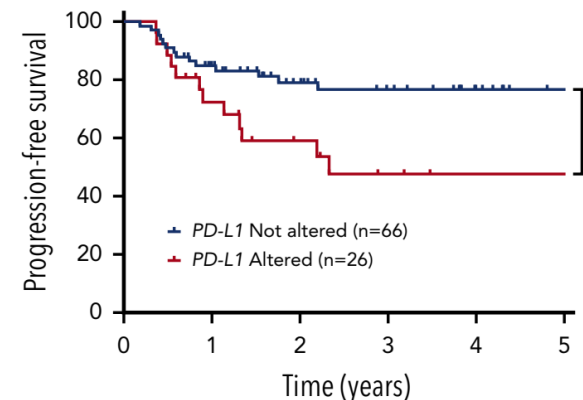
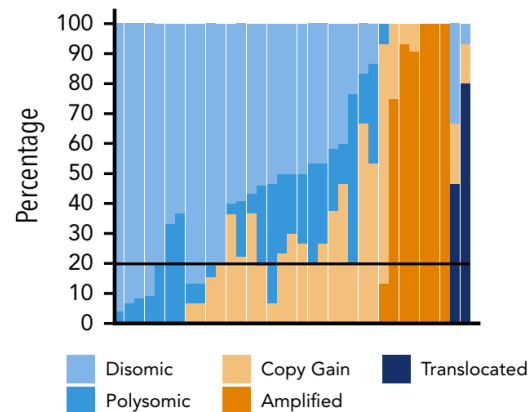


PD-L1 gene alterations identify a subset of diffuse large B-cell lymphoma harboring a T-cell-inflamed phenotype

Godfrey J et al, Blood, 2019

Checkpoint inhibitors:
Modest objective RR in DLBCL
(10-30%)

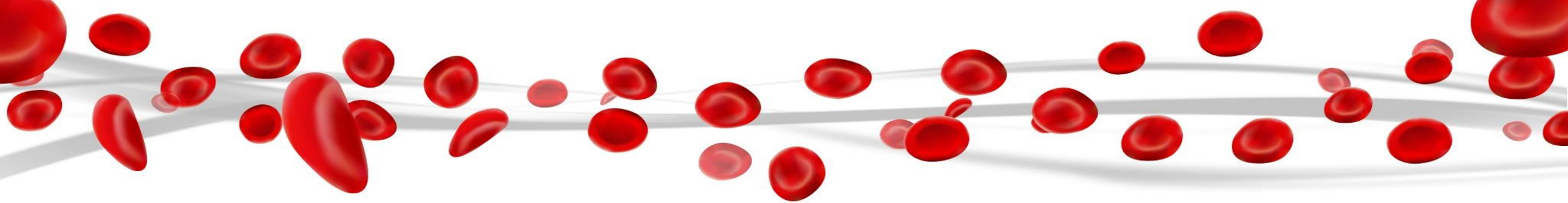
→ Select patients among PDL1/2 gene status?



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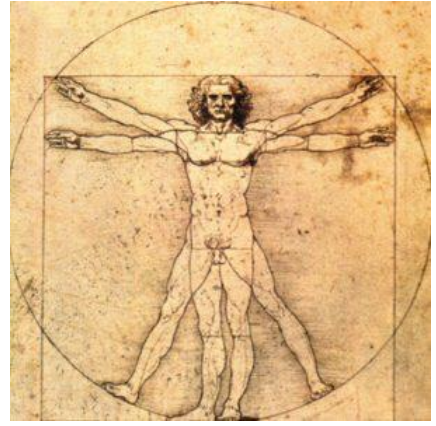
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1. PCDLBCL Leg type display typical clinical behavior and may be identified with **BCL2 +/ MUM1+ (> 50%) / CD10-/ FOXP1 + phenotype**
2. Advances in oncogenesis knowlegge : **MYD88 (70%), NFkB alterations and BCR pathway, Immune genes** are involved
3. Therapeutic perspectives have to be investigated: for who, when and how? **Lenalidomide, BTK inhibitors, NFkB targets, checkpoint inhibitors**



INSERM U1053 UMR BaRITON; Team 3 : Oncogenesis of cutaneous lymphoma

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Accueil | Aide

Se connecter | Mot de passe oublié

Actualité

- Accueil
- Actualité
- Autres événements

Objectifs & Fonctionnement

- Objectifs
- Fonctionnement
- Charte d'utilisation
- Statuts du GFELC

Recommandations

- Protocoles
- Engagement
- Bibliographie GFELC
- Information du Patient

Contact & Support

- Contact & Support

Le GFELC rassemble 31 équipes pluridisciplinaires (les « Centres ») constituées, chacune, d'un dermatologue et d'un pathologiste et, pour certaines, d'un biologiste moléculaire. Il constitue, officiellement, un Centre de Référence National pour les Lymphomes Cutanés, chaque centre étant considéré comme un Centre de Compétence. Il est constitué en Association de type Loi 1901. Le GFELC est un groupe thématique de recherche de la Société Française de Dermatologie.



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