

Linfomi della Zona Marginale Basi Biologiche

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Key aspects

Post-GC mature B-cells residing in the marginal zone

Subtypes defined by the anatomical site of origin (i.e., microenvironment)

- Extranodal (MALT): 70%
- Splenic: 20%
- Nodal: 10%

Median age ~60-65 years

Slight female predominance, especially in MALT lymphoma

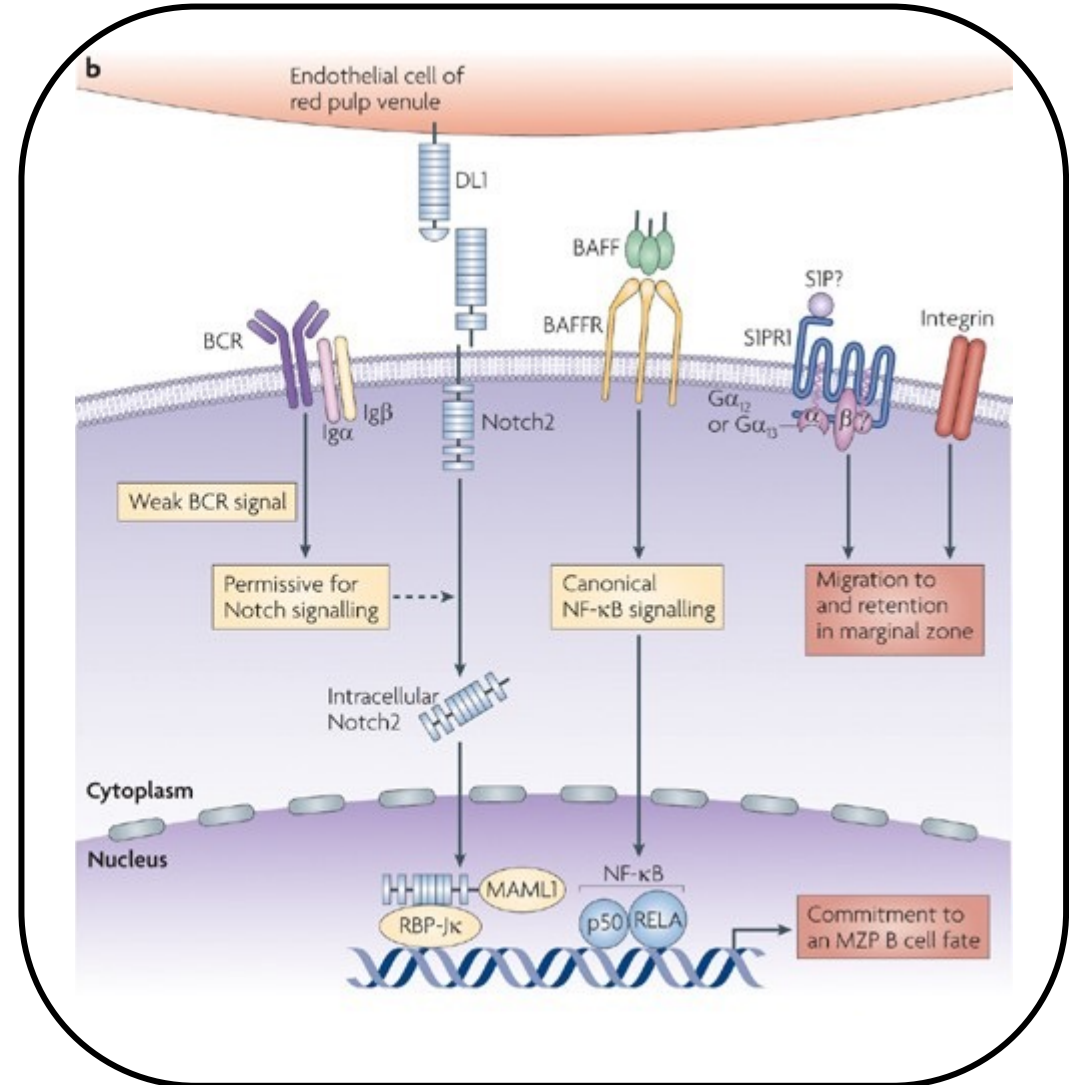
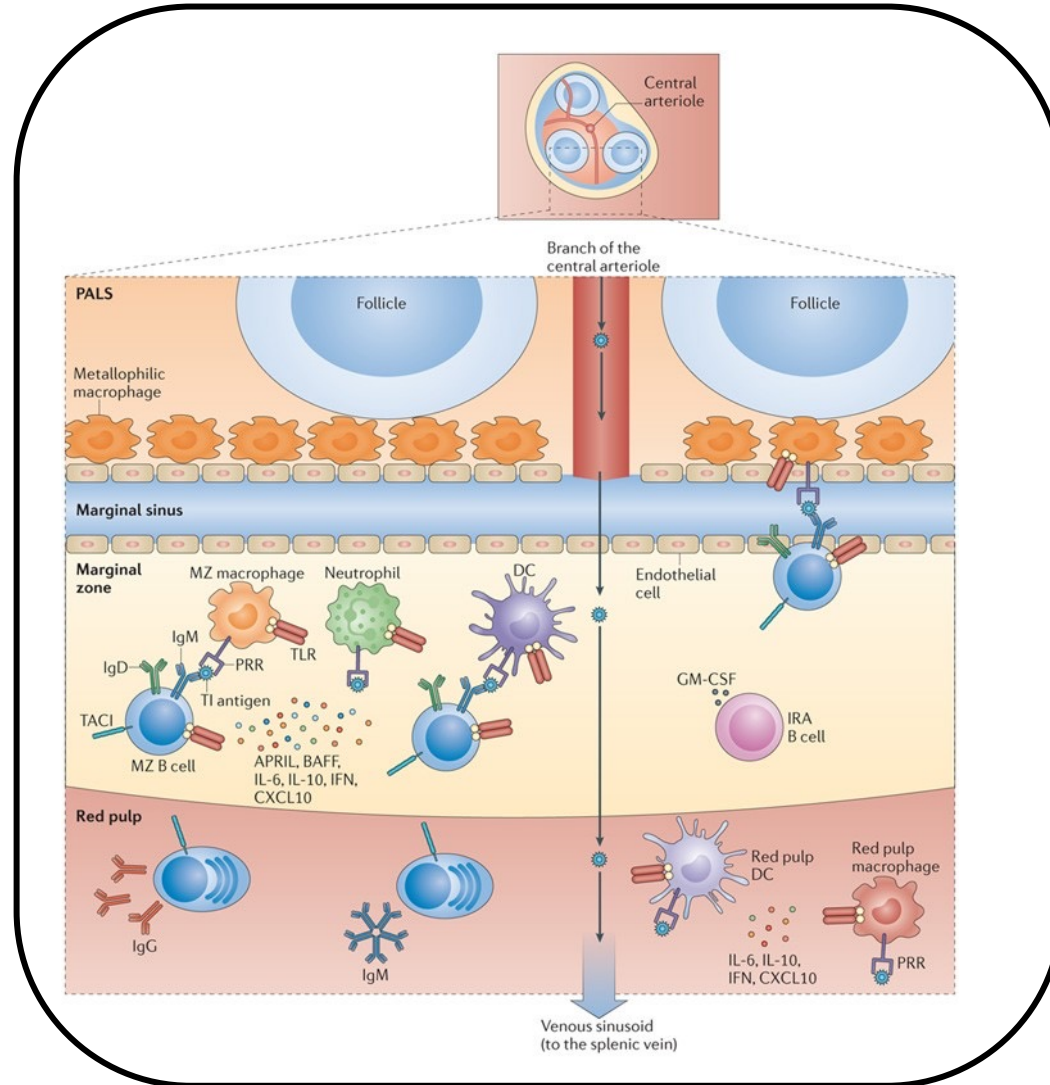
Risk factors

- Autoimmune diseases (Sjögren's syndrome, Hashimoto's thyroiditis)
- Chronic infections (*H. pylori*, *C. psittaci*, *B. burgdorferi*, *C. jejuni*, HCV)

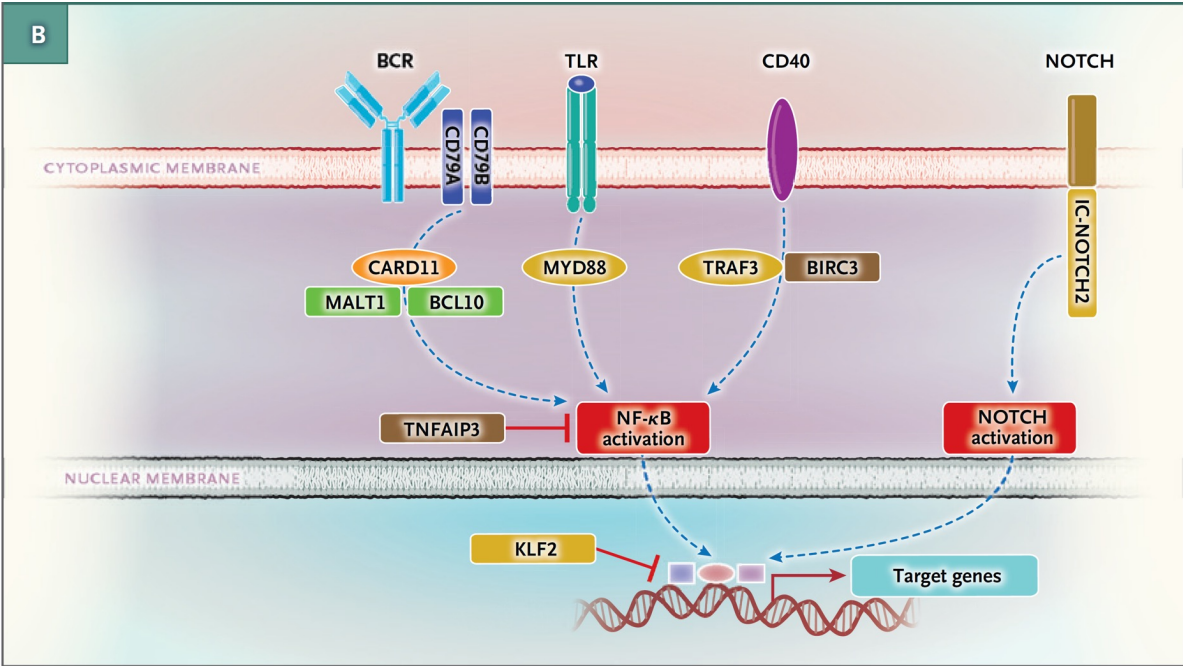
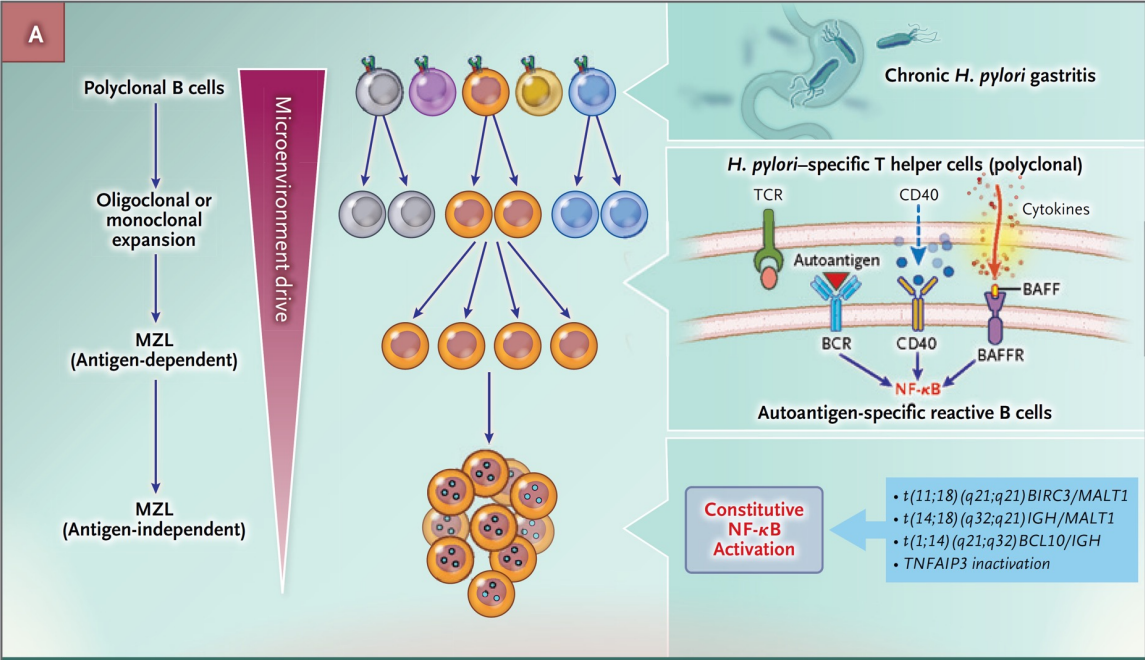
Indolent course but may progress and transform

Differential diagnosis may be challenging (nodal and splenic MZL)

Cell of origin of MZL

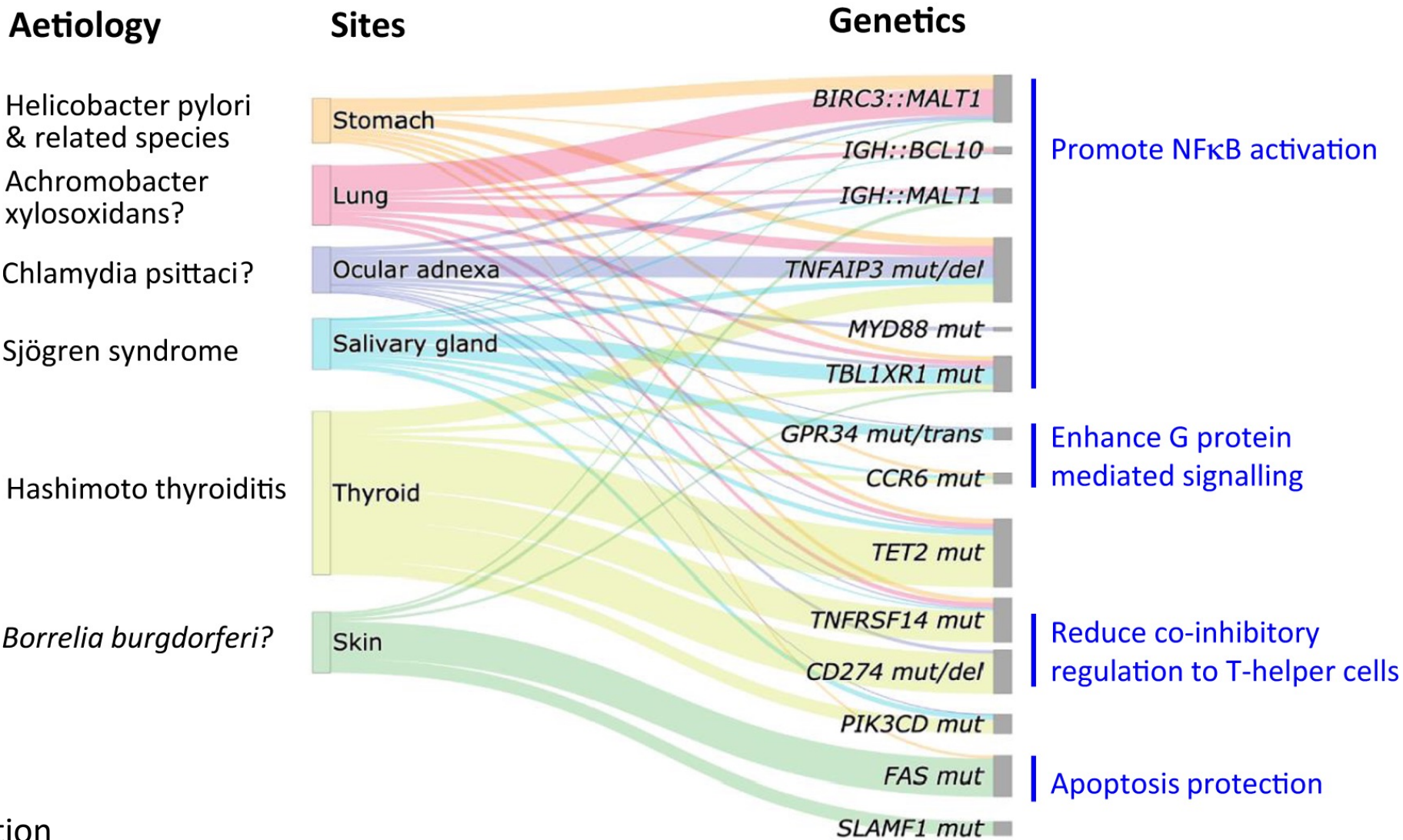


Pathogenesis of MZL

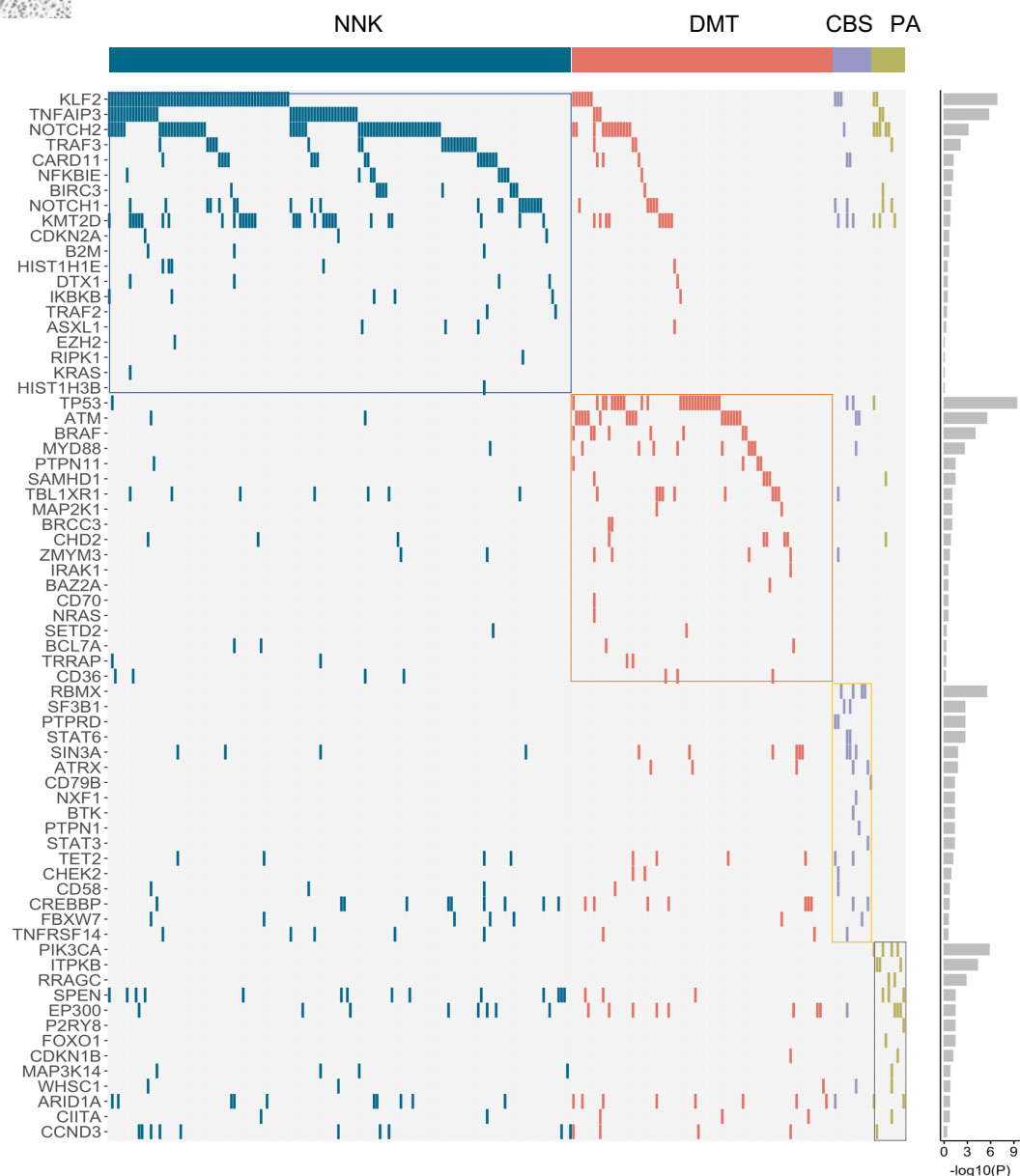


Site specific molecular heterogeneity of extranodal MZL

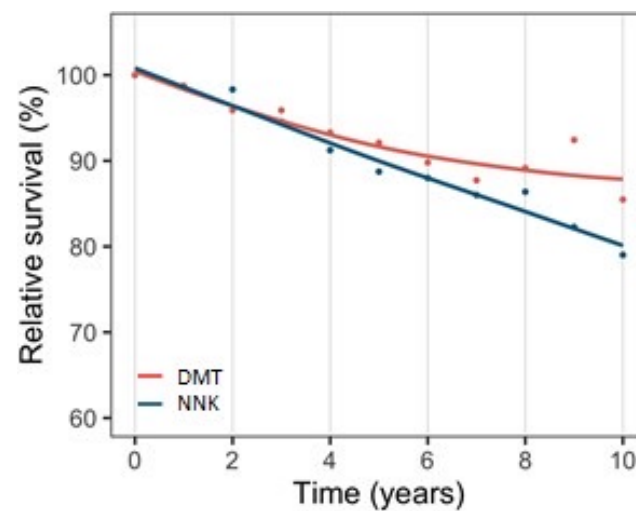
Extranodal marginal zone lymphoma



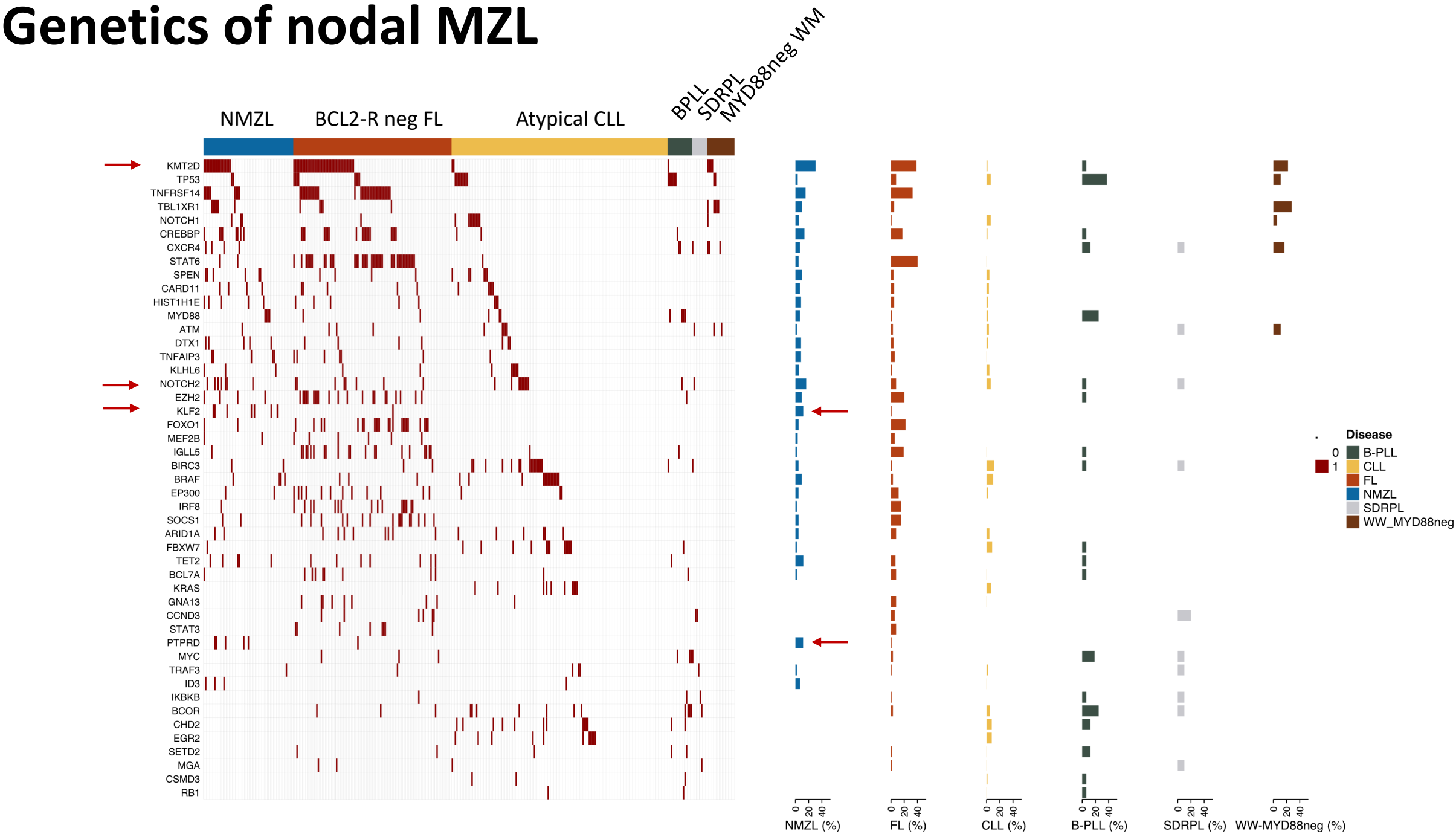
A Genetics of splenic MZL



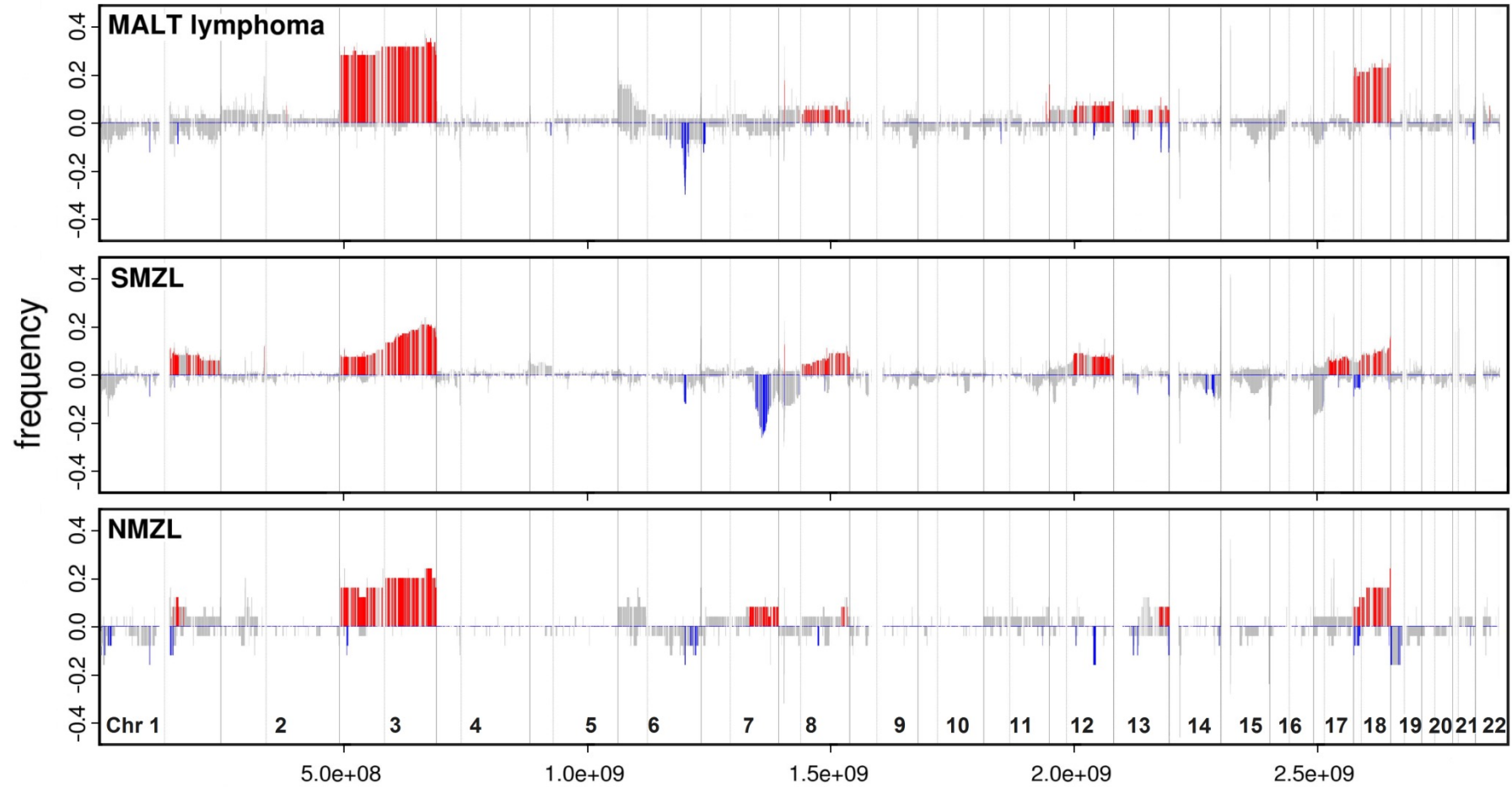
- Lack of *MYD88* L265P among SMZL confirmed on spleen histology
- NNK
 - 58% of cases
 - NF- κ B, NOTCH, and *KLF2* mutations
 - IGHV1-2*04 usage
 - Deletion 7q
- DMT
 - 32% of cases
 - DNA-damage response, MAPK, and TLR mutations



Genetics of nodal MZL



Cytogenetics of MZL



Implications for diagnosis

Small B-cell lymphomas without defining features

- **Marginal zone lymphomas**
- Follicular lymphoma **without** *IGH::BCL2*
- **Atypical** chronic lymphocytic leukemia
- Lymphoplasmacytic lymphoma **lacking** *MYD88* L265P
- Mantle cell lymphoma **lacking** *IGH::CCND1*
- Splenic B-cell lymphoma/leukemia unclassifiable
- (B-cell prolymphocytic leukemia)

Small B-cell neoplasms with unexpected phenotypes

MZL phenotype CD5-/CD23-/BCL6-/CD10- (MZL? LPL?)

Unexpected phenotypes

- CD5-/CD23+ (MZL? FL?)
- CD5+/CD23-/cyclin D1- (MZL? MCL?)
- CD5+/CD23+, but bright CD20+/slg+, CD79b+, FMC7+ (MZL? CLL?)

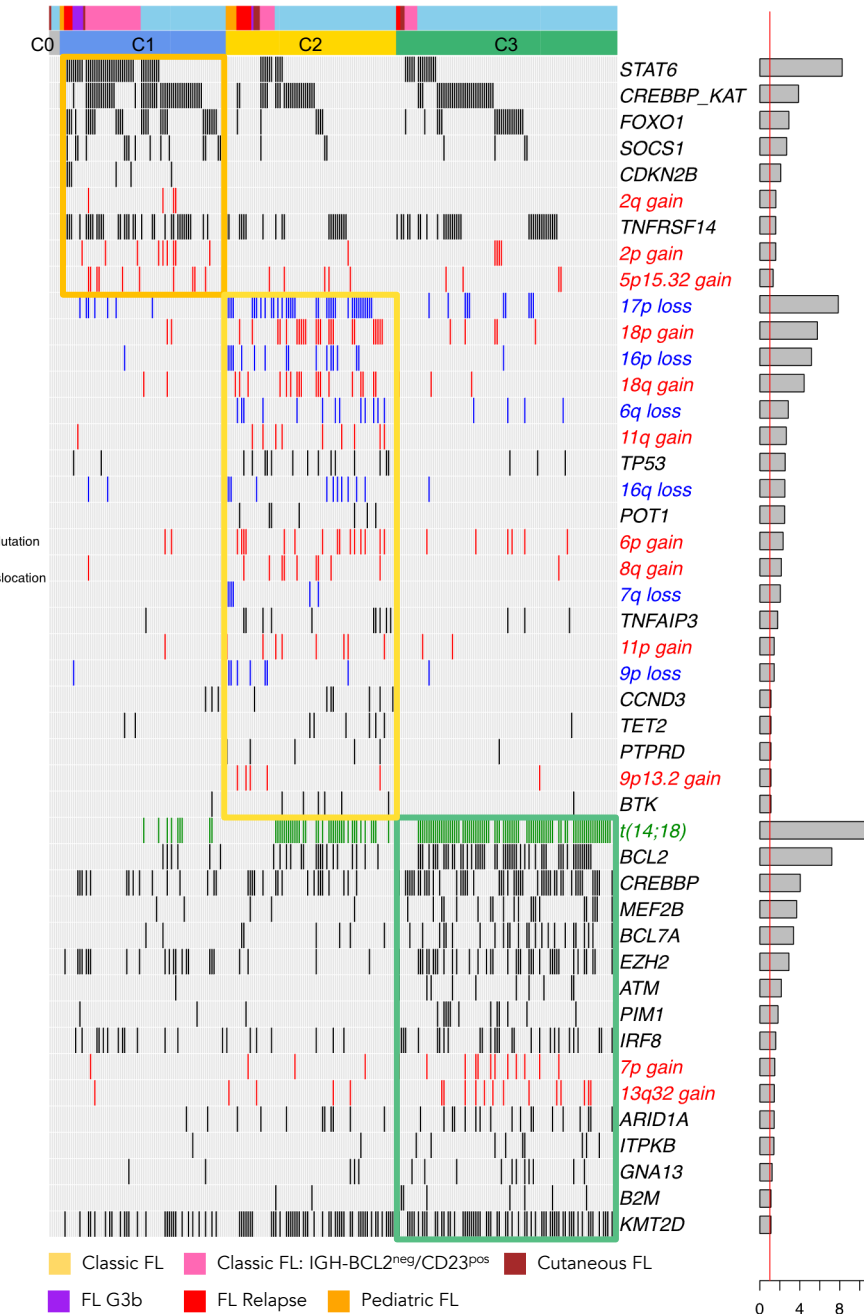
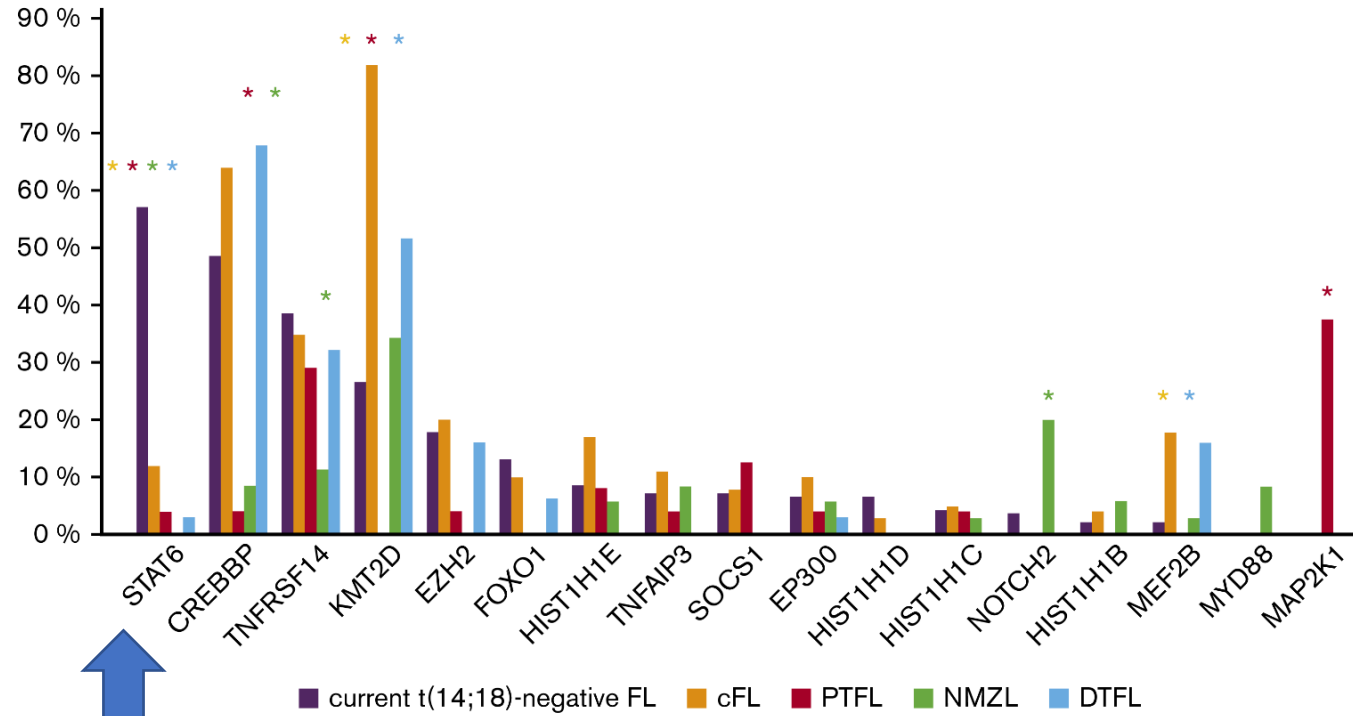
Dissemination of occult EMZL in 20% of NMZL presenting in limited stage

Wrong diagnosis in 20-30% of cases

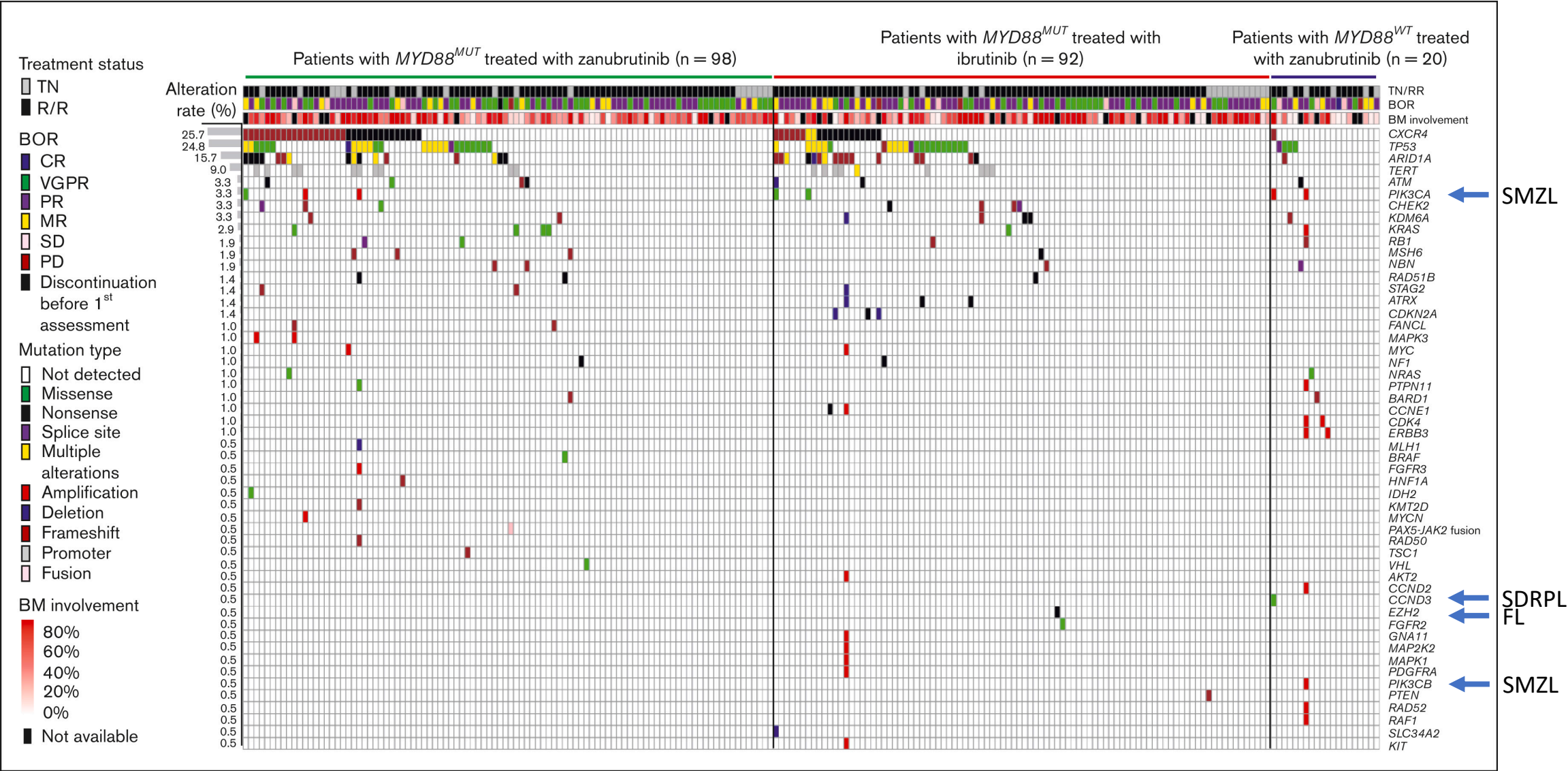
Diagnosis is challenging: wrong diagnosis affects outcome

Follicular lymphoma (without *IGH::BCL2*)

N=45

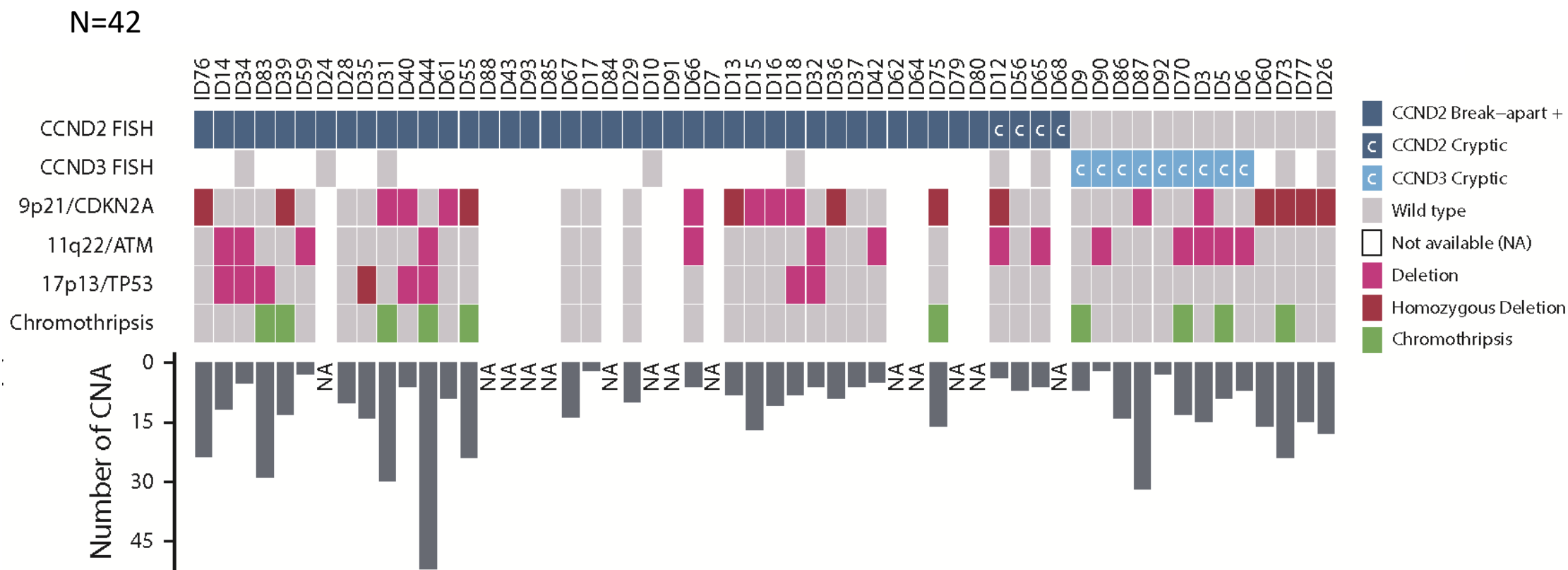


MYD88 wild type LPL



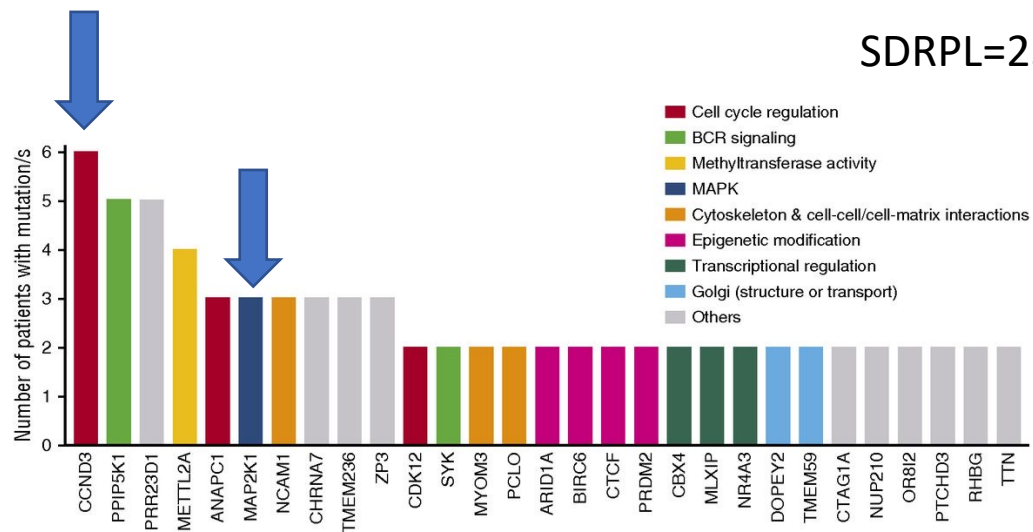
Does it really exist? No genetic biomarkers

Genetics of Cyclin D1 negative MCL



CCND2 and *CCND3* translocation are the best biomarkers

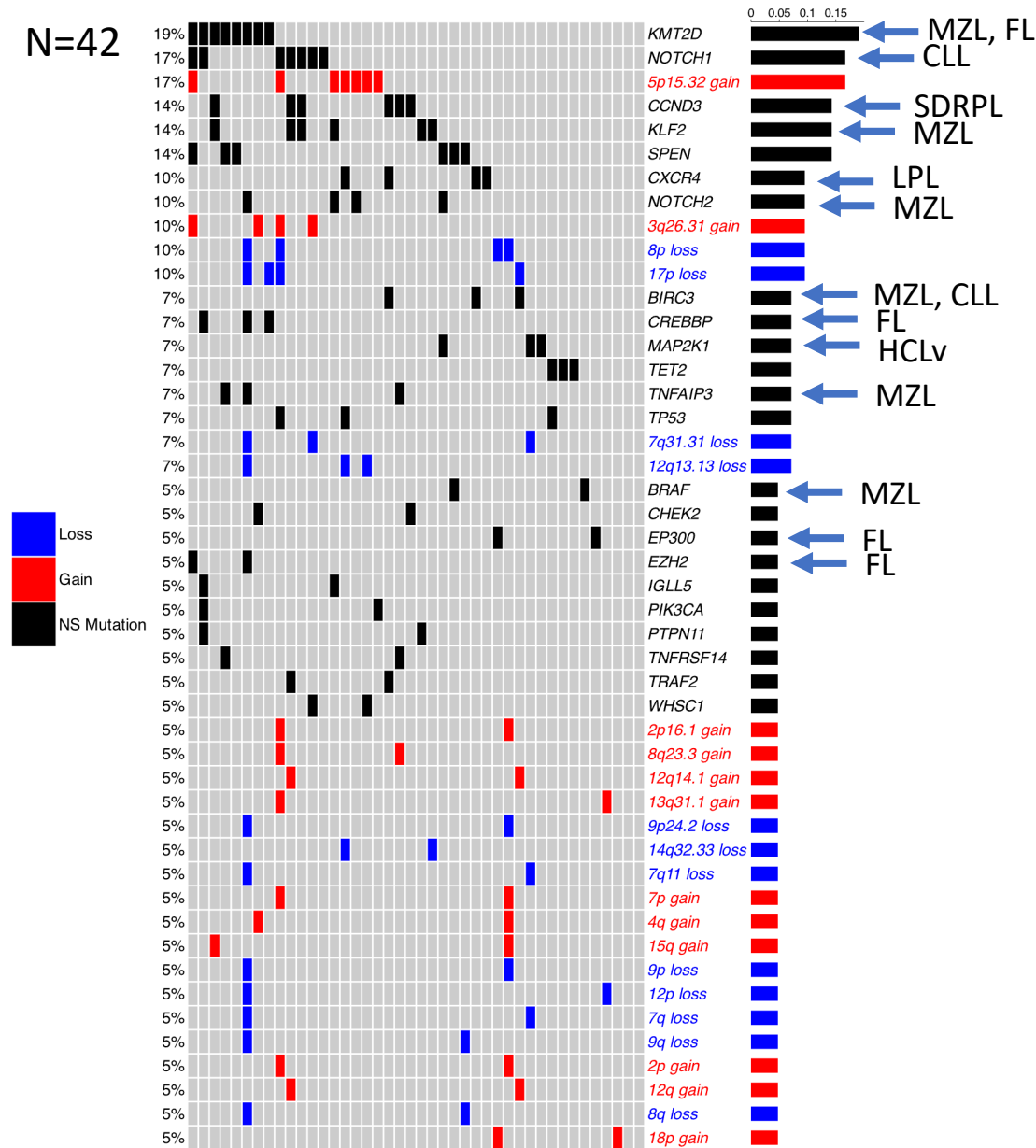
Splenic B-cell lymphomas unclassifiable



Curiel-Olmo S, et al. Blood. 2017

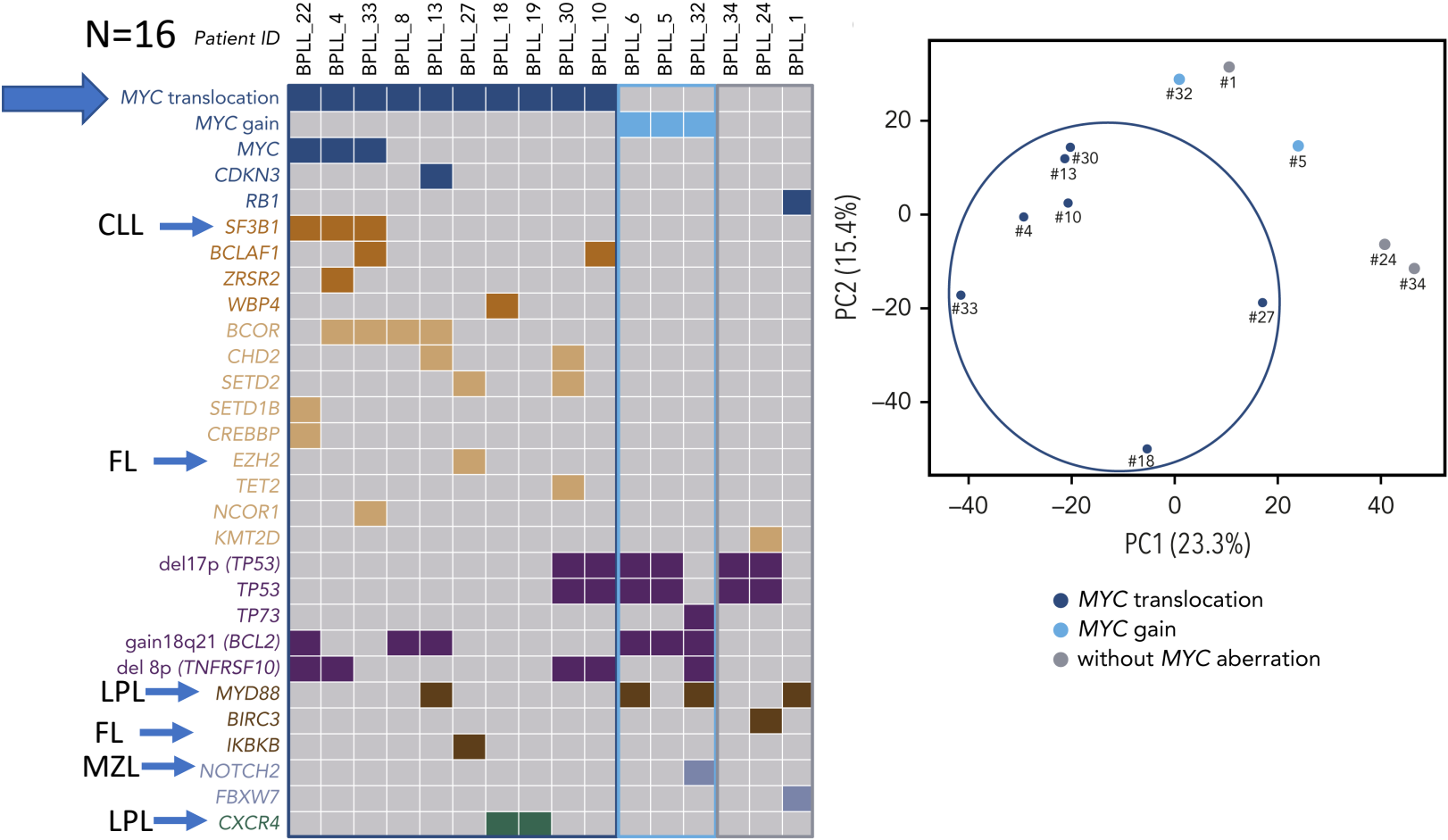
MAP2K1 mutations in 50% of HCLv (n=7)

Waterfall JJ, et al. Nat Gent. 2014



Unpublished

Genetics of B-PLL



MYC translocation (mostly with IG) is a biomarker of B-PLL

Extended diagnostics

- Marginal zone lymphomas → **+3, +12, +18, -7, *KLF2* m, *NOTCH2* m**
- Dissemination of an occult MALT MZL → **extensive imaging**
- Follicular lymphoma without *IGH::BCL2* → **GCB IHC marker, *STAT6* m**
- Chronic lymphocytic leukemia → **+12, -13q, *SF3B1* m, *NOTCH1* m**
- Lymphoplasmacytic lymphoma → ***MYD88* L265P**
- Mantle cell lymphoma lacking *IGH::CCND1* → **IHC *SOX11***
- Splenic B-cell lymphoma/leukemia unclassifiable → ***MAP2K1* m, *CCND3* m**

Clinico-pathologic variants

MZL non-CLL type/CBL-MZ

- Isolated lymphocytosis
- Matutes score <3
- No clinical evidence of MZL
- Morphologic heterogeneity
- Mixed patterns of BM infiltration
- Del(7q), *NOTCH*, NF-kB, KLF2, MYD88, CCND3, MAP2K1

Xochelli A, Blood 2014

Bruscaggin A, Br J Haematol. 2014

Kalpadakis C, Hematol Oncol. 2017

Defrancesco I, Hematol Oncol. 2020

Parker H, Blood Adv. 2018

SMZL with nodal dissemination

- Extra hilar lymph nodes: 20-30%
- Risk factor of the HPLL and MZL-IPI

Arcaini L, EClinicalMedicine. 2024
Bonfiglio F, Blood. 2022
Iannitto E, Br J Haematol. 2018
Arcaini L, Blood. 2006
Montalbán C, Abraira V, Arcaini L, Domingo-Domenech E, Guisado-Vasco P, Iannitto E, Br J Haematol. 2012
Berger F, Blood. 2000

Genetic lesion	Enrichment	P(Adj)
del7q31.33	1.13	0.9
KLF2	0.97	0.9
NOTCH2	1.20	0.9
amp12q24.13	1.38	0.9
del1p36.32	0.28	0.9
del11q12.3	0.55	0.9
KMT2D	1.65	0.9
TNFAIP3	1.37	0.9
amp22q12.1.	1.04	0.9
NOTCH1	0.52	0.9
CREBBP	0.44	0.9
CARD11	1.83	0.9
amp15q26.1.	0.48	0.9
SPEN	1.53	0.9
TP53	0.35	0.9
ARID1A	0.58	0.9
ATM	0.38	0.9
TBL1XR1	1.64	0.9
TRAF3	1.64	0.9
CCND3	0.82	0.9
EP300	1.51	0.9
TET2	0.26	0.9
MYD88	1.11	0.9
SIN3A	1.11	0.9

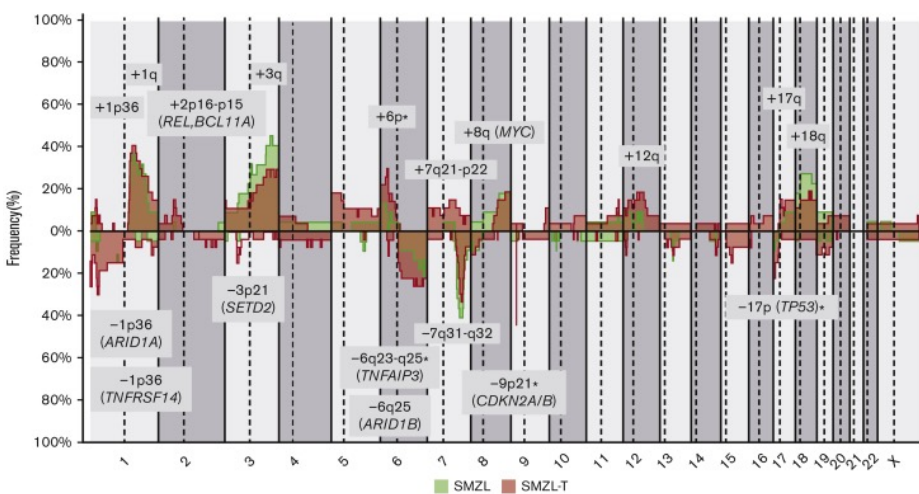
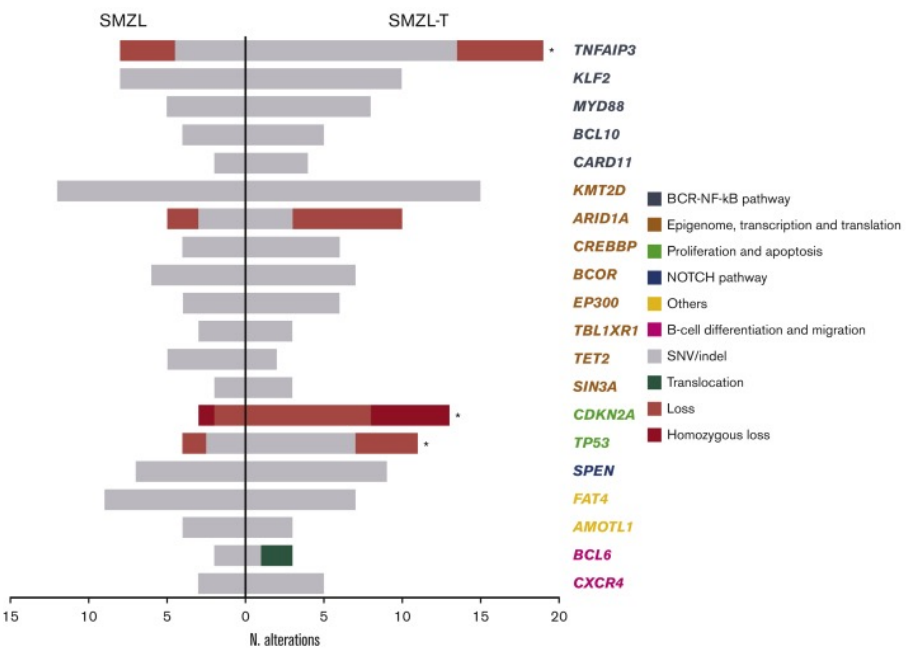
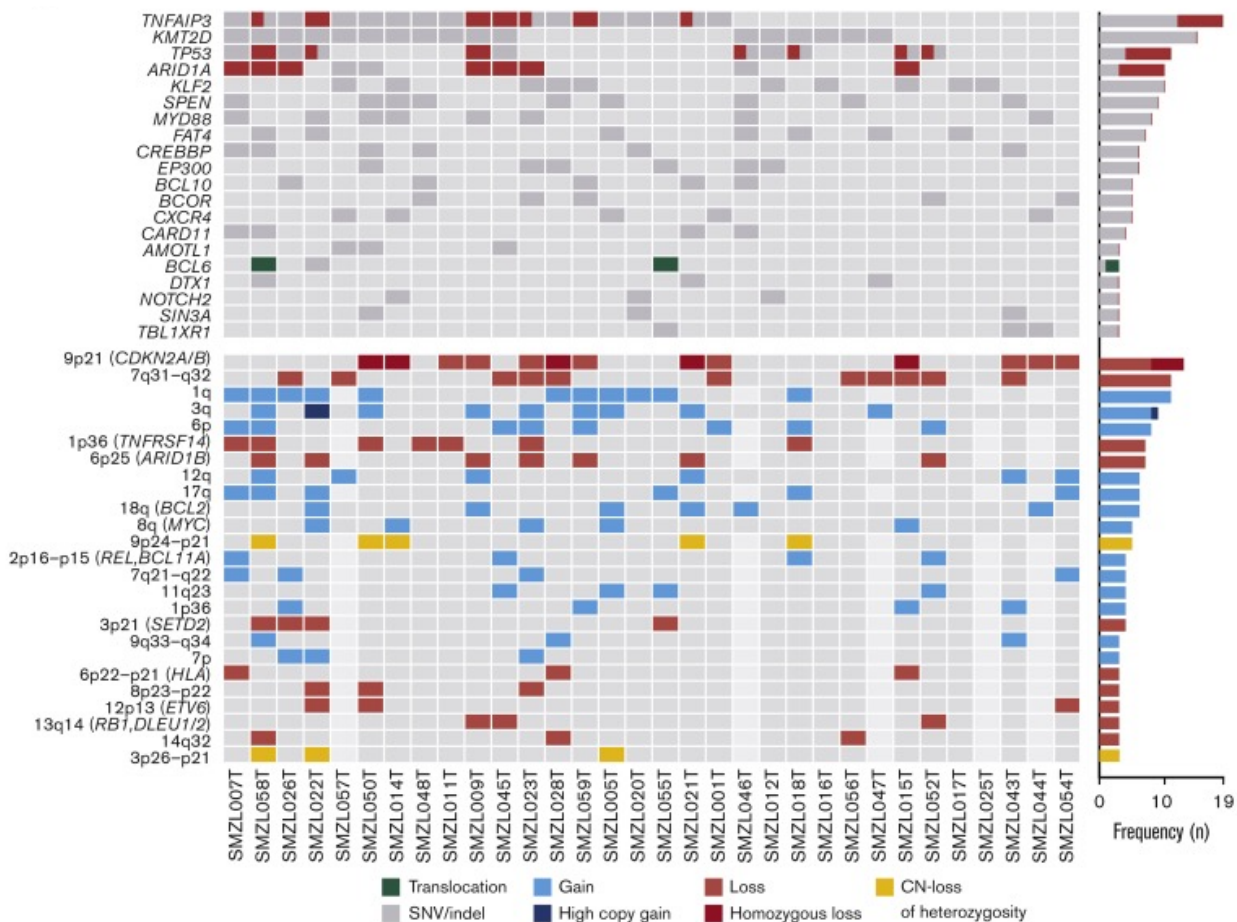
MZL with aggressive histologic presentation

- Increased number of scattered large cells
 - Definition: large cells >10 per HPF
 - Ki-67 expression $\geq 30\%$
- Prevalence: 20% (more common in NMZL)
- Presentation with advanced stage and increased LDH
- Shorter PFS, increased rate of relapse, increased rate of transformation

Histologic transformation

- 10-years cumulative incidence 8% (EMZL=3-5%, SMZL=7-13%, NMZL=9-13%)
- 5-years OS from transformation 55%
- Not well defined pathologic criteria (sheets of large cells): DD with MZL with increased scattered large cells

Genetics of transformed SMZL





IELSG52

Integrated molecular and clinical profiling to improve disease characterization and outcome prediction in nodal marginal zone lymphoma



IELSG54

Integrated Molecular and Clinical Profiling of Transformed Splenic Marginal Zone Lymphoma