



Marginal zone lymphomas: biology and molecular genomics

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Hematology

IOSI - Oncology Institute of Southern Switzerland

IOR - Institute of Oncology Research

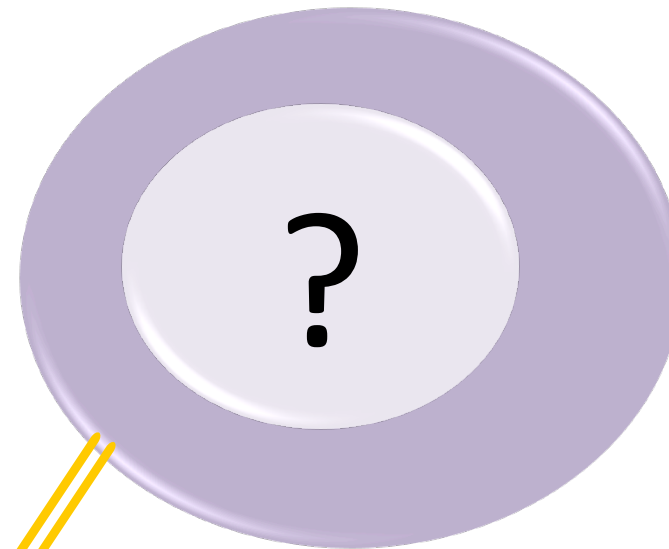
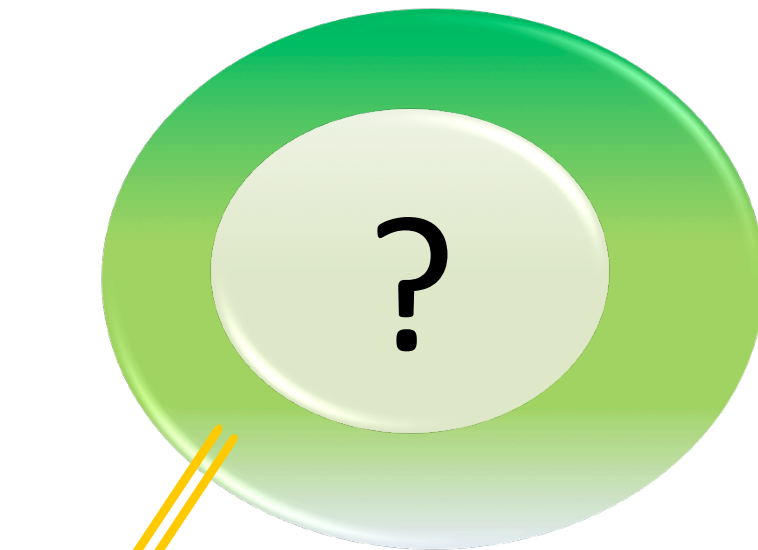
Bellinzona

- **Genetics of non-MALT MZL**
 - **SMZL**
 - **NMZL**

- **Clinical implications of non-MALT MZL genetics**
 - **Diagnosis**
 - **Prognosis**
 - **Therapy**

SMZL

NMZL

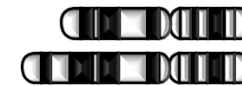


IGHV1-2*04
30%



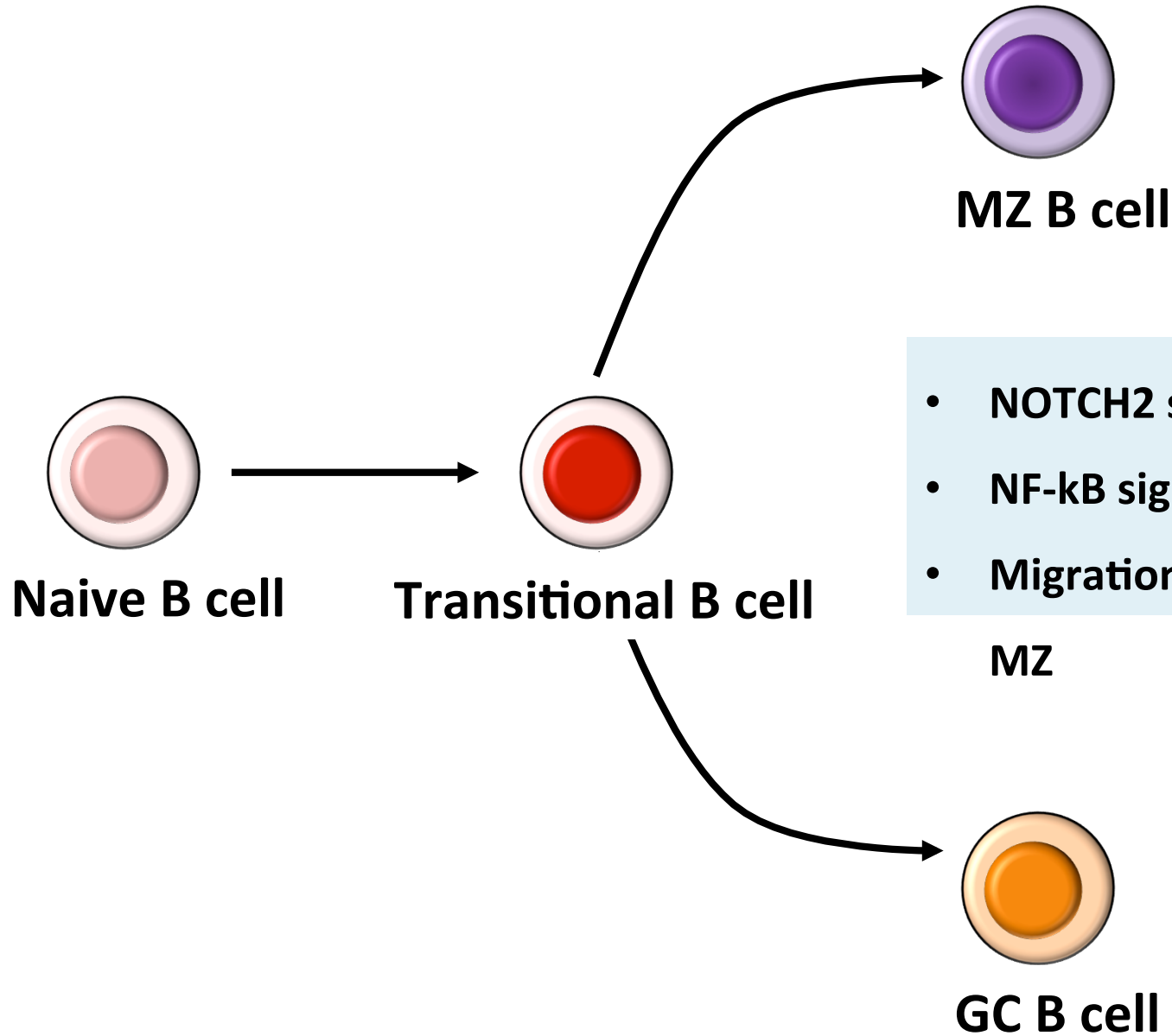
7q deletion: 30%

IGHV4-34
30%



+3: 14%
+12: 13%
+18: 10%

Pathways in the SMZL and NMZL signatures

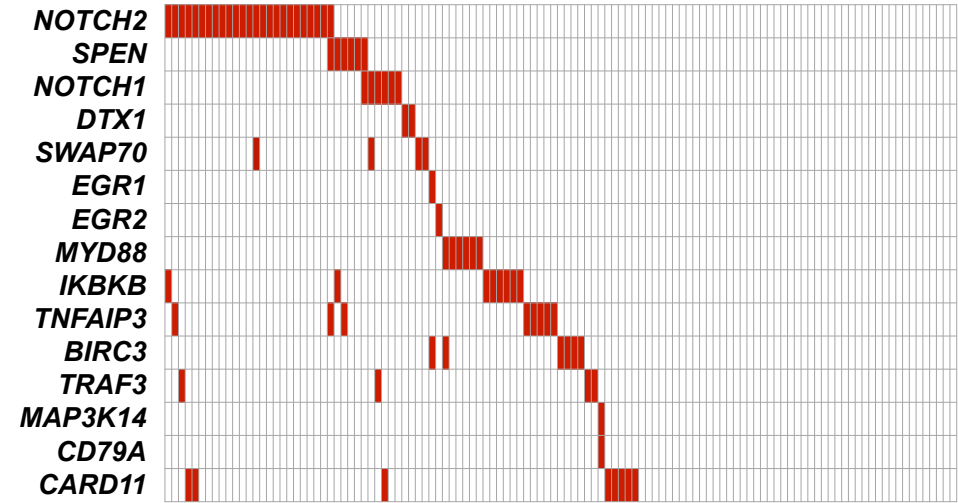
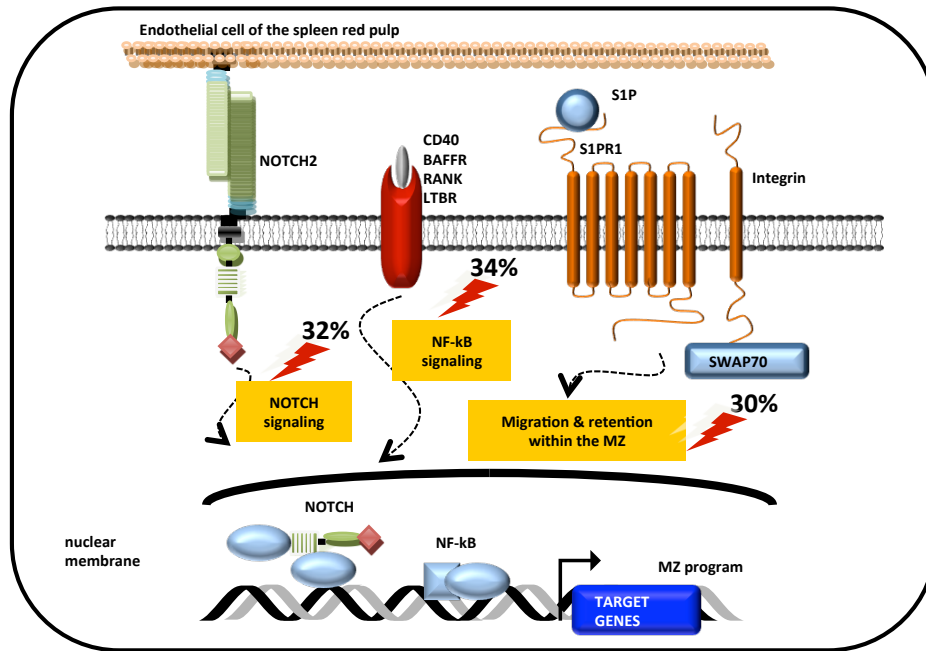


- **NOTCH2 signals**
- **NF-kB signals**
- **Migration to and retention in**

MZ

Trøen G, et al. J Mol Diag 2004
Ruiz-Ballesteros E, et al. Blood 2005

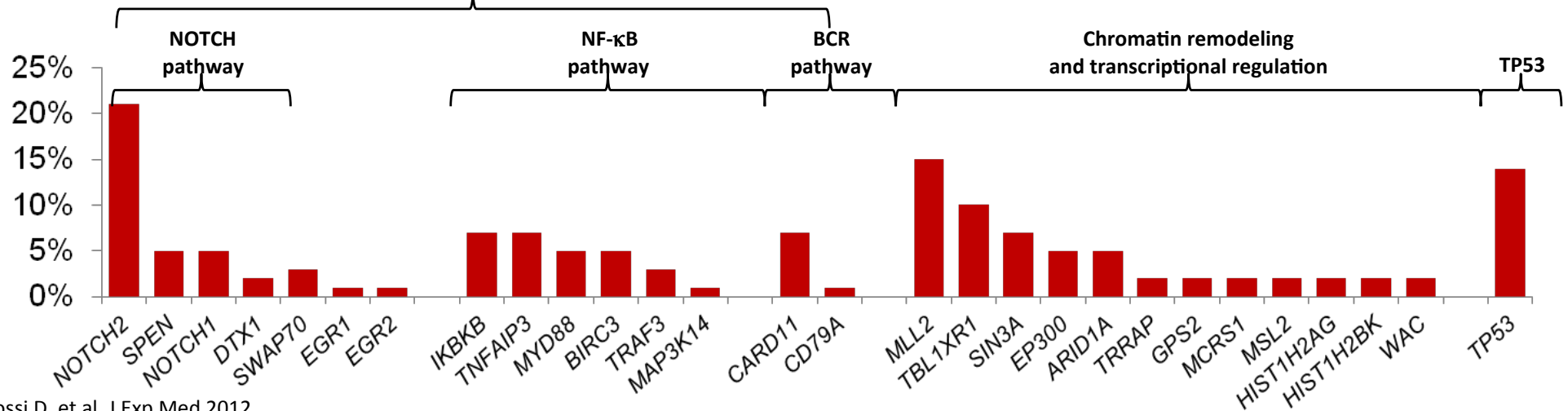
The coding genome of SMZL



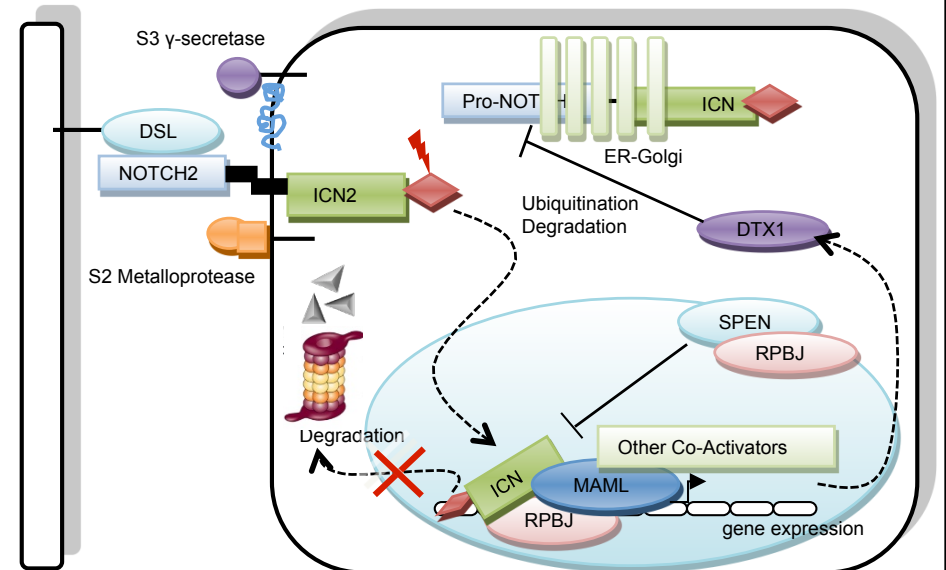
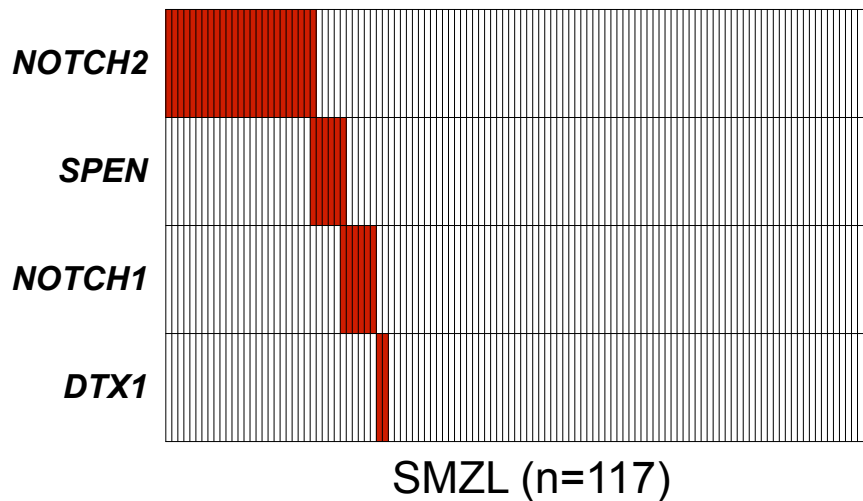
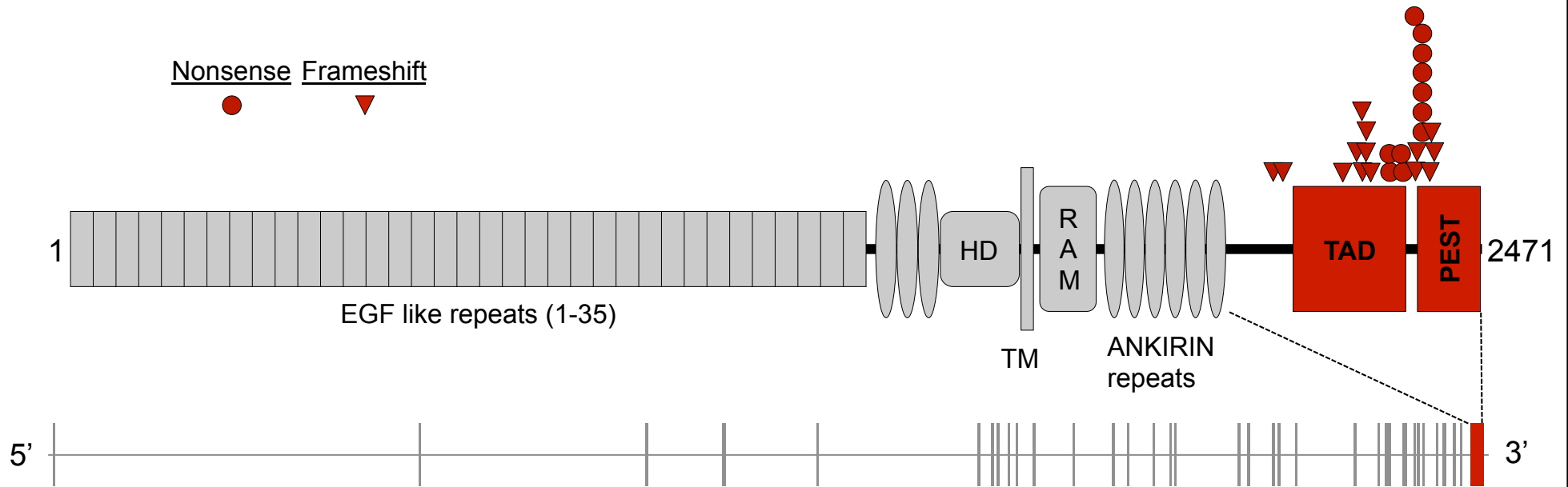
SMZL (n=117)

~60% of patients

Genes involved in marginal zone development

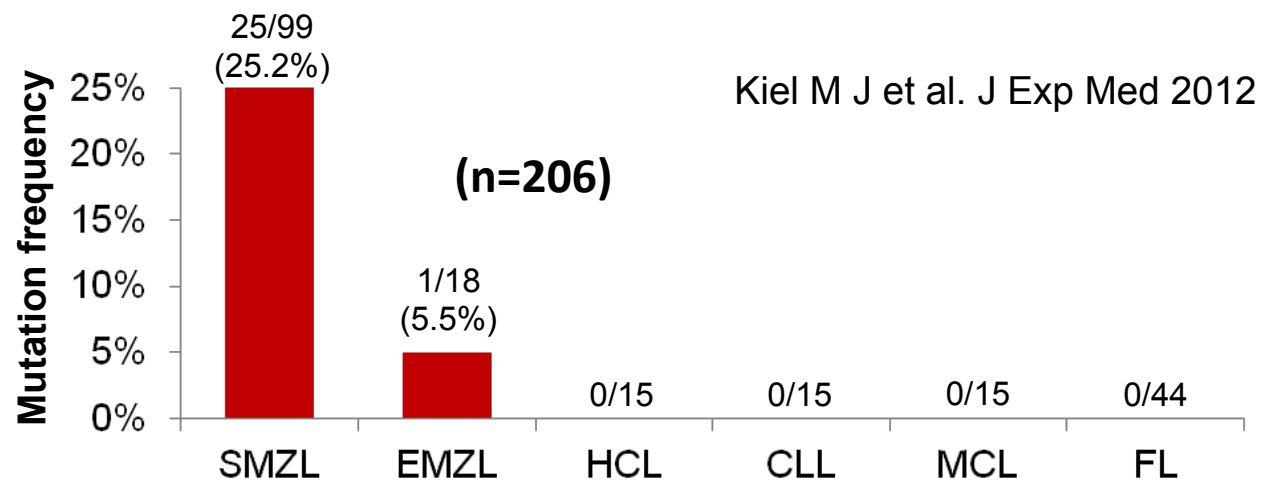
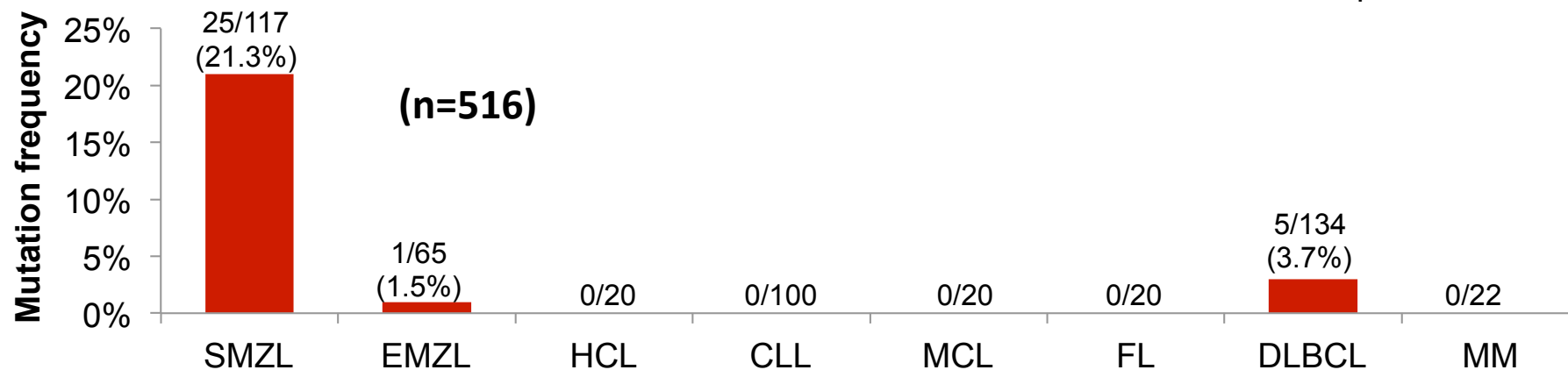


NOTCH2 is the most frequently mutated gene in ~20% SMZL

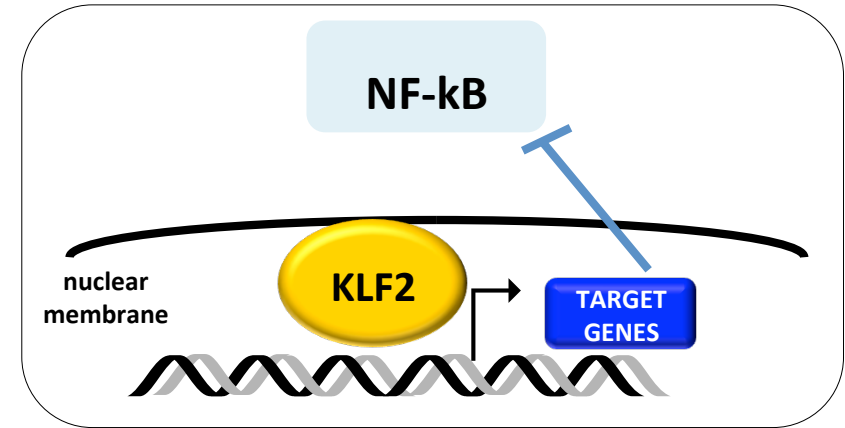
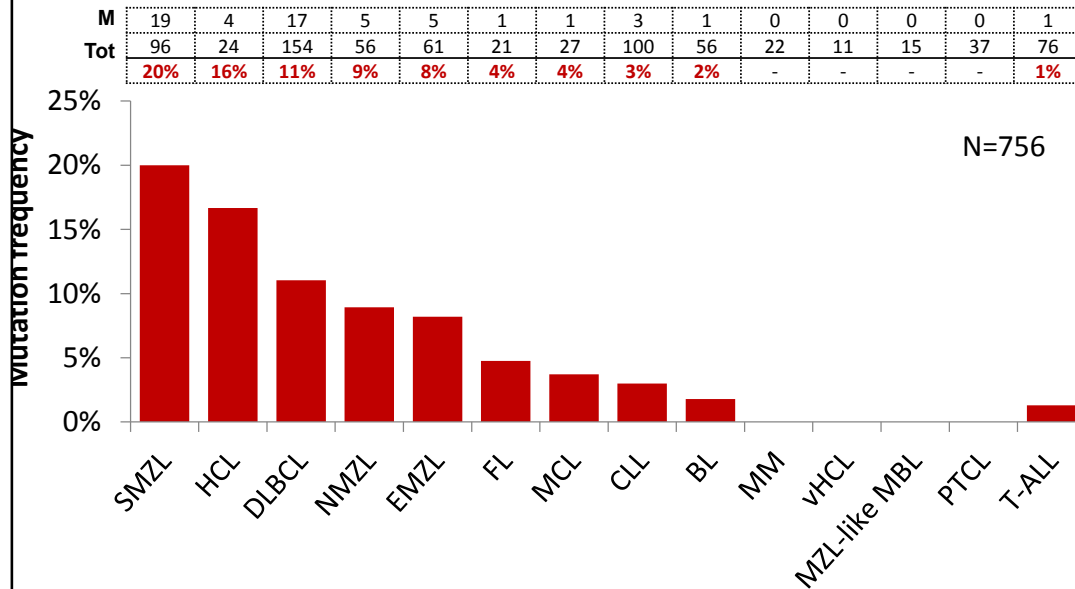


NOTCH2 mutations are restricted to SMZL across mature B-cell tumors

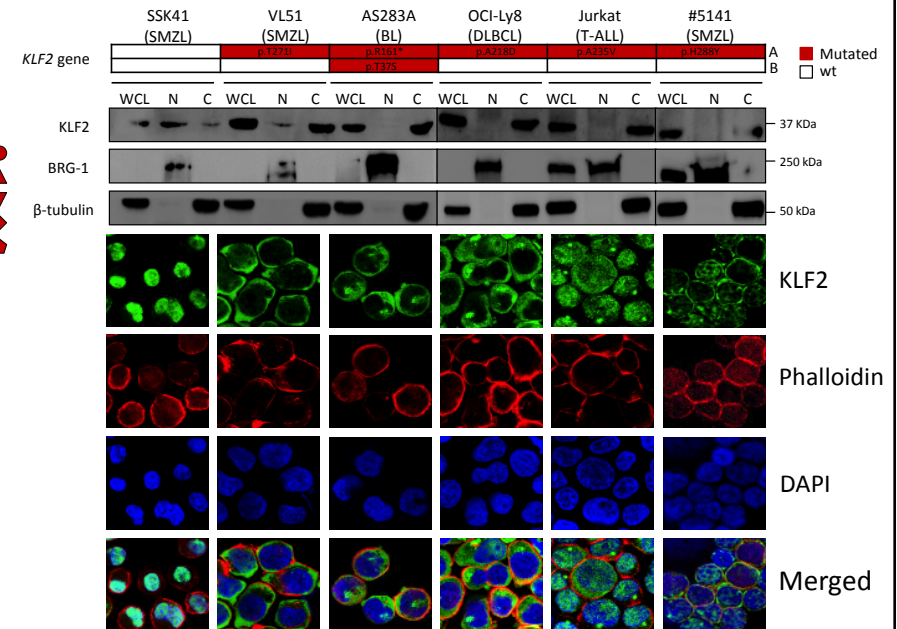
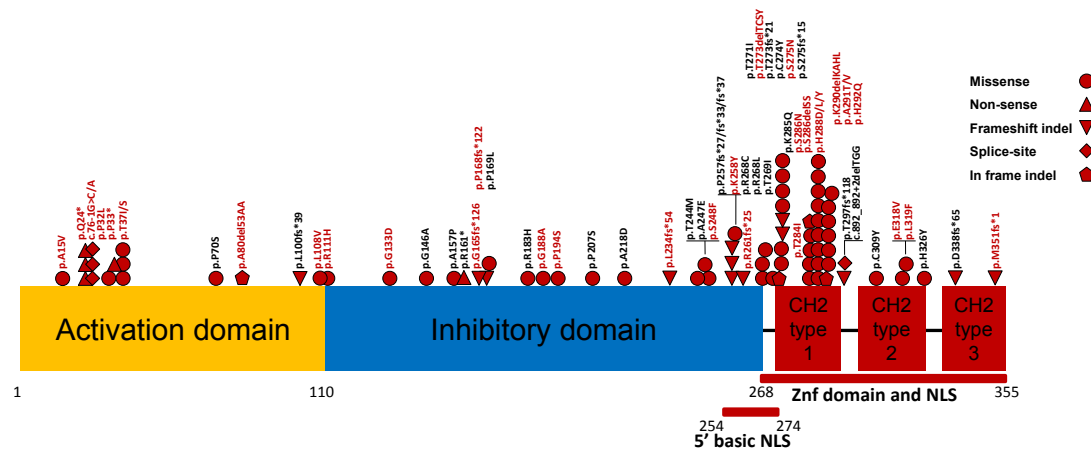
Rossi D, et al. J Exp Med. 2012



KLF2 mutations in SMZL

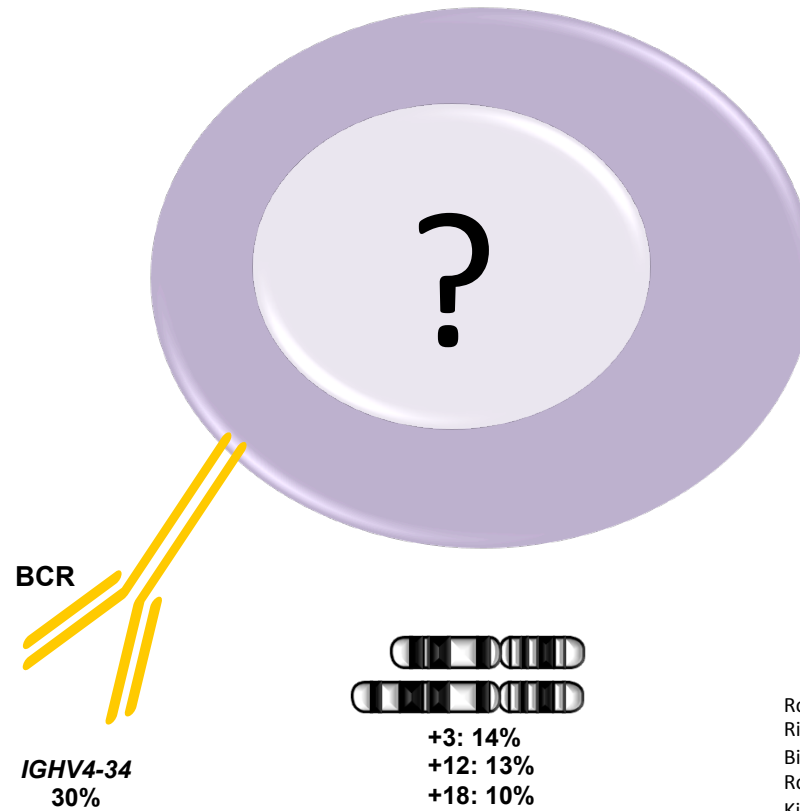
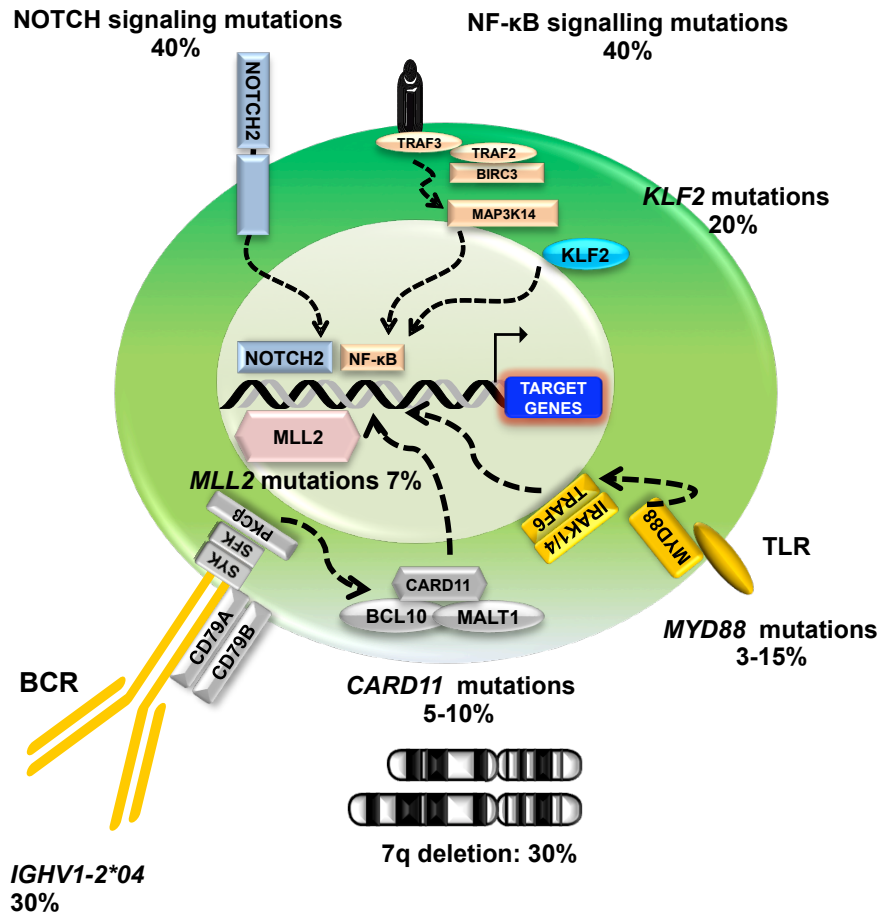


Clipson A, *Leukemia* 2015



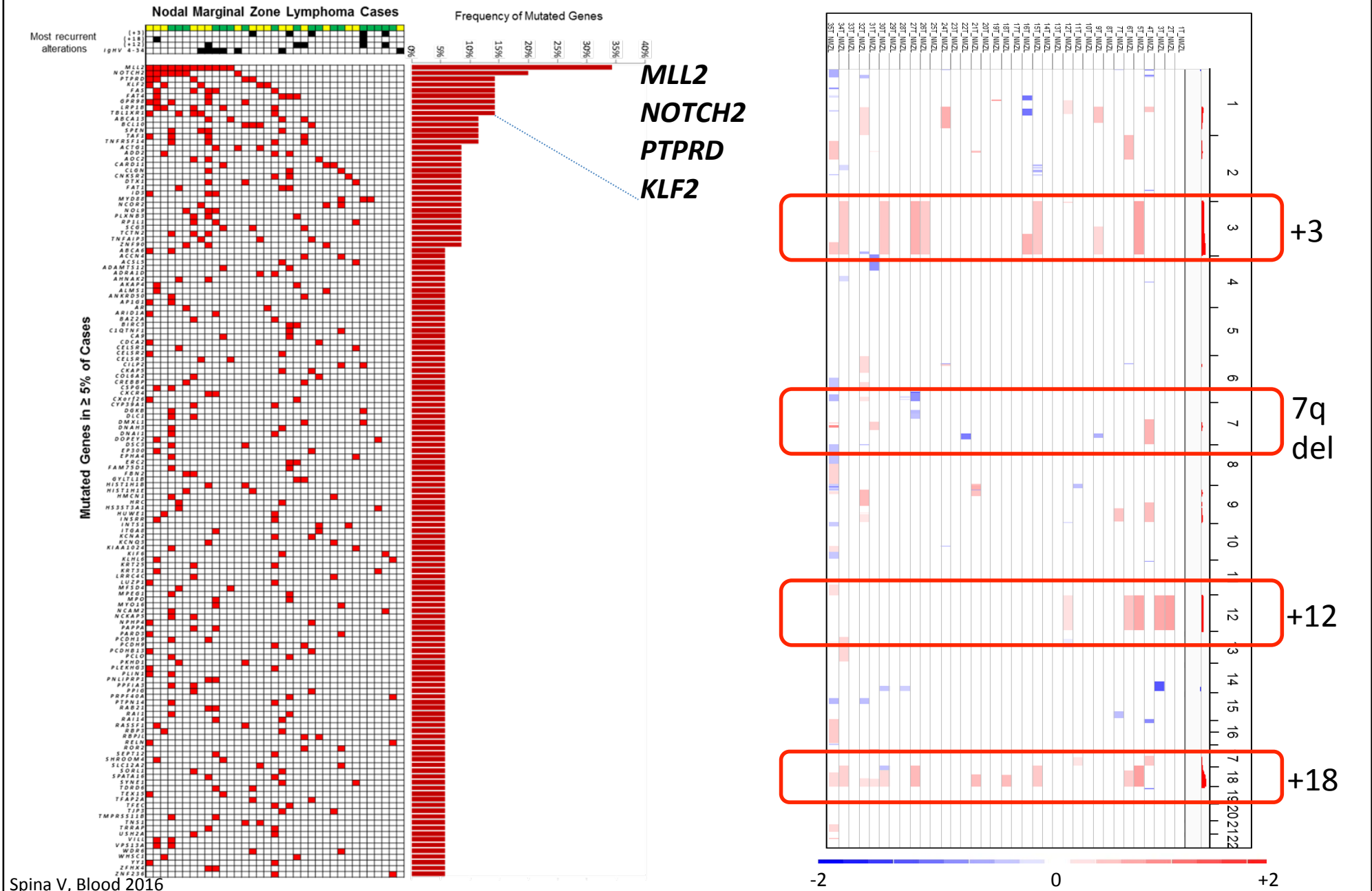
SMZL

NMZL

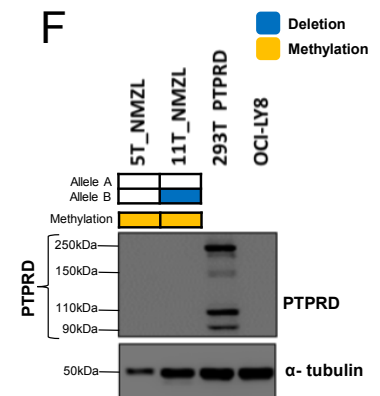
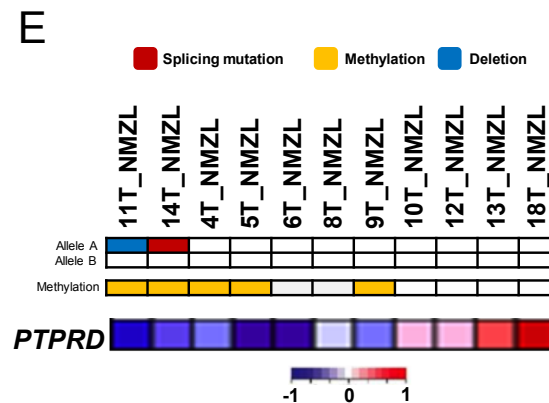
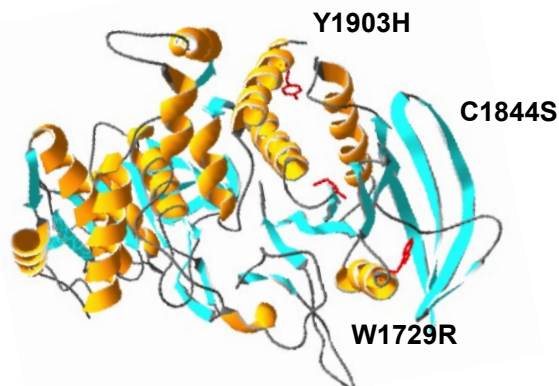
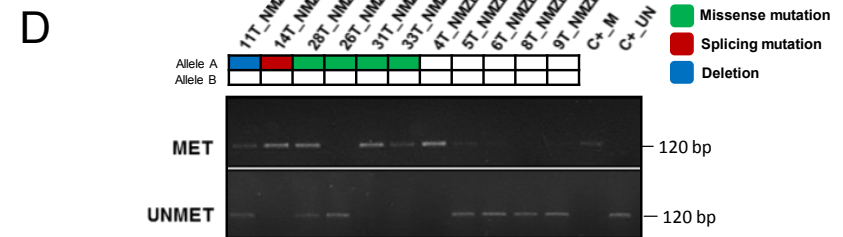
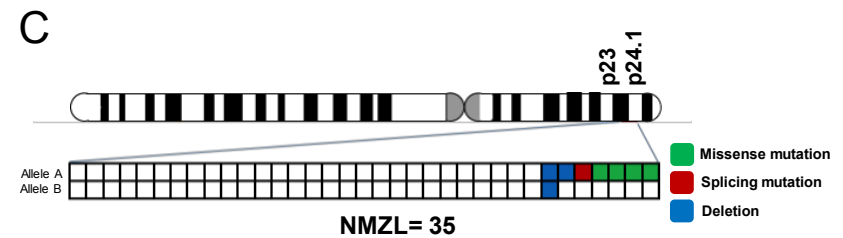
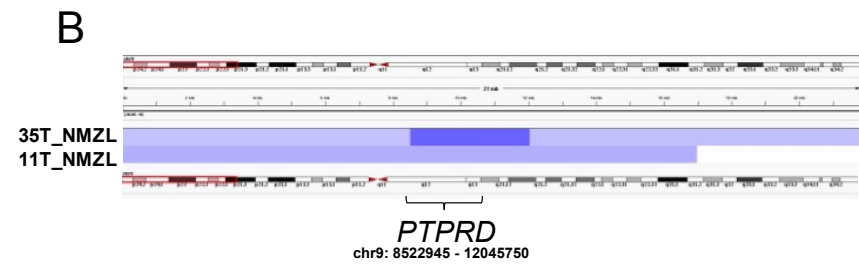
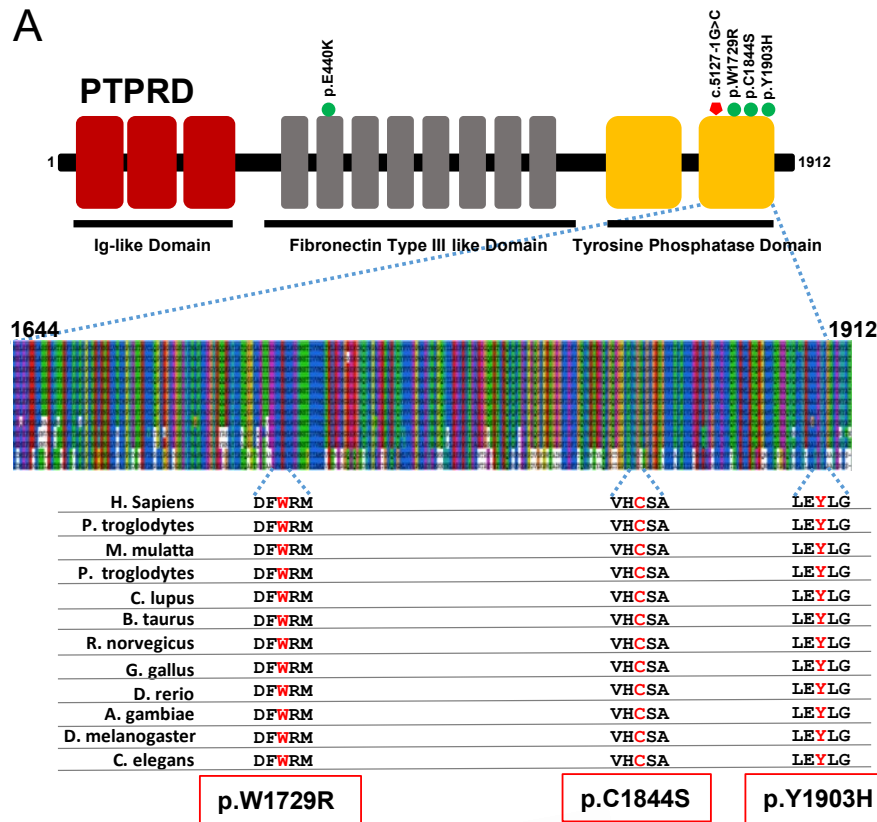


Rossi D, *Blood* 2011
 Rinaldi A, *Blood* 2011
 Bikos V, *Leukemia* 2012
 Rossi D, *J Exp Med* 2012
 Kiel MJ, *J Exp Med* 2012
 Parry M, *PLoS One* 2013
 Martinez N, *Leukemia* 2014
 Piva R, *Leukemia* 2015
 Clipson A, *Leukemia* 2015
 Parry M, *Clin Cancer Res* 2015
 Arribas A, *Blood* 2015

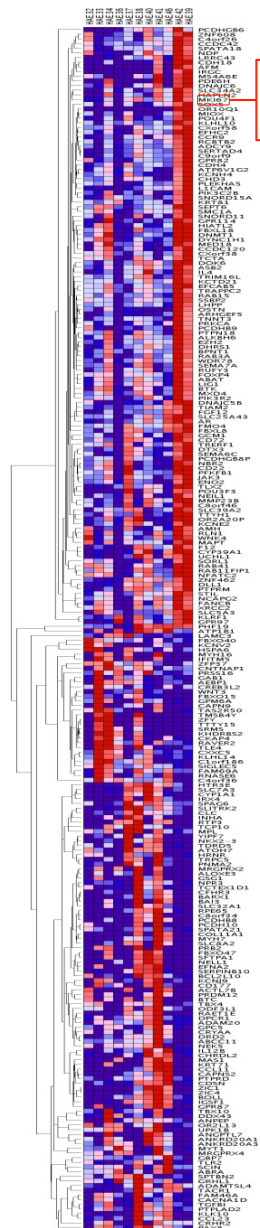
The coding genome of NMZL



PTPRD in NMZL

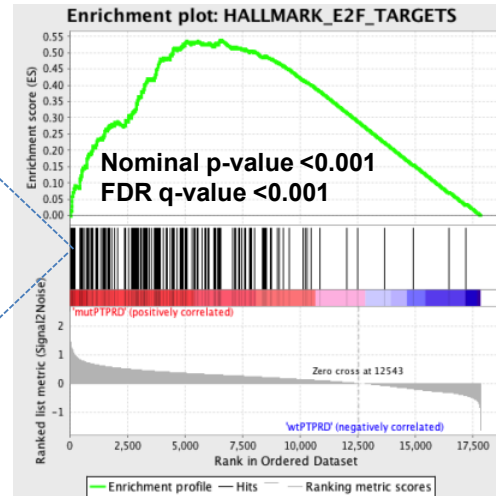


PTPRD mutations in NMZL associate with cell cycle signature and proliferation

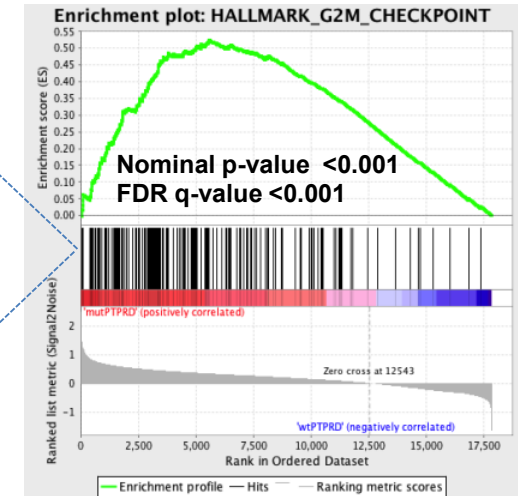


Ki67

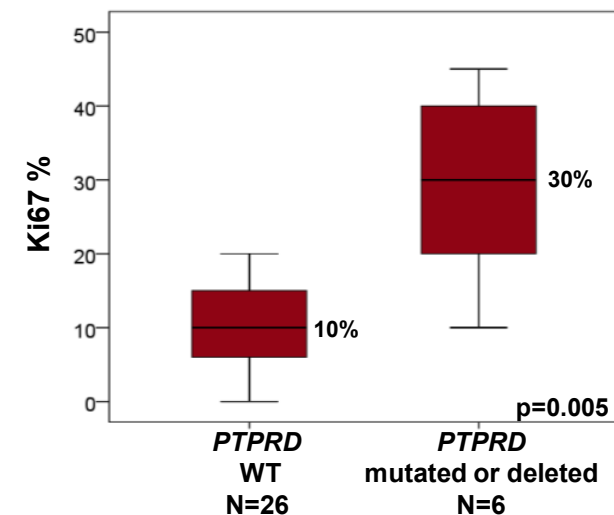
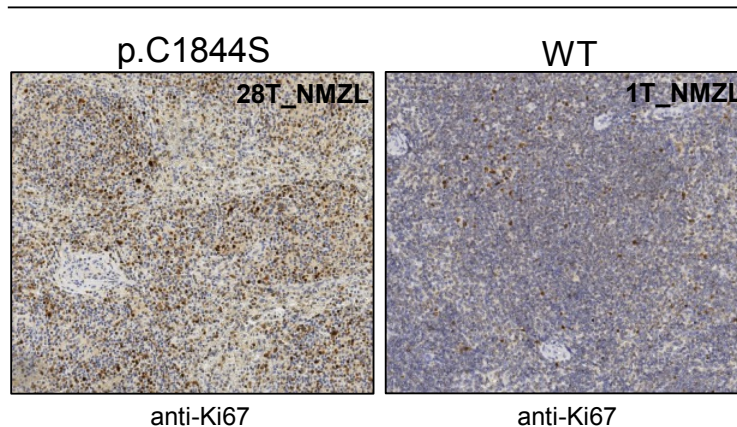
- EZH2
- DNMT1
- MKI67**
- SMC1A
- LIG1
- CIT
- ING3
- ILF3



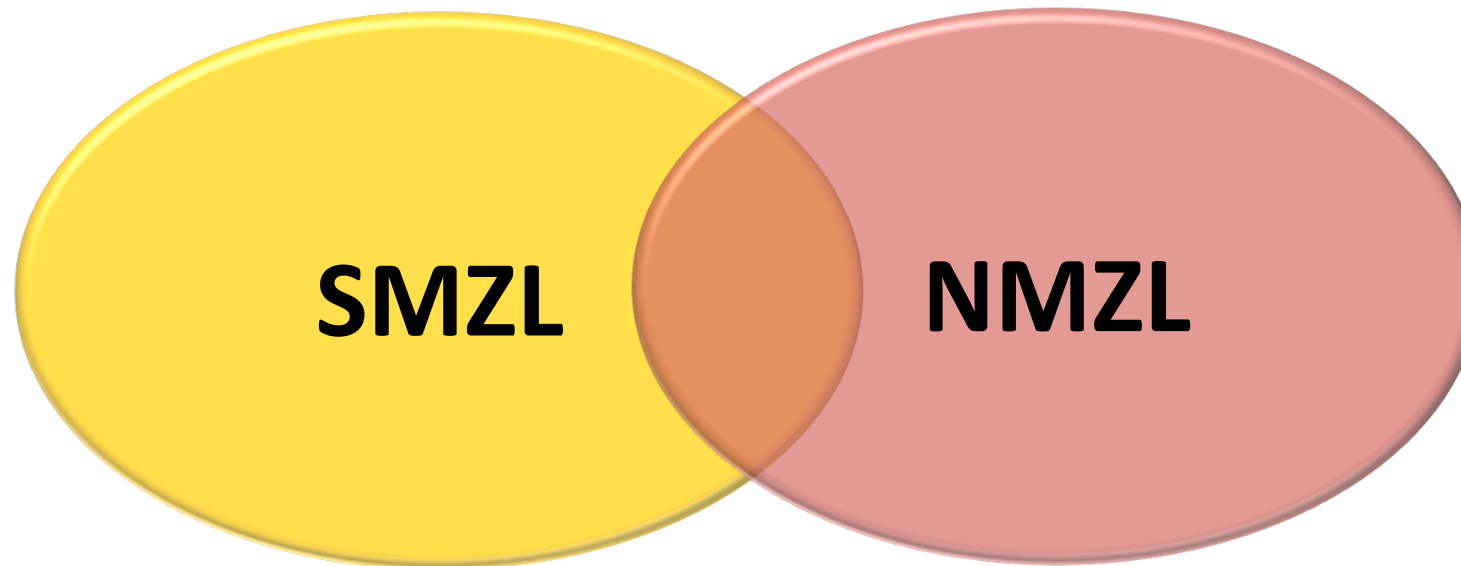
- EZH2
- STIL
- MKI67**
- SMC1A
- CDC7
- ILF3
- CENPE
- E2F3
- SUV39H1



PTPRD



Splenic and nodal marginal zone lymphoma are distinguished by the different involved sites



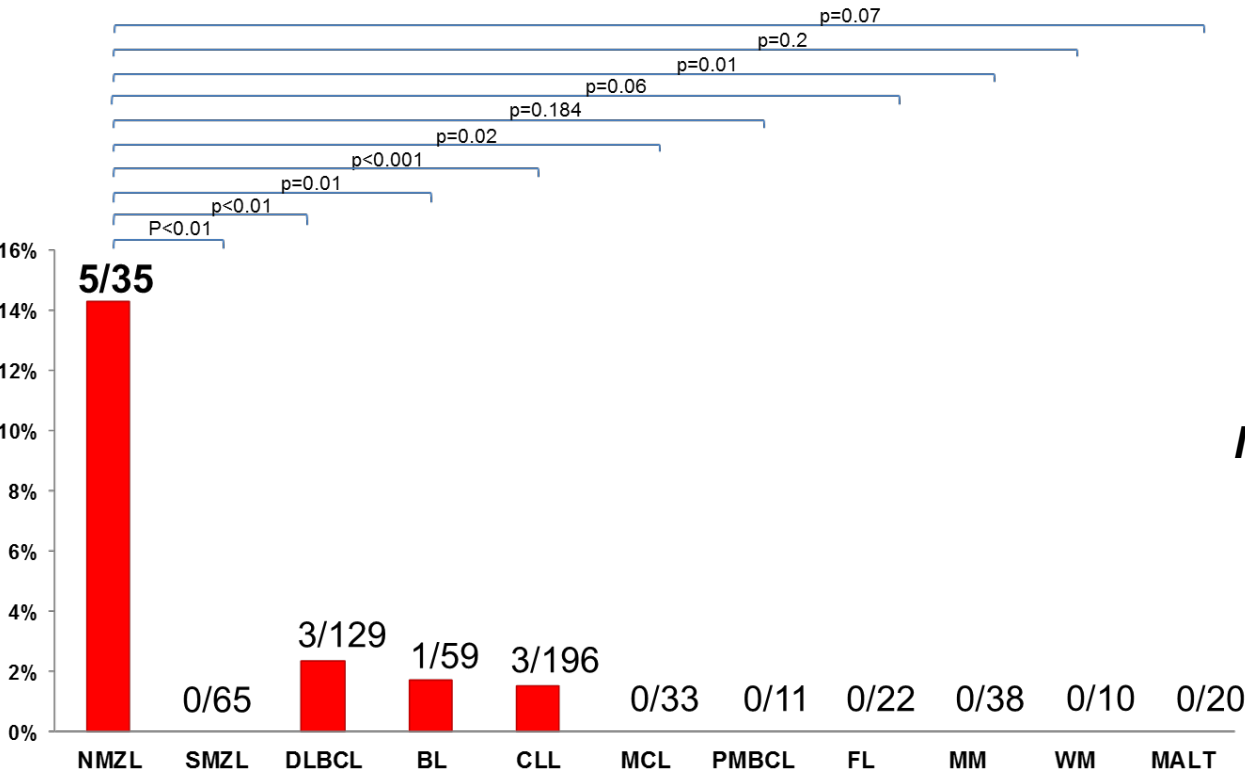
Shared features

Cytologic: clonal expansion of centrocyte-like and monocytoid-like B-cells

Histologic: marginal zone colonization with interfollicular expansion

Phenotypic: CD19+, CD20+, CD79a+, CD5-, CD10-, CD23-, BCL6-, cyclin D1-

PTPRD mutations are enriched in NMZL across mature B-cell tumors (n=619)



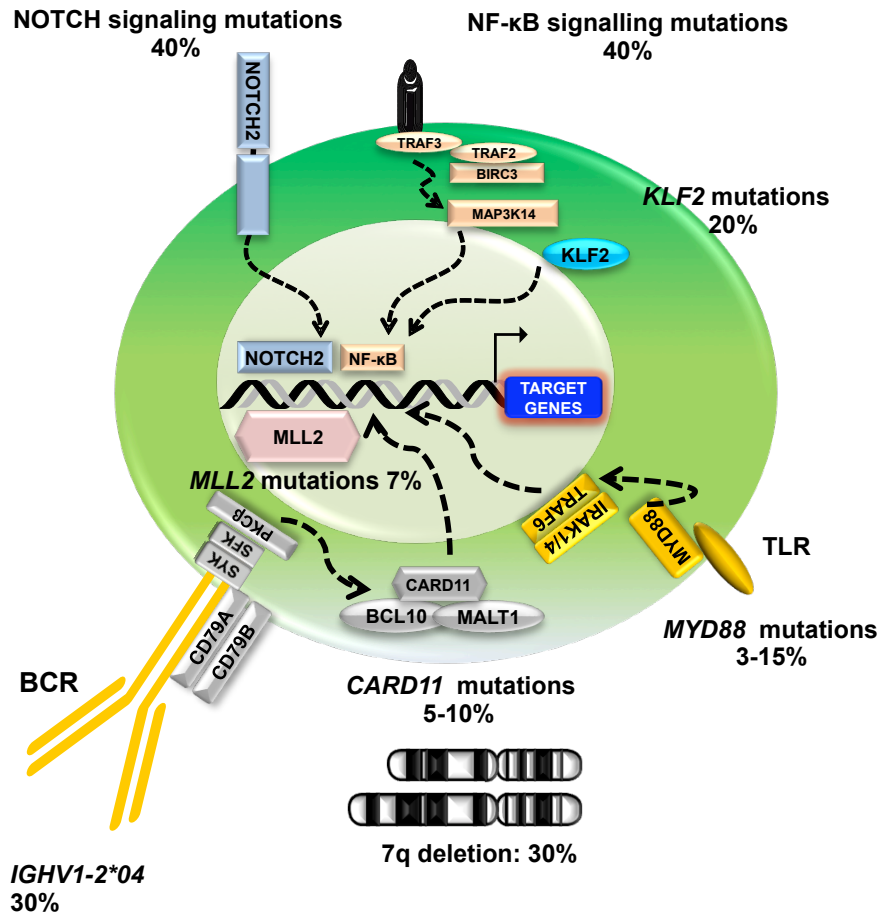
	NMZL	SMZL	p-value
<i>PTPRD</i>	14.3%	0%	p=.01
<i>MLL2</i>	34.3%	7.7%	p<.001
<i>NOTCH2</i>	20%	21.6%	p=1
<i>KLF2</i>	14.3%	21.1%	p=0.5

Mutation frequency: Low to High

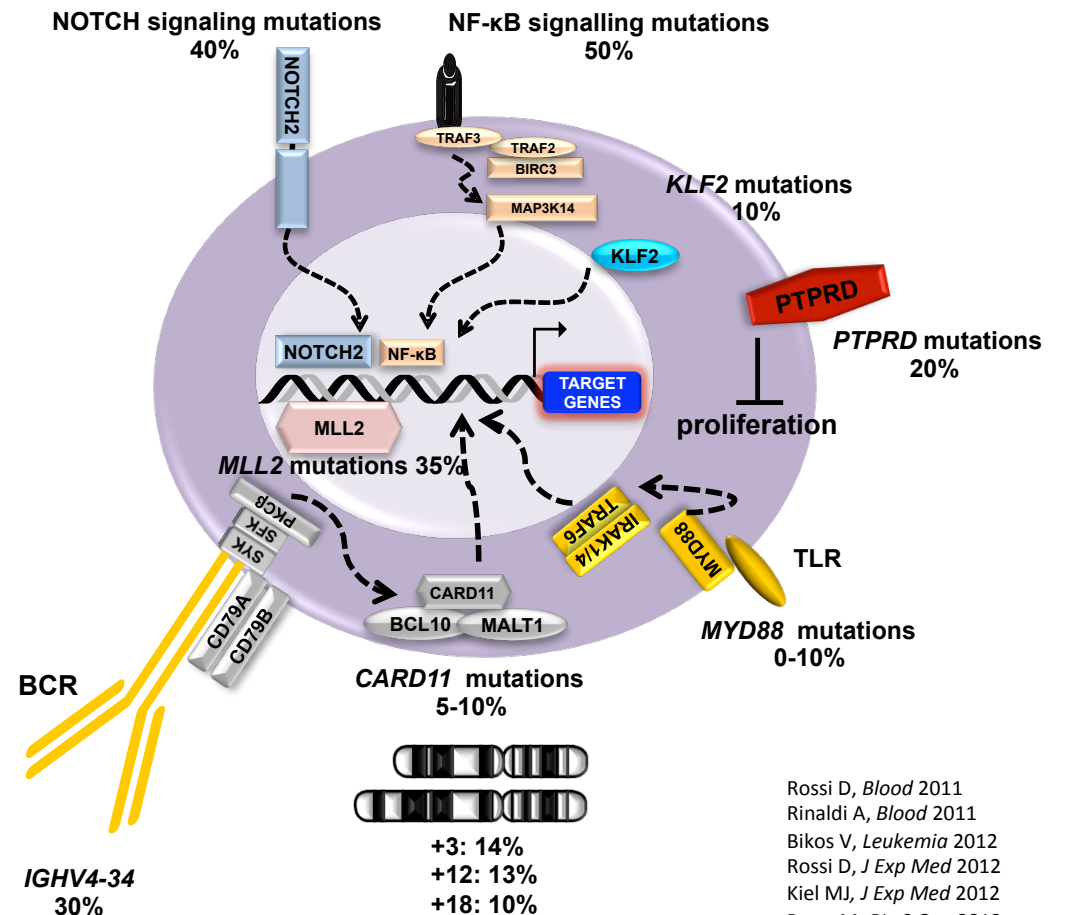
Genes mutated in $\geq 15\%$ of NMZL and/or SMZL

Molecular pathogenesis of non-MALT MZL

SMZL



NMZL

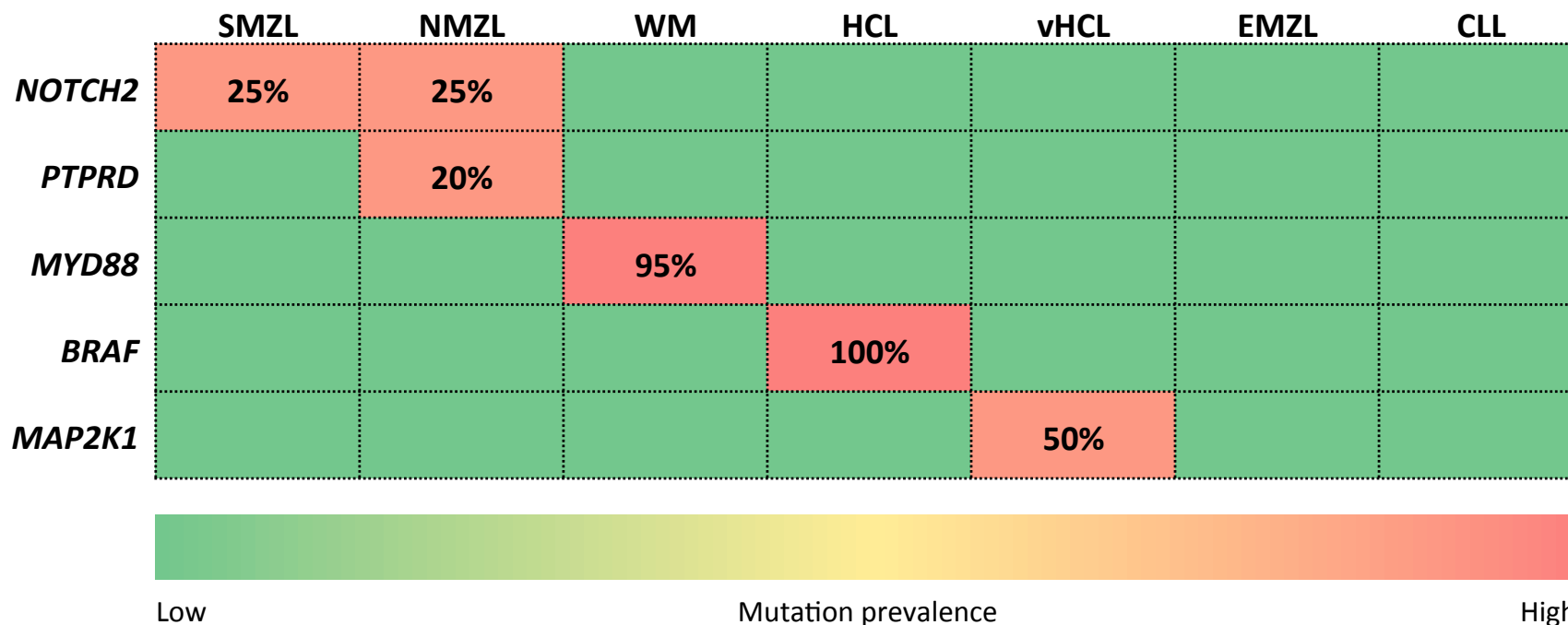


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 Arribas A, *Blood* 2015
 Spina V, *Blood* 2016

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Genetics can assist in the specification of histopathologically challenging B-cell LPDs



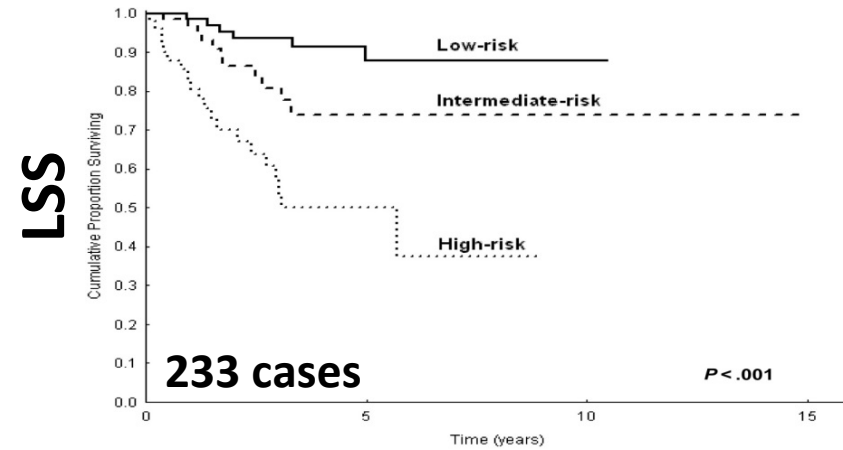
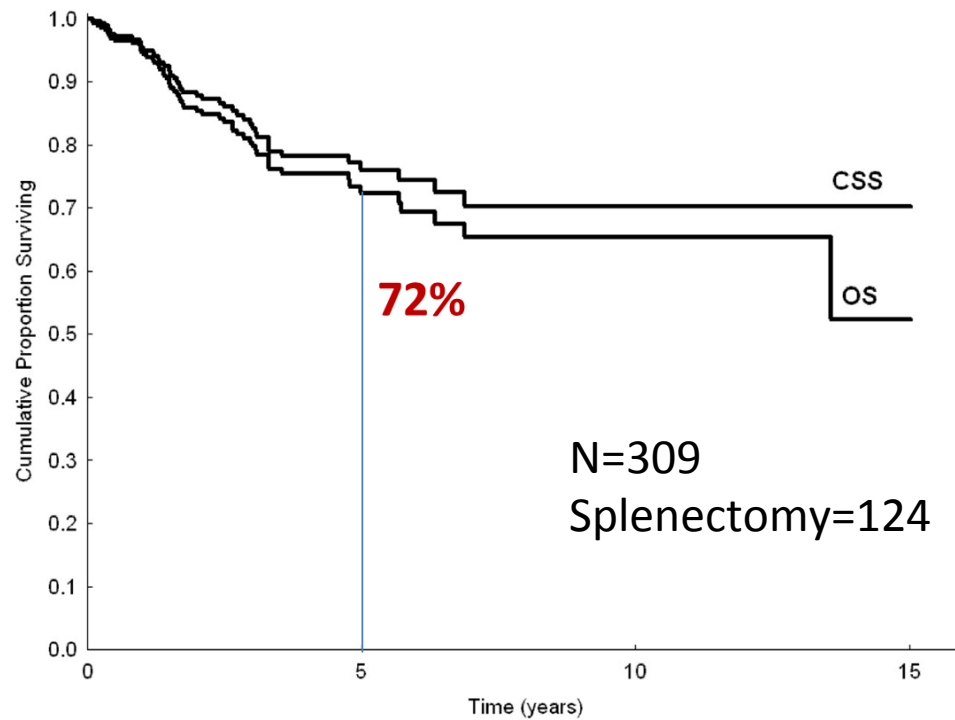
Tiacci et al, New Engl J Med 2011
 Treon et al, New Engl J Med 2012
 Landau et al, Cell 2013
 Rossi et al, J Exp Med 2013
 Rossi et al, ASH 2013
 Waterfall et al, Nat Genet 2013
 Spina et al, Blood 2016

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Background

SMZL shows an heterogeneous clinical course



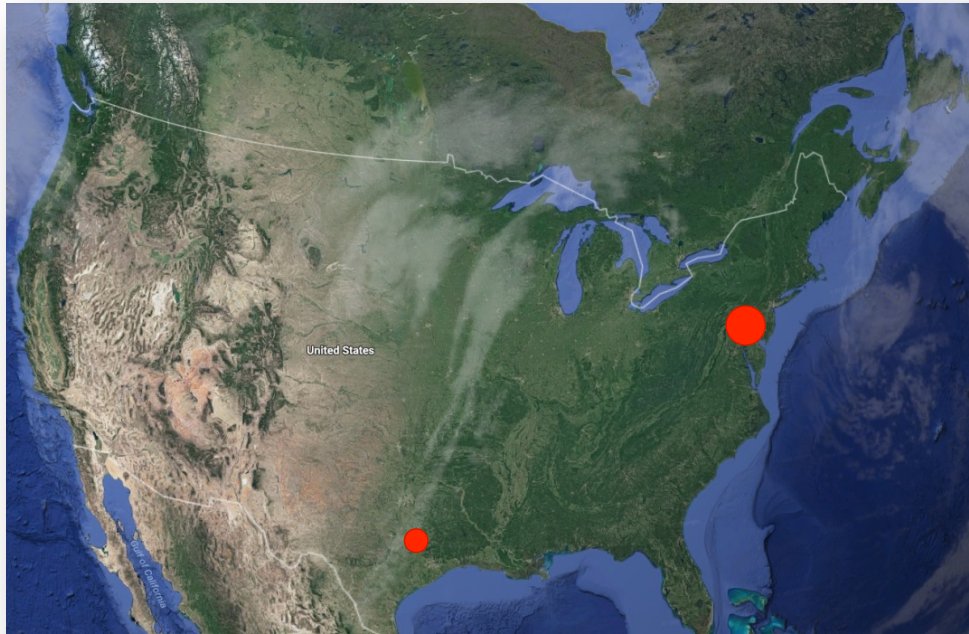
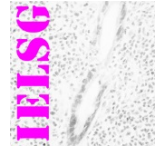
III score
Hemoglobin
LDH
Albumin

	Deaths	5-year LSS
Low	16%	88%
Int	30%	73%
High	54%	50%

Molecular features of SMZL associated with prognosis

- *NOTCH2* mutations
- *KLF2* mutation
- *TP53* disruption
- *IGHV1-2*04* usage
- 7q deletion

IELSG 46: INTEGRATED MOLECULAR AND CLINICAL PROFILING TO OPTIMIZE OUTCOME PREDICTION IN SMZL



New York

- Columbia University Medical Center
- Weill Cornell Medicine

Houston

ClinicalTrials.gov: NCT02945319



Basel
Bellinzona
Bern
St. Gallen



London
Bournemouth



Lisbon



Milano

- Ist tumori
- Niguarda
- San Raffaele
- Policlinico

Aviano
Alessandria
Bergamo
Bologna
Brescia
Modena
Novara
Pavia

Roma
Torino
Udine
Varese
Verona
Vicenza



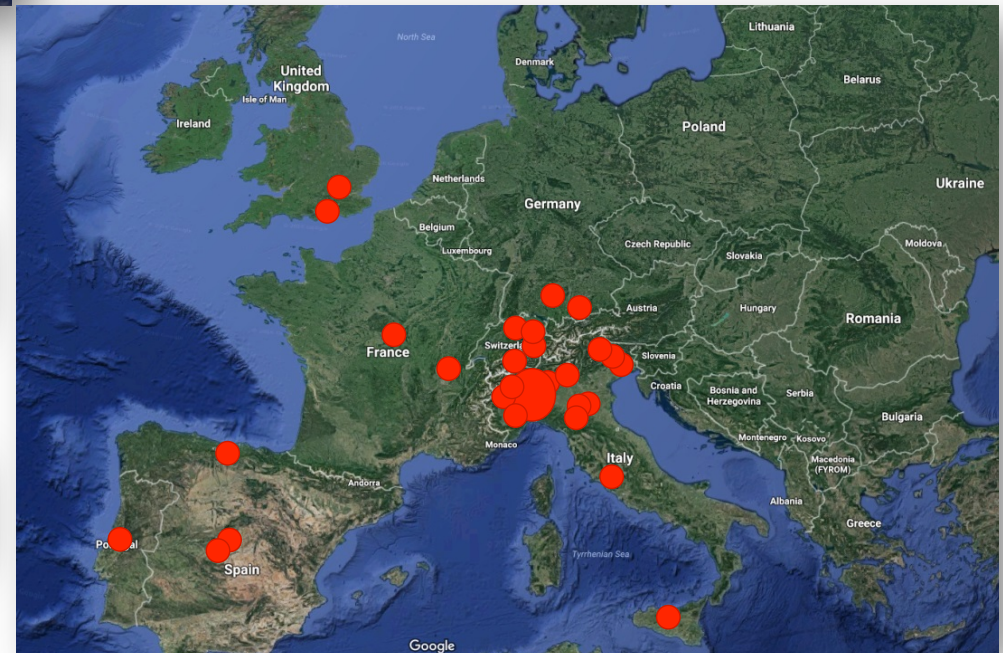
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Paris



Heidelberg
Kiel
Munich
Ulm



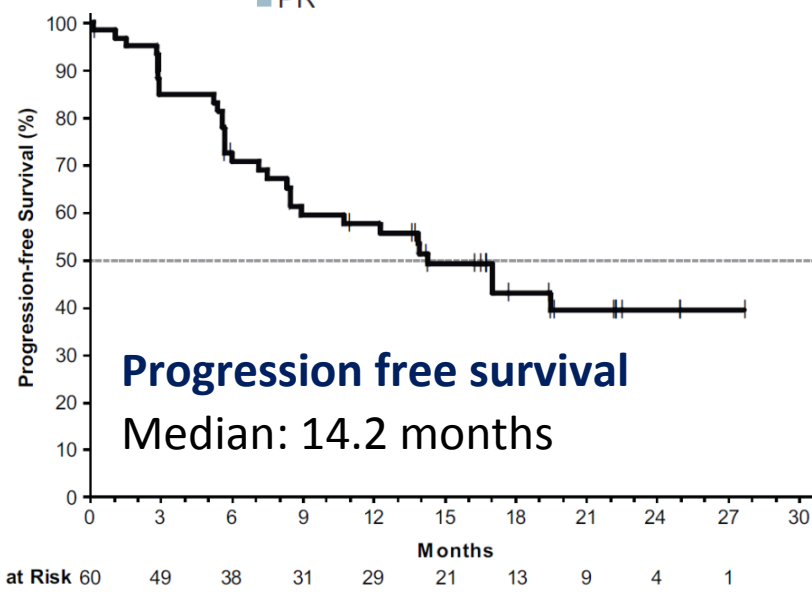
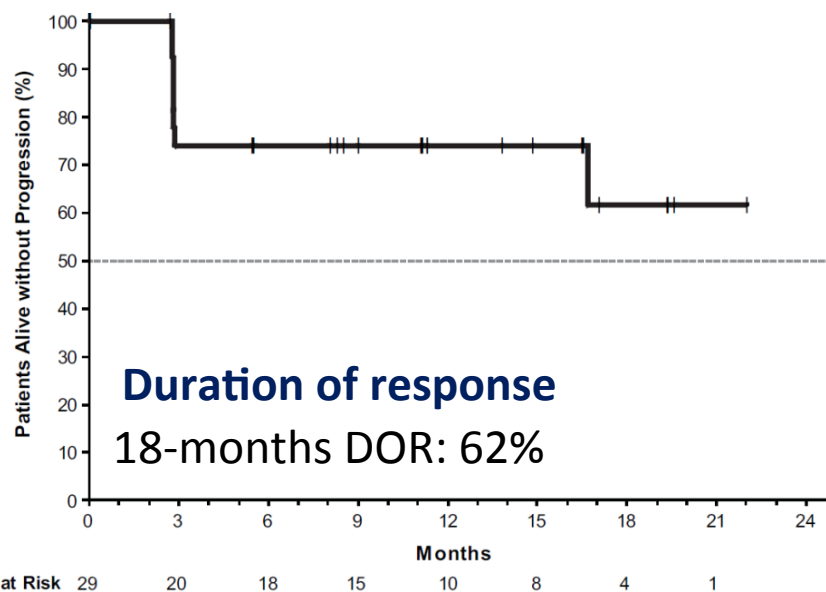
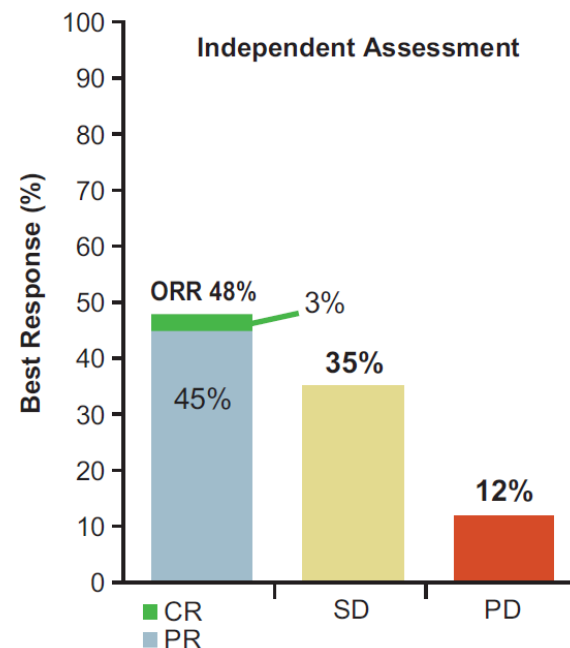
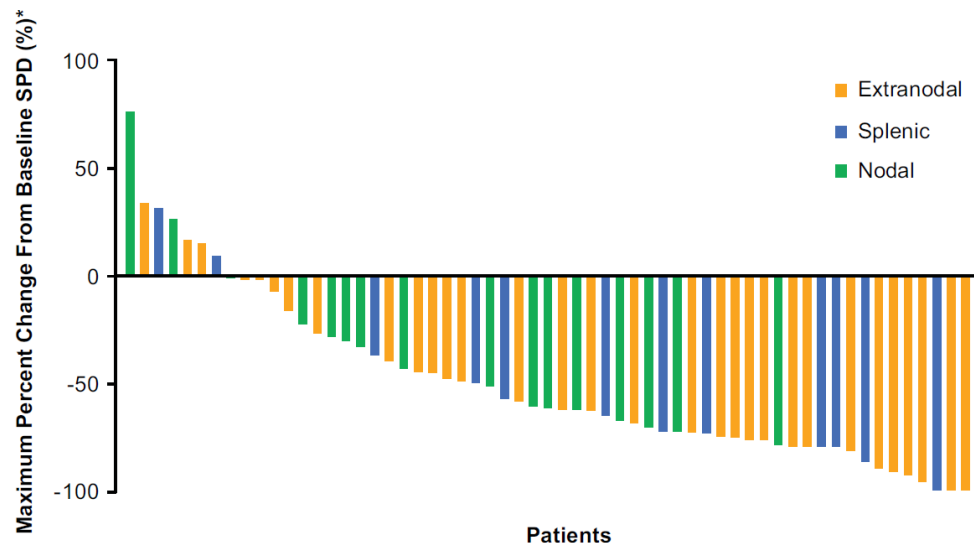
Barcelona
Madrid
Santander
Toledo



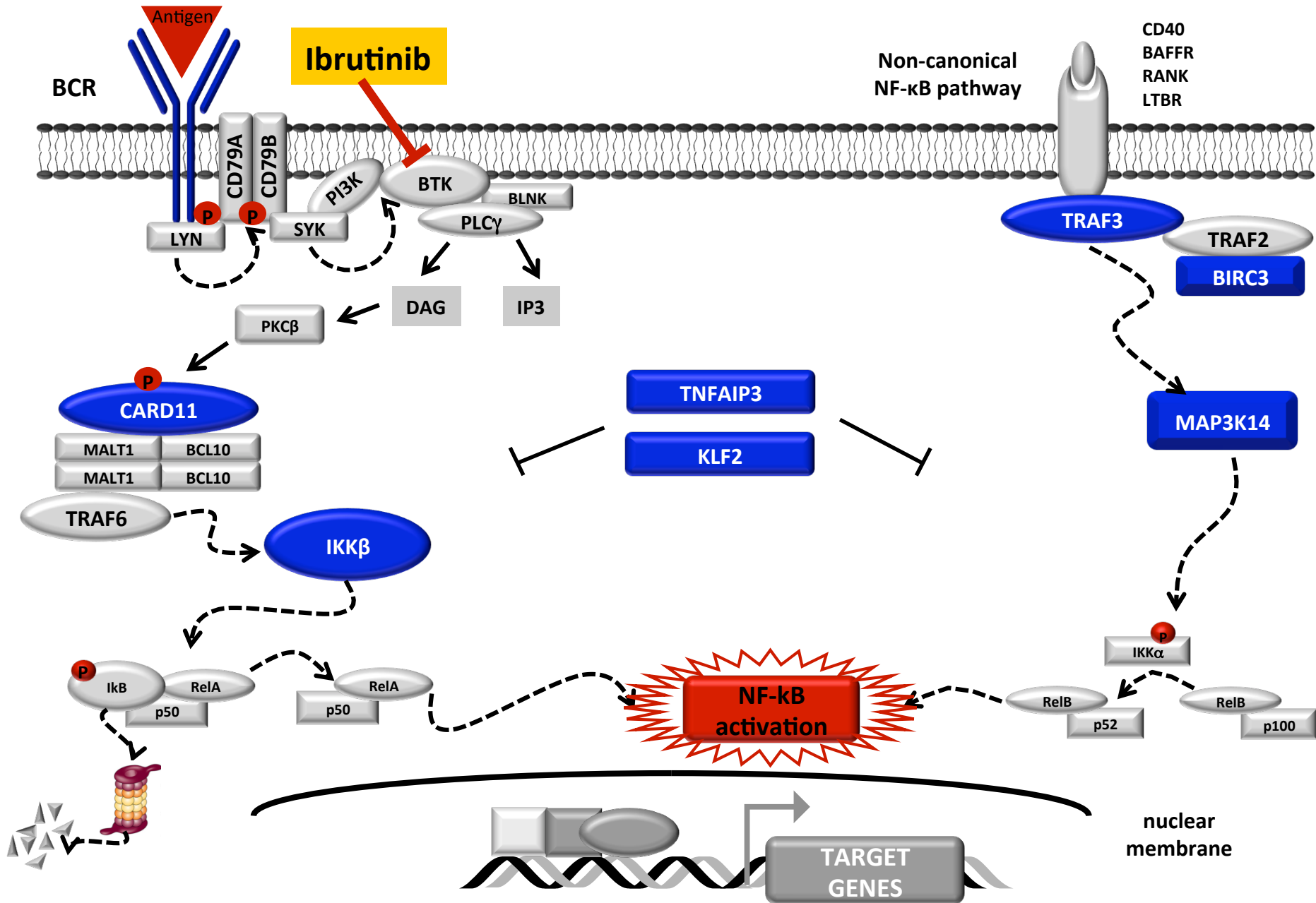
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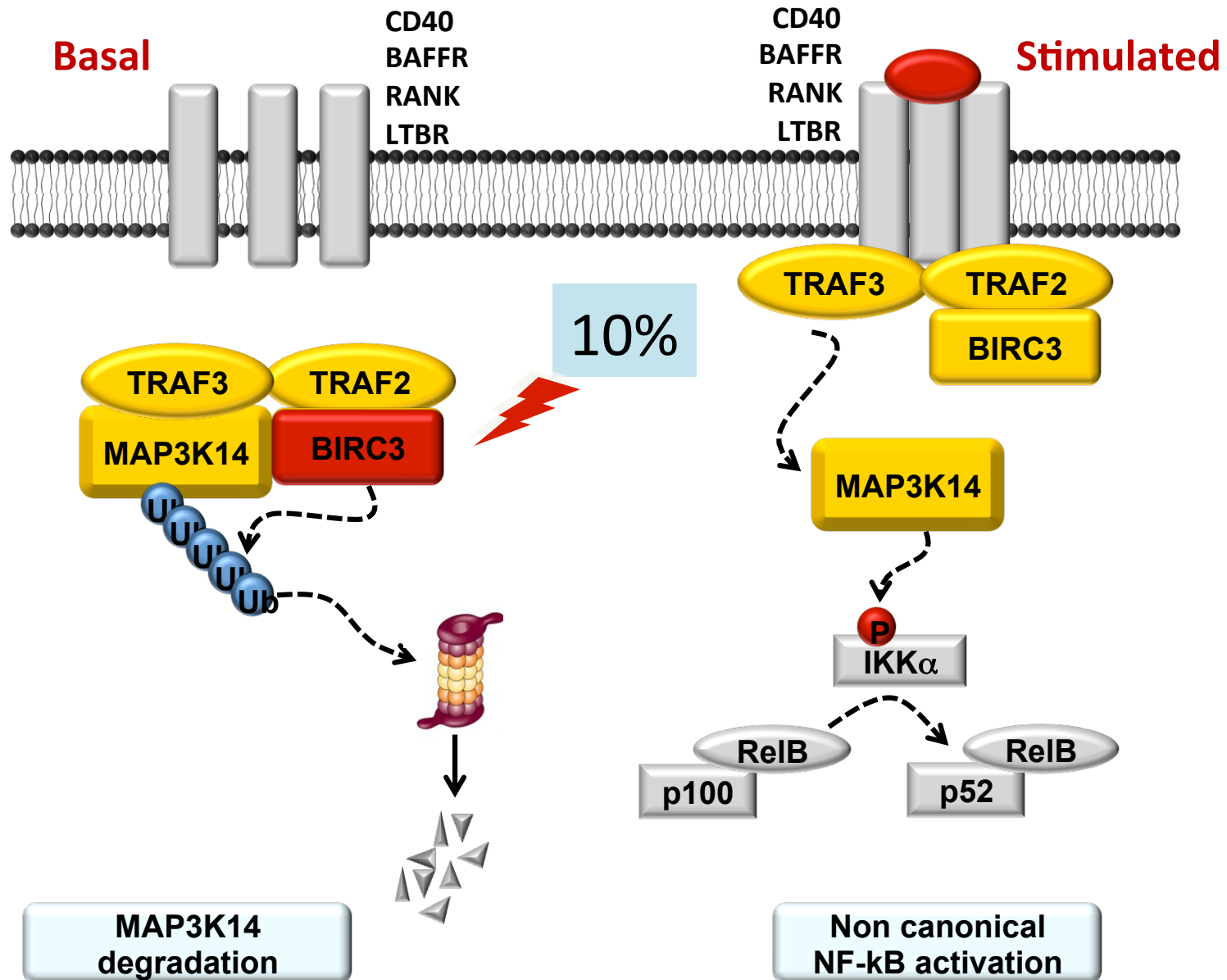
Ibrutinib monotherapy in R/R MZL



Signaling activation in MZL

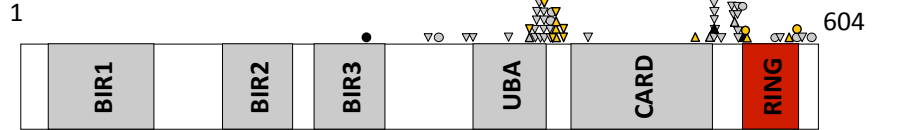


BIRC3 mutations in SMZL

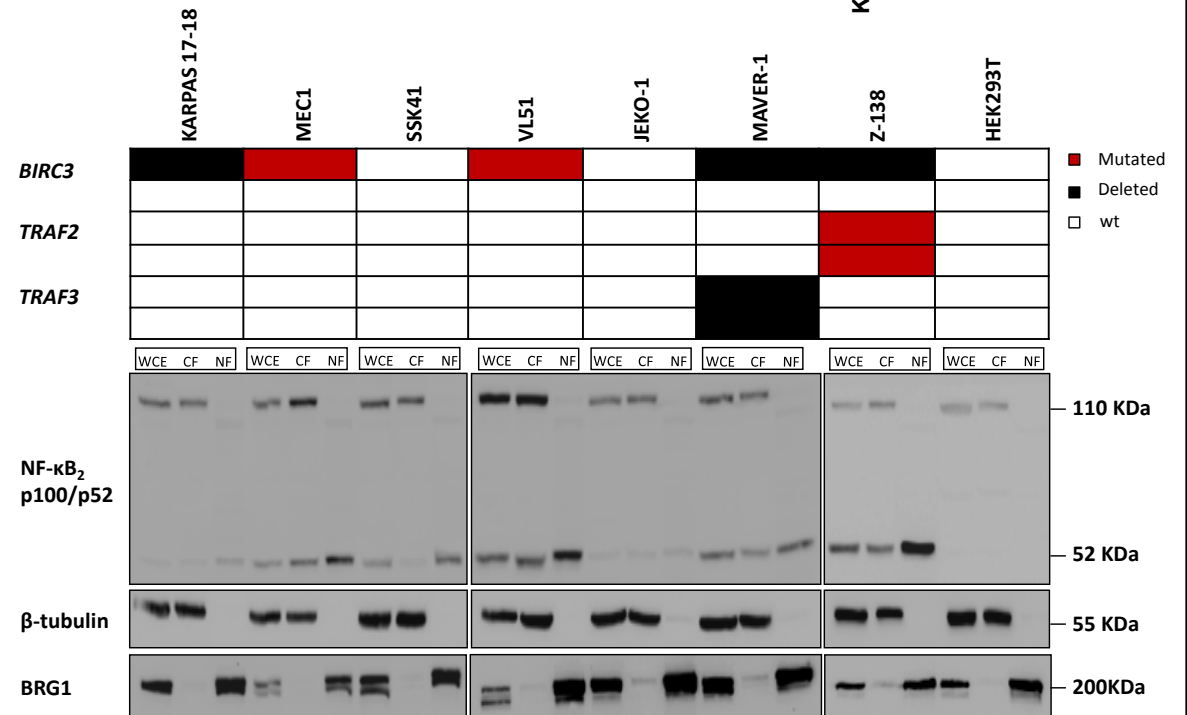
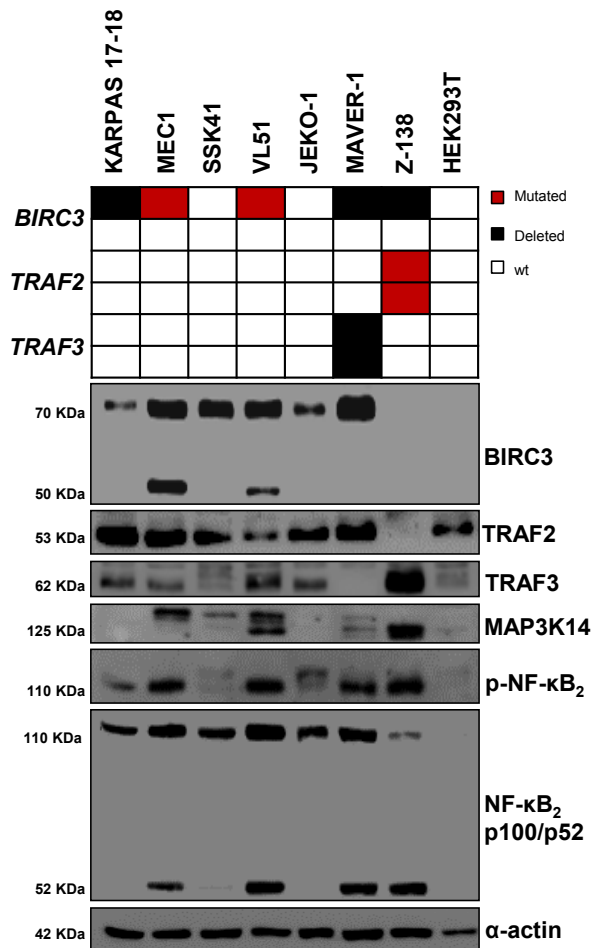
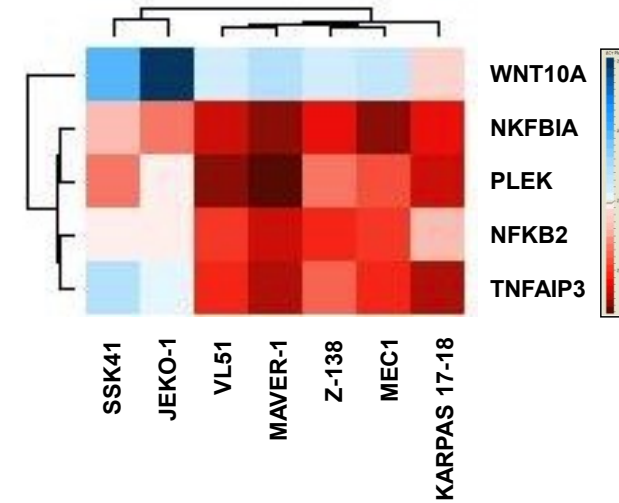


BIRC3 mutations activate non-canonical NF- κ B

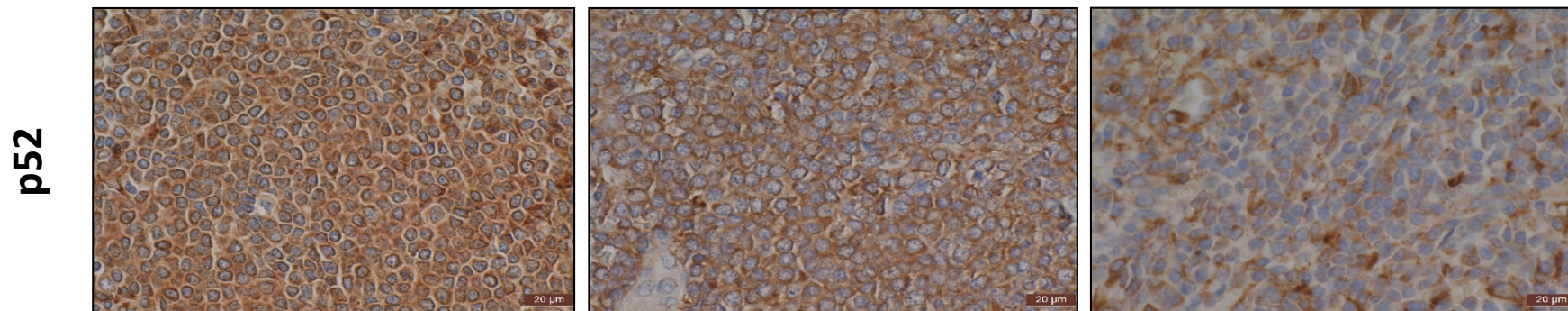
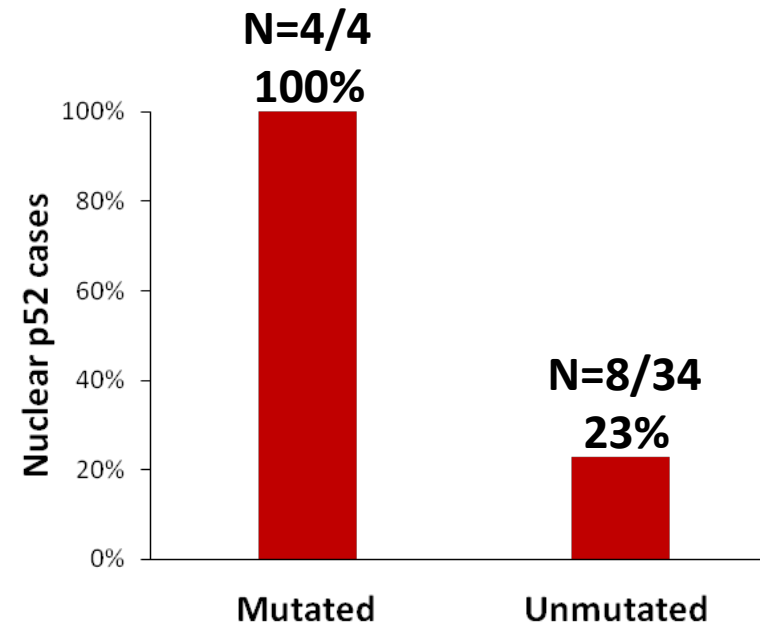
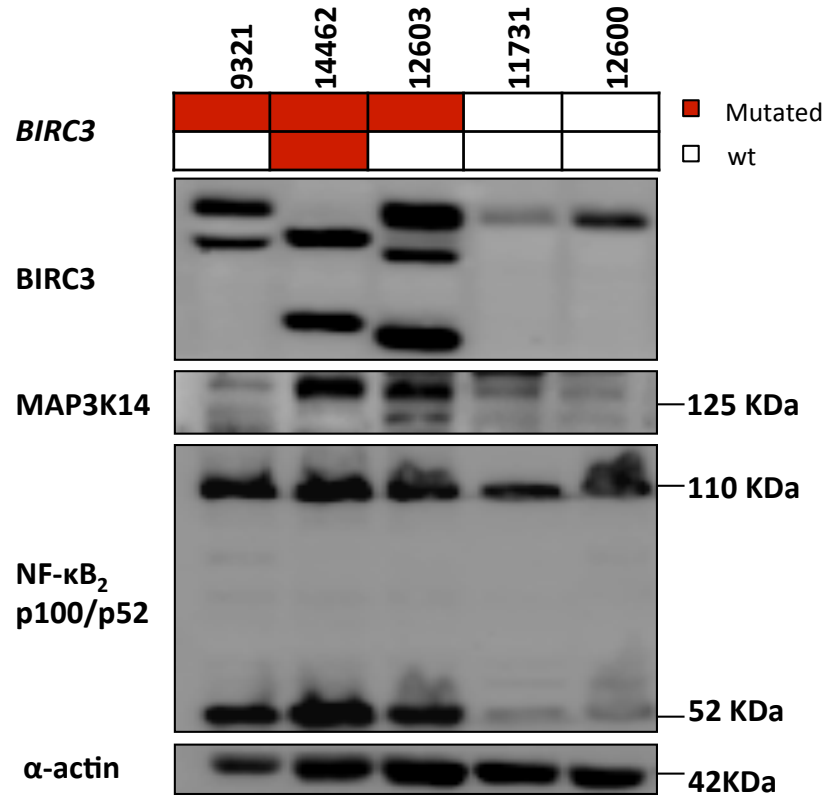
BIRC3



Non-canonical NF- κ B genes



Non-canonical NF- κ B pathway is active in *BIRC3* mutated primary samples

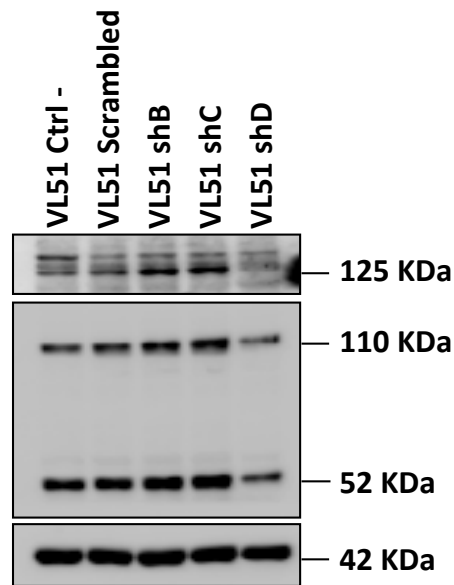
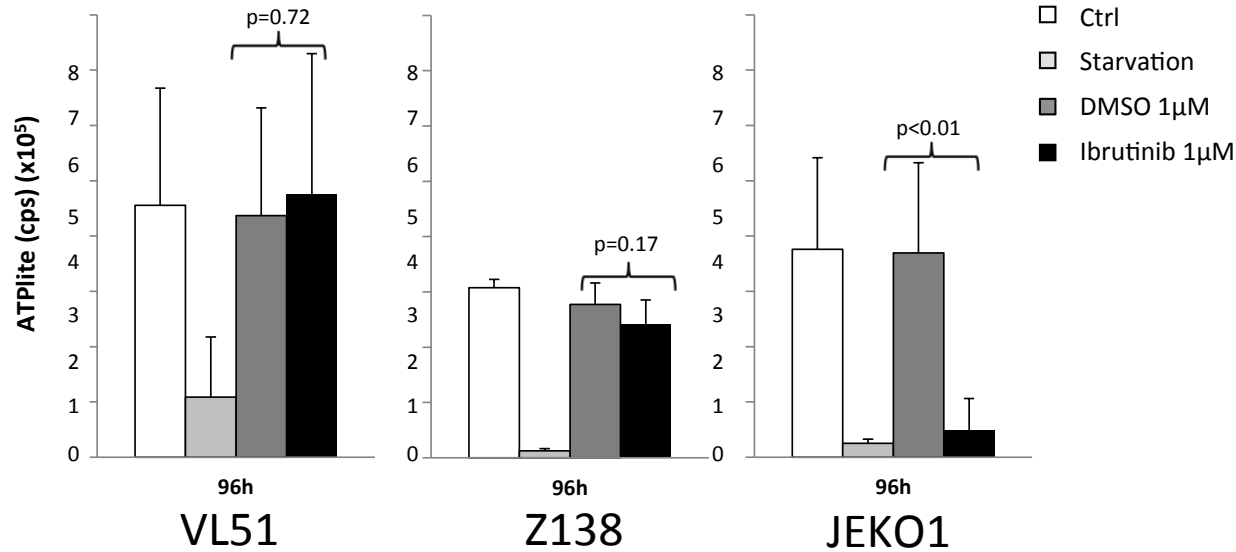


Nuclear+Cytoplasmic

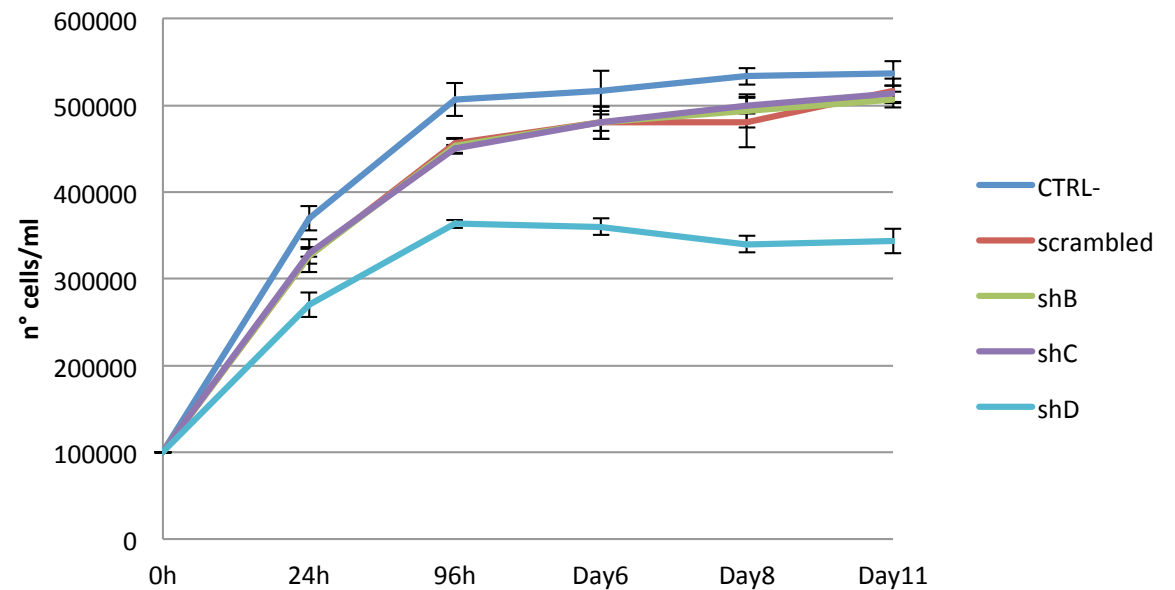
Cytoplasmic

p52 Negative

BIRC3 mutated cells are addicted of NIK



VL51 (Tripan Blue)



- NMZL and SMZL are enriched of mutations in genes involved in the physiological development of MZ B-cells
- *NOTCH2* mutations are specific for SMZL and NMZL across mature B-cell tumors (diagnostic implications)
- *PTPRD* mutations are specific for NMZL across mature B-cell tumors (diagnostic implications)
- Signals on the prognostic implications of mutations prompt the development of molecular biomarker-based score systems (pts counseling, clinical trial design)
- NF-kB mutations may mark resistance to ibrutinib (treatment tailoring)



Francesco Bertoni
Bernhard Gerber
Anastasios Stathis
Georg Stüssi
Emanuele Zucca

Franco Cavalli
Michele Ghilmini



Laura Pasqualucci
Riccardo Dalla Favera



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14-ICML

14th International Conference on Malignant Lymphoma

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SAVE THE DATE: June 14-17, 2017