

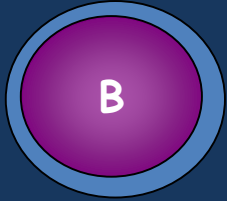
B cell Immunophenotyping for PID diagnosis and Classification

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Specific Reference Centre for Primary Immunodeficiencies-
Paediatric Immunology,
"Aghia Sophia" Children's Hospital, Athens. Greece

Basic B cell development pathway

Bone Marrow

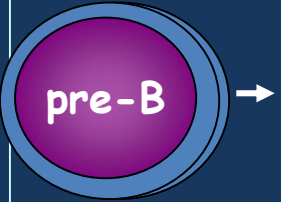


Spleen

Germinal Center

Basic B cell development pathway

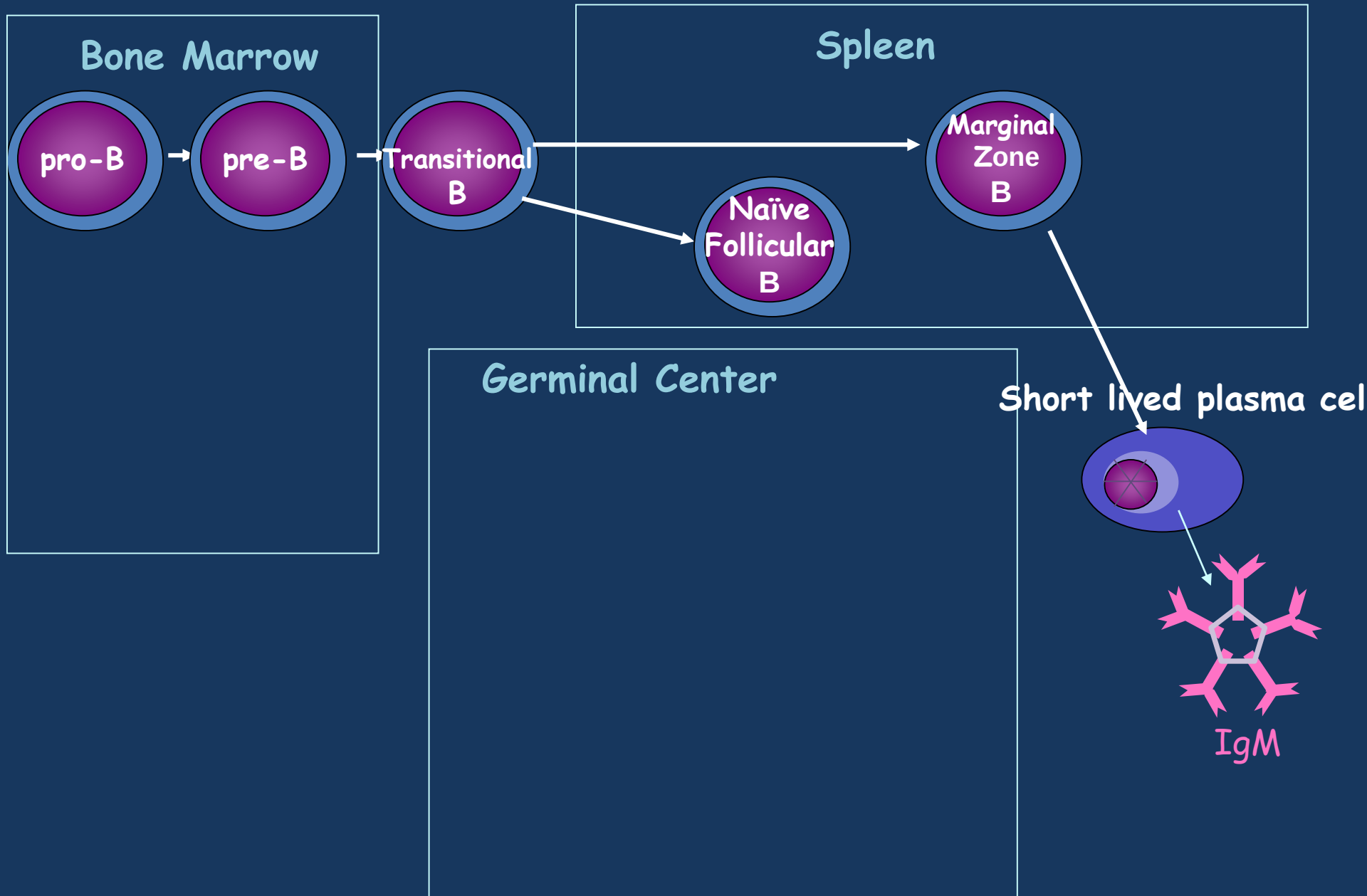
Bone Marrow

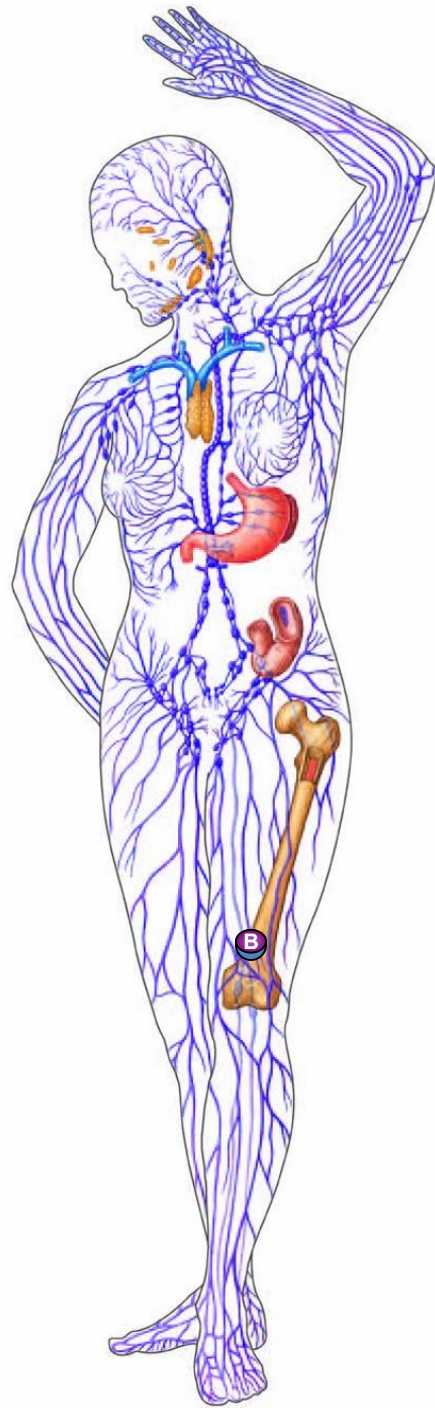


Spleen

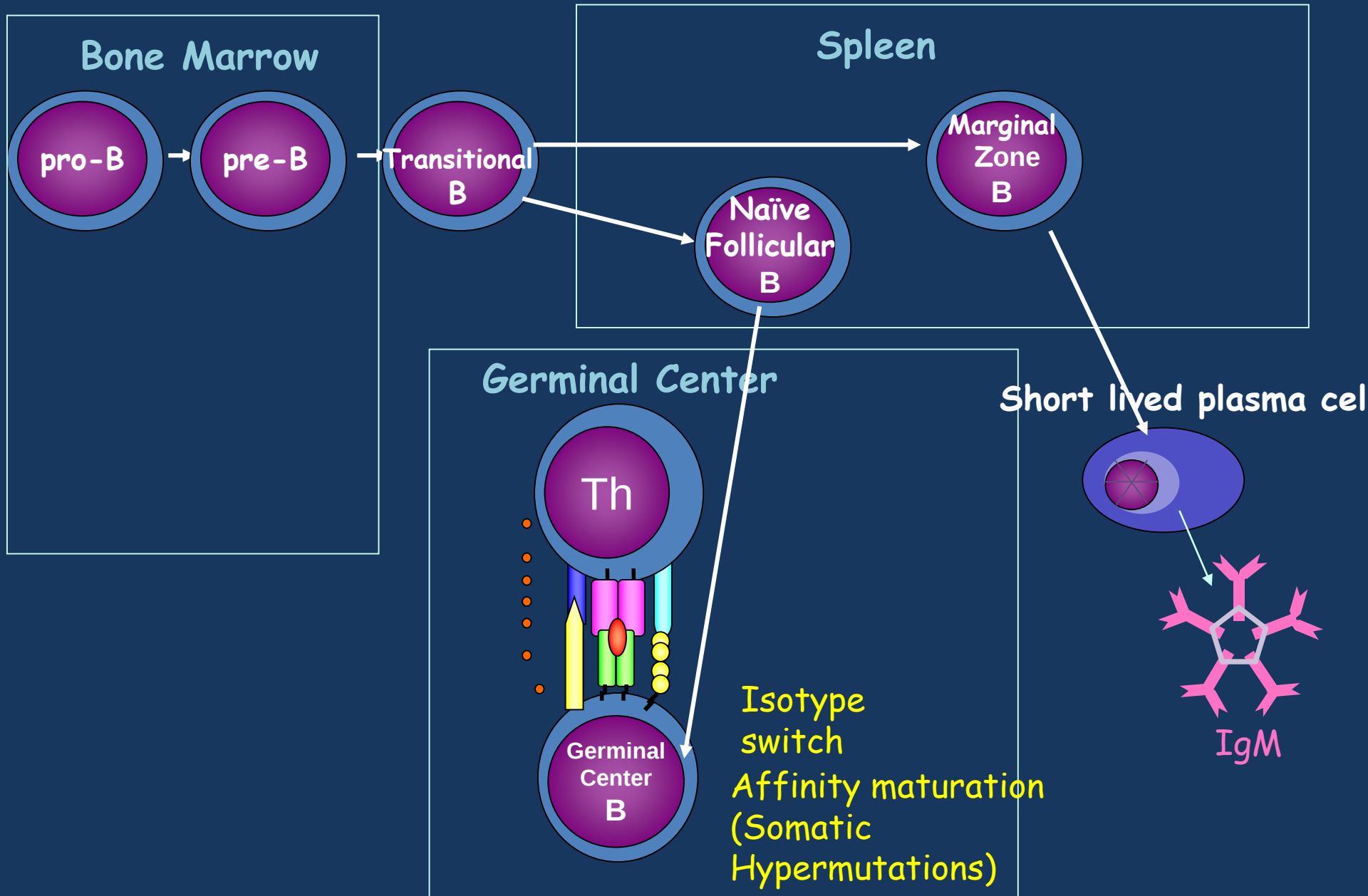
Germinal Center

Basic B cell development pathway

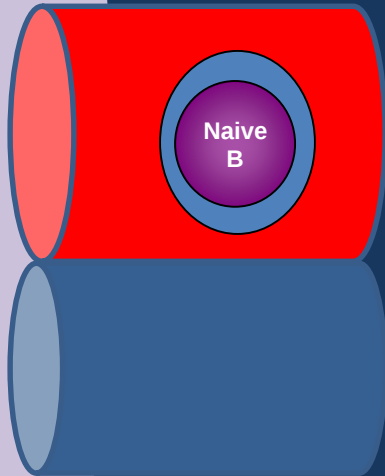




Basic B cell development pathway

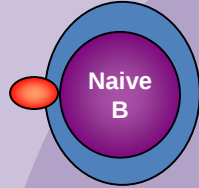


Lymph nodes

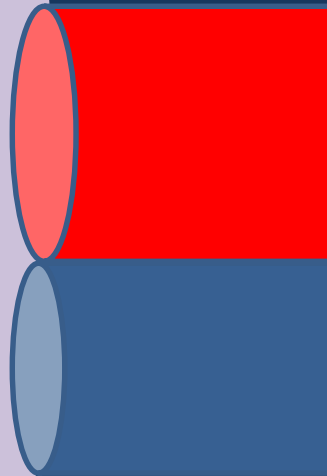


Lymph nodes

IgD+IgM+

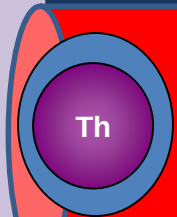
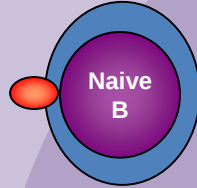


Naive
B



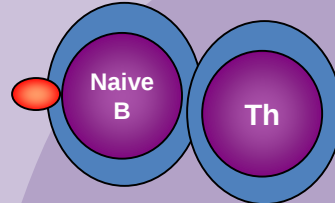
Lymph nodes

IgD+IgM+

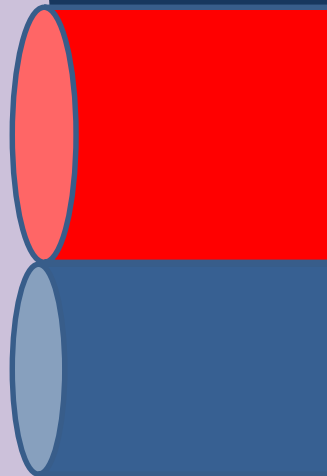


Lymph nodes

IgD+IgM+

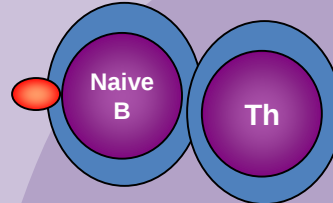


T cell help (CD40-CD40L)
Class switch



Fate depends on BCR
affinity!!!

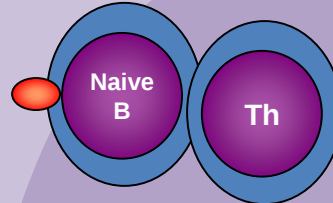
IgD-IgM+



T cell help (CD40-CD40L)
Class switch

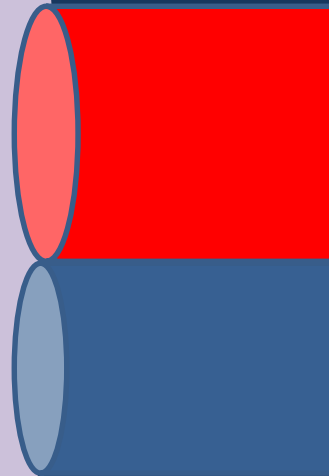
Fate depends on BCR
affinity!!!

IgD-IgM+



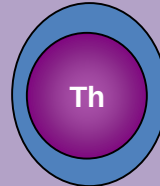
T cell help (CD40-CD40L)
Class switch

Low affinity BCR



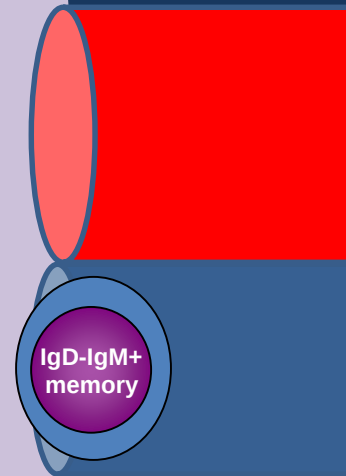
Fate depends on BCR
affinity!!!!

IgD-IgM+



T cell help (CD40-CD40L)
Class switch

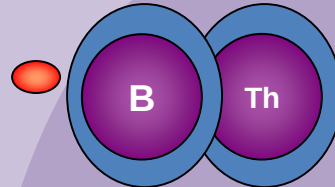
Low affinity BCR



IgD-IgM+
memory

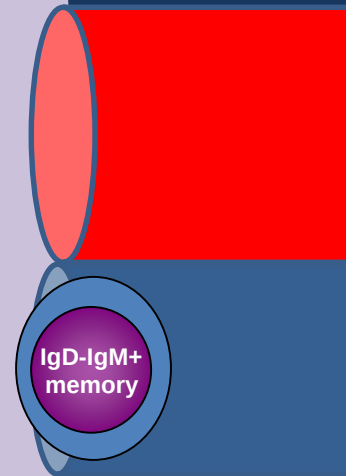
Fate depends on BCR
affinity!!!

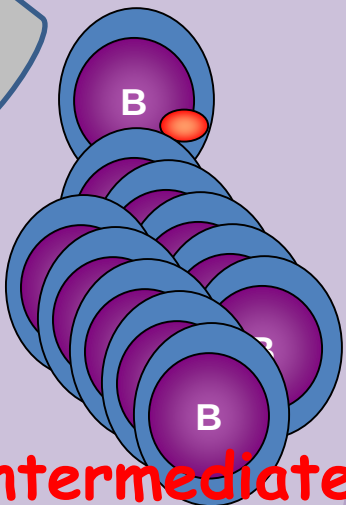
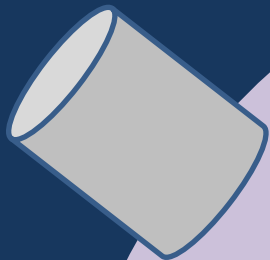
IgD-IgM+



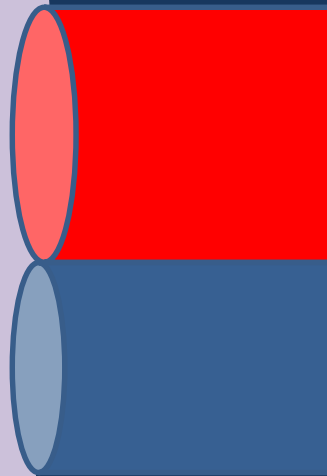
T cell help (CD40-CD40L)
Class switch

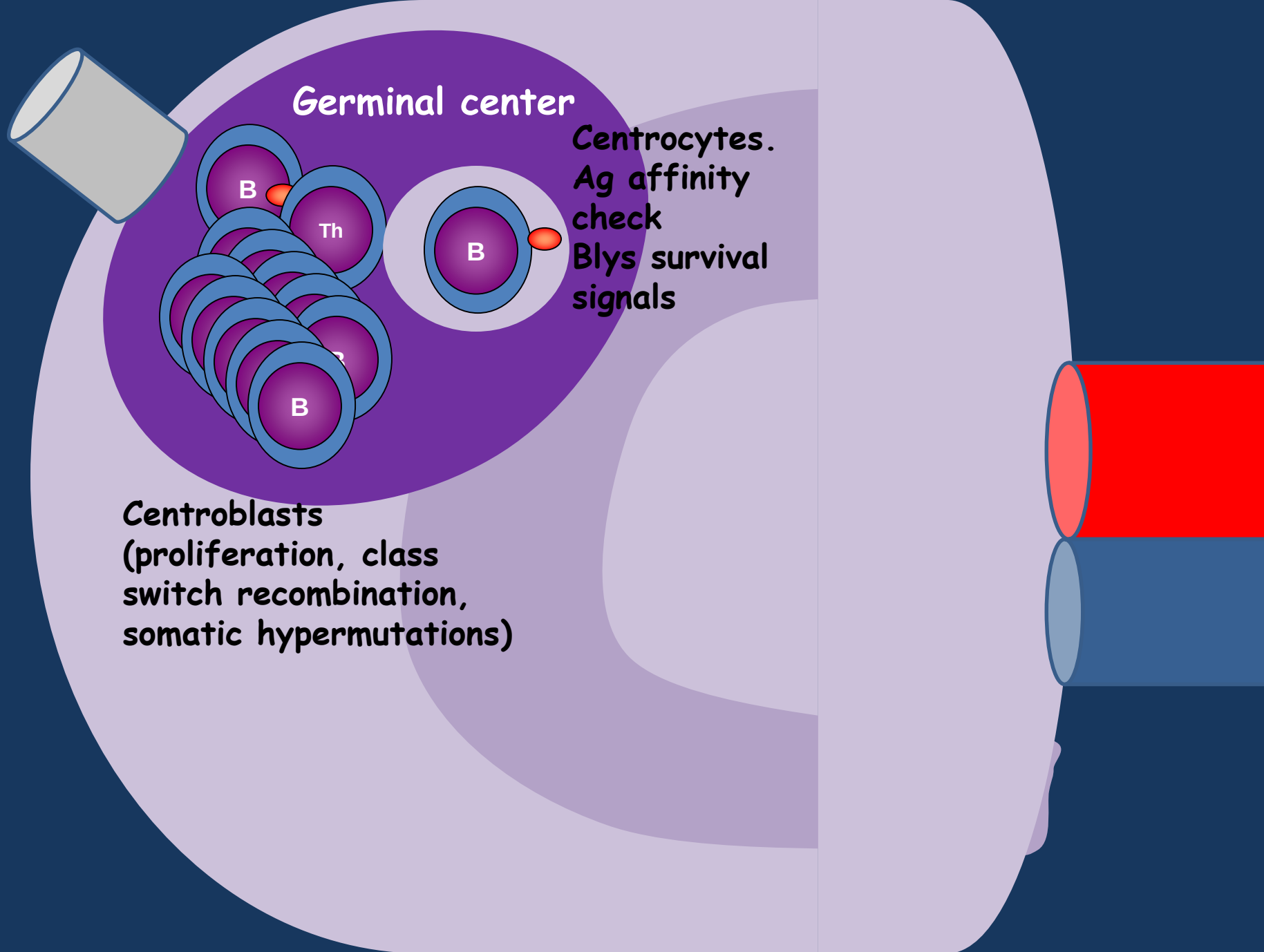
Intermediate affinity
+ IL21 signals





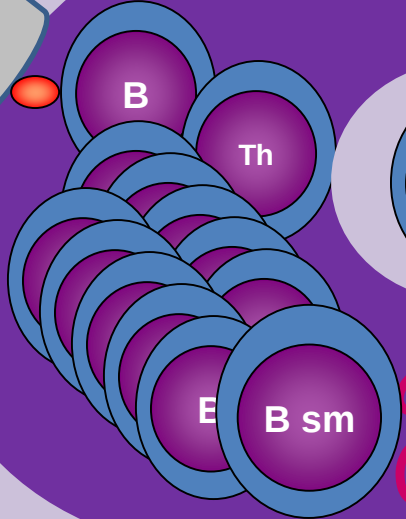
Intermediate affinity
+ IL21 signals





Germinal center

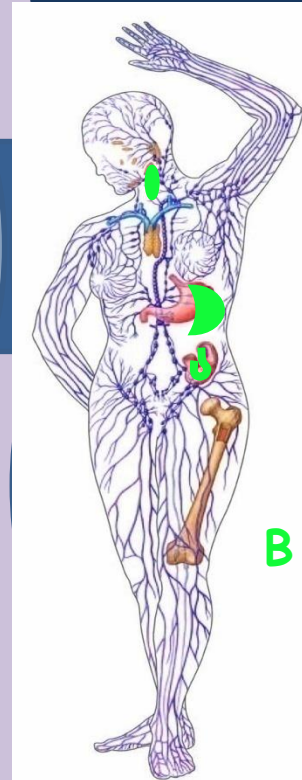
Centrocytes.
Ag affinity
check
Bly s
signal.



CD10+, CD19-
(survival)

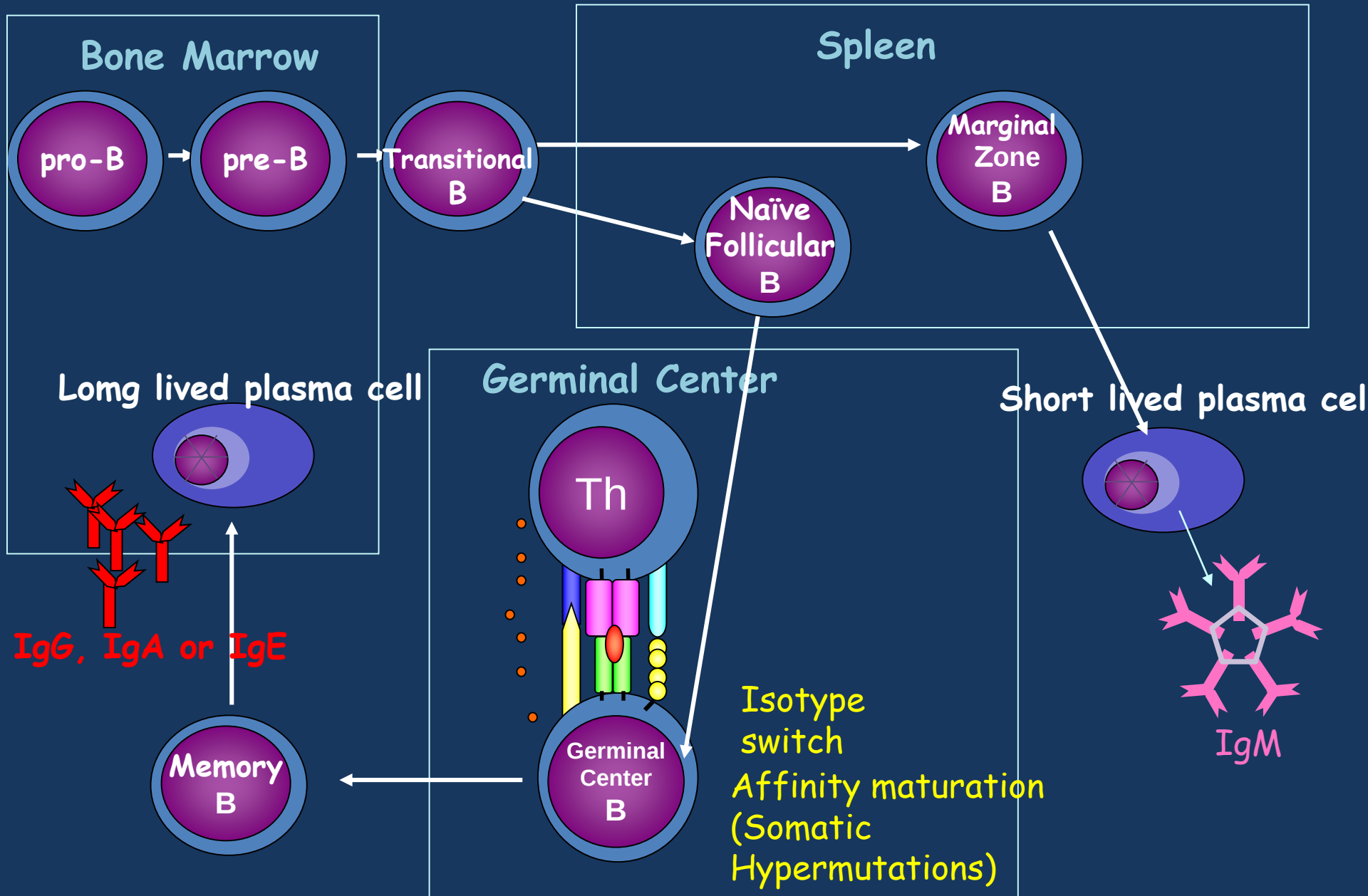
Centroblasts
(proliferation,
class switch
recombination,
somatic hypermutations)

CD27+ IgD-IgM-

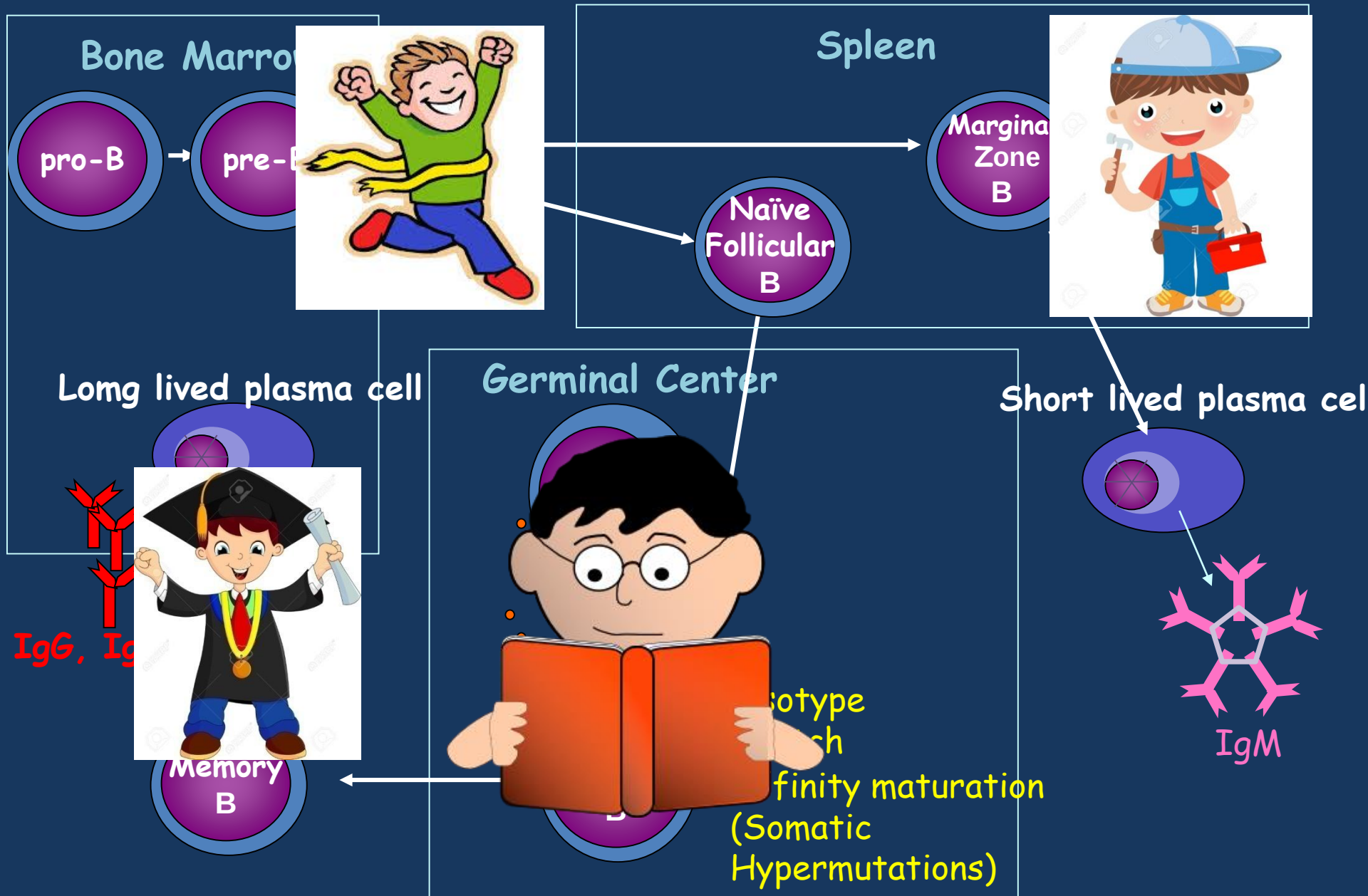


B

Basic B cell development pathway



Basic B cell development pathway

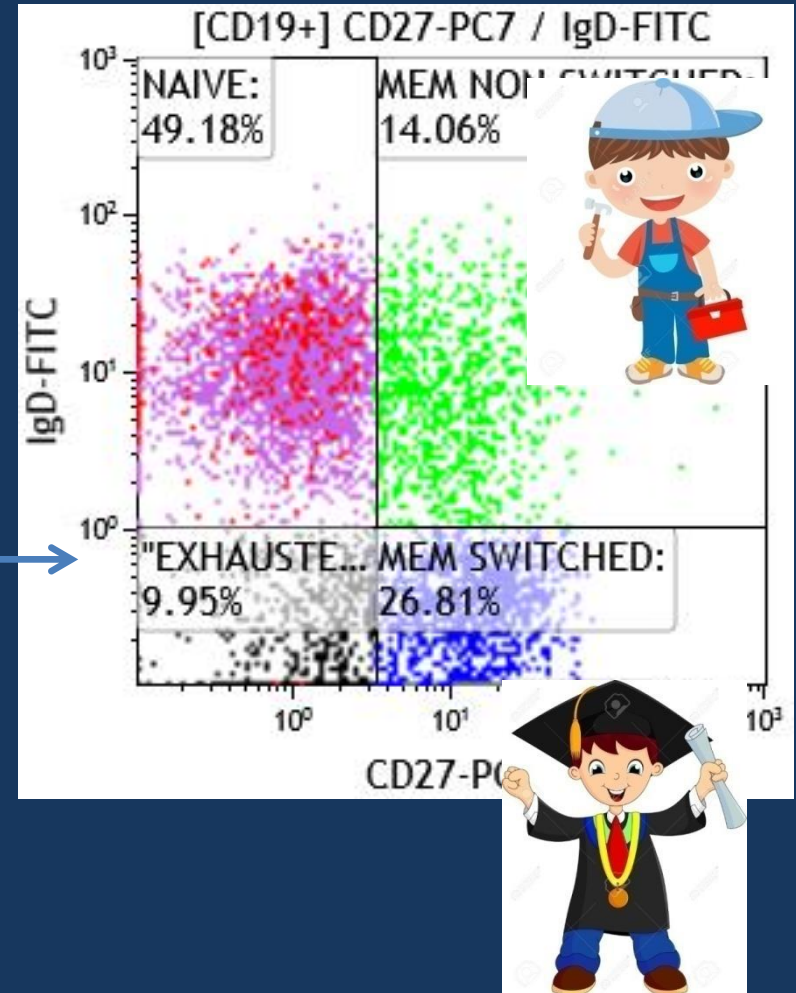
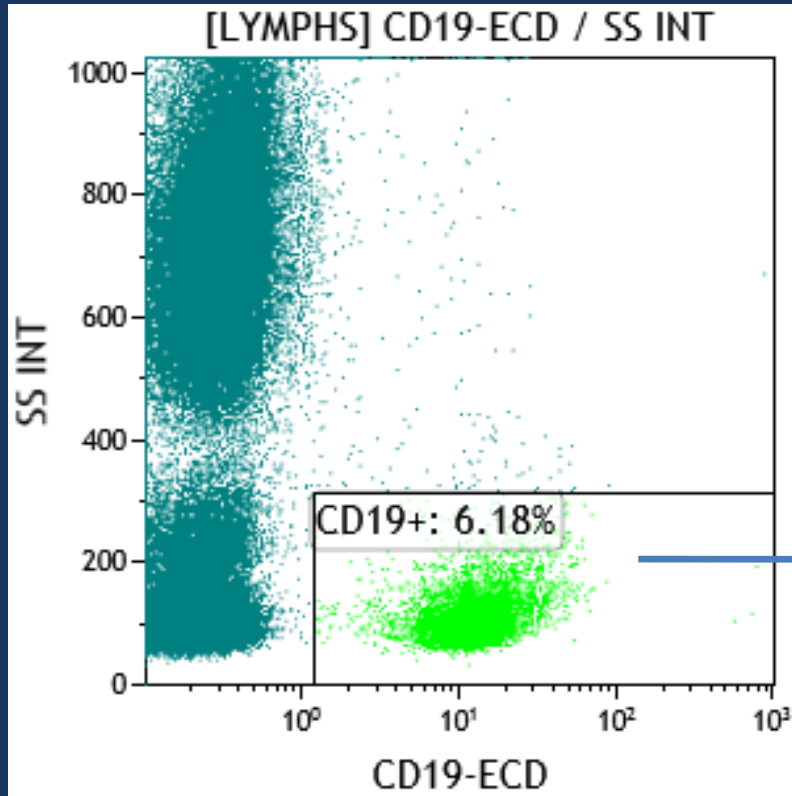


Which markers are useful
to identify these subsets??

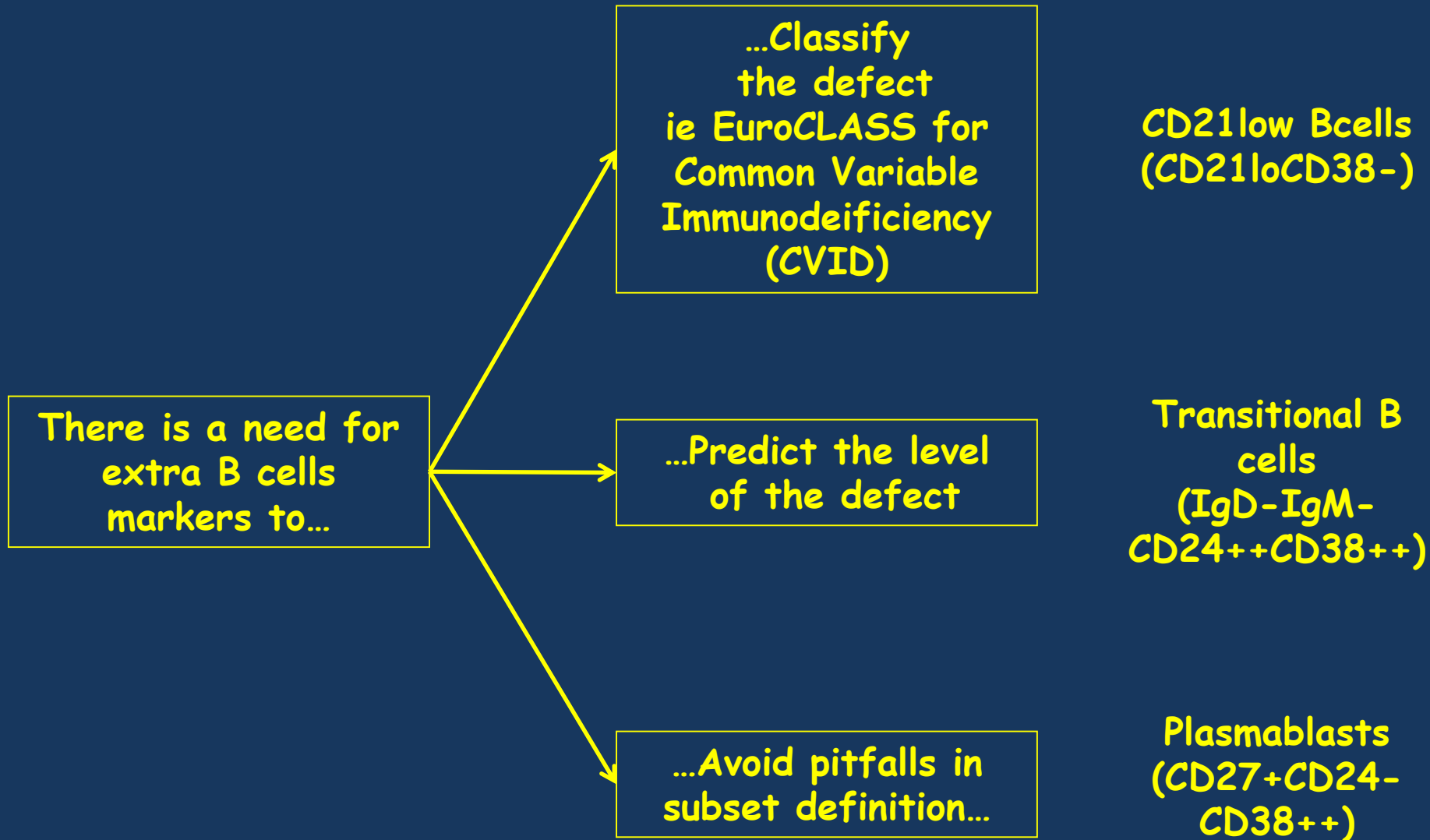
Basic immunophenotypic markers of Peripheral Blood B cell characterization

Marker	Role	B cell subset					
		 Activational	 Naïve Follicular	 Memory Switched	 Exhausted	Plasmablasts	 Marginal Zone like
CD19	forms complex with CD21 and CD81-co stimulation	+	+	+	+	+low	+
CD21	Complement Receptor-Co receptor	med	+	+	 low	med	 ++
CD27	TNF-R family-co stimulation	-	-	 +	-	 ++	 +
CD23	FcεR-IgE coated Ag capture in GC	-	 +	-	-	-	-
CD24	GPI anchored-Cell adhesion +...	 ++	 +	 ++		-	 ++
IgM		++	low/+	-	-	-	++
IgD		low	++	-	-	-	low
CD38	cyclic ADP ribose hydrolase-???	++	+	+		+++	-

Basic gating strategies on CD19+ cells (I)

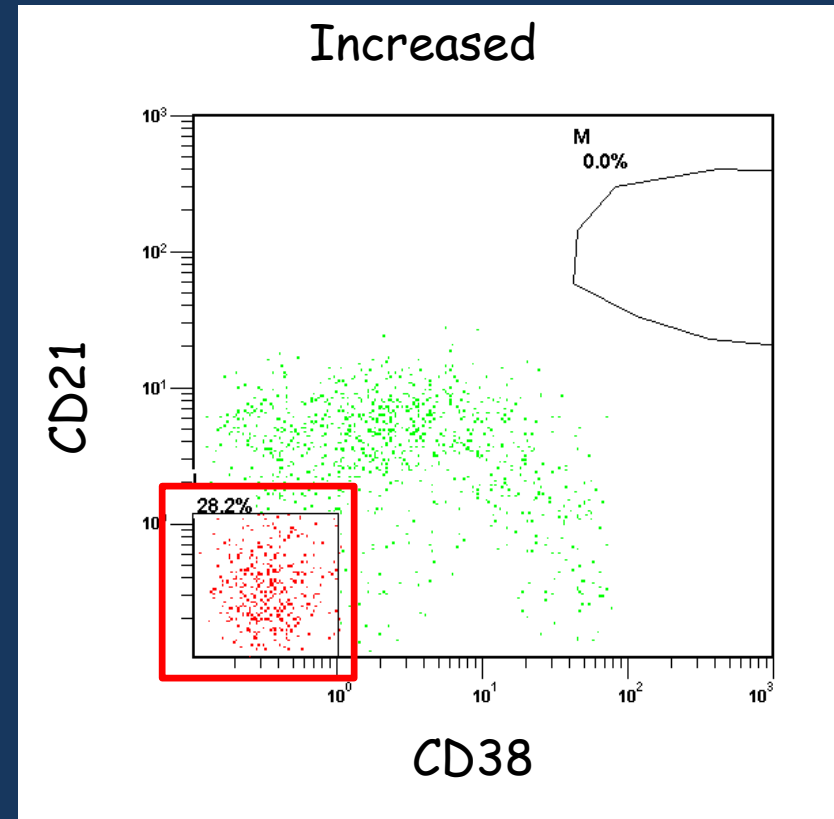
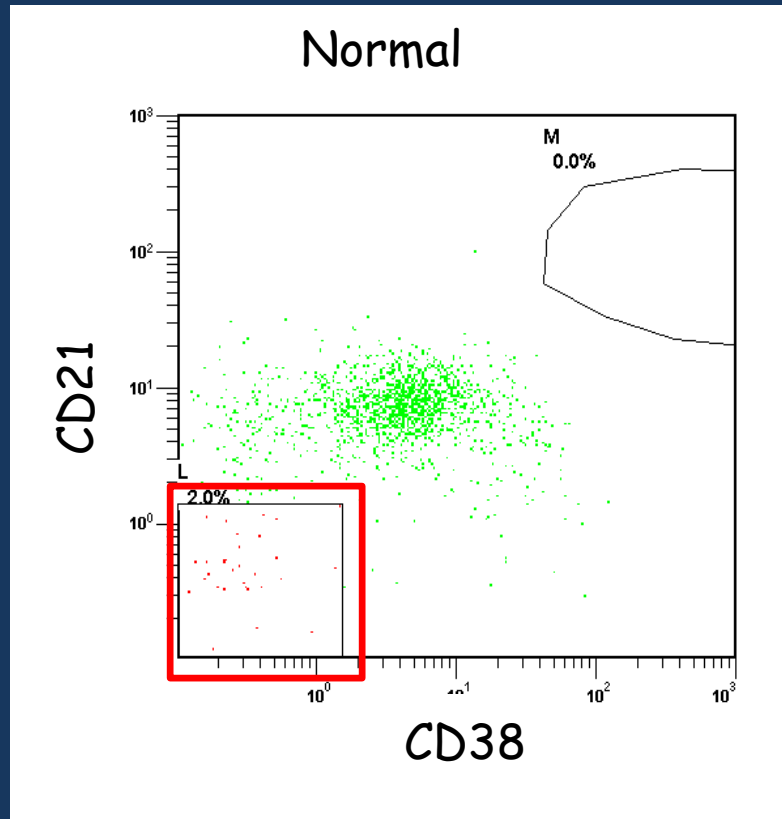


Is this enough??



Basic gating strategies on CD19+ cells (II)

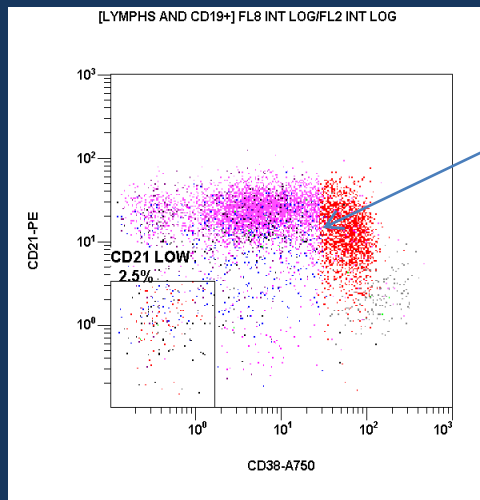
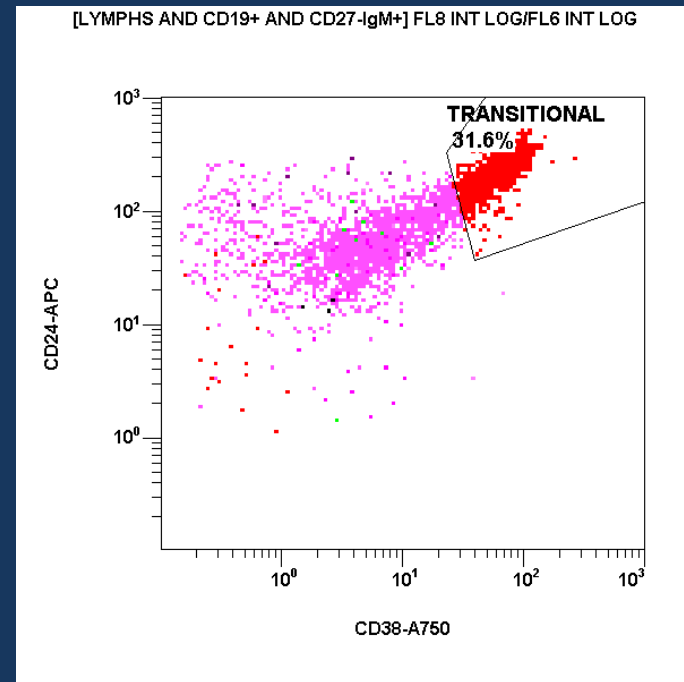
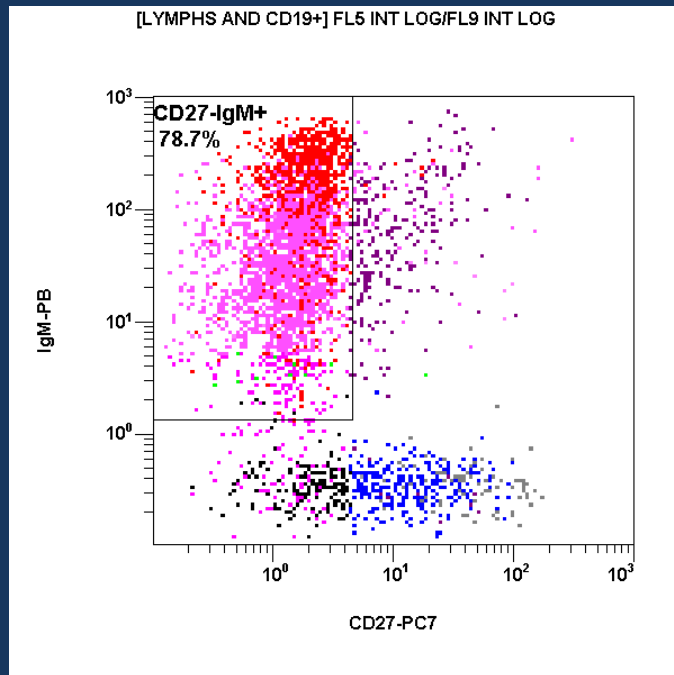
CD21^{low}



CD19⁺

Basic gating strategies on CD19+ cells (III)

Transitional B cells

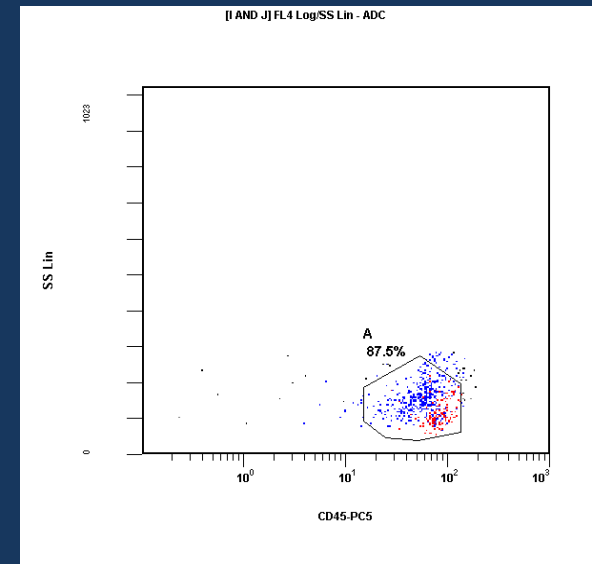
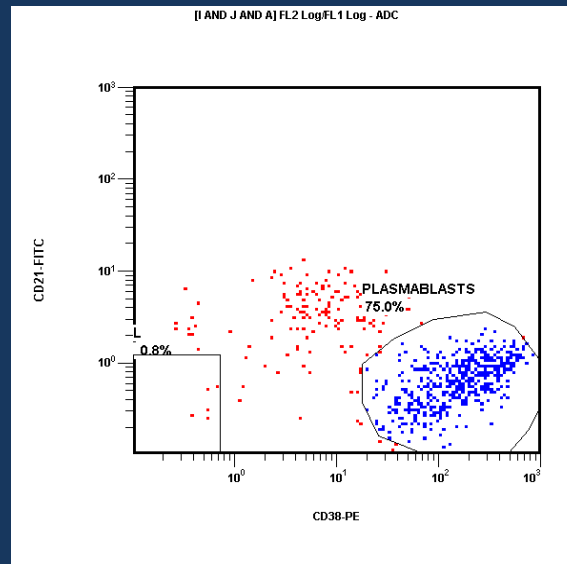
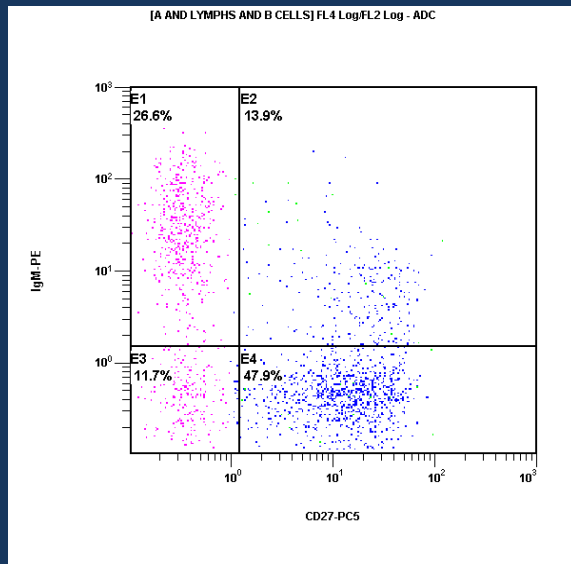


HELPS DEFINE
TRANSITIONAL
REGION

Gate	Number	Logic
All	399,696	Ungated
LYMPHS	66,565	LYMPHS
CD19+	7,915	CD19+ AND LYMPHS
CD27-IgM+	6,492	CD27-IgM+ AND CD19+ AND LYMPHS
TRANSITIONAL	1,788	TRANSITIONAL AND CD27-IgM+ AND CD19+ AND LYMPHS

Basic gating strategies on CD19+ cells (IV)

a quite common pitfall

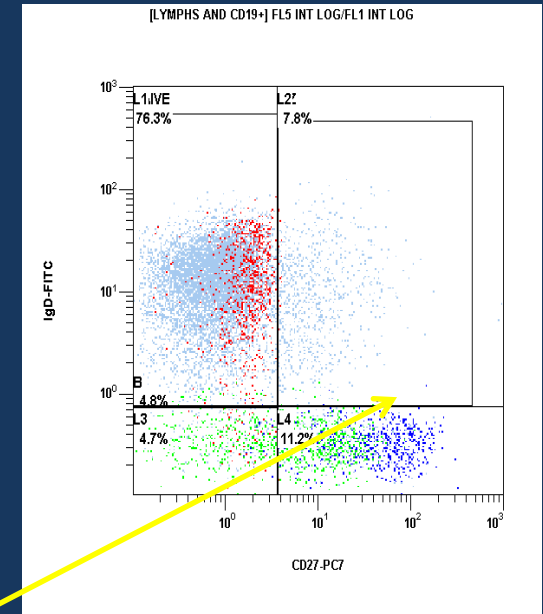
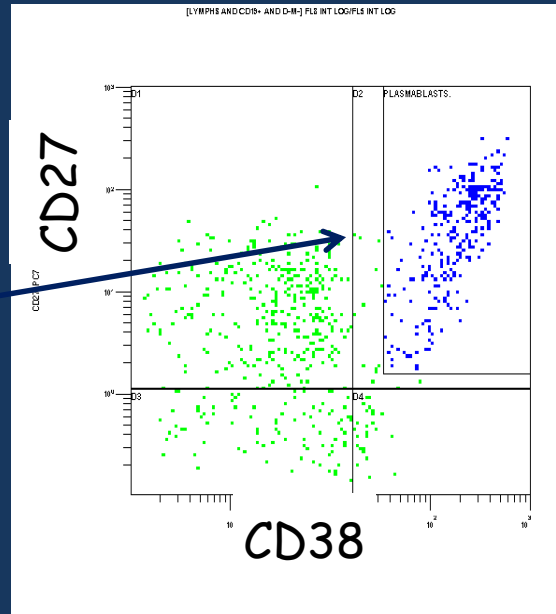
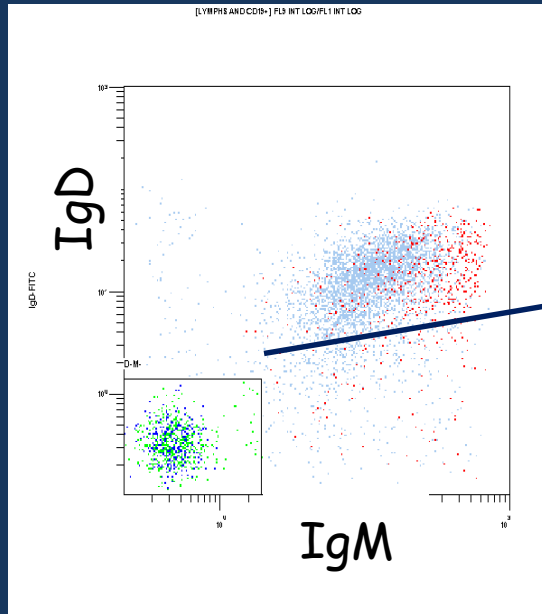


Does this patient have so many memory switched B cells?

Most of them are plasmablasts!

Basic gating strategies on CD19+ cells (IV)

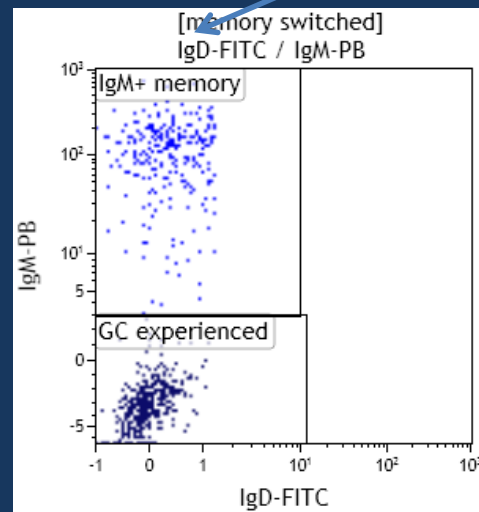
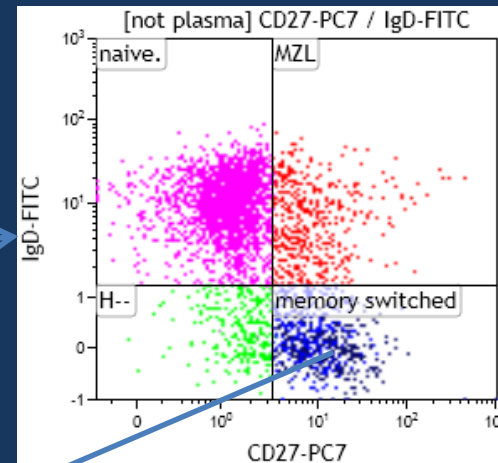
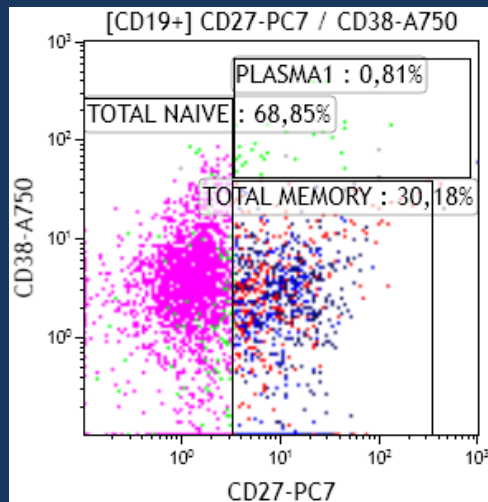
Plasmablasts



They increase the % of the Memory switched region

Basic gating strategies on CD19+ cells (IV)

Exclude Plasmablasts when estimating memory B cells



Be careful
with normal values!

B cell subset values for adults

Age and Gender Leucocytes Variances and References Values Generated Using the Standardized ONE-Study Protocol

Anders H. Kverneland,^{1,2} Mathias Streitz,¹ Edward Geissler,³ James Hutchinson,³ Katrin Vogt,¹ David Boës,¹ Nadja Niemann,¹ Anders Elm Pedersen,² Stephan Schlickeiser,^{1†} Birgit Sawitzki^{1†*}

ALL CELL SUBSETS FIRST COHORT			MEDIAN (RANGE)	
DESCRIPTION	PHENOTYPE	DENOMINATOR	PROPORTIONAL (%)	ABSOLUTE (CELLS/ μ L)
B cells	CD19 ⁺ CD3 ⁻	Lymphocytes	11.3 (3.8–21.5)	205 (51–728)
Naïve B cells	CD19 ⁺ CD27 ⁻ IgD ⁺	B cells	53.6 (17.9–75.1)	43 (5–401)
Transitional B cells	CD19 ⁺ CD27 ⁻ CD38 ^{high} IgM ⁺ CD24 ⁺	B cells	2.1 (0.1–6.3)	2 (0–20)
MZ B cells	CD19 ⁺ CD27 ⁺ IgD ⁺	B cells	12.5 (3.8–35.6)	8 (1–84)
Memory B cells	CD19 ⁺ CD27 ⁺ CD38 ^{dim}	B cells	21 (11–46.6)	18 (3–80)
Class-switched B cells	CD19 ⁺ CD27 ⁺ CD38 ^{dim} IgD ⁻ IgM ⁻	B cells	15.9 (4.6–35.5)	14 (1–53)
Non-switched B cells	CD19 ⁺ CD27 ⁺ CD38 ^{dim} IgM ⁺	B cells	6 (1.9–23.7)	5 (1–28)
Plasmablasts	CD19 ⁺ CD27 ^{high} CD38 ^{high} IgD ⁻ IgM ⁻	B cells	1.2 (0.3–7.8)	1 (0–6)
CD21 ^{low} B cells	CD19 ⁺ CD38 ^{low} CD21 ^{low}	B cells	6.2 (1.9–19.3)	5 (1–37)

OMIP-047: High-Dimensional Phenotypic Characterization of B Cells

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Grant sponsor: Swiss National Science Foundation, Grant number: SNF; #314730_152663 and #314730_172790

Grant sponsor: Clinical Priority Research Program of the University of Zurich (Viral infectious diseases: Zurich Primary HIV

PURPOSE AND APPROPRIATE SAMPLE TYPE

THIS 16-color, 18-parameter panel was designed to allow a detailed dissection of human B cell subsets and their phenotype in peripheral blood mononuclear cells (PBMC) in healthy donors and in the context of chronic viral diseases such as Human Immunodeficiency Virus 1 (HIV-1) infection. The panel encompasses a range of backbone markers for the accurate definition of common B cell subsets with a focus on memory B cells and a unique collection of phenotypic markers (chemokine receptors, cytokine receptor, B cell receptor isotypes, and proliferation marker) not combined in multicolor flow cytometry B cell phenotyping thus far. This new panel allows highly detailed phenotypic and functional investigations of B cell subsets. The panel was validated using cryopreserved PBMC from healthy and HIV-1 infected donors allowing the retrospective analysis of clinical samples (Table 1).

Review Article**Flowcytometric Phenotyping of Common Variable Immunodeficiency****Klaus Warnatz* and Michael Schlesier**

Division of Rheumatology and Clinical Immunology, University Medical Center Freiburg, Freiburg, Germany

Table 3
Reference Values of B-Cell Subpopulations in Peripheral Blood

B-cell population		Reference range ^a (%)
CD19+ in CD45+ lymphocytes	B cells	4.9–8.4
IgM– IgD– in CD19+ B cells	Class switched B cells	7.6–31.4
IgD+ CD27– in CD19+ B cells	Naive B cells	42.6–82.3
IgD+ CD27+ in CD19+ B cells	Marginal zone-like B cells	7.4–32.5
IgM–IgD– CD27+ in CD19+ B cells	Class switched memory B cells (and Plasmablasts)	6.5–29.1
CD21 ^{low} CD38– in CD19+ B cells	CD21 ^{low} B cells	0.9–7.6
M++ D++ CD24++ CD38++ in CD19+ B cells	Transitional B cells	0.6–3.4
IgM-(+) CD24-CD38+++CD27++ in CD19+ B cells	Plasmablasts	0.4–3.6

^a5. to 95. percentile, based on 54 healthy donors (age range 19–61 years).

B cell subset values for children

(Beware of subset definitions!!)

Clinical Immunology (2009) 131, 50–59



available at www.sciencedirect.com



www.elsevier.com/locate/yclim



Memory B-cells in healthy and antibody-deficient children

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Sven Bellert^a, Tim Niehues^{b,*}

^a Department of Pediatric Oncology, Hematology and Clinical Immunology, Centre for Child and Adolescent Health, Heinrich Heine University of Düsseldorf, Moorenstrasse 5, 40225 Düsseldorf, Germany

^b Centre for Child and Adolescent Health, HELIOS Klinikum Krefeld, Academic Hospital, Heinrich Heine University of Düsseldorf, Lutherplatz 40, 47805 Krefeld, Germany

Received 7 August 2008; accepted with revision 19 November 2008

Available online 21 January 2009

CD27-IgD ⁺ in %CD20	88	85	78.5	78	74.5
naive	88.4	84.2	77.8	77.03	75.27
	(84.5–93)	(77.4–90)	(70.7–85)	(66.1–87.6)	(63.3–87.9)
	(83.25–93.75)	(74.7–90.5)	(89.9–85.6)	(63.1–89.15)	(60.15–88.95)
CD27-IgD ⁺ in %CD20	7	8	9.5	10.75	11
non-switched memory	6.81	8.7	10.1	10.76	11
	(4–9.5)	(5–14.3)	(7–14)	(4.7–17)	(6.1–16.9)
	(3.25–10.75)	(4.9–14.2)	(7–15.2)	(2.93–19)	(5.05–17.95)
CD27-IgD ⁻ in %CD20	3	5	8	8	9
switched memory	3.03	5.2	8.6	8.72	10.32
	(1.35–5)	(3–8)	(5–12.3)	(4–14)	(4.1–18.7)
	(1–5)	(2.9–9.2)	(3.9–16.2)	(3.85–16.5)	(4–22.8)

Which molecules are essential
for the process??

Basic B cell development pathway

Bone Marrow

pro-B



pre-B

Spleen

Germinal Center

V

D

J



1 2 3 4 5 6 7 8 9... 1 2 3 4 5 6 7 8 9... 1 2 3 4 5 6 7 8 9...

V

D

J

1 2 3 4 5 6 7 8 9... 1 2 3 4 5 6 7 8 9... 1 2 3 4 5 6 7 8 9...

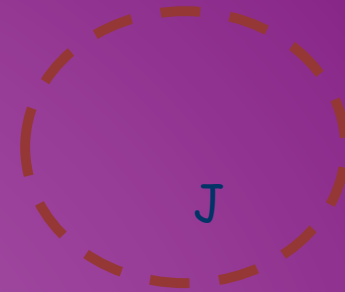
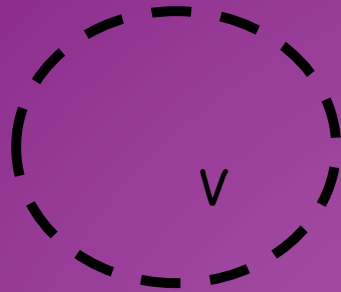
V

D

J

1 2 3 4 5 6 7 8 9... 1 2 3 4 5 6 7 8 9... 1 2 3 4 5 6 7 8 9...

1 2 3 4 5 6 7 8 9... 1 2 3 4 5 6 7 8 9...

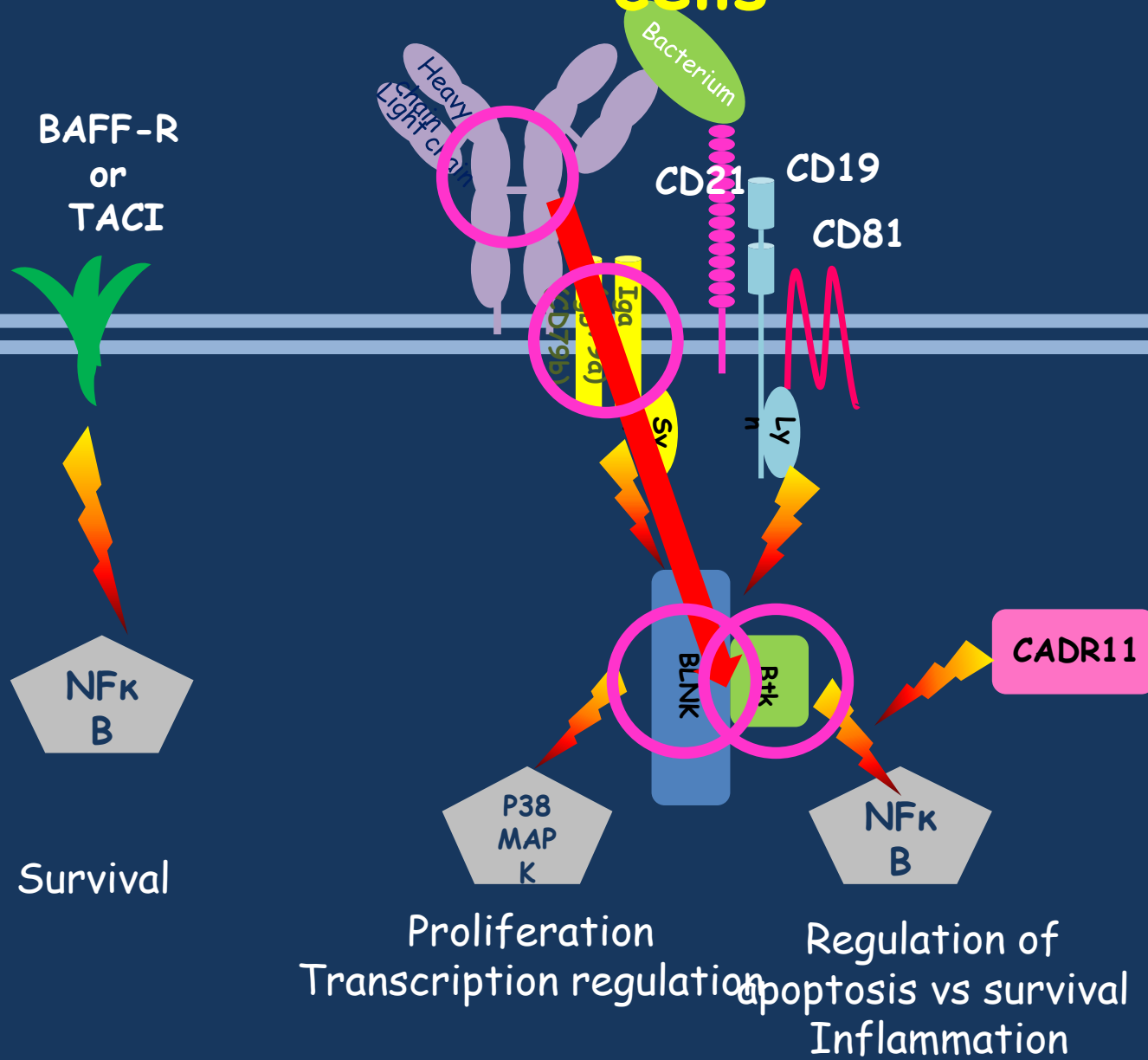


VDJ Recombination

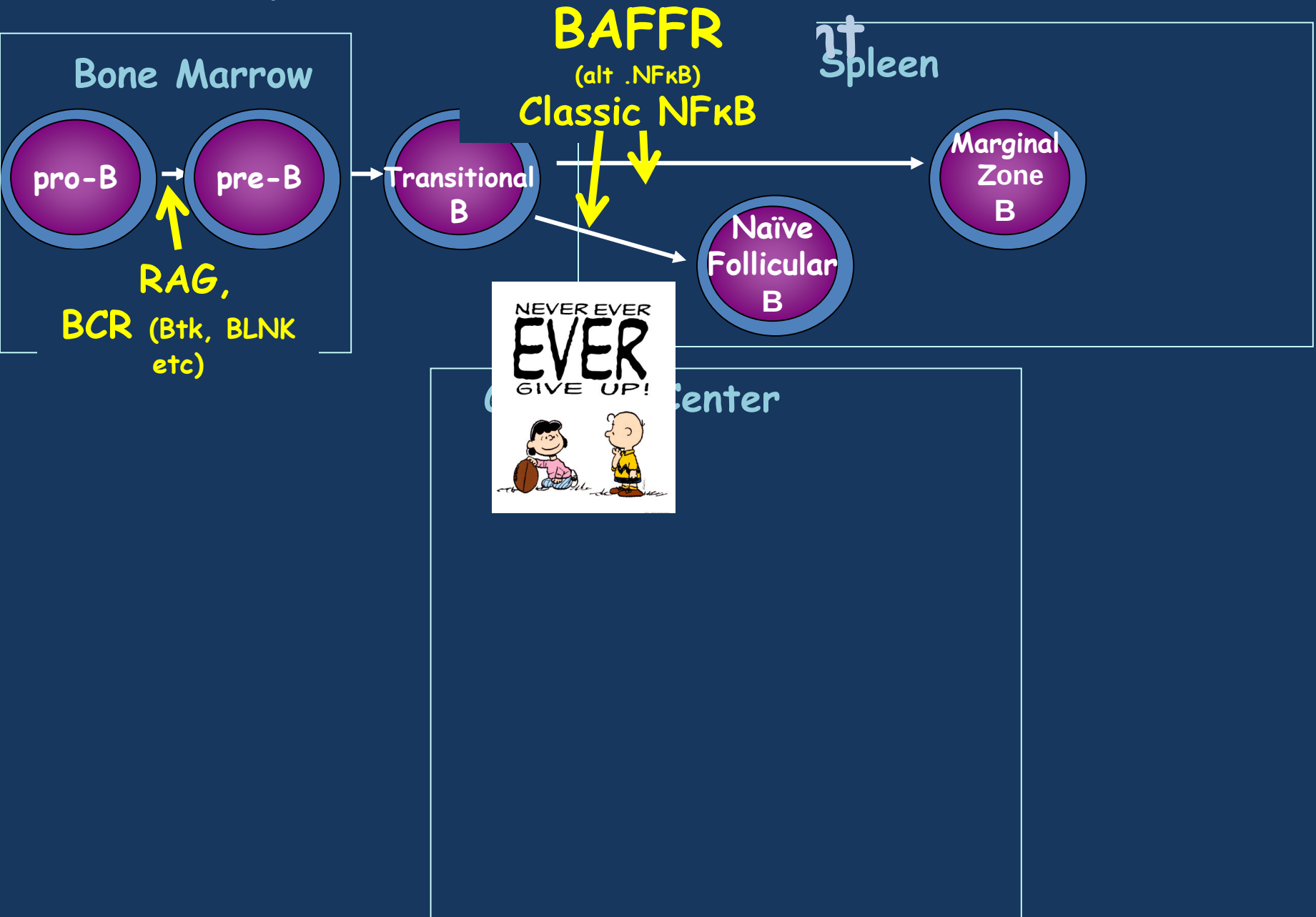
RAG!!!

Artemis, DNAligase IV, DNA PKcs...

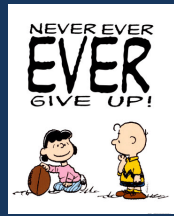
Basic receptors and signal transduction in B cells



Key molecules and signals in B cell



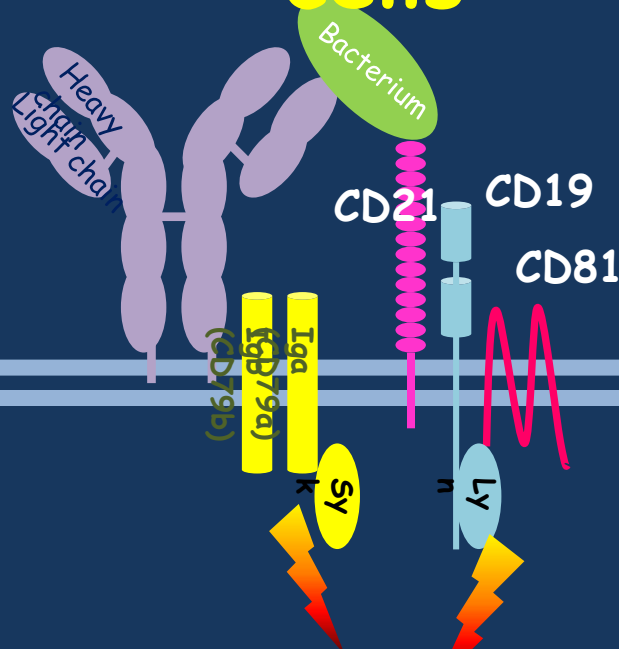
Basic receptors and signal transduction in B cells



BAFF-R
or
TACI

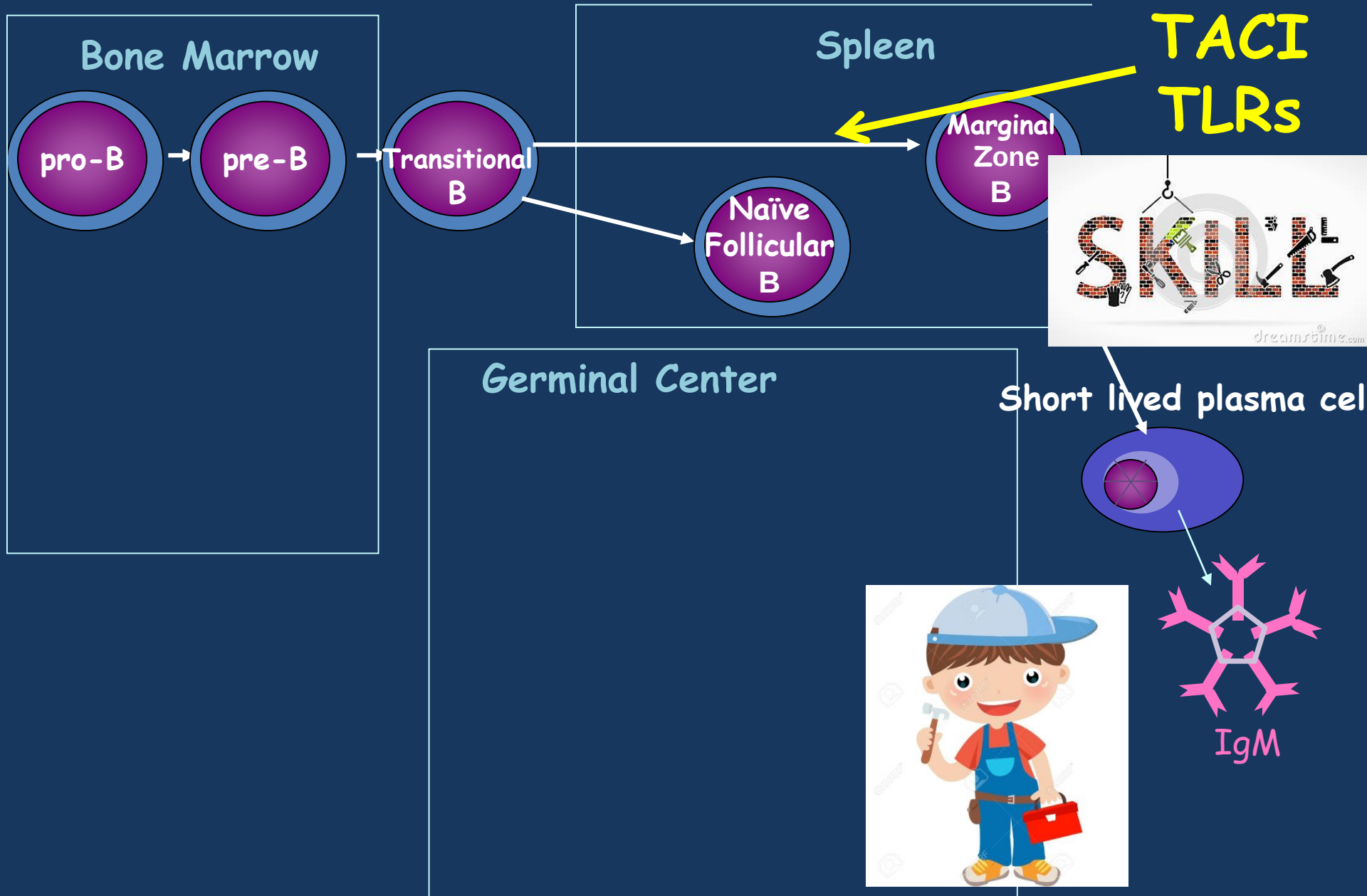


Survival

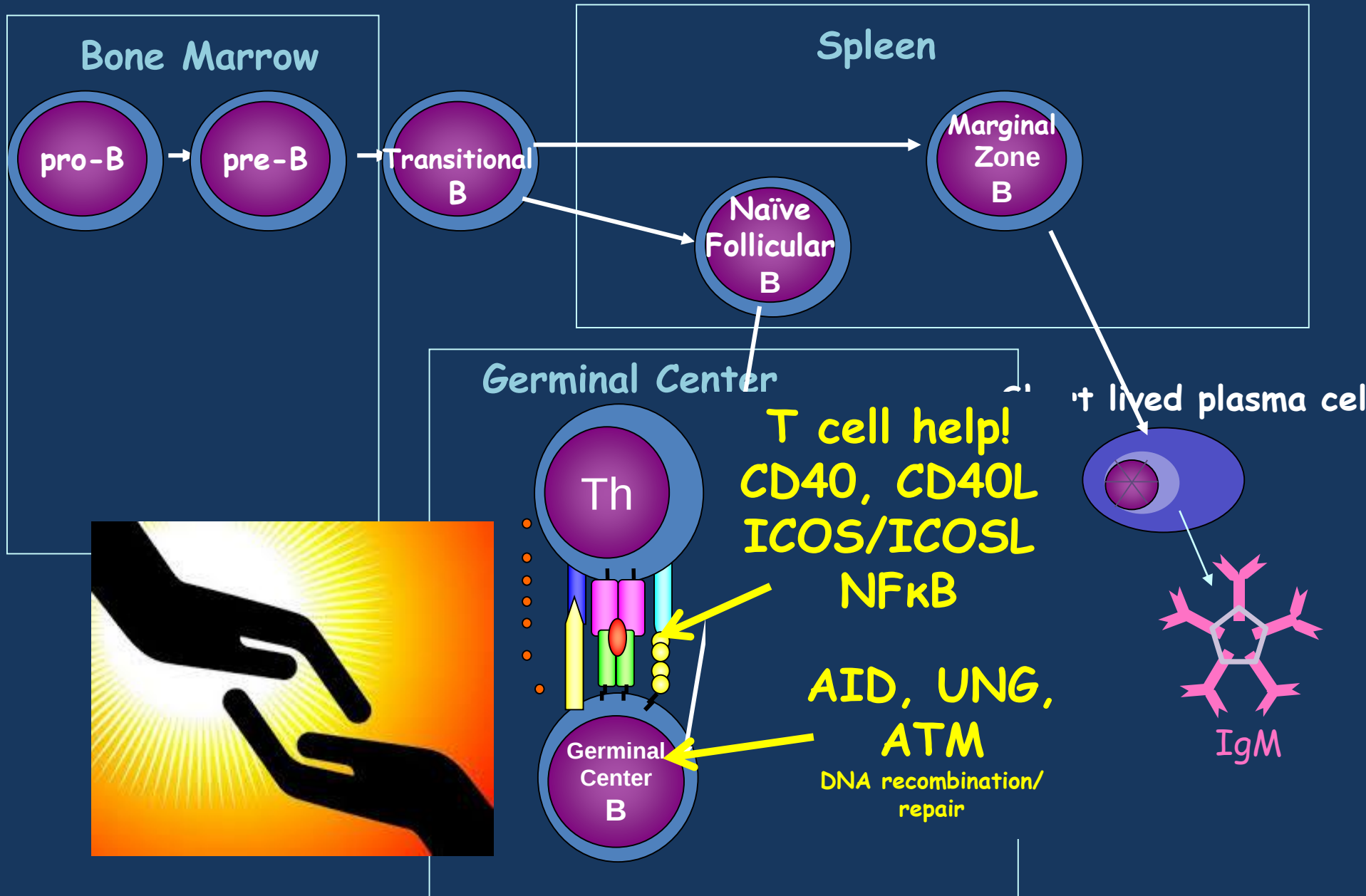


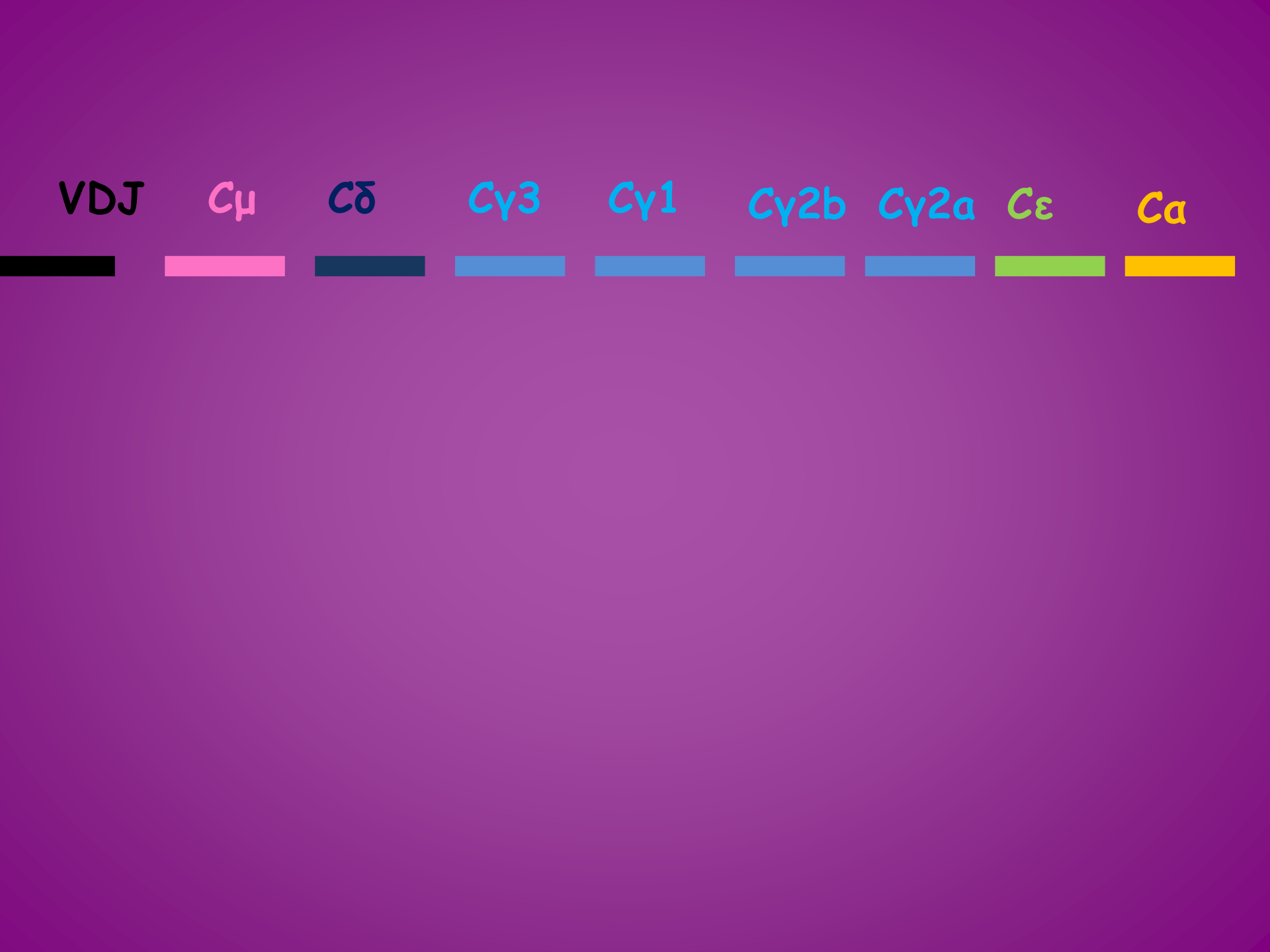
Proliferation
Transcription regulation
Regulation of
apoptosis vs survival
Inflammation

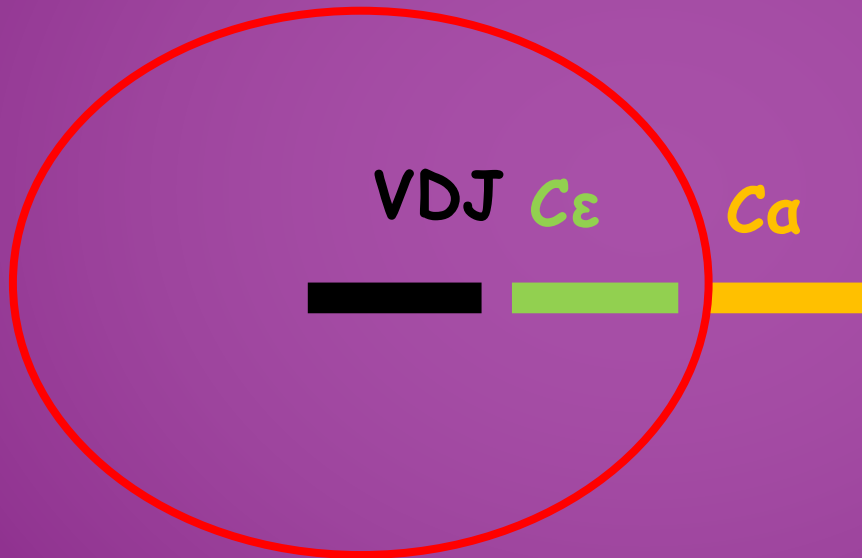
Basic B cell development pathway



Basic B cell development pathway







IgE



CS Recombination

AID!!!

UNG, ATM,

How can we spot the defect ?

No B cells in the Peripheral Blood

Block in

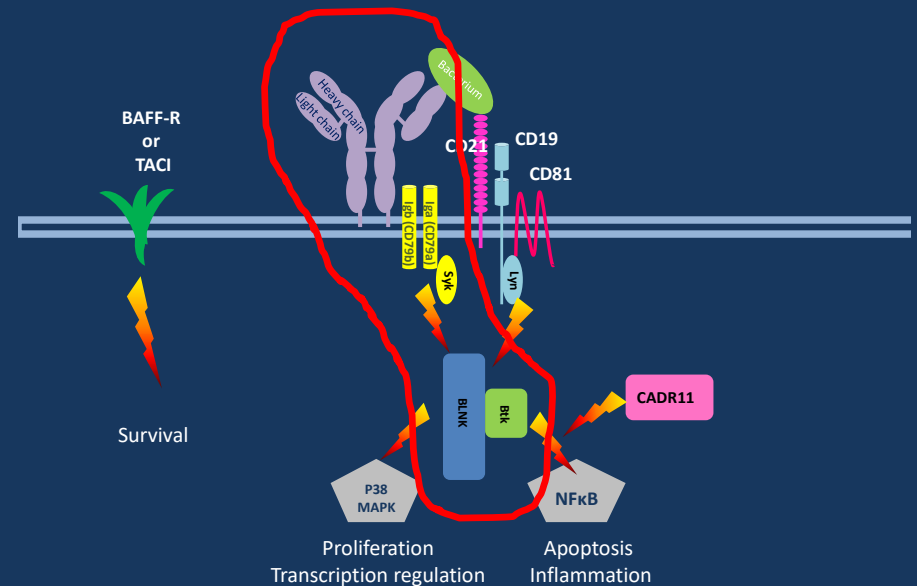
VDJ recombination
(RAG1/RAG2)

NHEJ

(Artemis Cernunnos,
LIG4,
DNA-PKcs)

SCID phenotype

BCR signals



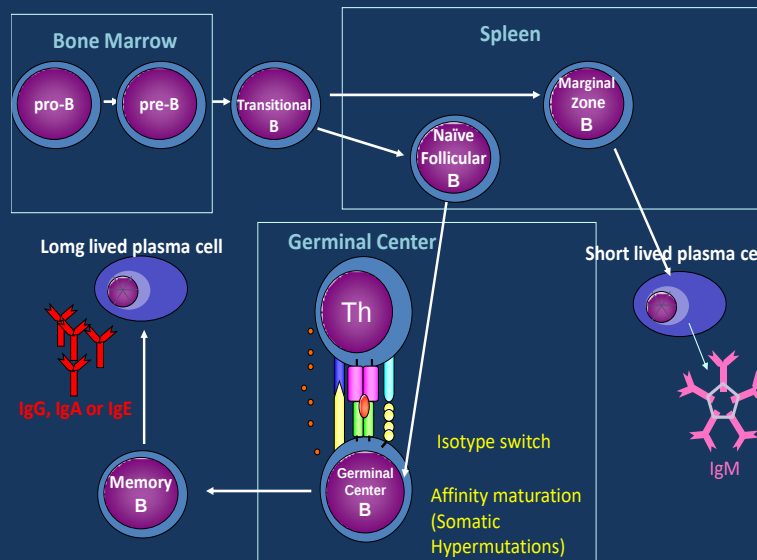
Agammaglobulinaemia

Detectable B cells in the Peripheral Blood with antibody deficiency

B cell subsets can give clues on underlying
mechanisms

5 Patterns have been proposed

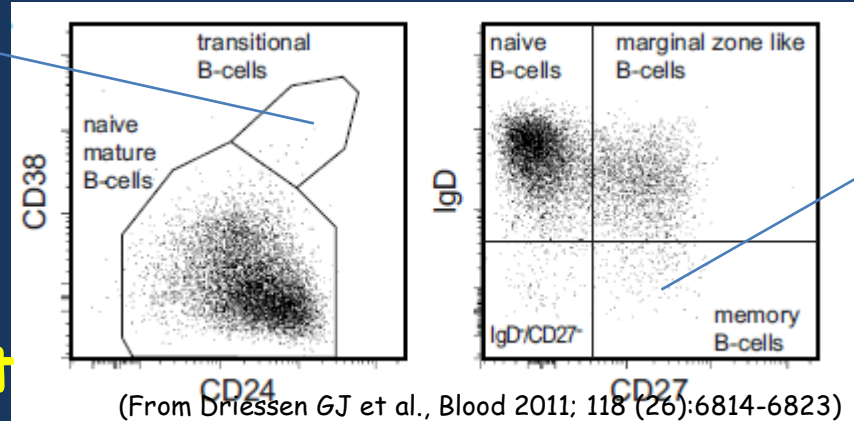
(Driessen GJ et al., Blood 2011; 118 (26):6814-6823)



Pattern 1

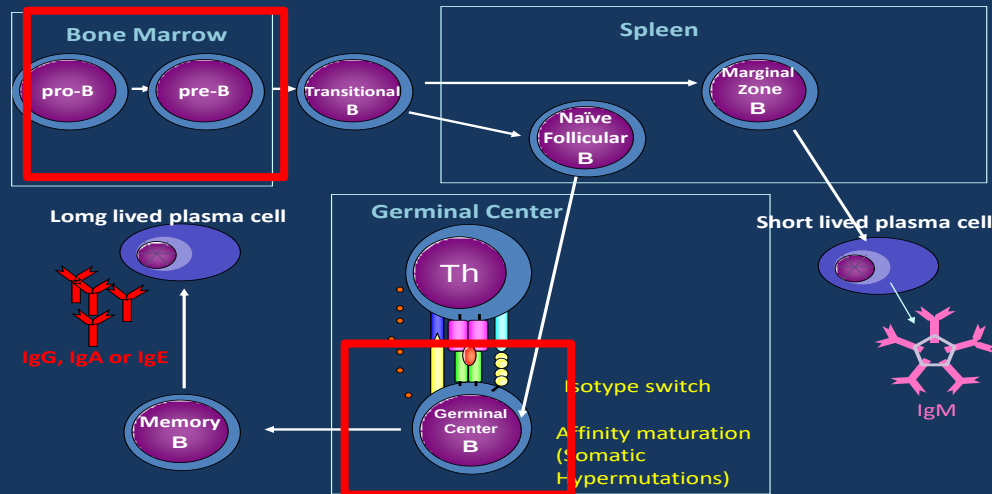
Low
transitional B

Low BM output



Low switched
memory B

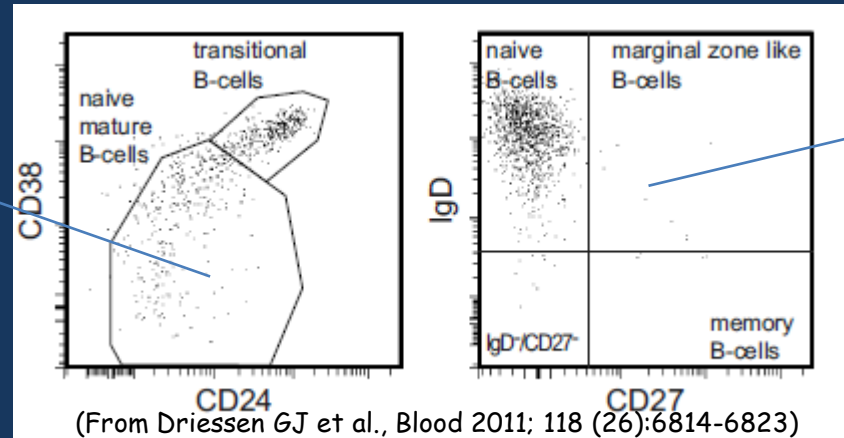
Germinal
center
defect



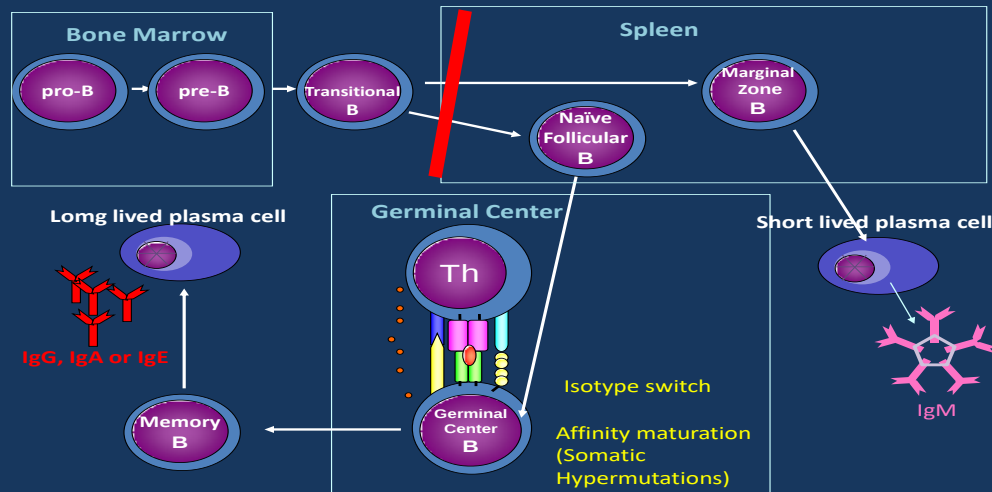
Common mechanism: DNA repair
(eg. Niimengen sy)

Pattern 2

Low % of cells beyond transitional B (incl, naïve)



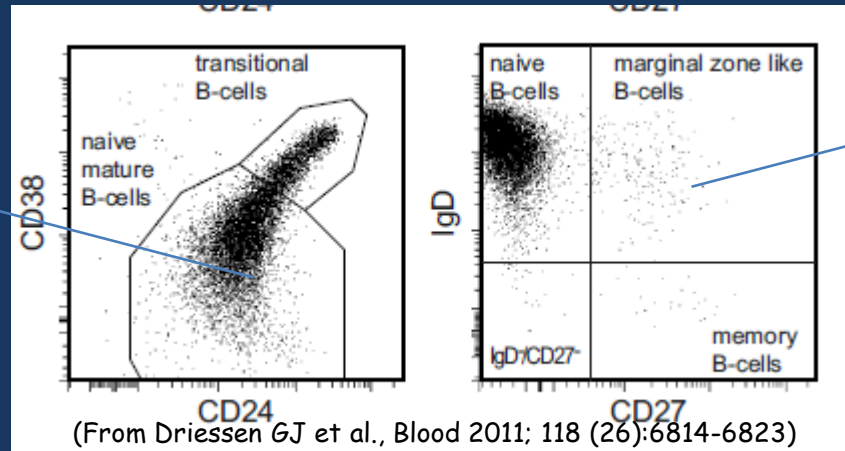
i.e. Low total memory B



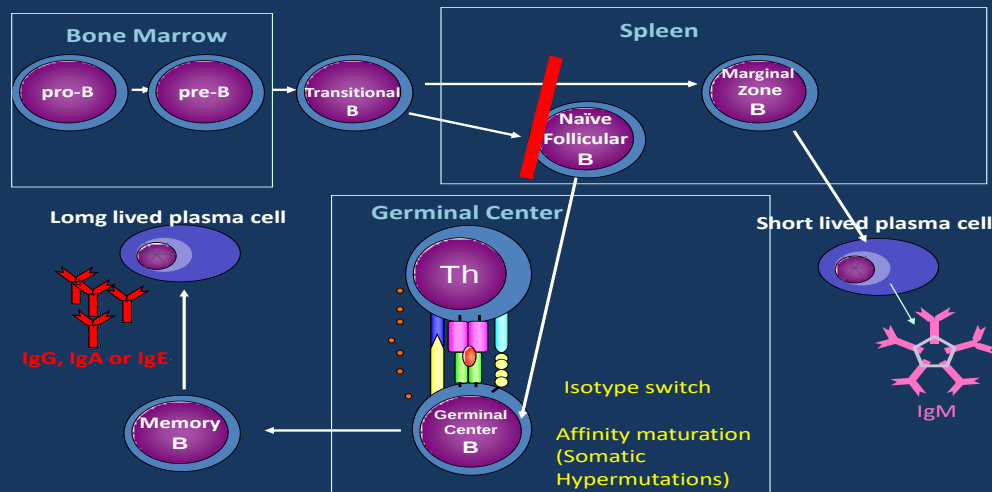
Possible mechanism: lack of survival signals (e.g. BAFFR def. CARD11 def.)

Pattern 3

Normal maturation from transitional B

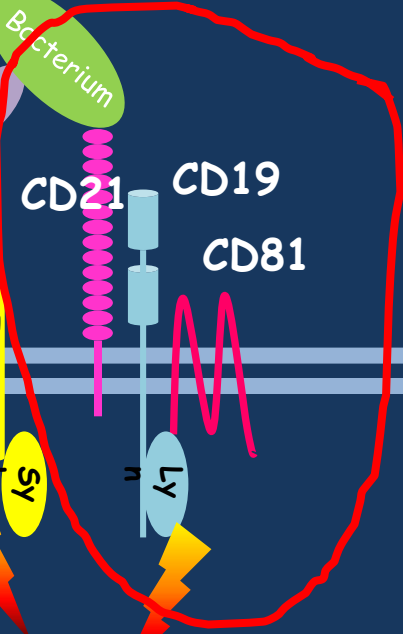


Low total memory B (MZ and switched)



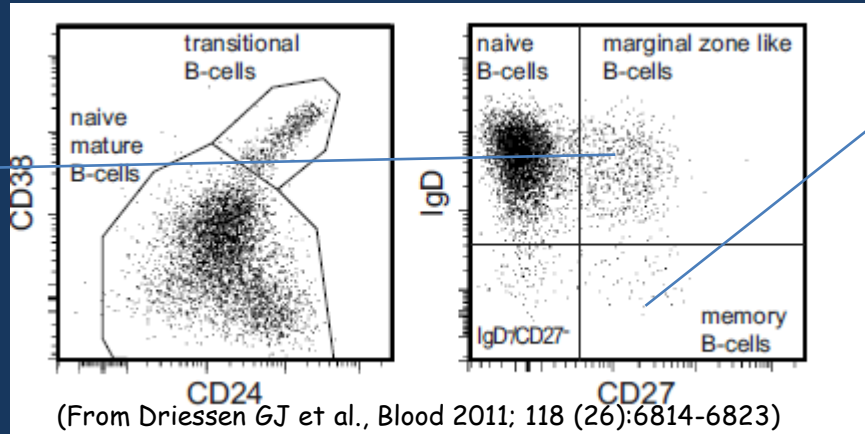
Possible mechanism: Impaired response to Ag:
lack of costimulatory signals
(e.g. CD19, CD81, TLR signal def.)

cells

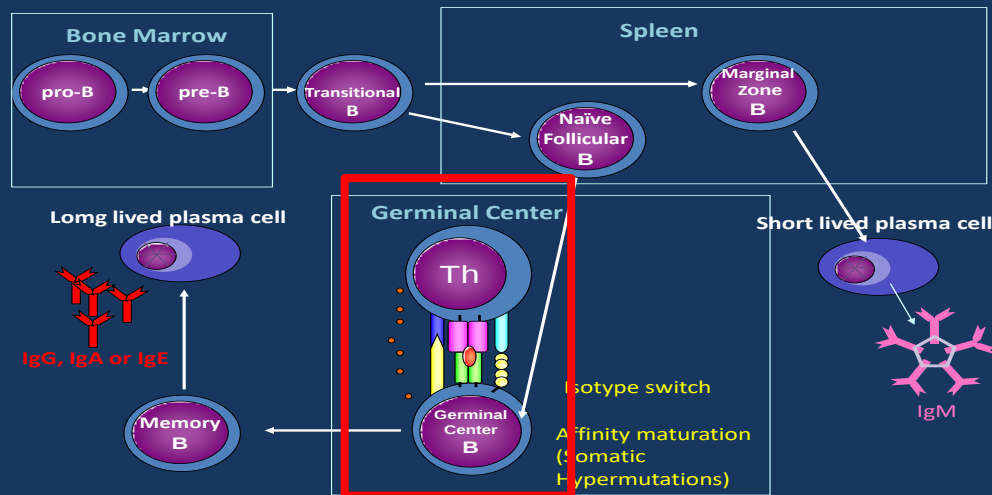


Pattern 4

Normal
MZ
B



Low memory
switched
B

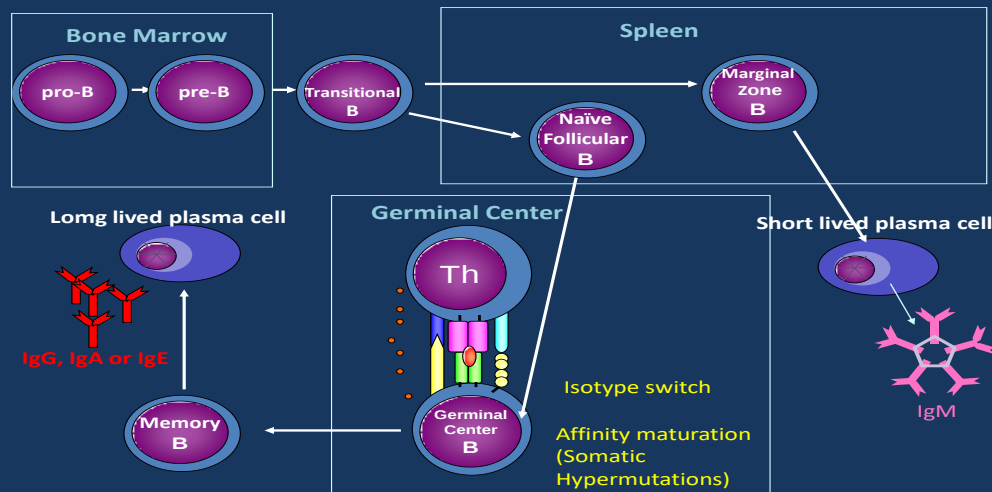
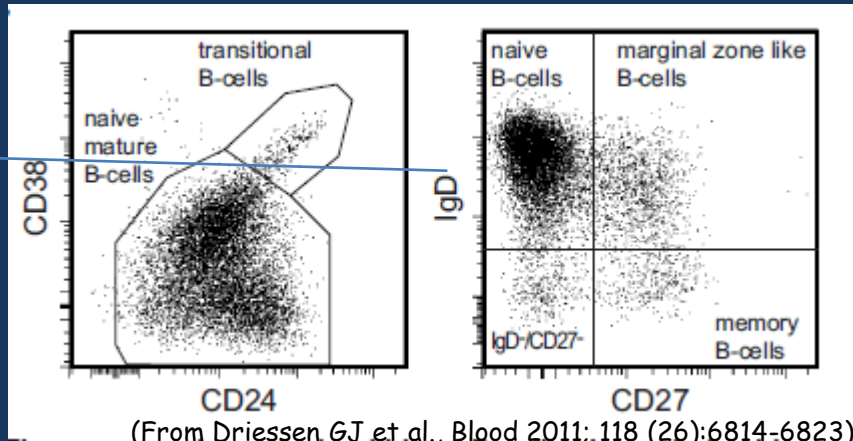


Possible mechanism: Germinal center dysfunction - NOT necessarily CVID -

(T cell deficiency, CD40def., DNA recombination and repair def. (AID, UNG, NEMO), PIK3δ GOF)

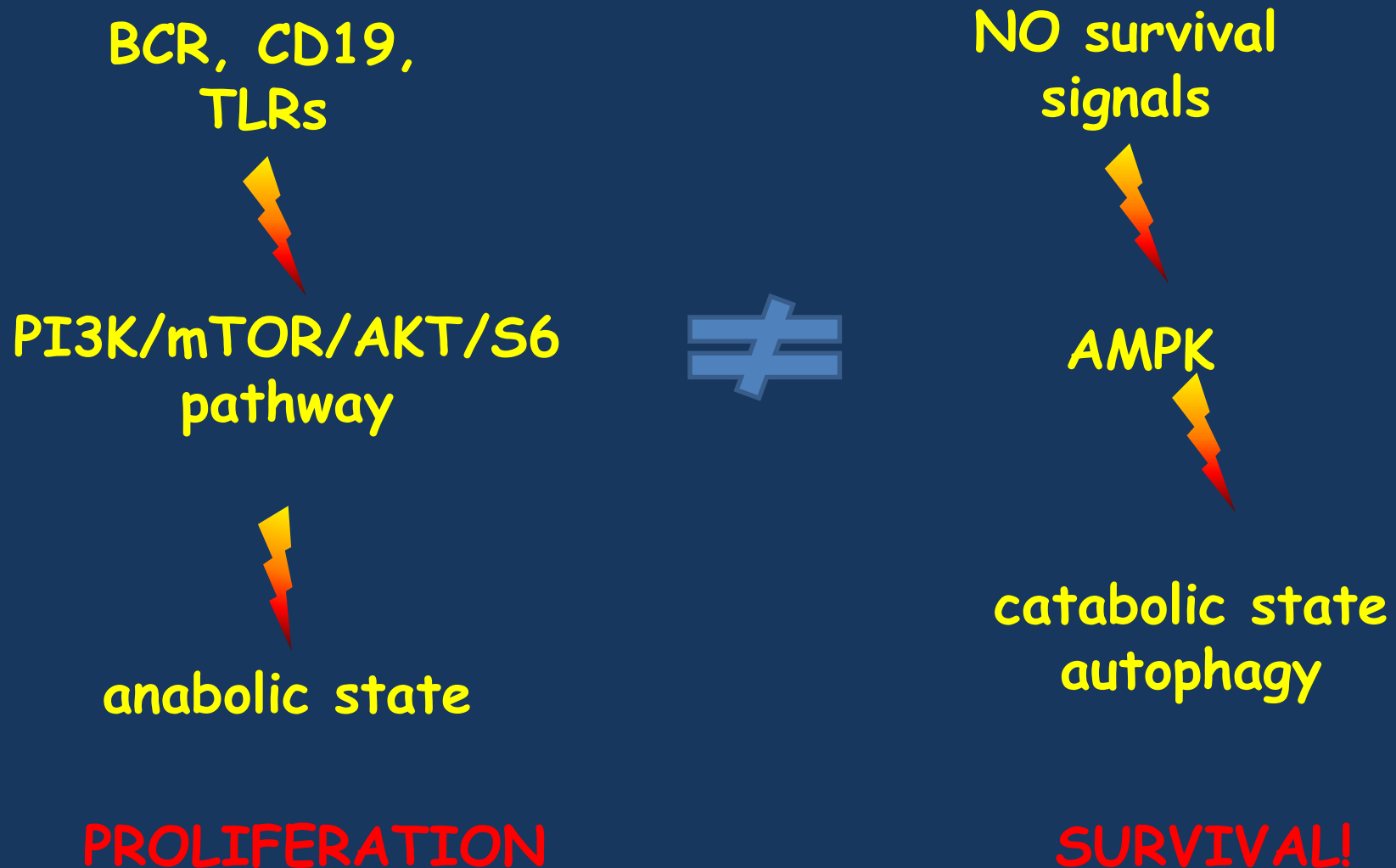
Pattern 5

Normal
B cell
subsets



Possible mechanism: ?
Post GC cell survival?

Generation-survival of memory B cells is correlated to metabolism regulation



Generation-survival of memory B cells is correlated to metabolism regulation

BCR, CD19,
TLRs



PI3K/mTOR/AKT/S6
pathway



NO survival
signals



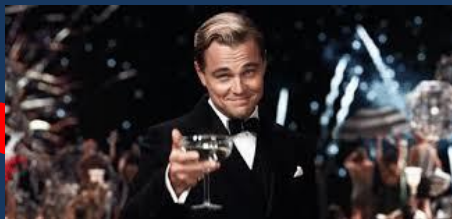
AMPK



anabolic state



PROLIFERATION



catabolic state
autophagy

SURVIVAL!





“Immune TOR-opathies,” a Novel Disease Entity in Clinical Immunology

Sophie Jung^{1,2,3}, Laura Gámez-Díaz^{3†}, Michele Proietti^{3†} and Bodo Grimbacher^{3}*

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TOR-opathies

BCR, CD19,
TLRs



PI3K/mTOR/AKT/S6
pathway



NO survival
signals



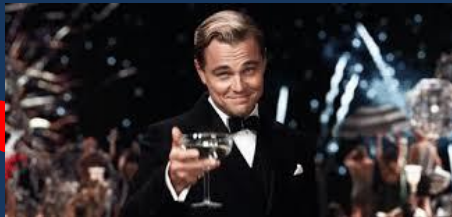
AMPK



anabolic state



PROLIFERATION



catabolic state
autophagy

SURVIVAL!



TORopathies

BCR, CD19,
TLRs



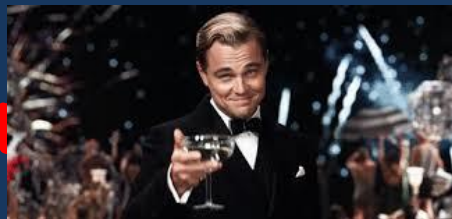
PI3K/mTOR/AKT/S6
pathway



NO survival
signals



anabolic state



PROLIFERATION

anabolic state



NO memory B cells

European Concensus Classification for CVID (EUROclass) 2008

>1% B
cells =
group B+

>2% switched
memory B cells
= group smB+

≤2% switched
memory B cells
= group smB-

≥10%
CD21lo B cells
= group
smB+21 lo

<10%
CD21lo B cells
= group
smB+21norm



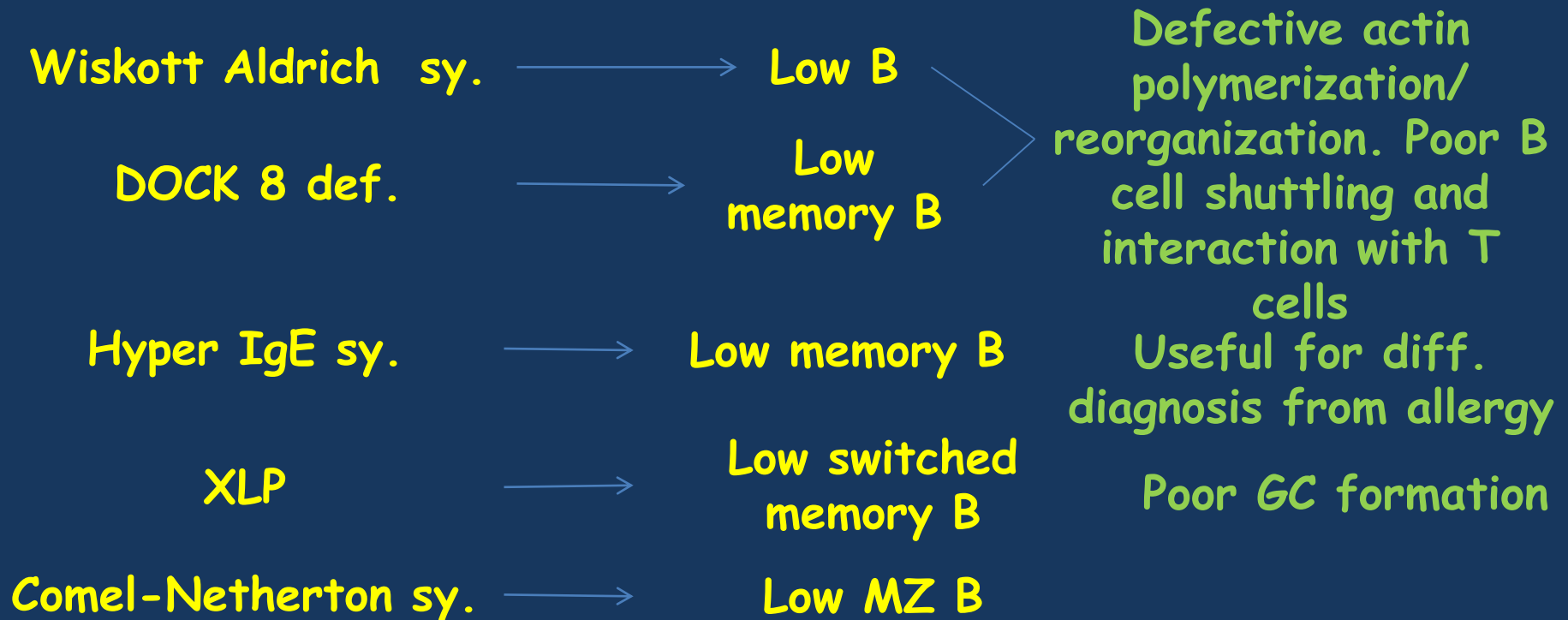
≥10%
CD21lo B cells
= group
smB-21 lo

<10%
CD21lo B cells
= group
smB-21norm

≥9
Transitional
B cells
= group
smB-Trhi

<9%
Transitional
B cells
= group
smB-Trnorm

Some, non CVID, PIDs correlated with abnormal B cell subsets













Спасибо !!!