Basic Immunology Dentistry

Lymphocyte groups. Genetics of immunoglobulins, organization and expression of antigen receptor genes. Central B-cell development. Central (thymic) T cell development.

Ferenc Boldizsar

Cells of the lymphoid lineage

Innate lymphoid cells (ILC)

Lymphocyte

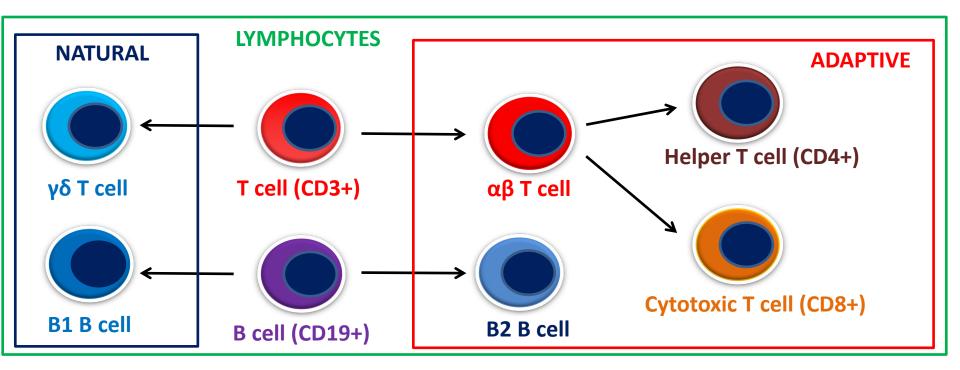


There is no difference in the morphology!

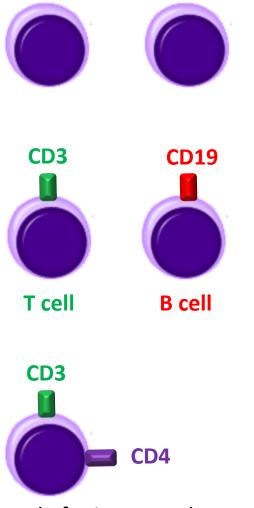


HAVE NO ANTIGEN-RECOGNITION RECEPTORS

HAVE ANTIGEN-RECOGNITION RECEPTORS



CD markers



Certain cells (e.g. lymphocytes) cannot always be distinguished based on their morphology.

Different cells can be identified and distinguished by the molecules they express on the cell surface or in the cytoplasm.

IMMUNOPHENOTYPE: The characteristic molecular pattern of a cell type determined with the use of antibodies.

Such SURFACE MOLECULES were given a standardized nomenclature:

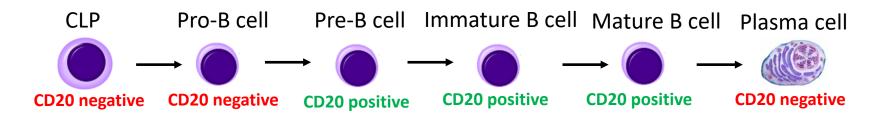
CD = **Cluster of differentiation**, usage: CD+number, e.g.: CD1, CD2, CD3, CD4, etc...

The structure and function of CD marker varies!

Example for immunophenotype: CD3+/CD4+/CD8- \rightarrow Helper T cell

Types of CD markers

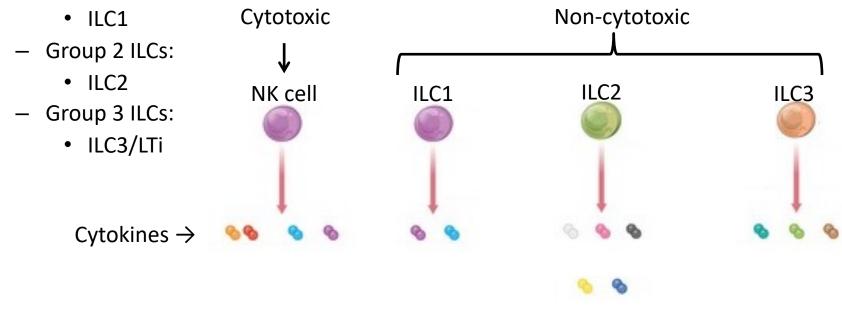
- Lineage markers: Molecules expressed exclusively on certain cell lineages.
 - − E.g.: CD3 \rightarrow found on all T cells CD19 \rightarrow found on all B cells
- **Maturation markers:** The immunophenotype might differ in the phases of cell maturation, certain molecules are only expressed on immature cells, others on mature, fully functioning cells, etc.
 - E.g.: CD20 (It is also a lineage marker of B cells, cannot be found on any other cells)



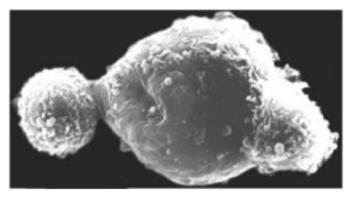
- Activation markers: Molecules expressed by activated cells, whereas resting cells either lack them completely or express them at low levels, e.g.:
 - CD25 (The alpha chain of the interleukin-2 receptor, IL-2R α , see later)
 - CD80 and CD86 (B7-1 and B7-2, so-called costimulatory molecules expressed by activated antigen presenting cells, see later)

Innate lymphoid cells (ILC)

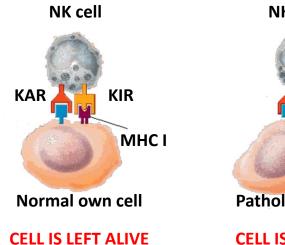
- They cannot be distinguished from lymphocytes based on their morphology but unlike adaptive lymphocytes they cannot recognize antigens. → They have no antigen recognition receptors.
- They are classified based on the cytokines they produce and the transcripition factors that are necessary for their formation. (see in the lectures):
 - Group 1 ILCs:
 - NK cells



Natural killer cells (NK cells)



Two NK cells kill a cancerous cell. (Scanning electron microscopy image)

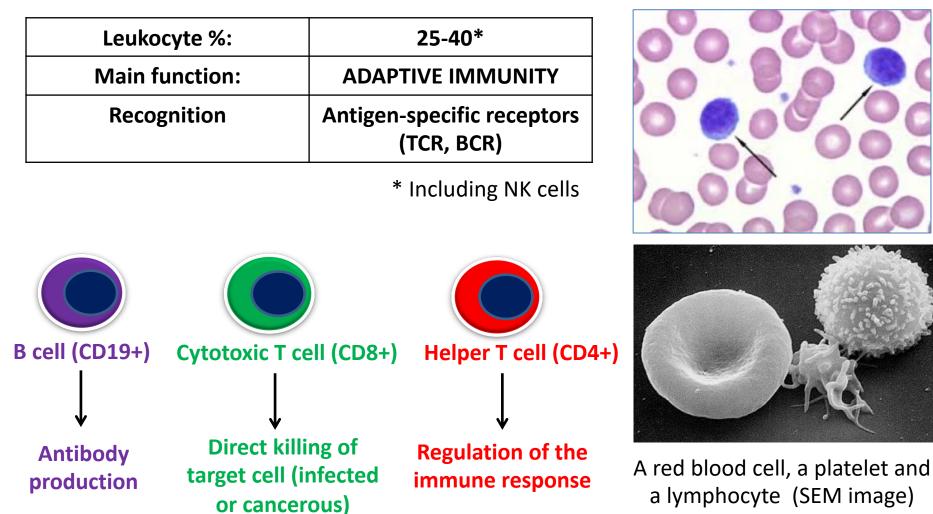


NK cell
033
Pathological cell
CELL IS KILLED

Blood lymphoid cells %:	≈ 10
Main function:	Killing cells infected with intracellular pathogens, Killing cancer cells
Recognition:	KAR → killing the target KIR → sparing the target Fc receptor, Complement receptor
Cytotoxicity:	Fas-FasL, Perforin, Granzymes
Produced mediators:	Cytokines
Fc receptor:	FcγR (<mark>binds Ig</mark> G)
Characteristic marker:	CD56

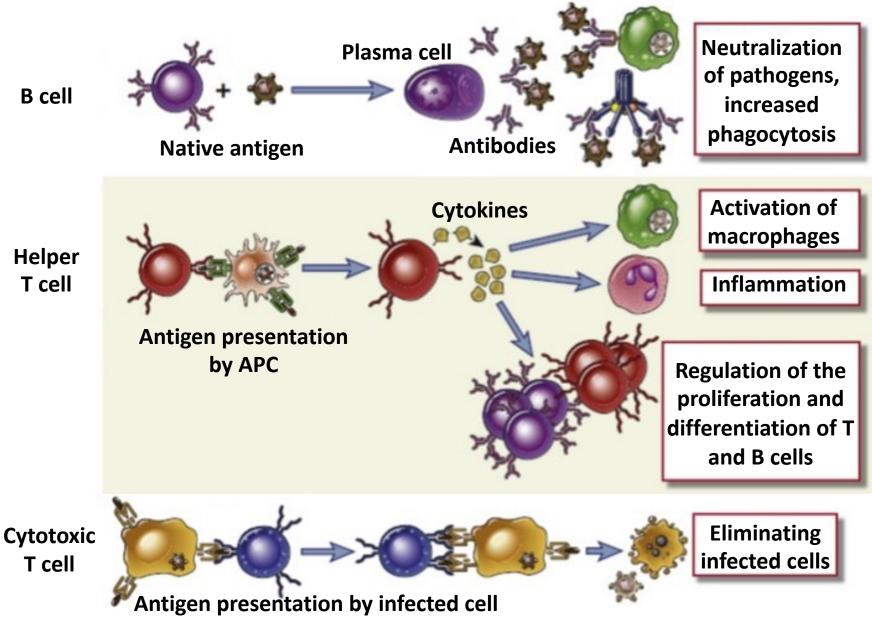
Red: Only possible after the activation of the adaptive immunity

Lymphocytes



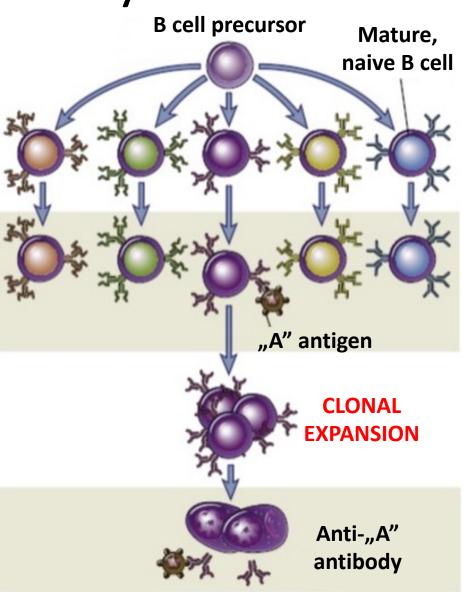
All of the above are done in an ANTIGEN-SPECIFIC manner!

Main groups of lymphocytes



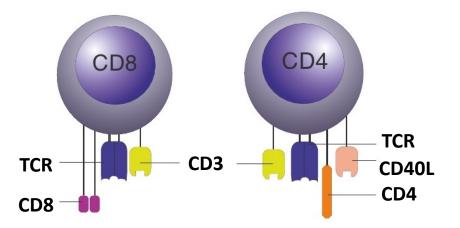
Clonality

- Each newly produced lymphocyte expresses a unique antigenbinding receptor.
- 2. Only those lymphocytes will become activated which recognize an antigen. These selected cells will proliferate and produce clones of themselves with each sister cell having the same antigenrecognition receptor.
- 3. These clones will differentiate into **effector cells** which will participate in the immune response. (e.g. effector plasma cells produce antibodies)



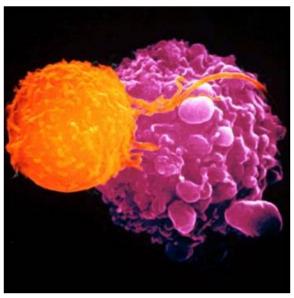
T cells

Main function:	Antigen-specific killing of target cell (CD8+), Regulation of the immune response through cytokines (CD4+)	
Recognition:	Through MHC, antigen-specific TCR	
Possible type of TCR:	αβ and γδ	
Produced mediators:	Cytokines	
Main types of $\alpha\beta$ T cells:	CD4+ Helper CD8+ Cytotoxic	
Site of production:	Bone marrow, thymus	
Characteristic marker:	CD3 (Makes a complex with the TCR)	



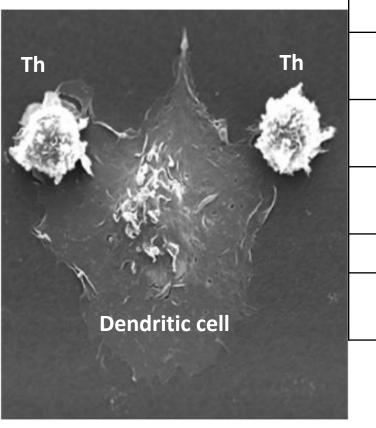
Cytotoxic T cells (Tc or CTL)

Blood T cells:	1/3
Main function:	Effector cell of the cellular immunity
Recognition:	Through MHC I, antigen- specific TCR
Target cells to kill:	Infected with IC pathogens, Cancerous, Foreign (transplantations!)
Recognized antigens:	Endogenous (from the cytoplasm of the target cell)
Cytotoxicity:	Fas-FasL, Perforin, Granzyme
Immunophenotype:	CD3+/CD8+/CD4-



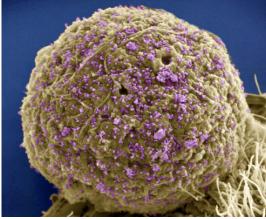
A cytotoxic T cell kills a cancer cell. (SEM image)

Helper T cells (Th)



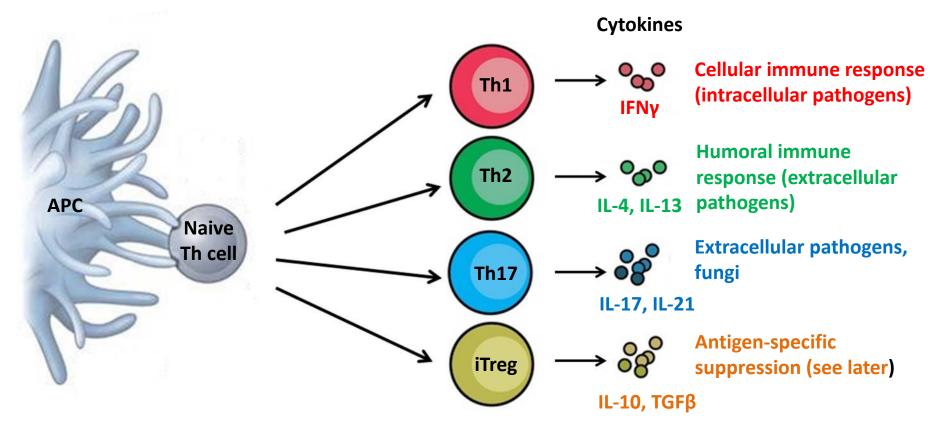
Two helper T cells attached to a dendritic cell. (Scanning electron microscopy image)

Blood T cells:	1/3
Main function:	Regulation of immune response
Recognition:	Through MHC II, antigen- specific TCR
Recognized antigens:	Exogenous (degraded in phagolysosomes)
Immunophenotype:	CD3+/CD4+/CD8-
Role in diseases:	Autoimmunity, HIV infection



Yellowish-brown: Th cell purple: **HIV** virions (SEM image)

Main subtypes of Th cells

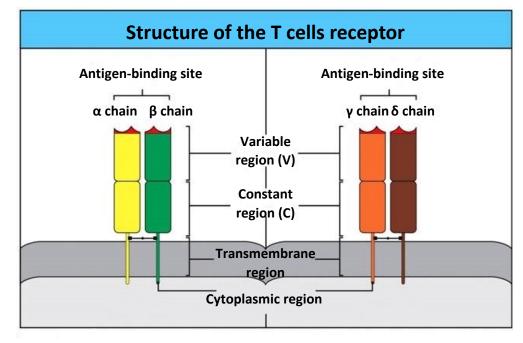


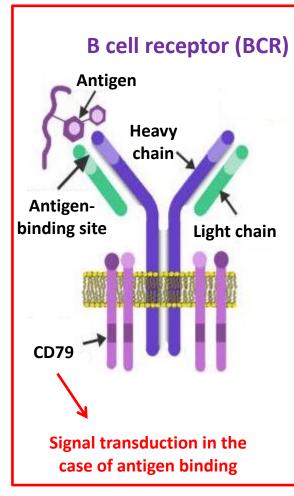
• Th17 cells play an important role in **inflammatory disorders**. (see later)

• Regulatory T cells (Treg): They can inhibit other immune cells (suppression, see later), their immunophenotype is: CD4+/CD25+/Foxp3+

$\gamma\delta$ T cells

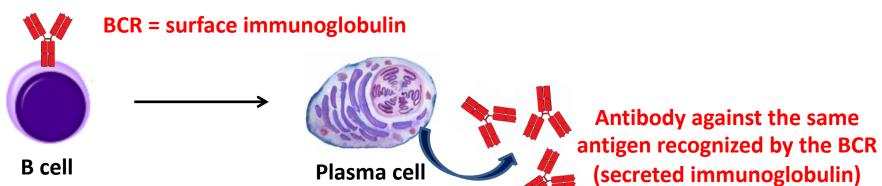
- They express TCRs that consist of γ and δ chains.
- They are **innate-like lymphocytes**, they are not as well-characterized as $\alpha\beta$ T cells.^[17.]
- They are mainly found in the **skin** and the **mucosa**; usually as intraepithelial lymphocytes (IELs). They can be detected in the peripheral blood in low numbers.
- They participate in the early phases of the immune response against invasive pathogens.
- Their antigen-recognition is **MHC-independent**.
- They mainly recognize **lipid antigens**.





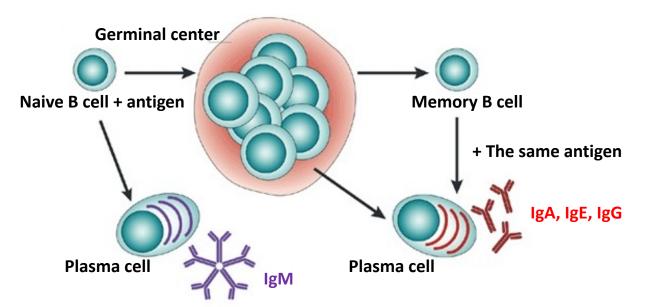
B cells

Blood lymphoid cells %:	10-15
Main functions:	Antibody production, Antigen presentation
Recognition:	Native antigens with antigen- specific BCR
Main types:	B1 and B2
Site of production:	Bone marrow
Characteristic marker:	CD19 (makes a complex with BCR)



B2 B cells

Found in:	Follicles in secondary Imyphoid organs, blood
Main functions:	Antibody production, Antigen presentation
Recognition:	Native antigens with antigen-specific BCR
Site of primary maturation:	Bone marrow
Site of antigen-dependent maturation:	Germinal center
Produced antibodies:	Monospecific, high-affinity, with varying isotype

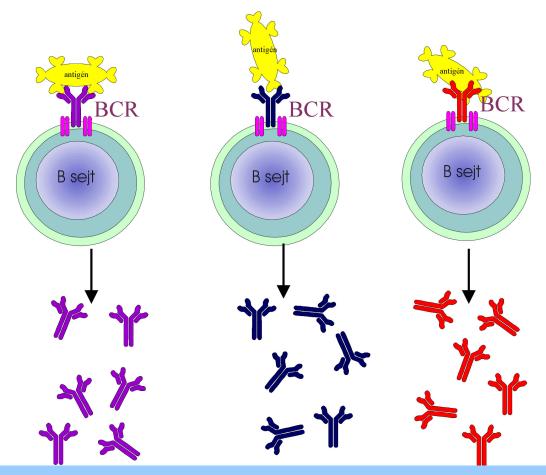


B1 B cells

- Only few can be found in the peripheral blood.
- They are innate-like lymphocytes, most of them reside on serous membranes. (e.g. peritoneum, pleura, pericardium)
- They are first produced in the fetus and later undergo self-renewal in the periphery, not in the bone marrow, as B2 cells do.
- They produce **natural autoantibodies** that can bind that can bind evolutionarily **conserved self-antigens**.
- They were first described as CD5+ B cells in mice.
- The immunophenotype of the human B1 cells is still controversial.

	B1 cells	B2 cells
Spontaneous antibody production	Significant	Minimal
Isotype of produced antibodies	IgM	IgM/IgG/IgA/IgE
Affinity and specificity of antibodies	Polyspecific with low affinity	Monospecific with high affinity
Affinity maturation, memory	No	Yes

Antibody – B-cell-Repertoire: **10**¹¹



Tonegawa (Nobel prize:1987)

During B cell differentiation Immunoglobulin genes are rearranged and somatic Hypermutations take place.

Compared to the large repertoire relatively few Ig V genes are inherited.

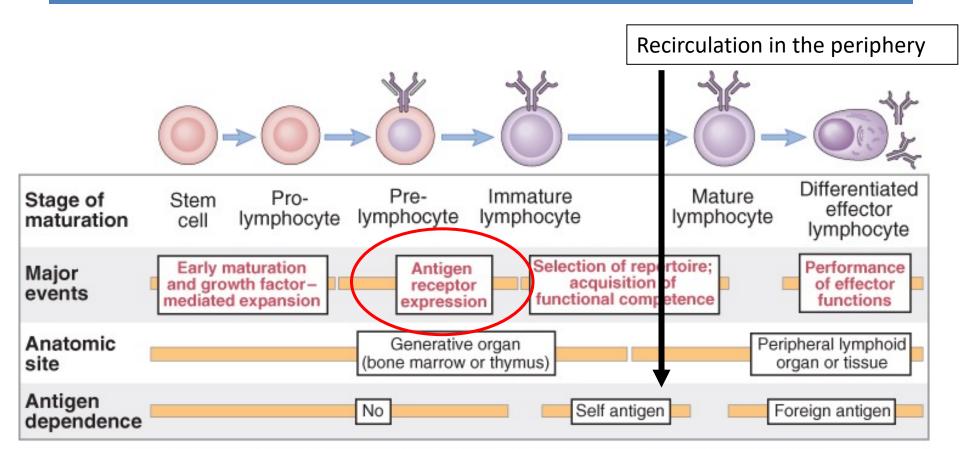
Aim of lymphocyte differentiation

- Expression of Antigenreceptors with different specificitites
- Production of B- and T cell repertoire = Number of antigen recognition molecules: 10⁹-10¹¹ BcR, 10¹⁵-10¹⁶ TcR;

"Lymphocyte production = Glove factory" – <u>Jan Klein</u>. The immune system produces antigen receptors for all potential antigens and is therefore ready to recognize those structures.

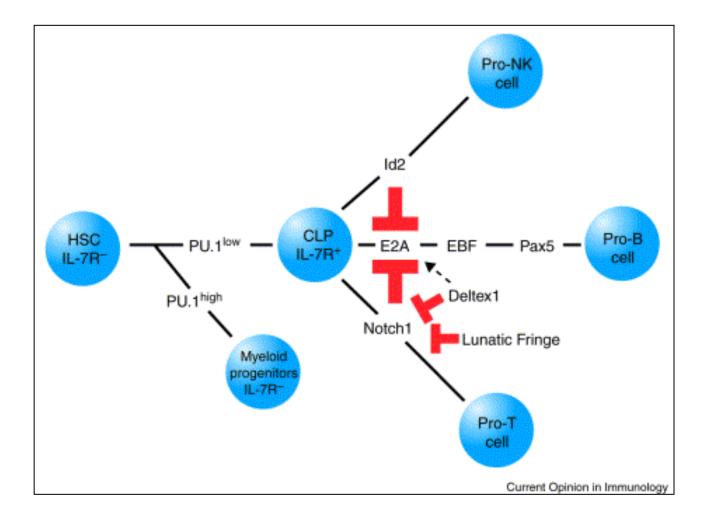
The genetic background of B- and T cell receptor production is the gene rearrangement of Ig- and TcR genes in the progenitor cells.

Steps of lymphocyte development



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B/T/NK commitment – default E2A (B) path overruled by Notch (T) and/or Id2 (NK) signals



Role of BM stroma

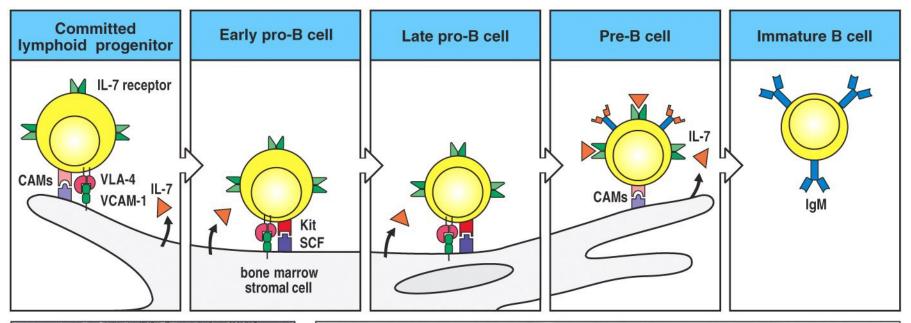
1. Adhesion: – CD44, VCAM-1

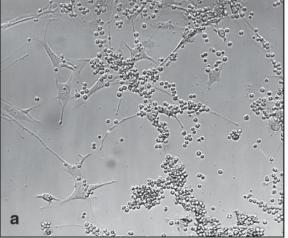
2. Growth factors: IL-7, IL-3, SCF.

3. <u>Response modifiers</u>: Wnt factors, ECM components.

4. Chemokine-production: SDF-1/CXCR4 ligand.

Elements of B:stromal interactions





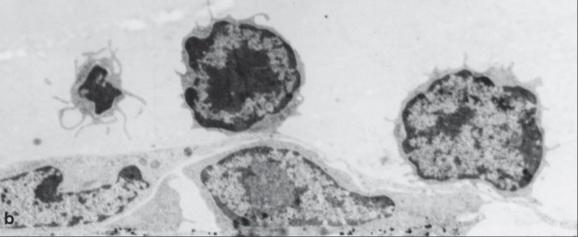


Figure 7-4 Immunobiology, 6/e. (© Garland Science 2005)

The antigen binding parts of the Immunoglobulins contain hypervariable (CDR) regions

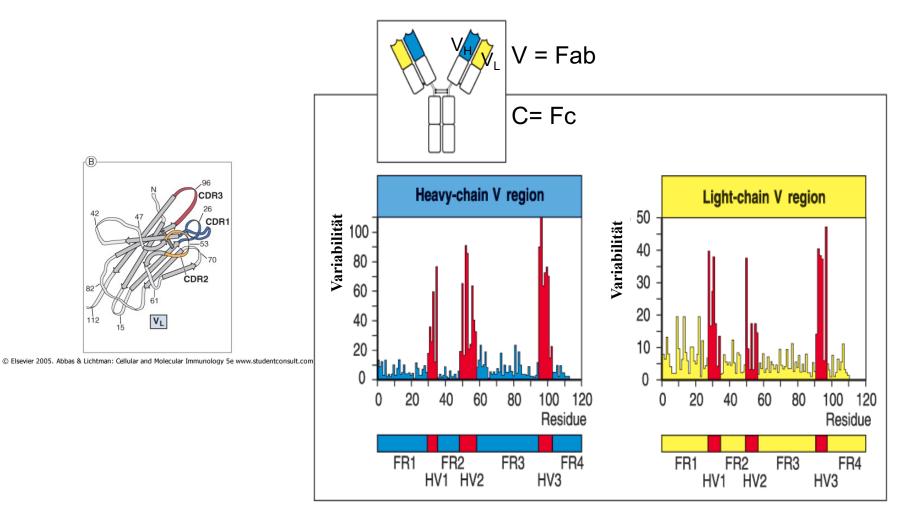
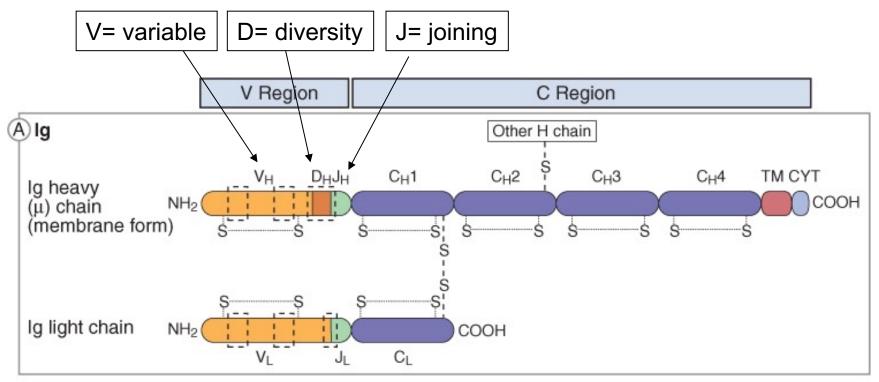


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Domains of the immunglobulin heavy- and light chains

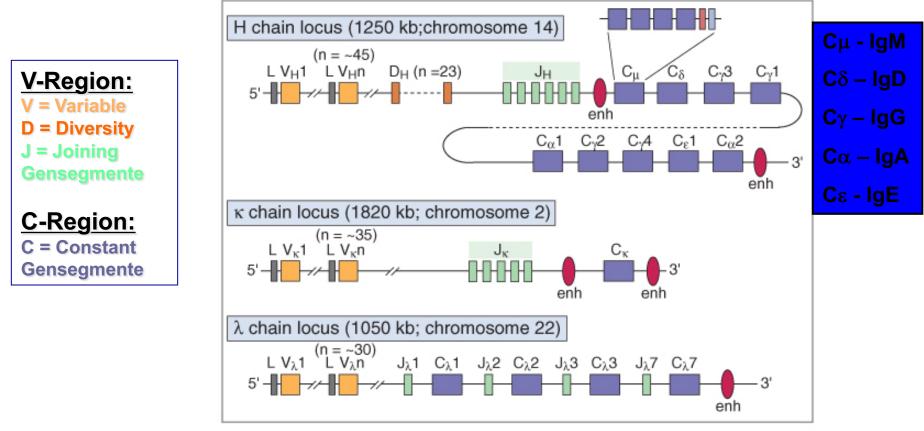


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- The <u>variable (V)</u> and <u>constant (C)</u> domains (units) of the heavy- and light polypeptide chains are encoded by different gene segments.

- The genes of the Ig heavy- and light polypeptide chains are located in different chromosomes.

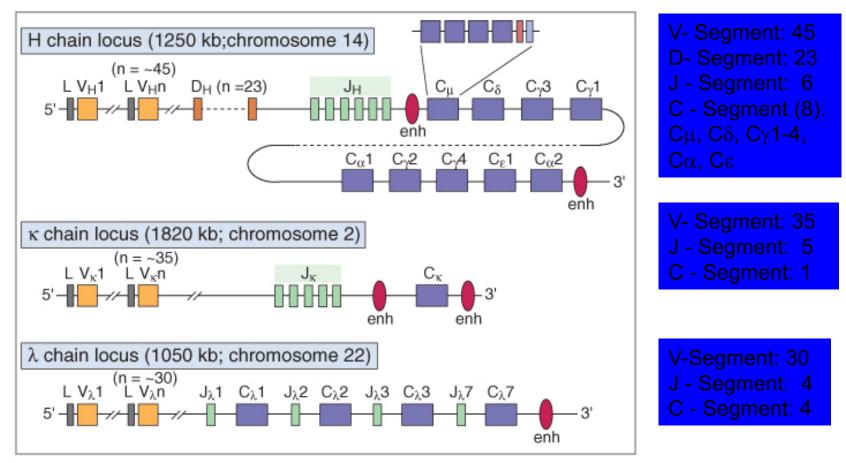
Gene organisation of the immunglobulin heavy- and light chain loci



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The germline-DNA \rightarrow the basic, not-recombined form of the immunoglobulin genes

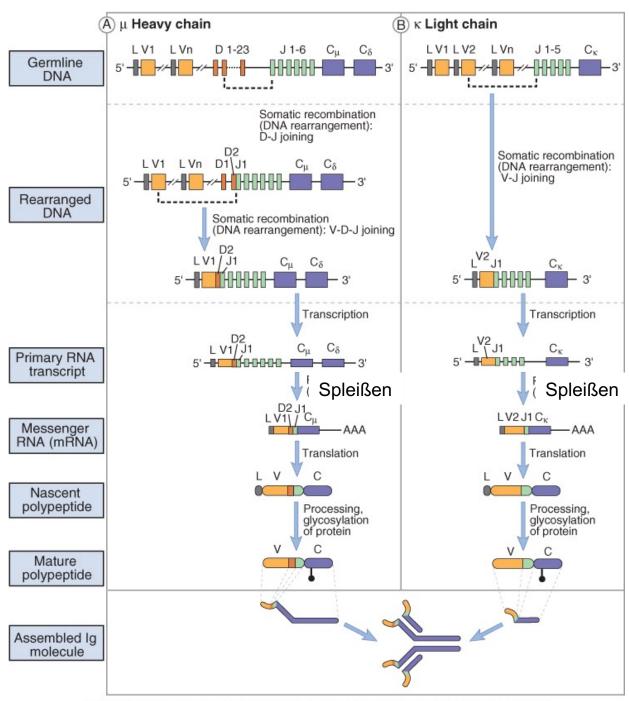
The germline Ig DNA: number of V-D-J-gene segments



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In lymphocyte precursors the germline DNA will be rearranged by somatic recombination. = **Rearrangement**

Steps of the gene rearrangement



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Molecular mechanism of the gene rearrangement

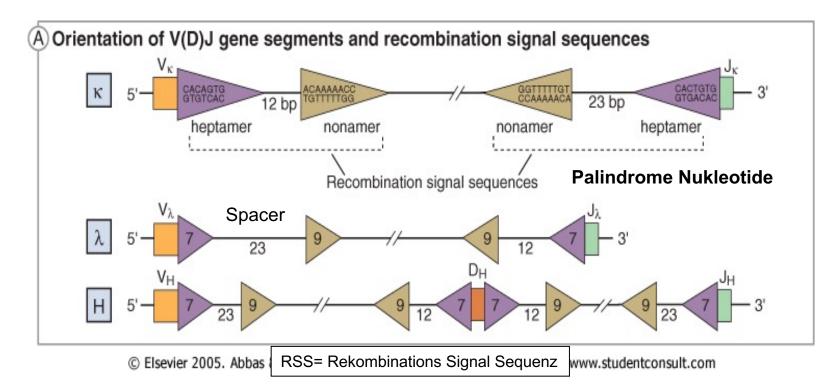
- 1. DNA loop formation
- 2. DNA cutting Deletion
- 3. Ligation of the free DNA ends

Enzymes:

- VDJ-Recombinase: RAG1 and -2
- Heteromeric Proteincomplex: **DNA-Ligase, DNA-PK, Artemis-Protein**
- Terminale Deoxynukleotidyl-Transferase (TdT): \rightarrow

N-Nukleotide-addition – random addition of nucleotides

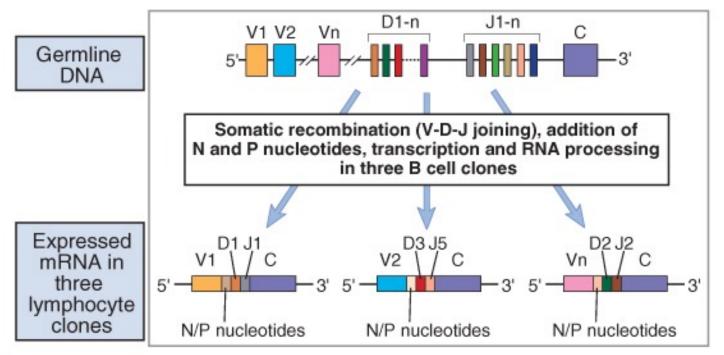
The 12/23-base-pair rule during the recombination of Ig gene segments:



Recombination-Signal-Sequence (RSS):

Contains a conserved heptamer and nonamer sequences which are divided by a non-conserved spacer sequence of either 12 or 23 basepairs.

Heavy chain gene rearrangement in three pro-Bcells

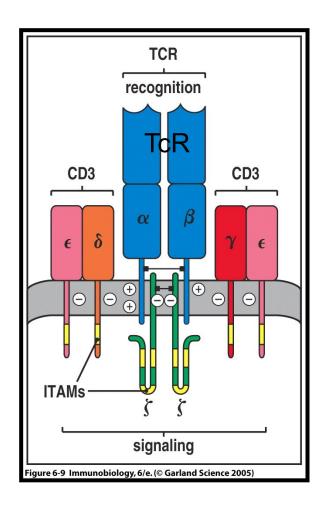


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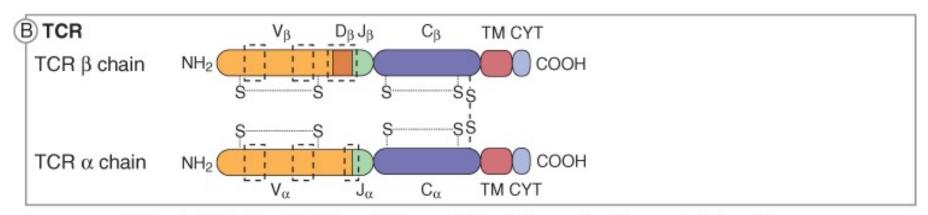
Random gene rearrangement — Diversity

T-cell-receptor (TcR)

T-cell-types: 1. αβ TcR+ 2. γδ TcR+

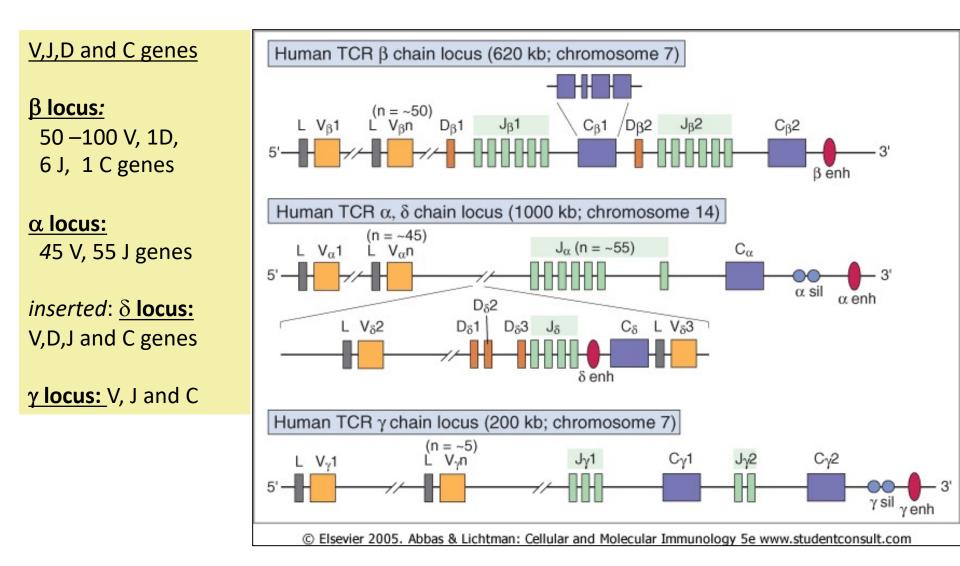


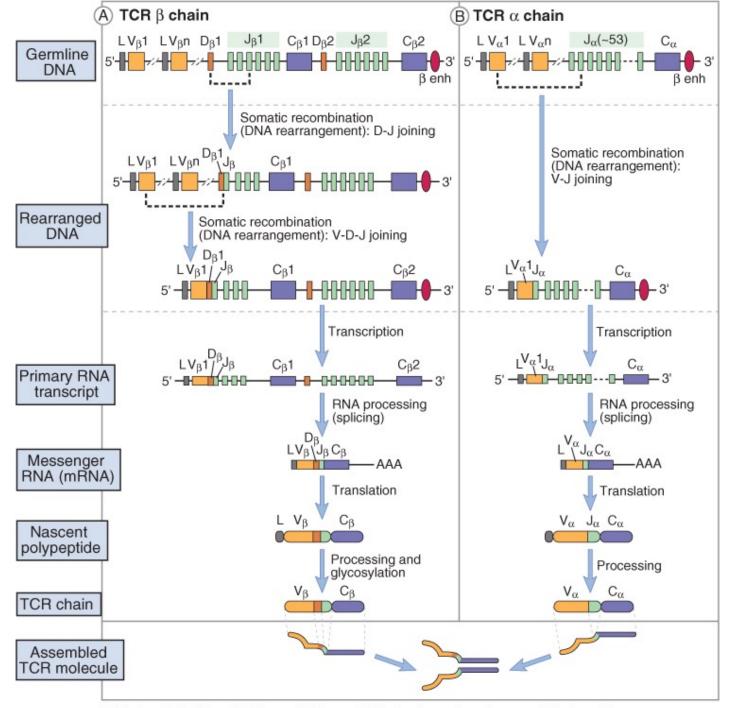
TcR α - β chains



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Human TCR encoding genes





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The basis of TcR and BcR diversity

- The genes encoding the TcR α/β and γ/δ chains have similar structure (multiple V, D, J and C segments) than that of the Ig genes and the steps of the gene rearrangement is also the same (role of RAG1 and RAG2)
- The large number of V, D and J segments and their free recombination
- The effect of TdT (terminal deoxynucleotidyl transferase) during recombination → CDR3 variability is higher
- Random combination of TcR α/β and γ/δ chains (like Ig H/L chains)

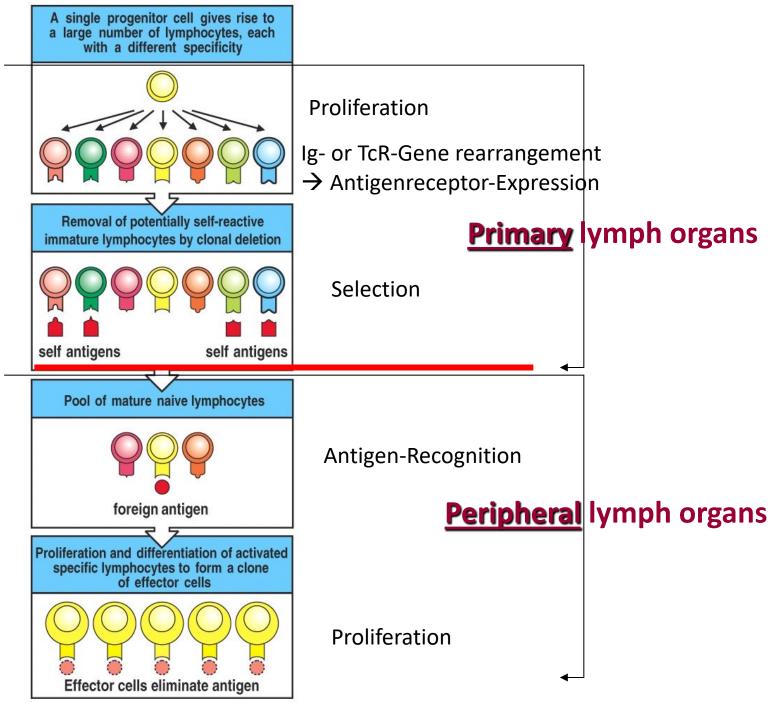


Figure 1-14 Immunobiology, 6/e. (© Garland Science 2005)

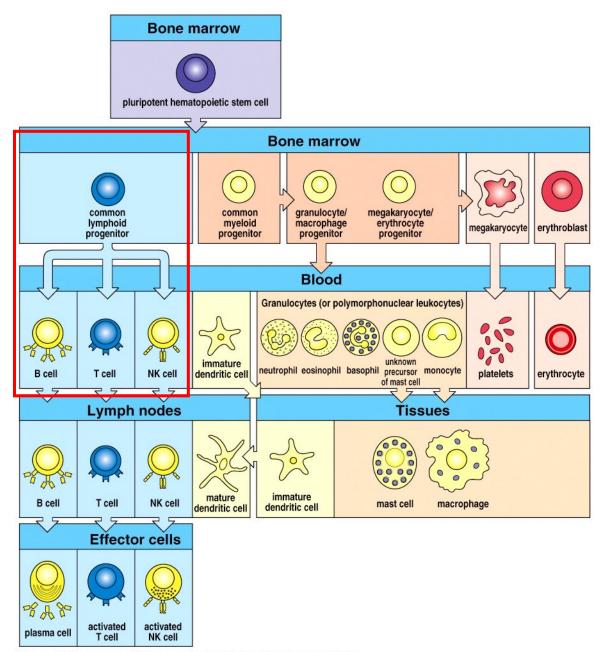
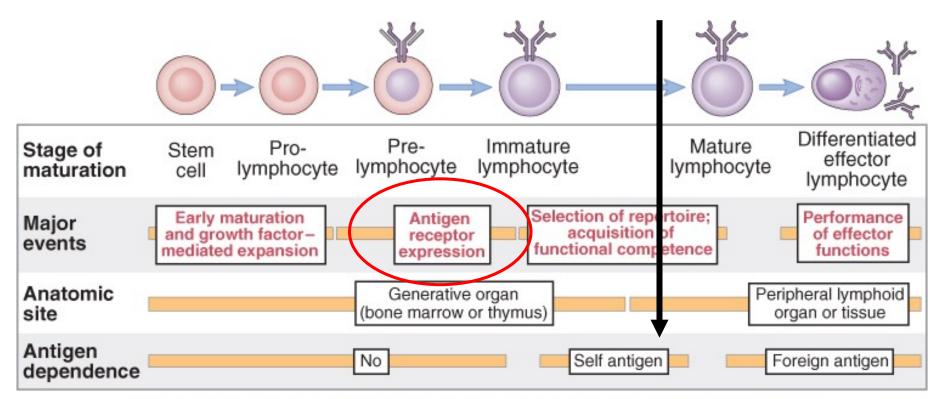


Figure 1-3 Immunobiology, 6/e. (© Garland Science 2005)

General characteristics of lymphocyte differentiation

- 1. Proliferation
- 2. Receptor-Gene rearrangement
- 3. Migration
- 4. Selection
- 5. Apoptosis

Stages of Lymphocyte differentiation



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B-cell development I: HSC > "Large pre-B"

	Stem cell	Early pro-B cell	Late pro-B cell	Large pre-B cell	
				pre-B receptor	
H-chain genes	Germline	D-J rearranging	V–DJ rearranging	VDJ rearranged	
L-chain genes	Germline	Germline	Germline	Germline	
Surface Ig	Absent	Absent	Absent	μ chain transiently at surface as part of pre-B-cell receptor. Mainly intracellular	
	CD34, c- <i>kit</i>	(CD34), CD19, CD45RA, IL-7R	CD9, CD19 CD45RA, IL-7R		

Fig 7.5 part 1 of 2 © 2001 Garland Science

B-cell development II "Small pre-B" > "mature B"

	Small pre-B cell	Immature B cell	Mature B cell		
_		IgM	IgD IgM		
H-chain genes	VDJ rearranged	VDJ rearranged	VDJ rearranged		
L-chain genes	V–J rearranging	VJ rearranged	VJ rearranged		
Surface Ig	intracellular µ chain	IgM expressed on cell_surface	IgD and IgM made from alternatively spliced H-chain transcripts		
		CD19, MHC-II, CD45RA	CD19/24, MHC-II, CD45RA		

Fig 7.5 part 2 of 2 © 2001 Garland Science

Ontogenic differences between B-cell subsets

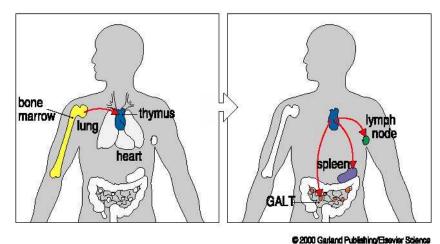
- B-1 B cell subsets: fetal origin, self-renewal, low-affinity autoantibody production, dominance in neonates and CLL, located in body cavities.
 (CD5+, CD43+, IgM++/IgD+)
- Marginal zone B cells: Ig phenotype similar to B-1 B cells, adult BM origin, distinct developmental regulation to Fo B cells, relatively sessile.

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(lgM++/lgD+, CD21++, CD23+/-)
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• **Conventional follicular B cells.** (IgM+/IgD++, CD21+, CD23++, recirculating).

T-cell development in the thymus.

Figure 5.1



Production of T cell repertoire

Total repertoire: TCR α , β : 10¹⁵ TCR γ , δ : 10¹⁶

T cell precursors are produced in the **bone marrow** from the common haemopoietic stem cell

They migrate through the blood circulation to the thymus

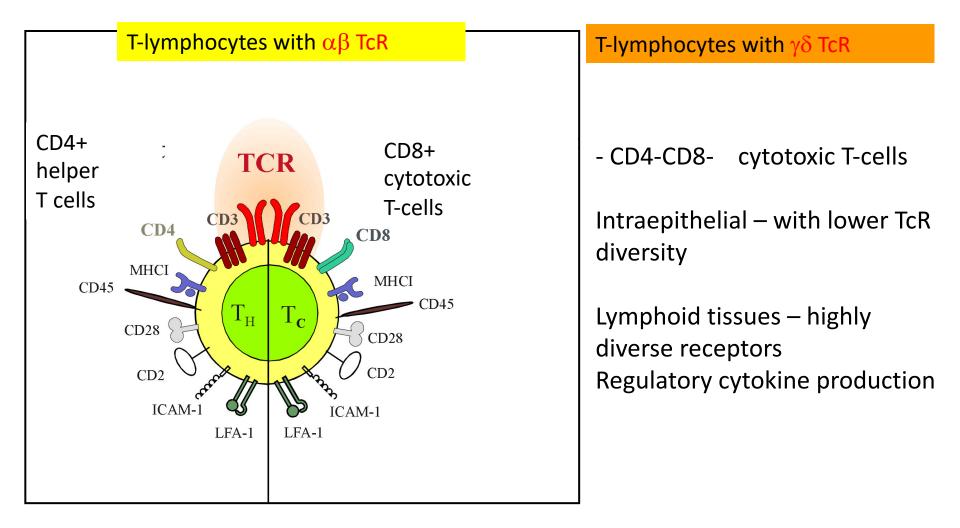
<u>In the thymus</u>: T cell maturation, educational steps ,,double recognition" (MHC and peptide)

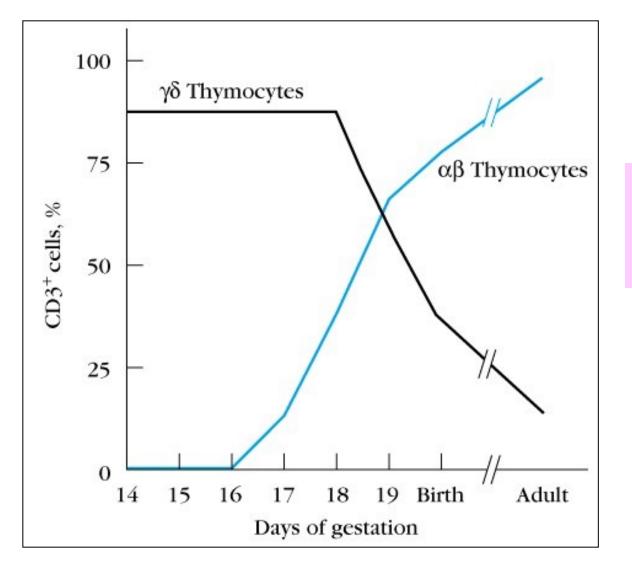
<u>Periphery</u>: mature, TCR expressing, CD4 or CD8 positive T cells

Self-MHC restricted

Self-tolerant T cells

Two different T cell lines with different receptor types (TcR)

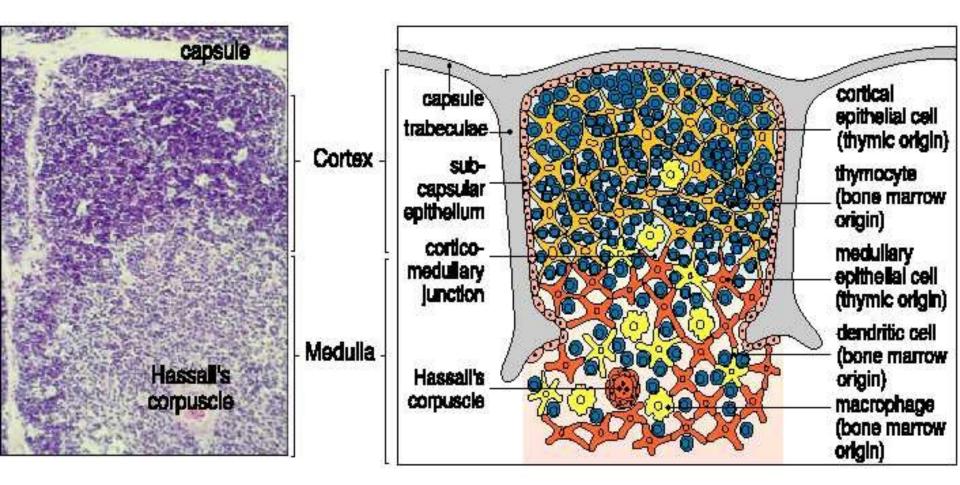




Full repertoir: TCR α , β : 10¹⁵ TCR γ , δ : 10¹⁶

Structure of the thymus

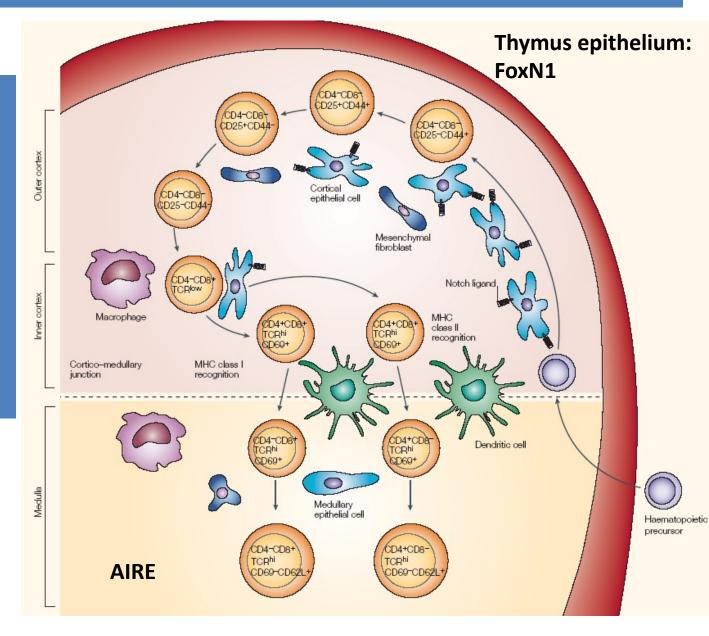
Figure 5.3



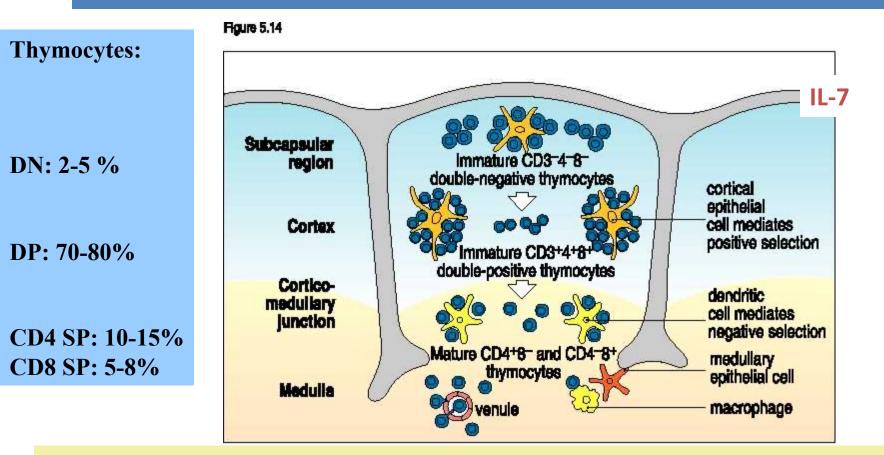
The thymic stoma creates the microenvironment that is essential for T-cell development

Thymic Microenvironment and T-cell Development

- 1. <u>Migration:</u> Chemokine effect
- 2. <u>Proliferation</u> IL-7
- 3. Differentiation
- TcR rearragement
- Phenotypes
- 4. <u>Selection</u> Apoptosis



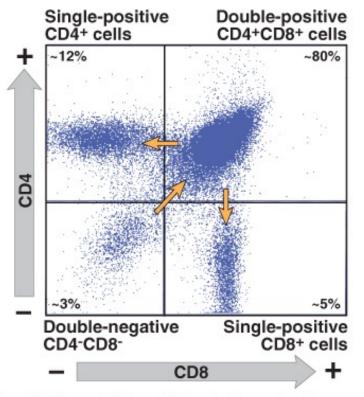
T-cell development in the thymus



Young mouse: 5x10⁷ T-cells dayly During selection 98 % of thymocytes die by apoptosis

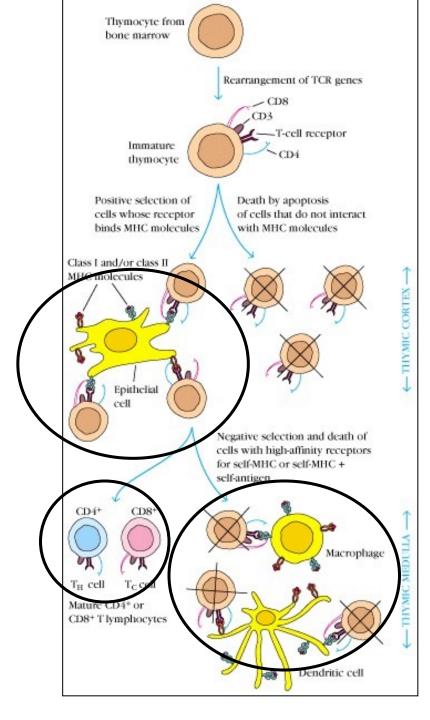
Dayly 1-2 x 10⁶ mature T-cell migrate to the periphery

Thymocyte populations based on their cell surface markers



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				DP-	SP -		
Stage of maturation	Stem cell	Pro-T	Pre-T	Double positive	Single positive (immature T cell)	Naive mature T cell	
Proliferation				1			
RAG expression							
TdT expression							
TCR DNA, RNA	Unrecombined (germline) DNA	Unrecombined (germline) DNA	Recombined β chain gene [V(D)J-C]; β chain mRNA	Recombined β, α chain genes [V(D)J-C]; β and α chain mRNA	Recombined β, α chain genes I [V(D)J-C]; β and α chain mRNA	Recombined β , α chain genes [V(D)J-C]; β and α chain mRNA	
TCR expression	None	None	Pre-T receptor (β chain/pre-T α)	Membrane αβ TCR	Membrane αβ TCR	Membrane αβ TCR	
Surface markers	c- <i>kit</i> + CD44+ CD25 ⁻	c- <i>kit</i> + CD44+ CD25+	c- <i>kit</i> + CD44+ CD25+	CD4+CD8+ TCR/CD3lo	CD4+CD8 ⁻ or CD4-CD8+ TCR/CD3 ^{hi}	CD4+CD8 ⁻ or CD4-CD8+ TCR/CD3 ^{hi}	
Anatomic site	Bone marrow		Thy	rmus		Periphery	
Response to antigen	None	None	None	Positive and negative selection	Negative selection	Activation (proliferation and differentiation)	
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Positive selection:

Epithelial cell - thymocyte interaction in the thymus cortex

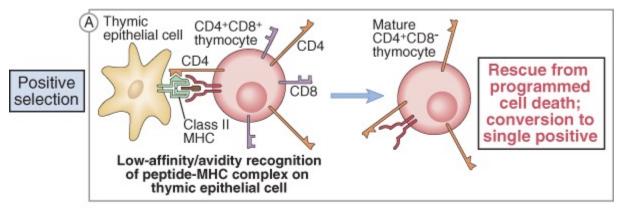
Survival of DP cells whose TcR is appropriate for self MHC recognition

Negative selection:

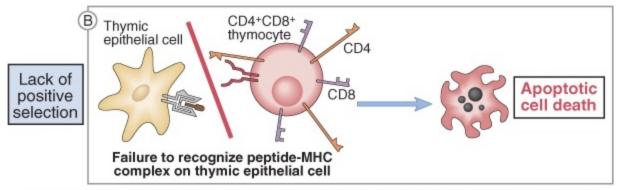
APC (macropahge or DC) – thymocyte Interaction in thymus medulla

Death of DP cells with high affinity TcR for self MHC + self peptide recognition

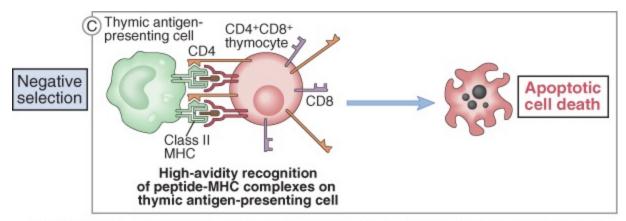
Differentiation into SP stage





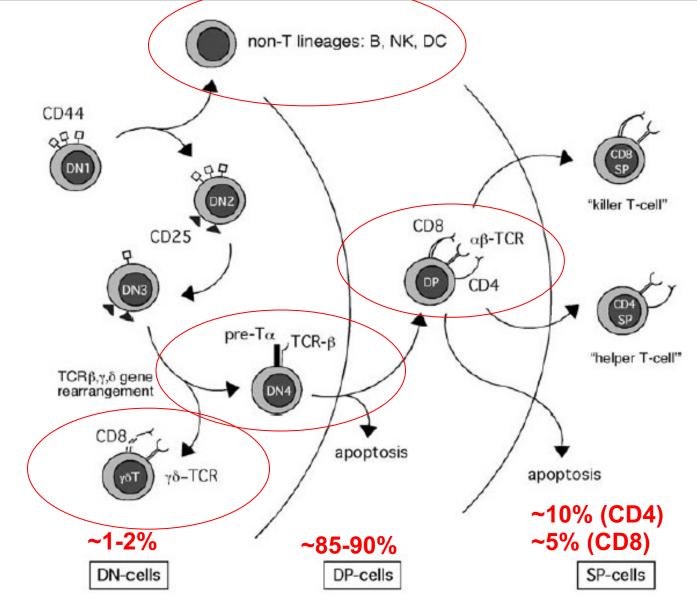


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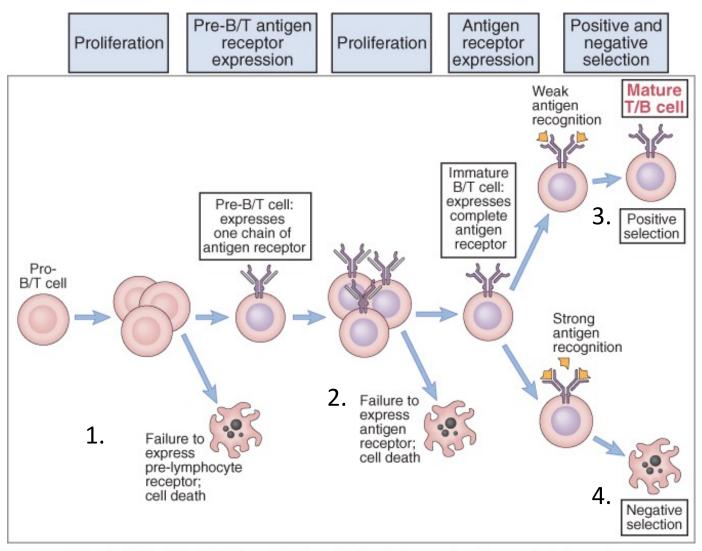
Decision-making during the development of T cells



Phases of T-cell maturation in the thymus.

- 1. Initiation of either TCR β or γ/δ chain gene rearrangement.
- 2. Formation of pT α /TCR β /CD3 (pTCR), allelic exclusion, IL-7-dependent proliferation β -selection.
- 3. Initiation of TCR α gene rearrangement.
- 4. Completion of TCR α/β gene rearrangement, co-expression of CD4/CD8 molecules.
- 5. Recognition of MHC/peptide complexes displayed by thymic cortical epithelium *positive selection*.
- 6. Binding to MHC/peptide complex displayed by thymic APC/medullary epithelial cells *negative selection*.
- 7. Influence of stronger/more persistent signal: commitment towards CD4 or Treg (CD4/CD25+) subset.

"Checkpoints" in central B/T-lymphocyte development



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