

Basic Immunology

Dentistry

Genetics of immunoglobulins, organization and expression of antigen receptor genes.

Central B-cell development.

Central (thymic) T cell development.

Ferenc Boldizar

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

NATURAL



$\gamma\delta$ T cell



B1 B cell

LYMPHOCYTES



T cell (CD3+)



B cell (CD19+)



$\alpha\beta$ T cell



B2 B cell

ADAPTIVE



Helper T cell (CD4+)

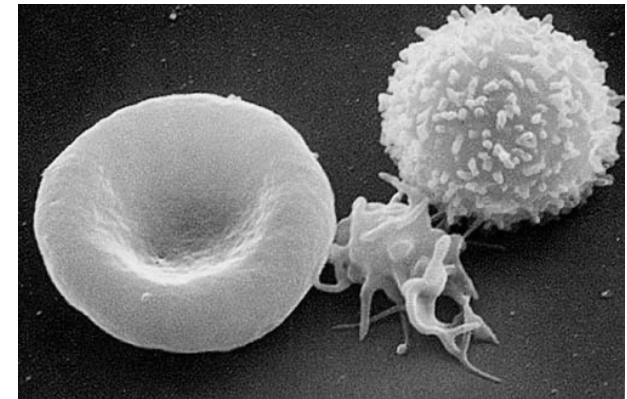
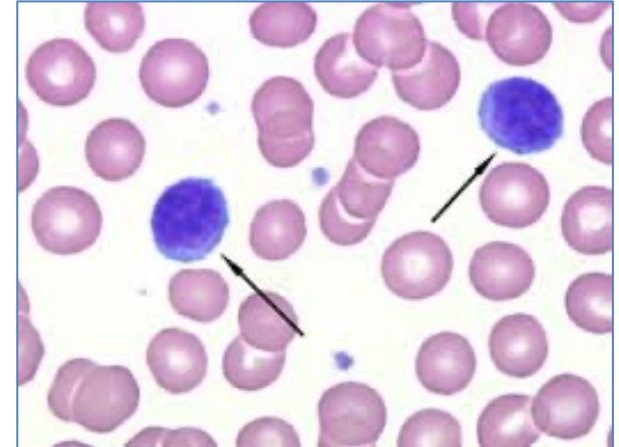


Cytotoxic T cell (CD8+)

Lymphocytes

Leukocyte %:	25-40*
Main function:	ADAPTIVE IMMUNITY
Recognition	Antigen-specific receptors (TCR, BCR)

* Including NK cells



A red blood cell, a platelet and a lymphocyte (SEM image)



B cell (CD19+)



Antibody production



Cytotoxic T cell (CD8+)



Direct killing of target cell (infected or cancerous)



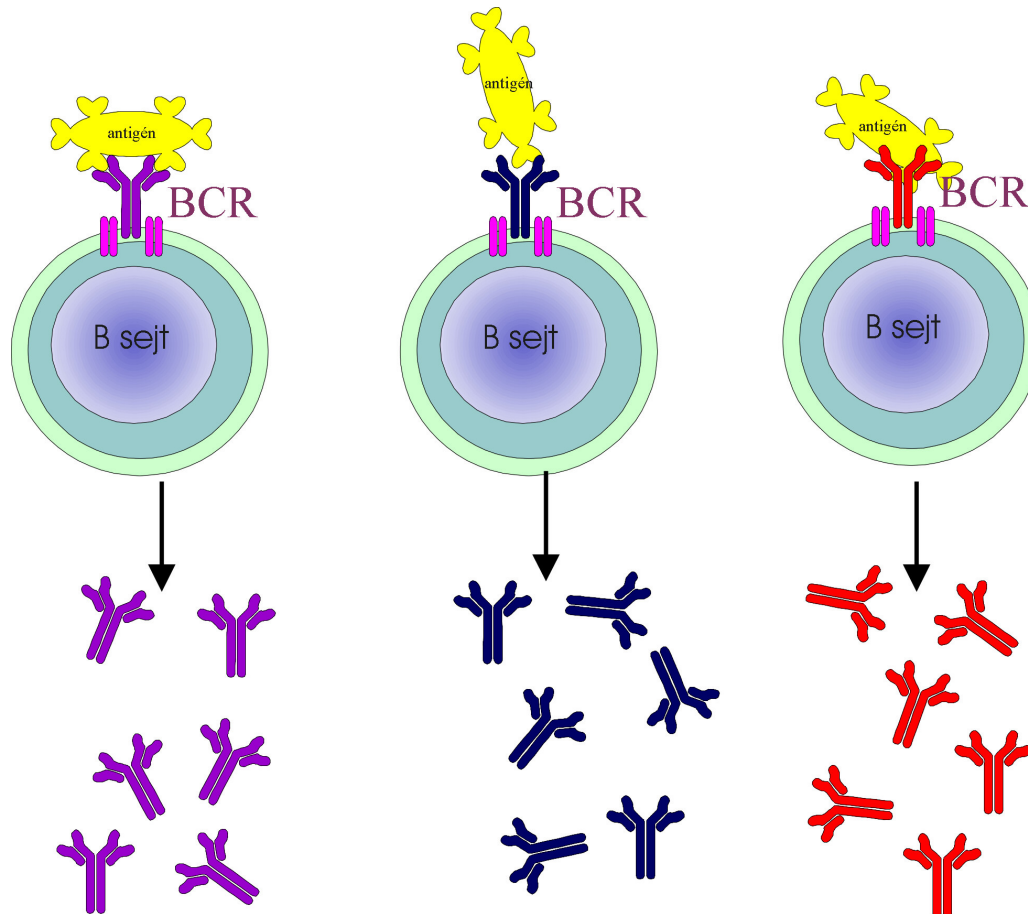
Helper T cell (CD4+)



Regulation of the immune response

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

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

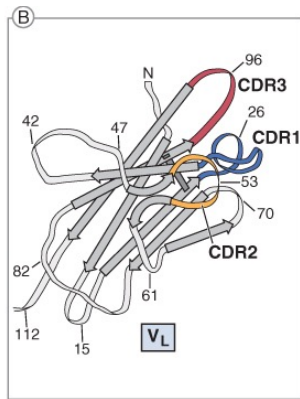
- Expression of Antigenreceptors with different specificities
- Production of B- and T cell repertoire = Number of antigen recognition molecules: 10^9 - 10^{11} BcR, 10^{15} - 10^{16} TcR;

„Lymphocyte production = Glove factory” – Jan Klein.

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.**

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



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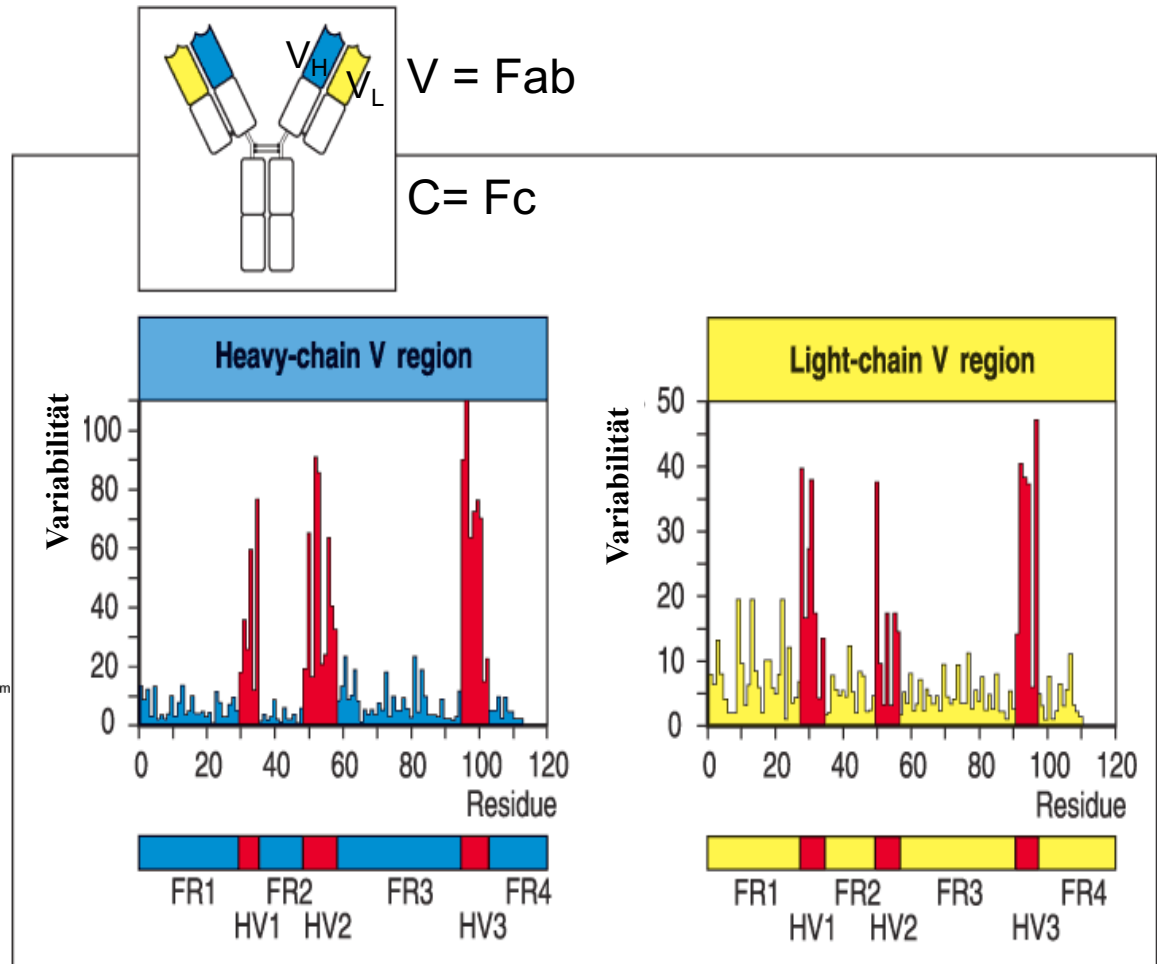
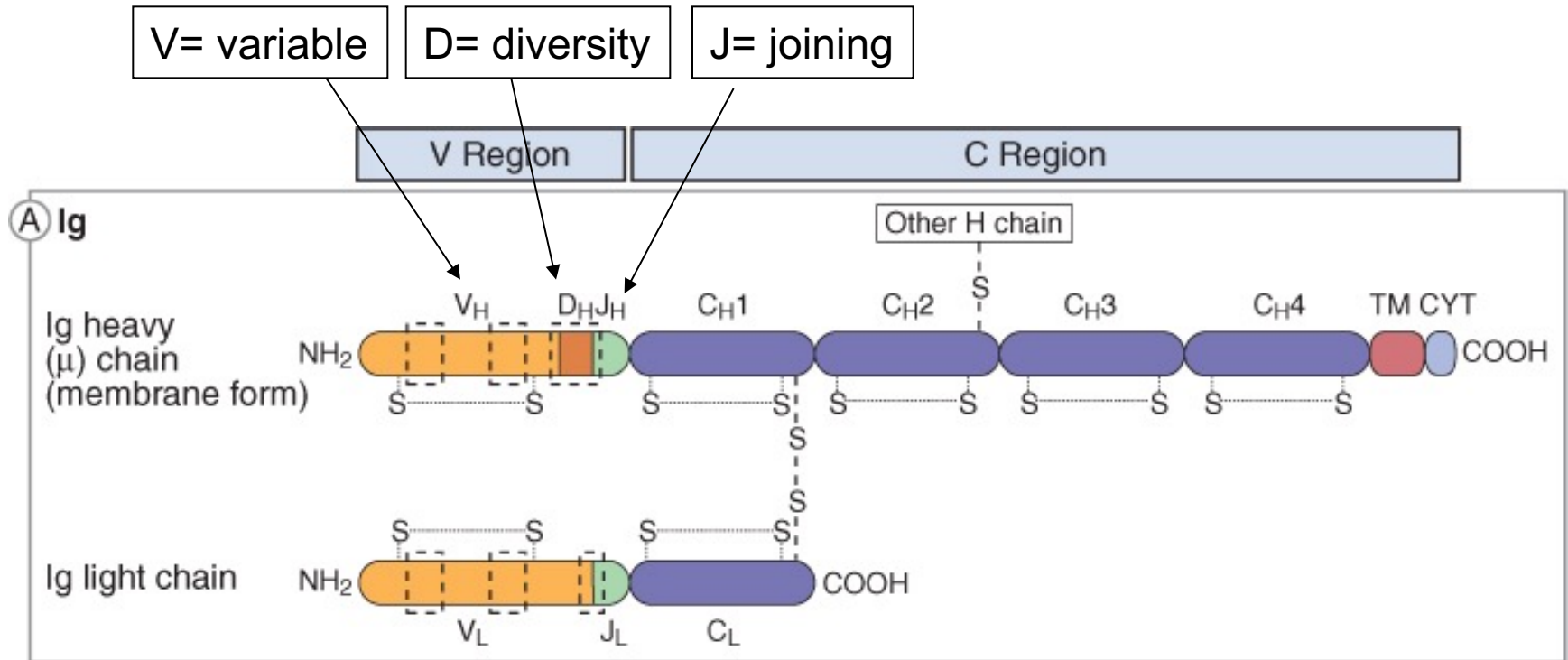


Fig 3.6 © 2001 Garland Science

Domains of the immunoglobulin heavy- and light chains



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- The **variable (V)** and **constant (C) 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 immunoglobulin heavy- and light chain loci

V-Region:

V = Variable

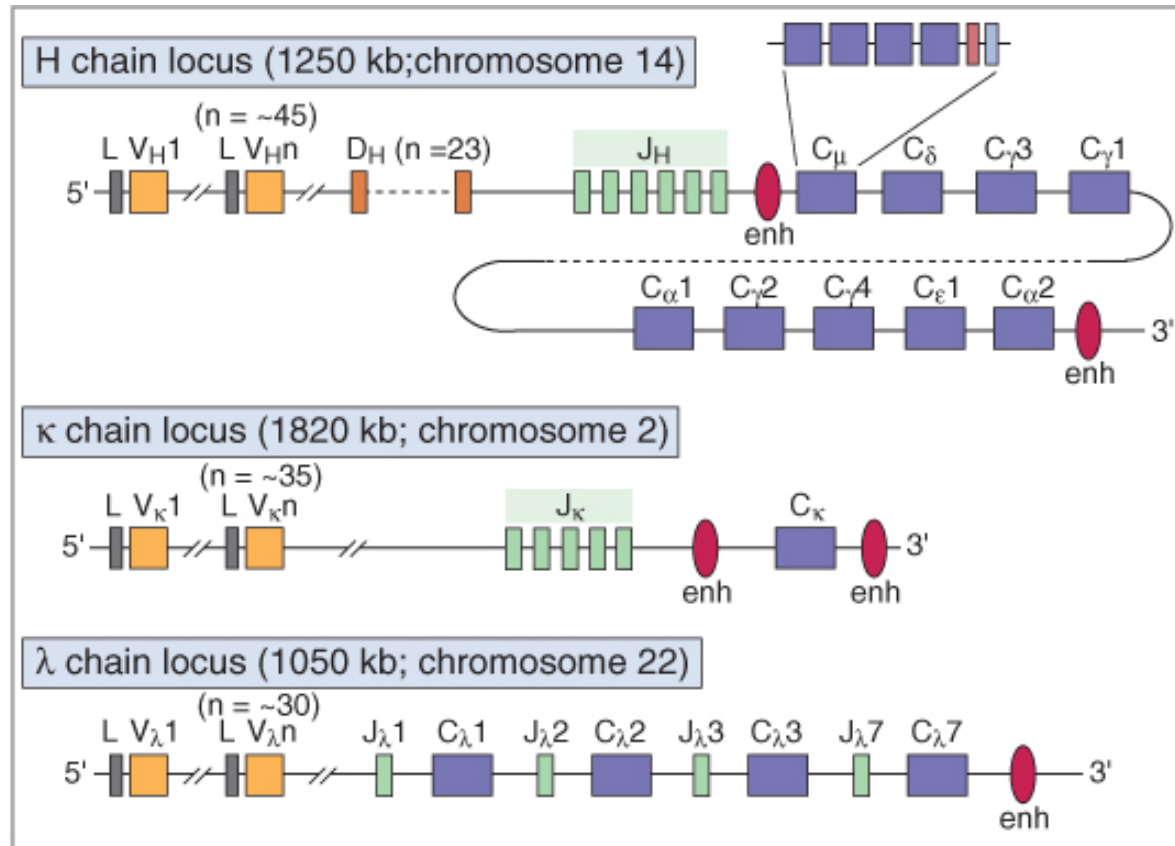
D = Diversity

J = Joining
Gensegmente

C-Region:

C = Constant

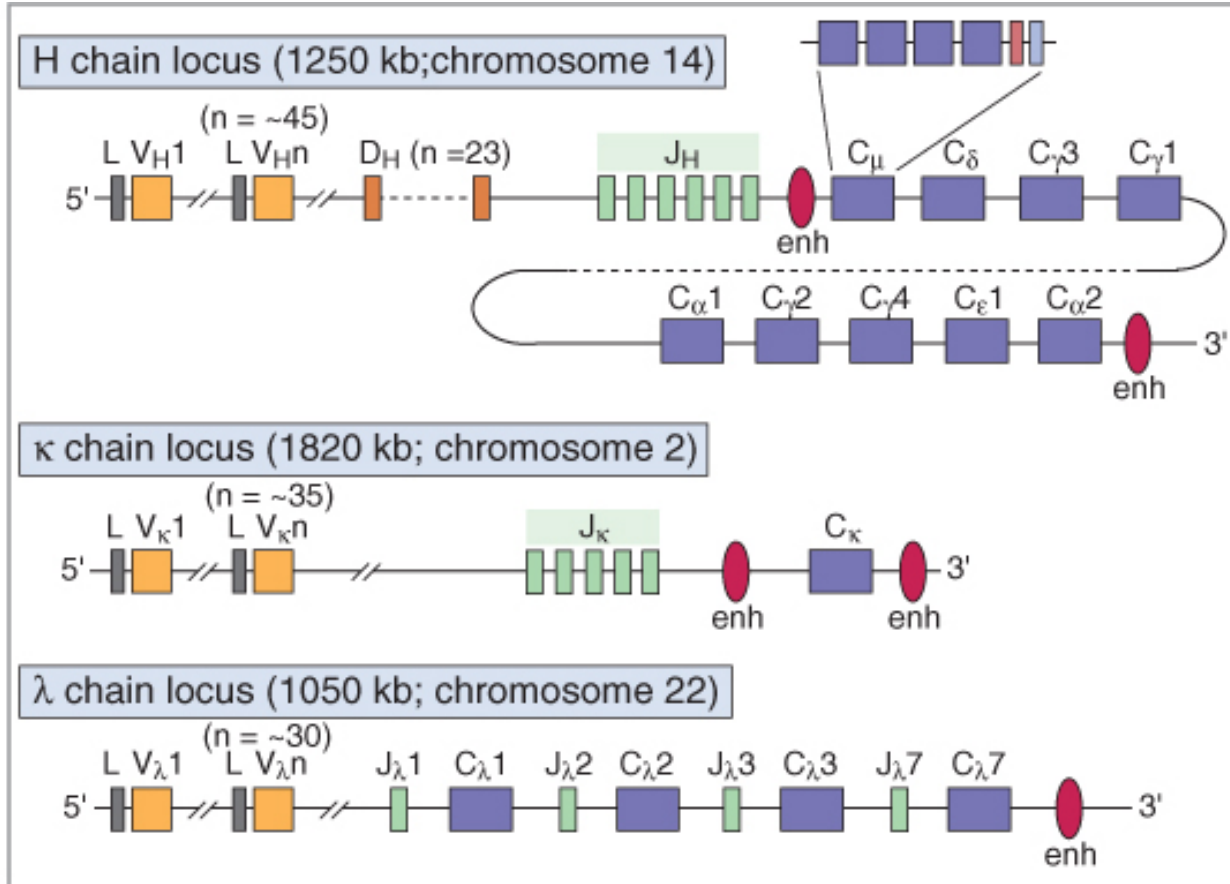
Gensegmente



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The **germline-DNA** → the basic, not-recombined form of the immunoglobulin genes

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



V- Segment: 45
 D- Segment: 23
 J - Segment: 6
 C - Segment (8):
 C_μ, C_δ, C_{γ1-4},
 C_α, C_ε

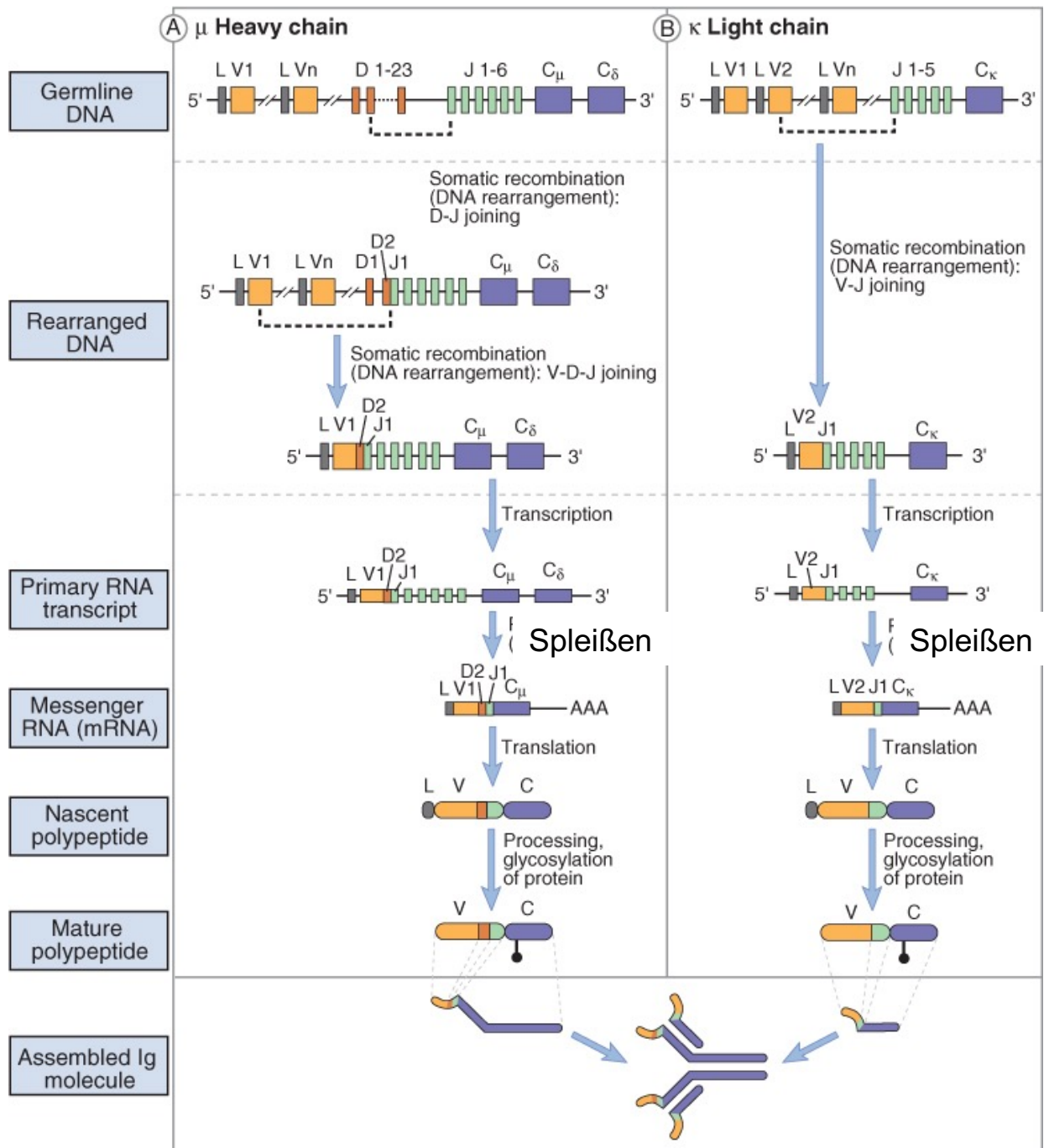
V- Segment: 35
 J - Segment: 5
 C - Segment: 1

V-Segment: 30
 J - Segment: 4
 C - Segment: 4

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

Steps of the gene rearrangement



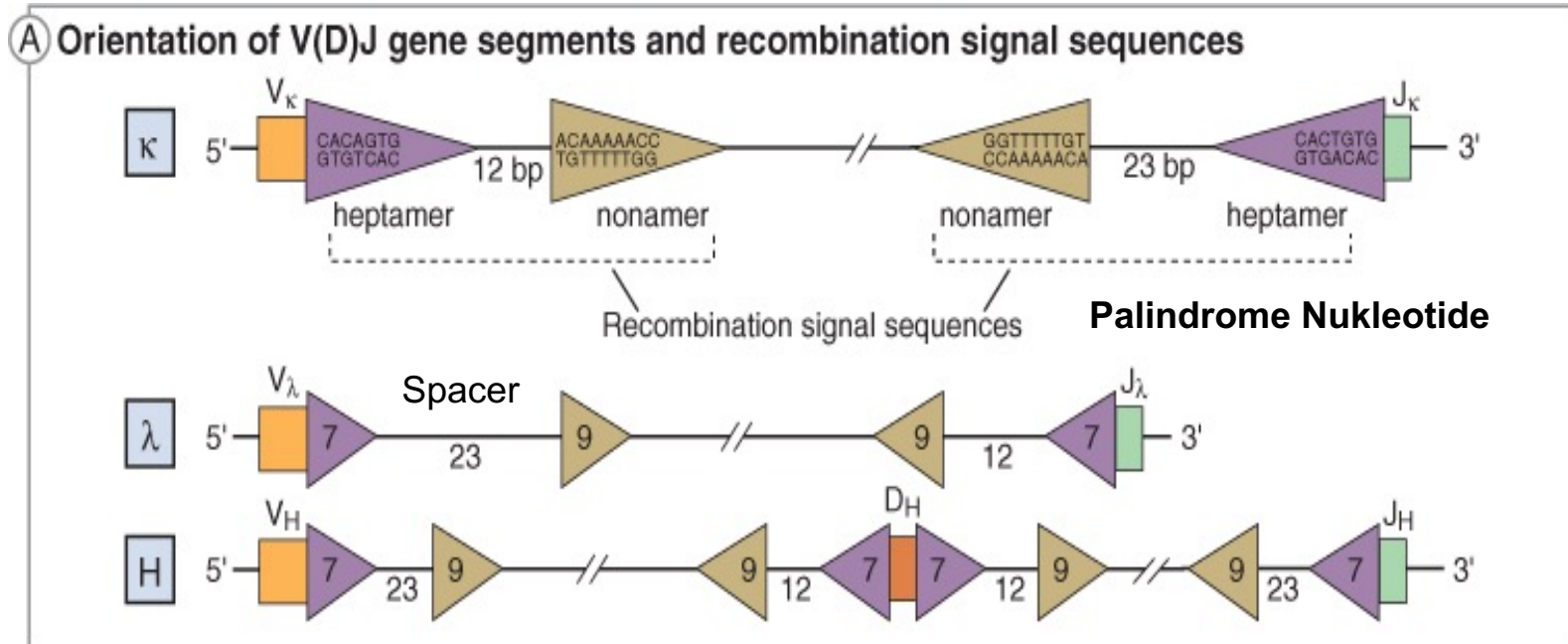
Molecular mechanism of the gene rearrangement

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): →
N-Nukleotide-addition – random addition of nucleotides

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

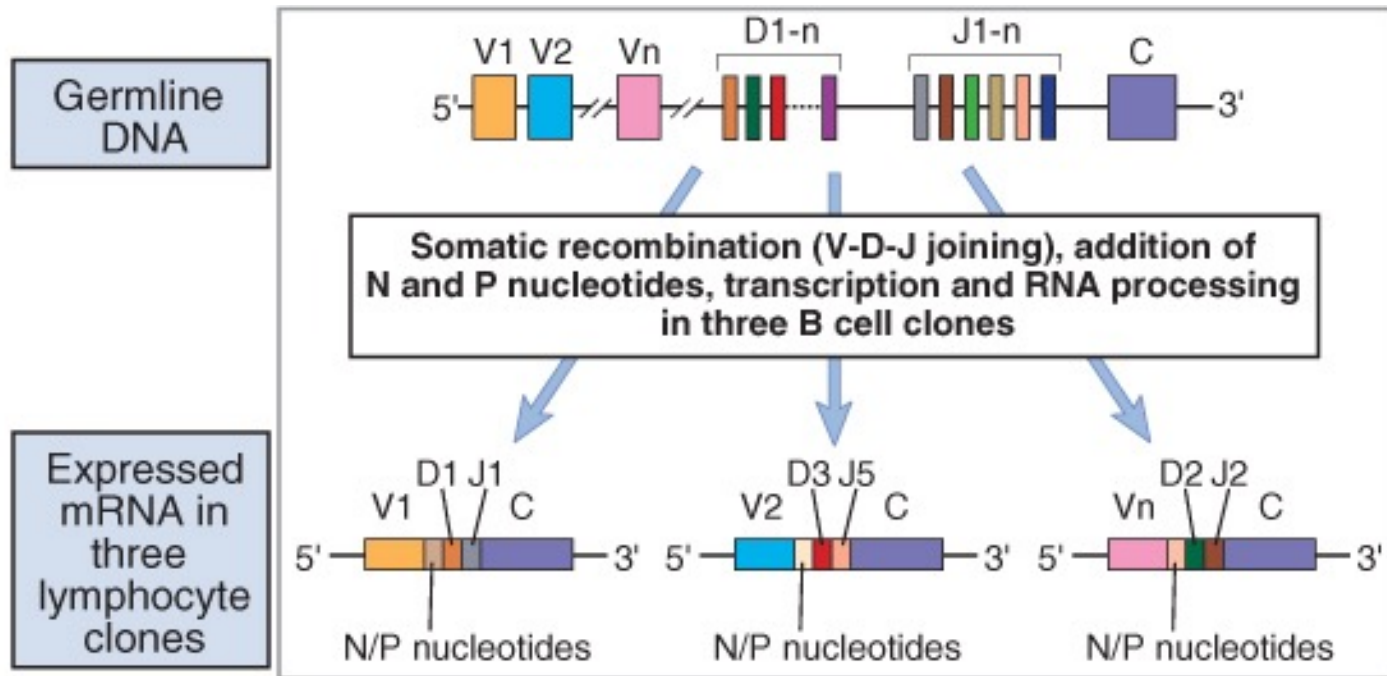


© Elsevier 2005. Abbas | RSS= Rekombinations Signal Sequenz | www.studentconsult.com

Recombination-Signal-Sequence (RSS):

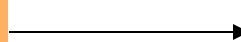
Contains a conserved heptamer and nonamer sequences which are divided by a non-conserved spacer sequence of either 12 or 23 basepairs.

Heavy chain gene rearrangement in three pro-B-cells



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Random gene rearrangement



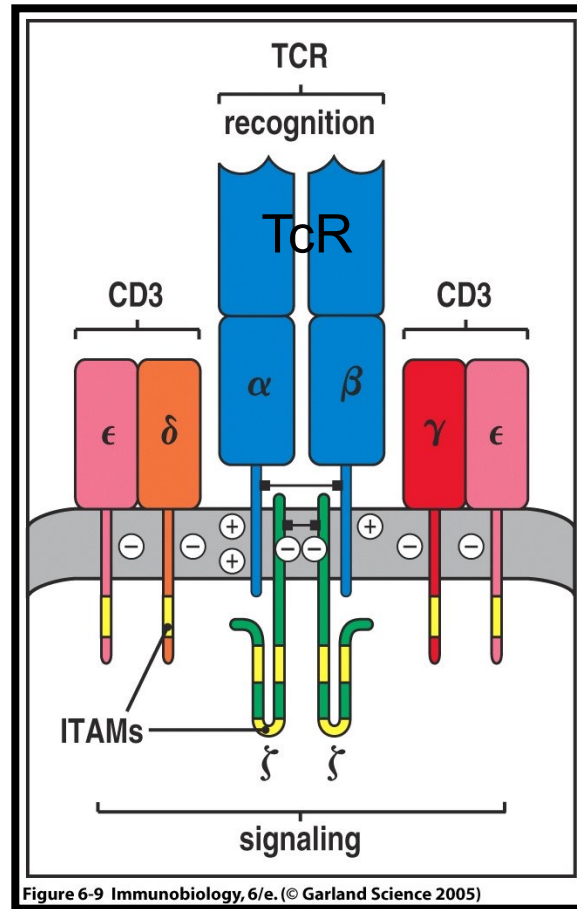
Diversity

T-cell-receptor (TcR)

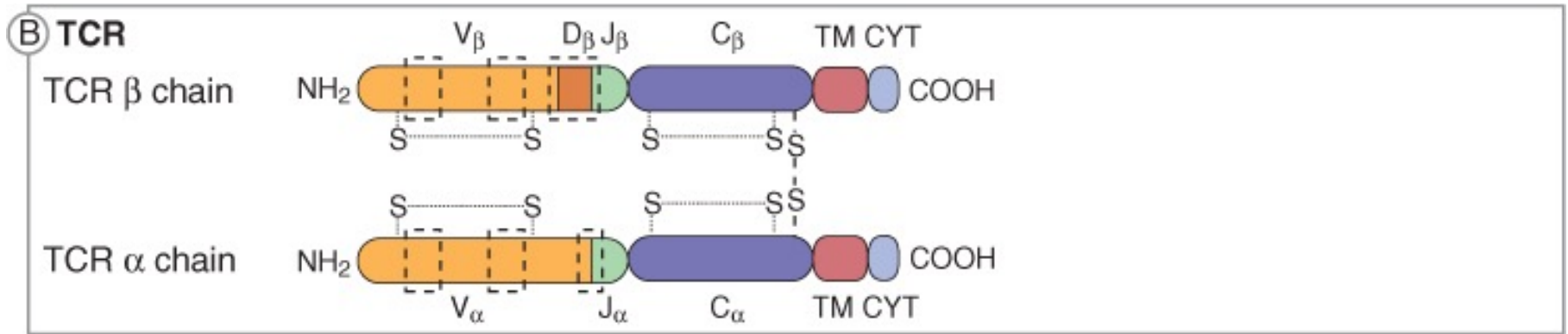
T-cell-types:

1. $\alpha\beta$ TcR+

2. $\gamma\delta$ TcR+



TcR α - β chains



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Human TCR encoding genes

V, J, D and C genes

β locus:

50 –100 V, 1D,
6 J, 1 C genes

α locus:

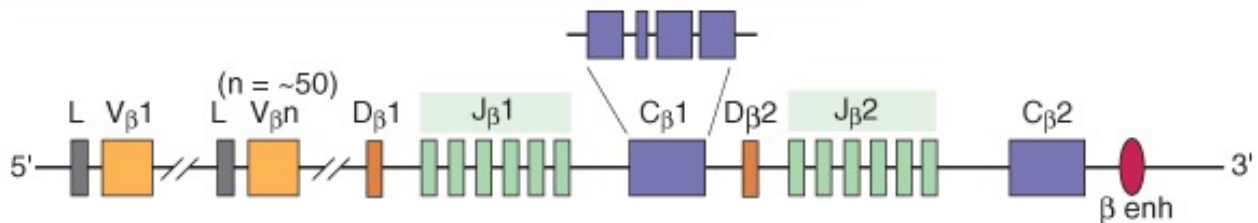
45 V, 55 J genes

inserted: δ locus:

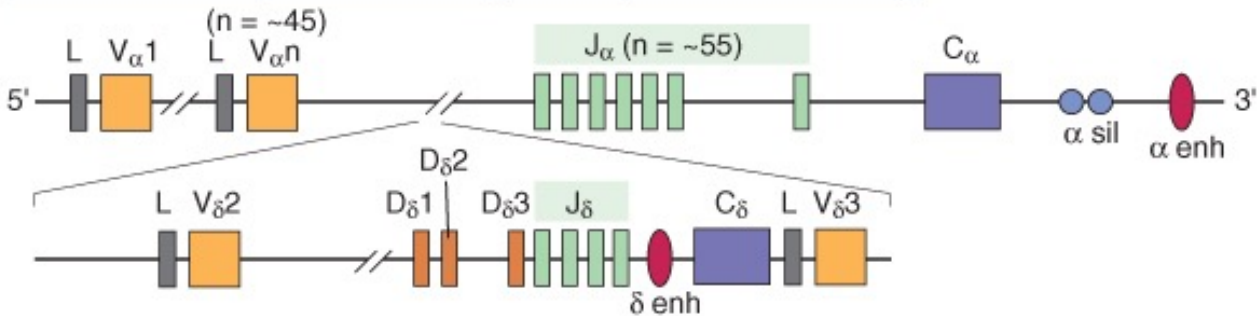
V, D, J and C genes

γ locus: V, J and C

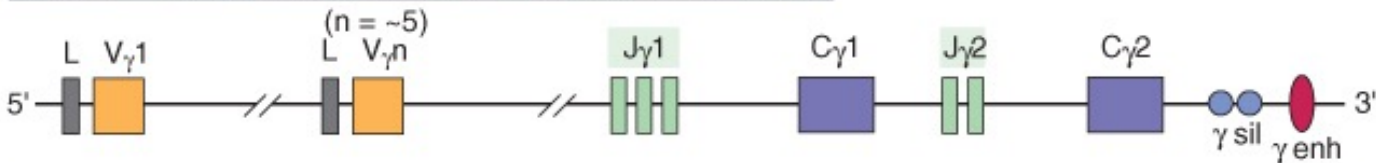
Human TCR β chain locus (620 kb; chromosome 7)

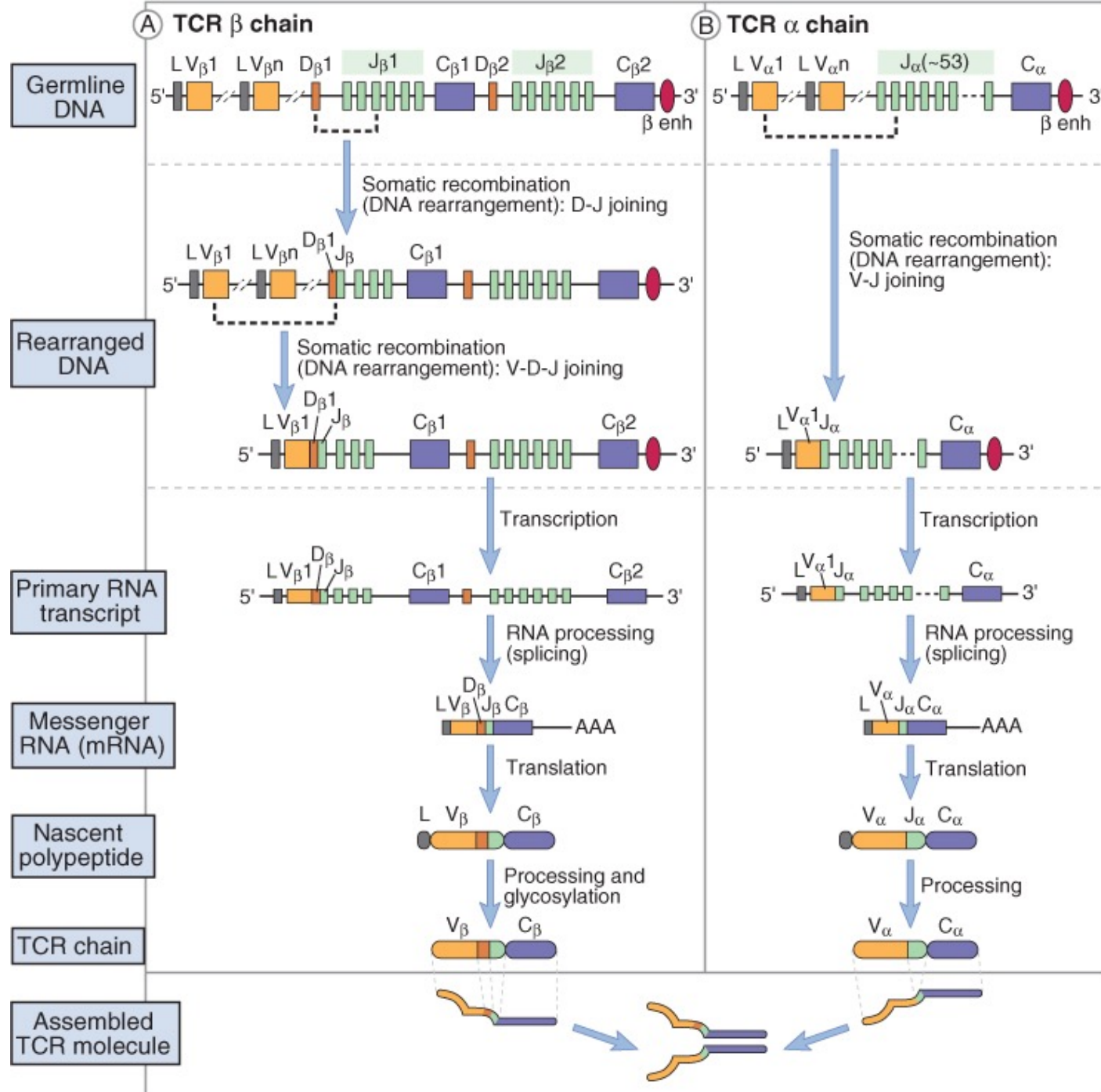


Human TCR α, δ chain locus (1000 kb; chromosome 14)



Human TCR γ chain locus (200 kb; chromosome 7)





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 \rightarrow CDR3 variability is higher
- Random combination of TcR α/β and γ/δ chains (like Ig H/L chains)

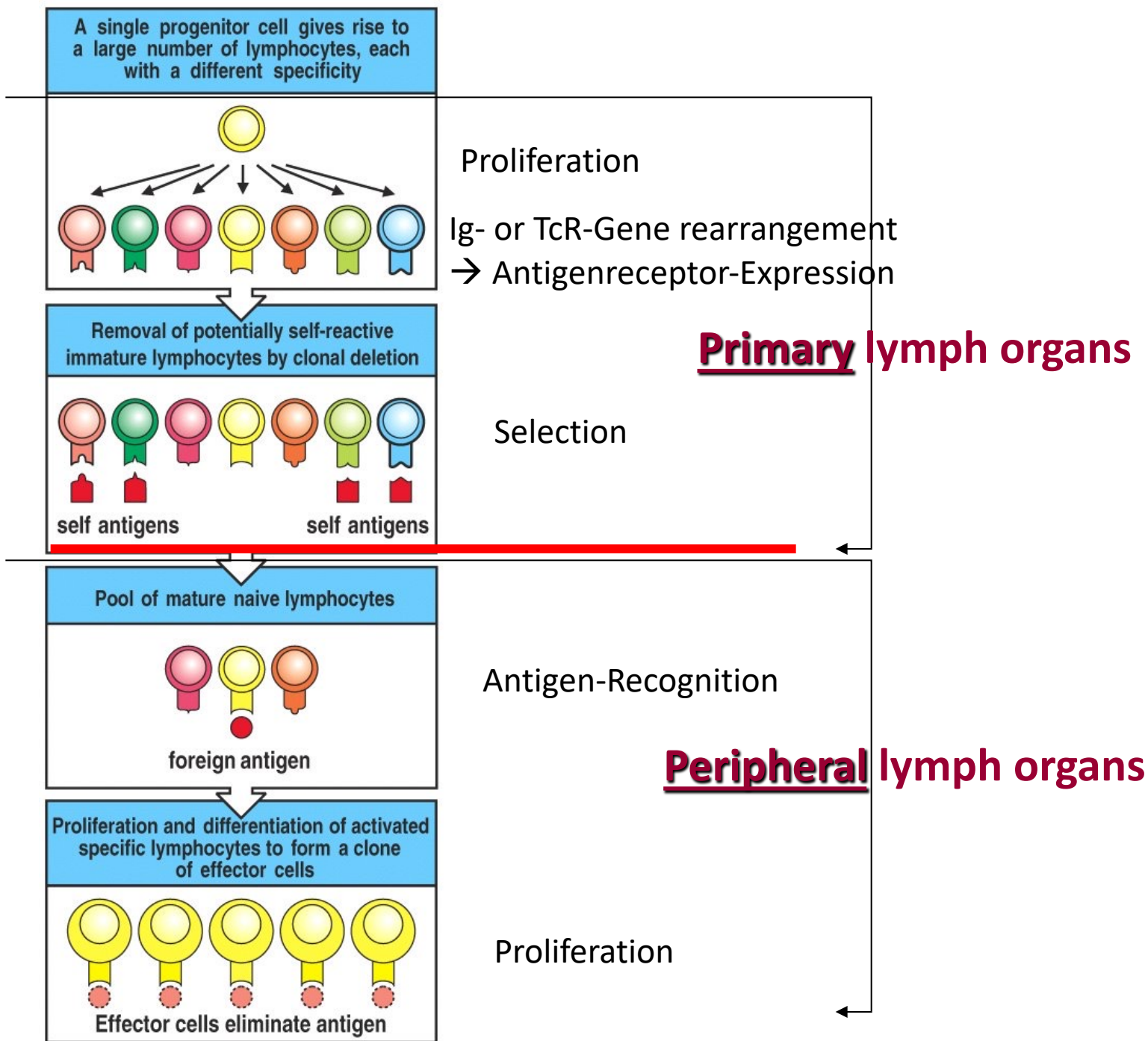


Figure 1-14 Immunobiology, 6/e. (© Garland Science 2005)

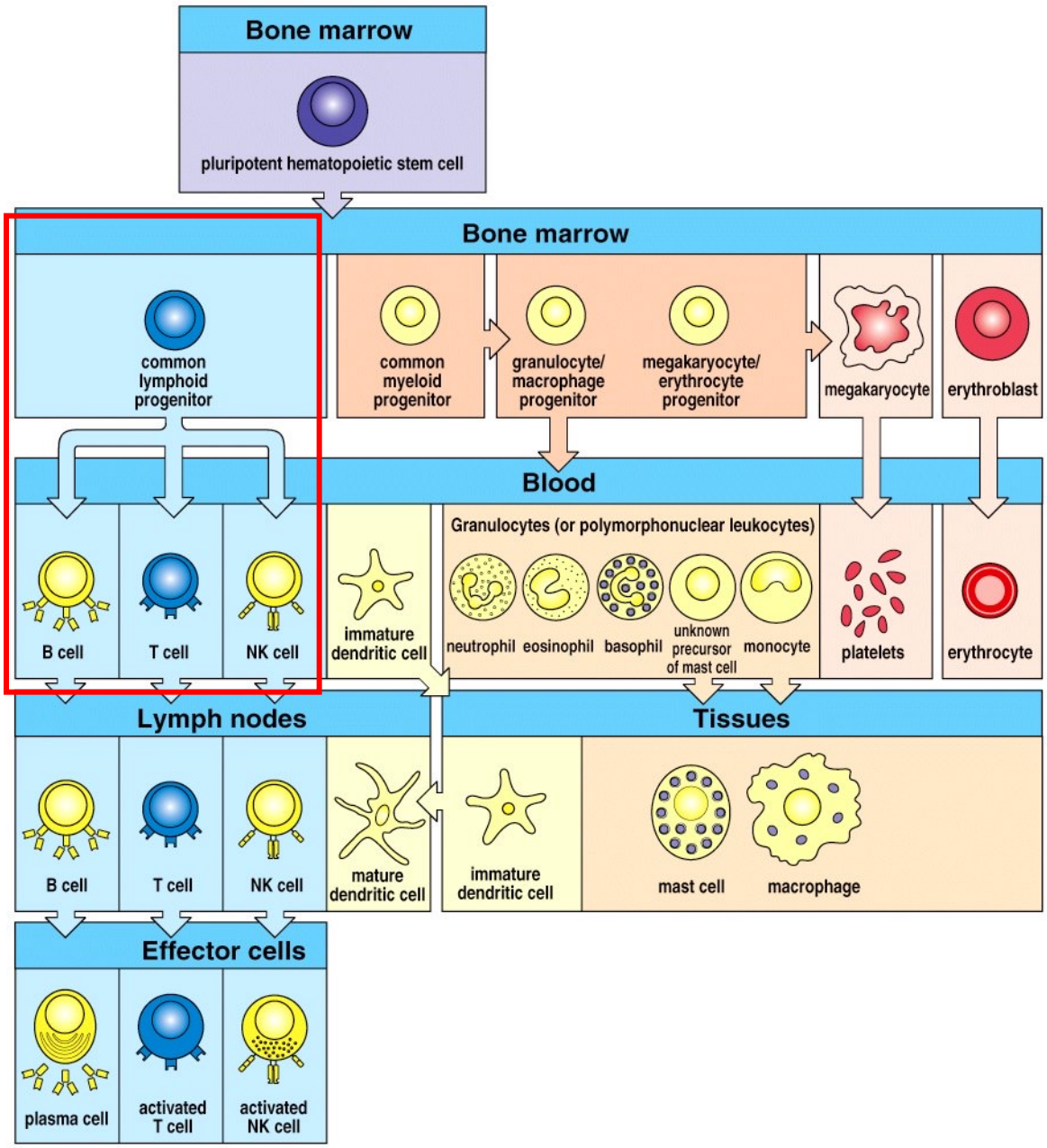
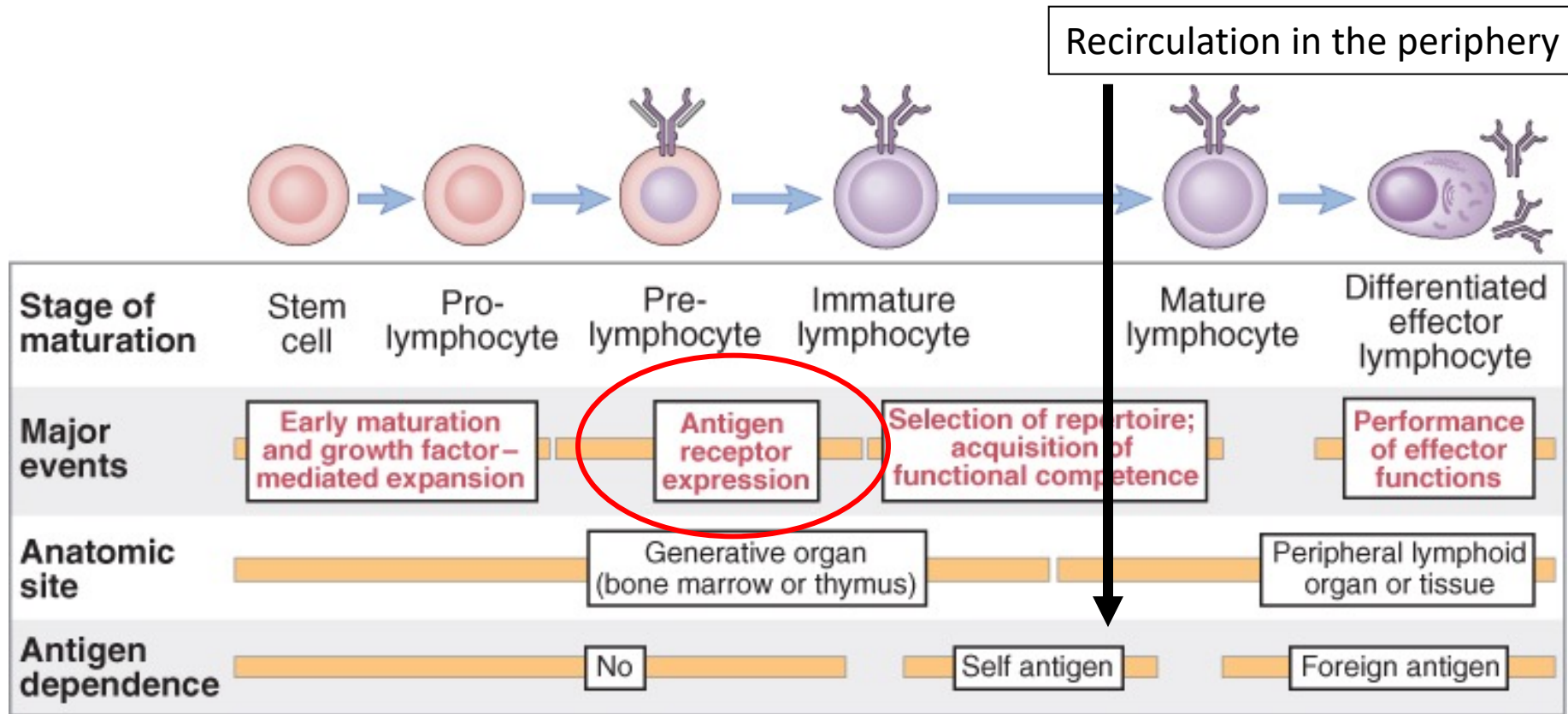


Figure 1-3 Immunobiology, 6/e. (© Garland Science 2005)

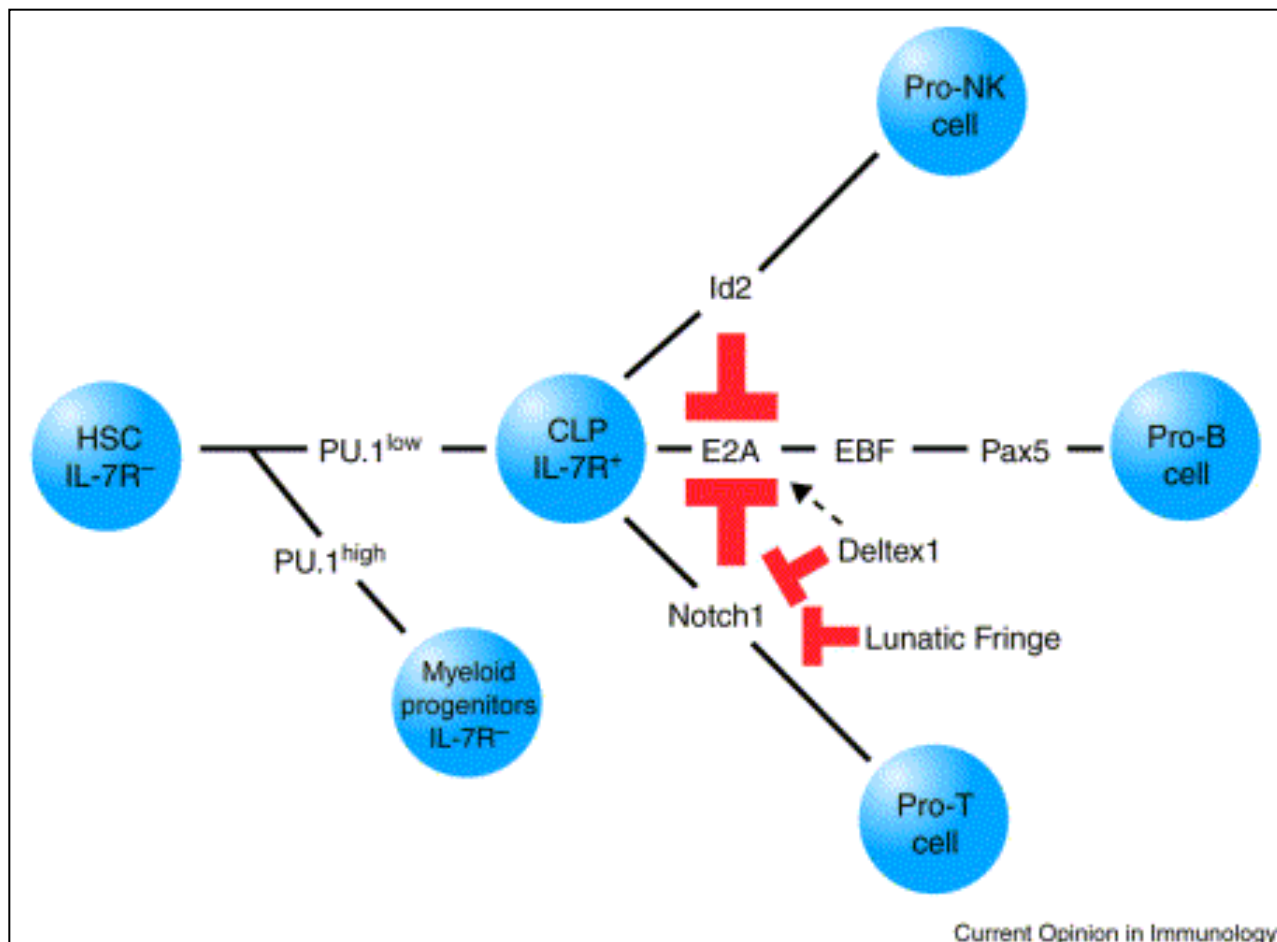
General characteristics of lymphocyte differentiation

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

Steps of lymphocyte development



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. Response modifiers: Wnt factors, ECM components.
4. Chemokine-production: SDF-1/CXCR4 ligand.

Elements of B:stromal interactions

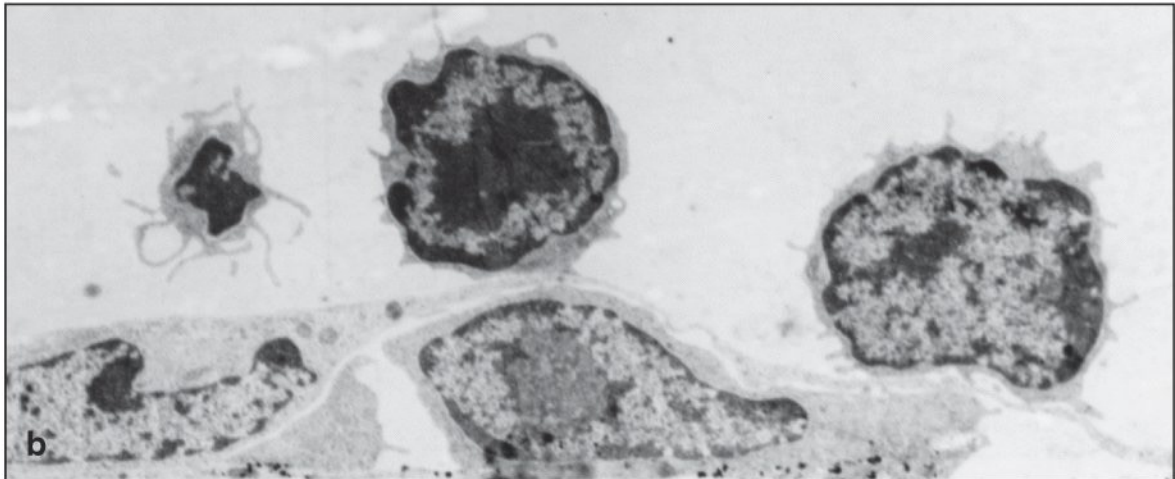
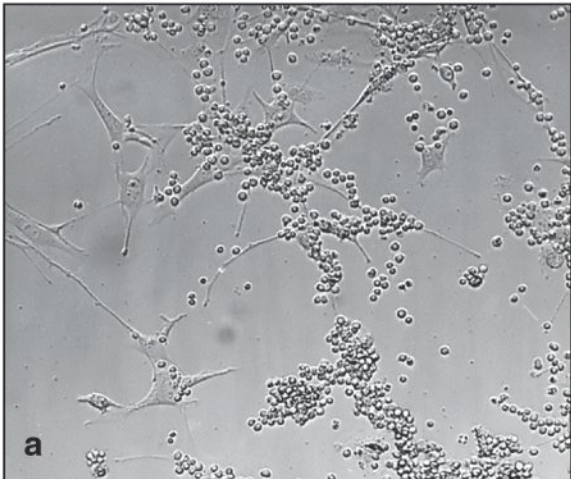
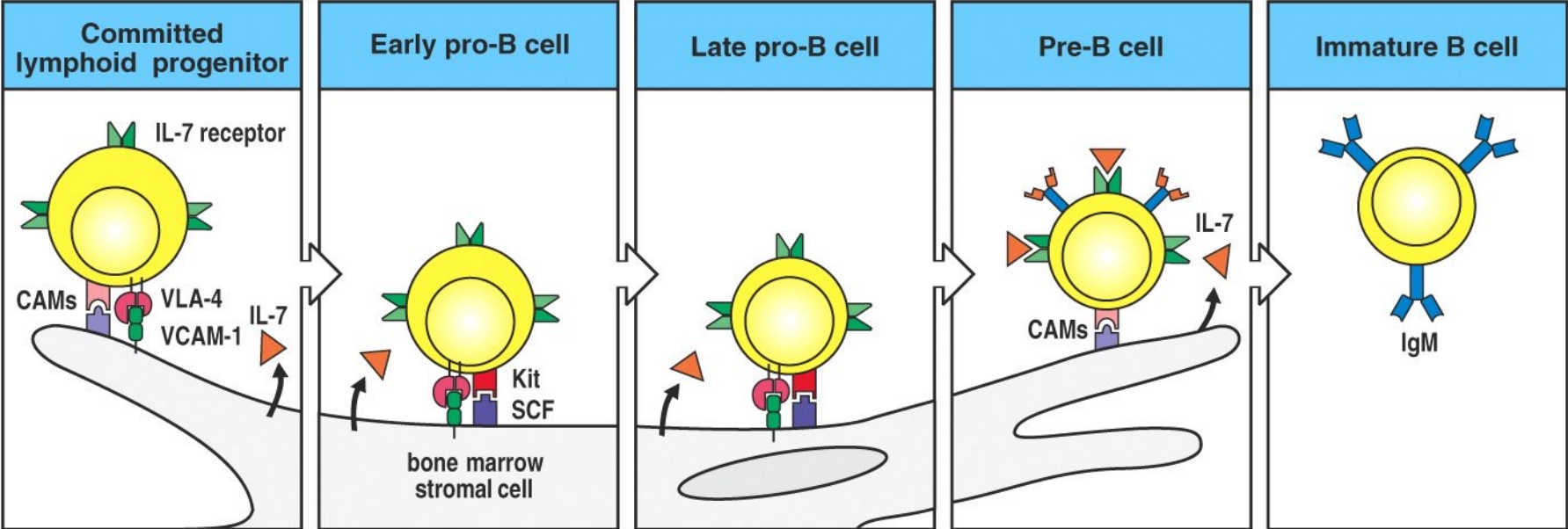


Figure 7-4 Immunobiology, 6/e. (© Garland Science 2005)

B-cell development I: HSC > “Large pre-B”

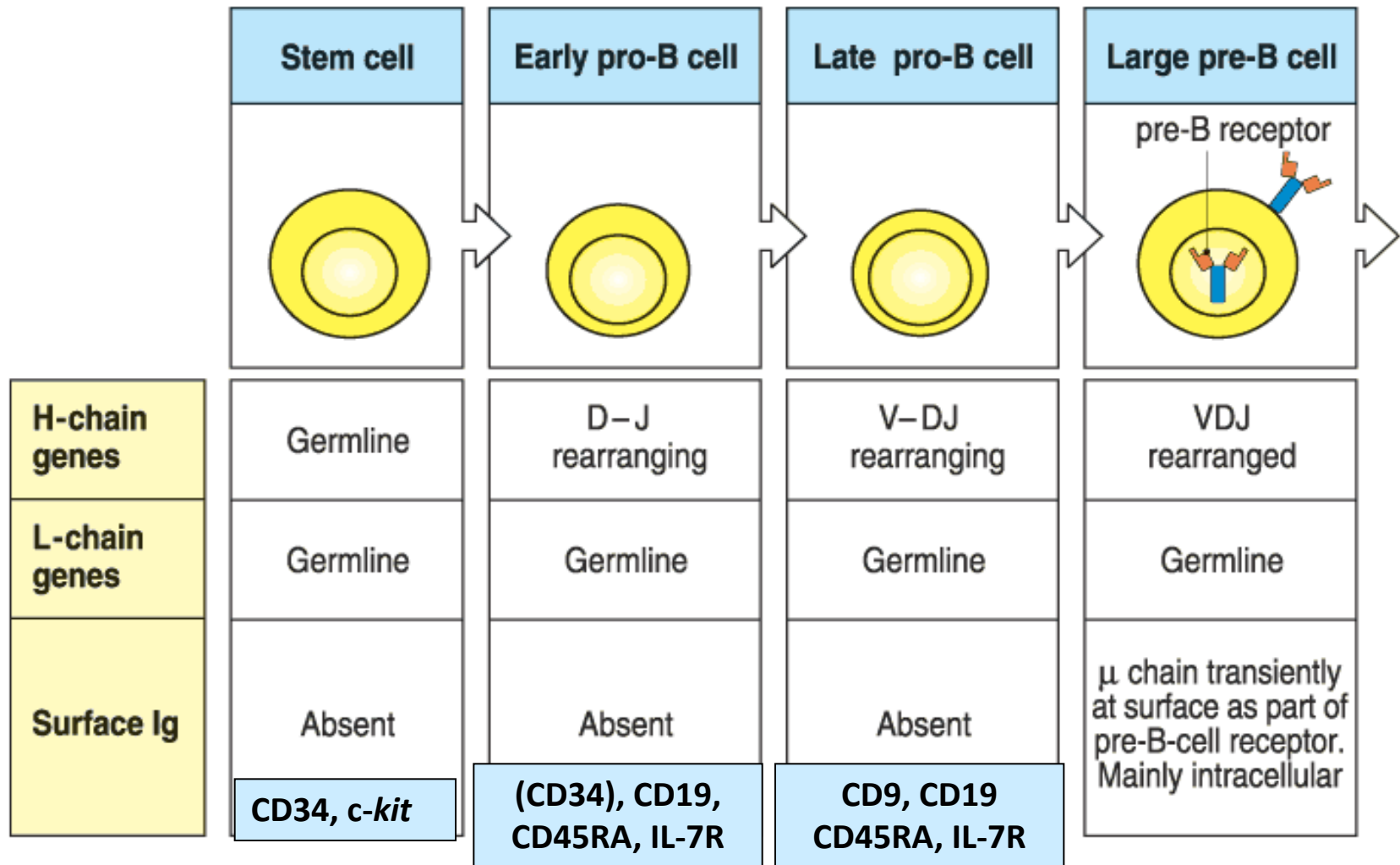


Fig 7.5 part 1 of 2 © 2001 Garland Science

B-cell development II “Small pre-B” > “mature B”

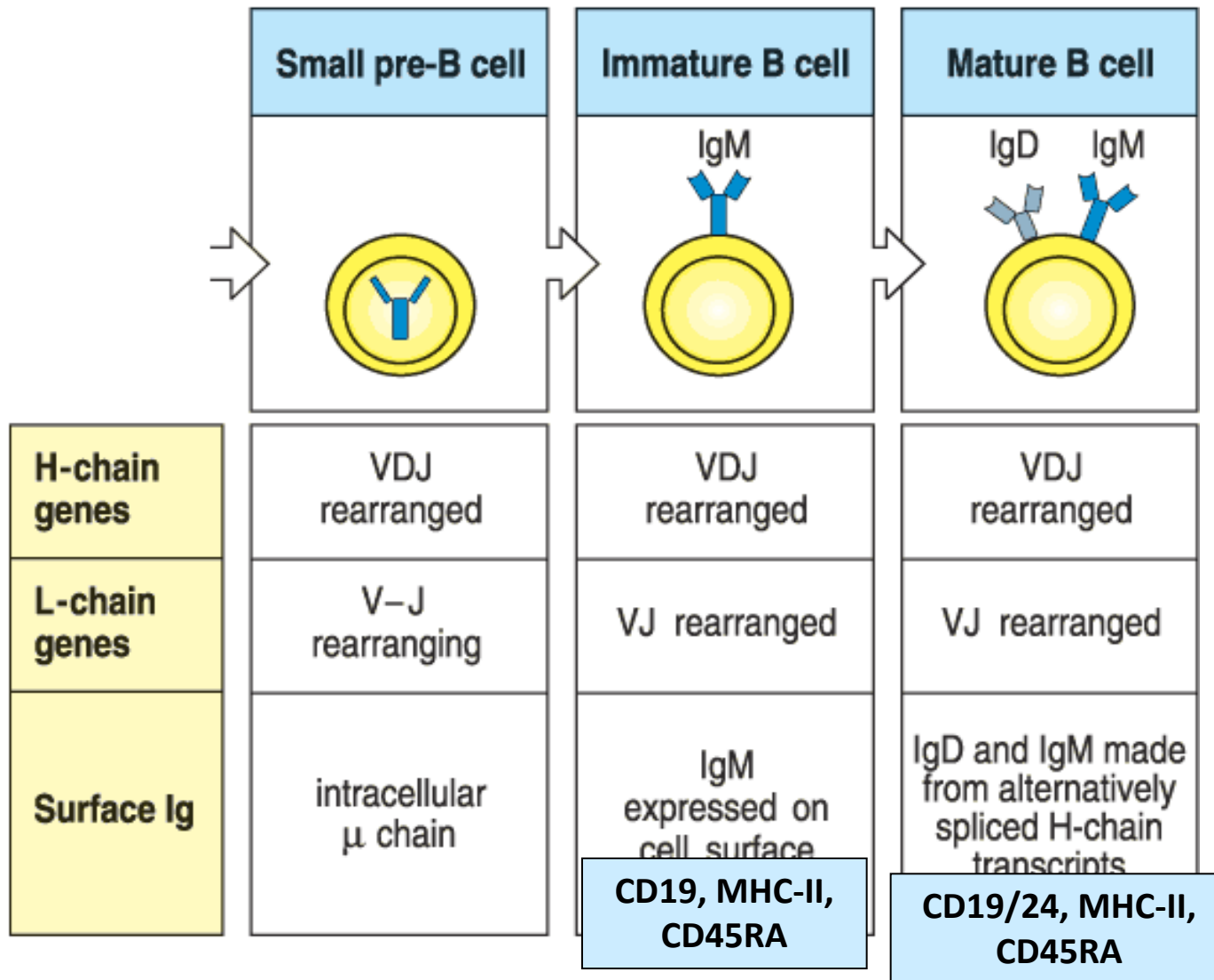


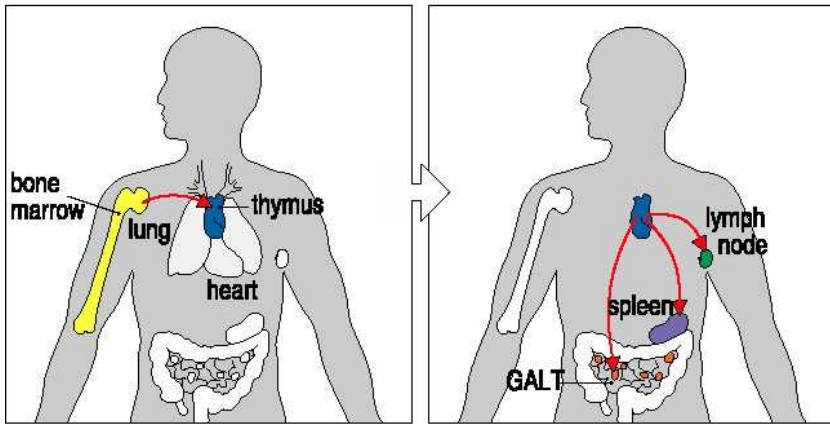
Fig 7.5 part 2 of 2 © 2001 Garland Science

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

T-cell development in the thymus.

Figure 5.1



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Production of T cell repertoire

Total repertoire:
TCR α , β : 10^{15}
TCR γ , δ : 10^{16}

T cell precursors are produced in the bone marrow from the common haemopoietic stem cell
They migrate through the blood circulation to the thymus

In the thymus: T cell maturation, educational steps
„double recognition” (MHC and peptide)

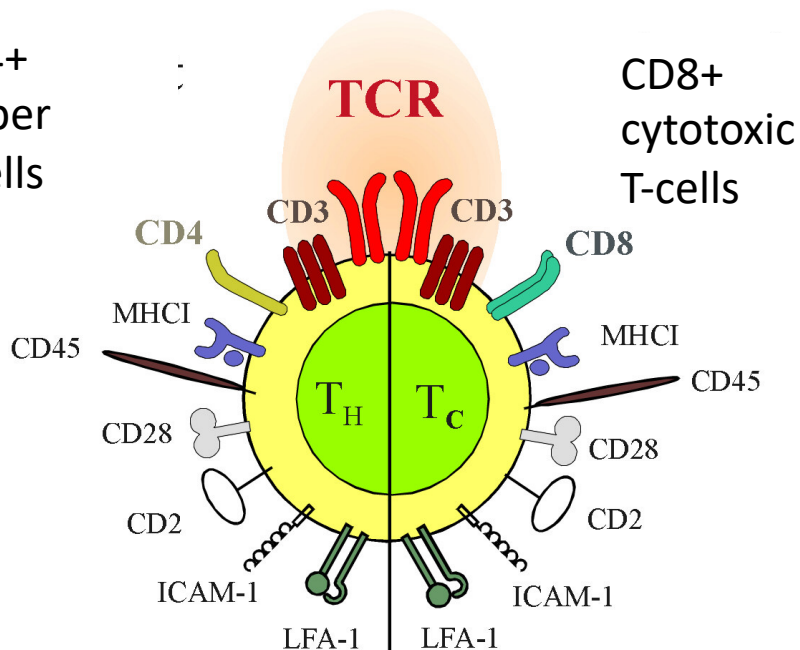
Periphery: mature, TCR expressing,
CD4 or CD8 positive T cells

Self-MHC restricted
Self-tolerant T cells

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

T-lymphocytes with $\alpha\beta$ TcR

CD4+
helper
T cells

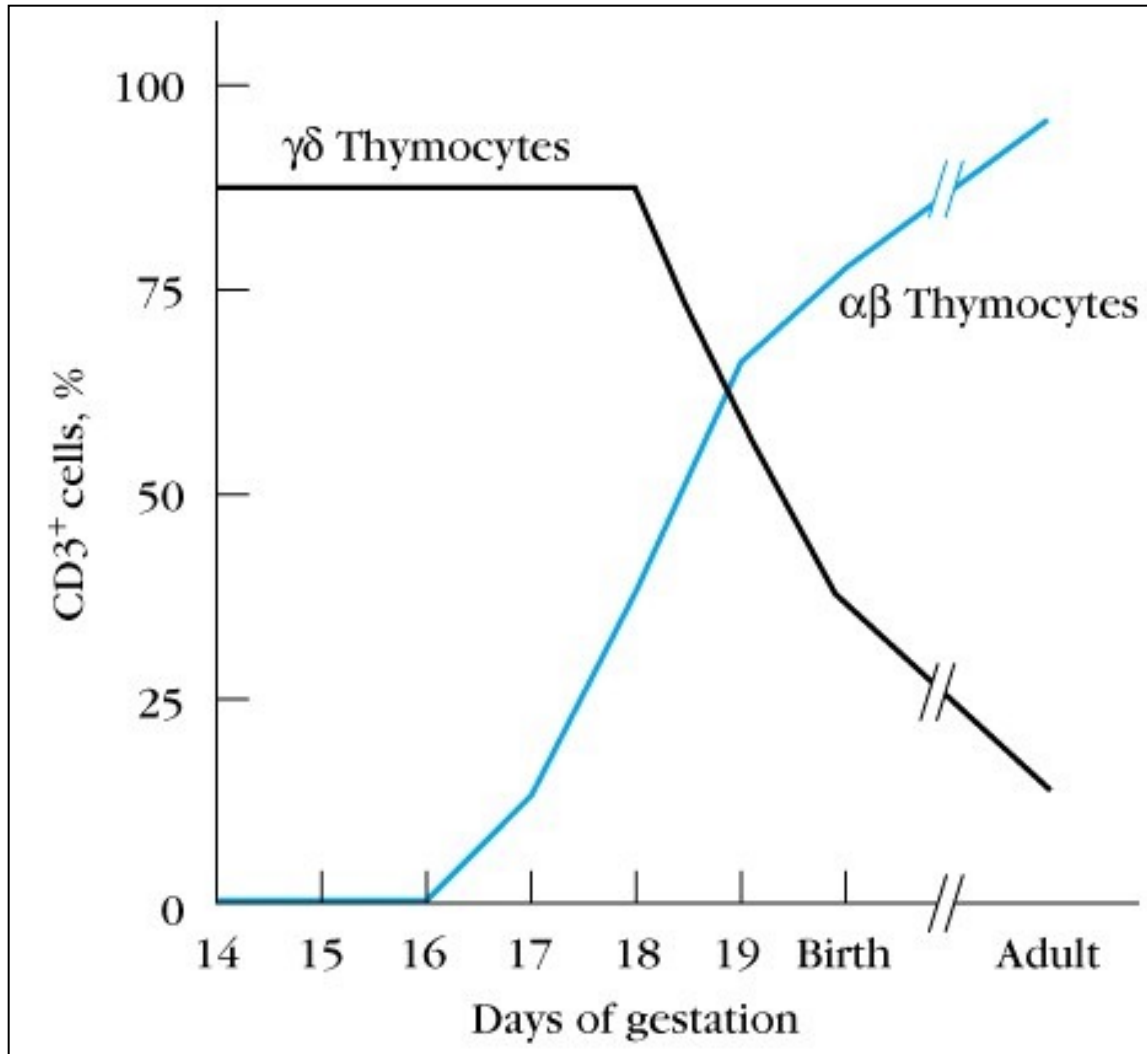


T-lymphocytes with $\gamma\delta$ TcR

- CD4-CD8- cytotoxic T-cells

Intraepithelial – with lower TcR diversity

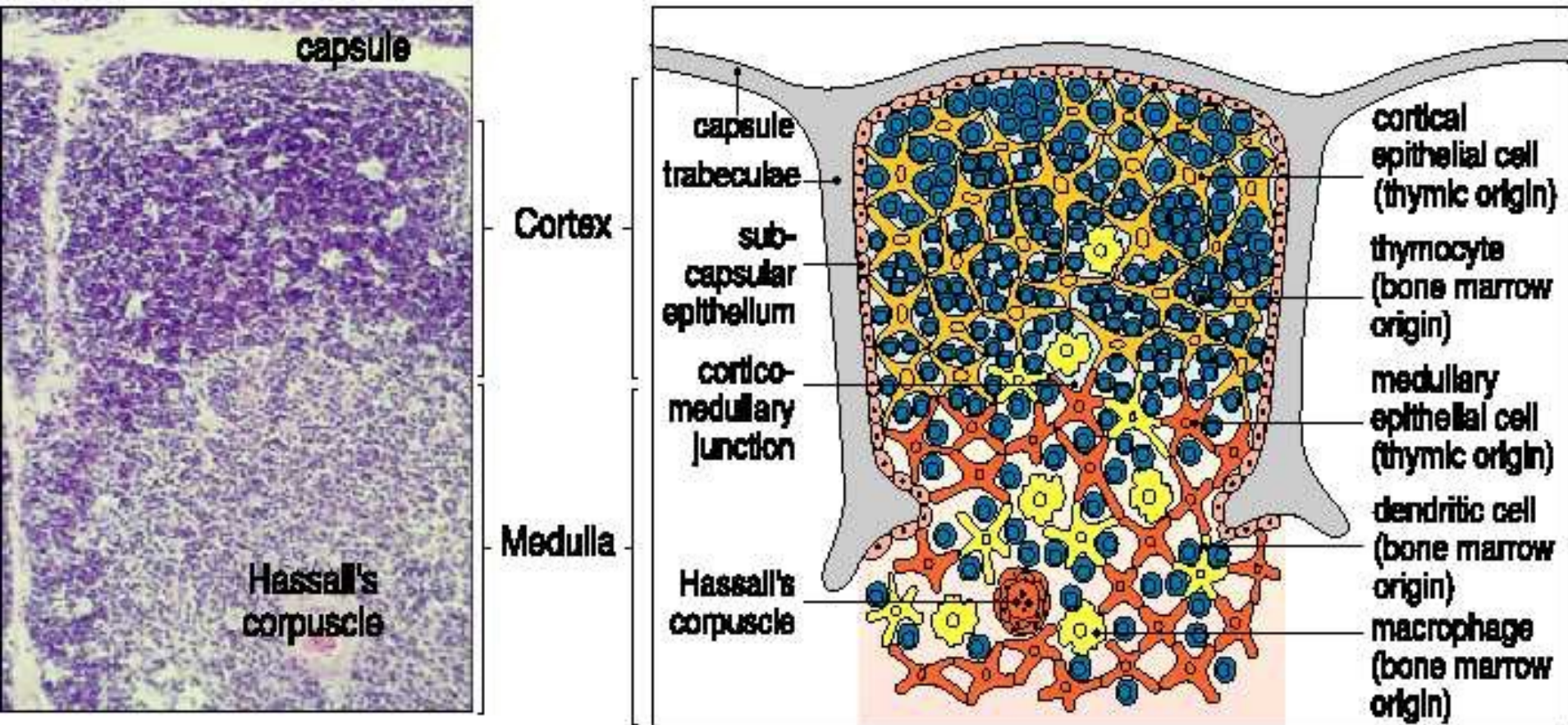
Lymphoid tissues – highly diverse receptors
Regulatory cytokine production



Full repertoire:
TCR α , β : 10^{15}
TCR γ , δ : 10^{16}

Structure of the thymus

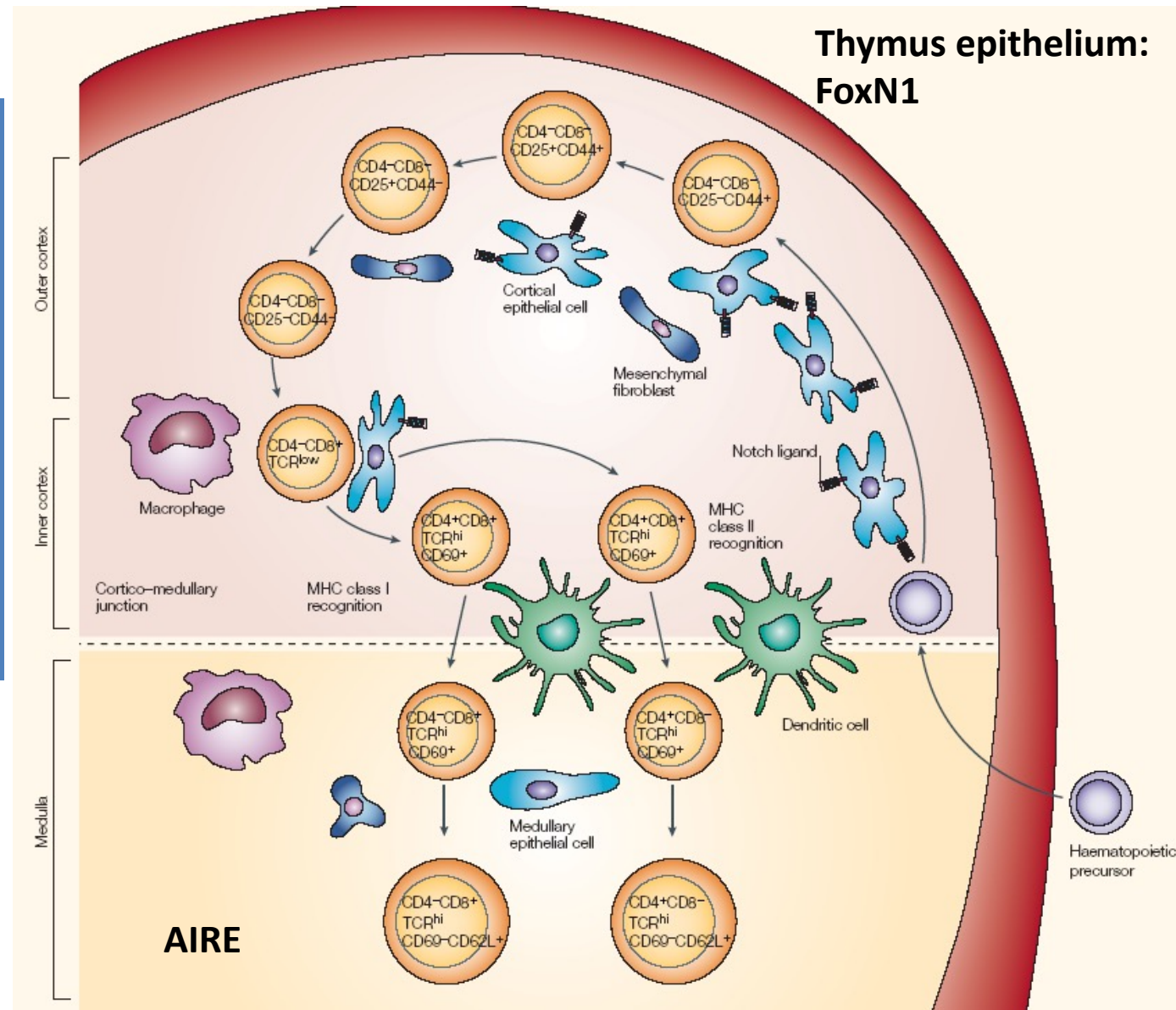
Figure 5.3



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

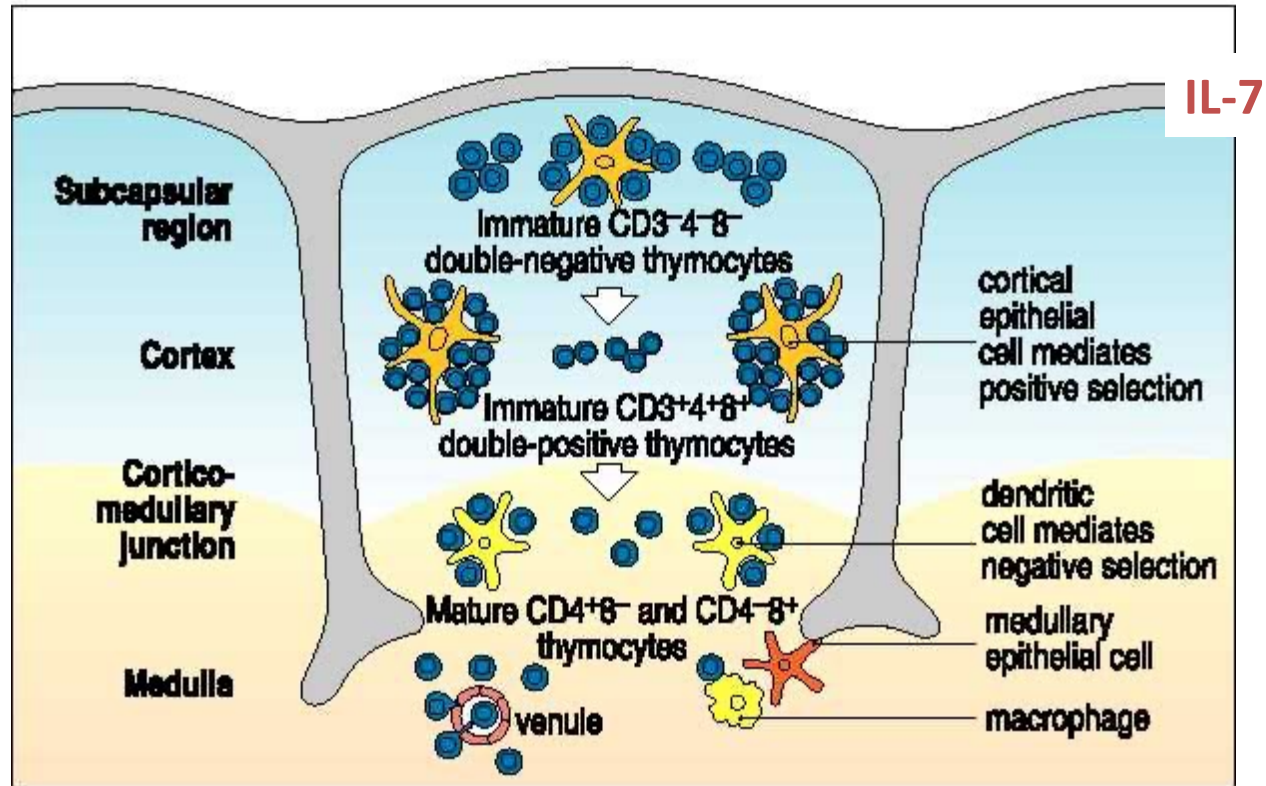
Thymic Microenvironment and T-cell Development

1. Migration:
Chemokine effect
2. Proliferation
IL-7
3. Differentiation
 - TcR-rearrangement
 - Phenotypes
4. Selection
Apoptosis



T-cell development in the thymus

Figure 5.14



Thymocytes:

DN: 2-5 %

DP: 70-80%

CD4 SP: 10-15%

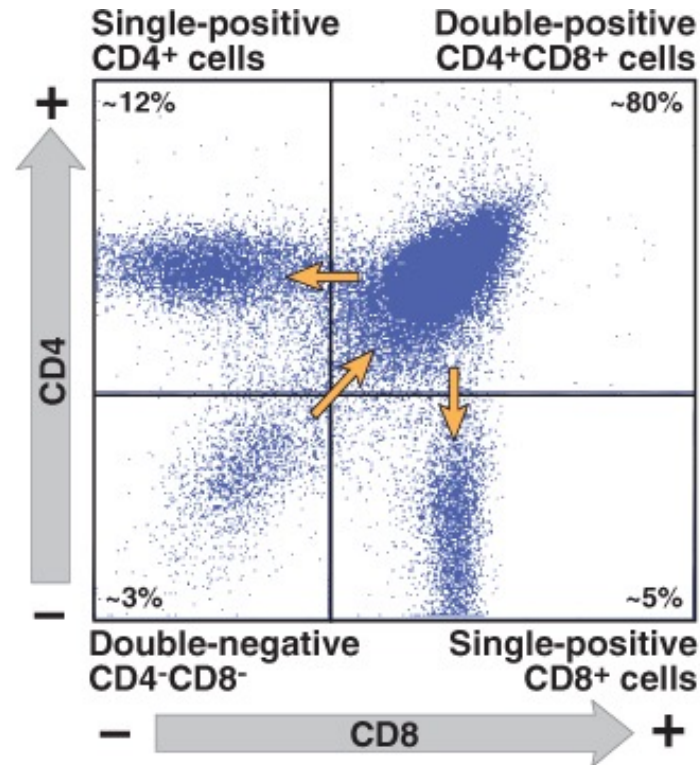
CD8 SP: 5-8%

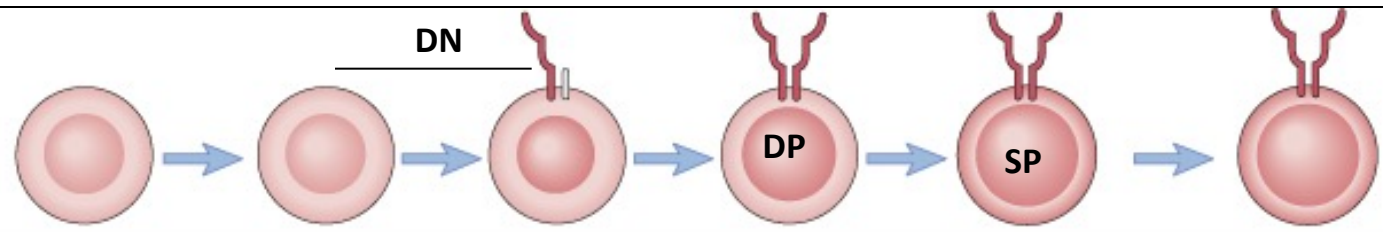
Young mouse: 5×10^7 T-cells daily

During selection 98 % of thymocytes die by apoptosis

Daily $1-2 \times 10^6$ mature T-cell migrate to the periphery

Thymocyte populations based on their cell surface markers





Stage of maturation	Stem cell	Pro-T	Pre-T	Double positive	Single positive (immature T cell)	Naive mature T cell
Proliferation	██████████		██████████			
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 [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	<i>c-kit</i> ⁺ CD44 ⁺ CD25 ⁻	<i>c-kit</i> ⁺ CD44 ⁺ CD25 ⁺	<i>c-kit</i> ⁺ CD44 ⁺ CD25 ⁺	CD4 ⁺ CD8 ⁺ TCR/CD3 ^{lo}	CD4 ⁺ CD8 ⁻ or CD4 ⁻ CD8 ⁺ TCR/CD3 ^{hi}	CD4 ⁺ CD8 ⁻ or CD4 ⁻ CD8 ⁺ TCR/CD3 ^{hi}
Anatomic site	Bone marrow	Thymus				Periphery
Response to antigen	None	None	None	Positive and negative selection	Negative selection	Activation (proliferation and differentiation)

Thymocyte from bone marrow

Rearrangement of TCR genes

Immature thymocyte

CD8
CD3
T-cell receptor
CD4

Positive selection of cells whose receptor binds MHC molecules

Death by apoptosis of cells that do not interact with MHC molecules

Class I and/or class II MHC molecules

Epithelial cell

THYMIC CORTEX

Negative selection and death of cells with high-affinity receptors for self-MHC or self-MHC + self-antigen

CD4⁺

T_H cell

CD8⁺

T_C cell

Mature CD4⁺ or CD8⁺ T lymphocytes

Macrophage

Dendritic cell

THYMIC MEDULLA

Positive selection:

Epithelial cell - thymocyte interaction in the thymus cortex

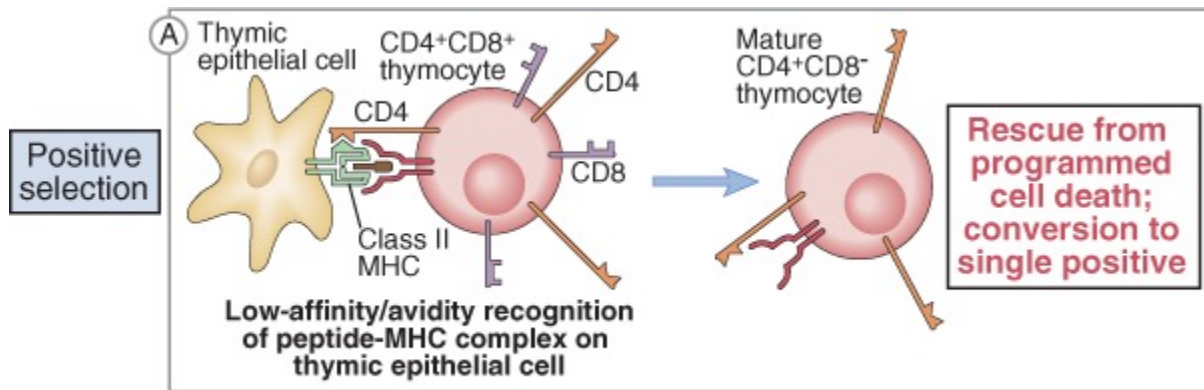
Survival of DP cells whose TcR is appropriate for self MHC recognition

Negative selection:

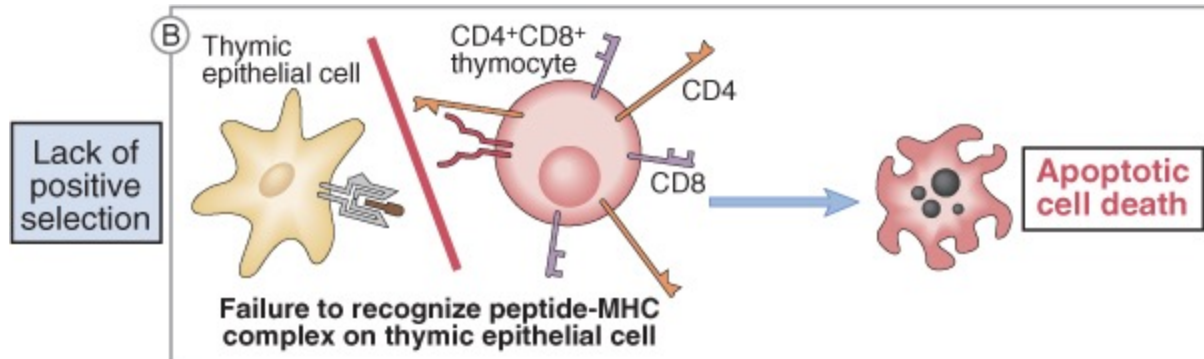
APC (macrophage or DC) – thymocyte interaction in thymus medulla

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

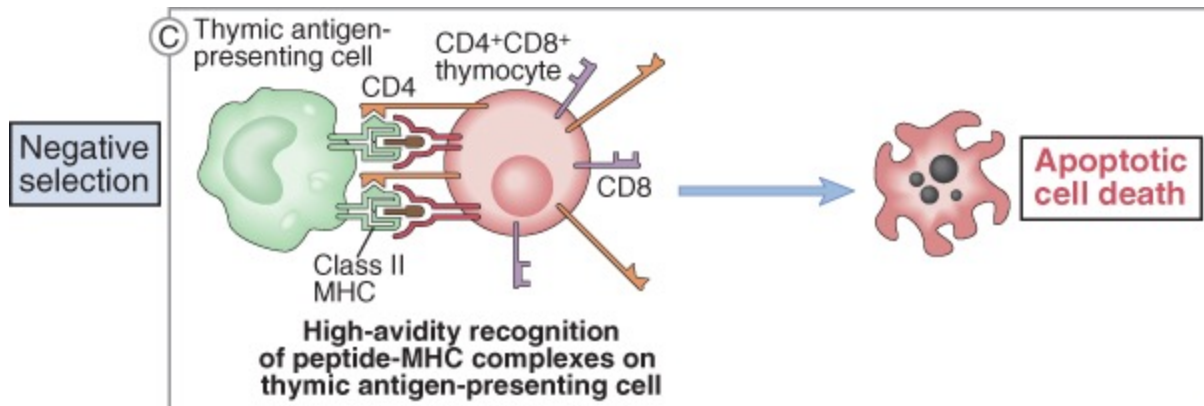
Differentiation into SP stage



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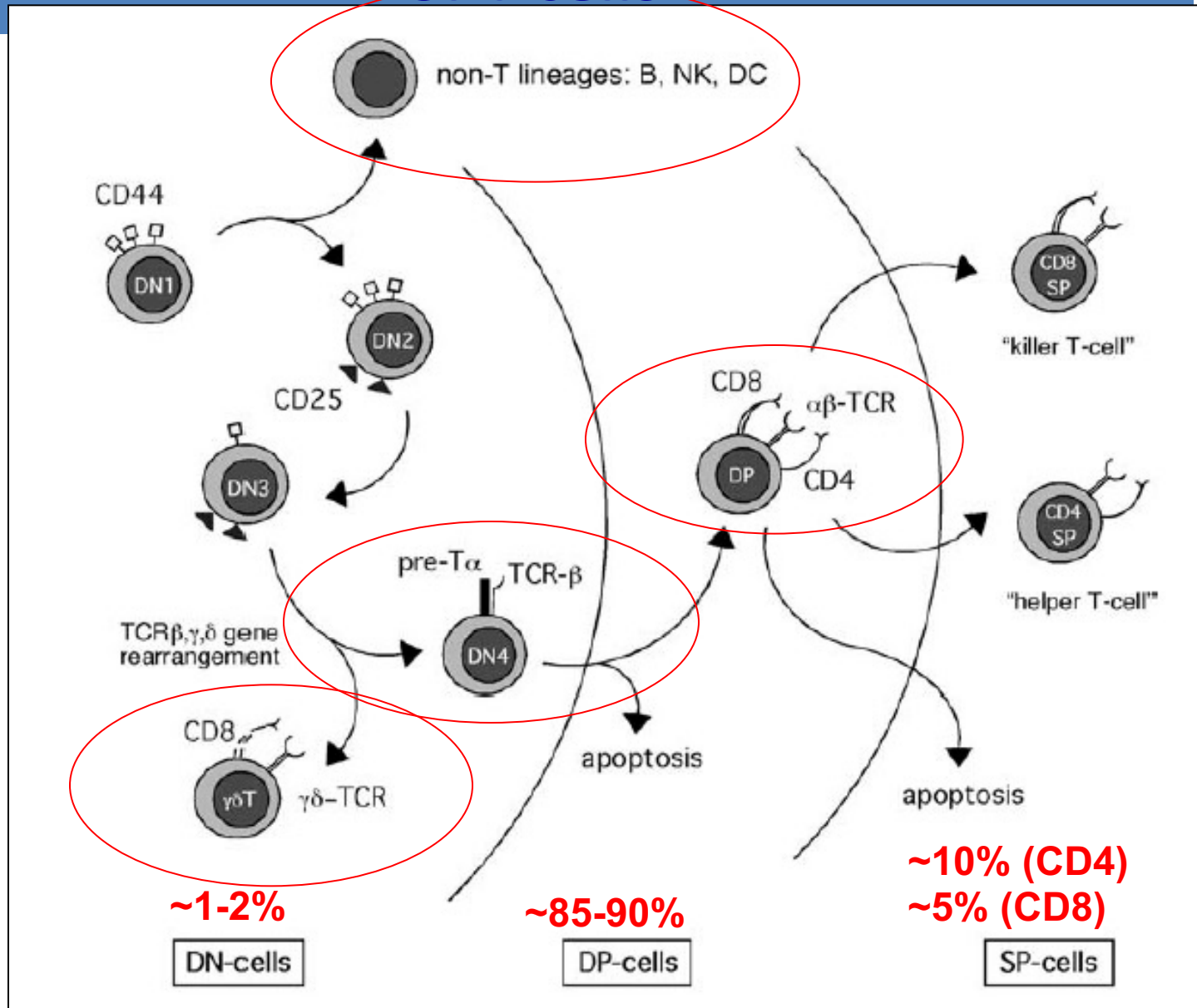


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Decision-making during the development of T cells



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

