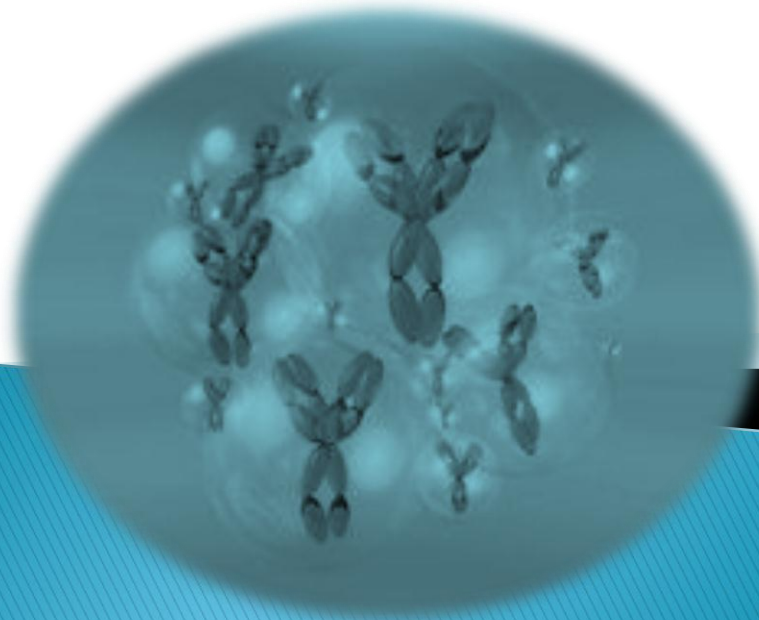


เอกสารประกอบการสอนวิชา Infection and Immunity (01419461)

เรื่อง immunoglobulin genes

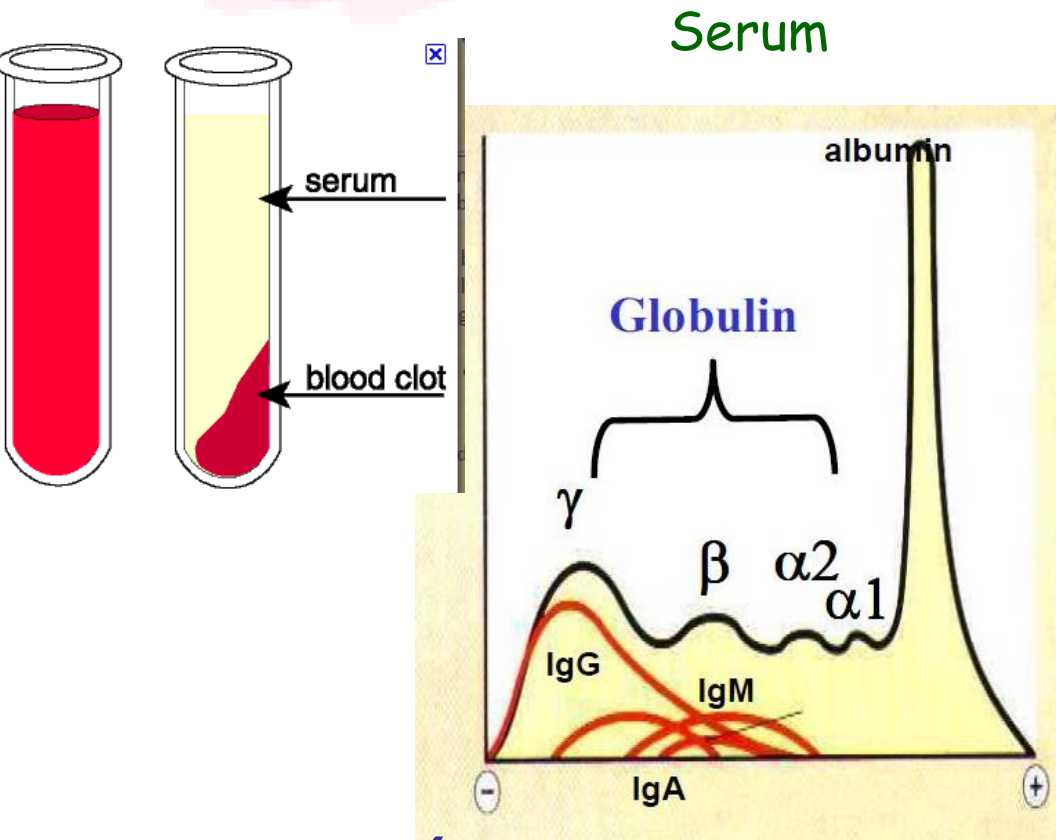
โดย อ.ดร. อิงอร กิมกง

ภาควิชาจุลชีววิทยา คณะวิทยาศาสตร์ มหาวิทยาลัยเกษตรศาสตร์

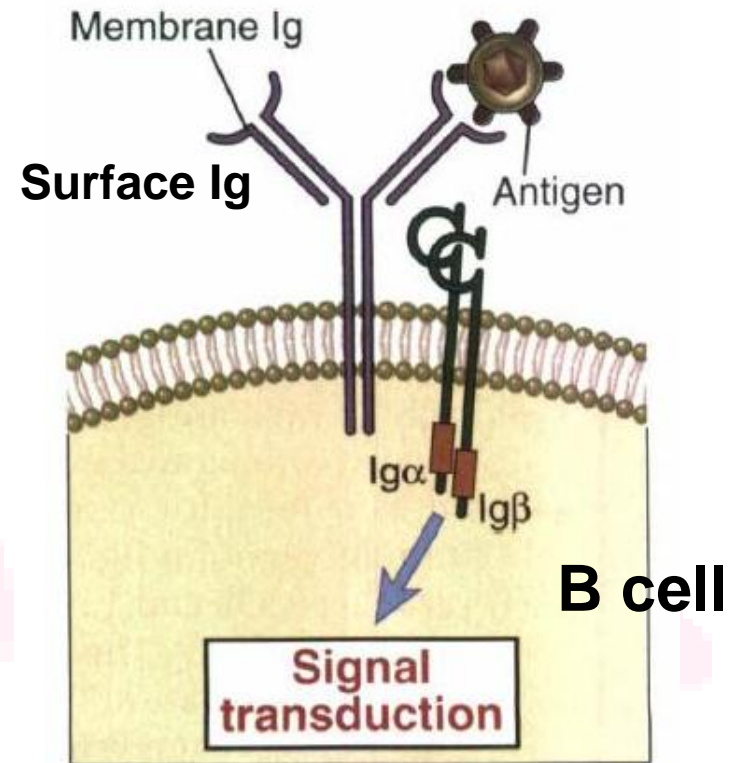


What are Immunoglobulins (Ig) ?

- ⇒ They are glycoproteins that are found in blood or other bodily fluids.
- ⇒ There are 2 forms of Ig: secreted and membrane-bound forms.

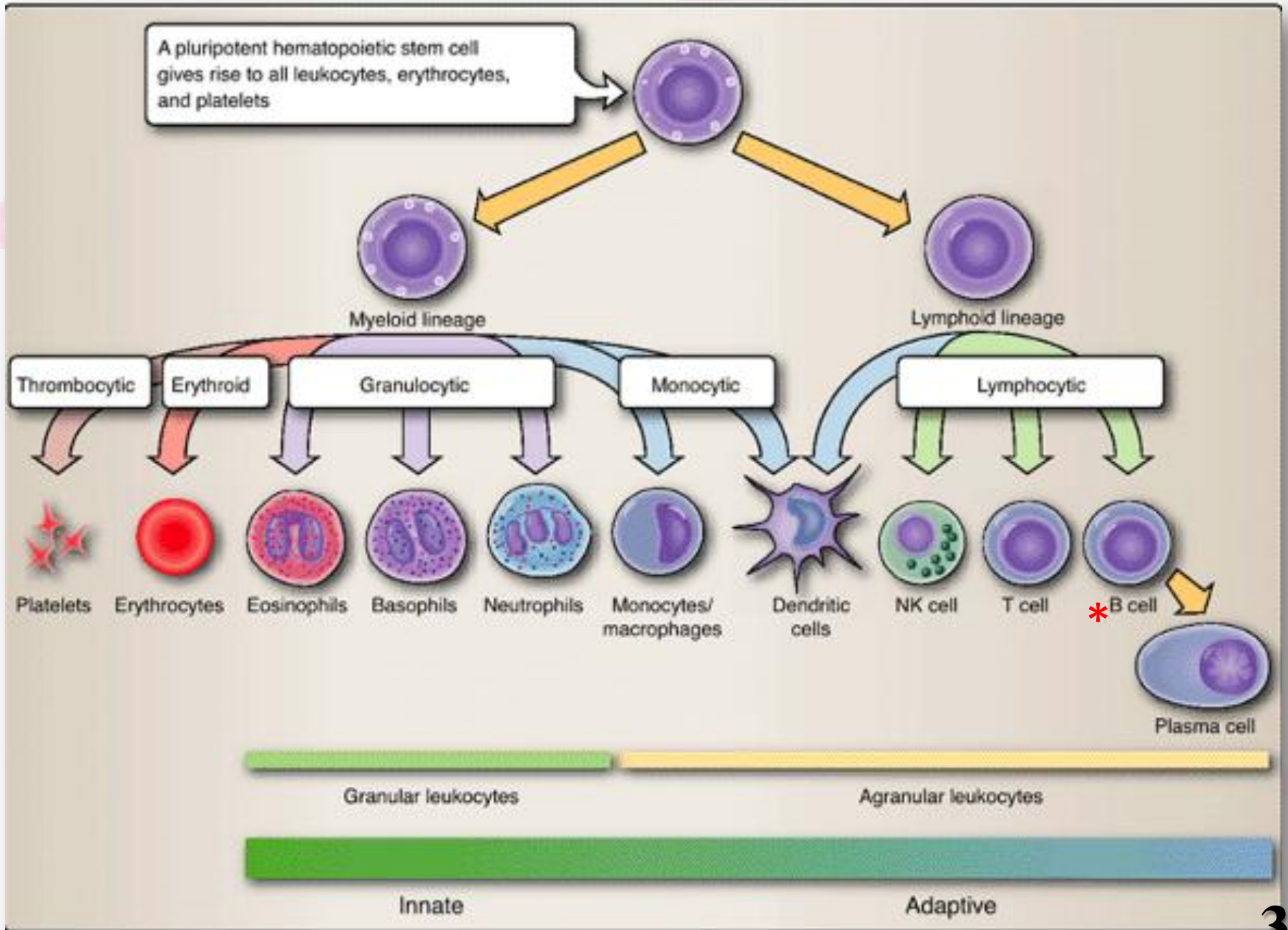


(A. Tiselius and E. A. Kabat, in 1939)



(Abbas AK; 6th ed, p138)

⇒ Ig are synthesized by B lymphocytes



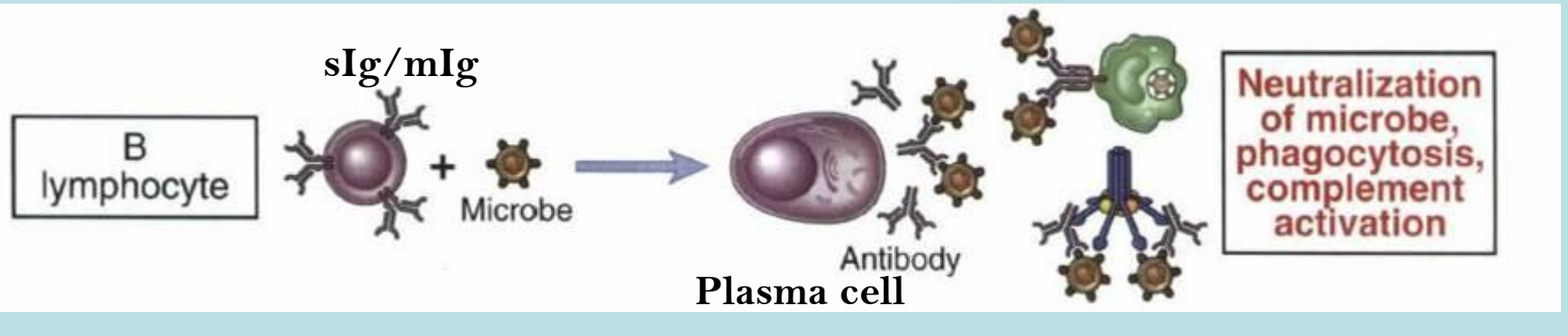
What are Immunoglobulins (Ig) ?

⇒ Ig are synthesized by B lymphocytes.

⇒ Its functions are ...

1. Surface/membrane Ig

2. Secreted Ig



(Abbas AK et al., Cellular and Molecular Immunology 6th Edition;2007;p12)

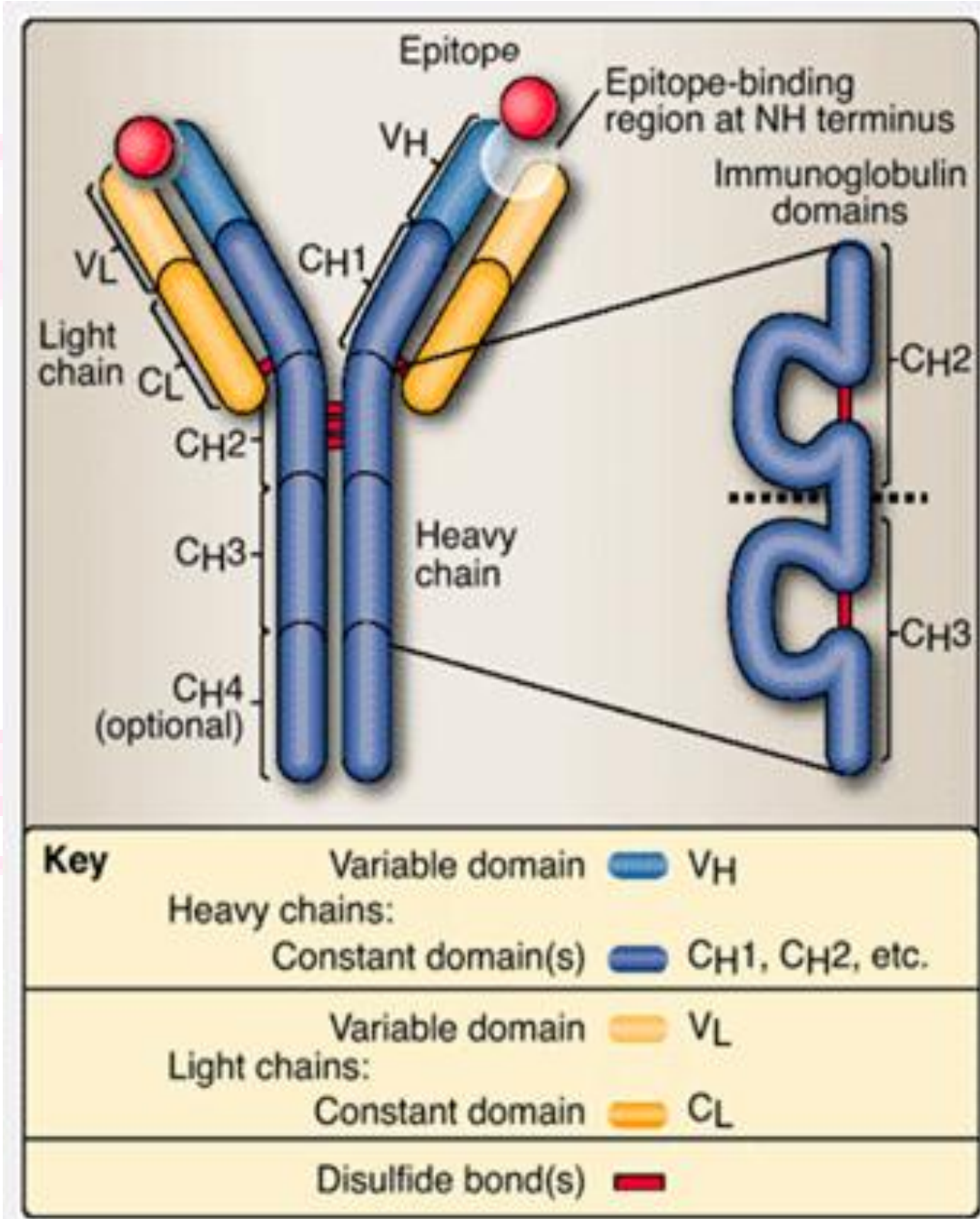
Basic structure of Ig

ประกอบด้วย polypeptide 4 สาย

- Light chain 2 สาย
- Heavy chain 2 สาย

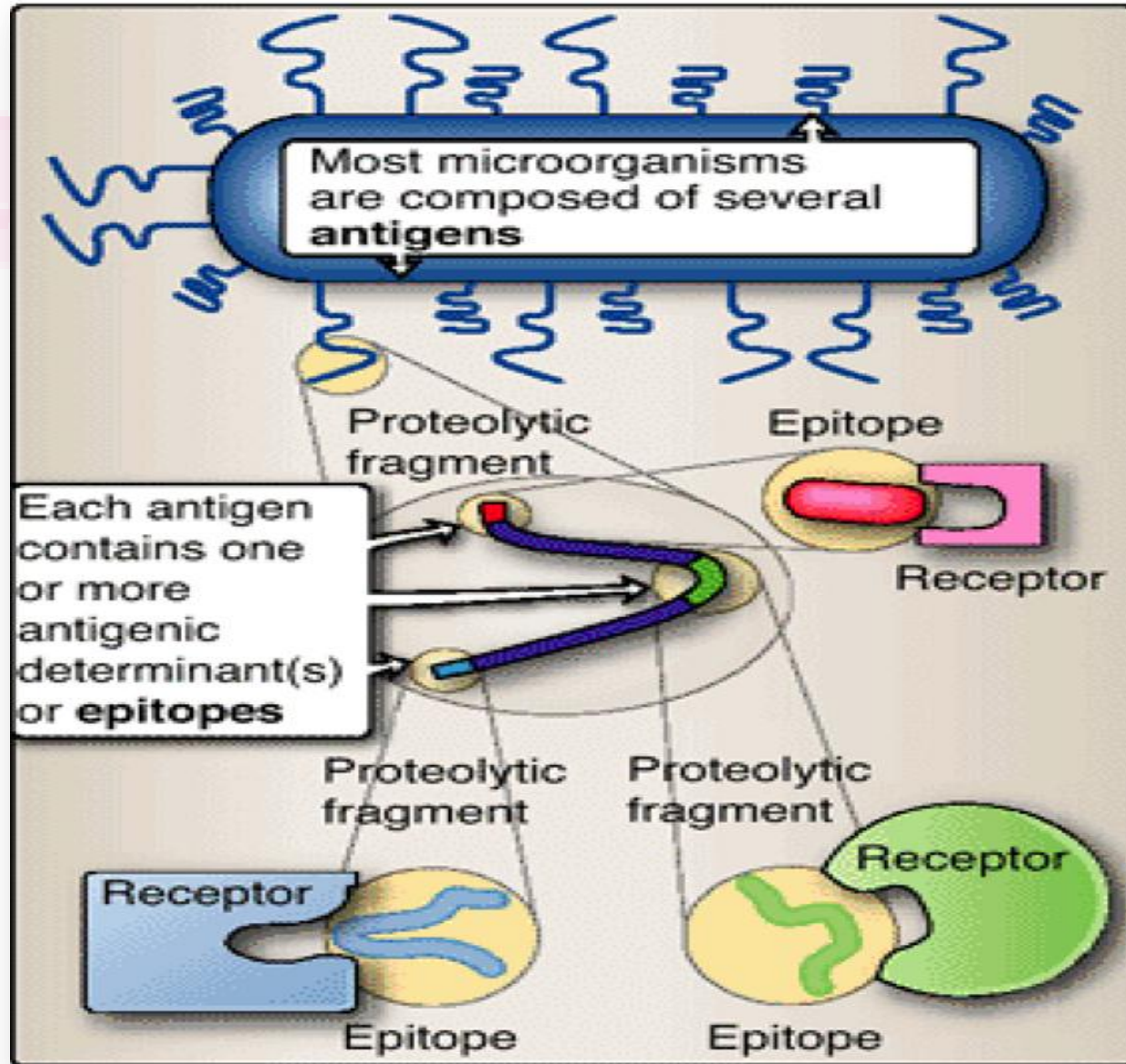
NH₂-variable region (V)

COOH-constant region (C)



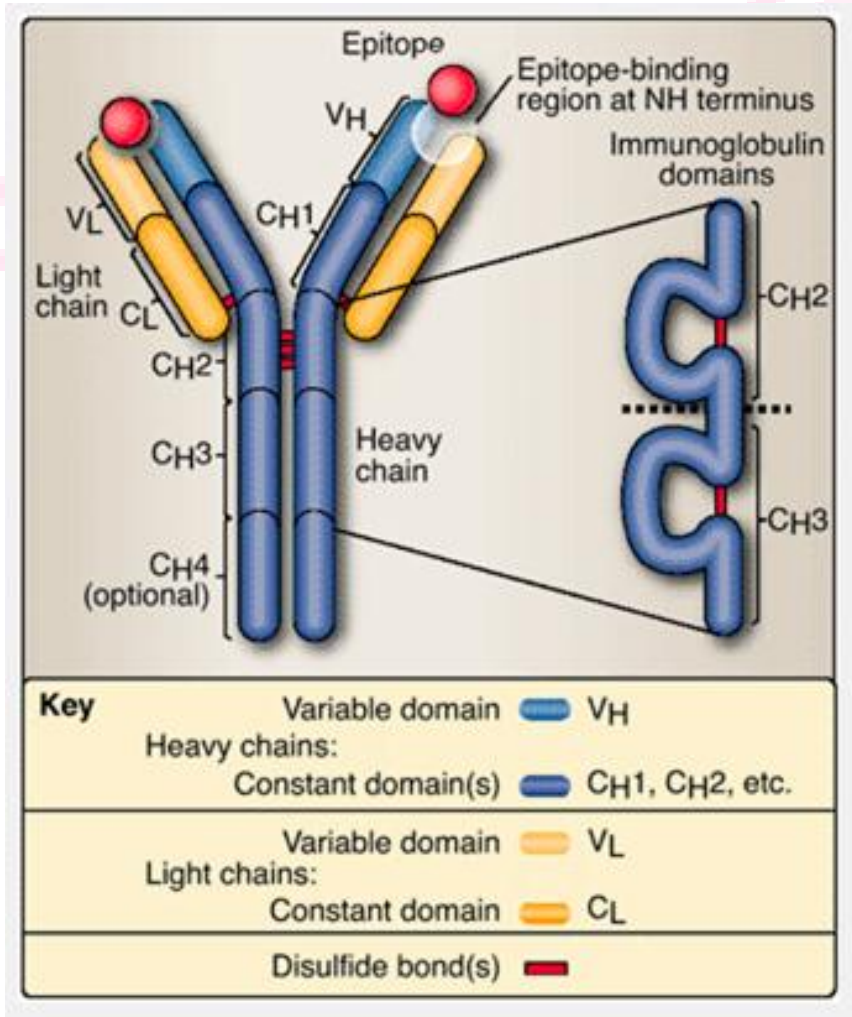
Epitopes (Antigenic or immunogenic determinants)

➤ ตำแหน่งย่อยบนโมเลกุลของแอนติเจนที่กระตุ้น immune response



8-22 aa

Basic structure of Ig



➤ B cell epitopes $> 10^{19}$

➤ 1 gene \rightarrow 1 transcript \rightarrow 1 protein

Ig $> 10^{19}$ genes

➤ Human genome : 20,000-25,000 genes

How can the immune system generate enough different Ig with different antigen specificities??

(Figure from Doan T et al., Lippincott's Illustrated Reviews: Immunology 1st Edition;2007;p57)

Hypotheses

1. Germ-line theory

: The genome contributed by the germ cells, egg and sperm, contains a large repertoire of immunoglobulin genes.

2. Somatic mutation theory

: The genome contains a relatively small number of immunoglobulin genes, from which a large number of antibody specificities are generated in the somatic cells by mutation or recombination.

3. Two-gene model

: in 1965 proposed by Dreyer & Bennett

Two-gene model

In 1965, *Dreyer and Bennett* proposed that two or more genes could control the production of one polypeptide chain.

{opposed the one gene, one polypeptide theory}

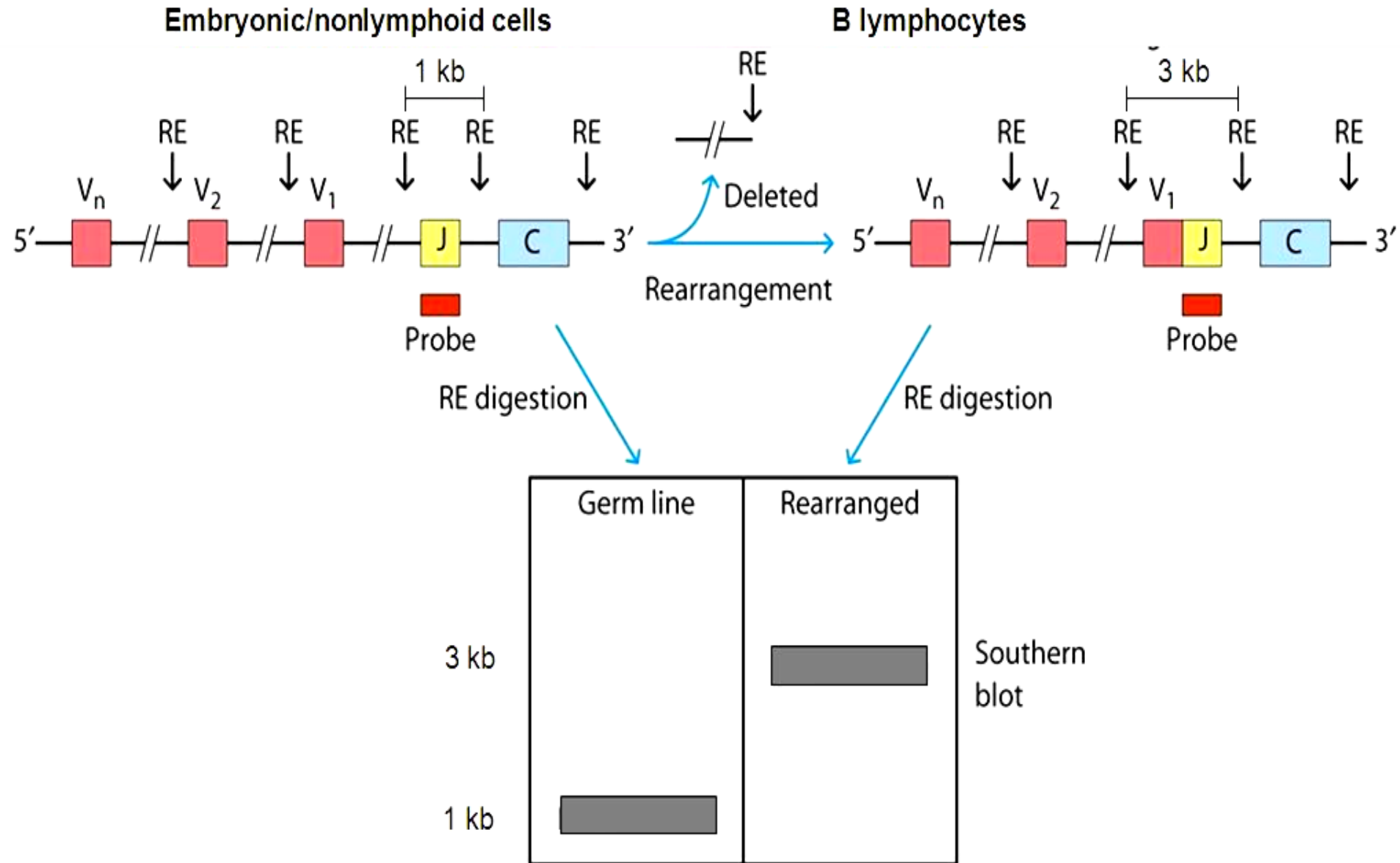


- A single C region gene encoded in the GERMLINE and separate from the V region genes
- Multiple choices of V region genes available
- A mechanism to rearrange V and C genes in the genome so that they can fuse to form a complete Immunoglobulin gene

(Dryer, W.J., and Bennett, J., Proc. Natl. Acad. Sci. USA 54, 864-869; 1965; Nandini Shetty; Immunology Introductory Textbook 2nd Ed; 2005; p33)

Proof of the Dreyer - Bennett hypothesis by Susumu Tonegawa in 1976

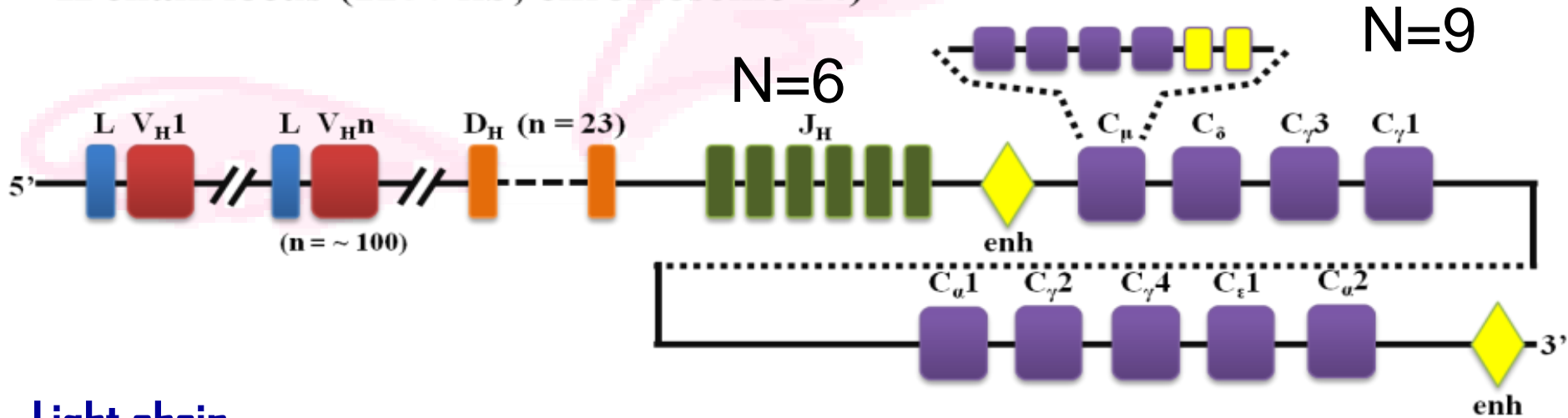
(In 1987, Tonegawa was awarded the Nobel prize)



Germline organization of human Ig genes

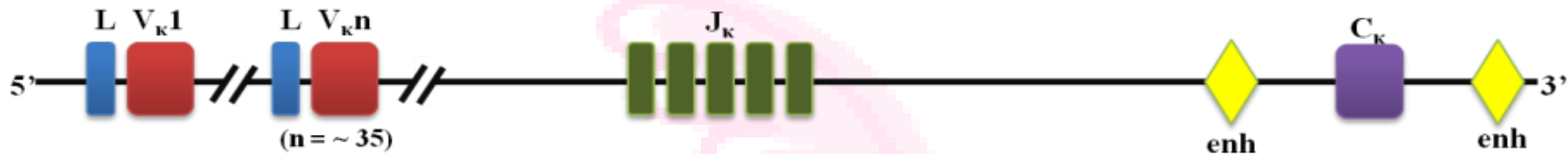
- Heavy chain

H chain locus (1250 kb; chromosome 14)

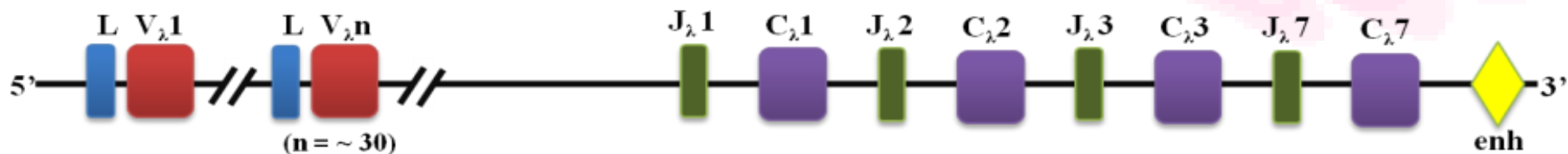


- Light chain

κ chain locus (1820 kb; chromosome 2)



λ chain locus (1050 kb; chromosome 22)



Germline DNA

Rearranged DNA

Primary RNA transcript

Messenger RNA (mRNA)

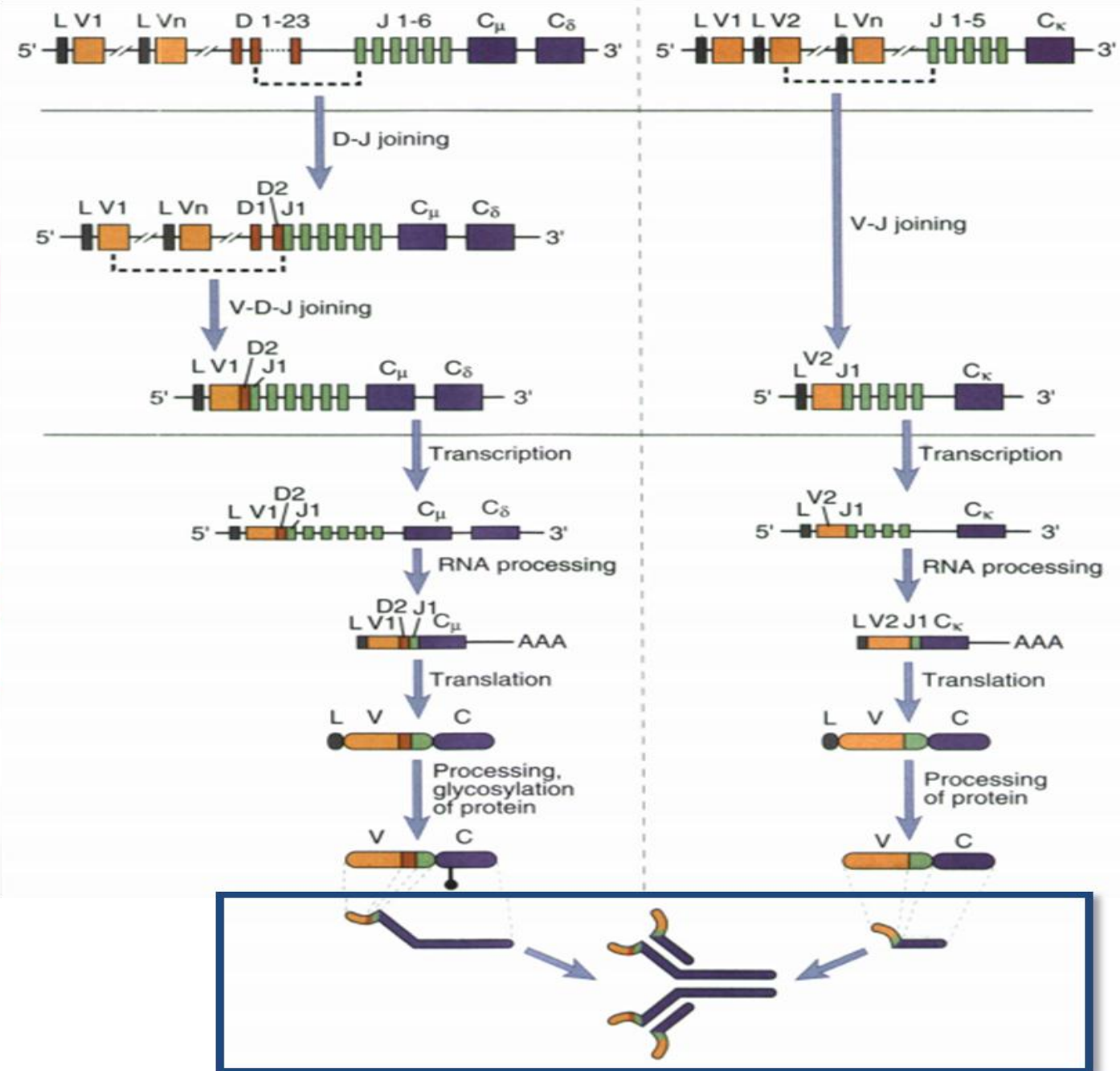
Nascent polypeptide

Mature polypeptide

Assembled Ig molecule

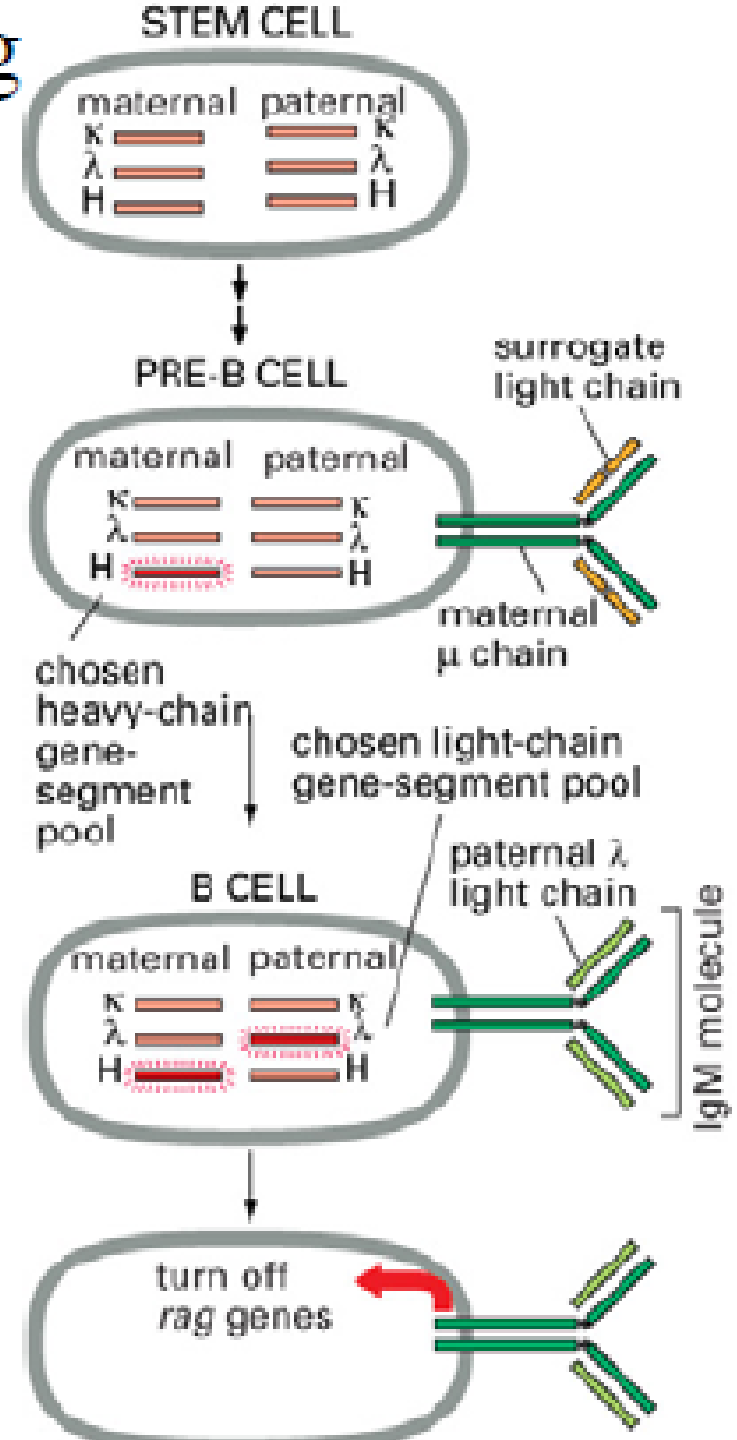
(A) μ Heavy chain

(B) κ Light chain



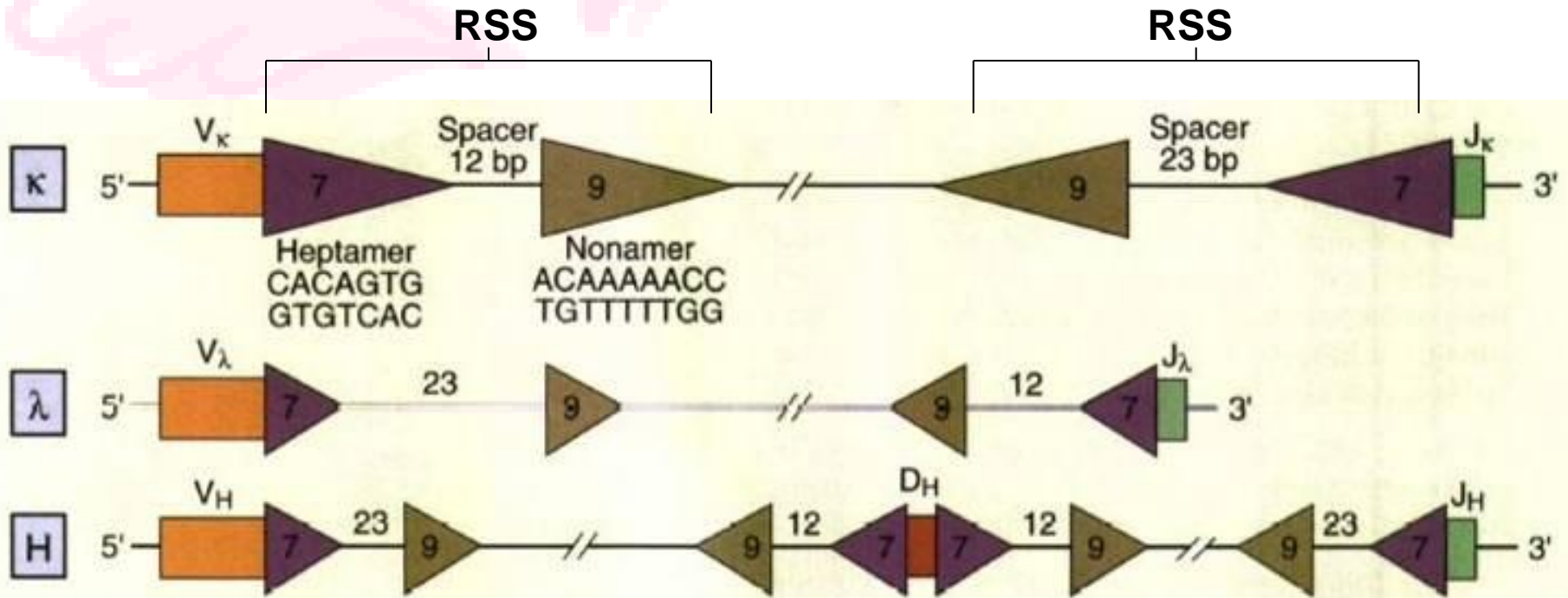
Allelic exclusion ensuring specificity

- Each B cell clone produces only antibody against one antigen. The dimeric antibody doesn't carry two different antigen recognition sites on two different monomers
 - Ensuring cross-linking of antigens for aggregation and elimination
 - Ensuring no "co-stimulation" of non-specific antigen binding due to the stimulation of the specific antigen binding on the other monomer
- However, there are two sets of chromosomes in each B cells
 - Will use only one each of the light chain and heavy chain genes



Mechanisms of V (D) J recombination

Orientation of V (D) J gene segments and recombination signal sequences (RSS)



Process of V (D) J recombination

(Abbas AK; 6th ed, p164)

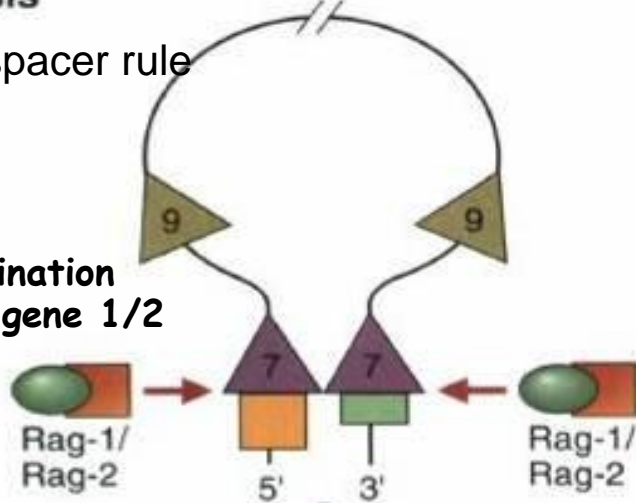
Unrearranged locus



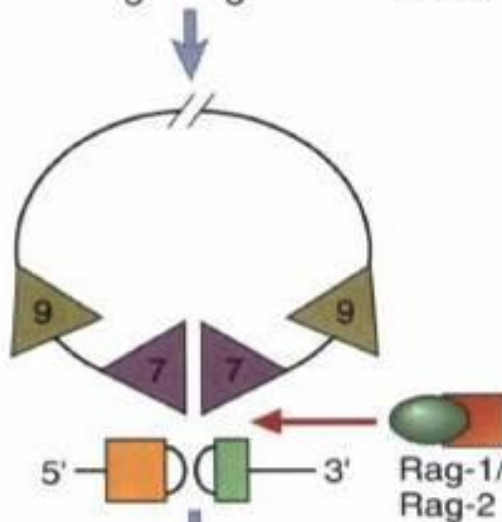
① Synapsis

12/23 bp spacer rule

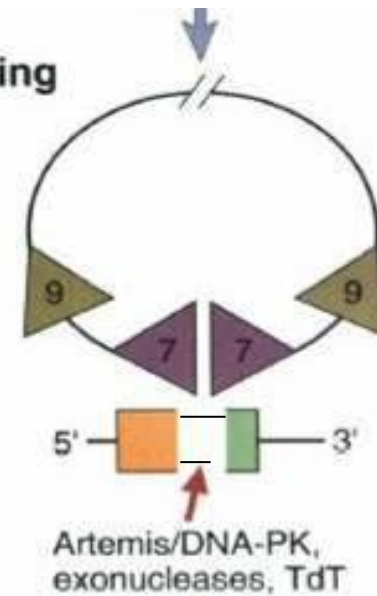
Recombination activating gene 1/2



② Cleavage

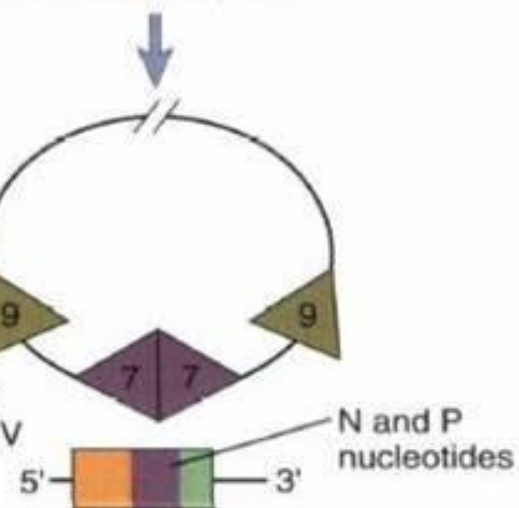


③ Hairpin opening and end-processing

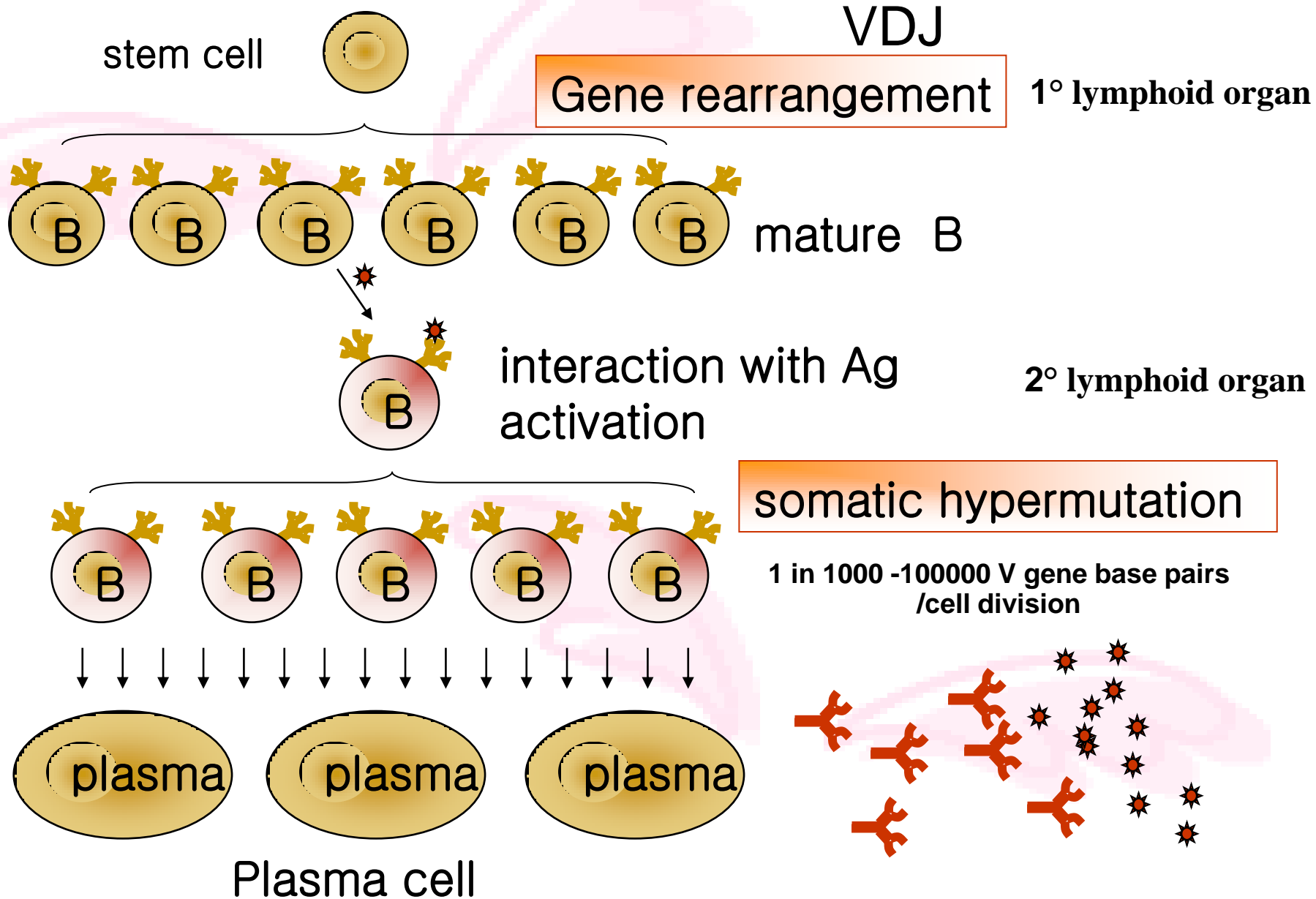


④ Repair/ligation

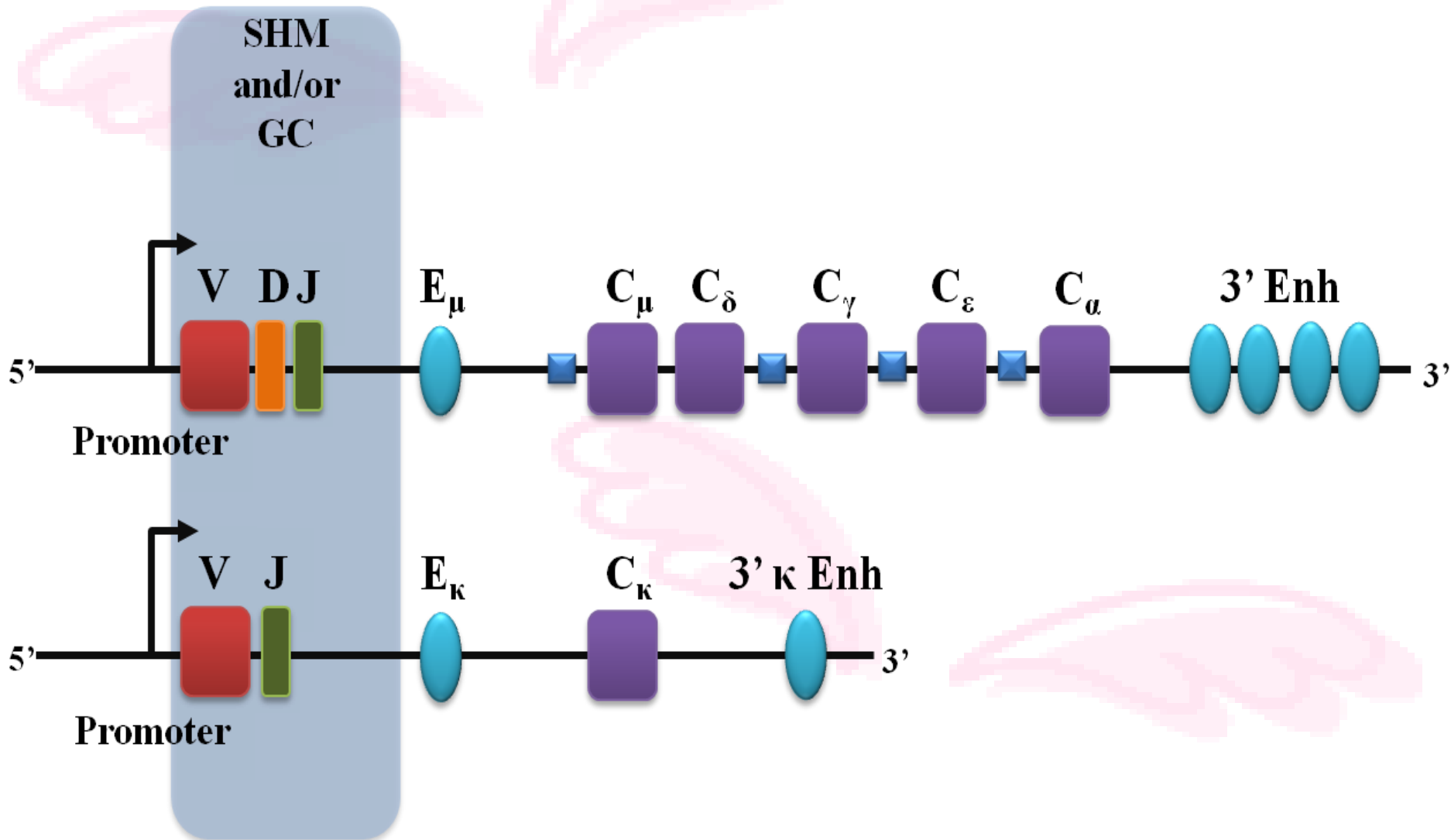
Ku70/Ku80/DNA-PK
XRCC4/DNA LigaseIV



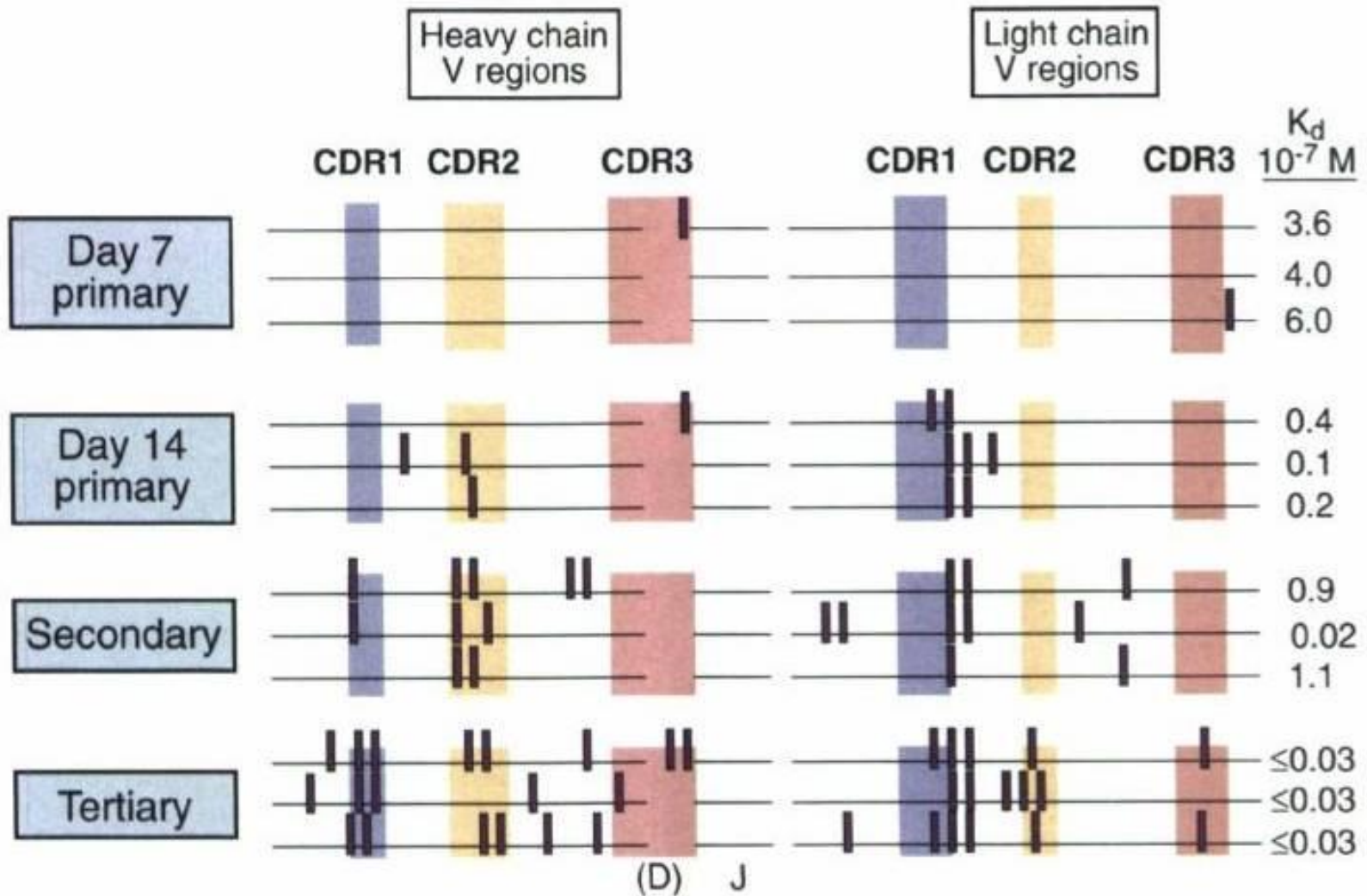
Somatic hypermutation



บริเวณการเกิด somatic hypermutation ในยีนของ Ig heavy-chain และ light-chain



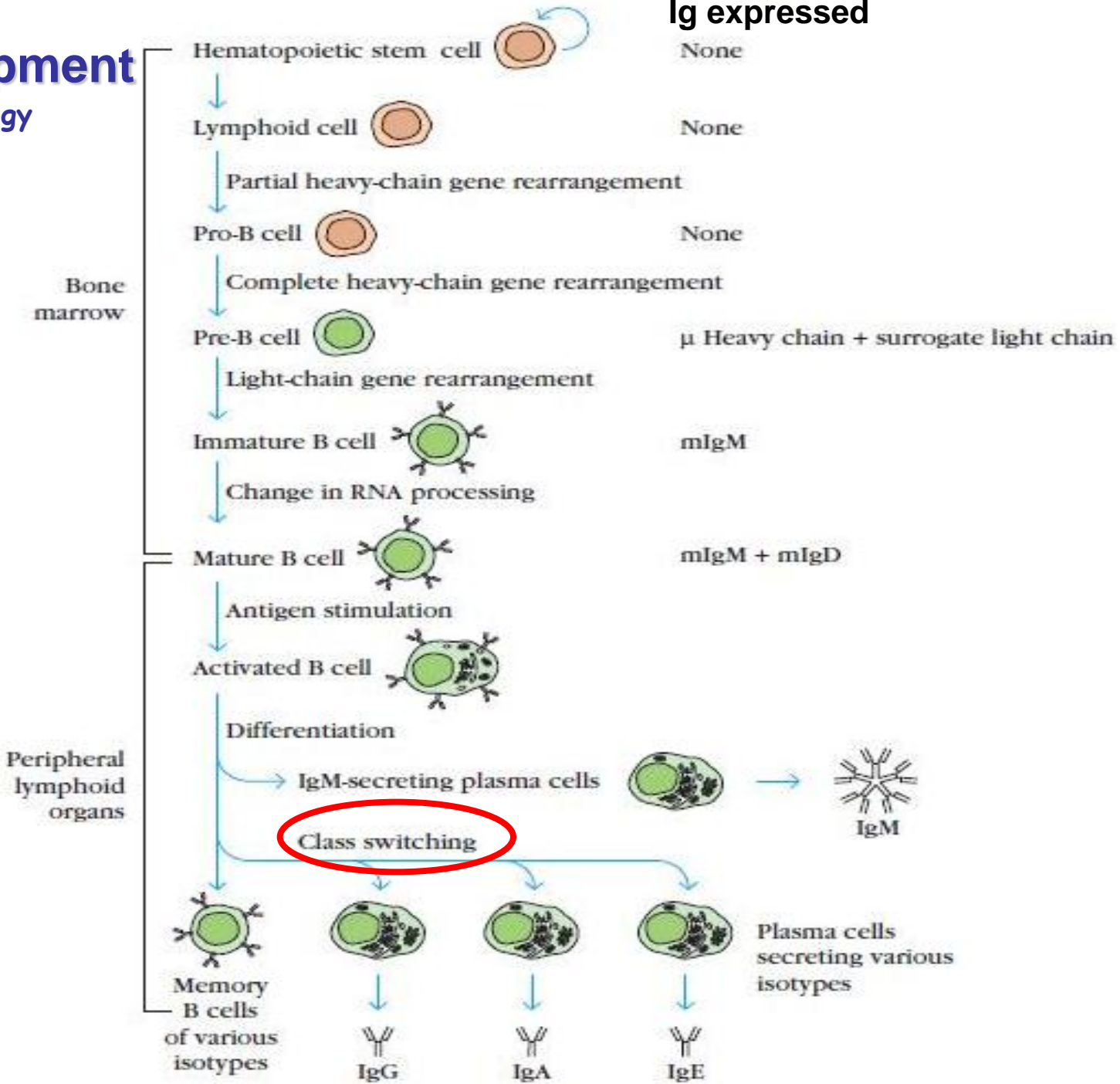
Evidence for somatic mutation in variable regions of Ig genes



Total potential repertoire $>10^{12}$

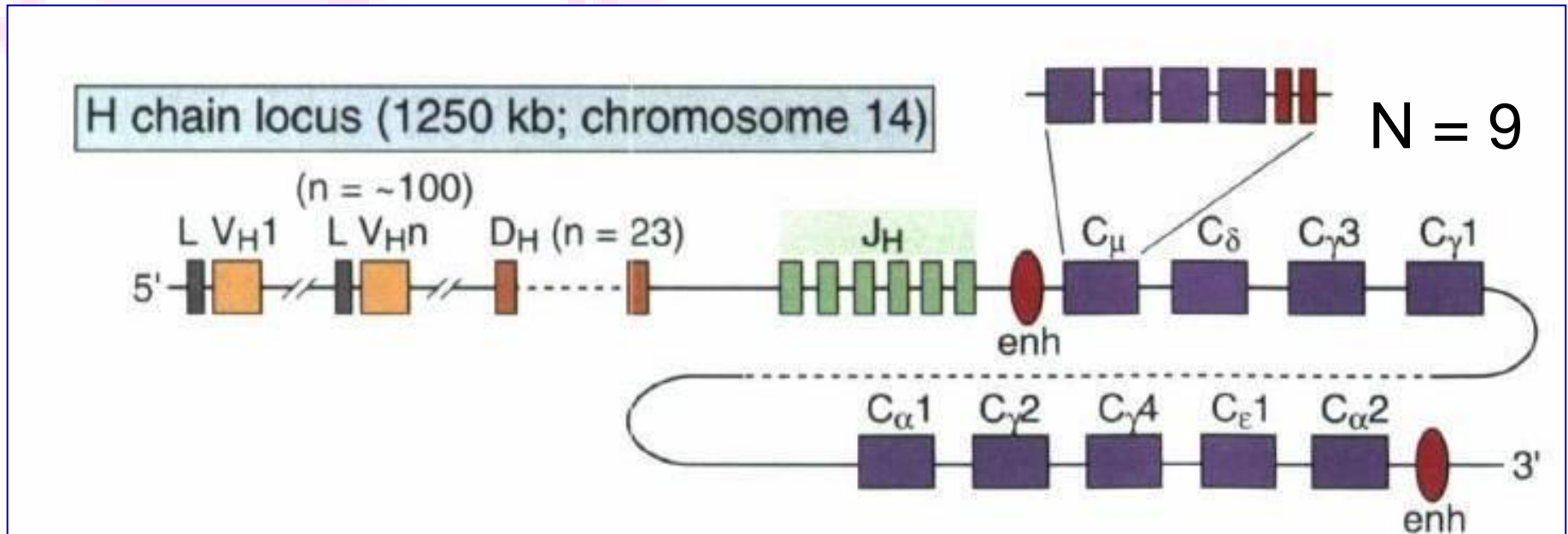
B-cell development

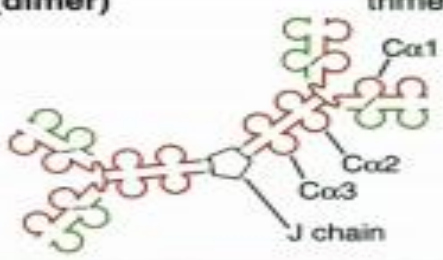
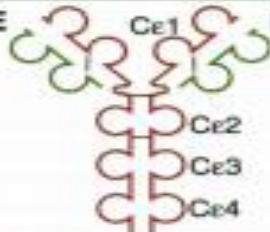
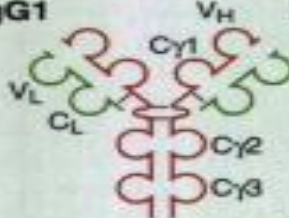
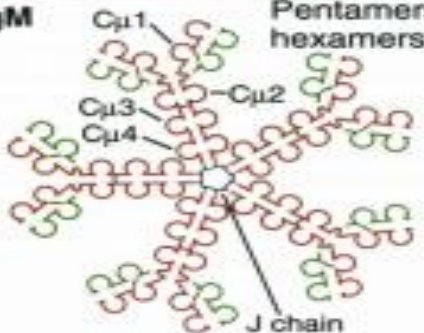
(Kuby J. Immunology
5th ed; p106)



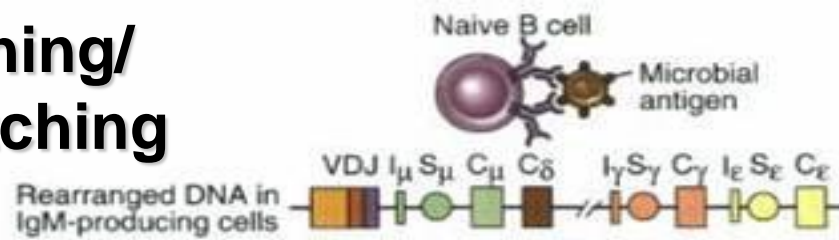
Germline organization of human Ig genes

Heavy chain



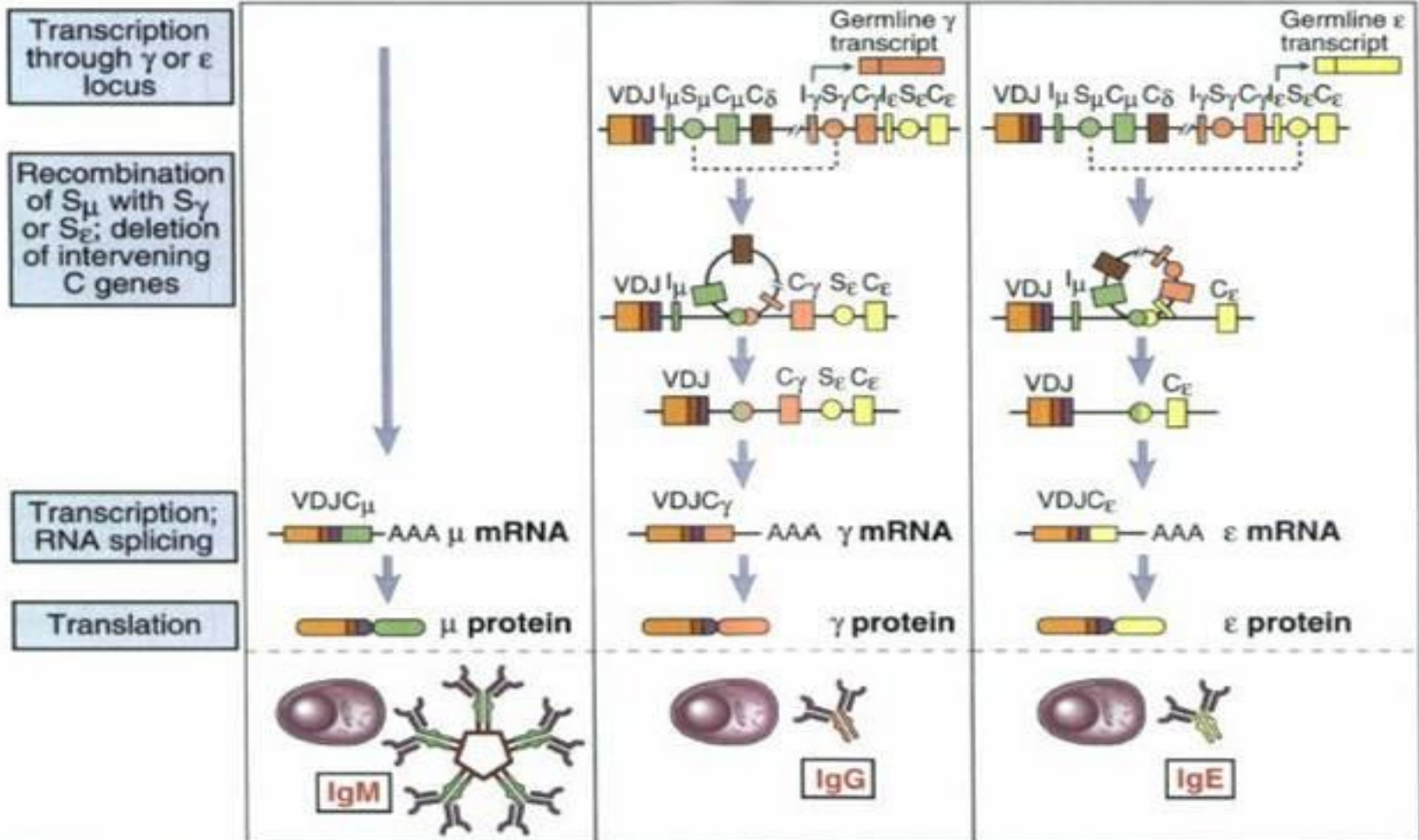
Isotype of antibody	Subtypes	H chain	Serum concentr. (mg/mL)	Serum half-life (days)	Secreted form	Functions
IgA	IgA1,2	α (1 or 2)	3.5	6	IgA (dimer) Monomer, dimer, trimer 	Mucosal immunity
IgD	None	δ	Trace	3	None	Naive B cell antigen receptor
IgE	None	ϵ	0.05	2	IgE Monomer 	Defense against helminthic parasites, immediate hypersensitivity
IgG	IgG1-4	γ (1,2,3 or 4)	13.5	23	IgG1 Monomer 	Opsonization, complement activation, antibody-dependent cell-mediated cytotoxicity, neonatal immunity, feedback inhibition of B cells
IgM	None	μ	1.5	5	IgM Pentamers, hexamers 	Naive B cell antigen receptor, complement activation

Class switching/ Isotype switching



No signals from helper T cells

Signals from helper T cells (CD40 ligand, cytokines)



สรุปปัจจัยที่ทำให้เกิดความหลากหลายของแอนติบอดี

1. Multiple germ-line gene segments

	Heavy chain	κ -light chain	λ -light chain
	Estimated number of segments		
V	100	35	30
D	23	0	0
J	6	5	4

2. Combinational V-(D)-J joining

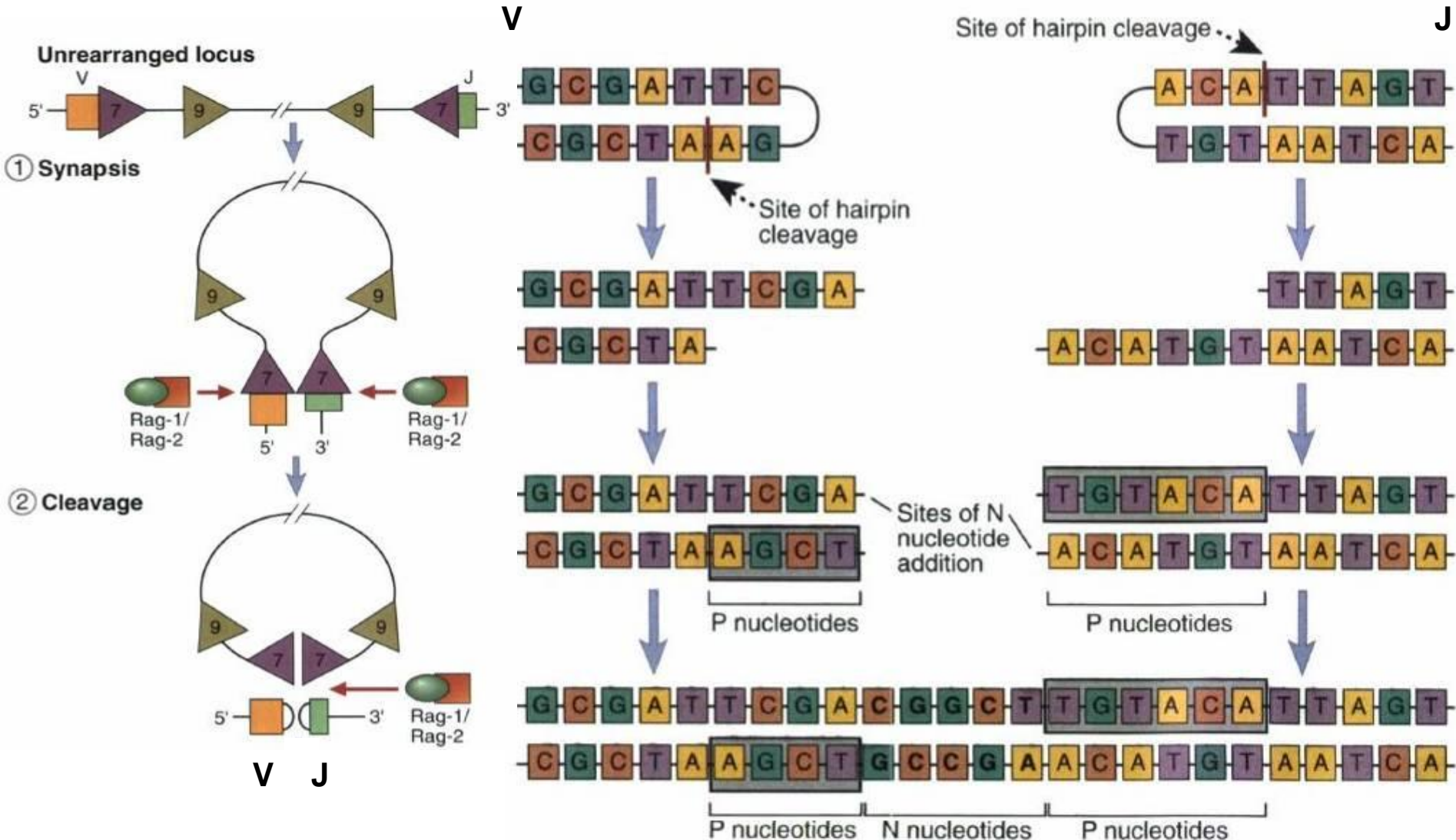
$$100 \times 23 \times 6 = 13800 \quad 35 \times 5 = 175 \quad 30 \times 4 = 120$$

$$[175 + 120 = 295 \text{ light chains}]$$

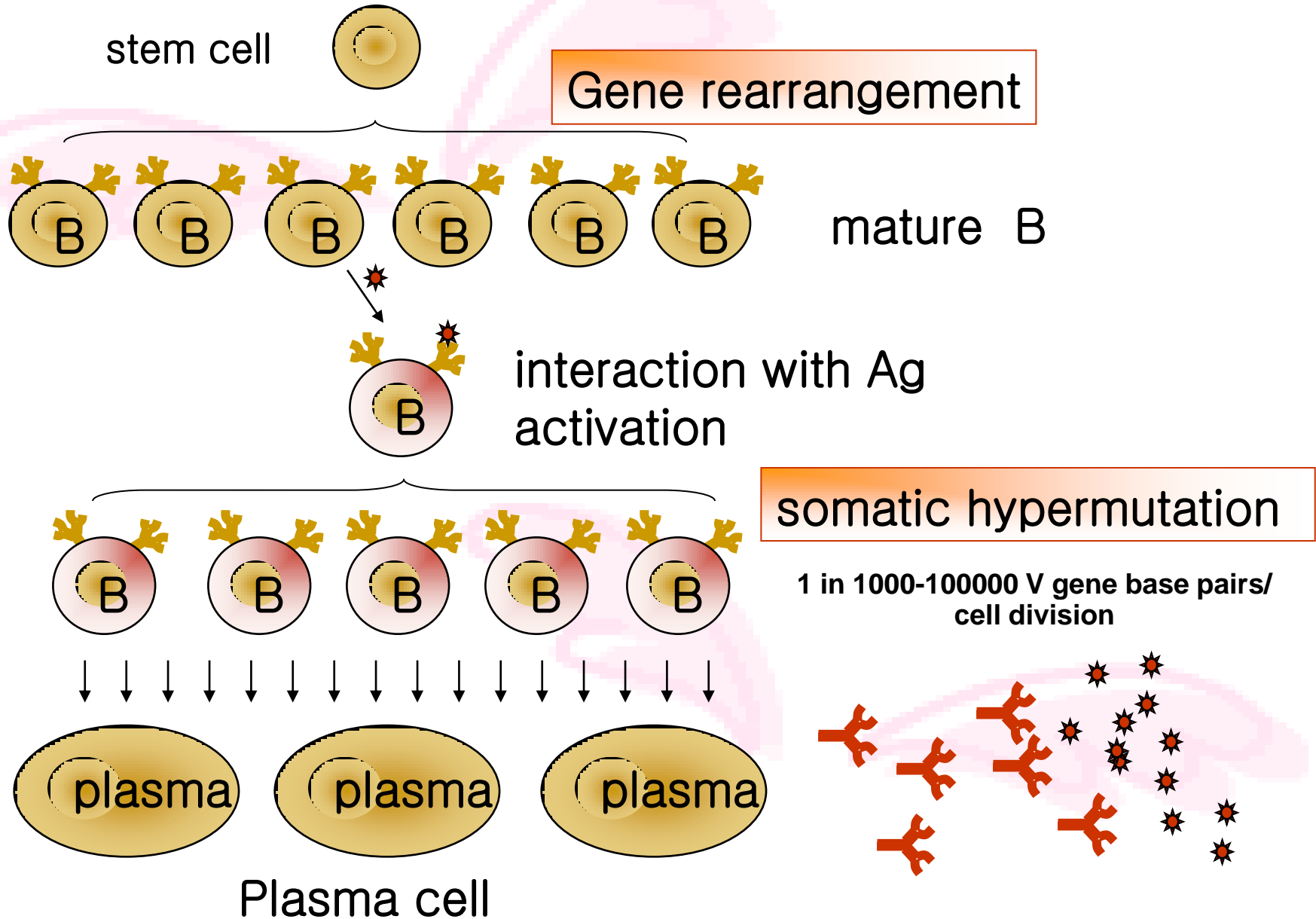
3. Combinational association of heavy and light chains

$$13800 \times 295 = 4.07 \times 10^6$$

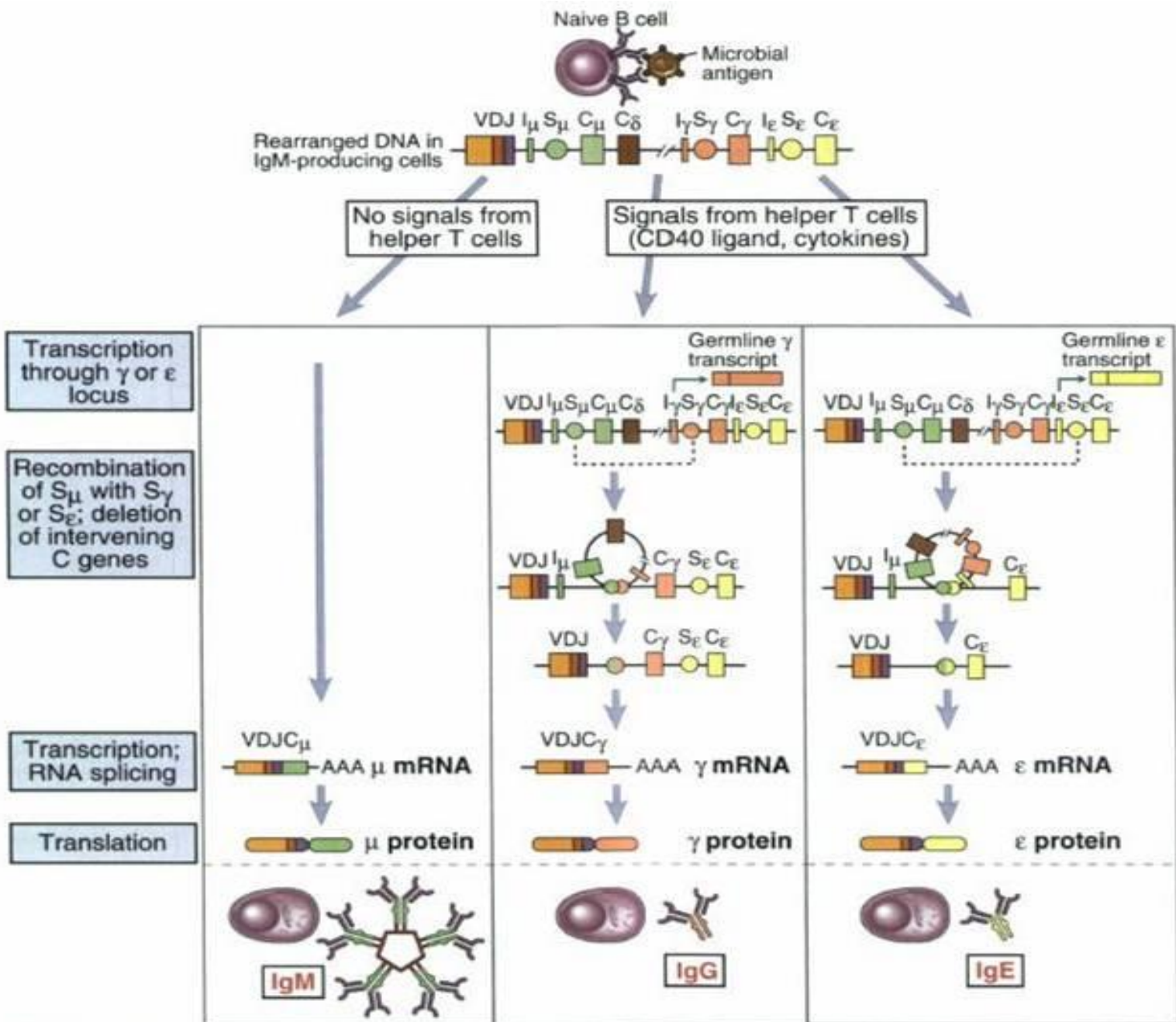
4. Junctional diversity



5. Somatic hypermutation



6. Class switch recombination (CSR)



C region is also important!!

Heavy chain

