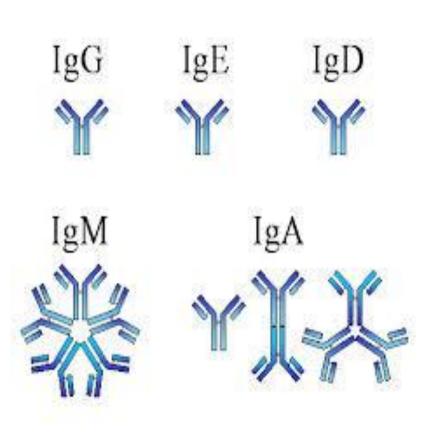
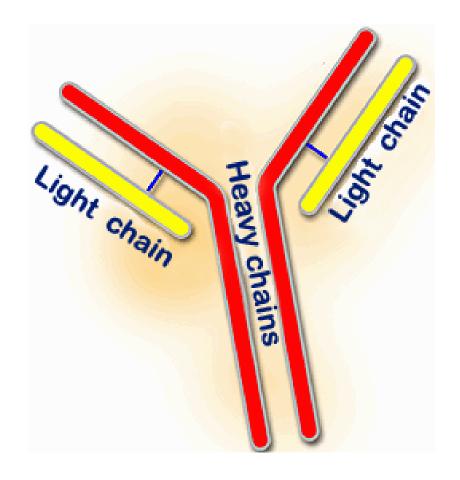


IMMUNOGLOBULINS

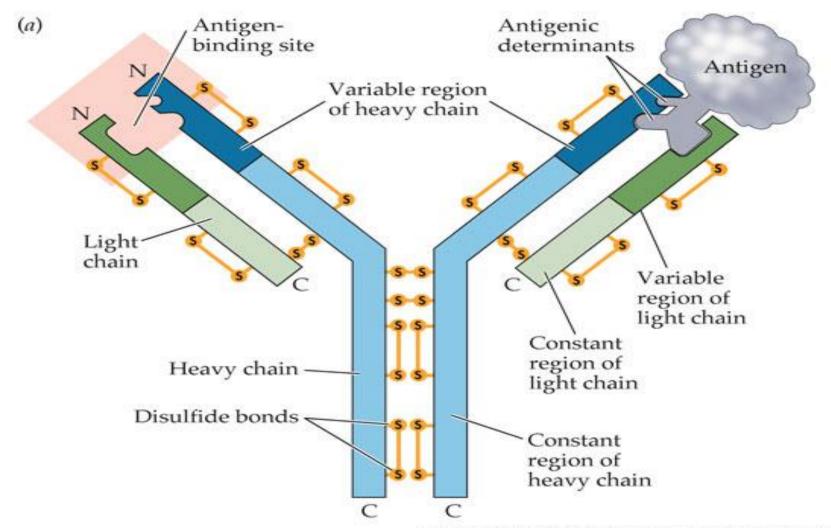
- are glycoproteins found in the serum portion of the blood
- Composed of 82% 96% polypeptide and 2% 14% carbohydrate
- Humoral branch of the immune response
- Primary role is antigen recognition and in biological activities related to immune response and complement activation
- Has five major classes (IgG, IgM, IgA, IgD, IgE)

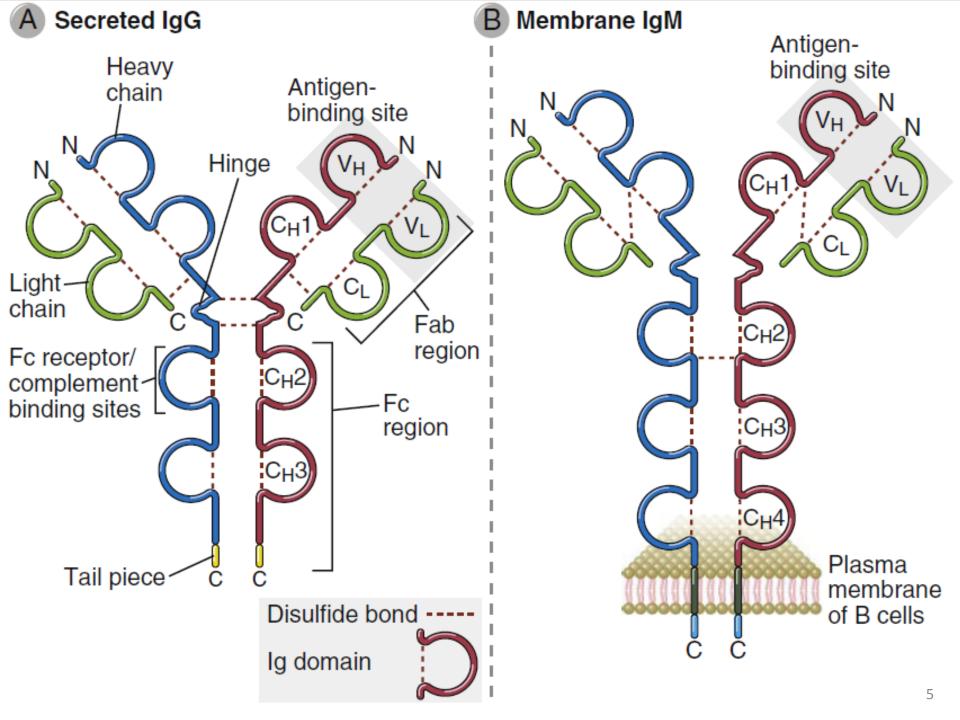
TETRAPEPTIDE STRUCTURE



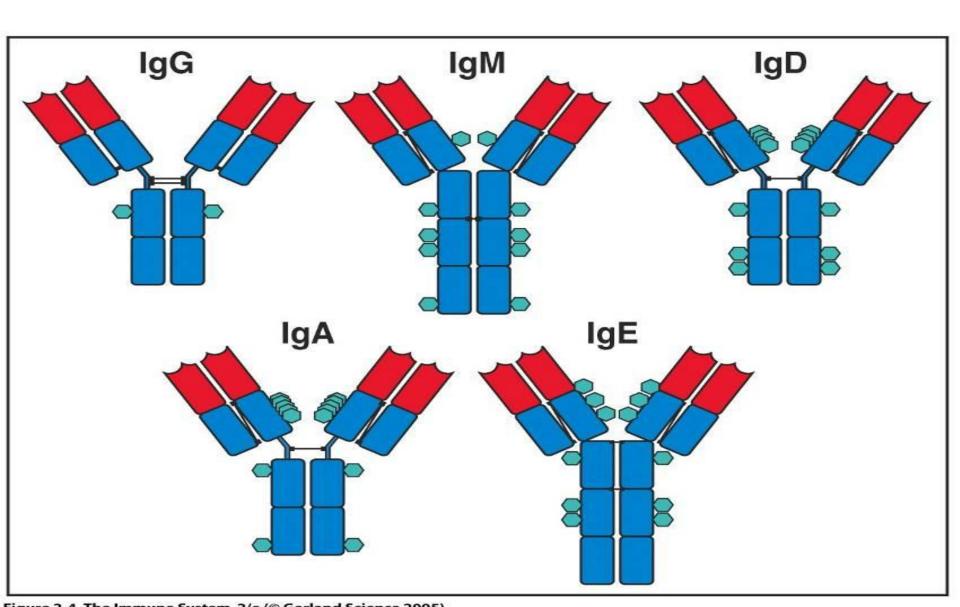


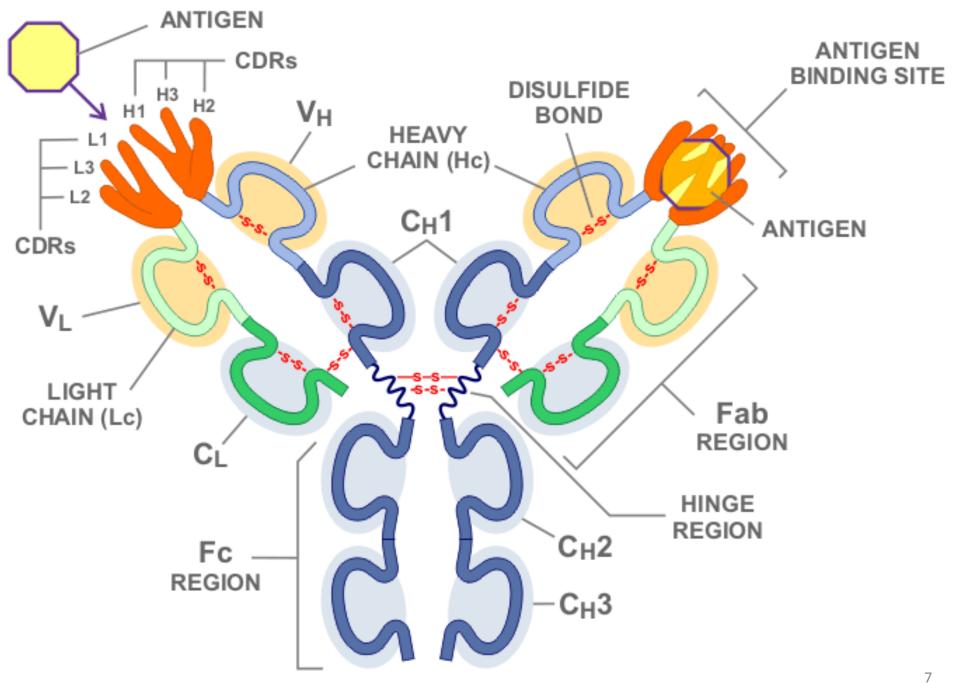
Structural configuration of Antibody





Different classes of Antibodies



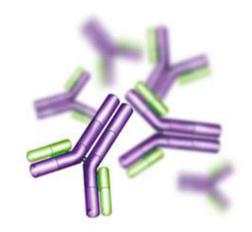


NATURE OF LIGHT CHAINS

- Bence-Jones Proteins
 - Kappa (€)
 - Lambda (λ)
 - Constant region
 - is the C-terminal end and contains similar amino acids for each class of antibody.
 - Variable region
 - includes 110-130 amino acids of the light and heavy chains, and is responsible for binding to antigen. This part of the antibody shows variations in amino acids when the specificity of the antibody for antigen is changed.

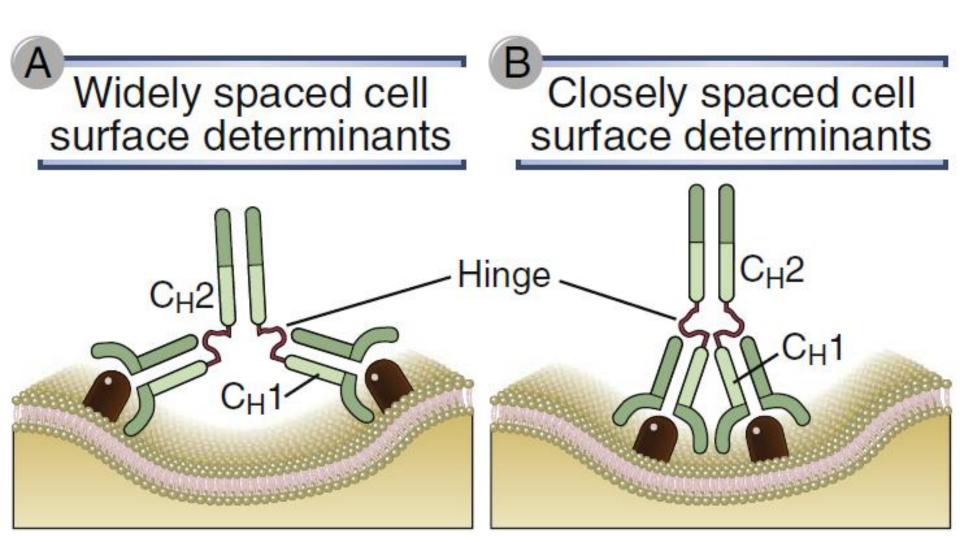
Heavy chain sequencing

- Variable domain first 110 amino acids at the aminoterminal domain
- The remaining amino acid can be divided up into three or more constant regions with very similar sequences, $C_H 1, C_H 2, C_H 3$.
- > IgG Gamma (γ) heavy chains
- ➤ IgM Mu (µ) heavy chains
- \triangleright IgA Alpha (α) heavy chains
- \triangleright IgD Delta (δ) heavy chains
- \triangleright IgE Epsilon (ϵ) heavy chains



HINGE REGION

- flexible amino acid stretch in the central part of the heavy chains of the IgG and IgA immunoglobulin classes, which links these 2 chains by disulfide bonds
 - √ High content of proline and hydrophobic residues
 - √ Flexibility assists initiation of the complement cascade
- rich in cysteine and proline amino acids, extremely variable in amino acid sequence,
- no resemblance to any other immunoglobulin region
- Gamma, delta and alpha chains have hinge region
- Mu and epsilon chains do not have hinge region



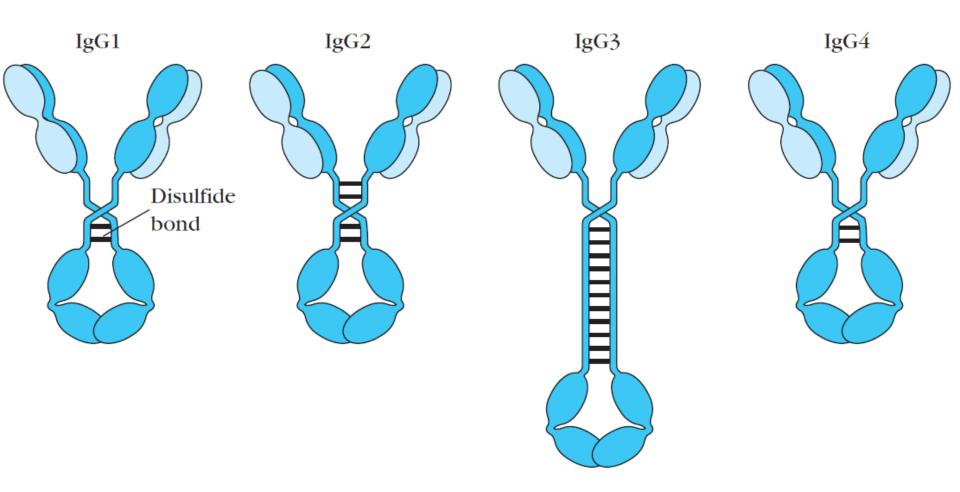
CARBOHYDRATE FUNCTIONS

- \triangleright All types of immunoglobulins contain a carbohydrate portion located between C_H 2 domains of the two H chains
 - They function as:
 - 1. Increases the solubility of the immunoglobulin
 - 2. Provides protection against degradation
 - 3. enhances functional activity of the F_C domains

- Structure:
 - ✓ All IgG's are monomers. The subclasses differ in the number of disulfide bonds and length of the hinge region.
- Properties
 - ✓ most versatile immunoglobulin because it is capable of carrying out all of the functions of immunoglobulin molecules.
 - a) IgG is the major Ig in serum 75% of serum Ig is IgG

SUBCLASSES

- a) IgG1 Gamma 1 heavy chains ► 67%
- b) IgG2 Gamma 2 heavy chains > 22%
- c) IgG3 Gamma 3 heavy chains
 - > 7%
 - Largest hinge region
- d) IgG4 Gamma 4 heavy chains > 4%

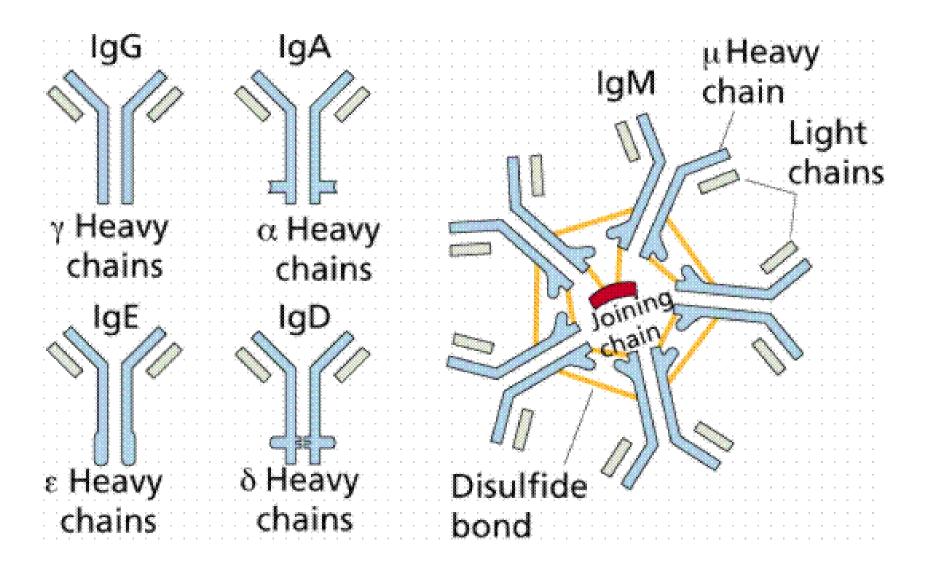


- b) Placental transfer IgG is the only class of Ig that crosses the placenta. Not all subclasses cross equally well; IgG2 does not cross well.
- c) Fixes complement Not all subclasses fix equally well; IgG4 does not fix complement
- d) The term opsonin is used to describe substances that enhance phagocytosis. IgG is a good opsonin.

> IgM

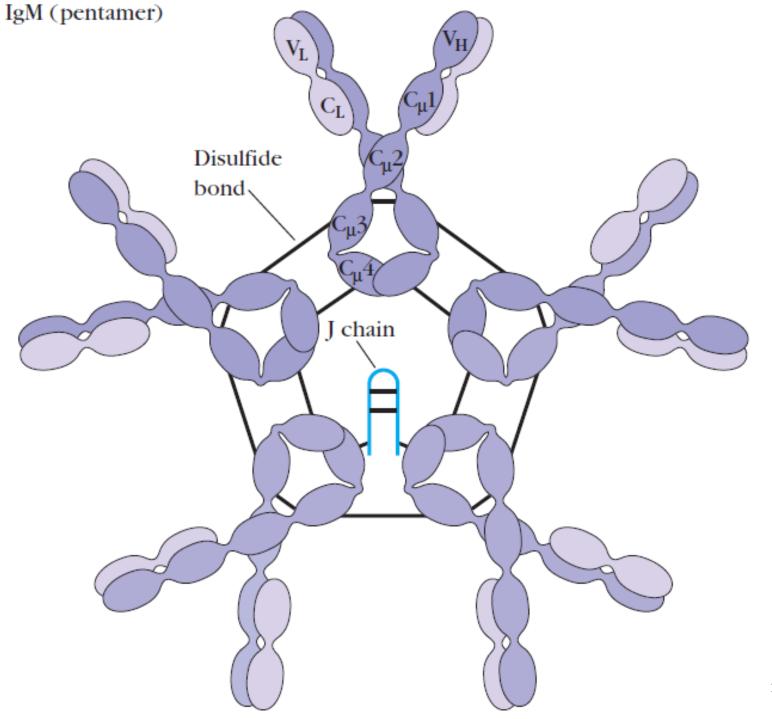
Structure:

- ✓ IgM normally exists as a pentamer but it can also exist as a monomer.
- ✓ IgM has an extra domain on the mu chain (C_{H4}) and it has another protein covalently bound via a S-S bond called the Joining chain or J chain. This chain functions in polymerization of the molecule into a pentamer.
- √ macroglobulin



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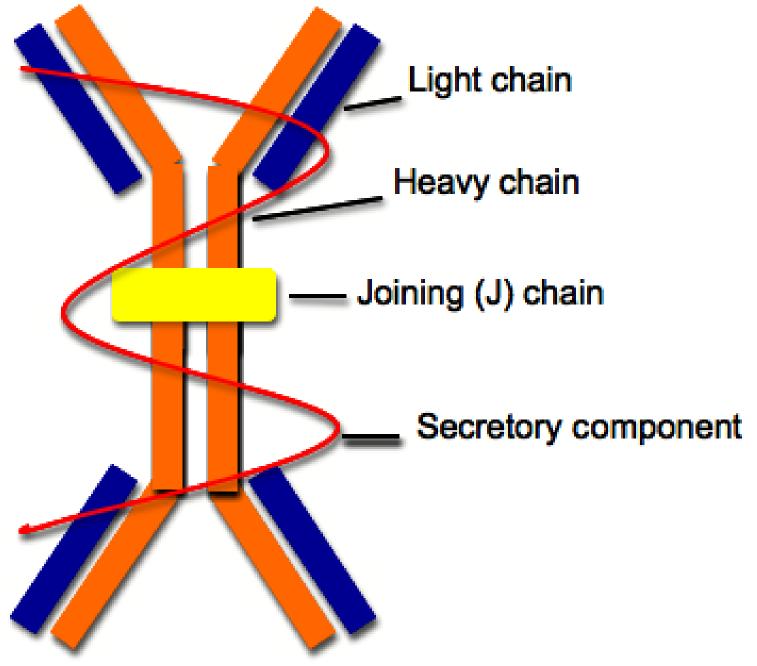
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Properties:

- √ a) IgM is the third most common serum Ig.
- ✓ b) IgM is the first Ig to be made by the fetus and the first Ig to be made by a virgin B cells when it is stimulated by antigen.
- √ c) As a consequence of its pentameric structure, IgM is a good complement fixing Ig.
- √ d) IgM is also a good agglutinating Ig .

- **>** IgA
 - Structure:
 - ✓ Dimer
 - ✓ When IgA is found in secretions is also has another protein associated with it called the secretory piece
 - ✓ The secretory piece helps IgA to be transported across mucosa and also protects it from degradation in the secretions.



Properties:

- a) IgA is the 2nd most common serum Ig.
- b) IgA is the major class of Ig in secretions tears, saliva, colostrum, mucus. Since it is found in secretions secretory IgA is important in local (mucosal) immunity.
- c) Normally IgA does not fix complement, unless aggregated.

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- ≽IgD
- Structure:
- ✓ Monomer
- ✓ Percentage serum antibodies: 0.2%
- √ Half-life in serum: 3 days
- ✓ Complement Fixation: No
- ✓ Placental Transfer: No
- ✓ Known Functions: In serum function is unknown. On B cell surface, initiate immune response.

- **≻**IgE
- Structure:
- ✓ Monomer
- ✓ Percentage serum antibodies: 0.002%
- ✓ Half-life in serum: 2 days
- ✓ Complement Fixation: No
- ✓ Placental Transfer: No
- ✓ Known Functions: Allergic reactions. Possibly lysis of worms.

TABLE 3-2 Chain composition of the five immunoglobulin classes

Class*	Heavy chain	Number of C _H Ig domains	Subclasses	Light chain	J chain	Molecular formula
IgG	γ	3	γ1, γ2, γ3, γ4 (human) γ1, γ2a, γ2b, γ3 (mouse)	κorλ	None	$\gamma_2 \kappa_2 \ \gamma_2 \lambda_2$
IgM	μ	4	None	κorλ	Yes	$(\mu_2 \kappa_2)_n$ $(\mu_2 \lambda_2)_n$ $n = 1 \text{ or } 5$
IgA	α	3	α1, α2	κorλ	Yes	$(\alpha_2 \kappa_2)_n$ $(\alpha_2 \lambda_2)_n$ $n = 1, 2, 3, \text{ or } 4$
lgE	€	4	None	κorλ	None	$\epsilon_2 \kappa_2$ $\epsilon_2 \kappa_2$
IgD	δ	3	None	κorλ	None	$\begin{array}{l} \delta_2\kappa_2 \\ \delta_2\lambda_2 \end{array}$

^{*}See Figure 3-22 for general structures of the five antibody classes.

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