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Antibodies as immunochemical tools

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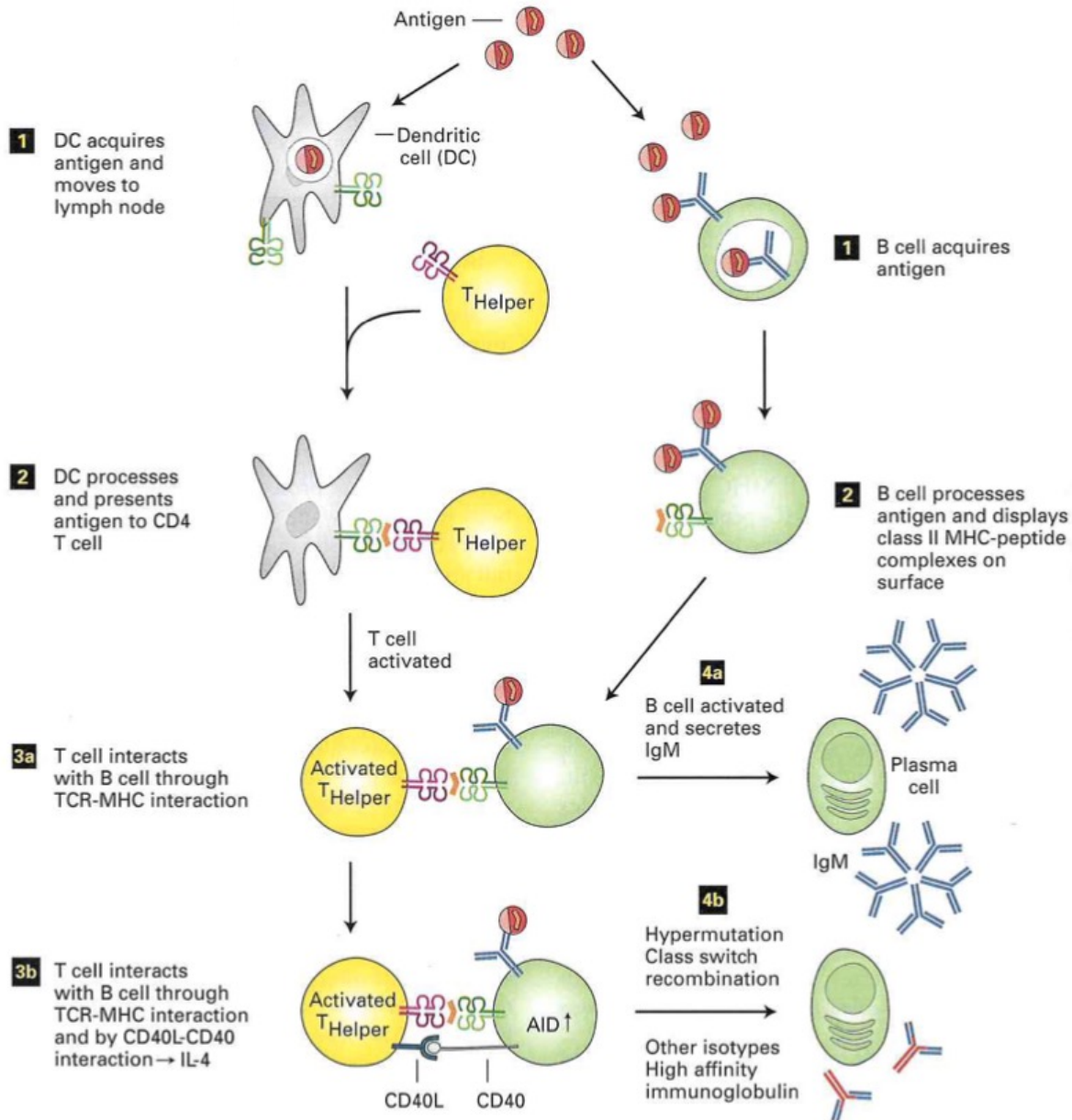
Summary of interplay between T_H and B cells

Antigen

T cell epitope
(binds to MHC,
recognized by TCR)

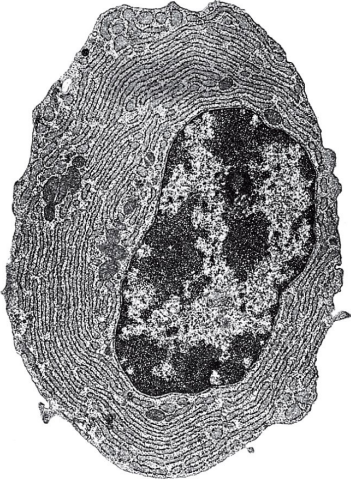


B cell epitope
(binds to BCR)



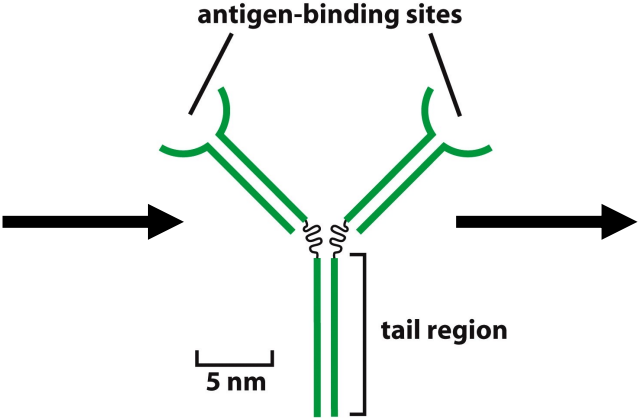
Antibodies as immunochemical tools

Immunology

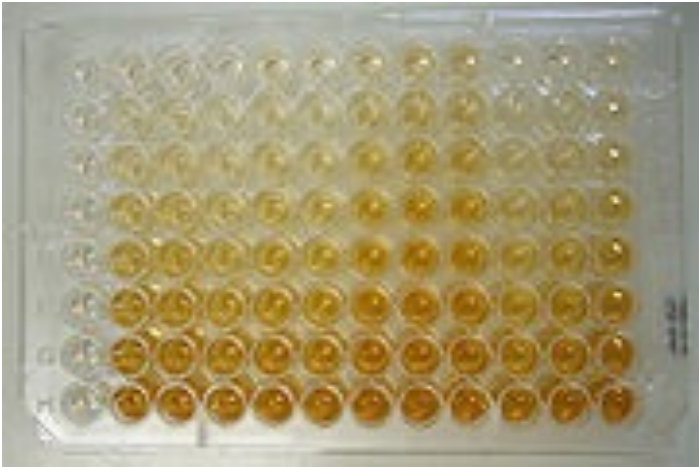


effector B cell (plasma cell) 1 μm

The “tools“:
antibodies



Immunoassay

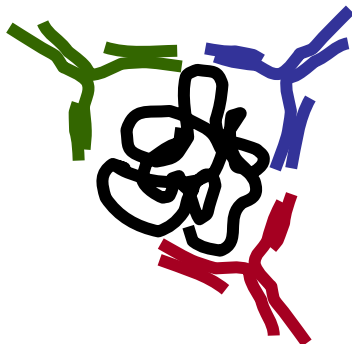


Polyclonal vs. monoclonal antibodies

polyclonal

Antibodies that are collected from sera of exposed animal

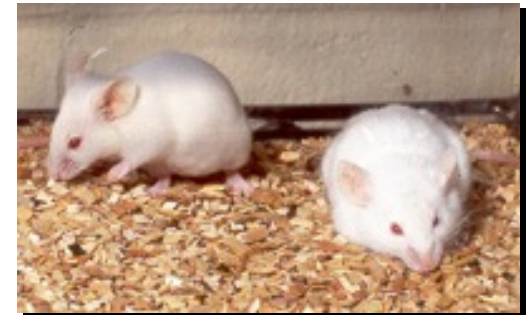
recognize multiple antigenic sites of injected substance



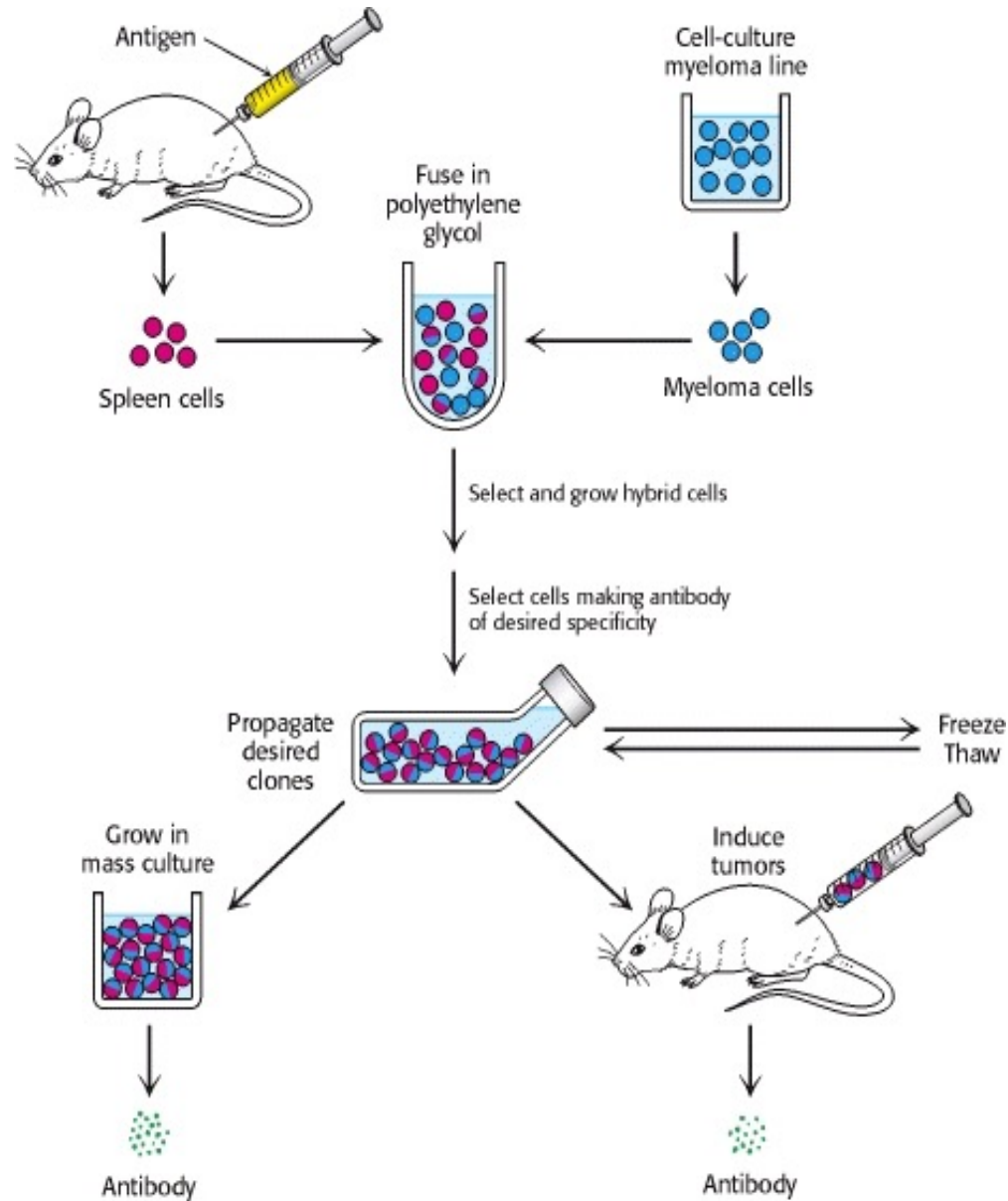
monoclonal

Individual B cell hybridoma is cloned and cultured.
Secreted antibodies are collected from culture media

recognize ONE antigenic site of injected substance



Generation of monoclonal antibodies



Antibodies: Definitions

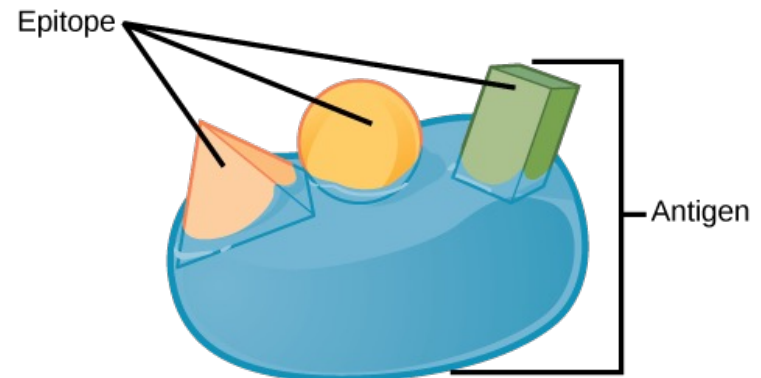
Antibodies, or immunoglobulins (Igs), are γ -globulin proteins folded into well defined three-dimensional structures synthesized by living organisms, e.g. mice, rabbits or goats, or by living cells, in response to the presence of a foreign substance known as the antigen.

Immunogen: Molecule that is capable of eliciting an immune response by the immune system of an organism.

Antigen: Molecule that is able to bind to the product of that immune response: the antibody.

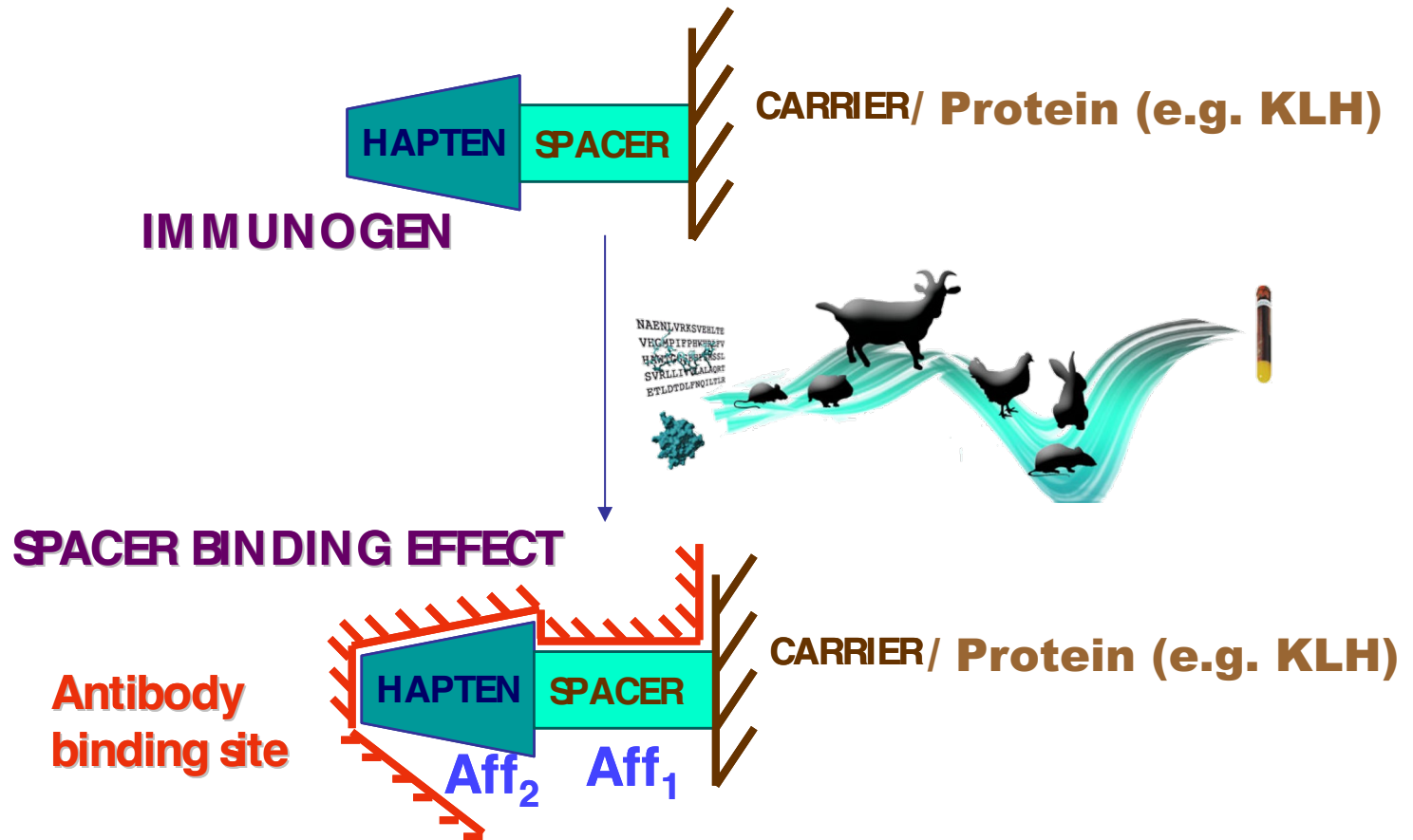
Epitope: An epitope is a specific location on the surface of an antigen that has a particular molecular structure and that is recognized by a particular antibody or a set of specific antibodies that the epitope elicits during the immune response.

Hapten: Small molecules (< 5000 Dalton) that need to be conjugated to a carrier protein (e.g. bovine serum albumin (BSA), keyhole limpet hemocyanin (KLH) or ovalbumin) to elicit the immune response.



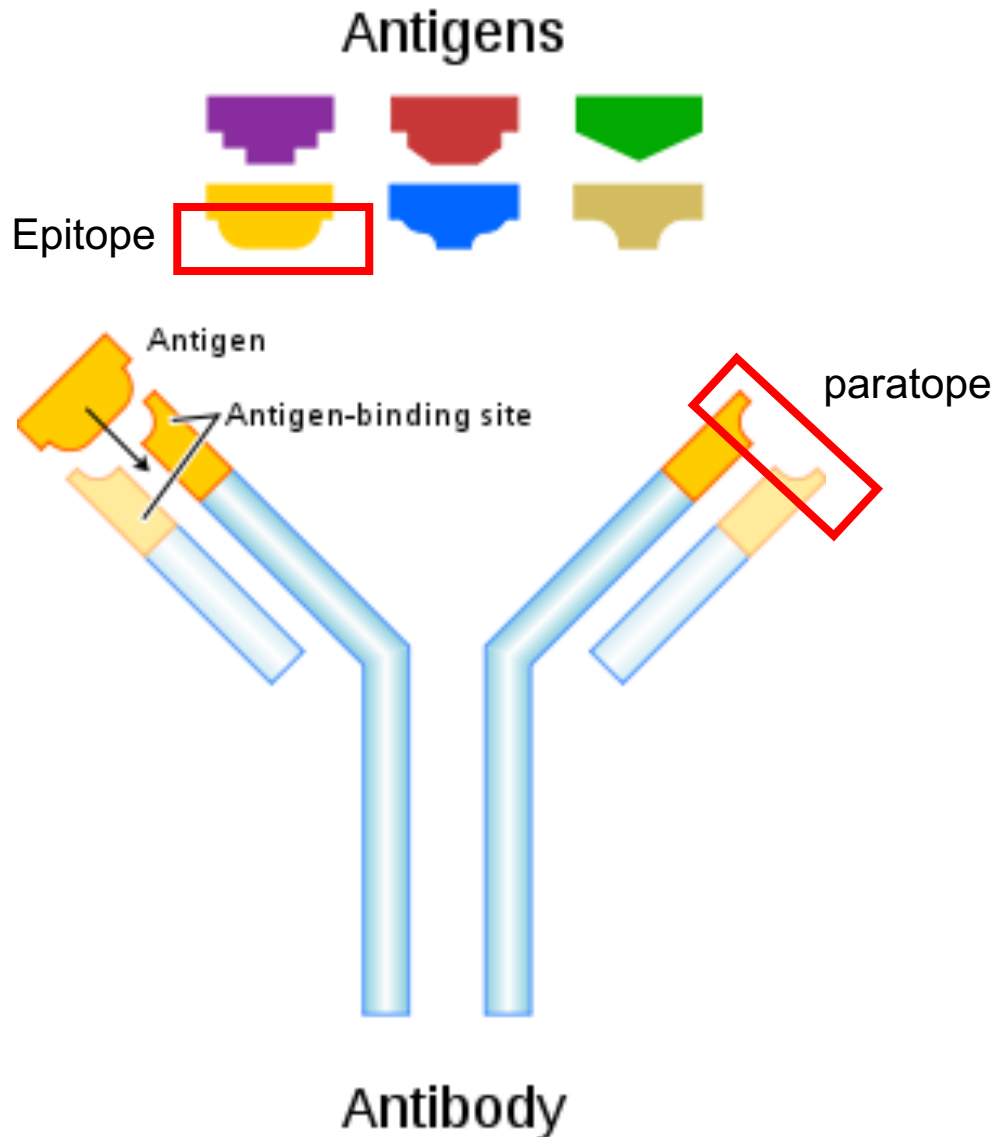
Antibodies: Definitions

Immunogens are always antigens but not all antigens are immunogens



Antibodies as immunochemical reagents

=> Antibodies are used as bioanalytical reagents to specifically detect and quantify other molecules

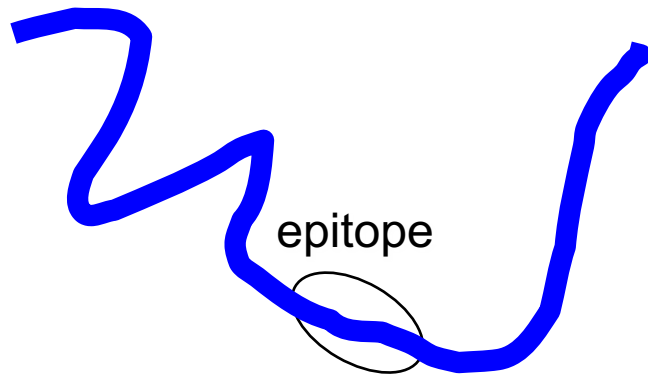


Continuous vs. discontinuous epitopes

Continuous epitope:

short peptide or denatured protein structure, epitope consist of **sequential** amino acids

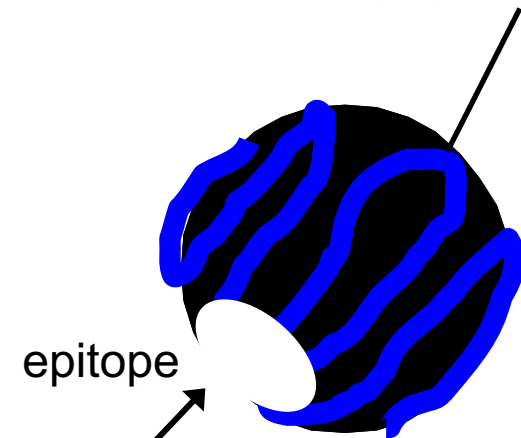
peptide chain



Discontinuous epitope:

present only in 3-dimensional protein structure, epitope comprises **non-sequential** amino acids

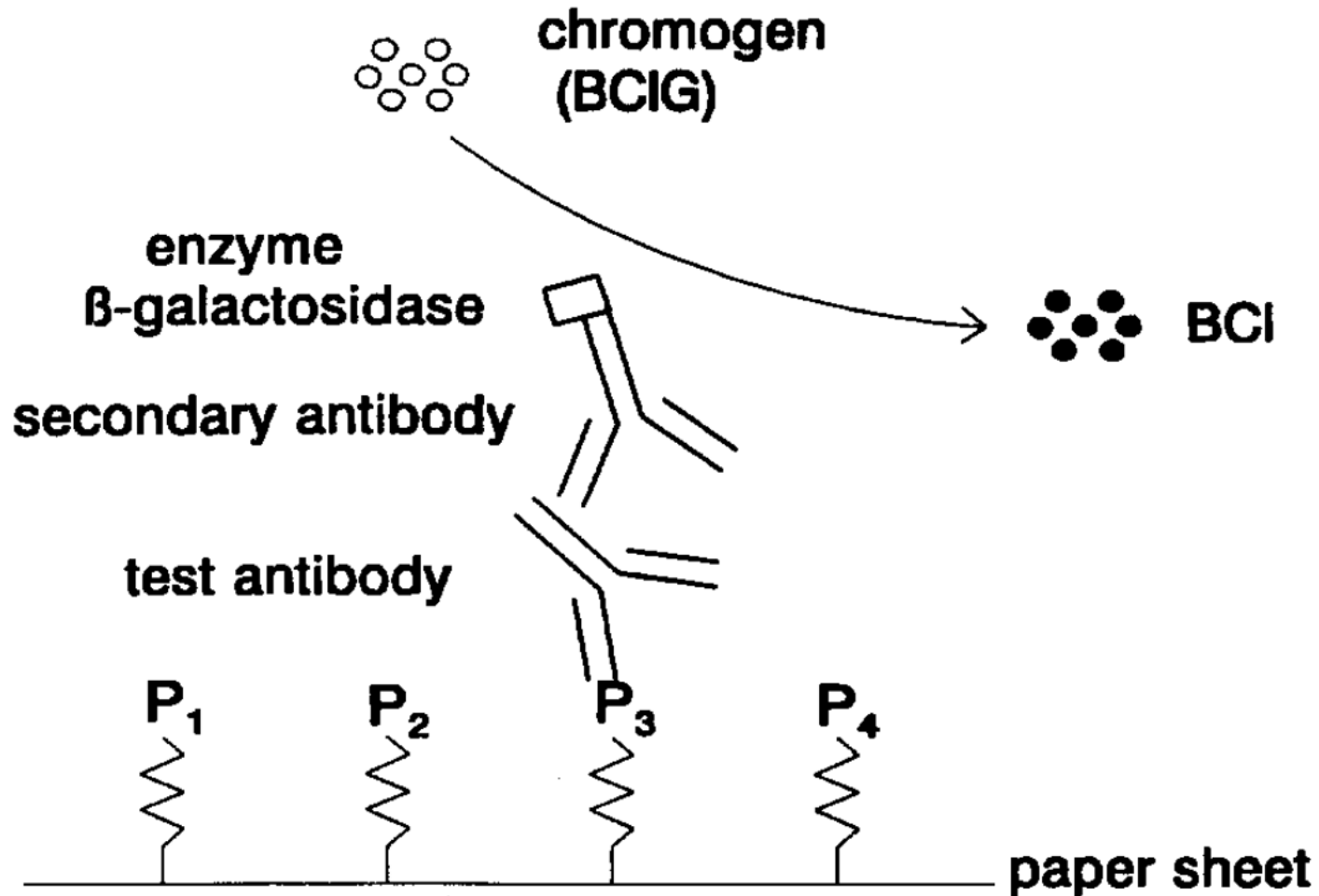
peptide chain



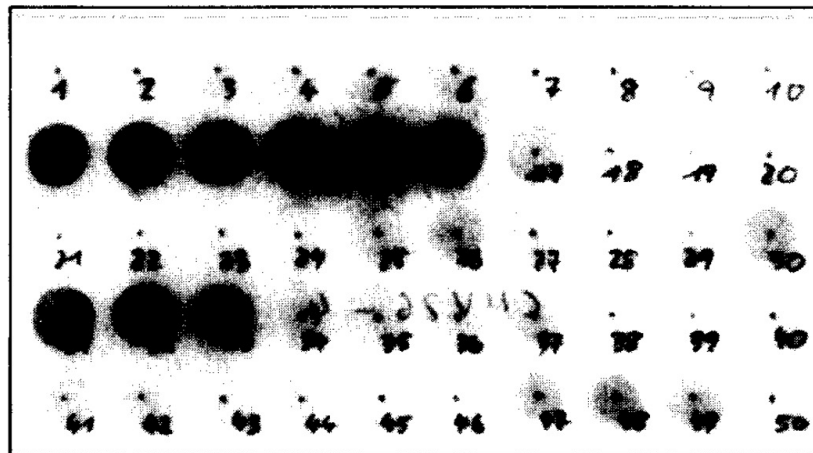
antibody binding/recognition site

Excursion: Epitope mapping

How do we know to what epitope an antibody binds?



Epitope mapping



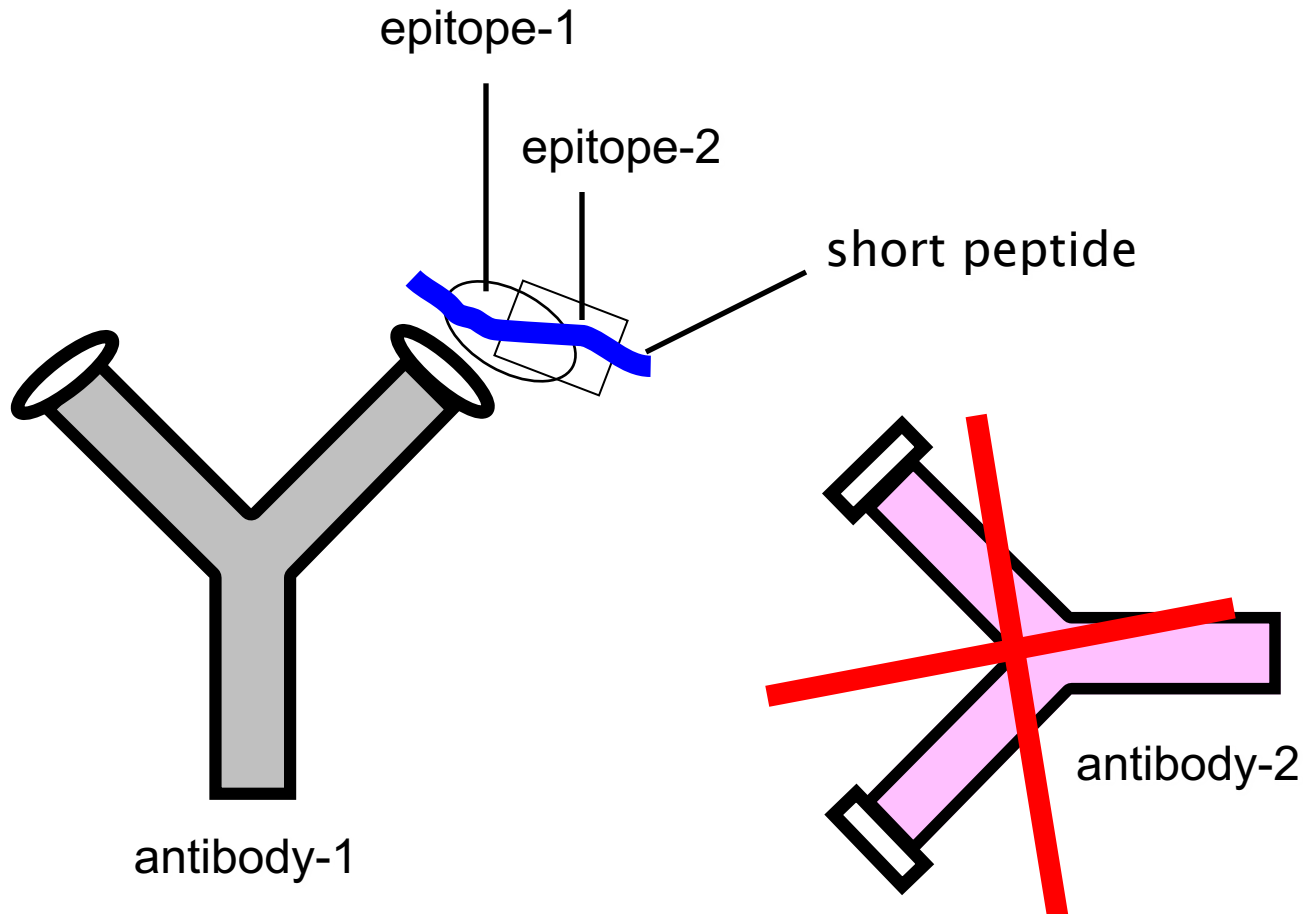
CMV26-decapeptide scan

10	20	30	40	50
IEGRGKSRGGGGGG <u>SLSS</u> LANAGGLHDDGPGLDNDLMNEPMGLGGLGGGGGGGGKKH				
1. IEGRGKSRGG	21. ANAGGLHDDG	41. PMGLGGLGGG		
2. EGRGKSRGGG	22. NAGGLHDDGP	42. MGLGGLGGGG		
3. GRGKSRGGGG	23. AGGLHDDGPG	43. GLGGLGGGGG		
4. RGKSRGGGGG	24. GGLHDDGPGL	44. LGGLGGGGGG		
5. GKSRRGGGGG	25. GLHDDGPGLD	45. GGLGGGGGGG		
6. KSRGGGGGGG	26. LHDDGPGLDN	46. GLGGGGGGGG		
7. SRGGGGGGGS	27. HDDGPGLDND	47. LGGGGGGGGK		
8. RGGGGGGGSL	28. DDGPGLDNDL	48. GGGGGGGGGK		
9. GGGGGGGSLS	29. DGPGLDNDLM	49. GGGGGGGGGK		
10. GGGGGGSLSS	30. GPGLDNDLMN			
11. GGGGGSLSSL	31. PGLDNDLMNE			
12. GGGGSLSSLA	32. GLDNDLMNEP			
13. GGGSLSSLAN	33. LDNDLMNEPM			
14. GGSLSSLANA	34. DNDLMNEPMG			
15. GSLSSLANAG	35. NDLMNEPMGL			
16. SLSSLANAGG	36. DLMNEPMGLG			
17. LSSLANAGGL	37. LMNEPMGLGG			
18. SSLANAGGLH	38. MNEPMGLGGL			
19. SLANAGGLHD	39. NEPMGLGGLG			
20. LANAGGLHDD	40. EPMGLGGLGG			

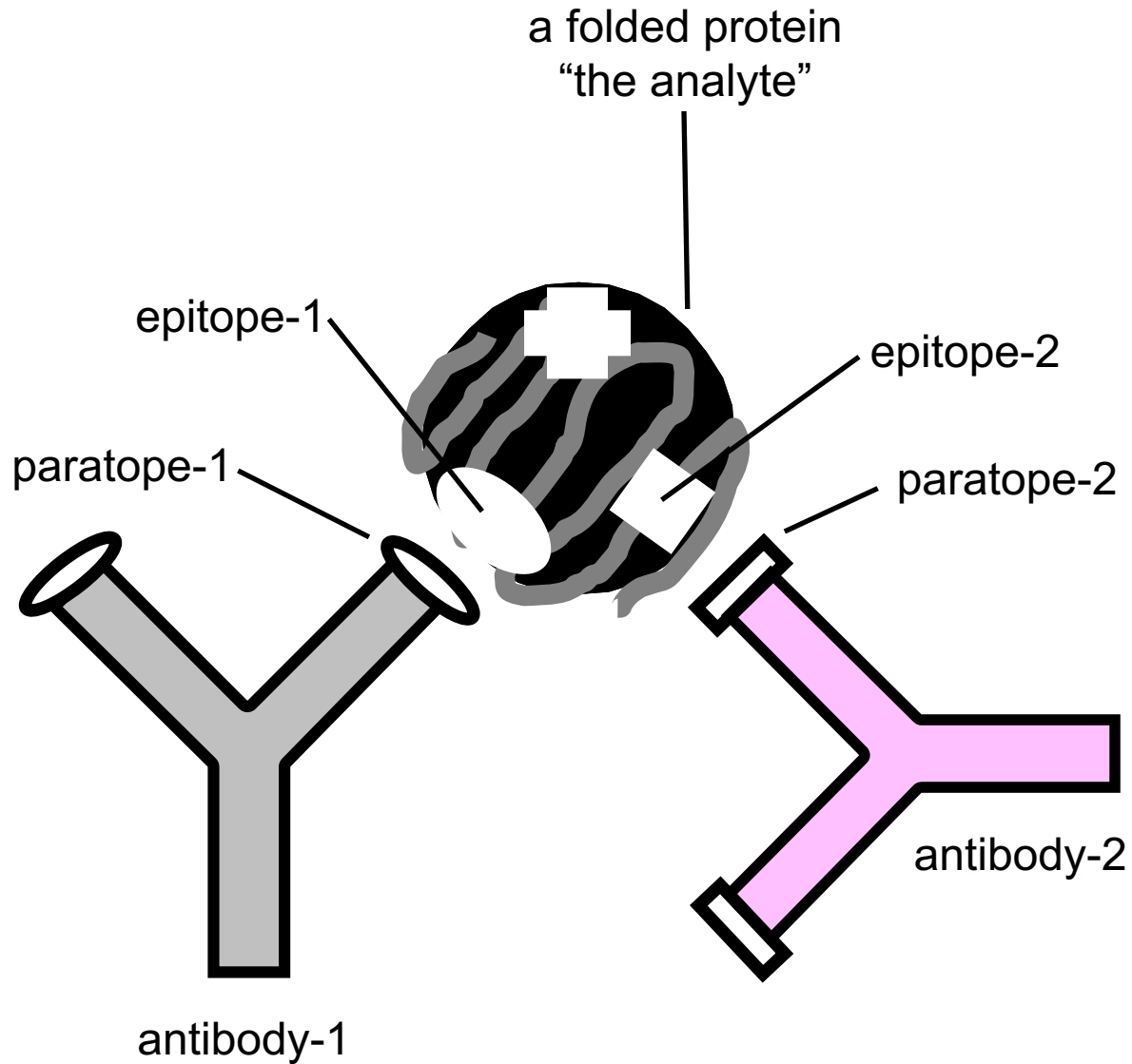
=> But only continuous epitopes

Overlapping epitopes

Even small analytes can have multiple epitopes, but antibody binding to one epitope **blocks** another epitope, i.e. these epitopes are **overlapping**



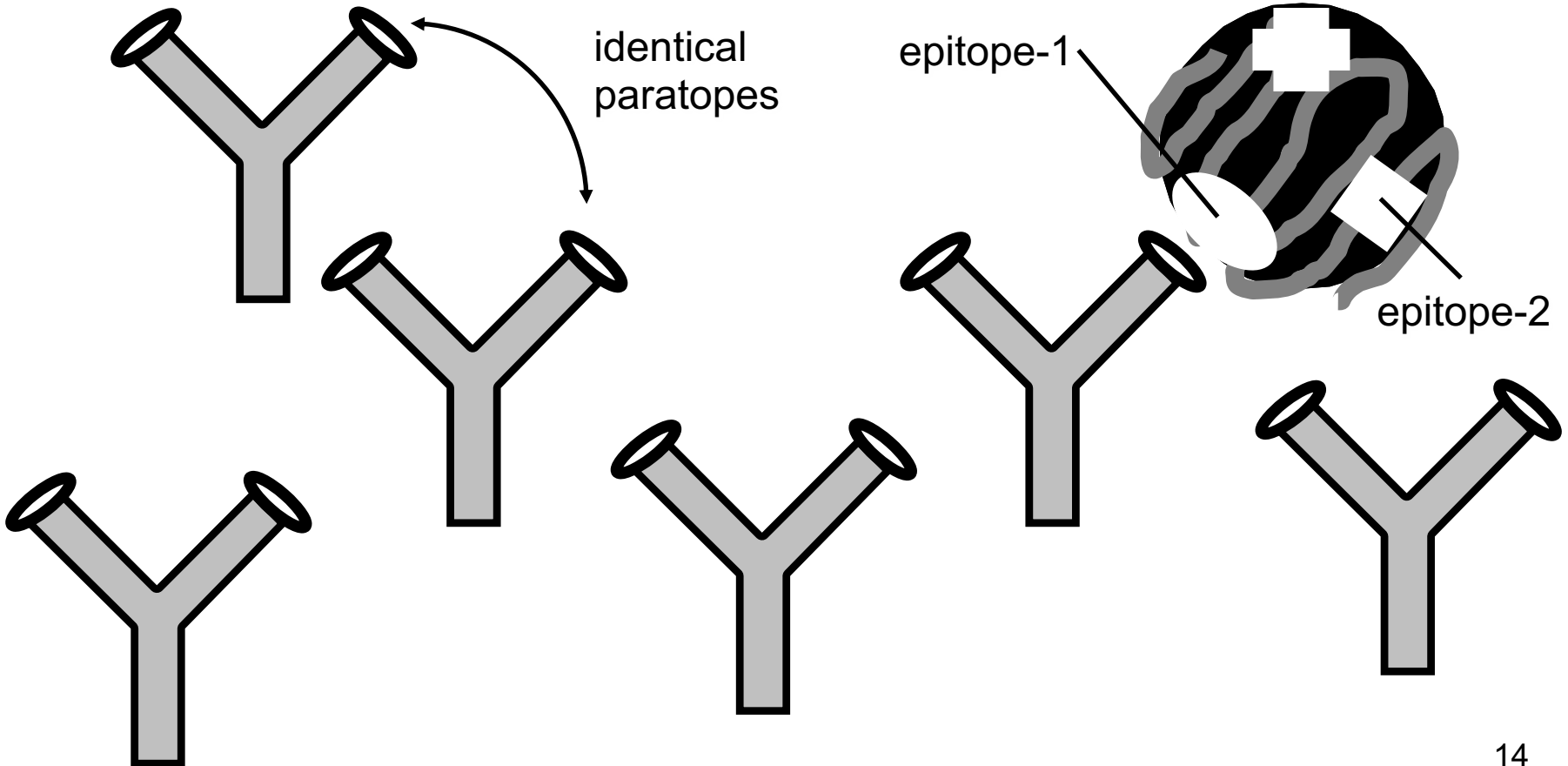
Non-overlapping epitopes



Monoclonal antibody reagent

all antibodies are from the **same B cell clone**
=> reagent consist of identical antibodies,
and all recognize and are specific
for only one identical epitope

.. will bind only to one specific
epitope in the analyte - unless
there are multiple identical
epitopes in the same analyte

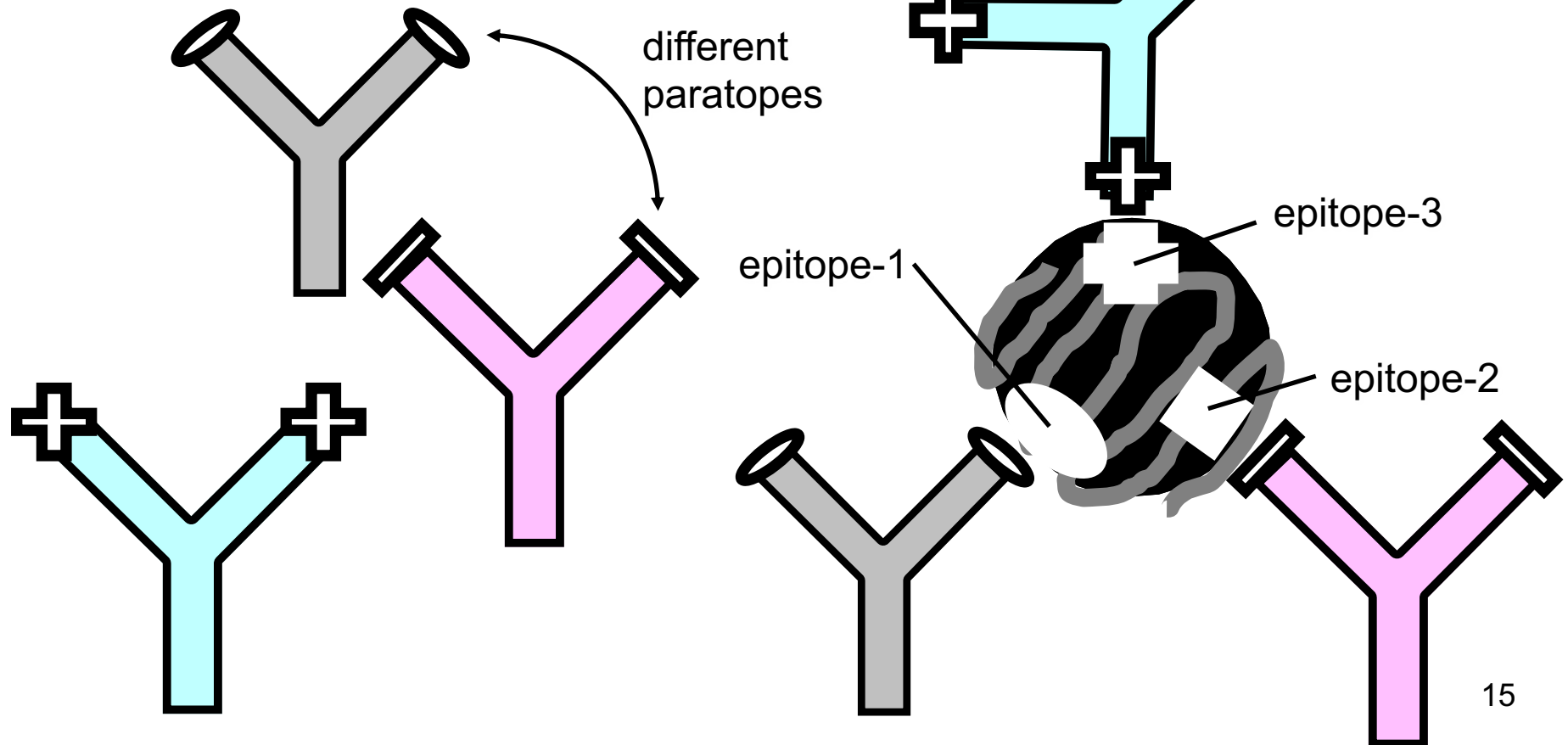


Polyclonal antibody reagent

Mix of **different B cell clones**

=> reagent consist of antibodies that have different paratopes and recognize different epitopes, but exact composition is not known

... will bind simultaneously to one or several non-overlapping epitopes in the analyte



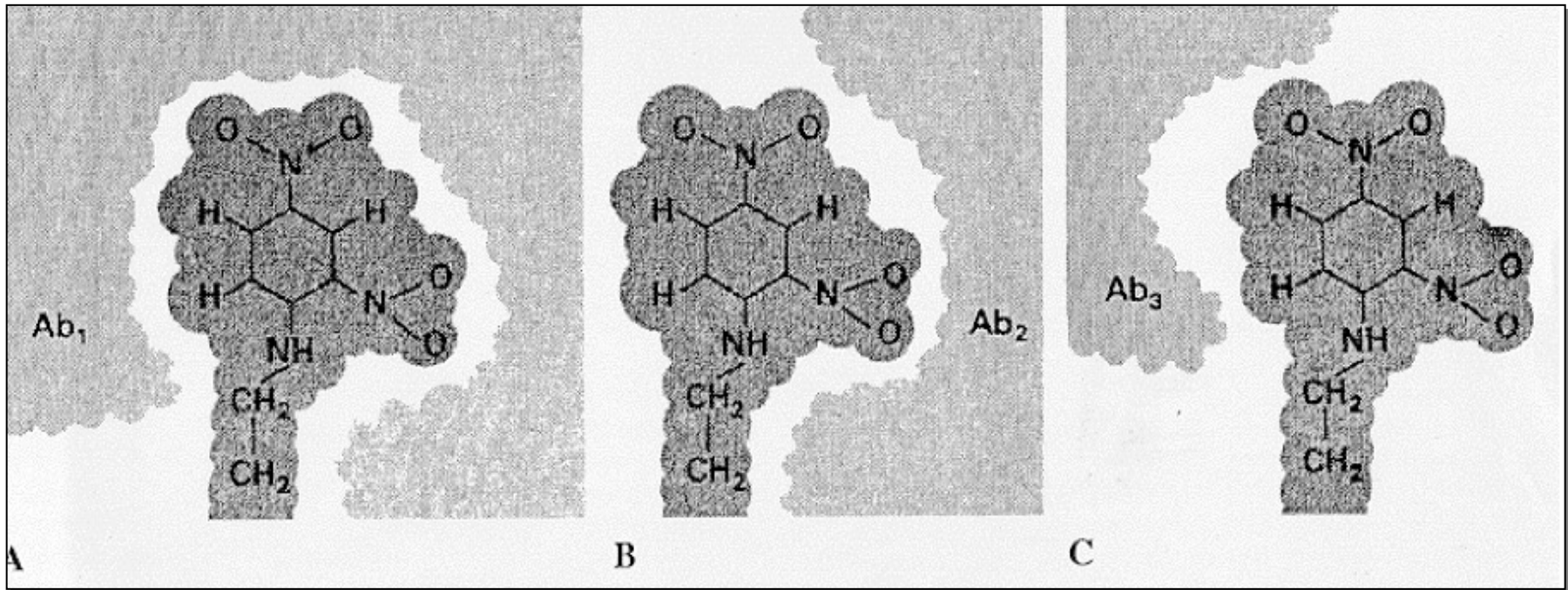
Antibody affinity

Affinity of an antibody

Tight fit /
high affinity

Less interaction
/ lower affinity

Little interaction /
very low affinity

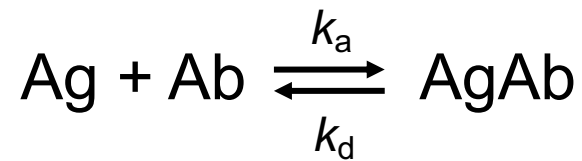
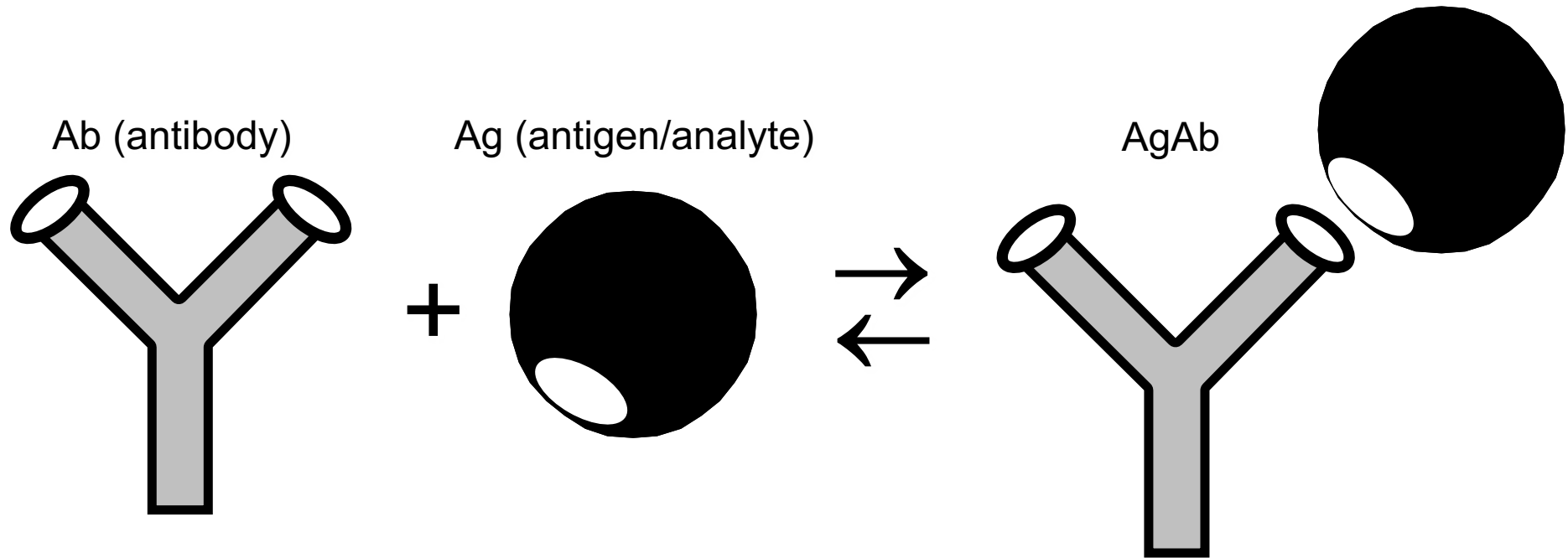


good epitope specificity

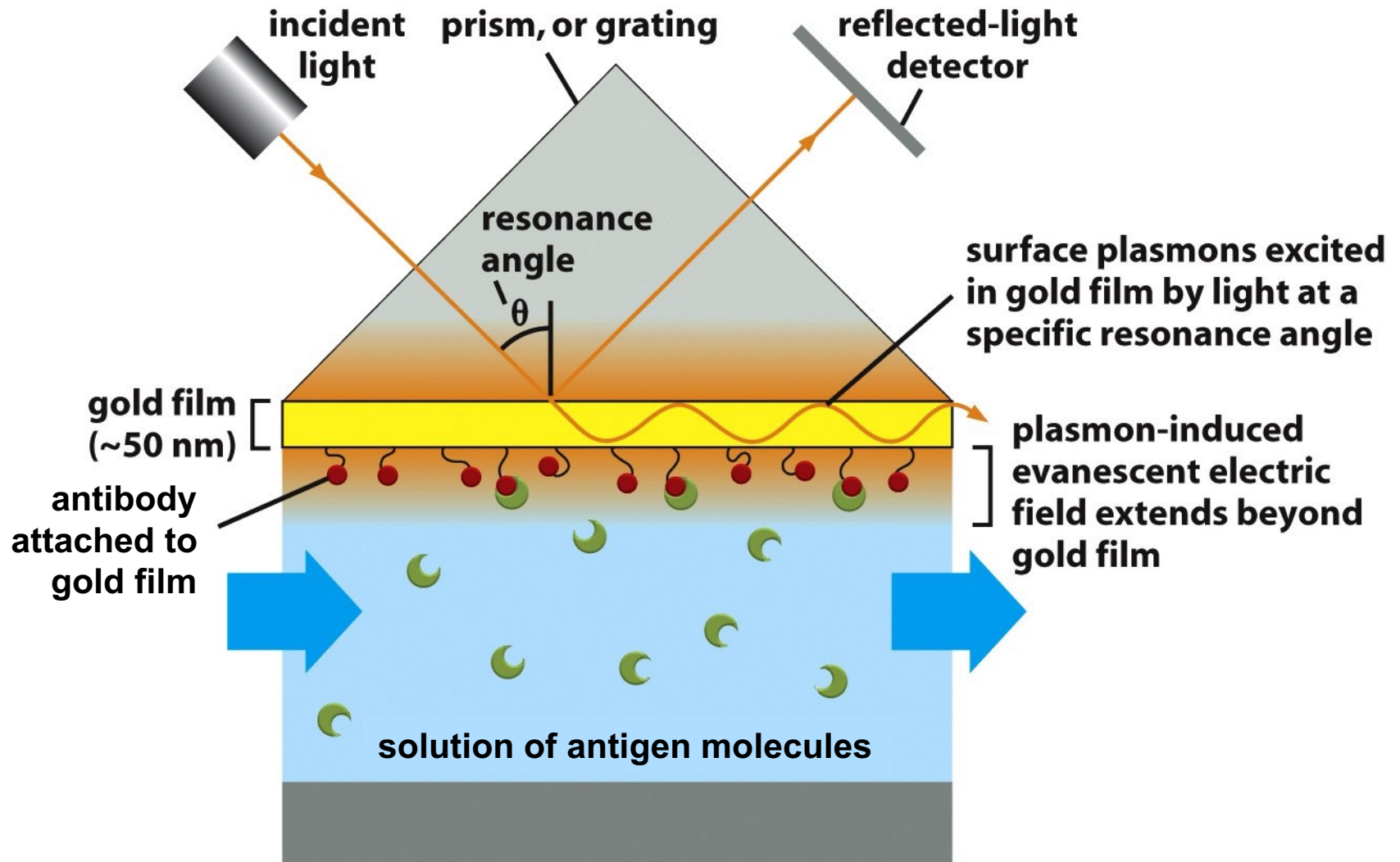


decreasing epitope specificity

Antibody-antigen binding reaction

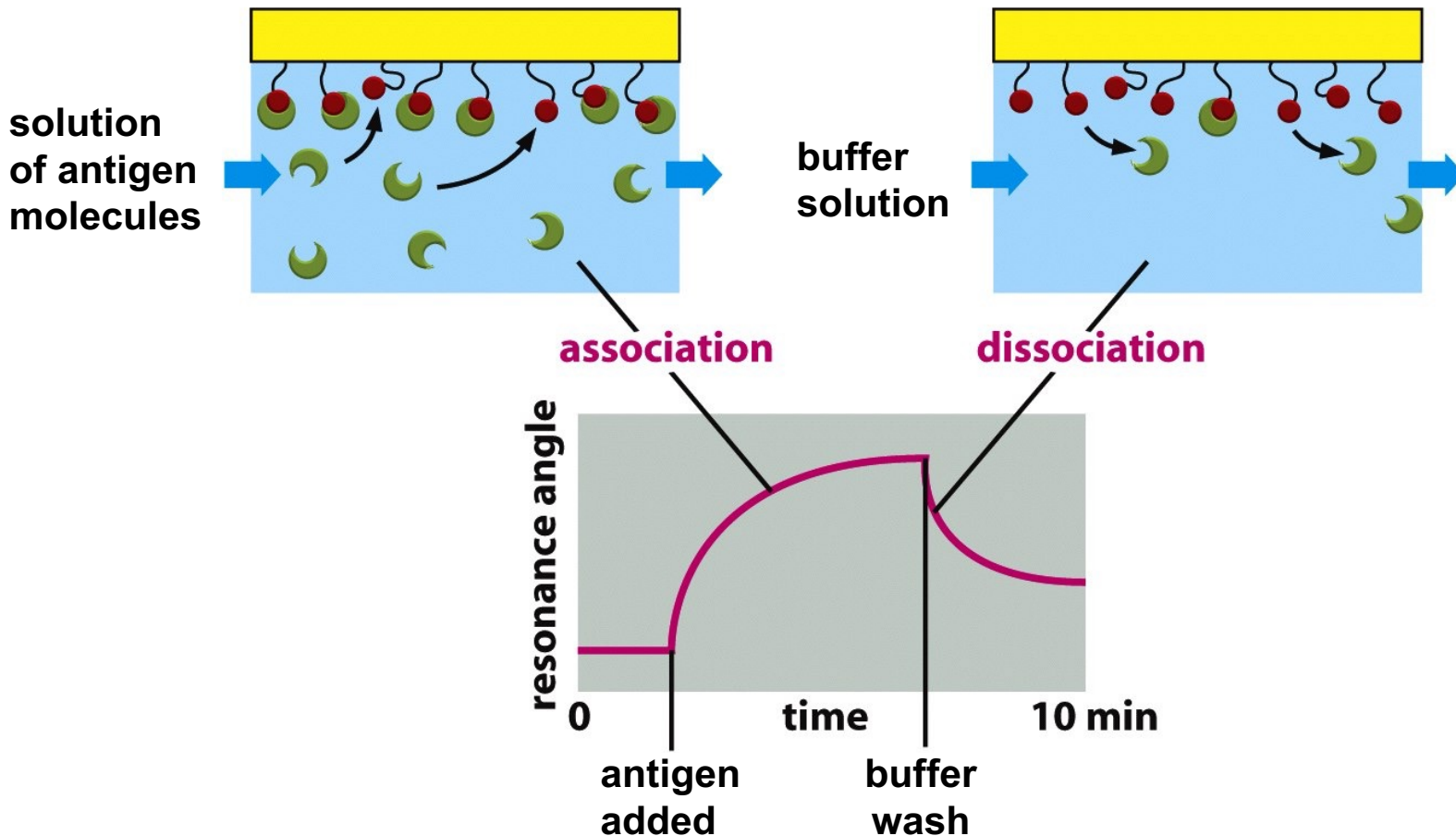


Surface plasmon resonance (SPR)

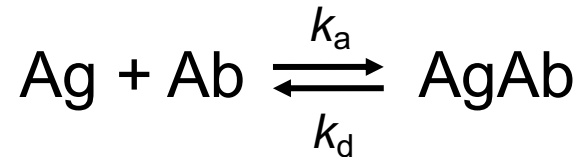


Determining the affinity of antibodies by SPR

- (1) Binding of antigen to surface immobilized antibodies increases the refractive index of the surface layer.
- (2) The resulting change of the resonance angle for plasmon induction can be measured by a photodetector.



Affinity of an antibody



$$k_a[\text{Ag}][\text{Ab}] = k_d[\text{AgAb}]$$

reaction velocities at equilibrium:

$$K = \frac{k_a}{k_d} = \frac{[\text{AgAb}]}{[\text{Ag}][\text{Ab}]}$$

← Bound antibody and antigen

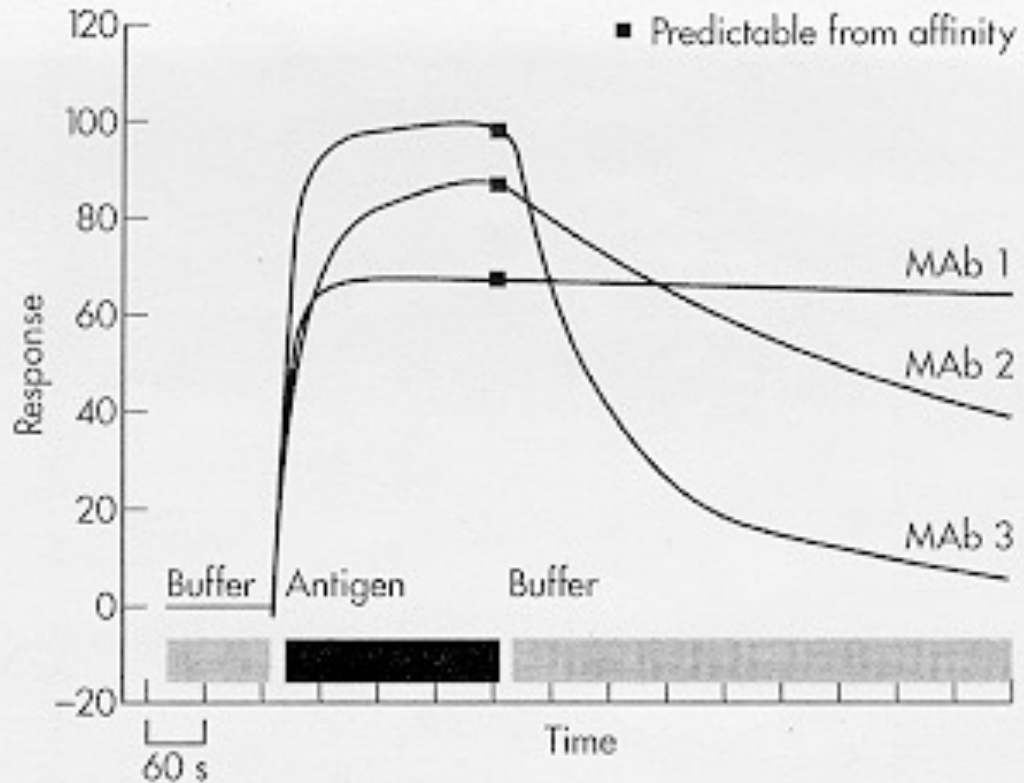
free antigen free antibody

k_a : association rate constant (on rate)

k_d : dissociation rate constant (off rate)

K: affinity constant

Affinity of an antibody



extremely stable binding:
off-rate is very slow

poorly suited for immuno-
assay due to dissociation
problems, e.g. during
wash or during 2nd step
if used as capture antibody

Fig. 1.6 Response curves illustrating the interaction of P24 antigen (125 nM) with three different monoclonal antibodies (MAbs).

81

Affinity of an antibody

$$K = \frac{k_a}{k_d} = \frac{[AgAb]}{[Ag][Ab]}$$

← Bound antibody and antigen
↑ free antigen ↑ free antibody

approximate calculation of concentrations in equilibrium:

if $[Ag]_{tot} \ll [Ab]_{tot}$, only a very small antibody fraction is present in the complex $[AgAb]$

$\Rightarrow [Ab] \approx [Ab]_{tot}$

$$\begin{aligned} [Ag]_{tot} &= [Ag] + [AgAb] \\ [Ab]_{tot} &= [Ab] + [AgAb] \end{aligned}$$



free (unbound) concentrations

$$[AgAb] = \frac{[Ab]_{tot} [Ag]_{tot} K}{([Ab]_{tot} K) + 1}$$

Affinity of an antibody

Calculating the equilibrium concentration

$$[AgAb] = \frac{[Ab]_{tot}[Ag]_{tot} K}{([Ab]_{tot} K) + 1}$$

$$[Ab]_{tot} = 1 * 10^{-9} \text{ M}$$

$$[Ag]_{tot} = 1 * 10^{-12} \text{ M (i.e. much smaller)}$$

$$K = 1 * 10^9 \text{ M}^{-1}$$

by calculating we get $[AgAb] = 0.5 * 10^{-12} \text{ M}$ (i.e. 50%)

"rule of thumb":

when	$[Ab]_{tot} = 1/K$	then	$[AgAb] = 50\% [Ag]_{tot}$
	$[Ab]_{tot} = 10/K$	then	$[AgAb] = 90\% [Ag]_{tot}$
	$[Ab]_{tot} = 0.1/K$	then	$[AgAb] = 10\% [Ag]_{tot}$

Affinity of an antibody

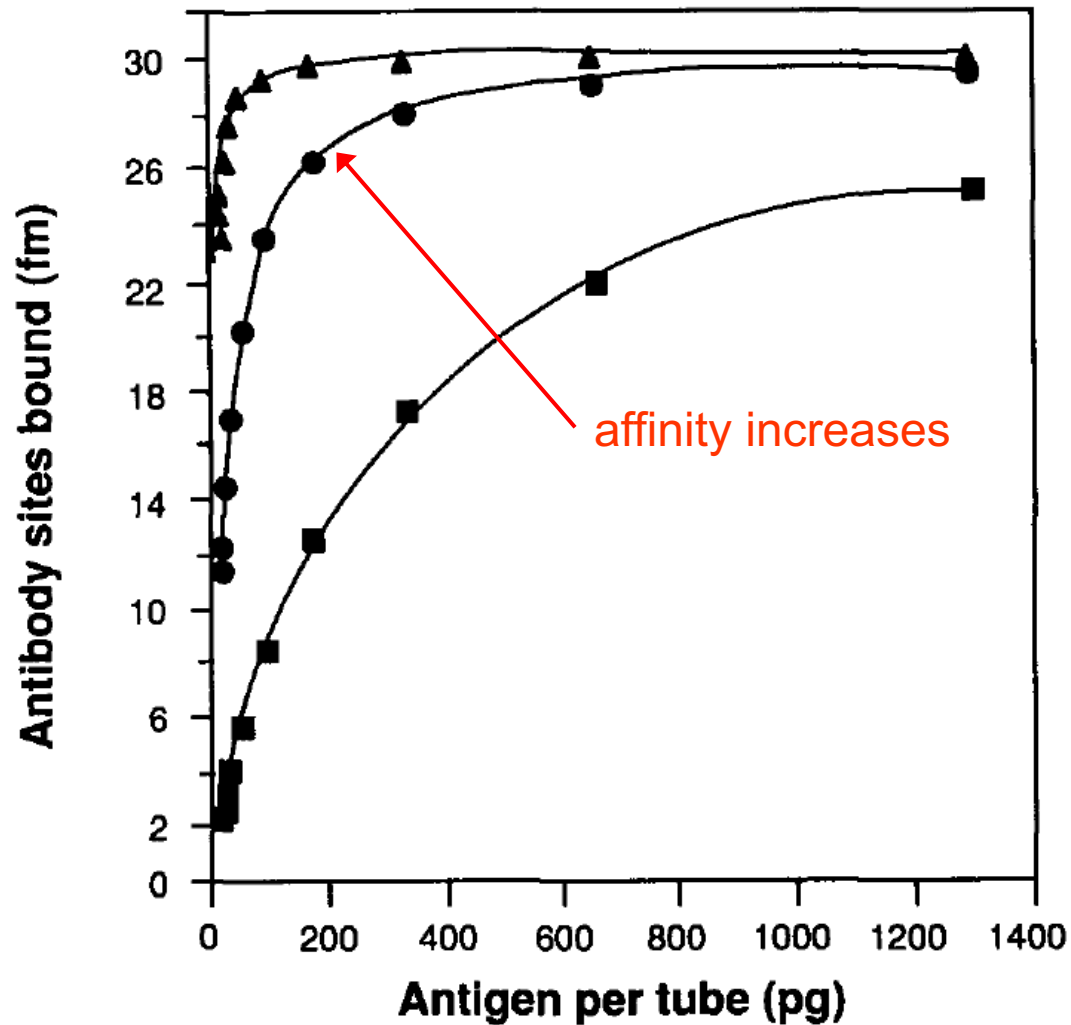
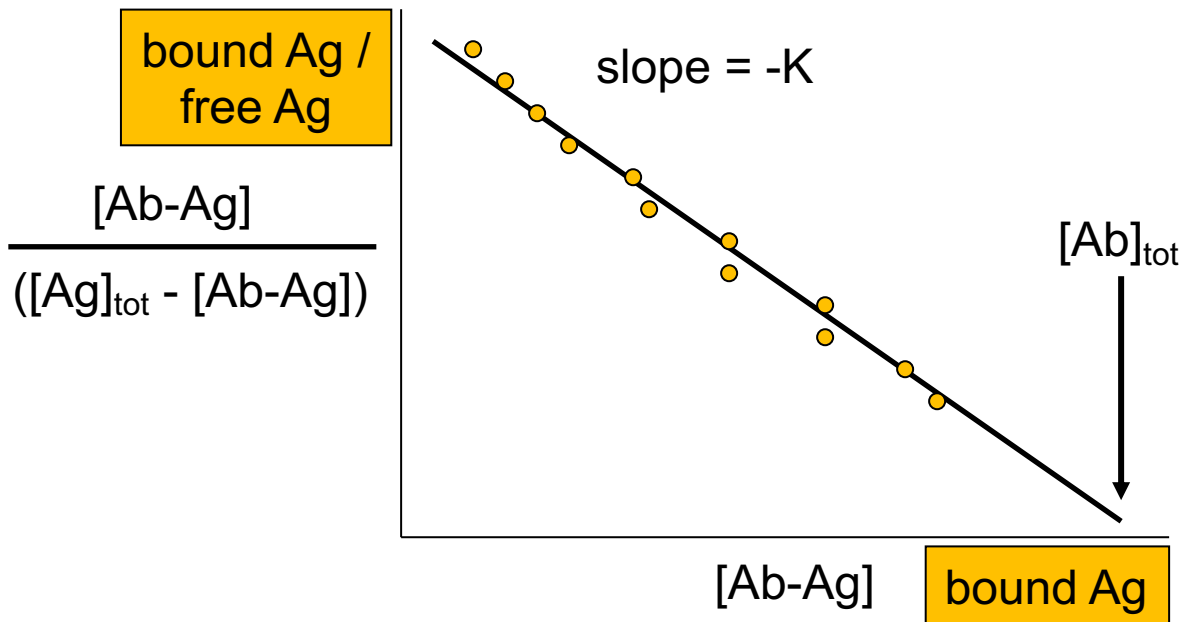


Figure 8.4 Estimation of filled antibody sites, at different concentrations of antigen, for three antibodies with different affinity constant (l/mol). ■ = 1×10^9 , ● = 1×10^{10} , ▲ = 1×10^{11} .

Affinity of an antibody: Scatchard plot

linearization:

$$K ([Ab]_{\text{tot}} - [Ab-Ag]) = \frac{[Ab-Ag]}{([Ag]_{\text{tot}} - [Ab-Ag])} = \frac{\text{bound Ag}}{\text{free Ag}}$$



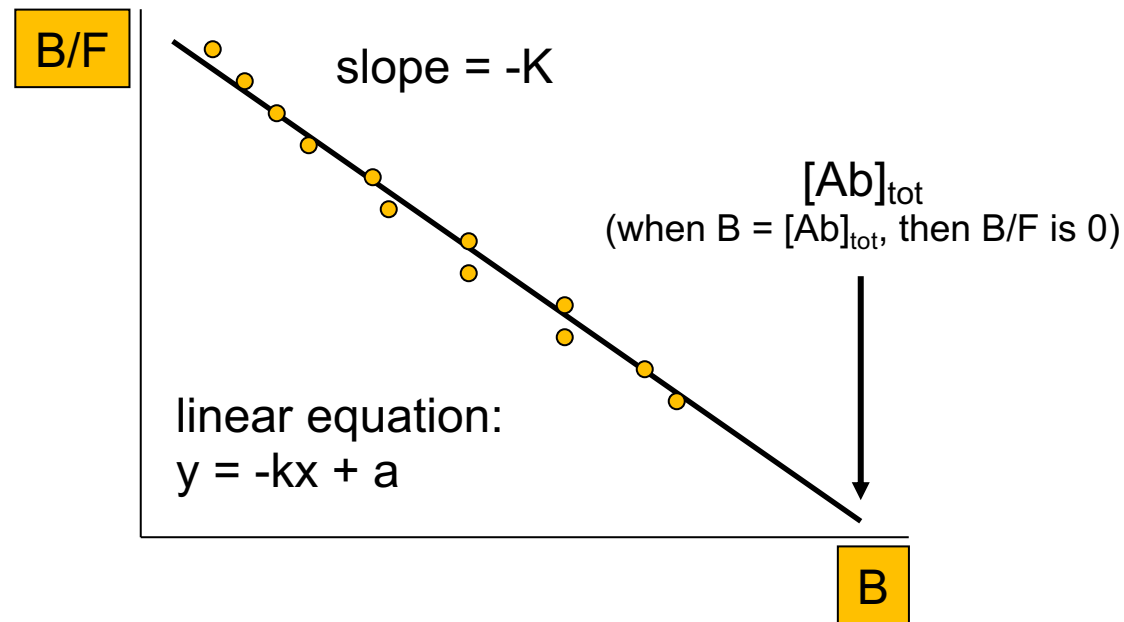
Affinity of an antibody: Scatchard plot

$$B = [Ag-Ab]$$

$$F = [Ag] = [Ag]_{tot} - [Ag-Ab]$$

$$K ([Ab]_{tot} - B) = B/F \Rightarrow B/F = -K B + K [Ab]_{tot}$$

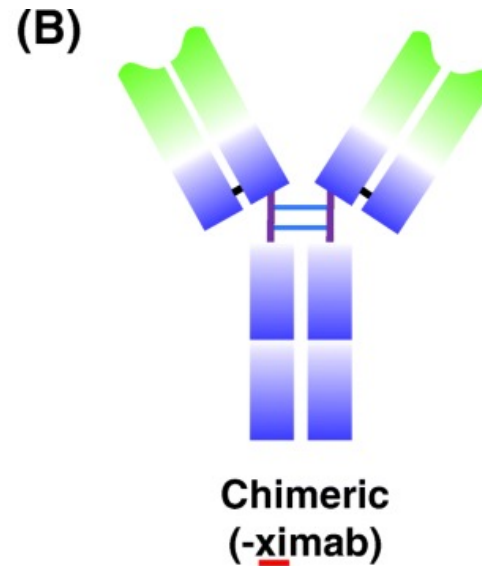
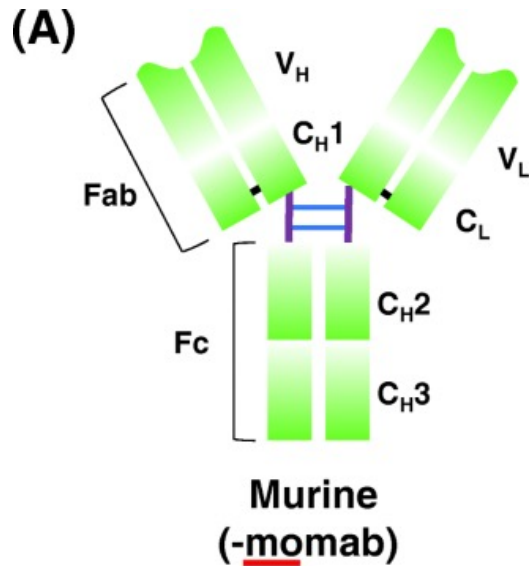
constant



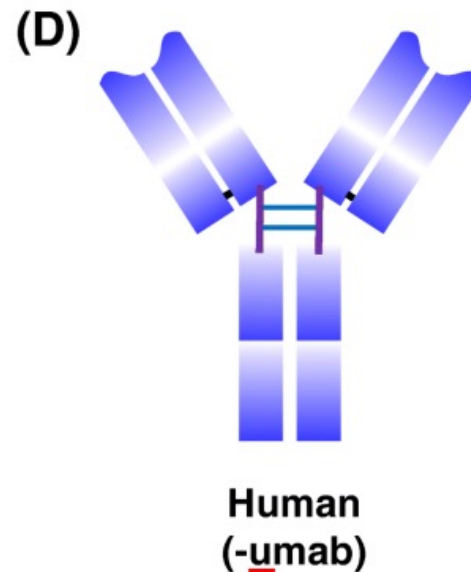
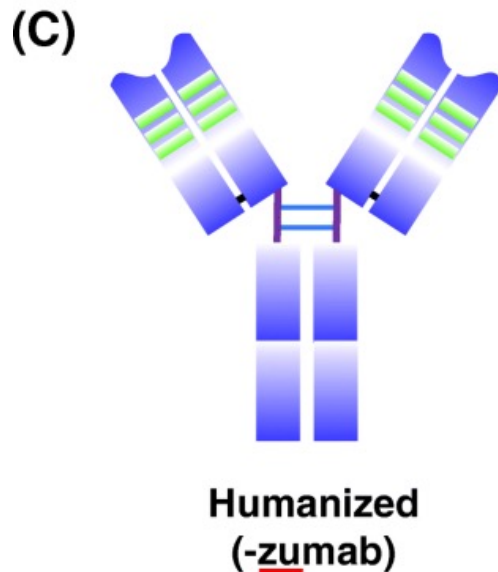
=> Typically replaced by non-linear fitting using computer programs

Antibody engineering

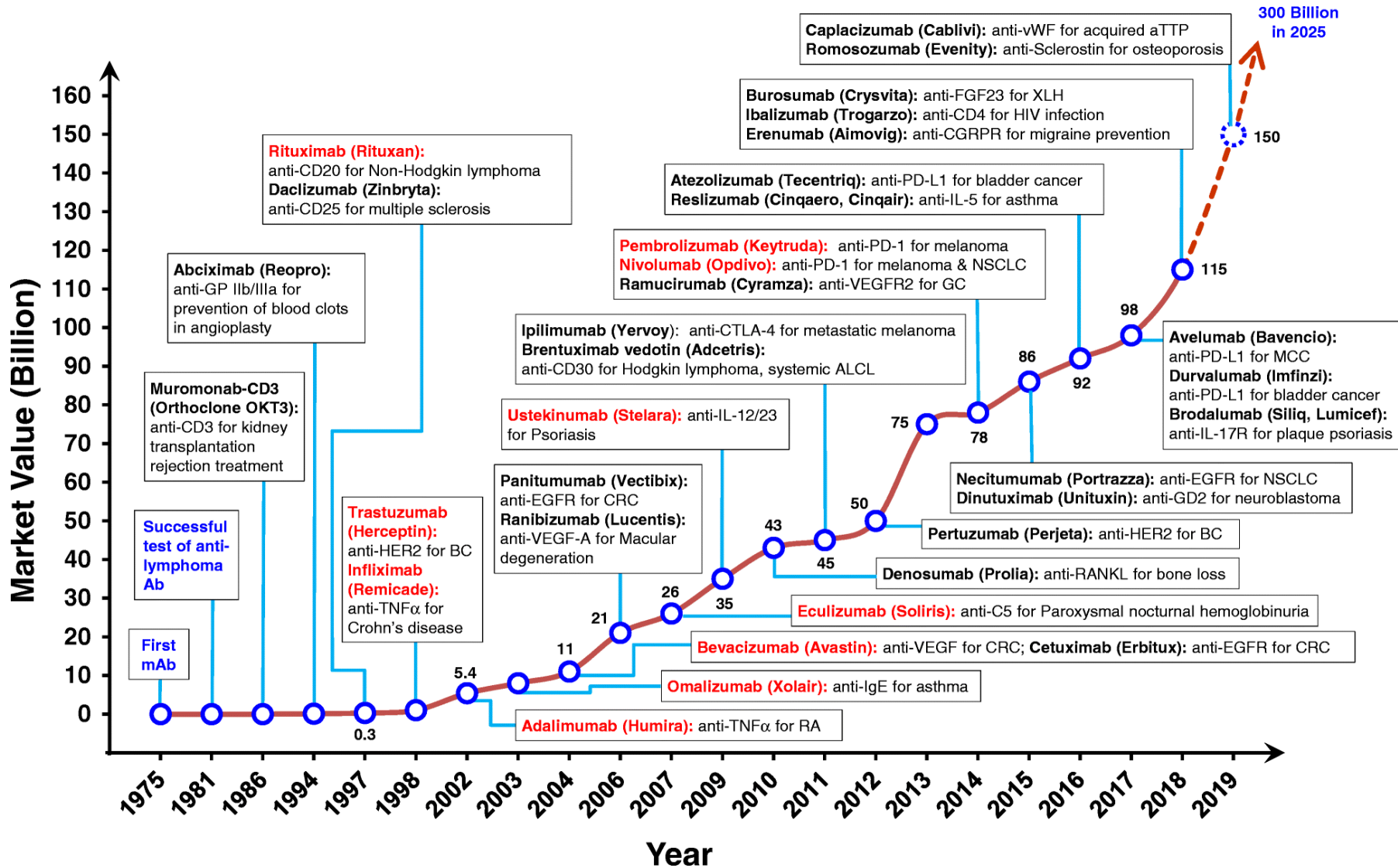
Excursion: Antibody engineering for therapy



Natural antibodies
(raised in mice) are
potentially immunogenic
=> Potential side effects



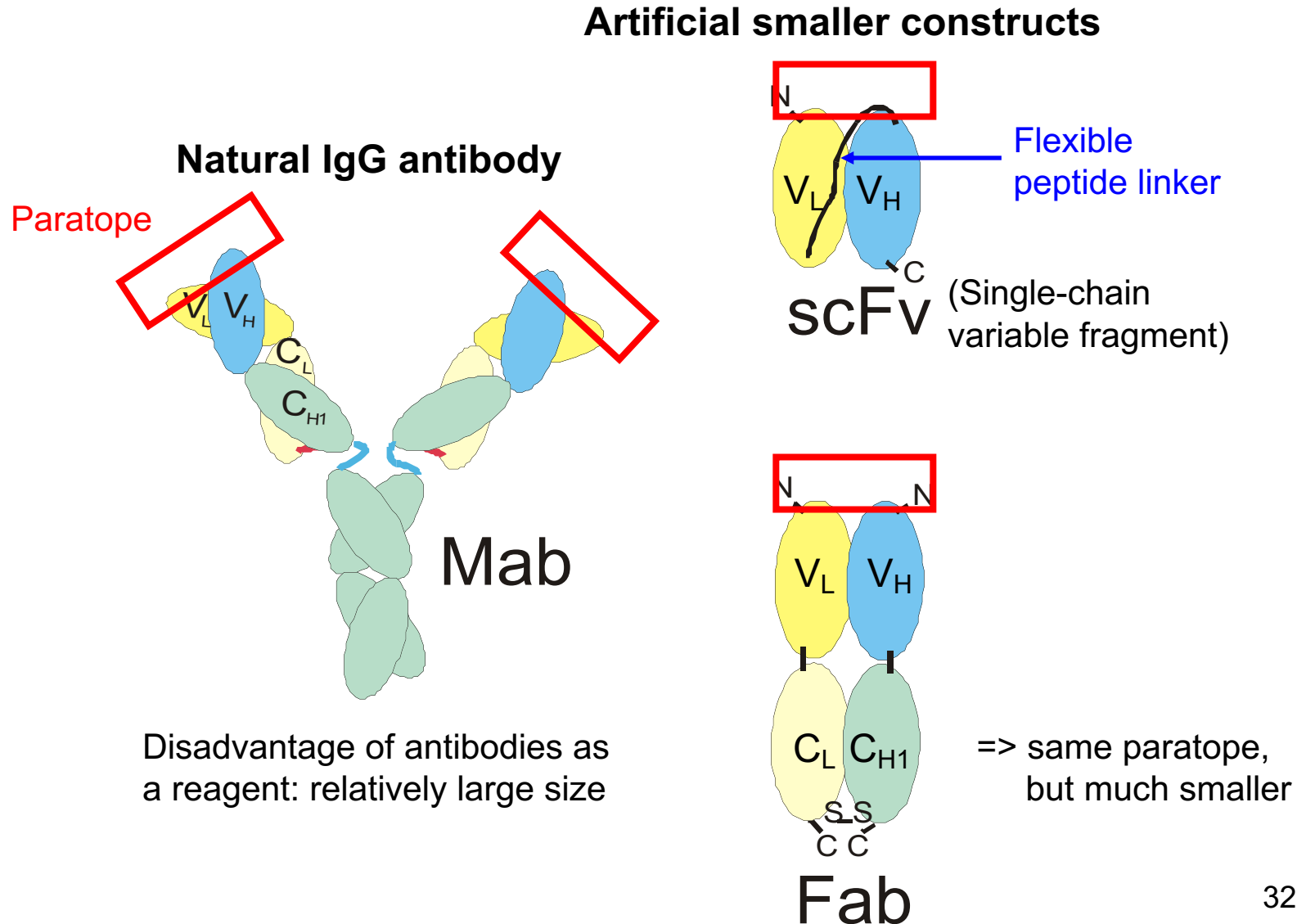
Therapeutic antibodies (market value)



Humanized antibodies

No.	Drug	Indication (1st US FDA Approval Year)	Company	2018 Revenue (USD, billion)
1	Adalimumab (Humira)	Rheumatoid arthritis (2002) Psoriatic arthritis (2005) Ankylosing spondylitis (2006) Juvenile Idiopathic Arthritis (2008) Psoriasis (2008) Crohn's disease (2010) Ulcerative colitis (2012) Hidradenitis suppurativa (2015) Uveitis (2018)	AbbVie	\$19.9 bn
2	Nivolumab (Opdivo)	Melanoma (2015) Non-small cell lung cancer (2015) Renal cell carcinoma (2015) Head and neck squamous cell (2016)	Bristol-Myers Squibb	\$7.6 bn
3	Pembrolizumab (Keytruda)	Melanoma (2014) Head and neck cancer (2016) Non-small cell lung cancer (2015) Lymphoma (2018) Cervical cancer (2018) Microsatellite instability-high cancer (2018)	Merck & Co	\$7.2 bn
4	Trastuzumab (Herceptin)	Breast cancer (1998) Gastric cancer (2010)	Roche	\$7.0 bn
5	Bevacizumab (Avastin)	Colorectal cancer (2004) Non-small cell lung cancer (2006) Breast ERB2 negative cancer (2008) Renal cell carcinoma (2009) Glioblastoma (2011)	Roche	\$6.8 bn
6	Rituximab,	Non-Hodgkin's lymphoma (1997)	Roche	\$6.8 bn

Recombinant antibody fragments



Recombinant antibody fragments

Immortalization of hybridomas through cloning

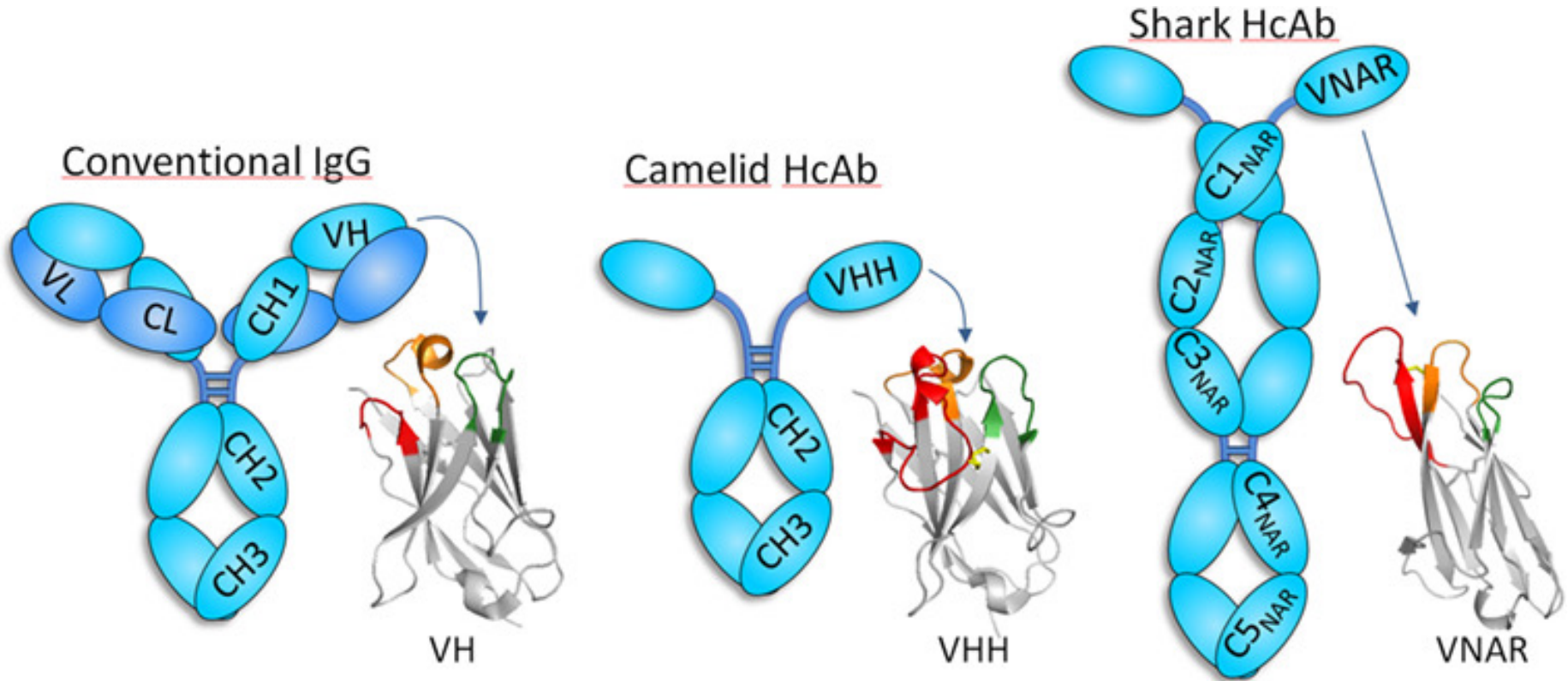
or

generation of new **antibodies without immunization**

- Greater speed of **production** (*E. coli batch fermentation*)
- New specificities especially for **poor immunogens**
- Possibility to **fine-tune** antibody specificity and affinity
- Possibility to **tailor make** the antibody to perform special tasks
 - tags, handles (for conjugation, immobilization)
 - fusing to other protein (e.g. enzymes)

Likely to be increasingly used in **miniaturised systems** to enable full control of antibody performance.

Heavy chain antibodies

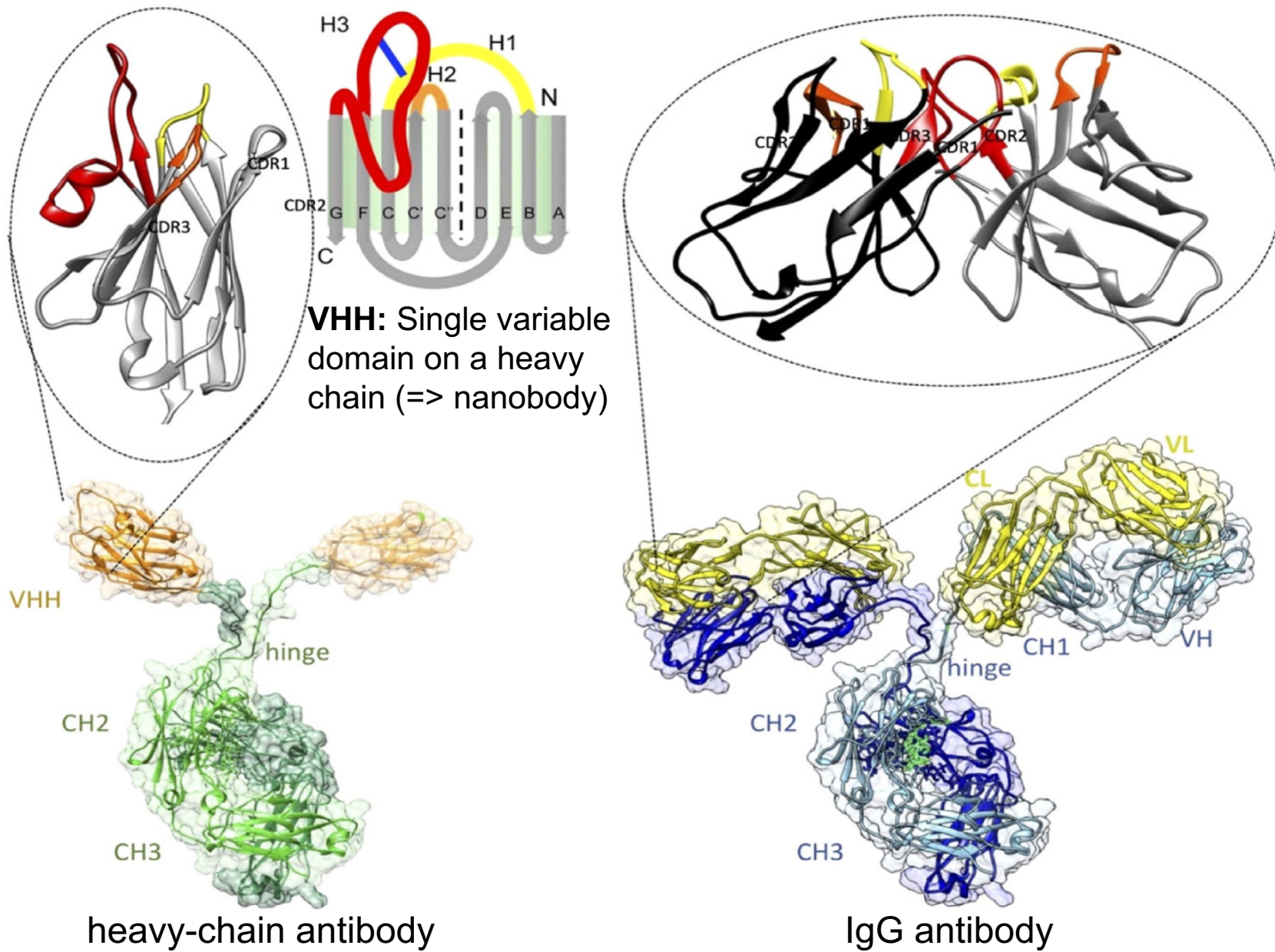


our own most
common antibody

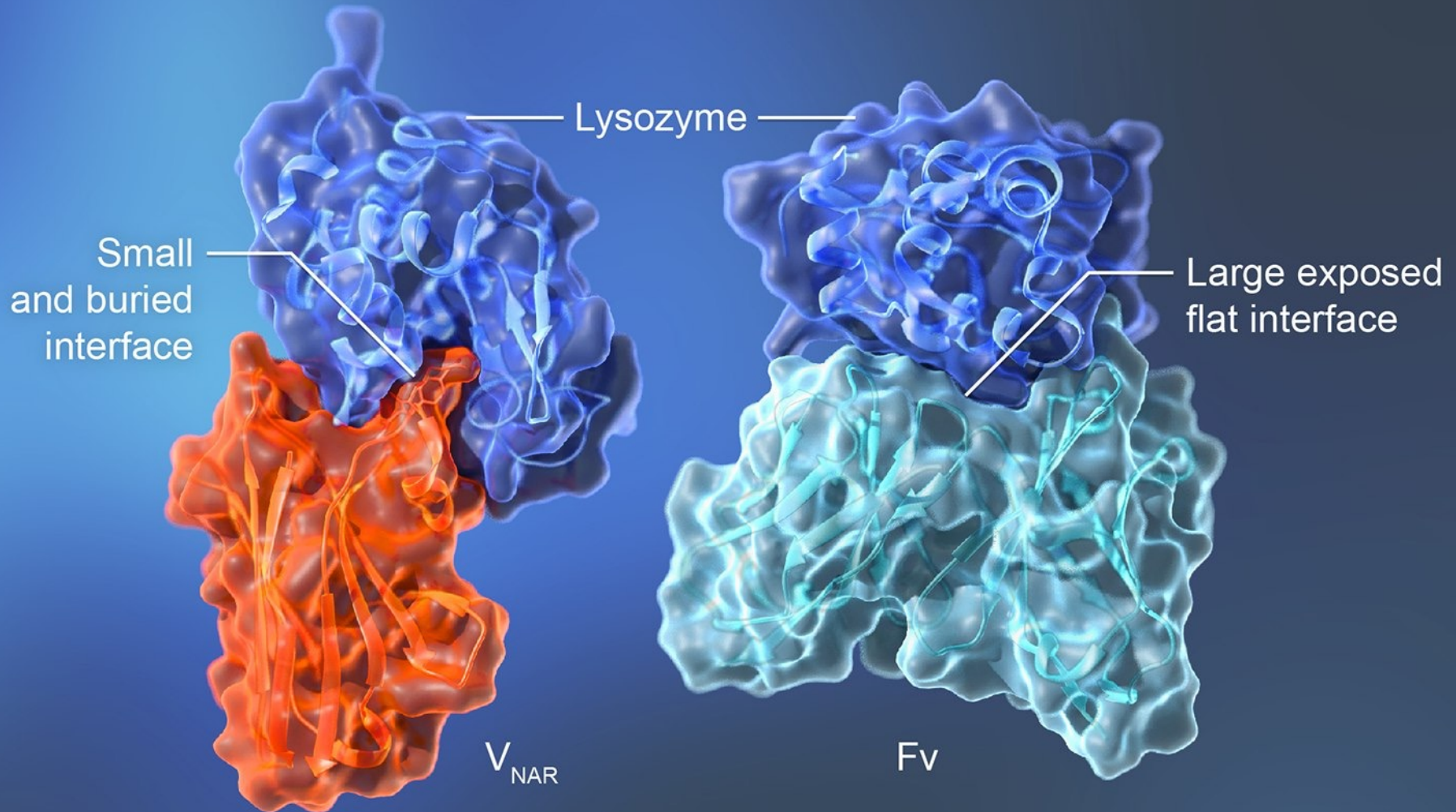
heavy chain antibodies
(velbloud, dromedár, lama)

(žralok)

From heavy chain antibodies to nanobodies



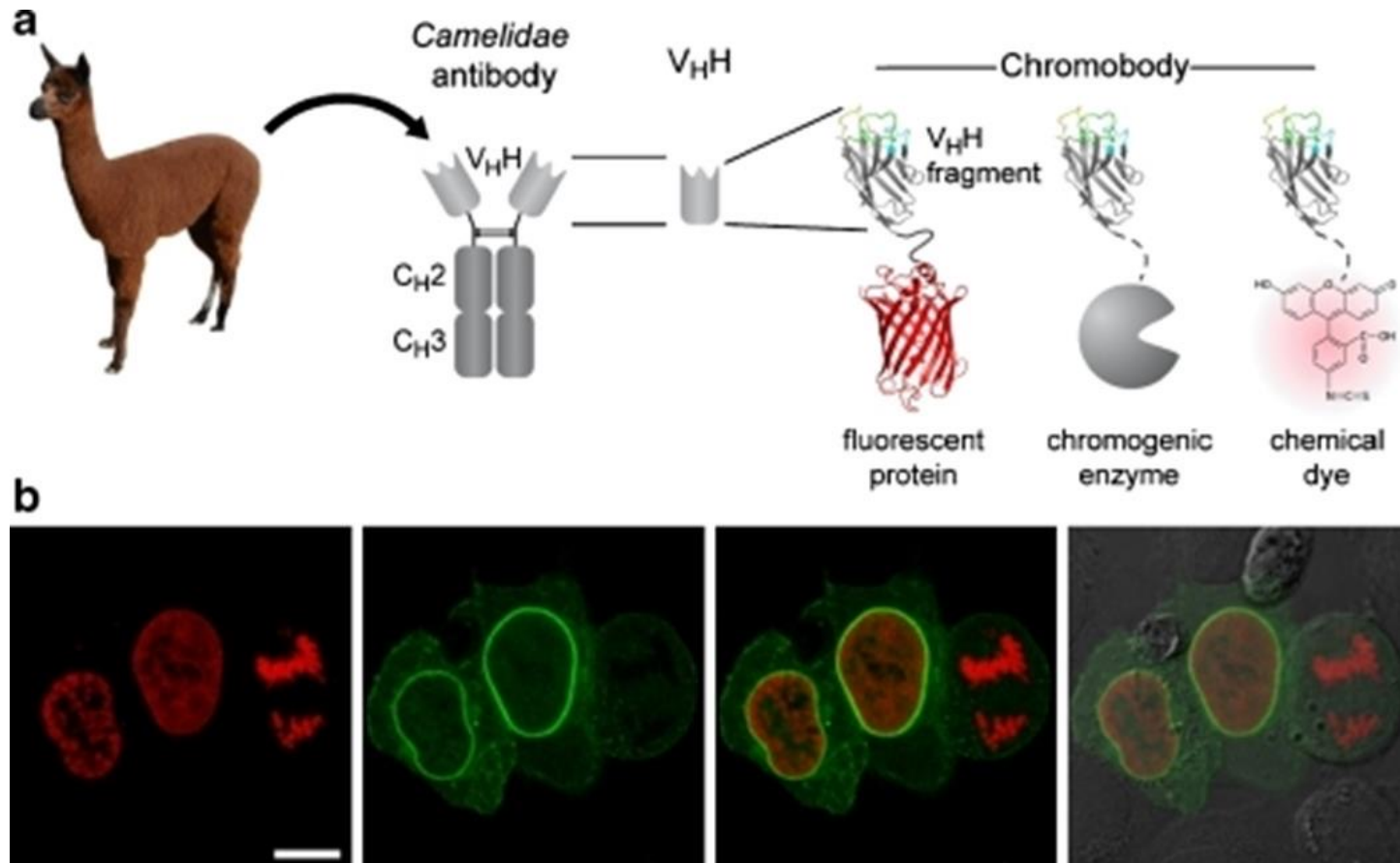
Nanobodies: Detection of hidden epitopes



Advantages of nanobodies

- Mass: ca. 15 kDa (IgG: 150 kDa), 2.5 nm diameter (IgG 15 nm)
- High solubility
- Rapid targeting and fast blood clearance
- Detection of “hidden“ epitopes
- Easy cloning: Recombinant engineering and protein expression *in vitro* in bacterial production systems are much simpler
- Very stable and heat resistant (no cold storage required)
- Simple genetic structure allows easy re-engineering of nanobodies to introduce new antigen-binding characteristics or attach labels

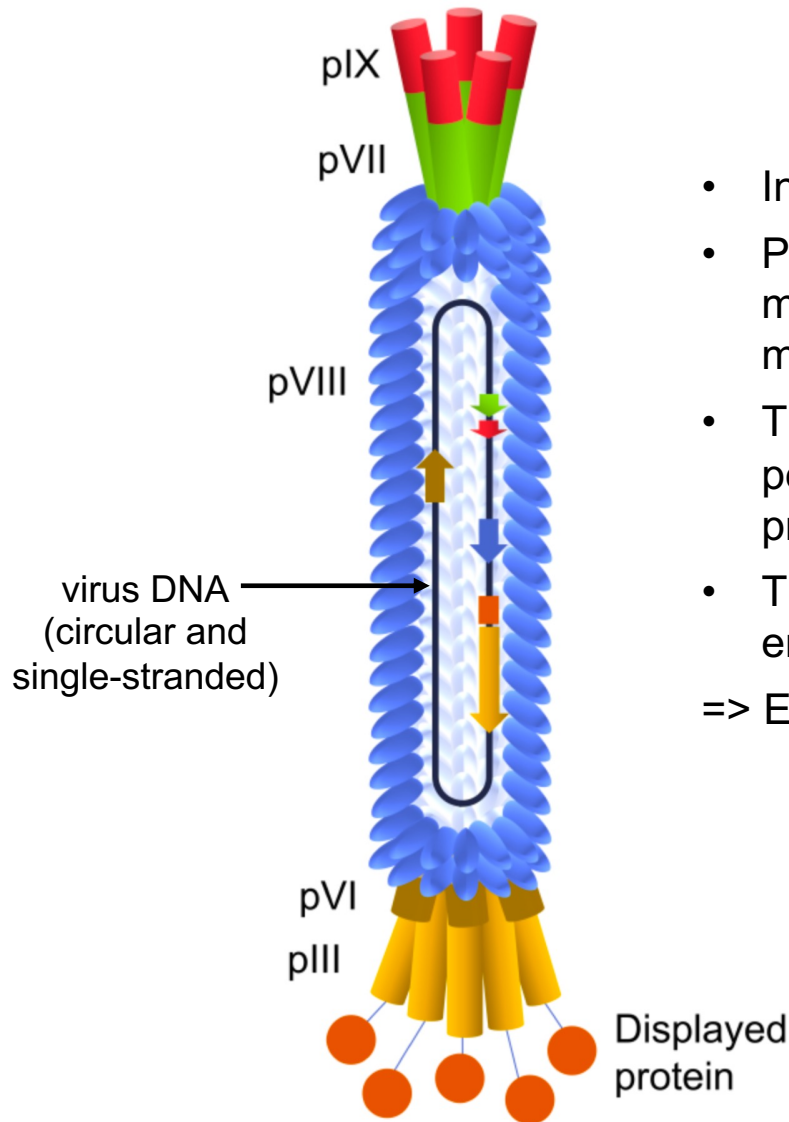
Recombinant nanobodies



a Chromobodies.

b Detection of the nuclear lamina with lamin chromobody in living cells. Confocal images of HeLa cells coexpressing lamin chromobody (*green*) and red fluorescent histone H2B as a mitosis marker. Scale bar: 10 μm

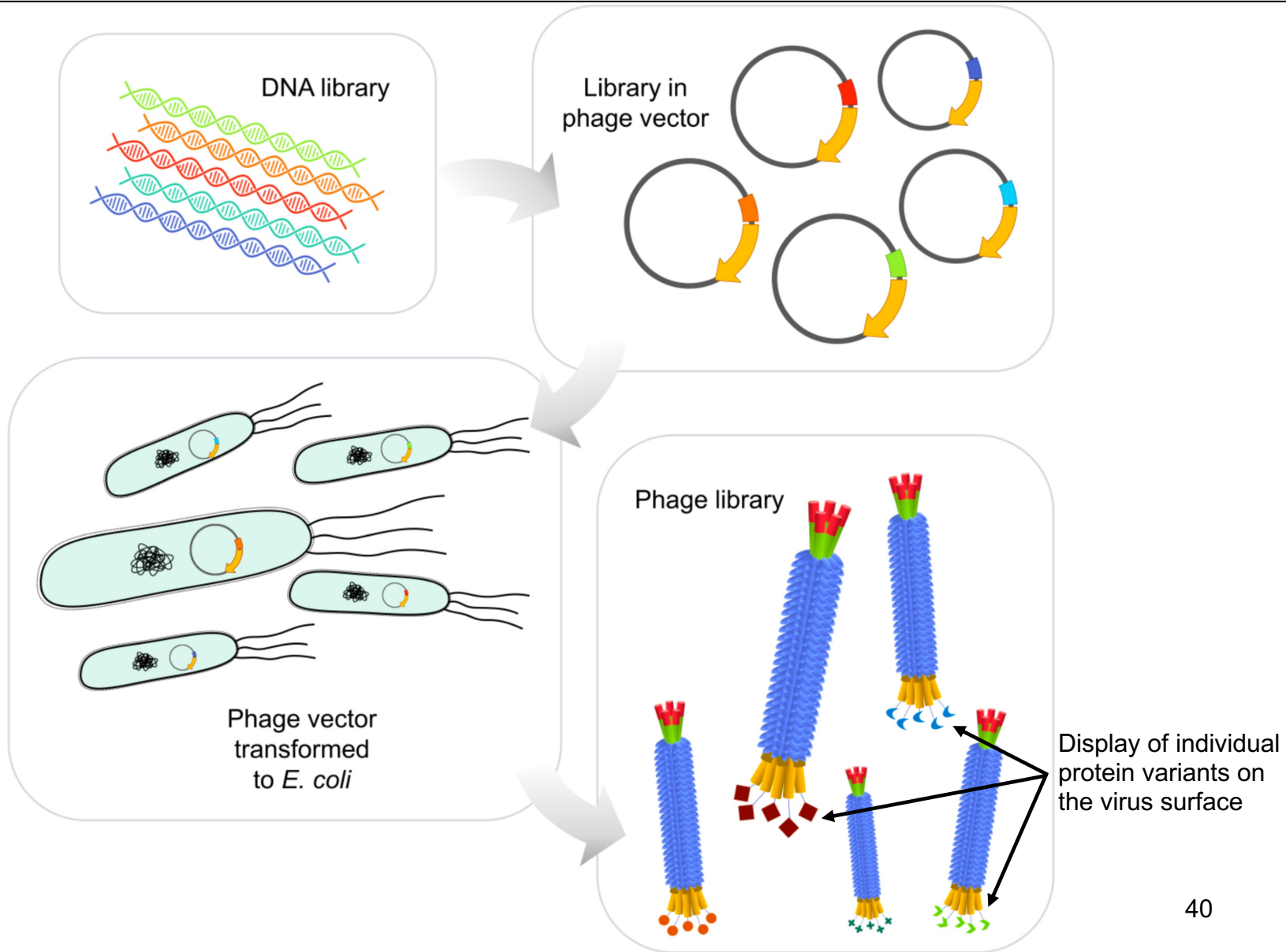
Phage display using filamentous phage M13



- Infects / replicates in *E. coli*
 - Protein coat:
major coat protein: pVIII
minor coat proteins: pIII, pVI, pVII, pIX
 - The phage can be engineered to display foreign peptides or proteins as a fusion with one of the coat proteins, most commonly pIII.
 - The genomic DNA encoding for the coat proteins is enclosed within the protein coat.
- => Each protein remains connected to its encoding DNA

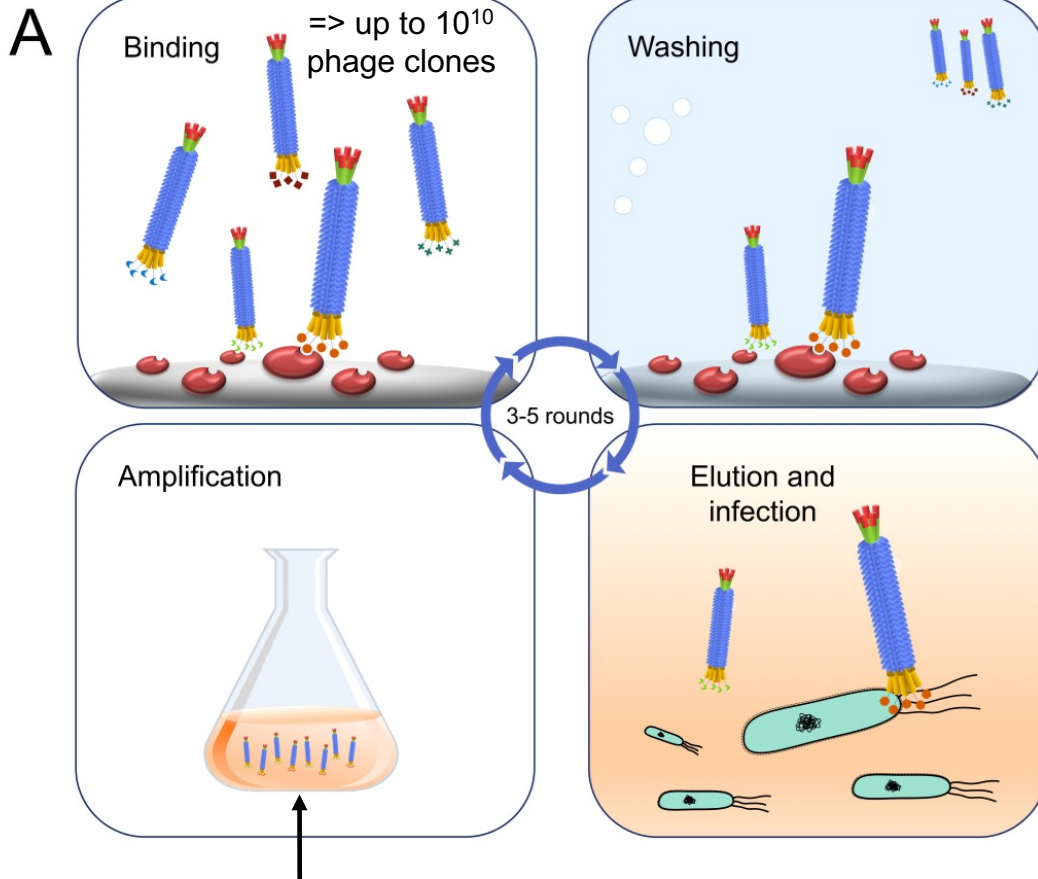
George Smith / Greg Winter:
Nobel prize in chemistry 2018

Construction of phage displayed protein libraries

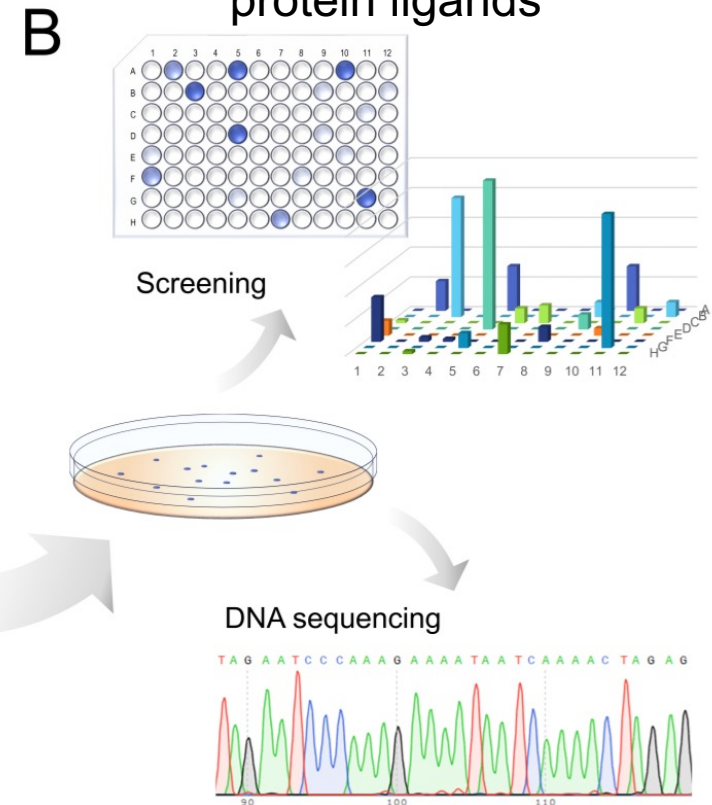


Protein engineering by *in vitro* evolution

Selection cycle

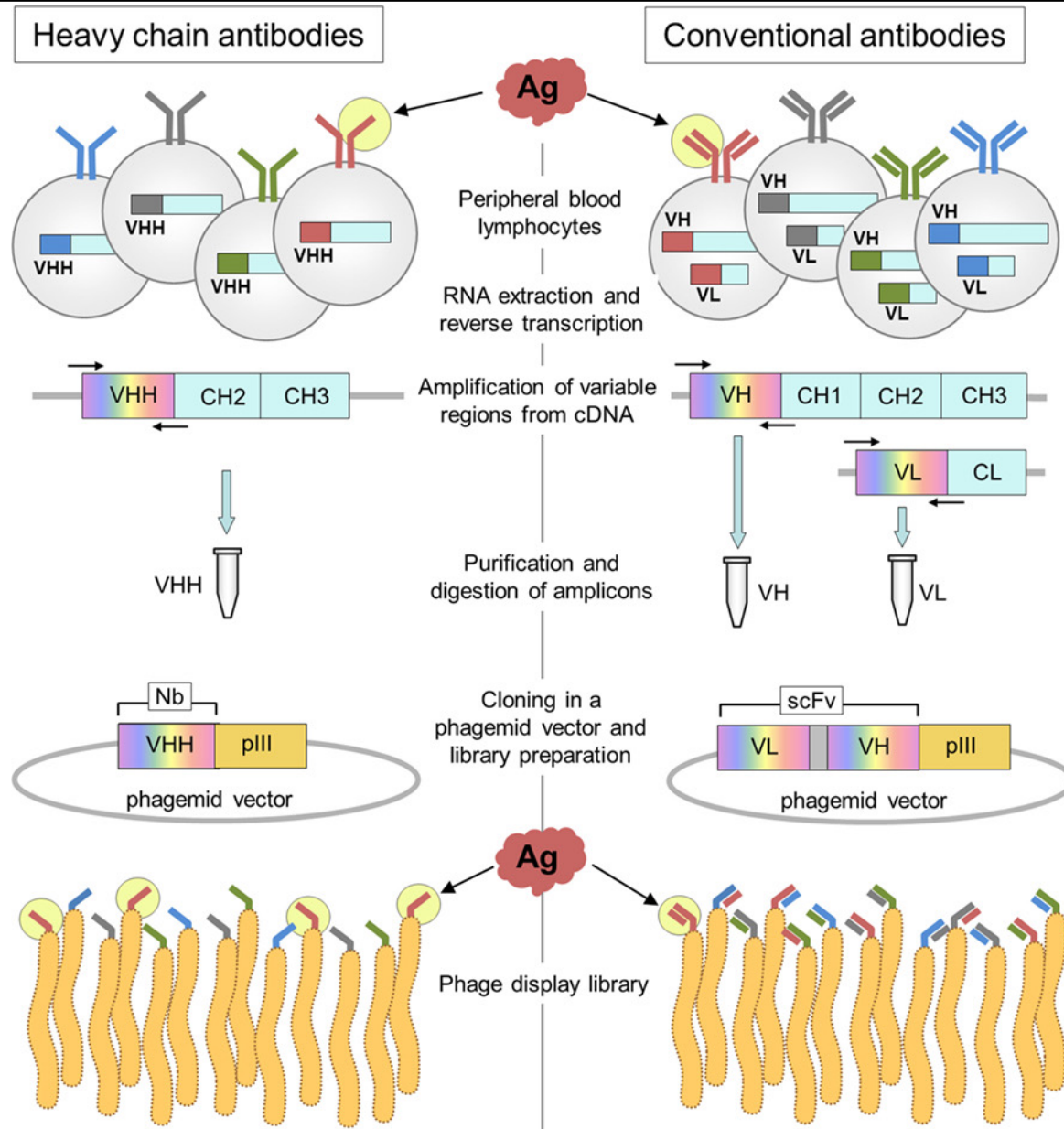


Identification of high affinity protein ligands



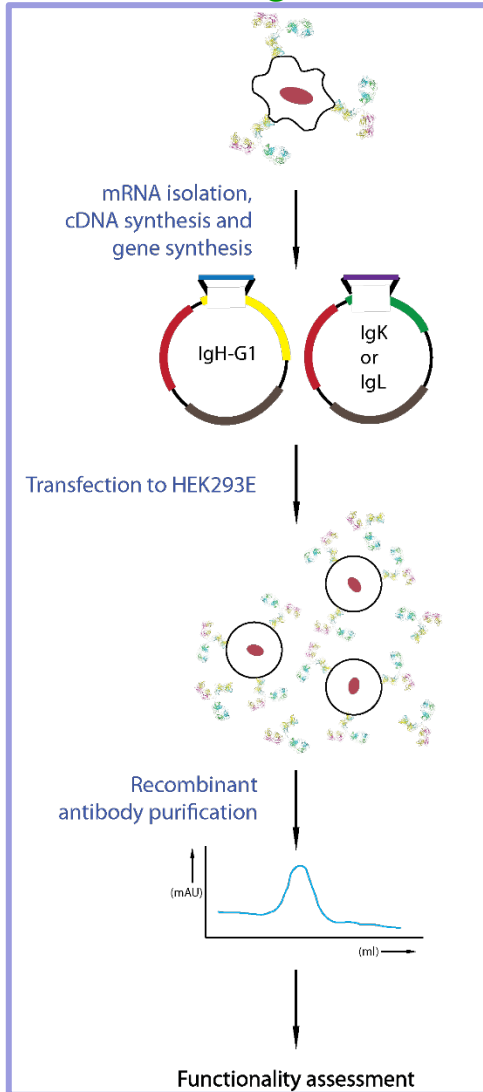
Option to introduce random mutations (e.g. error-prone PCR)

Single-domain antibody (nanobody)

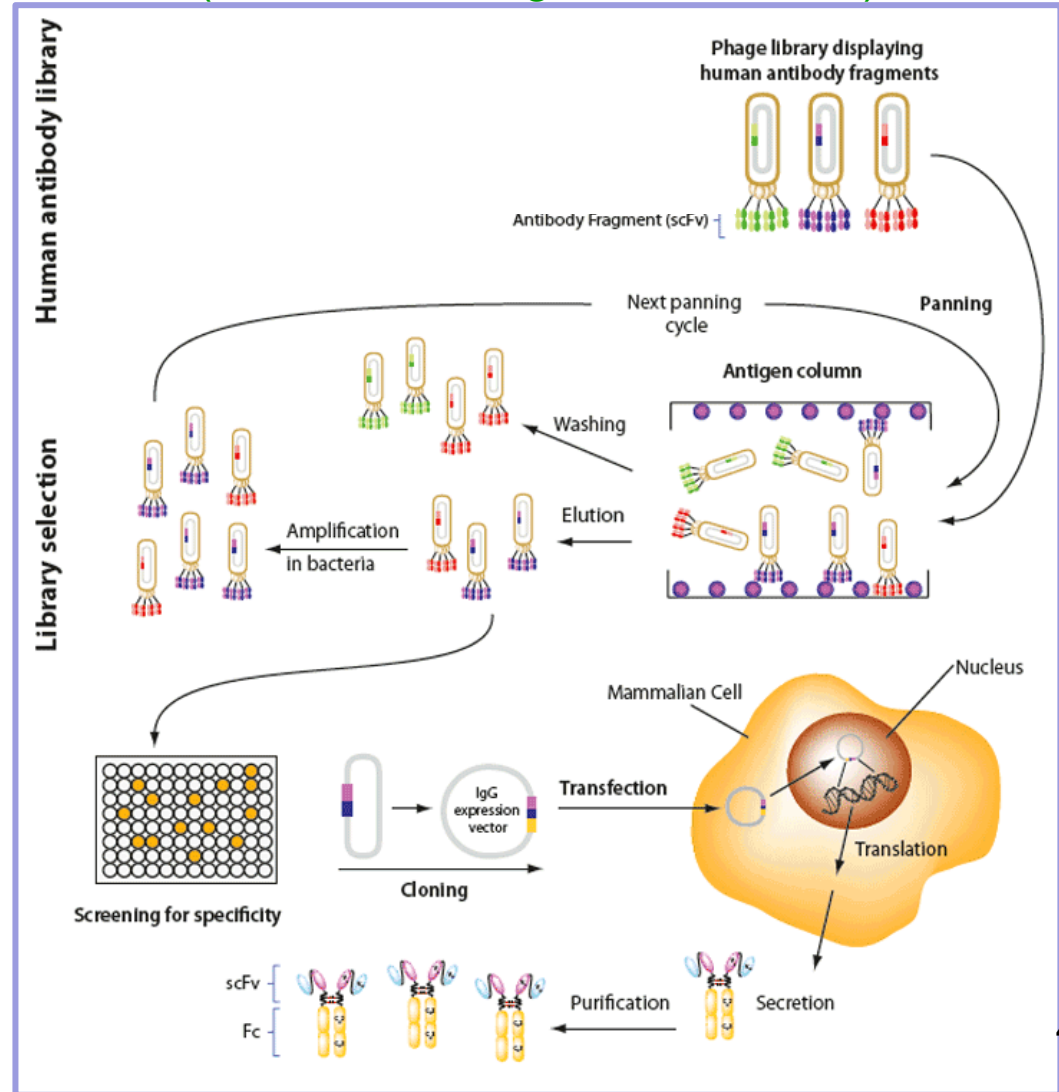


Production of recombinant antibodies

Expression system (Gene of Ab fragment known)

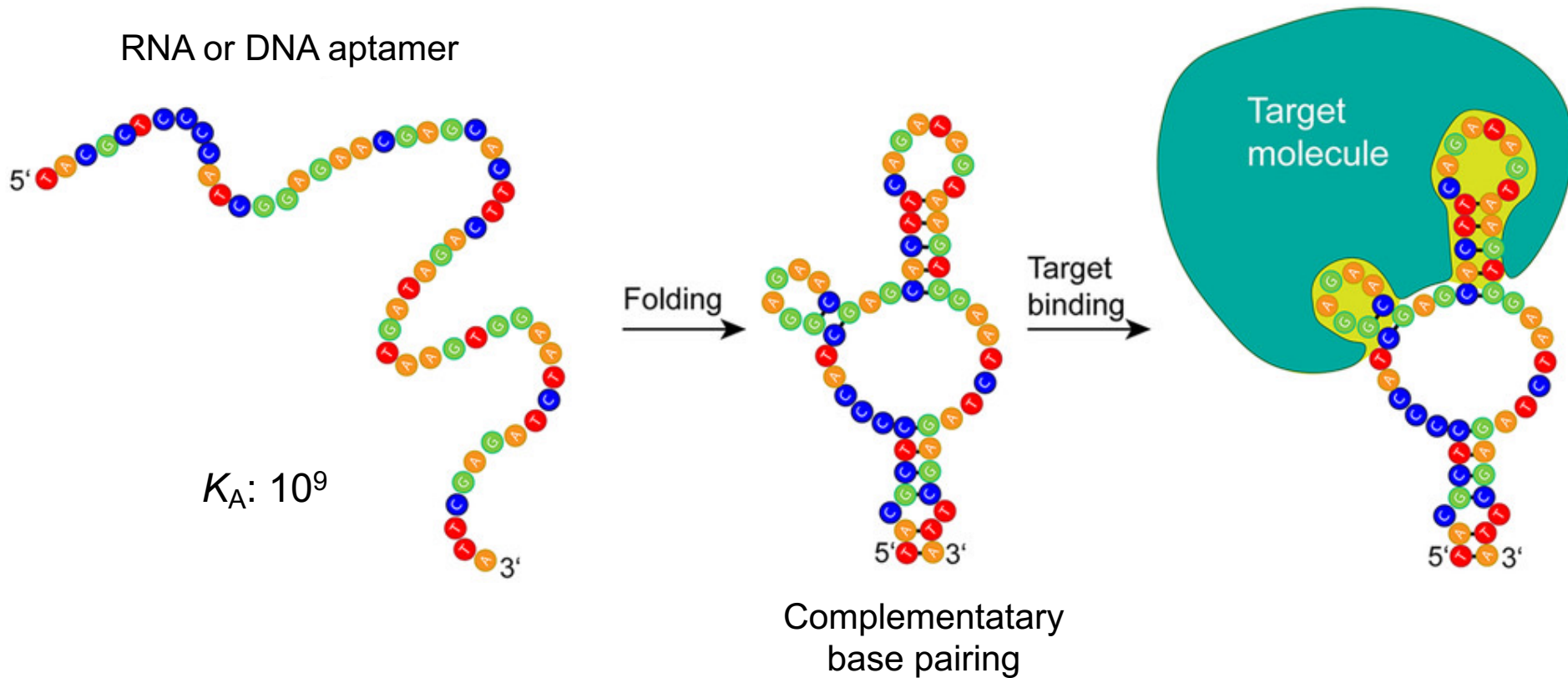


Phage display (Gene of Ab fragment unknown)



Alternatives for antibodies

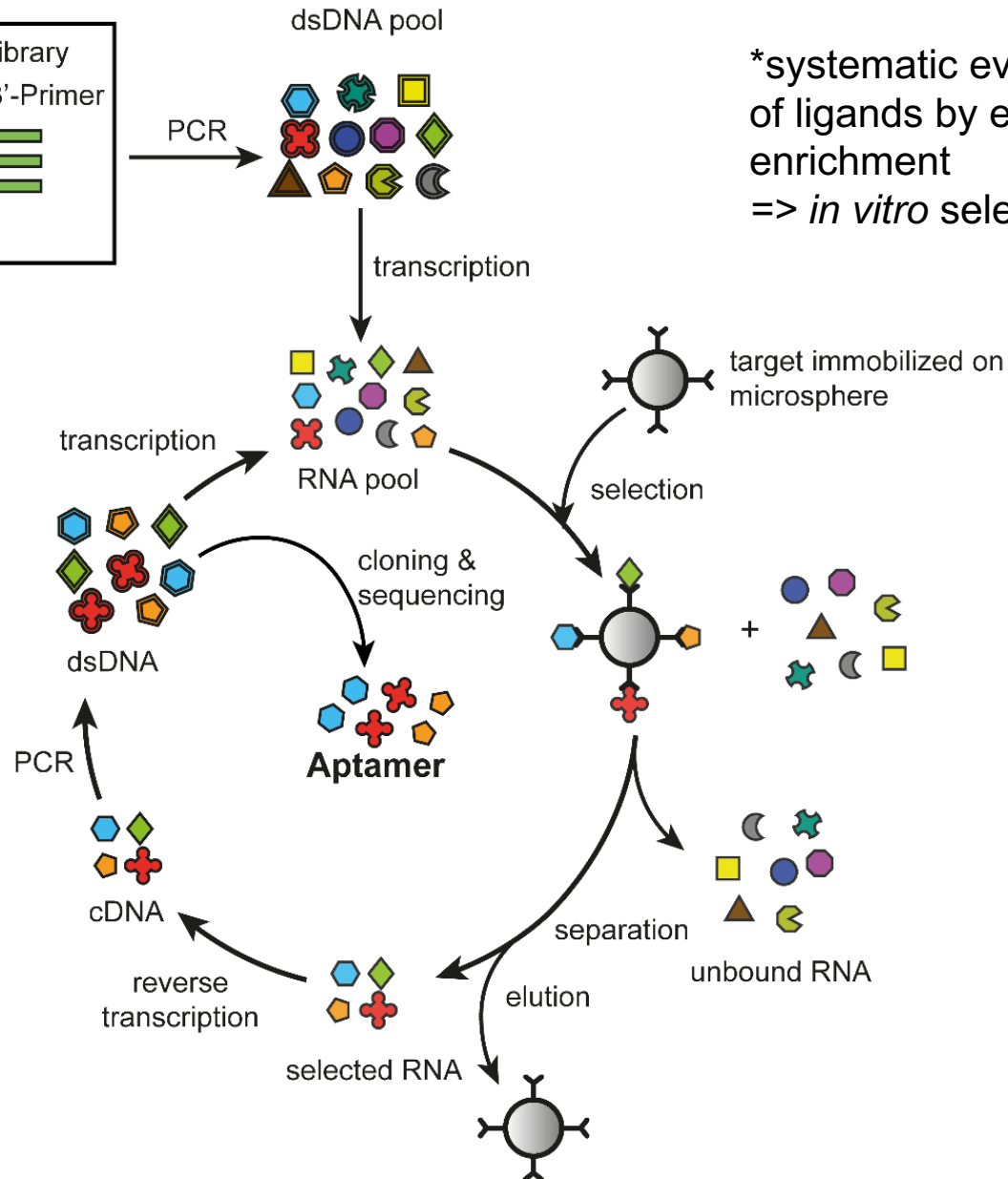
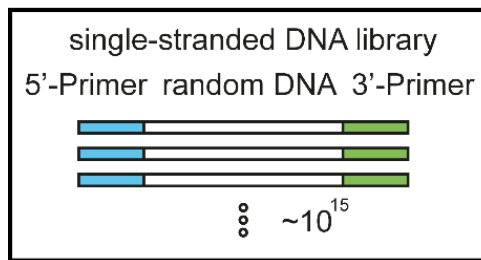
Aptamers



Binding through:

- (1) 3-dimensional, shape-dependent interactions
- (2) hydrophobic interactions, base-stacking, intercalation

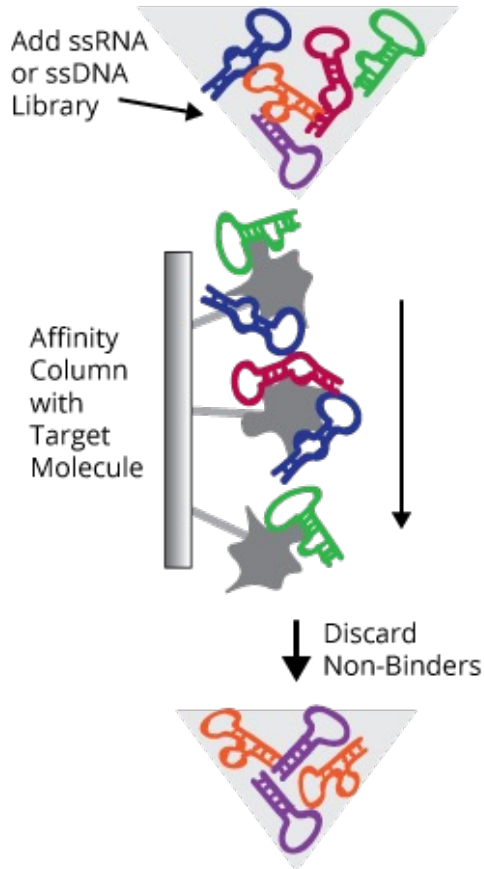
SELEX*



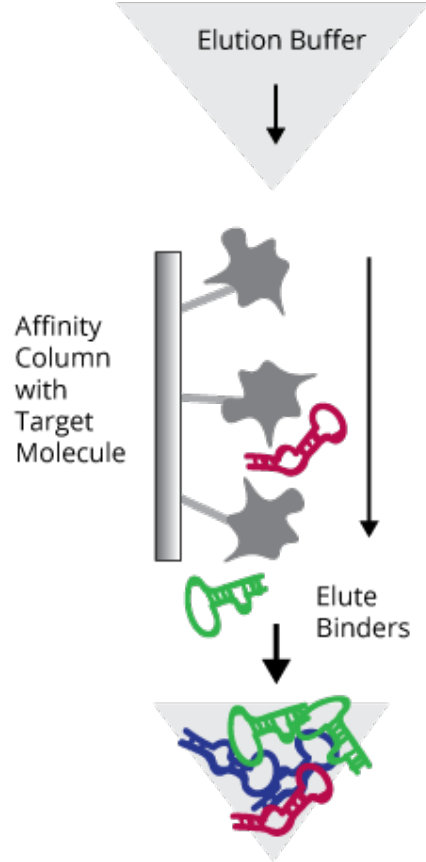
*systematic evolution
of ligands by exponential
enrichment
=> *in vitro* selection

SELEX*

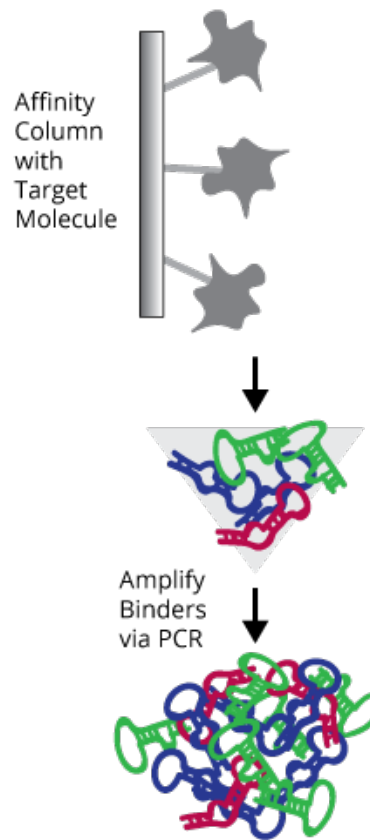
Step 1: Bind oligonucleotide library and discard non-binder



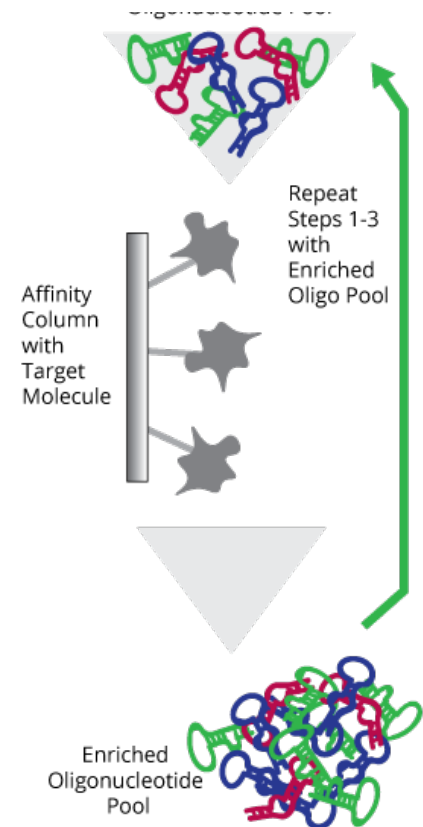
Step 2: Elute oligonucleotides that bind desired targets



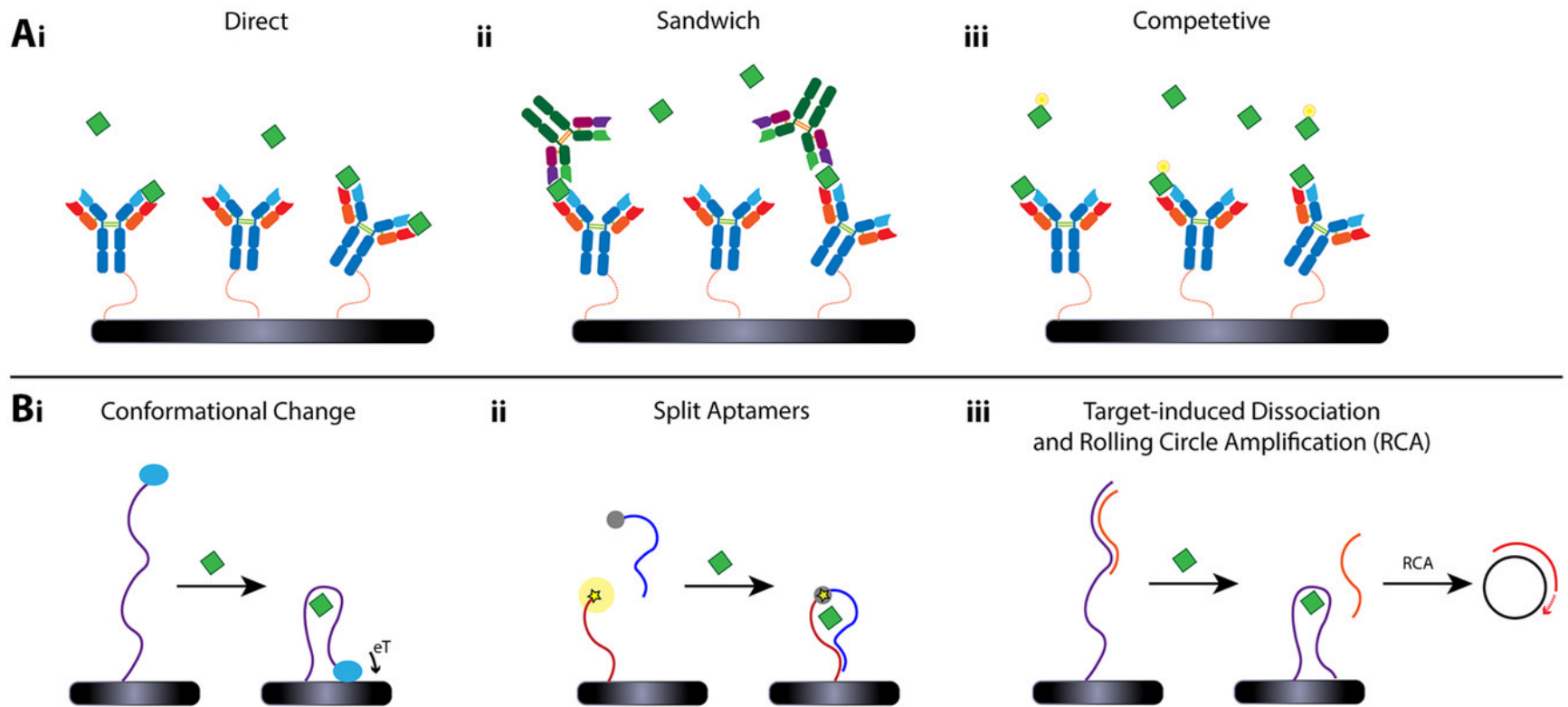
Step 3: Perform PCR to amplify eluted binders



Step 4: Repeat steps 1 through 3 using enriched oligonucleotide pool

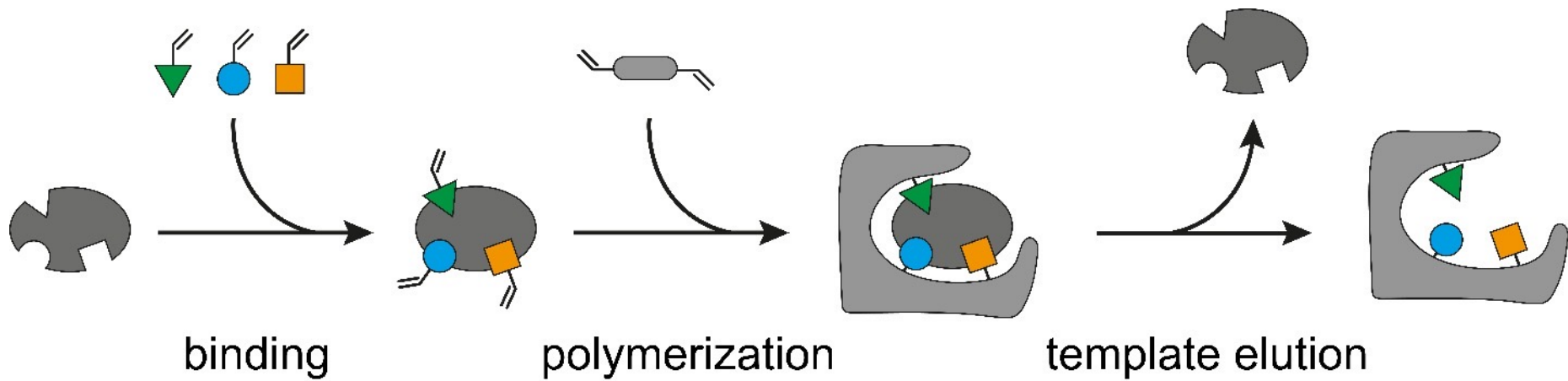


Aptamers: Assay designs



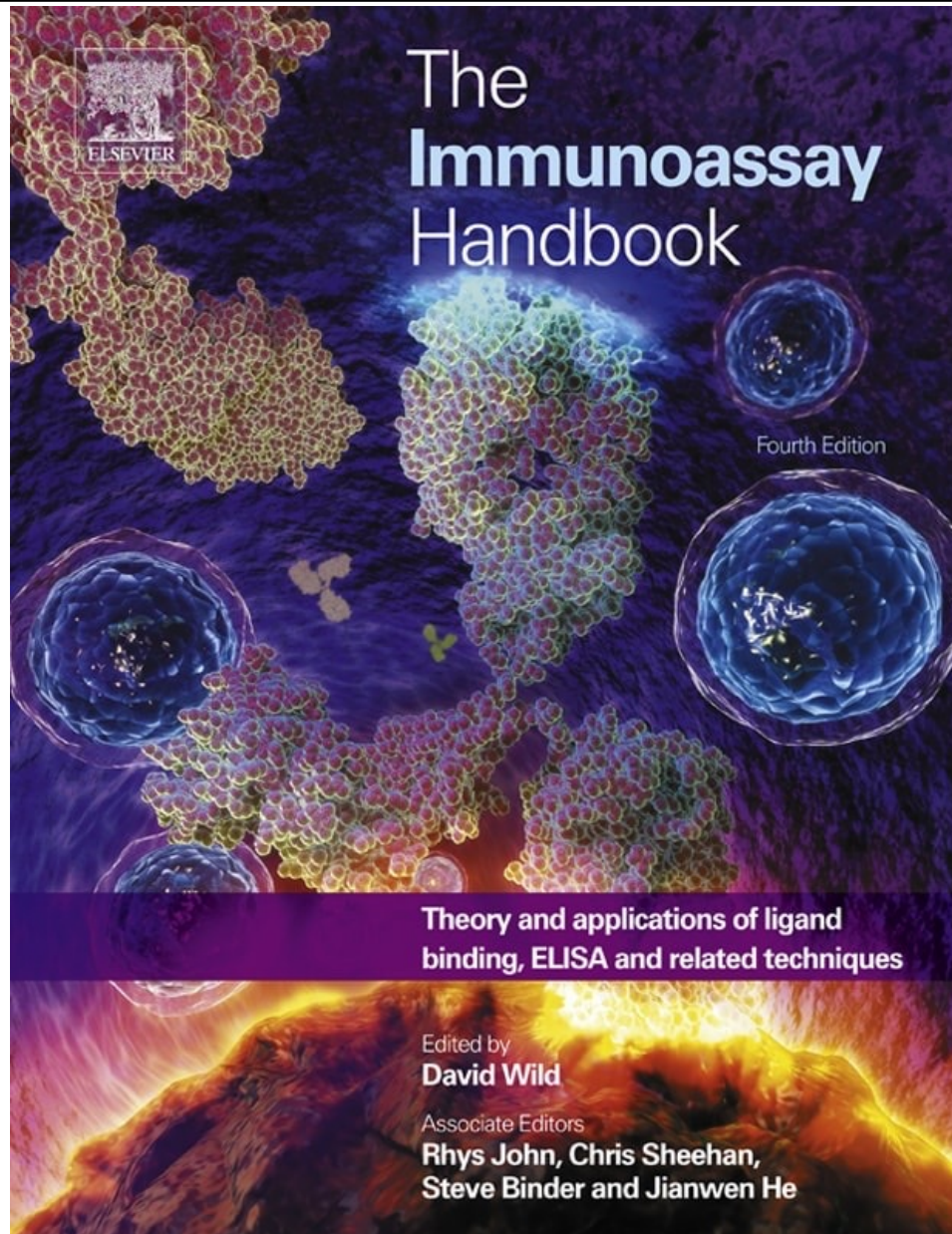
Molecularly imprinted polymer (MIP)

“Plastic antibodies”

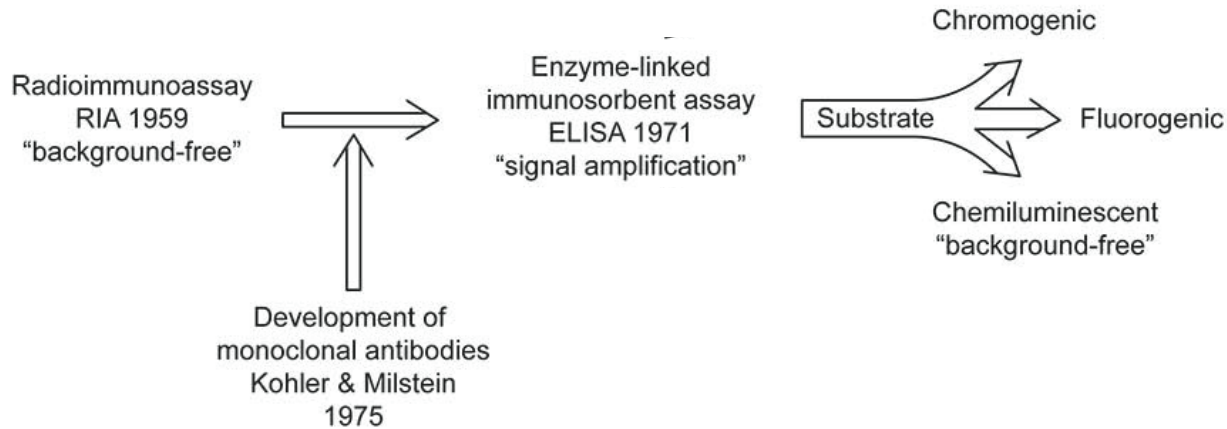


Immunoassays

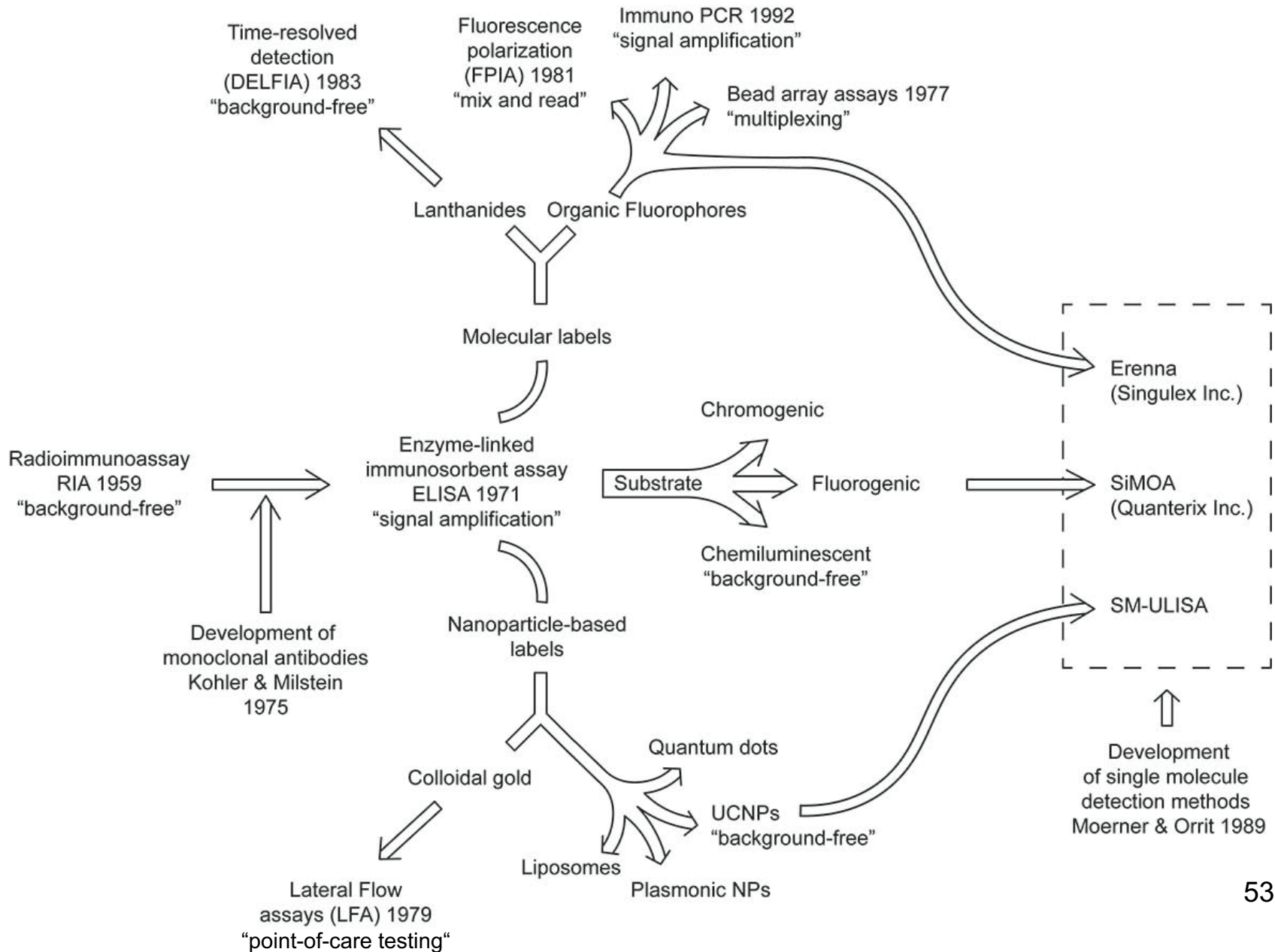
Literature for in-depth reading



History of immunoassays

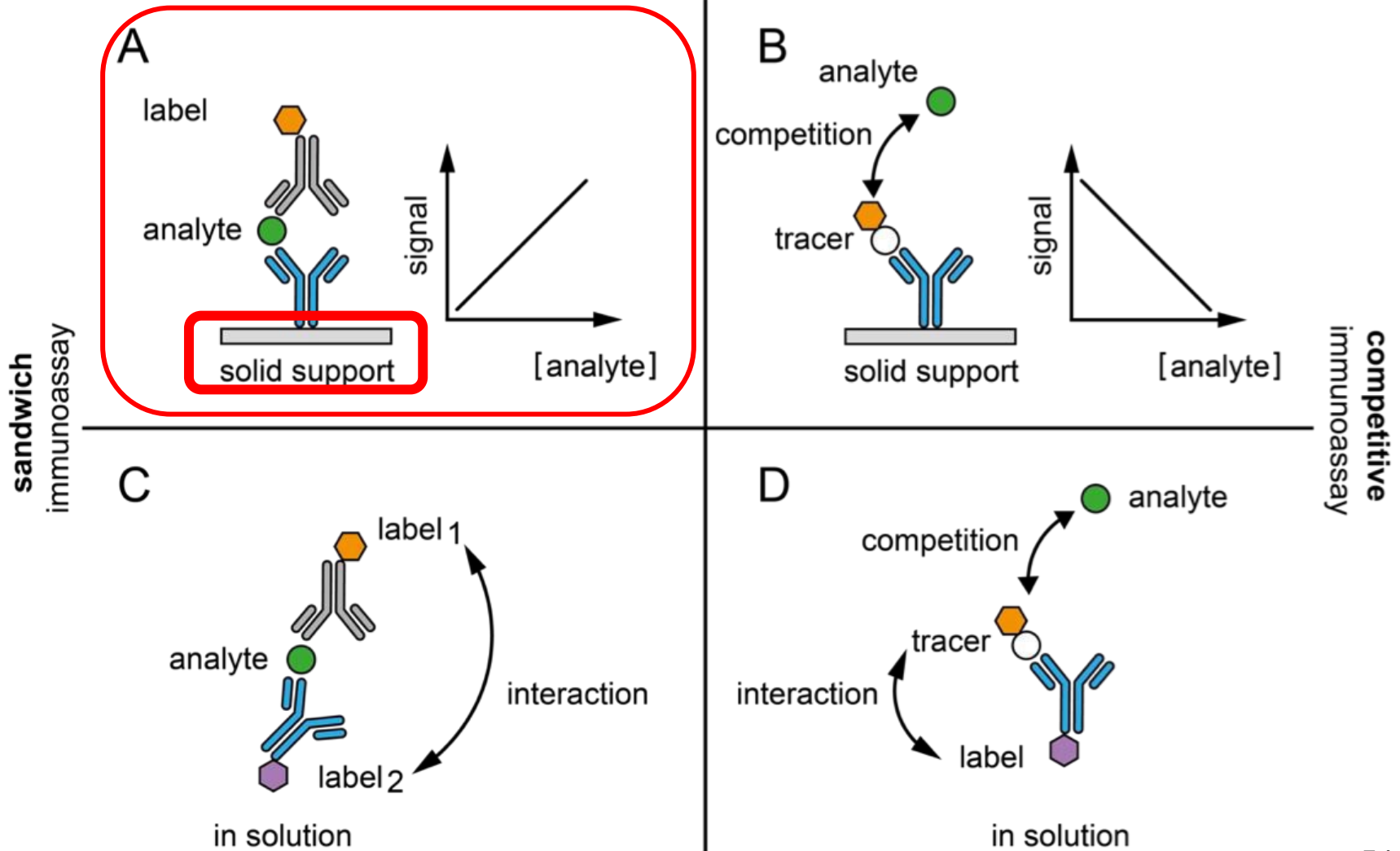


History of immunoassays



A rough categorization of immunoassays

HETEROGENEOUS



HOMOGENEOUS

Solid phase matrix

in heterogeneous non-competitive sandwich immunoassays

Performance-related issues:

- 1) low background in detection system
- 2) immobilization qualities:
 - high capacity
 - suitable and easy coupling chemistries
 - large surface
 - maintained reactivity of capture protein
 - no leakage
- 3) easy handling
- 4) inert in binding the labelled antibody/analyte => low background
- 5) effectively washed => low background
- 6) antibody excess through **high density** - surface measurement
- 7) antibody excess through **large surface** - integrating measurement

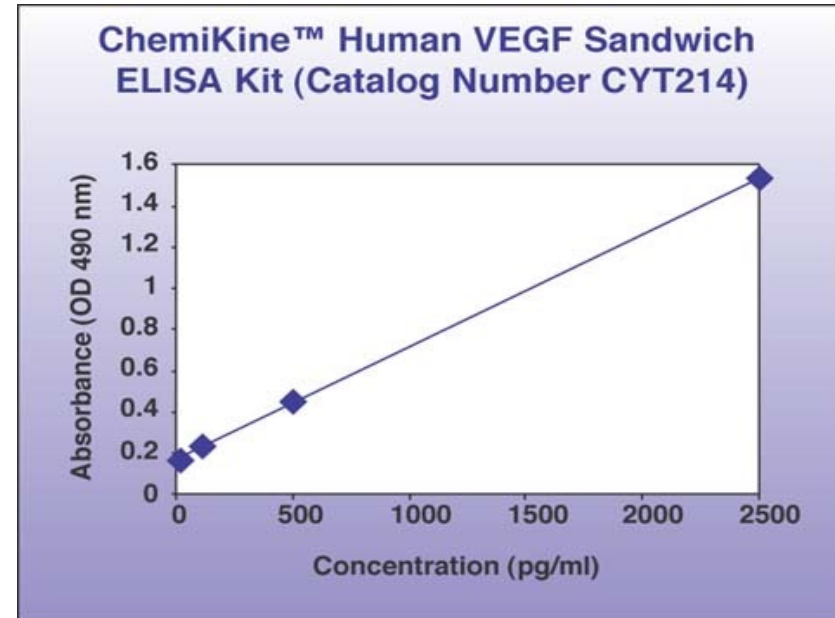
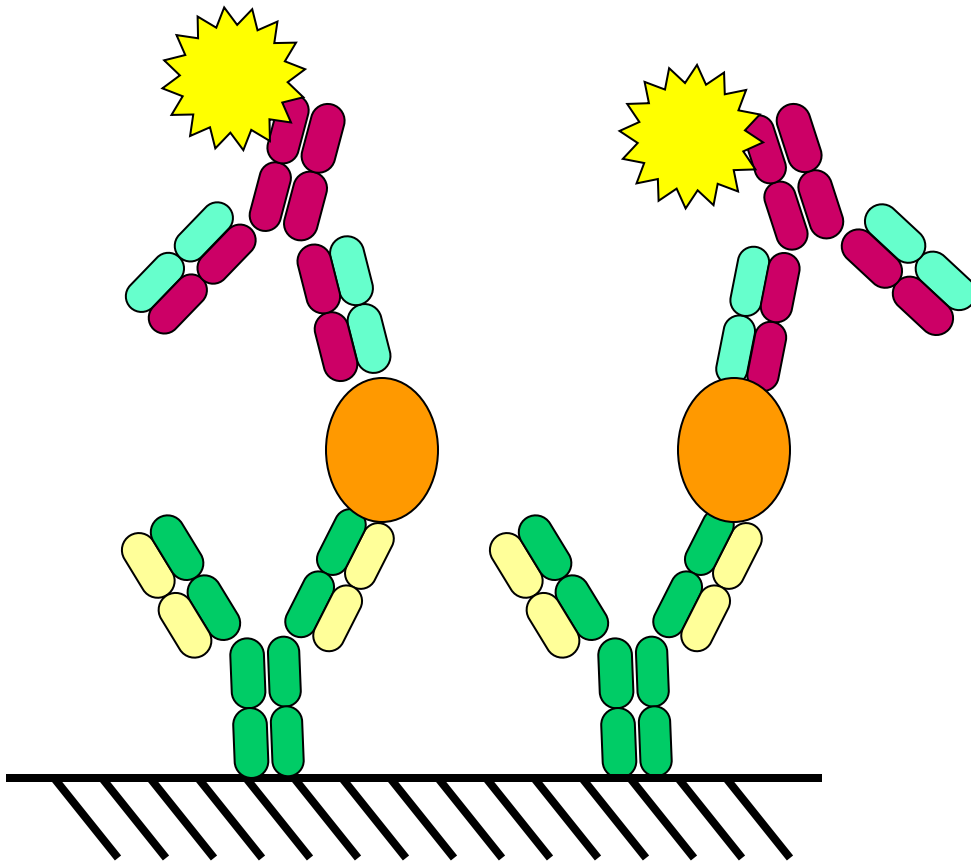
Solid phase matrices

Size	Examples	Advantages	Disadvantages
Small particle / "beads" (< 20 µm)	Latex Microcrystalline cellulose Fine porous glass Magnetic beads Liposomes Starburst™ dendrimers	Dispensing as for liquids Agitation not required High antibody binding capacity	Centrifugation required (unless used with a membrane capture) Long magnetic precipitation
Medium particle (< 1 mm)	Sepharose beads Sephacryl beads Sephadex beads	Centrifugation not required Short magnetic separation	Agitation required Slower binding kinetics than above Moderate antibody binding capacity
Small particle (< 20 µm)	Polystyrene Polystyrene-divinylbenzene	Centrifugation not required Agitation not required	Some variability in antibody coupling Lower antibody binding capacity Difficulty in dispensing Poor binding kinetics
Fibers	Membranes Glass fibers Nylon Silicon rubber	Centrifugation not required Agitation not required No dispensing of reagent Simple to use	Medium antibody binding capacity Can be fast binding kinetics
Solid surface	Coated tubes Dipsticks Microtiter plates (MTP)	Centrifugation not required Agitation rare No dispensing of reagent Simple to use	Variability in antibody coupling Lowest antibody binding capacity Slowest binding kinetics

Most frequently used solid phase matrices

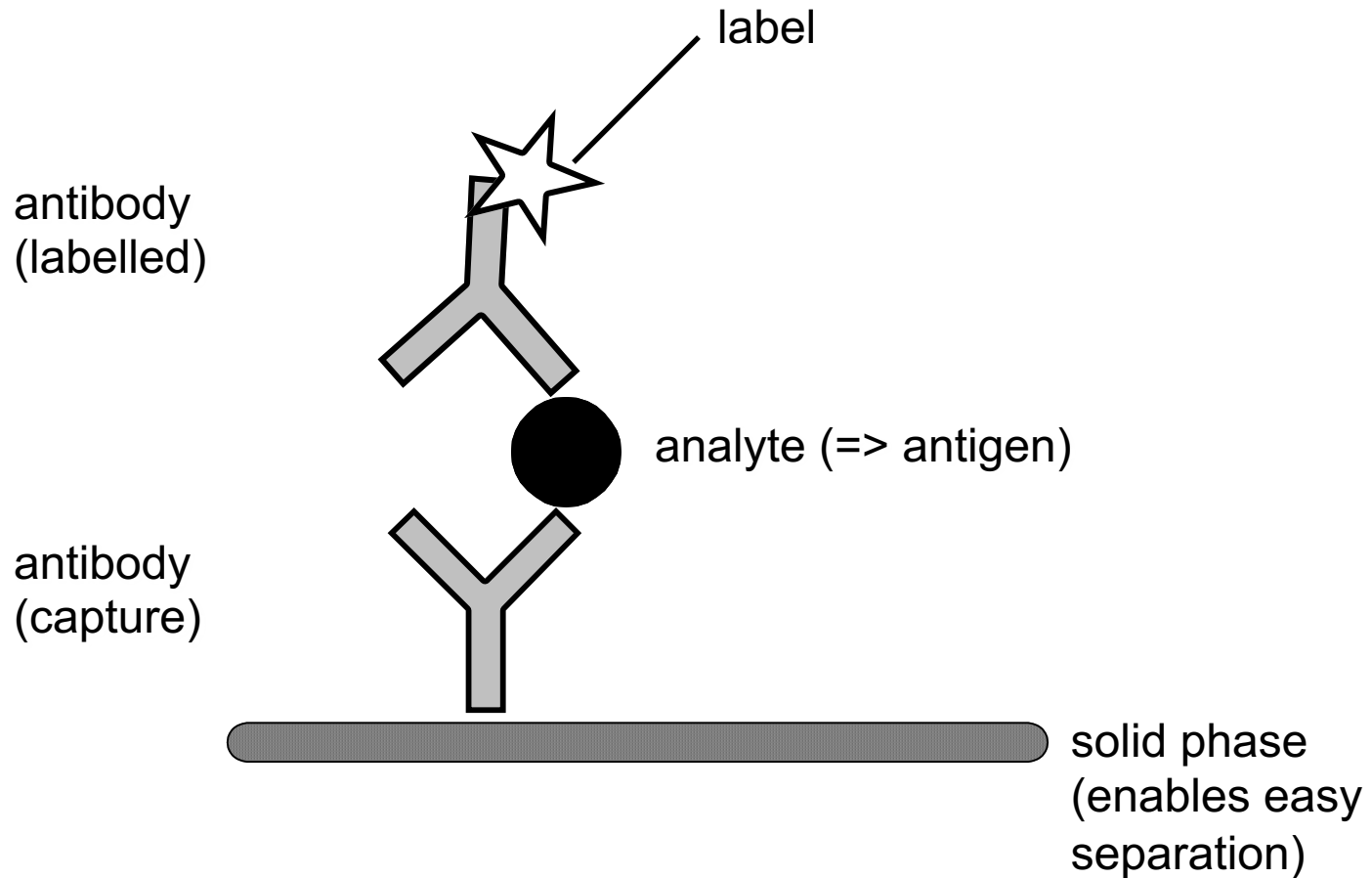
Non-competitive immunoassay

("sandwich" immunoassay)



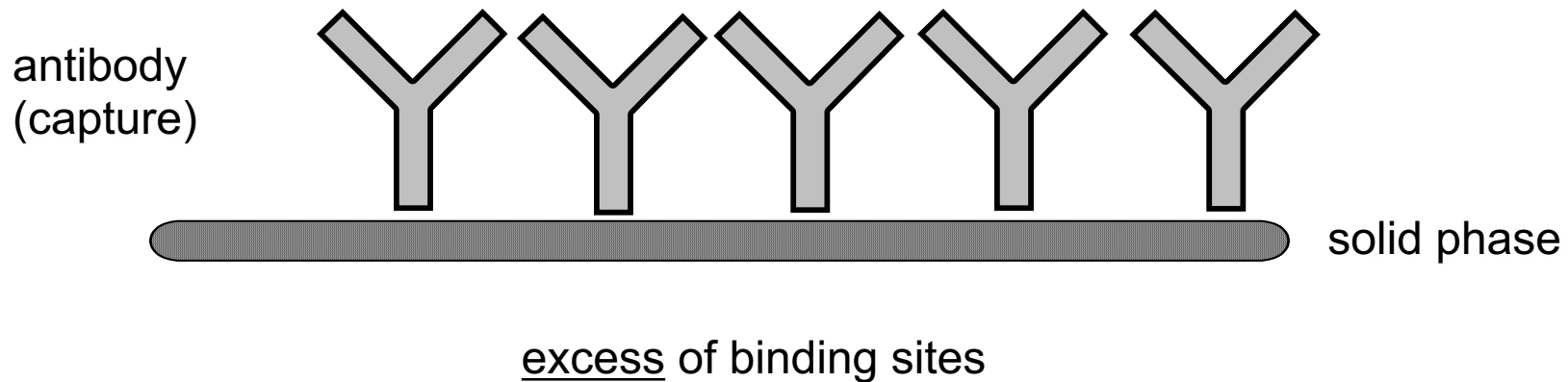
Non-competitive immunoassay

("sandwich" immunoassay)



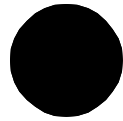
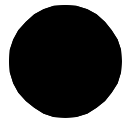
Non-competitive immunoassay

a capture antibody specific for a single epitope of the analyte is coated on a solid phase
(e.g. on a microtiter plate)
(=> monoclonal antibody preferred)



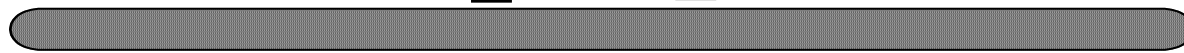
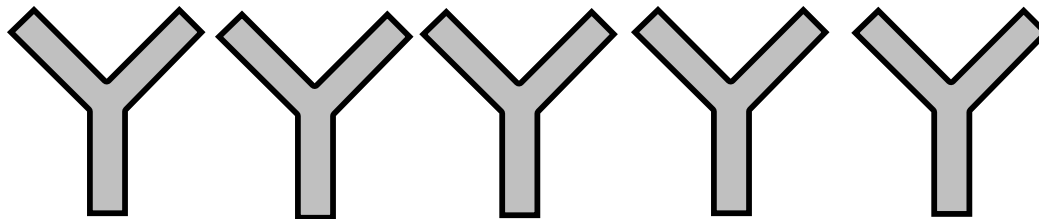
Non-competitive immunoassay

analyte



sample containing the analyte (at least two non-overlapping epitopes) is added; incubation for binding

antibody
(capture)

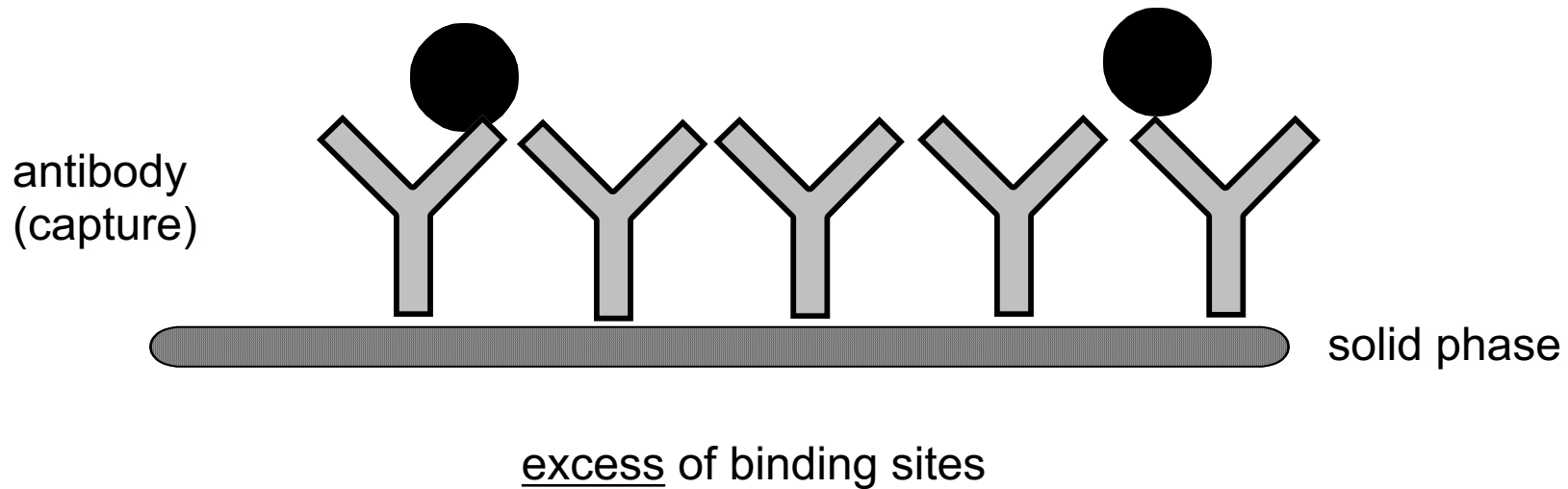


solid phase

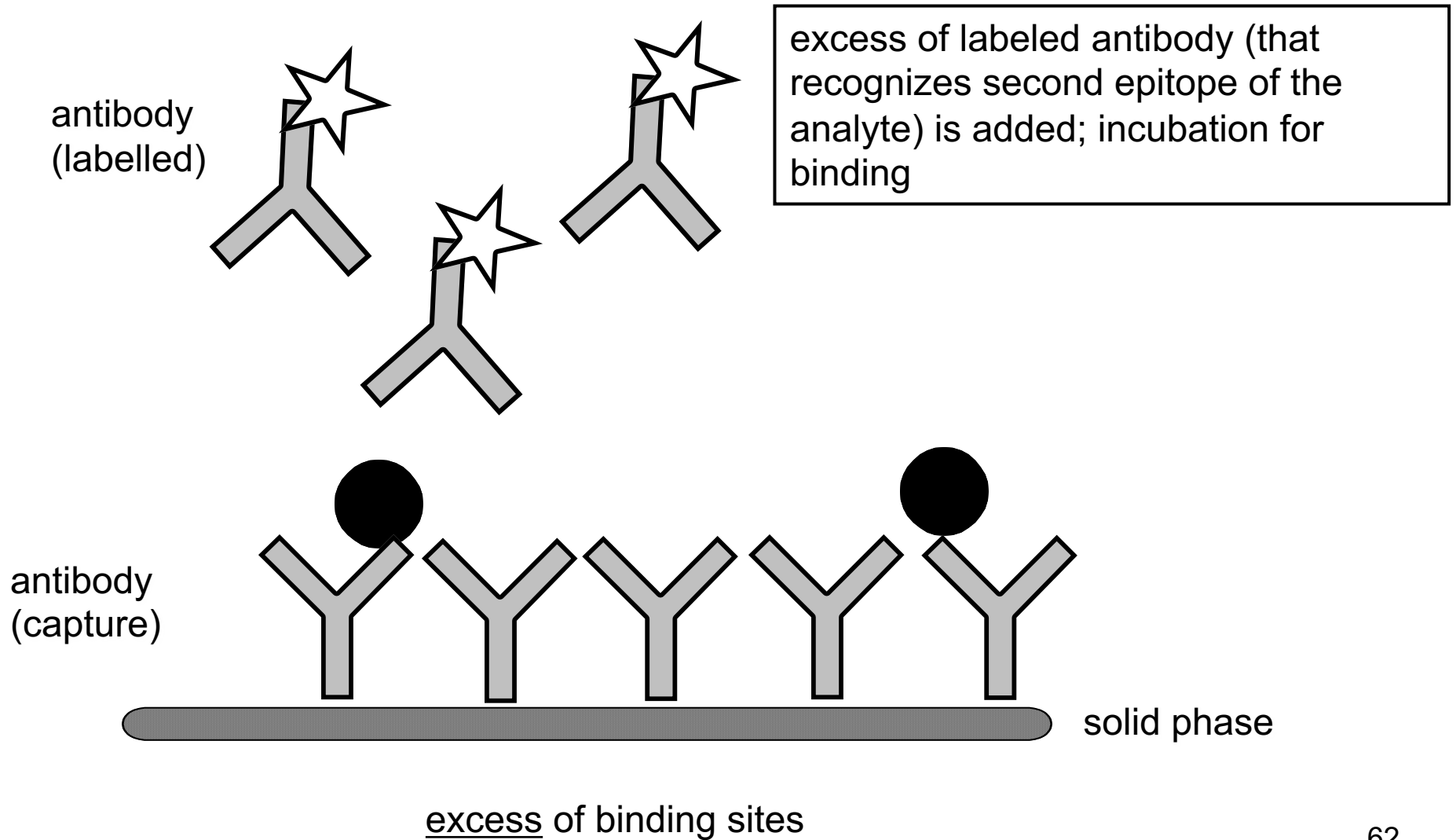
excess of binding sites

Non-competitive immunoassay

analyte is bound; in two-step assay:
sample is washed away with excess
of analyte

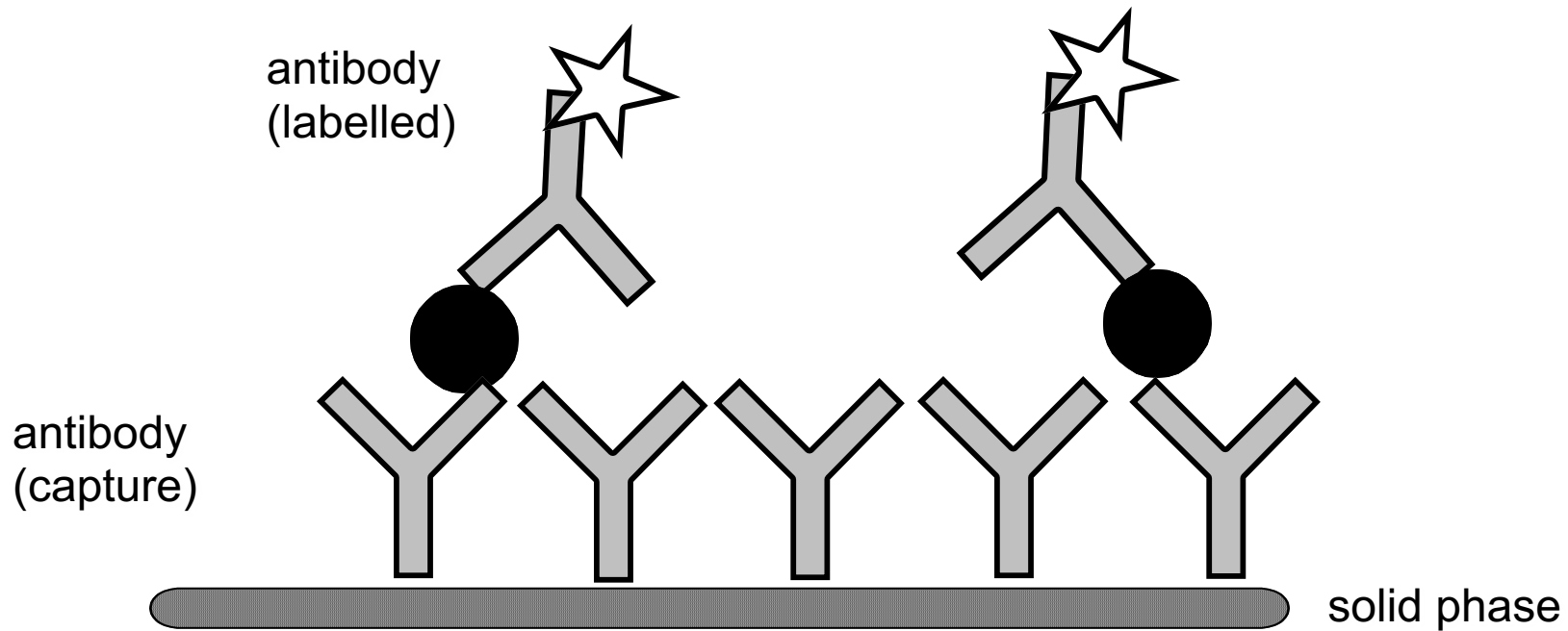


Non-competitive immunoassay

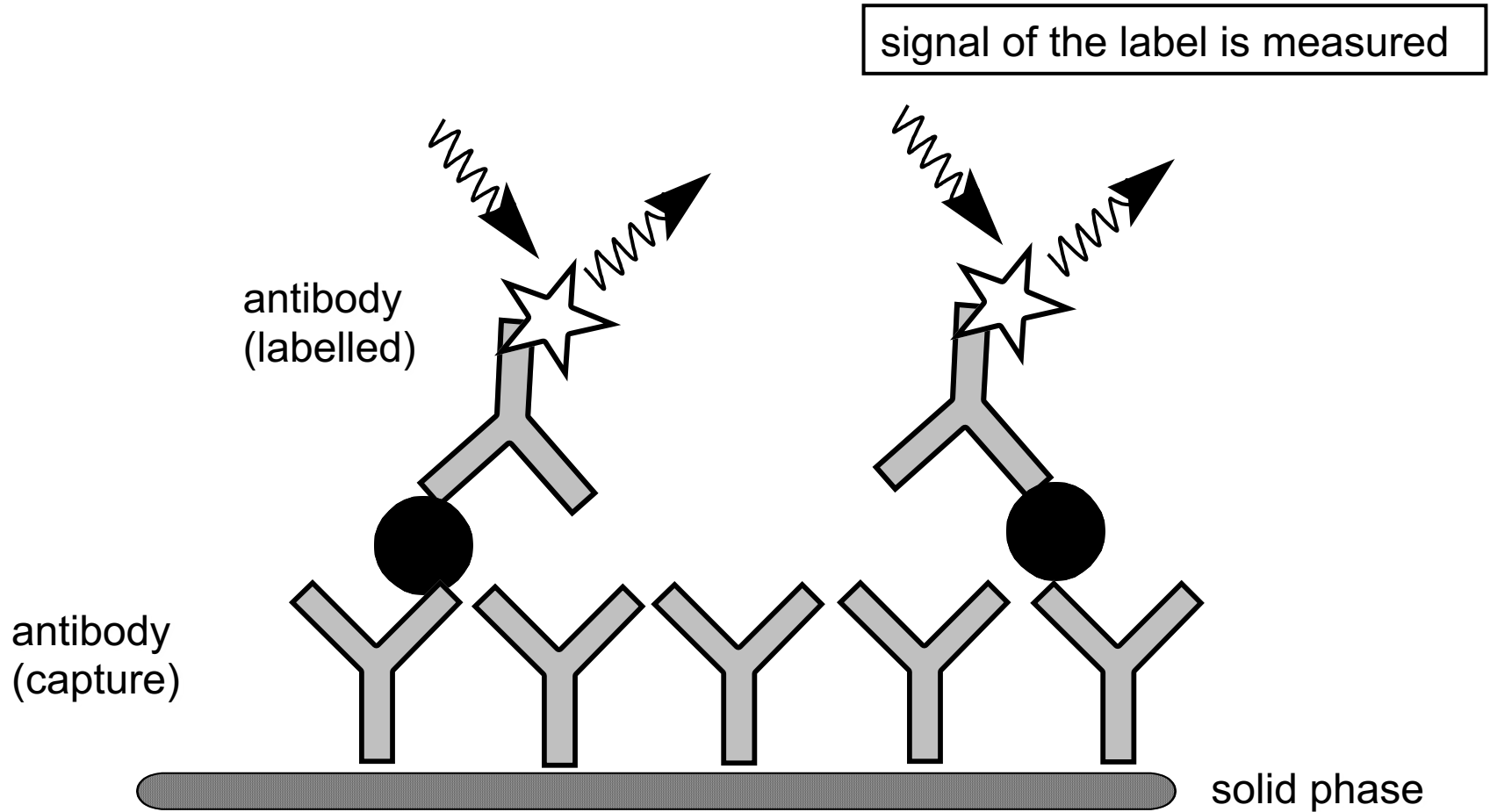


Non-competitive immunoassay

labeled antibody is bound;
excess is washed away

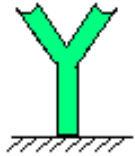


Non-competitive immunoassay



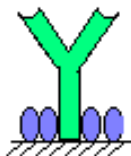
Enzyme-linked immunosorbent assay (ELISA)

Thorough **washing steps** required



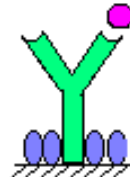
A. Coating anti-IgG

Non-covalent absorption of capture antibody to polystyrene surface (microtiter plate)



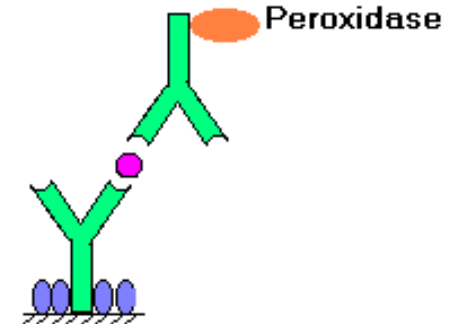
B. Blocking

Block surface with BSA or detergents to prevent non-specific binding of other proteins



C. React sample antigen

Add sample that contains the antigen (the analyte), e.g. tumor markers, viruses, or antibodies in serum.



D. React POx labelled secondary antibody

1. Add enzyme-labeled detection antibody (e.g. horseradish peroxidase); wash
2. Add chromogenic reagent (e.g. TMB)
3. Add "stop solution" (e.g. H_2SO_4)

Blocking is essential to avoid non-spec. binding

Normal serum

Normal serum (1-5% w/v) carries antibodies that bind to reactive sites and prevent non-specific binding of the secondary antibody. Serum is rich in albumin and other proteins that readily bind to non-specific protein binding sites of the sample.

Protein solutions

Blocking buffers often contain proteins such as bovine serum albumin (BSA), gelatin or nonfat dry milk (1-5% w/v). These inexpensive and readily available proteins are present in large excess compared to the antibody, so they compete with the latter for binding to nonspecific sites in the sample. Many labs developed homemade blocking buffers. It is important that blocking buffers are free of precipitates and other contaminants that can interfere with the detection.

Commercial buffers

Ready-made blocking buffers can contain highly purified single proteins or proprietary protein-free compounds. Many options are available that perform better than gelatin, casein or other proteins used alone, and they have improved shelf lives compared to homemade preparations.

Blocking tips

- Monitor both background (negative control) and signal strength (positive control) with various blocking reagents.
- Choose the blocking buffer that yields the highest signal-to-noise ratio.
- Ensure that there are no substances in the blocking buffer that interfere with a particular assay. Non-fat dry milk, for example, contains biotin and is inappropriate for use with any detection system that includes a biotin-binding protein.
- For optimal assay conditions, use the same blocking buffer for diluting the antibody that is used for the blocking step.