

Module Overview

Day	Lecture	Lab
1	Introduction	DNA library synthesis (PCR)
2	SELEX I: Building a Library	DNA library purification (agarose gel electrophoresis)
3	SELEX II: Selecting RNA with target functionality	RNA library synthesis (<i>In vitro</i> transcription = IVT)
4	SELEX III: Technical advances & problem-solving	RNA purification and heme affinity selection
5	Characterizing aptamers	RNA to DNA by RT-PCR
6	Introduction to porphyrins: chemistry & biology	Post-selection IVT Journal Club 1
7	Aptamer applications in biology & technology	Aptamer binding assay
8	Aptamers as therapeutics	Journal Club 2

Aptamer applications

20.109 Lecture 7
2 March, 2010

Today's Objectives

- Consider, through discussion, an increasingly important application area for aptamer technology
- Provide an overview of antibodies as affinity reagents to provide you with a context for deciding on using one over the other

Challenge

- As team leader at AptUs™ Biotechnologies (48-52 Mass Ave in Cambridge), you would like to develop a rapid and sensitive method for simultaneously detecting 50 proteins. These proteins have homologies ranging from 30-60% (i.e. they are highly similar to each other).
- The relative abundance of these proteins in human serum samples is correlated with diabetes risk.
- You are highly motivated to use your team's primary expertise to generate aptamers that can unambiguously distinguish these proteins.
- New Guy asks about using antibodies to tackle this problem. After the massive group laughter subsides, you promise to discuss this later.

Challenges

- *What are some key SELEX-related issues you'll need to address?*
- *What are your strategies for addressing these issues?*
- *How will you integrate your aptamers into a detection device?*
 - What will be your readout?
 - How will you calibrate your system?
 - Keep in mind that you will likely want to analyze hundreds of samples simultaneously.

Challenges

- *What are some key SELEX-related issues to address?*
 - Large number of proteins and selection optimization space to consider
 - High degree of similarity between the various targets => potential for cross-reactivity is high
 - Obtaining pure target proteins (all 50?!)

Challenges

- *What is your strategy for addressing these issues?*
 - Large number of proteins and selection optimization space to consider
 - Automation (options?)
 - High degree of similarity between the various targets => potential for cross-reactivity is high
 - Counter-selection (how might you efficiently do this?)
 - Perform SELEX on the isolated, non-homologous protein regions only
 - Advantages and disadvantages?
 - Obtaining pure target proteins (all 50?!)
 - You'll learn all about this in Module II!

Challenges

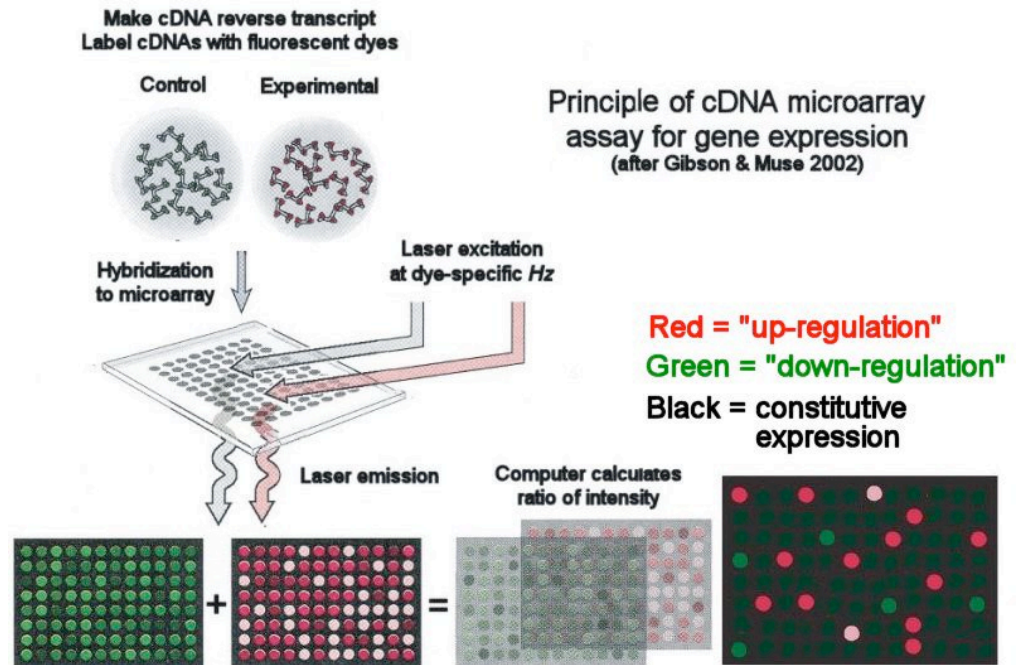
- *You've successfully obtained aptamers for 45/50 targets!*
 - How do you verify their respective specificities?
 - What cross-reactivities are you concerned about?
 - Between highly related classes of proteins
 - “Non-specific” interactions with other unrelated proteins
 - Especially high abundance proteins (e.g. albumin)

Challenges

- You now start thinking about how you'll use these aptamers to test the levels of these various proteins in blood samples.
 - *What test format do you choose?*
 - Solution phase
 - Immobilized (which component do you immobilize?)
 - How do you detect a binding reaction?
 - You expect to be doing high volume testing. Which formats are most compatible with having to run tests for 100 samples/day?

Microarray format

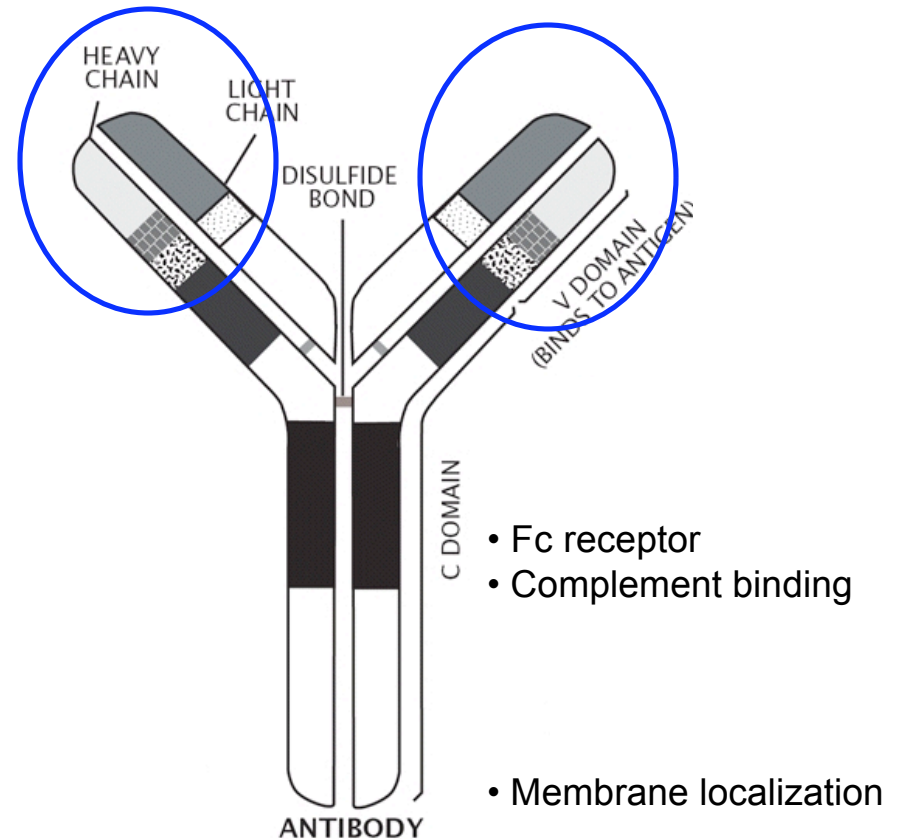
- Immobilized aptamer reagent
- Expose to fluorescently labeled sample
- Washing step to remove unbound material
- Image to quantify sample amount bound
- Same format as “gene chips” used to profile RNA



Antibodies: General

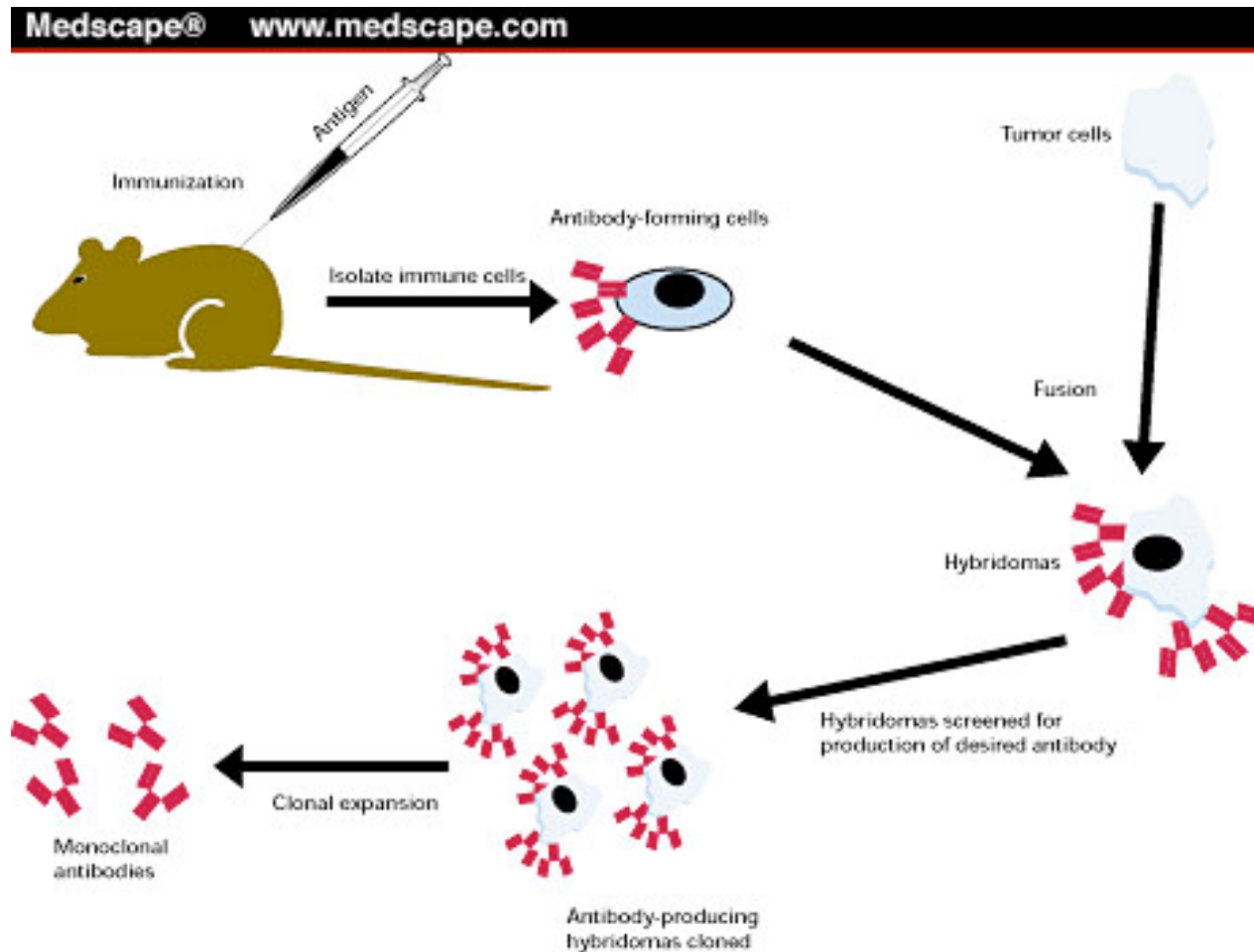
- Antibodies are proteins
 - Produced by B cells
 - Each B cell produces one type of antibody
- Overall structure
 - 2 heavy chains
 - 2 light chains
 - Both intra- and inter- chain disulfide bonds important for maintaining structure and function
- Functional Regions
 - Variable
 - Antigen binding
 - Both heavy & light chains contribute to binding site
 - Constant regions

Diversity: $\geq 10^{11}$



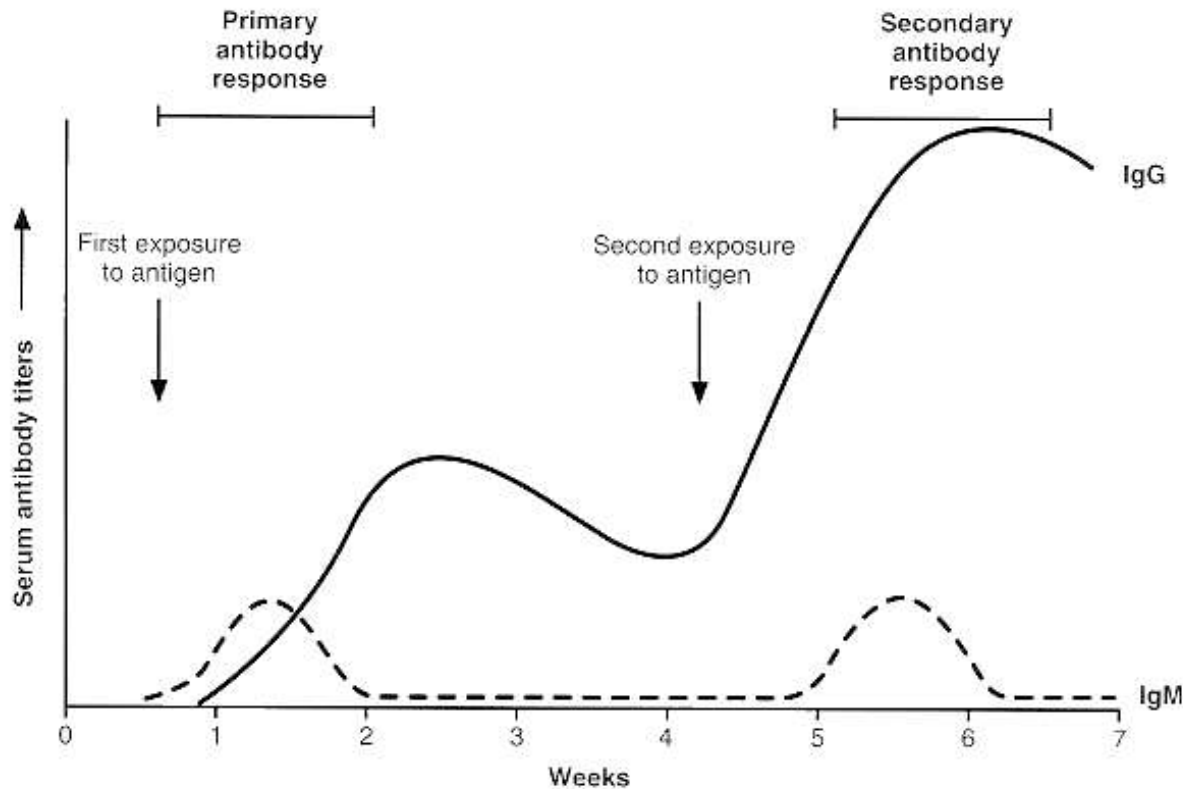
Antibodies

- How are antibodies produced?
 - Immunize an animal (e.g. mouse, goat, rabbit)
 - Several months later, antibodies can be isolated (if immunization successful)



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Typical time course for antibody production after immunization

Antibodies

- Are there any similarities between antibody and aptamer selections?
 - An animal makes large, randomized antibody pools (B cell clones)
 - Antibodies to the target antigen are **selected** from this pool
- Do you expect an antibody to every possible target will be present in an animal's antibody repertoire?
 - Consider representation within space considerations
 - Any other factors?
 - Yes--Antibodies that interact strongly with self-proteins are stringently selected against.
 - The B-cell clones capable of making these antibodies are eliminated from an animal's antibody repertoire

Antibodies

- What are some advantages to using or working with antibodies for binding reactions?
 - Lots of existing expertise!
 - Stability
 - Nature has optimized the selection process
 - Robustness of antibody production process
 - Potentially higher success rate of identifying an antibody
 - Selection stringency high ==> skewed towards identifying high affinity antibodies
 - Selection occurs (unattended) in immunized animal

Antibodies

- What are some **disadvantages** to using or working with antibodies for binding reactions?
 - Limited to targets that are not toxic to the animal!
 - Cannot easily tune the selection stringency
 - No guarantee that antibody will function in non-physiological conditions
 - Limited antibody reuse
 - Requires using animals (costly to house)
 - Batch-to-batch variability can be high

Aptamers versus antibodies

- Limited to targets that are not toxic to the animal!
- Aptamers to toxic substances can be developed

- Cannot easily tune the selection stringency
- Stringency easily tuned

- No guarantee that antibody will function in non-physiological conditions
- Aptamers can be selected under conditions in which they will be used

- Limited antibody reuse
- Aptamers can be refolded and reused

- Requires using animals (costly to house)
- No live animals required

- Batch-to-batch variability can be high
- Chemically well-defined aptamers can be reproducibly synthesized

Summary

- Aptamers are a viable strategy for binding and distinguishing closely related protein family members
- For multi-target, high throughput protein identification studies, there is a need to:
 - Increase the throughput for producing aptamers to distinct targets
 - Use aptamers in a format compatible with processing many samples in parallel
- Antibodies are the gold standard affinity reagents in biology/biotechnology
 - Many desirable characteristics
 - Aptamers can rival antibodies
 - Must carefully consider your applications to decide which affinity agent is more suitable