

NAME _____ TA _____ SEC _____

7.012 Problem Set 5 FRIDAY October 29, 2004

Due to popular demand, and a congenial staff due to the recent performance of the Red Sox, this Problem set upon completion must be inserted into the box

Problem sets will NOT be accepted late. Solutions will be posted on the web.

Question 1

Viruses have evolved several mechanisms to evade recognition and elimination by the immune system. Our immune systems have likewise evolved ways to counteract these methods of evasion.

- a) What type(s) of cells of the immune system recognize virally infected cells?

- b) How do these cells do so? Explain briefly.

- c) Besides viruses "hiding-out" from most of the immune system within cells, how might viruses further prohibit the previously mentioned immune cells from identifying them in host cells? ****Hint**, our immune system has **retaliated** to this evasion technique with special immune cells called natural killer cells (NK cells for short). These NK cells recognize and kill cells that have low or no MHC class I expression at their surface.

It turns out NK cell activity is a little more sophisticated. NK cells do not kill cells that have MHC I receptors on their surface. Cells without MHC I receptors on their surface are candidates for killing by NK cells. But NK cells also need to be activated for killing by specific receptors on the target cell's surface.

- d) Knowing more about how NK cells function, suggest a way in which the virus may now evade NK cell recognition and thus avoid its subsequent demise.

- e) In addition, all cells have mechanisms to protect themselves from double stranded RNA (dsRNA) viruses. What is this innate mechanism of protection? Briefly explain how it works.

Question 2

As a UROP in the Weinberg lab, one of your duties is to take care of the mice used for experimentation. During your tenure in the mouse facilities you notice a colony of mice that is highly susceptible to bacterial infections. In order to figure out why, you first take a blood sample and look at the levels of serum proteins. You are shocked to find that these mice lack antibodies in their serum! You are now aware of why they are so susceptible to infection.

a) What cell type do you suspect is affected by this mutation?

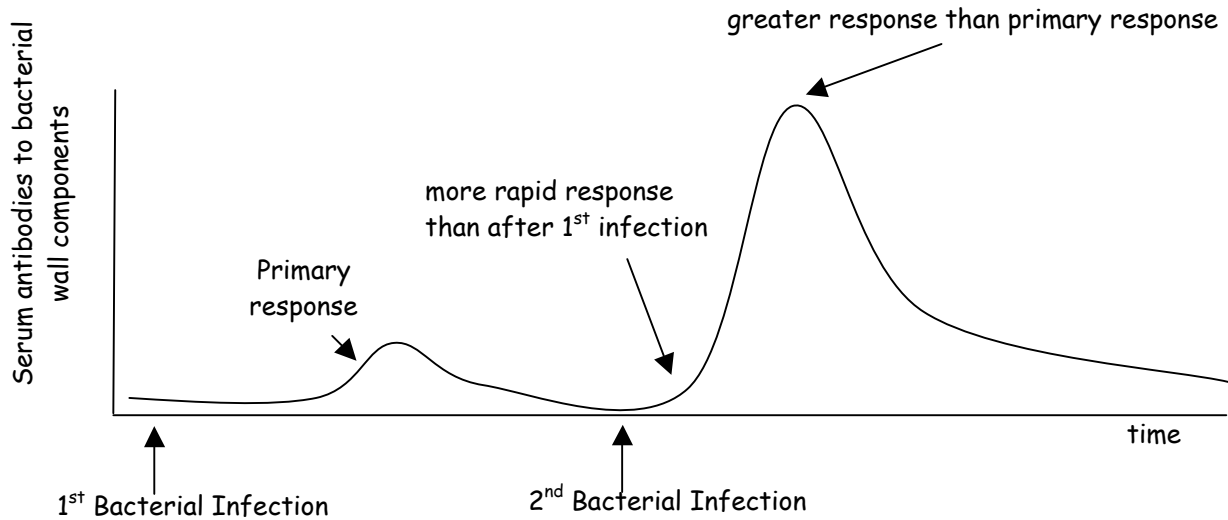
With what little spare time you have from your lab duties and school work, you decide to try to figure out why there are no antibodies circulating in these mice. You find that the mutation that leads to the disease is within the immunoglobulin heavy chain locus. Sequencing the locus reveals that the mutation is a deletion of the C-terminal transmembrane domain of the heavy chain. You're not an immunologist and frankly you were kind of surprised to learn that the antibody heavy chain has a transmembrane domain, after all you always have been taught simply that antibodies are secreted. However, based on your knowledge of cell signaling that you learned in 7.012, you begin to think of the antibody as a receptor and thus can make sense of why it has a transmembrane domain.

b) How does the transmembrane domain of the antibody enable the cell to signal from an "antibody receptor" to the nucleus? (Answer by comparing the alternative possibility in which the antibody is just secreted.)

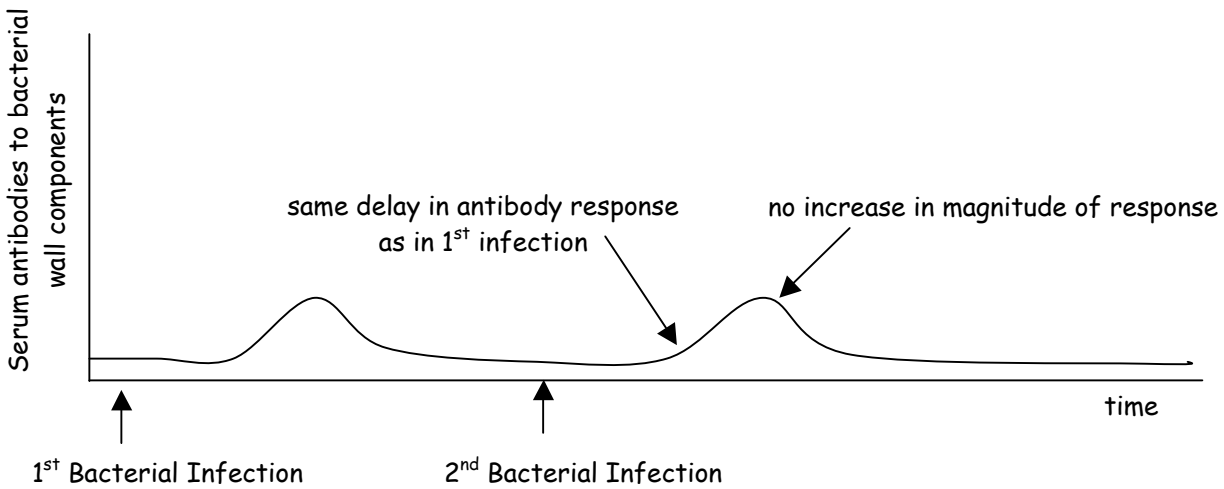
c) Why might this signalling be important for antibody selection prior to secretion from the cell?

d) Now hypothesize why these mutant mice lack antibodies in their circulation?

Question 3



The diagram above depicts a typical antibody response to a primary and secondary bacterial infection in wildtype mice. Being that it is the week following Halloween and you have an abundance of candy, you decide to feed your mice some of the “more suspect” candy that you got from some creepy old ladies whom you think may have tricked rather than treated. Indeed, after putting the mice on the candy diet you find that their antibody response has changed! It now looks like this...



You postulate that the candy must contain some kind of chemical or agent that is causing this phenomenon. What specific cell does this hypothetical chemical most likely target? Be very specific and explain your reasoning.

Question 4

You are isolating new viruses, one of which you set aside to study further. You measure the amounts of each nucleotide in this virus and determine the following...

A=20%, G=30%, T=0%, C=30% and U=20%

a) What does this suggest about this viral genome?

You find a very interesting virus that can infect *E. coli* that you name *Soxrule* bacteriophage. You measure the amounts of each nucleotide in the *Soxrule* viral genome and determine the following...

A=18%, G=32%, T=18%, C=32% and U=0%

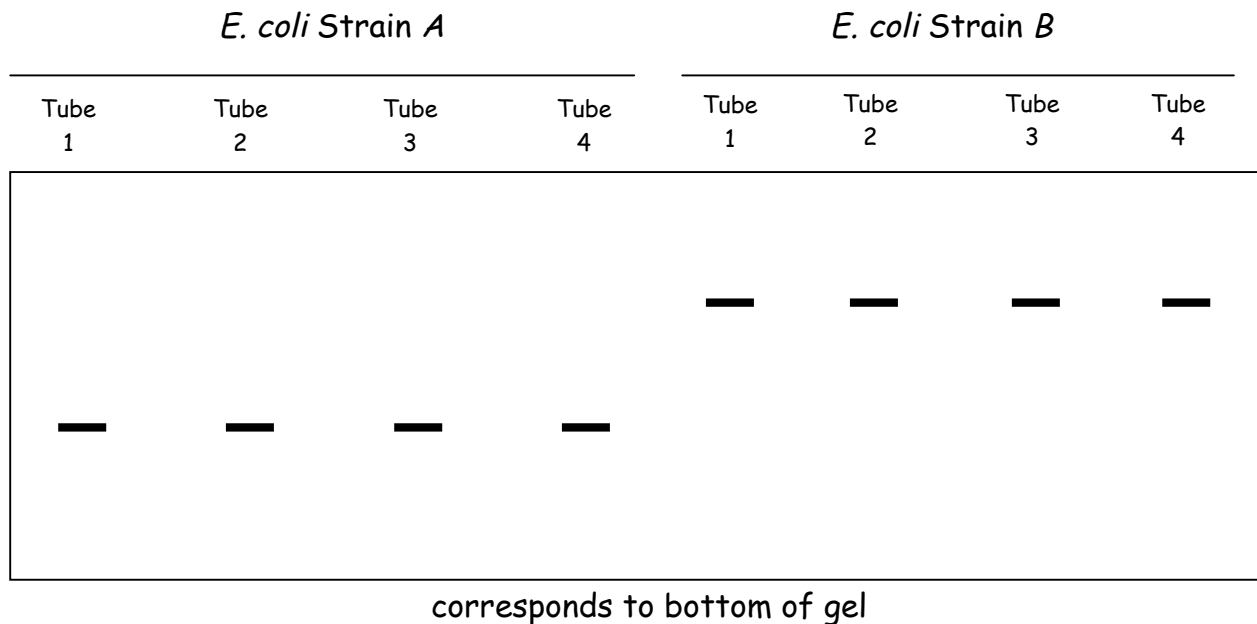
b) What does this suggest about this *Soxrule* genome?

c) After infecting *E. coli* with the *Soxrule* virus, you don't see any lysis until after many rounds of cell division. What is going on and why is it advantageous for the virus?

You perform the following experiment:

- 1) Dilute a culture of *E. coli* strain A into four test tubes.
- 2) Dilute a culture of *E. coli* strain B into four test tubes.
- 3) Infect all eight test tubes having the two different bacterial strains with the *Soxrule* bacteriophage and incubate them for several generation times.
- 4) Wash the cells in each tube to remove unattached bacteriophage particles.
- 5) Purify the **chromosomal** DNA from the cells in each tube and cut with *Cla* I restriction enzyme. (You've already determined that the *Soxrule* genome has no *Cla* I restriction sites.)
- 6) Load the chromosomal DNA digests onto an agarose gel and perform electrophoresis.
- 7) Transfer the separated DNA fragments from the gel to a nitrocellulose filter.
- 8) Probe the filter with radioactively labeled viral DNA.
- 9) Place the filter on X-ray film and develop after overnight exposure.

You observe the following autoradiogram that you label with corresponding information from how you loaded the gel:



d) Does the *Soxrule* genome integrate into the bacterial genome?

In strain A? yes / no

In strain B? yes / no

e) If the *Soxrule* genome integrates into the bacterial chromosome, does the integration occur randomly or at specific site?

In strain A? random / specific

In strain B? random / specific

f) Explain the differences in the data derived from the two strains that is highlighted in the autoradiogram above.

Question 5

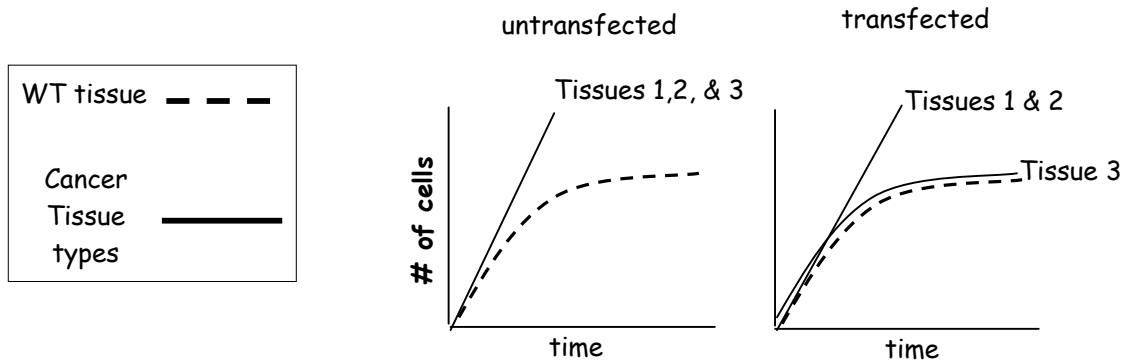
You work in a lab that studies cancer. Your lab receives a package from a hospital containing a letter that says they sent...

- breast tissue (from a patient with breast cancer)
- bladder tissue with a known p21^{RAS} mutation (from a patient with bladder cancer)
- eye tissue (from a patient with retinoblastoma)

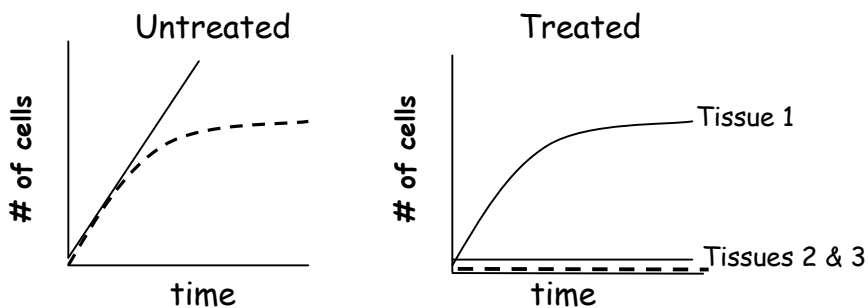
Unfortunately the labels fell off the tubes, and you take on the task of determining which sample is which.

In order to distinguish between the tissues, you name them 1, 2, and 3, and you perform the following experiments:

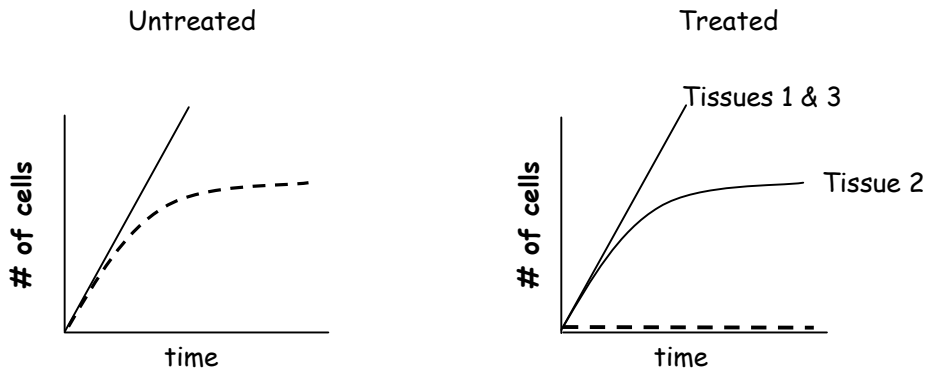
- Transfect a wild-type copy of the Rb gene into each tissue type and wild-type tissue as a control. You measure the number of cells in a tissue culture plate over time.



- Because you know that Cyclin D is highly upregulated in breast cancer patients, you wonder what will happen if you add low levels of Cyclin D inhibitors to the growth medium. When you perform the experiment, you find that wild type tissue and tissues 2 and 3 do not grow with Cyclin D inhibitors. The results are as follows.



° You know that the Ras protein activates a kinase called Raf. To see if any of the tissues are affected, you add low levels of an inhibitor of Raf to the medium. You find that wild type tissue does not grow at all with this addition. You measure the number of cells in a cell culture plate over time.



a) Based on the results above you can conclude that...

Tissue 1 is _____ Tissue 2 is _____ Tissue 3 is _____

b) Fill in the blanks.

In a cell, one functional copy of the *rb* gene is sufficient to prevent tumor formation.

i) The *rb* gene is therefore referred to as _____ gene.

ii) A different gene involved in tumor formation is the *ras* gene. What kind of gene is

the *ras* gene in terms of cancer biology? _____

iii) Upon binding to cyclins, _____ change their conformations to become active allowing cell cycle progression.

c) Biologically, how does your answer in b i) differ from your answer b ii)?