NAME

7.012 Problem Set 4 FRIDAY October 15, 2004 Answers to this problem set must be inserted into the box outside

TA

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

Question 1

NASA astronauts found a mysterious gene from the H zone of outer space. They needed your help to know more about the gene, so they cloned it into a plasmid vector and told you they would send you the plasmid, that they named *p*SET (plasmid "Sequence Extra-Terrestrial"). When the package arrived, however, you found two unlabeled tubes.

Not sure of what they sent you, you decide to make restriction maps of what's in each tube, assuming that there is plasmid DNA in each.

You perform digestions with the restriction enzymes *Bg*/II (B), *Not*I (N) and *Sa*/I (S) either alone or in combination and obtain the following gel-electrophoresis data.



a) Draw the restriction maps of the plasmids in the space provided below. Write the distances between the restriction sites. Also write down the total sizes of the plasmids in the boxes below, in kilobases.



b) Confused, you call NASA and ask them what's going on and why they sent you two tubes. They do not know why you got two tubes, but they know that the gene of interest had been cloned into the *Not*I and *Sal*I restriction sites of the vector. Which of the plasmids has the gene? Circle your answer.

Plasmid #1

Plasmid #2

c) What is the length of the cloned gene in bases?

d) NASA also told you that the plasmid they used is a bacterial expression vector. Which one of the following elements should be in the vector? Circle all that apply.

centromere	multiple cloning sites	nuclear localization signal
telomere	inducible terminator	bacterial origin of replication
selectable marker	nucloid insertion sites	cytosolic targeting sequences
inducible promoter	topoisomerase gene	gene encoding restriction enzyme

Question 2

You rename the gene GOSH for "gene from the outer space H." For some reason, NASA wants you to study the effect of expressing GOSH in amphibians. To express GOSH in amphibian cells, you have to clone it into an amphibian expression plasmid (pEA) with an amphibian promoter. Below is the map of pEA.



a) Since *p*EA does not contain *Not*I and *Sal*I sites, you decide to *PCR*-amplify *GOSH* with primers that will replace the *Sal*I and *Not*I sites with *BamHI* and *XhoI* sites. You know the sequences of the 5' and 3' ends of *GOSH* as they are in the bacterial plasmid:

Start Stop 5'GTCGACatggtcgccatgcga.....tgctcgatatcgtaaGCGGCCGC3' Sall NotI

Which of the following primer pairs would you use to successfully amplify the gosh gene with the desired restriction sites at either end? All primer sequences are written in the $5' \rightarrow 3'$ direction. Circle all that apply.

Pair1: GTCGACATGGTCGCCATGCGA	and	GCGGCCGCTTACGATATCGAG
Pair2: TCGCATGGCGACCATGTCGAC	and	GGATCACTACGAGATCGAGCA
Pair3: GGATCCATGGTCGCCATGCGA	and	CTCGAGTTACGATATCGAGCA
Pair4: TCGCATGGCGACCATCTCGAG	and	TGCTCGATATCGTAAGGATCC
Pair5: CTCGAGATGGTCGCCATGCGA	and	GGATCCTTACGATATCGAGCA

Now you try to express the new construct (*pEA-GOSH*) in frog (Xenopus) embryos. Hmm... Something seems wrong and the expected peptide is not expressed. Since PCR can introduce errors, you sequence *pEA-GOSH* from its 5' end and obtain the following sequencing gel:



b) What is the sequence in the gel? (in $5' \rightarrow 3'$ direction)

- c) Based on the sequence data, why is GOSH not expressed? Circle all that apply.
- i) Restriction enzyme recognition site is lost, resulting in an immature, nonfunctional peptide.
- ii) A nonsense mutation is introduced, introducing an early stop codon.
- iii) The promoter in the plasmid is lost, so GOSH cannot be expressed.
- iv) A frameshift mutation is introduced, so the encoded protein has a different primary structure from *Gosh*.

Question 3

After fixing the mutation, you express *p*EA-*GOSH* in frog embryos. The transformed frogs show abnormal eye-tongue coordination. You call this phenotype *Helpless*. (See an artist's depiction below.)



Figure by MIT OCW.

To better understand the function of Gosh, you construct two more plasmids. The first one is pEA-GFP, which will express the green fluorescent protein in amphibians. The second one is pEA-GOSH-GFP, which will express a fusion protein of Gosh and GFP. You examine the impact of expressing these plasmids in frogs.



Figure by MIT OCW.

a) According to the result above, does the presence of GFP affect the function of *Gosh*? Briefly explain your answer.

b) Under a microscope, you compare the green fluorescence signals in frog neuronal cells that express either GFP alone, or the *Gosh*-GFP fusion protein. (If your P-set is printed black and white, please look at this colored figure in the PDF file on a computer screen.)



Figure by MIT OCW.

Legend:

A and B are two independent microscopy pictures of the same neuronal cell that expresses GFP.

In A, the green signal comes from GFP molecules. In B, the blue signal comes from DAPI molecules. DAPI is a blue dye that binds (and therefore stains) DNA.

Likewise, C and D are two pictures of another neuronal cell, which expresses the *Gosh-GFP*. Green signal comes from GFP molecules and blue signal comes from DAPI molecules.

c) What do these images suggest? Circle all that apply.

- i) GOSH contains an NLS.
- ii) GFP contains an NLS.
- iii) GOSH-GFP is transported into the central vacuole.
- iv) GOSH-GFP is synthesized by ribosomes on rough ER.
- v) GOSH-GFP is a part of the nuclear pore complex.

d) You intend to find out which part of the *Gosh* protein causes the phenomenon seen in part B. How would you do it? (Please keep your answer within 20 words. Do NOT go into technical details. Just write down the <u>principle</u> of your experiment briefly.)

Question 4:

A human geneticist next door comes to you to discuss a hyperactive human mutant, who in his bone marrow can convert a small amount of adenosine into caffeine. After doing some research, you find out that this person grew up in a region where *Coffea mirus* (a coffee tree) was abundant. At age 10, he had a viral infection during a viral epidemic in *Coffea mirus*.

a) You want to find out if a known enzyme that makes caffeine in *Coffea mirus* is **expressed** in the human mutant's bone marrow. Which of the following methods would you use?

i) Create a cDNA library from the mutant's bone marrow. Then probe for the cDNA encoding the enzyme in the library.

ii) Create a genomic DNA library from the mutant's bone marrow. Then probe for the gene encoding the enzyme in the library.

iii) Create a genomic DNA library from the *Coffea mirus* tree. Then probe for the gene encoding the enzyme in the library.

iv) Create an rDNA library from the mutant's bone marrow. Then probe for the rDNA of the enzyme in the library.

v) Create an rRNA library from the mutant's bone marrow. Then probe for the rRNA of the enzyme in the library.

b) Suppose you do find the coffee-tree enzyme expressed in the mutant's bone marrow. What is a likely vehicle that transferred the gene from the coffee tree to the human mutant? [Give a very short yet concise answer, please.]

We hope you liked this p-set. Please remember to hand it in by 4:40pm on Oct 21.