Biology 13 2024 Winter Term - Exam 1

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Name: Juson Jiao	NetID _	FU06 VMY
Directions : Please put your name on ALL pages. Point values for each question are indicated and should serve as a rough guide for the amount of time you spend and/or the amount of writing you do. There are 50 points total . You have 90 minutes to complete this exam. Good luck.		
You must confine your answers to the page on which each question appears. DO NOT WRITE OF THE BACK OF ANY EXAM PAGE (answers on the back will not be graded). The space provided for each question should be sufficient for your answer.		
You cannot hand in extra pages that contain answ	ers.	
You have 90 minutes to complete this exam. Please pace yourselves accordingly.		
Remember the ideals of academic honor, as exemplified by the Dartmouth Honor Principle. Please keep these tenets in mind while completing this exam. Thank you.		
NOTE: Use of ANY electronic device is prohibited during the exam.		
In accordance with the Dartmouth Honor Principle examination paper as totally my own, having neith during the course of the examination. My signature (computers, phones, watches, calculators) have been of the exam.	her received n re also indicate een placed in r	es that all electronic devices my bag or backpack for the duration
Signature Jousen Jian	Da	ate 1)16)24

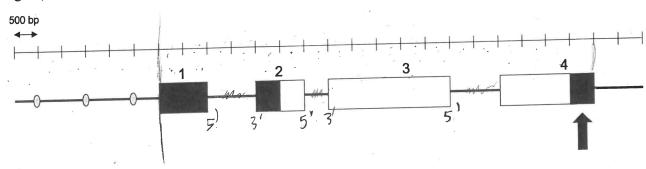
(o) Repetitive DNA in intergenic regions is important for controlling the transcription of adjacent talse, intergeniz DIVA does not

genes.

Tras Jason

Short Answer. Point values vary.

1) The figure below shows a diagram of a region of a eukaryotic chromosome. The exons (boxes) for a single gene are shown. The white boxes indicate the coding region. The location of all of the important transcriptional regulatory elements (enhancers) in the control region for this gene are indicated by ovals. The ruler above provides a size scale. The gene is transcribed left to right (i.e. the 5' end of the mRNA is in exon 1). (10 pt total)



a) How many introns does this gene have? (0.5 pt)

b) How large is the mature mRNA for this gene? (2 pt)

c) What is the arrow on the right indicates? (1 pt)

d) How large (in amino acids) is the protein? (1 pt)
$$(500) - (500) / 3 = 1500$$

e) How large (in base pairs) is the gene? (2 pt)

f) Using the numbering system we discussed for genes, define the position of:

a) the AUG (the start codon for the protein) (0.5 pt)

b) the stop codon (0.5 pt)

$$17.500 = 8500 + 8500$$

b) location of three of the enhancers (0.5 pt)

(g) Assume this gene is capable of alternative splicing involving exons 2 and 3. What spliced forms

Form 1: Pon't use 3' or 5' splice junction so Exon 2 becomes part of intron

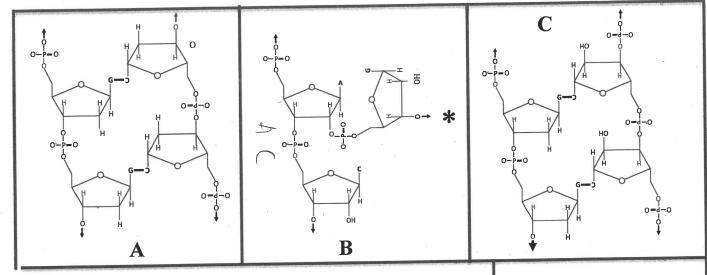
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Form 2: Pon't use exon 2 5' splice junction and exon 3 3' splice junction

Intron' between exon 2+3 becomes part of one big exon

D

2) Below are four structures: A, B, C, and D. Arrows in the figure indicate that the polynucleotide chain continues in that direction. (11 pt total)



a) What is structure A? (1 pt)

DNA

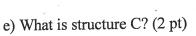
b) Give one example of where you might find structure A in cells. (1 pt)

Nucleus

c) What is structure B? (1 pt)

Powt of sphreosome where branch-pt sequence A
3 bound to the bat free end of intron.

d) In B, what is the next nucleotide in the sequence at the position of



RNA-DNH hybrid

f) Give one example of where C occurs in cells? (2 pt)
Nucleus during transcription when DNH and RNH our e bound

g) Give one example of where you might find structure D in cells. (2 pt)

Nucleus as mRNH 12 beines polynerized in transcription bubble and NTPs are being added Name:

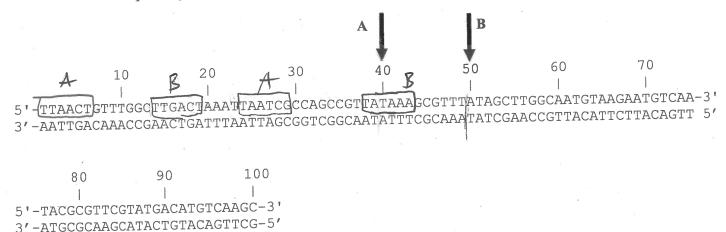
Jason Jao

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January 16, 2024

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3) You are studying a region of the *E. coli* genome. Two RNAs (A and B) are transcribed from the 100 base pair region, and the transcription start sites are indicated with arrows. (14 pt total)



Note: The sequence above is one sequence, drawn on two lines because it will not fit on one line.

a) What are the first ten nucleotides of the RNA produced from transcription start site A? (3 pt) I/AHHGCGUUU

b) What are the first ten nucleotides of the RNA produced from transcription start B? (3 pt)

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- c) Put a box around the key promoter elements for both A and B? Label each element and make it clear whether it corresponds to A or B. (4 pt)
- d) Which promoter (A or B) directs higher levels of transcription? Clearly explain your reasoning. (4 pt)

B b/c both promoters clearly match the conserved

Sequence much more than A. In total B matches 10

sequence much more than A. In total B matches 10

out of 12 bases while A matches roughly 6 out of

12. B/c B matches more closely, the RIVA polymense

will hind more effectively, directing higher Leve to

of transcription