## 科目 分子生物學 科號 0805 共 7 頁第 1 頁 \*請在試卷【答案卷】內作答

- I. Choose a correct answer from each of the following questions. (40%, 2% each)
- 1. Which of the following RNA polymerases is most resistant to α-amanitin?
  - (A). RNA polymerase I
  - (B). RNA polymerase II
  - (C). RNA polymerase III
  - (D). B and C
- Which of the following RNA polymerase II subunits is homologous to the β subunit of E.coli RNA polymerase.
  - (A). RPB1
  - (B). RPB2
  - (C). RPB3
  - (D). RPB4
- 3. Which of the following genes has an internal promoter?
  - (A). β-globin gene
  - (B). 7S rRNA gene
  - (C). 5S rRNA gene
  - (D). U2 RNA gene
- 4. Which of the following transcription factors contains a TATA binding protein?
  - (A). TFII A
  - (B). TFII B
  - (C). TFII C
  - (D). TFII D
- Phage promoters that have been engineered into pBluescript to transcribe in vitro by one phage polymerase to produce strand specific RNA are:
  - (A). T3 and SP6 promoters
  - (B). SP6 and T7 promoters
  - (C). T7 and T3 promoters
  - (D). All above

### 科目 分子生物學 科號 0805 共 7 頁第 2 頁 \*請在試卷【答案卷】內作答

- Select a correct statement about bacterium B. subtilis.
  - (A). The sporulation-specific sigk rearranged with sigH during sporulation.
  - (B). The rearrangement process is irreversible.
  - (C). The rearrangement process usually occurs during development.
  - (D). All above.
- The co-crystal structure of a λ repressor fragment with an operator fragment shows the information about protein-DNA interaction.
  - (A). The most important contacts occur in the major groove.
  - (B). The most important contacts occur in the minor groove.
  - (C). The hydrogen bonds are not very important in the interaction.
  - (D). All above.
- Select a correct statement.
  - (A). The phage σ55 factor, like σ54 of E. coli, is defective.
  - (B). The phage σ<sup>55</sup> factor needs σ<sup>70</sup>.
  - (C). The T4 enhancer is a fixed region.
  - (D). All above.
- 9. Which of the following itemsdoes not induce apoptosis?
  - (A). E2F-1
  - (B). p53
  - (C), RB
  - (D). Myc
- 10. Which of the following statements about transcription factors is <u>not</u> true?
  - (A). Myc dimerized with Mad that interacts with mSin3 to repress transcription through E-box.
  - (B). CBP/p300 is a co-activator for transcription.
  - (C). R-Smad/Co-Smad complex translocates to the nucleus in response to TGF-β signaling.
  - (D). p53 can function as both a transcription activator and a transcription repressor.

### 科目\_分子生物學\_科號\_0805\_共\_7\_頁第\_3\_頁 \*請在試卷【答案卷】內作答

- 11. Which of the following properties about the signaling pathways is not true?
  - (A). Nitric oxide can induce cGMP.
  - (B). PKB/Akt is phosphorylated and activated by PI-3 kinases.
  - (C) Phosphorylation of EGF receptor by PKC decreases its affinity for EGF.
  - (D). G<sub>BT</sub> can direct inhibit the activity of some isoforms of adenylyl cyclase.
- 12. What is the direction of DNA synthesis by DNA polymerases?
  - (A) 3' to 3'
  - (B) 2' to 5'
  - (C) 3' to 5'
  - (D) 5' to 3'
  - (E) 5' to 5'
- 13. Which of the following is <u>not</u> required by DNA polymerase for in vitro synthesis of DNA?
  - (A) template
  - (B) ATP
  - (C) primer
  - (D) deoxynucleoside triphosphates
  - (E) Amino acids
- 14. Which of the following values is significantly unreasonable?
  - (A) An Okazaki fragment is 1,000 to 2,000 bases long.
  - (B) An RNA primer is 5 to 15 bases long.
  - (C) The E. coli genome is 5,386 base pairs long.
  - (D) The human genome is 3 billion base pairs long.
  - (E) A DNA primer is 500 to 1500 bases long.
- 15. Which one of the following proteins and enzymes do <u>not</u> function at the origin of replication in E. coli?
  - (A) DNA ligase
  - (B) SSBPs
  - (C) DNA gyrase
  - (D) DnaA, DnaB and DnaC proteins
  - (E) helicase

# 科目 分子生物學 科號 0805 共 7 頁第 4 頁 \*請在試卷【答案卷】內作答

- 16. Who discovered the first transposable elements in a study of maize (Corn) in the late 1940s?
  - (A). Rosalind Franklin
  - (B). Thomas Morgan
  - (C). Barbara McClintock
  - (D). James D. Watson
  - (E). Kary B. Mullis
- In E. coli glucose repressed the gene expression of many sugar utilization genes (ie. lactose operon). Choose the correct statement(s)
  - I. The glucose effect is mediated by the operator site of lactose operon.
  - II. The glucose effect is mediated by the lactose repressor.
  - III. The glucose effect is by a negative regulation.
  - IV. The glucose effect is mediated by the concentration of cAMP.
  - (A). III
  - (B). IV
  - (C). I and II
  - (D). III and IV
  - (E). I, II, and III
- 18. Which of the following enzymes or enzyme activities is <u>not</u> needed for the synthesis of Okazaki fragment during replication of E. coli chromosome.
  - I. DNA polymerase I
  - II. DNA polymerase III
  - III. primase
  - IV. Helicase
  - (A). I
  - (B). III
  - (C). I and III
  - (D). II, III and IV
  - (E). I, III, and IV

## 科目 分子生物學 科號 0805 共 7 頁第 5 頁 \*請在試卷【答案卷】內作答

- 19. Which of the following is not used in the regulation of eukaryotic gene expression?
  - I Inactivating or altering DNA
  - II. Controlling transcription with specific transcription factors
  - III. Splicing precursor mRNAs differently
  - IV Operons with polycistronic mRNA
  - (A). I
  - (B). II
  - (C). IV
  - (D). I and III
  - (E). III and IV
- 20. Which of the following statements concerning the telomeres is not true?
  - I Hairpin DNA configurations
  - II Repeated sequence
  - III Template-free sequence additions
  - IV G:G base pairs
  - (A). I
  - (B). III
  - (C). IV
  - (D). II and IV
  - (E). III and IV

# 科目 分子生物學 科號 0805 共 7 頁第 6 頁 \*請在試卷【答案卷】內作答

11	. Fill the blanks in the following questions.	(30%, 2% each)
1.	Transcription activator GAL4 stimulates transcription by facil to the preinitiation complex.	litating binding of
2.	Classical class III genes require two factors and to form a preinitiation complex with the poly	ymerase.
3.	The eukaryotic initiation factor eIF-4F is a protest production of eIF-4F and has	ein. eIF-4A is a
4.	Phage $\lambda$ establishes lysogeny by causing production of enough	
	to bind to the early operators.	
5.	Centrosome is the primary microtubule-organizing center that	contains a pair of
6.	Binding of ligand causes dimerization and auto-phosphorylation tyrosine kinases, and the phospho-tyrosines subsequently serve for proteins having or domains to signaling.	as docking sites
7.	The N-terminal glycine of Src protein is modified by bind to the cytosolic face of the cell membrane.	in order to
8.	Margaret Kidwell and her colleagues investigated the inserted into the white locus of dysgenic flies. They found the had great similarities in base sequences but differed cons Furthermore, they had direct repeats and were flanked by short bost DNA, both signatures of transposons.	at these elements siderably in size.
9.	Technically, can occur by either independ chromosomes or by crossing over between chromosomes. Husually refers to the latter.	ent assortment of lowever, the term
10.	A structure at the end of a eukaryotic chromosome, containing of a short DNA sequence is called	ng tandem repeats

### 科目 分子生物學 科號 0805 共 7 頁第 7 頁 \*請在試卷【答案卷】內作答

11.	Assuming wobble base pairing can occur, which two codons could be recognized by the anticodon 3'-GAG-5'? Write the codon in the 5'-3' direction.
12.	According to which hypothesis is protein degradation controlled by regions rich in proline, glutamic acid, serine, and threonine?
13.	An E. coli repair system utilizes adenine methylation to remove the unpaired nucleotide of the newly synthesized DNA strand. What is the name of this repair system?
Ш.	Short Essays (30%, 6% each),

- Explain why can histone acetylation enhance transcription of many genes.
- (1). How do we know that mRNA are read in 5' → 3' direction?
  - (2). Describe the similar and different parts in translation termination within prokaryotes and eukaryotes?
- Describe the functions of mitosis-promoting factor (MPF) and anaphasepromoting complex (APC) in the regulation of cell cycle progression.
  - 4. Describe the mismatch repair mechanism in E. coli.
  - 5. You have cloned a gene from tobacco that is regulated by the presence or absence of light. In the presence of light, the gene is turned on and in the absence of light the gene is off. You want to determine what upstream sequences in your clone are involved in light regulation. You have an Agrobacterium Ti-plasmid with a β-glucuronidase gene (whose expression is easily detected in plant tissue) without a promoter. This plasmid also contains a multiple cloning site near the start site of the β-glucuronidase gene. How would you determine what upstream sequences of your cloned gene are important for light-regulated gene expression?