编號:	
	图止风切入学102字十度调工项指生与武武速 茶 / 東 第7 奥
	别: 熱帶植物科學研究所
考試科目	目: 分子生物學 考試日期:0224,節次:3
※ 考生	主請注意:本試題不可使用計算機 請勿在本試題紙上作答,否則不予計分
``	DEFINITION OF THE FOLLOWING TERMS 解釋名詞 (two points for each question.每題2分) (20%)
1.	intron
2.	ribosomal RNA (rRNA)
3.	Wobble hypothesis
4.	transposon
5.	nucleosome
6.	restriction enzyme
7.	enhancer
8.	zinc finger
9.	bioinformatics
10.	Western blot
二、	SIMPLE-CHOICE QUESTIONS 單選題 (1~22 two points for each question; 23~28 one point for each question. 第1~22題,每題2分; 第23~28題,每題1分)(50%) (以A、B、C、D、E五種選項作答,其餘一律不計分)
1.	Which of the following statements is NOT true concerning homologous recombination?
	A. It can be intramolecular.
	B. It can be bimolecular.
	C. Only one cross-over event can occur.
	D. It results in genetic variation.
	E. It is a reciprocal event.

- 2. Which of the following is a technique used for genomic functional profiling?
 - A. DNA microchips
 - B. RNAi analysis
 - C. SAGE

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- D. transcription activation
- E. RFLP

背面仍有題目,請繼續作答)

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系所	新組	引: 熱帶植物科學研究所		
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	3.	Which of the following techniques was developed by Velculescu and colleagues t	to determine	which
		genes are expressed in a given tissue?		
		A. DNA microarray		
		B. DNA microchips		
		C. SAGE		
		D. RFLP		
		E. deletion analysis		
	4.	are homologous genes that have evolved from gene duplication withi	n a species.	
		A. Orthologs		
		B. Paralogs		
		C. Conservation		
		D. Conjugation		
		E. none of the choices are correct		
	5.	Molecular modeling of RuvA and a Holliday junction showed that they form a	th	at favors
		rapid branch migration.		
		A. square planer conformation		
		B. L-shape conformation		
		C. clover leaf conformation		
		D. D loop conformation		
		E. Y-shape conformation		
	6.	Put the following steps of <i>E. coli</i> primosome assembly in the correct order.		
		(1) Primase binds.		
		(2) DnaA binds to oriC at dnaA boxes.		
		(3) DnaB binds to the open complex.		
		(4) DnaA, RNA polymerase, and HU protein melt the DNA.		
		A. 2, 4, 1, 3		
		B. 2, 4, 3, 1		
		C. 4, 1, 2, 3		
		D. 4, 2, 1, 3		
		E. 3, 4, 1, 2		

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7. Which of the following repair mechanisms is a damage bypass mechanism, not an actual repair mechanism?
A. DNA photolyase
B. base excision repair
C. nonhomologous end joining
D. mismatch repair
E. recombination repair

 Insertion of a Group II intron of a particular gene into an intronless version of that same gene is called_____.

A. retrohoming

B. pseudo-genes

C. migration

D. homeostasis

E. none of the choices are correct

9. The correct folding of proteins in the cell is accomplished with the help of ______

A. proteosome.

B. molecular chaperones.

C. glyoxysome.

D. ribosome.

- E. none of the choices are correct
- 10. Trypanosome mitochondria contain two types of circular DNA linked together into large networks. These mitochondria are called
 - A. maxicircles

B. kinetoplasts

- C. leucoplasts
- D. plastids
- E. maxi-mitochondria

背面仍有題目,請繼續作答)

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11. miRNA synthesis involves the cleavage of the double stranded stem portion of a precursor RNA by the enzyme

A. Drosha

B. Dicer

C. RISC

D. AGO

- E. none of the choices are correct
- 12. Which of the following regions on calf-thymus poly(A) polymerase has been shown experimentally_ **NOT** to be necessary for its activity *in vitro*?

A. RNA binding domain

B. polymerase module

C. nuclear localization signals

D. serine/threonine rich regions

E. All domains are necessary.

13. Which of the following complexes found in the spliceosome cycle is mismatched with its snRNP(s)?

A. commitment complex: U6/U4

B. A complex: U1 and U2

C. B1 complex: all snRNPs

D. B2 complex: U2, U5 and U6

E. I complex: U2, U5 and U6

14. Which of the following histone proteins is the most highly conserved from one organism to another?

- A. H1
- B. H2A
- C. H2B
- D. H3
- E. H4

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15. An experiment is planned to look at the effect of blocking the thyroid hormone receptor. Which of the following is a good technique to employ?

A. phosphorylation

B. methylation

C. ubiquitination

D. sumoylation

- E. all of the choices are correct.
- 16. A mixture of TAF was purified and separated on an SDS-PAGE gel. Select the correct order in which they will appear from the top to the bottom of the gel.
 - A. TAF_{II}150, TAF_{II}110, TAF_{II}40, TAF_{II}60, TAF_{II}80, TAF_{II}250

B. TAF₁₁150, TAF₁₁110, TAF₁₁80, TAF₁₁60, TAF₁₁40, TAF₁₁250

C. TAF₁₁600, TAF₁₁110, TAF₁₁40, TAF₁₁150, TAF₁₁80, TAF₁₁250

D. $TAF_{II}250$, $TAF_{II}150$, $TAF_{II}110$, $TAF_{II}80$, $TAF_{II}60$, $TAF_{II}30$

E. TAF₁₁30, TAF₁₁110, TAF₁₁40, TAF₁₁60, TAF₁₁80, TAF₁₁250

17. Protein crystals are used in x-ray diffraction studies because

A. the molecules of proteins in a powder or solution are too loose.

B. the diffraction power of a single protein molecule is too low.

C. several molecules of proteins are in a crystal leading to strong diffraction patterns.

D. the molecules of proteins in a powder or solution are too loose and the diffraction power of a single protein molecule is too low only.

E. the molecules of proteins in a powder or solution are too loose, the diffraction power of a single protein molecule is too low, and several molecules of proteins are in a crystal leading to strong diffraction patterns.

- 18. Which of the following would be effective in blocking the transcription from the *lac* operon in the presence of permease?
 - A. antibody to RNA polymerase
 - B. antibody to the repressor protein
 - C. mutation in the operator
 - D. an antibody to β -galactosidase
 - E. both antibody to RNA polymerase and mutation in the operator

背面仍有題目,請繼續作答)

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- 19. Which of the following enzymes cannot catalyze the formation of a phosphodiester bond?
 - A. endonuclease

B. RNA polymerase

C. DNA polymerase

D. ligase

- E. none of the choices are correct
- 20. What is the RACE technique?
 - A. screening a genomic library
 - B. extending incomplete cDNA sequences
 - C. generating polynucleotide probes
 - D. rapid amplification of genomic DNA
 - E. screening a genomic library and generating polynucleotide probes
- 21. Which of the following is the best way to determine the base content of DNA?
 - A. density gradient centrifugation
 - B. gel electrophoresis
 - C. spectrophotometry
 - D. x-ray diffraction
 - E. deoxyribonuclease treatment
- 22. A new mutant cell line was developed and was found to be defective in polymerase III activity. Which of the following is likely to be observed in this cell line?
 - A. There will be an overabundance of secreted proteins.
 - B. Splicing function is impaired.
 - C. There will be an overproduction of 7 SL RNA.

D. There will be an overabundance of secreted proteins and splicing function is impaired are correct.

E. There will be an overabundance of secreted proteins, splicing function is impaired, and there will be an overproduction of 7 SL RNA are correct.

- 23. Experimental evidence has shown that it is necessary for the promoter and the enhancer to be on the same chromosome in order to function together.
 - A. True

B. False

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- 24. The estimated total number of genes in the Arabidopsis genome is 30,000-35,000.
 - A. True
 - B. False

25. Antibiotic resistance in bacteria can be transferred using insertion sequences.

- A. True
- B. False

26. The control regions of actively transcribed genes usually contain methylated CpG islands.

- A. True
- B. False
- 27. Phosphorylation of eukaryotic initiation factors can play both an inhibitory and stimulatory role in translational control of gene expression.
 - A. True
 - B. False
- 28. rRNA genes are transcribed as large precursor molecules and then spliced to produce the mature rRNAs.
 - A. True
 - B. False

三、 SHORT ESSAY 簡答題(Five points for each question.每題5分)(30%)

- 1. What is the difference between reverse transcriptase PCR (RT-PCR) and standard RCR? For what purpose would you use RT-PCR?
- 2. Describe the principle and method used in making a transgenic plant using Ti plasmid.
- 3. Describe principle of the yeast two-hybrid assay.
- 4. Proper function of messenger RNAs requires capping and polyadenylation. Please explain the function capping at the 5'-end of a pre-mRNA.
- 5. What is posttranscriptional control of gene expression by RNA interference?
- 6. What do you know about next-generation sequencing technology?