

臺北醫學大學 九十 學年度第一學期期末考試題紙 p1

系級	科 目	授課教師	考 試 日 期	學 號	姓 名
保四	分子生物學	趙振瑞	91 年 1 月 8 日 第 3,4 節		
保研一					

I. 簡答題：(30%)

- 請寫出 the reaction of aminoacyl-tRNA synthetase (包括 2 steps and overall reactions、反應物與生成物名稱及最後主要產物之結構)。(10%)
- 請寫出此 prokaryotic DNA antisense strand(不含 introns)經 translation 後，nascent peptide 之 amino acid sequences，並註明 N 端與 C 端(假設於 5' 端已出現 Shine-Dalgarno sequence)。(10%)
5'-----GTGATCATGTTCAGTGCGAATGACCATCTAATAACGAC-3'
- 請簡述 SOS response turns on 之機轉(hint: Lex A 之功能與 Rec A 之角色)。(10%)

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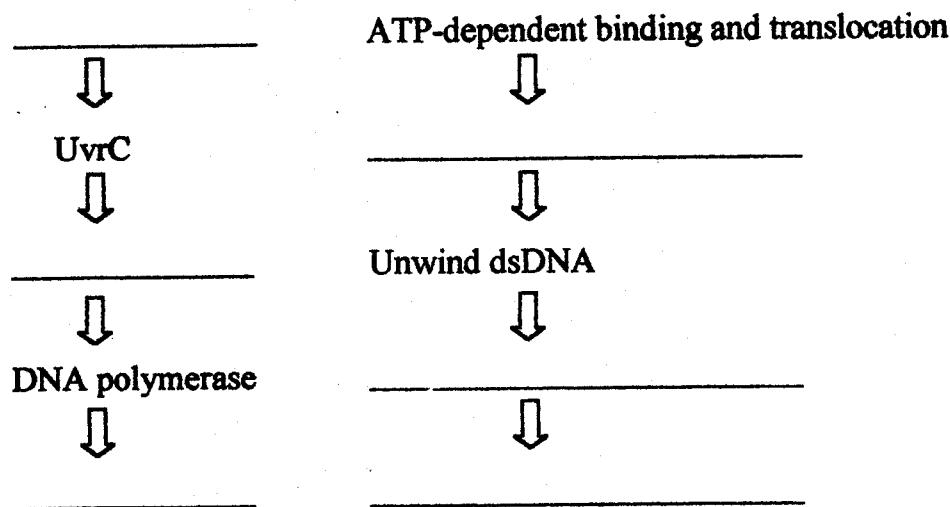
II. 填充題：(26%)(每格一分)

1. Comparisons of prokaryotic and eukaryotic ribosomes and rRNAs :

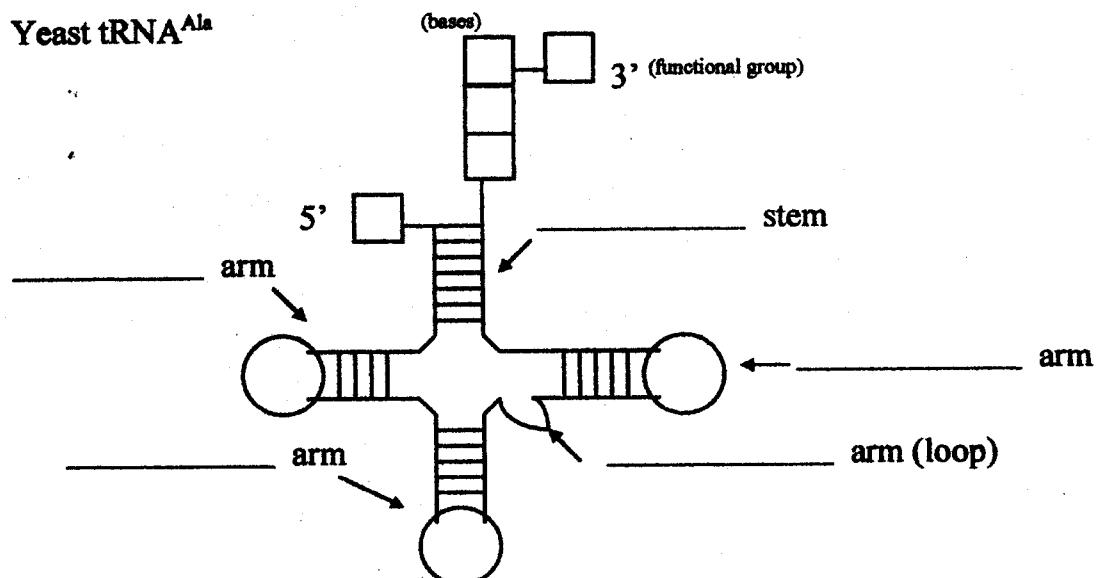
Characteristics	Eukaryotes	Prokaryotes
size of ribosome complex	_____ S	_____ S
size of large subunit	_____ S	_____ S
rRNA(s) in large subunit	_____	_____
size of small subunit	_____ S	_____ S
rRNA (s) in small subunit	_____	_____

2. Mismatch repair mechanism:

Protein(s) Function(s)



3. Yeast tRNA^{Ala}



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III. 配合題：(44%)(每題 2 分)(無重複配對且為單一選項)

- | | |
|---|---|
| <input type="checkbox"/> a. tRNA nucleotidyl transferase | 1. binds at or near 5' cap of mRNA |
| <input type="checkbox"/> b. peptidyl transferase | 2. binds to EF-Tu |
| <input type="checkbox"/> c. donor of formyl Met in prokaryote | 3. binds to GTP and aminoacyl tRNA |
| <input type="checkbox"/> d. IF-2 | 4. binds to GTP and initiator tRNA |
| <input type="checkbox"/> e. EF-Ts | 5. binds to GTP and protein signal sequence |
| <input type="checkbox"/> f. RF-1 | 6. CCA and CCG |
| <input type="checkbox"/> g. eIF-4F | 7. CCA and CGA |
| <input type="checkbox"/> h. synonyms | 8. enzyme to repair alkylated bases |
| <input type="checkbox"/> i. diphtheric toxin | 9. enzyme to directly repair thymine dimers |
| <input type="checkbox"/> j. heme-controlled inhibitor | 10. forms eIF-2 • GTP |
| <input type="checkbox"/> k. puromycin | 11. forms eIF-2-P • eIF-2B |
| <input type="checkbox"/> l. SRP | 12. hydrolyzes the glycosidic bond |
| <input type="checkbox"/> m. KDEL receptor | 13. hydrolyzes the hydrogen bond |
| <input type="checkbox"/> n. DNA glycosylase | 14. inhibits chain elongation in eukaryotes |
| <input type="checkbox"/> o. interferon | 15. inhibits chain initiation in eukaryotes |
| <input type="checkbox"/> p. wobble hypothesis | 16. inhibits chain elongation in prokaryotes |
| <input type="checkbox"/> q. photolyase | 17. inhibits chain initiation in prokaryotes |
| <input type="checkbox"/> r. Holliday recombination model | 18. N ¹⁰ -formyl tetrahydrofolate |
| <input type="checkbox"/> s. Shine-Dalgarno sequence | 19. N ⁵ ,N ¹⁰ -methylene tetrahydrofolate |
| <input type="checkbox"/> t. alkyltransferase | 20. purine-rich sequences on mRNA |
| <input type="checkbox"/> u. Rec A | 21. pyrimidine-rich sequences on mRNA |
| <input type="checkbox"/> v. χ site | 22. sorting ER proteins |
| | 23. sorting secretory proteins |
| | 24. the site of increased DNA mutation |
| | 25. the site of increased recombination |
| | 26. transfers a peptide to aminoacyl tRNA |
| | 27. transfers an amino acid to tRNA |
| | 28. transfers nucleotides to tRNA |
| | 29. UAG recognition |
| | 30. UGA recognition |
| | 31. via 3' end invasion to cross strand |
| | 32. via 5' end invasion to cross strand |
| | 33. with ATPase activity |
| | 34. with endonuclease activity |
| | 35. with helicase activity |
| | 36. wobble position at the first codon on mRNA |
| | 37. wobble position at the third codon on mRNA |

【附表一】 p4
The Standard Genetic Code

First Position (5' end)	Second Position				Third Position (3' end)
	U	C	A	G	
U	UUU Phe			UGU Cys	U
	UUC Phe	UCC Ser		UGC Cys	C
	UUA Leu	UCA Ser			A
	UUG Leu				G
C	CUU Leu	CCU Pro		CGU Arg	U
	CUC Leu	CCC Pro		CGC Arg	C
	CUA Leu		CAA Gln	CGA Arg	A
	CUG Leu		CAG Gln	CGG Arg	G
A	AUU Ile	ACU Thr			U
	AUC Ile	ACC Thr			C
	AUA Ile	ACA Thr	AAA Lys		A
			AAG Lys		G
G	GUU Val		GAU Asp		U
	GUC Val		GAC Asp		C
			GAA Glu		A
			GAG Glu		G