

UNIVERSITI SAINS MALAYSIA

Second Semester Examination
Academic Session 2004/2005

March 2005

BTT 202E/3 - Techniques in Biotechnology
[Teknik-Teknik Bioteknologi]

Duration: 3 hours
[Masa : 3 jam]

Please ensure that this examination paper contains NINE printed pages.

Answer FIVE out of SIX questions. Candidates are allowed to answer all question in English or Bahasa Malaysia or combination of both.

Each question carries 20 marks.

Sila pastikan bahawa kertas peperiksaan ini mengandungi SEMBILAN muka surat yang bercetak sebelum anda memulakan peperiksaan ini.

Jawab LIMA daripada ENAM soalan yang diberikan, dalam Bahasa Inggeris atau Bahasa Malaysia atau kombinasi kedua-duanya.

Tiap-tiap soalan bernilai 20 markah.

1. Answer the following questions based on the information extracted from the NCBI database. Phage T4 is a double-stranded DNA virus.

LOCUS NC_000866 168903 bp DNA linear PHG 13-JAN-2004
DEFINITION Enterobacteria phage T4, complete genome.

gene complement(9410..10729)
/gene="dda"
/locus_tag="T4p017"
/note="synonym: sud"
/db_xref="GeneID:1258784"
CDS complement(9410..10729)
/gene="dda"
/locus_tag="T4p017"
/note="DNA-dependent ATPase, DNA helicase"
/codon_start=1
/transl_table=11
/product="Dda DNA helicase"
/protein_id="NP_049632.1"
/db_xref="GI:9632724"
/db_xref="GeneID:1258784"

```
9301 cagtttcacg atttttgcta taagctatth cataaatttc gcttacttta tcttgaagaa
9361 gataaaactg atcttttagtt atttccacga atagcttctt caaatttaaat catacataaa
9421 acacatcata acgaccacgg gtgacaccaa cataaagaag ttgttgagct aattcaacat
9481 ctgcataatg aatacaaggc gtataaatga aagcacggtc tacagacata ccctgcgctt
9541 tatggaatgt tgatgcagga agtgctttca ctttactaaa ctgtgattta gcaccccaaa
9601 aatcactcca cggagctttt cgcctttgt tccaattttt ataagtttct gctgttttag
9661 ctaaaaatag gttaaaacta tacaattctt cgtcagatga aattatttta atcttttcac
9721 gataaatatt atcatcgcca taagtttcta ctgttaaatc ccaatgacga attagatatt
9781 ctccaggaac accacgggct ttaacaaaac ttgatgtata ctctgcttct ataatacga
9841 ctaattgtcc gttatataaa ataatttctg acacaggctt tccatcaatt ttatatgttt
9901 taaataatgg ttctctgcat acaataatth caccgacaat aaaatcttta tcagtttcaa
9961 aaatcttttt acgaataatg ctatttaact tgtcaacaga tttattcgta aatgccatta
10021 cgcgattttc aaacaaatca tctagtgatt tgacgattga aaaataatth accataaaat
10081 cgcgtaaagc ggtatcacca gtaaatccac gtactccatg cccgtcaaca actttatcat
10141 aattccactt accggttgcga acgtcagtag ctacatcaat aataggagca ttactgcgct
10201 taacttcagt gagttcacac tgataaaaat ctttatgtgt aaagaatgga ctgatataag
10261 cagtattttc tcctggttca acaggtctga tttgcttatt atccccctatt ccaattatag
10321 tacaccaagg tggaaatagtt gaaagcagaa ttttaaatag ctttctatca tacattgaca
10381 ctctcgtegca gattaatact ctgcatttgg ctaaatcagg tacttctttt tgttcaaaaa
10441 gaacattttc ttcatatggt actgggttaa ttttaagaat actatgaata gtactcgctt
10501 ctttccctga tagttttgaa agaatctttt tagctgcatg tgtaggagct gctaaaataa
10561 taccagttcc acccgtagat attaaagctt caatgatgaa cttagtaaga gtagtcttac
10621 cggtagcagc aggtccatta atagttacat gatgtttctt ttctttaata gccttcataa
10681 caatgttaaa ggcatttttc tggccttcgg tcaaatcatc aaatgtcatc gtaaatcccc
10741 tgcaattggg atactaacia tacgcccagt atctaaaatt cgctgatata atctttgcgt
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[BTT 202E/3]

- (a) Design a pair of primers to amplify the helicase gene by PCR. Be sure to specify the forward and the reverse primers and the bases where they are complemented. You are also required to introduce *NdeI* (CATATG) site at the beginning of the gene and *BamHI* (GGATCC) site at the end of the gene.
(5 marks)
- (b) With a help of a diagram, explain the principles of PCR.
(10 marks)
- (c) What would be the expected size of the amplified PCR fragment?
(1 marks)
- (d) If the genome was RNA, instead of DNA, what would you do in order to amplify the helicase gene?
(4 marks)

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1. *Jawab soalan-soalan berikut berdasarkan maklumat dari pengkalan data NCBI. Phage T4 adalah virus yang mempunyai benang DNA ganda dua.*

| | | | | | |
|-------------|---|-------------|------------|------------|------------------------|
| LOCUS | NC_000866 | 168903 bp | DNA | linear | PHG 13-JAN-2004 |
| DEFINITION | Enterobacteria phage T4, complete genome. | | | | |
| <u>gene</u> | complement(9410..10729) /gene="dda" /locus_tag="T4p017" /note="synonym: sud" /db_xref="GeneID:1258784" | | | | |
| <u>CDS</u> | complement(9410..10729) /gene="dda" /locus_tag="T4p017" /note="DNA-dependent ATPase, DNA helicase" /codon_start=1 /transl_table=11 /product="Dda DNA helicase" /protein_id="NP_049632.1" /db_xref="GI:9632724" /db_xref="GeneID:1258784" | | | | |
| 9301 | cagtttcacg | atTTTTgcta | taagctattt | cataaatttc | gcttacttta tcttgaagaa |
| 9361 | gataaaaactg | atcttttagtt | atTccacga | atagcttcct | caaatttaat catacataaa |
| 9421 | acacatcata | acgaccacgg | gtgacaccaa | cataaagaag | ttgttgagct aattcaacat |
| 9481 | ctgcataatg | aatacaaggc | gtataaatga | aagcacggtc | tacagacata ccttgcgctt |
| 9541 | tatggaatgt | tgatgcagga | agtgctttca | ctttactaaa | ctgtgattta gcatcccaaa |
| 9601 | aatcactcca | cggagctttt | ccgcctttgt | tccaattttt | ataagtttct gctgttttag |
| 9661 | ctaaaaatag | gttaaactta | tacaattctt | cgtcagatga | aattatttta atcttttcac |
| 9721 | gataaatattc | atcatcgcca | taagtttcta | ctgttaaatc | ccaatgacga attagatatt |
| 9781 | ctccaggaac | accacgggct | ttaacaaacg | ttgatgtata | ctctgcttct ataatacga |
| 9841 | ctaattgtcc | gttattaaaa | ataatttctg | acacaggctt | tccatcaatt ttatatgttt |
| 9901 | taantaatgg | ttcctgcatt | acaataattt | caccgacaat | aaaatcttta tcagtttcaa |
| 9961 | aaatcttttt | acgaataatg | ctatttaact | tgtaacaga | tttattcgta aatgccatta |
| 10021 | cgcgattttc | aaaceaatca | tctagtgtt | tgacgattga | aaaataattt accataaaat |
| 10081 | cgcgtaaagc | ggtatcacca | gtaaatccac | gtactccatg | cccgtcaaca actttatcat |
| 10141 | aattceactt | accgttgcca | acgtcagtag | ctacatcaat | aataggagca ttactgcgctt |
| 10201 | taacttcagt | gagttcacac | tgataaaaat | ctttatgtgt | aaagaatgga ctgatataag |
| 10261 | cagtattttc | tcctggttca | acaggtctga | tttgcttatt | atcccctatt ccaattatag |
| 10321 | tacacceagg | tggaatagtt | gaaagcagaa | ttttaaatag | ctttctatca tacattgaca |
| 10381 | cttcgctcgca | gattaatact | ctgcatttgg | ctaaatcagg | tacttctttt tgttcaaaaa |
| 10441 | gaacattttc | ttcatatggt | actgggttaa | ttttaagaat | actatgaata gtactcgctt |
| 10501 | ctttccctga | tagttttgaa | agaatctttt | tagctgcatg | tgtaggagct gctaaaataa |
| 10561 | taccagttcc | accgtagat | attaaagctt | caatgatgaa | cttagtaaga gtagtcttac |
| 10621 | cggtaccagc | aggtccatta | atagttacat | gatgtttctt | ttctttaata gccttcataa |
| 10681 | caatgttaaa | ggcatttttc | tggccttcgg | tcaaatcadc | aaatgtcadc gtaaatccc |
| 10741 | tgcaattggt | atactaacaa | tacgcccagt | atctaaaatt | cgctgatata atctttgcgt |

[BTT 202E/3]

- (a) *Reka sepasang pencetus untuk mengamplikasi gen helikase dengan menggunakan PCR. Pastikan anda tentukan pencetus hadapan dan pencetus kebelakang dan juga bes di mana pencetus-pencetus tersebut berkomplimen. Anda juga dikehendaki mewujudkan tapak pembatasan NdeI (CATATG) pada permukaan gen dan tapak pembatasan BamHI (GGATCC) pada penghujung gen.*
(5 markah)
- (b) *Terangkan prinsip PCR dengan bantuan gambarajah.*
(10 markah)
- (c) *Apakah size produk PCR yang dijangkakan terhasil?*
(1 markah)
- (d) *Sekiranya genom adalah RNA dan bukannya DNA, apakah yang patut anda buat untuk mengamplikasikan gen helikase? .*
(4 markah)
2. (a) Describe the functions and the applications of the following enzymes:
- (i) T4 DNA ligase
 - (ii) T4 polynucleotide kinase
 - (iii) Ribonuclease
 - (iv) Terminal deoxynucleotidyl transferase
 - (v) Dnase I
- (10 marks)
- (b) How would you label the following:
- (i) Oligonucleotide
 - (ii) 10 kb double-stranded DNA
- (10 marks)

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2. (a) *Terangkan fungsi dan kegunaan enzim-enzim berikut:*

- (i) *T4 DNA ligase*
- (ii) *T4 polynucleotide kinase*
- (iii) *Ribonuclease*
- (iv) *Terminal deoxynucleotidyl transferase*
- (v) *Dnase I*

(10 markah)

(b) *Bagaimana anda melabelkan yang berikut:*

- (i) *oligonuklutida*
- (ii) *10 kb DNA ganda dua*

(10 markah)

[BTT 202E/3]

3. (a) Explain the principles of ethanol precipitation in plasmid DNA preparation protocol.

(6 marks)

(b) A prepared sample of plasmid DNA could not be digested with restriction enzymes. What are the possible explanations to this problem?

(6 marks)

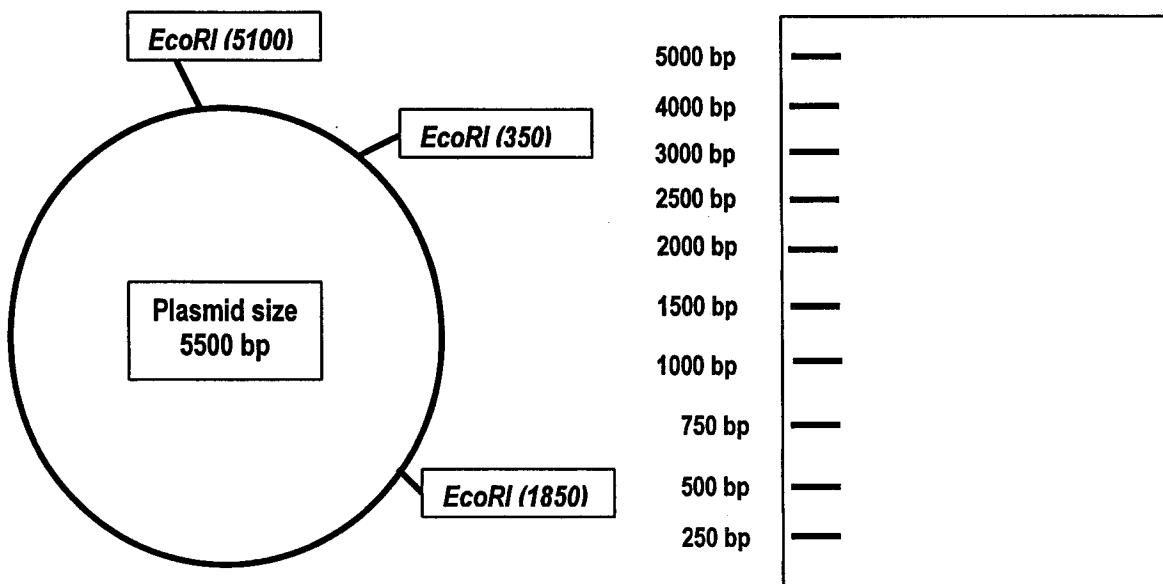
(c) After cleaning-up, the plasmid DNA sample was digested with *EcoRI* and *BamHI*. This was followed by alkaline phosphatase treatment and finally the sample was run on agarose gel electrophoresis.

(i) Draw the fragments resulting from the restriction enzymes digestion on the agarose gel provided.

(2 marks)

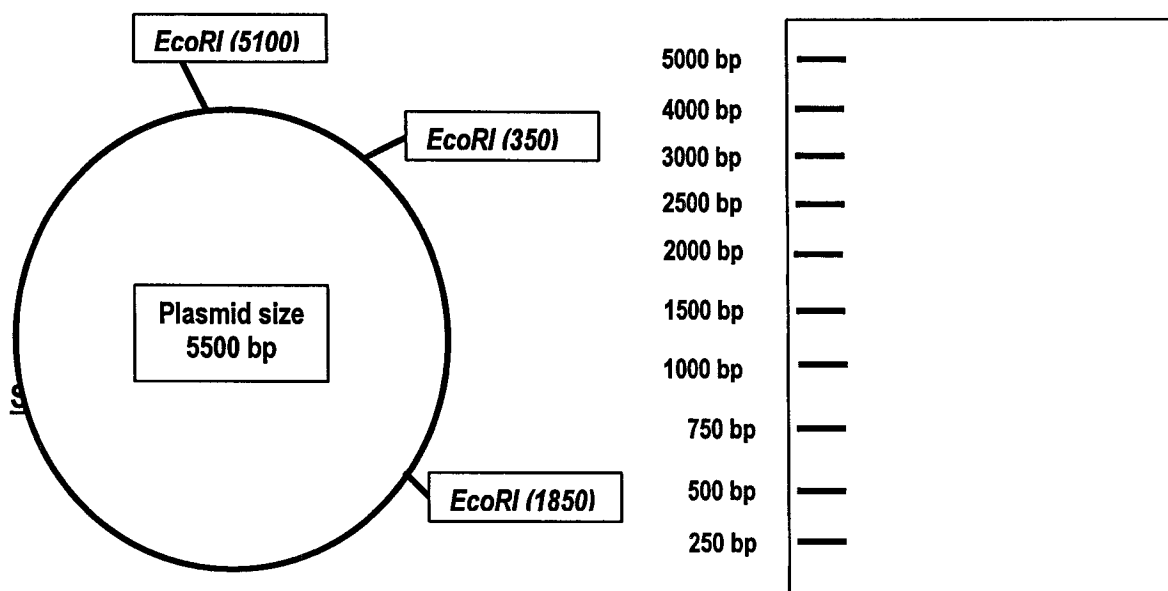
(ii) The biggest fragment was used as a vector in a ligation reaction but failed to ligate to the insert fragment. What was the problem and how would you solve it?

(6 marks)



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3. (a) *Terangkan prinsip pemendakan etanol dalam penyediaan DNA plasmid.*
- (6 markah)
- (b) *Didapati DNA plasmid yang telah disediakan tidak boleh dihadamkan oleh enzim pembatasan. Apakah penjelasan-penjelasan yang mungkin kepada masalah ini?*
- (6 markah)
- (c) *Selepas dicuci, DNA plasmid dihadamkan dengan enzim pembatasan EcoRI dan BamHI. DNA plasmid kemudiannya ditindakbalas dengan enzim fosfatase alkali dan akhirnya dilarikan di atas gel agarosa.*
- (i) *Lakarkan produk-produk yang terhasil dari penghadaman enzim pembatasan di atas gel agarosa yang disediakan.*
- (2 markah)
- (ii) *Serpihan yang terbesar telah digunakan sebagai vektor dalam tindakbalas ligasi tetapi gagal untuk diligasi. Apakah masalah yang dihadapi dan bagaimana untuk mengatasinya?*
- (6 markah)



4. Describe the ultrafiltration technology and its application in downstream processing.
(20 marks)
4. *Terangkan teknologi penurasan ultra serta kegunaannya dalam proses hiliran*
(20 markah)
5. Draw the flow-chart of the purification of a bacterial intracellular protease. Discuss each unit-process involved.
(20 marks)
5. *Lakarkan carta-alir proses penulenan enzim protease yang dihasilkan dalam sel bakteria. Bincangkan setiap proses unit yang terlibat.*
(20 markah)
6. Write notes on gel-filtration and ion-exchange chromatography.
(20 marks)
6. *Tulis nota berkenaan kromatografi penurasan-gel dan pertukaran ion.*
(20 markah)