

Code No: R05412301

R05

Set No. 2

IV B.Tech I Semester Examinations, November 2010
COMPUTATIONAL MOLECULAR BIOLOGY
Bio-Technology

Time: 3 hours

Max Marks: 80

Answer any FIVE Questions
All Questions carry equal marks

1. (a) What are the differences between the initial phases of BLAST and FASTA?
(b) Briefly describe how the PAM and BLOSUM scoring matrices are derived and how they are different. [8+8]
2. Explain about word or k-tuple methods, used by programs FASTA & BLAST? [16]
3. Transmembrane proteins are important in drug discovery
What are their properties and how can we generate their 3-D structures. [16]
4. What are the different methods used for detecting SNPs and give its applications? [16]
5. Write short notes on the following :
(a) Weighted parsimony
(b) Unweighted Parsimony. [8+8]
6. Describe the process of developing spotted arrays? [16]
7. How can you predict protein structure using Ramchandran's plot? [16]
8. Describe about Jukes - Cantor model for estimation of substitution numbers. [16]

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R05

Set No. 4

IV B.Tech I Semester Examinations, November 2010
COMPUTATIONAL MOLECULAR BIOLOGY
Bio-Technology

Time: 3 hours

Max Marks: 80

Answer any FIVE Questions
All Questions carry equal marks

1. Explain about word or k-tuple methods, used by programs FASTA & BLAST? [16]
2. Describe about Jukes - Cantor model for estimation of substitution numbers. [16]
3. Write short notes on the following :
 - (a) Weighted parsimony
 - (b) Unweighted Parsimony. [8+8]
4. What are the different methods used for detecting SNPs and give its applications? [16]
5. Transmembrane proteins are important in drug discovery
What are their properties and how can we generate their 3-D structures. [16]
6. Describe the process of developing spotted arrays? [16]
7. (a) What are the differences between the initial phases of BLAST and FASTA?
(b) Briefly describe how the PAM and BLOSUM scoring matrices are derived and how they are different. [8+8]
8. How can you predict protein structure using Ramchandran's plot? [16]

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R05

Set No. 1

IV B.Tech I Semester Examinations, November 2010

COMPUTATIONAL MOLECULAR BIOLOGY

Bio-Technology

Time: 3 hours

Max Marks: 80

**Answer any FIVE Questions
All Questions carry equal marks**

1. Describe about Jukes - Cantor model for estimation of substitution numbers. [16]
2. What are the different methods used for detecting SNPs and give its applications? [16]
3. Transmembrane proteins are important in drug discovery
What are their properties and how can we generate their 3-D structures. [16]
4. (a) What are the differences between the initial phases of BLAST and FASTA?
(b) Briefly describe how the PAM and BLOSUM scoring matrices are derived and how they are different. [8+8]
5. Write short notes on the following :
(a) Weighted parsimony
(b) Unweighted Parsimony. [8+8]
6. How can you predict protein structure using Ramchandran's plot? [16]
7. Describe the process of developing spotted arrays? [16]
8. Explain about word or k-tuple methods, used by programs FASTA & BLAST? [16]

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R05

Set No. 3

IV B.Tech I Semester Examinations, November 2010

COMPUTATIONAL MOLECULAR BIOLOGY

Bio-Technology

Time: 3 hours

Max Marks: 80

**Answer any FIVE Questions
All Questions carry equal marks**

1. Describe the process of developing spotted arrays? [16]
2. Transmembrane proteins are important in drug discovery
What are their properties and how can we generate their 3-D structures. [16]
3. Describe about Jukes - Cantor model for estimation of substitution numbers. [16]
4. How can you predict protein structure using Ramchandran's plot? [16]
5. Write short notes on the following :
 - (a) Weighted parsimony
 - (b) Unweighted Parsimony. [8+8]
6. Explain about word or k-tuple methods, used by programs FASTA & BLAST? [16]
7. What are the different methods used for detecting SNPs and give its applications? [16]
8. (a) What are the differences between the initial phases of BLAST and FASTA?
(b) Briefly describe how the PAM and BLOSUM scoring matrices are derived and how they are different. [8+8]
