BIF501 11:00

Difine RNa sequencing write a comprehensive note

2.Define RNA sequencing

3.define DNA carry

4. Write a comprehensive note on drug discovery

5

Write two methods of tree approach

6.write a note on splicing

7. Homology modeling

8.how drugs are discover

BIF ....12:30

READSCQ?2

What are long common subsequence? Write purpose.

What is Edit distance?give its example.

Describe the project for sequencing the genome of Haemophilus influenza.

Write the classes of biological database.

What are coding and non.coding regions on DNA sequence?

if501 msqz

differ b/supervsed and unsupervised machine learning

limitation of ab initio

challenge of ab initio

two target discovery approaches

pharmacology and pharmacogenomic

homology modeling folding recognition step

rna structure,type and process

: Today Bif 501 paper 8.00

Mcqs all from past papers

Q1.what is prob

2 explain primary secondary tertiary and qtrnry structure of protein

3 RNA discuss long q for 10 marks

4. Supervised and unsupervised machine learning

5 swiss port

6 secondary structure prediction process

: Bif501 8 am paper

supervised and unsupervised learning

mate and strategy

homology paralogy ortholgy

homology modeling steps

For More Visit VU Students

rna types structure secondary structure

what method used for sequence

Today Bif 501 paper 8.00

Q1.what is prob

2 explain primary secondary tertiary and qtrnry structure of protein

3 RNA discuss long q for 10 marks

4. Supervised and unsupervised machine learning

5 swiss port

6 secondary structure prediction process

BIF501 TIME= 5:00 DATE=27-08-2019

Maximum MCQs were from the past files.

Write and explain 3 approaches of target discovery.

Pharmacology and pharmagenomics and its 3 applications.

What is Edman degradation reaction strategy.

Challenges of Ab intio modelling.

Human T cells have how many fragments of tryptosine.

Explain RNA and its structure with 4 types and process of RNA forming.

What is fold recognition process.

What is substitution matrices write down its three types.

2 approaches of target discovery.

BIF501 VU Current Final Term Papers Spring 2020 (10 September onward)

Bif501 11:00

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BIF501 VU Current Final Term Papers Spring 2020

Bif 501 8:00 am

Define profiles

How Sanger solved edman degradation problem

3: how to calculate ratio of likelihoods

4: define two types of mapping

5: restrictions mapping uses, benefits

6: how stimulate folding process

7: work of neural network

8: gene mapping

9: after gaps and mismatches which step used in homology

10: fold recognition

11: needs of ab initio modelling

12: work flow of structural. Modelling

BIF501 current paper | 11-09-2020

Insertion deletion substitution (3)

Ab initio (5)

Drug discovery(10)

Homology modeling (3)

Bif501

11 sept

Insertion deletion substitution (3)

Ab initio (5)

Drug discovery(10)

Homology modeling (3)

Bif 501 5:00pm

Drugs?(2)

Pharmacogenomics and applications?(10)

MAPGIE and gene quiz?(2)

Advantages and disadvantages of abinitio method(5)

For More Visit VU Students

Chanlenges of ab-initio method?(3)

Splicing signel?(5)

Steps of (bhool gay or kisi ky steps ay thy 3mrkx ky

Mcqs past and ppts mix

Regards

Current Paper BIF501. (08:00 AM, 17-02-2020)

What is protein fold 2marks

When fold recognition technique employ 2 marks

Which is the first alogrithm for protein sequence 2 marks

What are two discovery aproach 3 marks

Difference between supervised and unsupervised machine learning 3 marks

Fold recognition technique 5 marks

Which the sequence involved in four different group expalin it 5 marks

What is RNA describe its structure 10 marks Mcq's all from past

BIF501 Bioinformatics II Current Papers Final Term Fall 2019

Current Paper BIF501. (08:00 AM, 17-02-2020)

What is protein fold 2marks

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