

ODD SEMESTER

BIO INFORMATICS (SAC5B)-SESSION PLAN

| SESSION NO | TOPICS TO BE COVERED | TEACHING METHOD |
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| UNIT-I | | |
| 1 | Genomic sequences-DNA,RNA,Peptide sequences,sequence analysis and sequencing | BB |
| 2 | Output management from different biological output sources-Biological databases,DDBJ,Gen cord,NCBI,PIR | BB |
| 3 | Gene prediction rules –Introduction, About codons,Exons | PPT |
| 4 | Gene finding rules in Prokaryotes and Eukaryotes | BB |
| 5 | Gene finding programs in Prokaryotes and Eukaryotes-based on algorithm | PPT |
| 6 | Gene prediction software – GenScan, GenView 2, NetGene 2 ORF Gene - Gene Structure Prediction using Homologous Proteins. | BB |
| 5 | Human Genome Project – Introduction,History,Applications,Ethical and social Issues | BB |
| 6 | Human Genome Project analysis Software at NCBI- RefSeq Proteins, ClinVar | PPT |
| 7 | Mutations-Introduction, Mis Sence databases, HGMD | BB |
| 8 | Population studies –Introduction, Demographic Model, HFD,HFC,DAT life | BB |
| 9 | cDNA sequences –Definition ,Retrieving Datas, EMBL,EBI | BB |
| 10 | Revision of unit-I | Group discussion |

| UNIT-II | | |
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| 11 | Gene therapy- Introduction, Potential uses of Gene therapy | BB |
| 12 | Types of gene therapy-Exvivo, Invivo, Methods for inserting and expressing a gene in target cell,Virus mediated gene transfer systems | BB |
| 13 | Applications of gene therapy-Advantages and Disadvantades | BB |
| 14 | Analysis of genomic and proteomic information with respect to biological systems –Gene expression studies,Determination of tumor marker genes. | BB |
| 15 | Genome application – Transgenic animals –Introduction, Methods of production of transgenic animals, Gene transfer methods | BB |
| 16 | Examples of transgenic animals-Transgenic fish, Transgenic Pig, Transgenic mouse, Transgenic live stock | PPT |
| 17 | Transgenic Plants –Introduction, Methods of production of transgenic plants, Gene transfer methods. | BB |
| 18 | Examples of transgenic animals- High glyisine corn, Herbicide tolerant plants, BT cotton | BB |
| 19 | Drug design - Introduction, History ,targets , Types ,Mechanism based design, Structure based, ligand based | Group discussion |
| 20 | Drug discovery - Introduction, History ,Therapeutic development pipeline, Target Selection | BB |
| 21 | Drug identification- Introduction, Drug Schedules, Drug categories, Stimulants | BB |
| 22 | Synthesis of new drugs-Stages, Preclinical trails, Clinical trails, Drug discovery phase | BB |

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| 23 | Animal Studies in Synthesis of new drugs-Pharmaco dynamic studies, Toxicological studies | BB |
| 24 | Management of diverse chemical libraries-Definition, Purpose, Generation, Design and optimization, Storage and development | BB |
| 25 | Revision of unit-II | BB |
| CIA TEST-I | | |
| UNIT-III | | |
| 26 | Gene expression- Introduction, Central dogma, Replication of DNA, Steps in replication, Enzymes in Gene expression | BB |
| 27 | Gene expression-Translation, Termination, Pribnow box, Steps in Translation &Termination | BB |
| 28 | Classification of Cancer types-Definition, Types of cancer, Examples of cancer, Diagnosis &Treatment | BB |
| 29 | Genes; Oncogenes -protooncogenes –Regulatory genes, Factors activating genes, Tumour suppressor genes | BB |
| 30 | Application of Microarrays in Drug toxicity testing-Microarray technology, Hybridization, Detection | BB |
| 31 | Microarrays in Drug toxicity testing –Microscope slide, Applications & limitations | BB |
| 32 | Metabolic pathways-Introduction, Metabolism-Catabolism, Anabolism, Energy Production, Enzymes action | PPT |
| 33 | Revision of unit-III | BB |
| CIA TEST-II | | |
| UNIT-IV | | |

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| 34 | Sequence analysis(Proteins)- Sequencing, Alignment, Fluorescence based sequencing, Methods- Edman degradation, Mass Spectroscopy | BB |
| 35 | Protein Sequencing methods-Pair wise Alignment, Multiple sequence Alignment | BB |
| 36 | Sequence analysis (Nucleic acids) –Data base –BLAST, NCBI, DDBJ, Overview of Gen bank | BB |
| 37 | Sequence alignment methods -Introduction, Types, Methods, Applications | BB |
| 38 | Sequence alignment methods-DOT matrix, Dynamic Programming method, Work method or K-tuple method | BB |
| 39 | Proteomic-Definition ,Proteome, Types, Structure of protein, Tools of Proteomics | BB |
| 40 | Proteins analysis – structural comparisons – 2D gel-Introduction, Principle, Overview, Steps, Sample Preparation, SDS-PAGE | Group discussion |
| 41 | Proteins analysis – structural comparisons –Mass spec-Introduction, Principle, Overview, Steps, Instrumentation, Applications | BB |
| 42 | Protein arrays related disease-Analytical Protein array, Instrumental Protein array, Biochemical activities | BB |
| 43 | Antibody arrays related disease-ELISA, DNA, Tissue microarray | PPT |
| 44 | Preventive arrays related disease-Concepts, Timeline approaches, Prevention approaches, Types of arrays | BB |
| 45 | Revision of unit-IV | Group discussion |
| CIA TEST-III | | |

| UNIT-V | | |
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| 46 | Protein Database: Definition, Examples, Applications. | BB |
| 47 | Protein Database -SWISS-PROT, Tr EMBL, NRL, PDB, CATH | BB |
| 48 | Comparison of Protein sequences and Database searching – Introduction, Pro CKSI, Results, Conclusion | BB |
| 49 | Methods for protein structure prediction –levels of Protein Structure | BB |
| 50 | Methods for protein structure prediction-NMR, X ray Crystallography, Electron Microscopy | BB |
| 51 | Conserved patterns in protein sequences and structures – | BB |
| 52 | Comparison of protein 3D structures –Different types of methods | BB |
| 53 | Predicting functions based on DNA and protein sequences- FASTA programs, Primary, Secondary, Composite | BB |
| 54 | Predicting functions-Prediction Strategies, Methods, GOR Method | PPT |
| 55 | Revision of unit-V | Group discussion |
| Revision | | |
| 56 | Revision of unit-I | BB |
| 57 | Revision of unit- II | BB |
| 58 | Revision of unit-III | BB |
| 59 | Revision of unit-IV | BB |

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| 60 | Revision of unit-V | BB |
| MODEL EXAM | | |

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