

LSSSDC (GOVT. OF INDIA) CERTIFIED 28 CREDIT COURSE BIOINFORMATICS ANALYST

Empowering Your Career through Practical Training and Industry Recognition

www.dromicslabs.com











Why DrOmics Labs





+91 9310870544, +91 7888118030, +91 7843088966



info@dromicslabs.com



www.dromicslabs.com





Key Features of Our Course:

- 1.28 Credit worth Comprehensive Learning: Our intensive course is equivalent to 28 Indian Education System Credits, ensuring you receive in-depth training and expertise in bioinformatics.
- 2. Practical Approach: We believe in learning by doing. Our curriculum emphasizes handson experience, allowing you to apply theoretical concepts to real-world scenarios, enhancing your problem-solving abilities and practical skills.
- 3. Exclusive Interview Opportunities: As part of our commitment to your success, we provide opportunities for you to interview with well-established companies in the bioinformatics domain. Gain insights into industry practices and secure your dream job.
- 4. **Third-Party Examination:** To validate your expertise, we offer a third-party examination option. For a nominal fee of INR1650/-, you can opt for this examination to further enhance your credentials and stand out in the competitive job market.
- 5. Government of India Recognized Certificate: Upon successful completion of the course, you will receive a prestigious certificate recognized by the Government of India, affirming your proficiency in bioinformatics analysis.
- 6. Guidance for Research Publication: We understand the importance of contributing to the field of bioinformatics. Our expert faculty members provide guidance and support for your research endeavors, facilitating publication in reputable journals and conferences.

+91 9310870544, +91 7888118030, +91 7843088966







Are You Interested in Scholarship for This Course ?

ELIGIBLITY

- Masters in any life science field / MSc (2nd yr pursuing)
- Those holding higher degrees in life science disciplines.
- Professionals with experience in Bioinformatics.

SCHOLARSHIP EXAM DATE 26 MAY 2024

SCHOLARSHIP

90%+ Marks

60000/- INR *Scholarship (only for top 3 students)

80-90% Marks

20000/- INR Scholarship

70-80% Marks

15000/- INR Scholarship

60-70% Marks

10000/- INR Scholarship



MODULE: 1 ORIENTATION FOR BIOINFORMATICS OCCUPATION

- Introduction to Bioinformatics and its Applications in Life Sciences
- Career Pathways and Opportunities in Bioinformatics
- The Interdisciplinary Nature of Bioinformatics: Bridging Biology and Computer Science
- Organizational Structure and Employment Benefits in the Life Sciences Industry
- Regulatory Framework and Compliance in Bioinformatics
- The Role of Bioinformatics Scientists in Advancing Life Sciences Research
- Essential Skills and Competencies for Bioinformatics Professionals
- Ethical Considerations and Responsibilities in Bioinformatics Practice
- Bioinformatics Tools and Technologies: A Landscape Overview
- Emerging Trends and Future Directions in Bioinformatics Research and Industry



MODULE:2 INTRODUCTION TO BIOINFORMATICS

- Importance of bioinformatics in modern biology
- Detailed explanation for central dogma of cell : Replication, Transcription, Translation, strand
- Detailed study about structure of gene, transcript,upstream,downstream regions,CDS,UTR
- Detailed study about protein primary/secondary and tertiary structure
- Detailed study about enzymes,Bonds and Interactions
- Basics of nucleotide and protein sequence, and FASTA format,fastq,SAM/BAM
- Pairwise sequence alignment techniques (local, global)and Introduction to multiple sequence alignment
- Introduction to genomics and Proteomics in Bioinformatics
- What is NGS ? Genome assembly and sequencing techniques (e.g., Sanger sequencing, Next-Generation Sequencing)
- Different applications of NGS(ex. DNAseq, RNAseq, CHIPseq, metagenomics, Methlyseq etc.)

MODULE : 3 INTRODUCTION TO BIOINFORMATICS DATABASES

- Understanding Data Sources in Bioinformatics: Open Source vs. Paid
- Utilizing Tools for Data Import from Public and Private Databases
- Overview of Bioinformatics Databases: Types and Categories
- Introduction to Types of Databases: primary/secondary/data structure/types of data etc..
- Navigating Genomic Databases: GenBank Database
- Protein Databases: Structure, Function, and Interaction Databases : PDB , UniProt Database
- Data Retrieval Techniques: Querying Databases Using Keywords, IDs: UCSC Database
- Literature Database: PubMed Database
- ClinVar Database
- Integrated Databases: Resources Combining Multiple Data Types (e.g., KEGG, Reactome)
- Ensemble Database



MODULE: 4 BIOINFORMATICS TOOLS

• Introduction to Sequence Alignment

- Types of Alignment(Pairwise & Multiple)
- Local & Global Alignment
- Online Blast
- Standalone BLAST
- MEGA
- ClustalW
- Visualization tools Pymol / Jmol(optional)

MODULE:5 INTRODUCTION TO LINUX

- Overview of Linux
- Package Management
- Basic Commands for file handling
- Advanced Linux commands
- Introduction to Bash Scripting



- Getting Ready with R introduction and installtion
- Data Types, Variables, and Basic R Operations
- Function-buit-in and User defined
- Conditional statments
- Data Wrangling and Cleaning :Importing data into R(e.g., FASTA, GenBank)
- Package installation from CRAN repository and Bioconductor
- Data manipulation with dplyr for biological datasets
- Working with Strings:Sequence Analysis with seqinr and biostring
- Statistical Test-t-test ,z-test ,chiSquare and ANOVA
- Data Visualization

return NULL; roup_info->ngroups = gidsetsize; roup_info->nblocks = nblocks; atomic_set(&group_info->usage, 1);

MODULE:7 INTRODUCTION TO PYTHON LANGUAGE

- Introduction to Python language
- Data types and data structure
- Control statements: if -else, If-elif-else, for loop, while loop
- Python data structure : List, Set, Tuple, Dictionary
- Methods of List, Slicing and indexing in List and Tuple
- Functions : Function introduction and its requirement, Defining a function, Calling a function
- File handling :file handling, OS module
- Pandas library: Reading different file formats such as csv, tsv and excel files
- Biopython
- SeqIO and visualization

MODULE: 8 MACHINE LEARNING AND IMAGE ANALYSIS

- Introduction to Machine Learning Fundamentals for Bioinformatics
- Linear Models and Nearest Neighbors: Learning Algorithms and Regularization
- Basics of Probabilistic Machine Learning and Its Applications in Bioinformatics
- Implementing Support Vector Machines (SVM) and Kernel SVM in Python
- Introduction to Naive Bayes Classifier and Its Use in Bioinformatics
- Decision Tree Classifier and Random Forest Classifier: Theory and Implementation
- Logistic Regression in Bioinformatics: Concepts and Practical Applications
- Introduction to Clustering Algorithms: K-Means and Its Application in Bioinformatics
- Validation of Machine Learning Models: Techniques and Accuracy Metrics
- Theoretical Concepts and Practical Aspects of Machine Learning for Image Analysis in Bioinformatics



MODULE : 9 STATISTICAL METHODS AND TOOLS FOR DATA EXTRACTION AND PREPARATION

- Introduction to Statistical Methods for Data Extraction and Preparation in Bioinformatics
- Exploring Data Characteristics and Distribution: Descriptive Statistics and Data Structures
- Understanding Correlation and Regression Analysis in Bioinformatics
- Probability and Bayes Theorem: Foundations for Statistical Inference
- Sampling Techniques and Distribution Theory in Bioinformatics
- Hypothesis Testing: Concepts and Methods for Data Analysis
- Statistical Tools for Data Management, Analysis, and Visualization in Bioinformatics
- Inferential Statistics: Making Valid Generalizations from Sample Data
- Interpreting Statistical Outputs for Informed Decision Making in Bioinformatics
- Practical Applications: Applying Statistical Methods to Solve Bioinformatics Problems



MODULE: 10 DATA MINING

- Introduction to Data Mining in Bioinformatics: Concepts and Applications
- Understanding Data Warehousing: Life Cycle and Implementation
- Classification and Clustering Techniques for Data Analysis in Bioinformatics
- Outlier Analysis: Identifying Anomalies in Bioinformatics Data
- Overview of Forecasting Techniques in Bioinformatics
- Introduction to Hadoop and its Role in Big Data Analytics
- Exploring the R Language for Statistical Computing and Data Analysis
- Data Analytics Project Life Cycle: Planning, Execution, and Evaluation
- Strategies for Importing Data from Different Databases for Analysis
- Practical Applications: Performing Data Mining from Large Data Sources in Bioinformatics



MODULE : 11 BASICS OF ALGORITHM DEVELOPMENT AND IMPLEMENTATION

- Introduction to Program Design: Principles and Methods
- Basic Structures for Algorithm Development
- Pros and Cons of Efficient and Naïve Algorithms
- Structured Programming Rules
- Divide and Conquer Technique for Problem Solving
- Algorithm Definition in Structured Language
- Algorithm Correctness Verification
- Data Validation and Error Handling in Algorithm Design
- Optimization Techniques for Algorithm Efficiency
- Practical Application of Program Design Principles in Bioinformatics

GTAGGAAACCATGCTCATGTGAAGCTCCCTCGACGAGTCGAGTAG GTAGGAAACCATGCTCATGTGAAGCTCCCTCGACGAGTCGAGTAG GTAGGAAACCATGCTCATGTGAAGCTCCCTCGACGAGTCGAGTAG GTAGGAAACCATGCTCATGTGAAGCTCCCTCGACGAGTCGAGTAG GTAGGACACCCAGCCCGTGTGACGCTCCCTCGACGAGTCGAGTAG GTAGGACACCCAGCCCGTGTGACGCTCCCTCGACGAGTCGAGTAG

MODULE: 12 CHEMINFORMATICS IN BIOINFORMATICS

- Drug Discovery and Development Process: Understanding QSAR Principles
- Introduction to Drug Discovery Process-drug discovery pipeline
- Role of Computational Methods- The significance of computational tools in drug design Examples of computational methods in drug discovery
- Utilizing Biological Databases and Good Clinical Practices (GCP) Standards
- Chemical Structure Visualization-ChemDraw / ChemSketch, Basics of chemical structure visualization
- Visual Representation of Biological Processes and Structures in Data Analysis
- Biomolecules- Properties and function
- Molecular Docking and Molecular Dynamics: Outcomes in Visualization and Evaluation
- Pharmacophore Modeling and Applications
- Pharmacophore Modelling



MODULE: 13 VARIANT CALLING ANALYSIS

- Introduction to NGS and DNAseq
- Basic Terminologies in NGS
- Understanding of SRA database
- Tools installation in Linux for Variation Calling
- Quality control (FastQC)
- Trimming of Reads (Trimmomatic)
- Indexing of Genome (BWA) and Alignment of Reads (BWA)
- Variation calling using GATK
- Variant Effect Prediction(VEP)
- Variation Visualization (IGV)



MODULE: 14 BIOLOGICAL DATA ANALYSIS

- Introduction to NGS and its's applications
- Introduction to RNAseq and it's basic terminologies
- Basic Terminologies in NGS
- Understanding of SRA database
- Tools installation in Linux for Gene Expression analysis
- Quality control (FastQC)
- Trimming of Reads (Trimmomatic)
- Indexing of Genome (STAR) and Alignment of Reads (STAR)
- Normalization of Data (Cufflinks)
- Merging of Data (Cuffmerge) and Differential expression of genes (Cuffdiff)



- Understanding of DEG results
- Annotation of DEG (Uniprot/DAVID)
- Functional and Pathway Enrichment Analysis
- Network Analysis
- Visualization of Differential expressed genes in R (Heatmap & Volcano Plot)

agccgcctgaaat tcccaaagcacaaaat aatcgaatatagggctta ggagacgacaccgagatc aatccatgctgataagga tatgaaagccagcggag

MODULE: 15 INTRDOCTION TO METAGENOMICS

Overview of Metagenomics: Concepts and Applications

Historical Perspective and Evolution of Metagenomics

• Sampling and Sample Preparation Techniques in Metagenomics

- DNA Extraction and Sequencing Technologies for Metagenomics
- Metagenomic Data Analysis Pipeline: From Raw Reads to Biological Insights
- Taxonomic Profiling in Metagenomics: Identifying Microbial Communities
- Functional Annotation and Pathway Analysis in Metagenomics
- Applications of Metagenomics in Biomedical and Environmental Research
- Challenges and Limitations in Metagenomic Data Analysis
- Future Directions and Emerging Trends in Metagenomics



MODULE : 16 GENE EXPRESSION ANALYSIS USING MICROARRAY

- Introduction to Microarray
- Data Downloding using GEO dataset
- Microarray Pipeline using R programming language
- Annotation of DEG
- Functional and Pathway Enrichment Analysis
- Network Analysis
- Visualization of DEG using R plot (Volcano Plot and Heatmap)



MODULE: 17 AWS

- Introduction to Cloud Computing for Bioinformatics: Concepts and Advantages
- Overview of Amazon Web Services (AWS) for Bioinformatics Data Analysis
- Setting up an AWS Account and Access Management for Bioinformatics Workflows
- Deployment Strategies for Bioinformatics Workflows on AWS: EC2, Lambda
- Utilizing AWS Services for Data Storage: S3, EBS, and Glacier
- Leveraging AWS Compute Services for Bioinformatics Analysis
- Implementing Data Analysis Pipelines on AWS: Using Step Functions and Data Pipeline
- Cost Optimization Techniques for Bioinformatics Workloads on AWS
- Security Best Practices for Bioinformatics Data in AWS: IAM Policies and Encryption
- Monitoring and Management Tools for Bioinformatics Workflows on AWS



MODULE: 18 SQL

- Introduction to SQL (Structured Query Language) for Bioinformatics
- Basic SQL Syntax and Data Types: Queries, Statements, and Operators
- Retrieving Data from Relational Databases: SELECT Statements and Filtering
- Manipulating Data in Relational Databases: INSERT, UPDATE, and DELETE Statements
- Joining Tables: Understanding INNER JOIN, LEFT JOIN, and other Join Types
- Aggregating Data: Using GROUP BY and Aggregate Functions in SQL
- Database Management Tasks: Creating, Modifying, and Dropping Tables
- Indexing and Optimization Techniques for Database Performance
- User Management and Security in Relational Databases
- Backup and Recovery Procedures: Ensuring Data Integrity and Availability



MODULE: 19 EMPLOYABILITY SKILLS

MODULE : 20 WORK MANAGEMENT

MODULE : 21 MANAGE YOUR WORK TO MEET REQUIREMENTS

MODULE : 22 WORK EFFECTIVELY WITH COLLOGUES

MODULE : 23 BUILD AND MAINTAIN RELATIONSHIP AT WORKPLACE

MODULE : 24 BUILD AND MAINTAIN CLIENT SATISFACTION





Join Us in Shaping the Future of Bioinformatics!

Embark on a journey of discovery and innovation with our LSSSDC certified Bioinformatics Analyst course. Whether you're a seasoned professional or a recent graduate, our program caters to individuals at all stages of their career. Take the first step towards a rewarding career in bioinformatics and unlock a world of opportunities.

Why Choose Our Course?

- Accredited Certification: Our program is certified by the Life Sciences Sector Skill Development Council (LSSSDC), ensuring that your credentials are recognized and respected by industry professionals worldwide.
- 2. Expert Faculty: Learn from seasoned experts in bioinformatics analysis who bring realworld experience and cutting-edge insights to the classroom.
- 3. Hands-On Learning: Gain practical skills through interactive workshops, case studies, and projects that simulate real-life scenarios, allowing you to apply theoretical knowledge to practical applications.
- 4. **Comprehensive Curriculum:** Covering key topics such as sequence analysis, genomics, proteomics, and more, our curriculum is meticulously crafted to provide you with a well-rounded understanding of bioinformatics principles and techniques.
- 5. Career Support: Receive personalized career guidance and support from our dedicated team, including resume building, interview preparation, and job placement assistance, to help you kickstart your career in bioinformatics.
- 6. **Networking Opportunities:** Connect with fellow aspiring bioinformatics analysts, industry professionals, and mentors to expand your professional network and stay updated on the latest trends and developments in the field.





+91 9310870544, 7888118030, 7843088966

info@dromicslabs.com



www.dromicslabs.com

602/e, W No 3, G/f, L/side, Seqno-m, H
 3/727 Gadaipur, New Delhi , South-West
 Delhi-110030

Thank you!

OUR CERTIFICATIONS & GRANTS













