

1 month  
*Summer*  
INTERSHIP

EARLY BIRD  
DISCOUNT  
OPEN TILL  
6<sup>TH</sup> JUNE

# Decode the Genome: Comprehensive Bioinformatics Bootcamp

Explore the Language of Life  
Through Data This Summer

1 Month | 13 June to 11 July 2025  
Registration Closes: 12 June 2025  
3:30 - 4:30 PM (IST) | Monday to Friday

## MORE DETAILS



## COURSE INSTRUCTORS

Dr. Shikha Mittal, Bioinformatics Trainer, CBIRT  
Dr. Tamanna Anwar, Training Organizer &  
CoFounder, CBIRT

## PROGRAM BENEFITS

- Certificates
- Access to Live Sessions
- Recordings
- Workshop Materials
- Online Technical Support Post Training

## Learning Modules

- 1 Introduction to Bioinformatics & Biological Databases
- 2 Sequence Alignment
- 3 Genomics & NGS
- 4 RNA-Seq Data Analysis
- 5 Bioinformatics with R
- 6 Genome Assembly & Annotation
- 7 Population Genomics & GWAS

## REGISTER NOW



Phone Number:  
**+91 6398142849**



E-mail  
**info@cbirt.net**



Website:  
**www.cbirt.net**

*1 month  
Summer*  
**INTERSHIP**

## **Decode the Genome: Comprehensive Bioinformatics Bootcamp**



### **MODULE I**

June 13-16, 2025

#### **Introduction to Bioinformatics & Biological Databases**

- Basics of Bioinformatics: Scope, Applications, Tools
- DNA, RNA, and Protein Structure Refresher
- NCBI, UniProt, Ensembl, KEGG, PDB, Pfam
- Data retrieval using Entrez, BioMart

#### **Hands-on:**

- BLAST and sequence similarity search
- Using NCBI/UniProt for gene/protein data

### **MODULE 2**

June 17-18, 2025

#### **Sequence Alignment**

- Pairwise & Multiple Sequence Alignment
- Global vs Local Alignments (Needleman-Wunsch, Smith-Waterman)
- MSA Tools: Clustal Omega, MUSCLE, MAFFT

#### **Hands-on:**

DNA/Protein sequence alignment  
Alignment visualization with Jalview

### **MODULE 3**

June 19-24, 2025

#### **Genomics & NGS**

- Overview of NGS platforms
- Quality Control: FastQC, Trimming: Trimmomatic
- Mapping reads: BWA, Bowtie2
- Variant calling: Samtools, Freebayes

#### **Hands-on:**

- NGS analysis pipeline using a small public dataset (DNA-Seq)

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## MODULE 4

June 25-30, 2025

### RNA-Seq Analysis

- RNA-Seq Workflow & Differential Expression Analysis
- Tools: HISAT2, STAR, StringTie, DESeq2, edgeR  
Functional Annotation: DAVID, g:Profiler

#### Hands-on:

- RNA-Seq pipeline on a sample dataset

## MODULE 5

July 1-3, 2025

### Bioinformatics with R

- Introduction to scripting (Bioconductor)
- Data manipulation, plotting, and workflow creation

#### Hands-on:

- Basic scripts for sequence handling and data analysis
- Exploratory plots and automation

## MODULE 6

July 04-07, 2025

### Genome Assembly & Annotation

- De novo vs Reference-based Assembly
- Assemblers: SPAdes, Velvet

#### Hands-on:

Bacterial genome assembly and annotation

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**MODULE 7**

July 08-11, 2025

**Population Genomics & GWAS**

- SNP Calling, Genotype Imputation
- GWAS: TASSEL, RGAPIT, Manhattan plots
- Population structure: PCA, STRUCTURE

**Hands-on:**

- Population genomics pipeline on sample data
- GWAS and structure analysis

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## REGISTRATION PROCESS

**REGISTER NOW**

Registration Form Link:  
<https://forms.gle/qB8CGeb7LGJR79gt7>  
or Scan the QR Code



To register for the workshop, please submit the registration form along with your payment details. We accept multiple payment methods to ensure a convenient and secure registration process.

## REGISTRATION FEES



**STUDENTS (UPTO PHD LEVEL)**

INR-4999



**POSTDOC/RESEARCH SCHOLARS**

INR-5499



**STAFF/FACULTY**

INR-5999



**INDUSTRY PROFESSIONALS**

INR-6499



**INTERNATIONAL PARTICIPANTS**

**STUDENTS  
(UPTO PHD LEVEL)**

USD 65

**POSTDOC/  
RESEARCH SCHOLARS**

USD 70

**STAFF/FACULTY**

USD 80

**INDUSTRY  
PROFESSIONALS**

USD 90

Early Bird Registration (Closes 6 <sup>th</sup> June, 2025)	Students (Upto PhD Level)	Postdoc/Research Scholars	STAFF/Faculty	Industry Professionals
<b>Indian Participants</b>	INR-3999	INR-4499	INR-4999	INR-5499
<b>International Participants</b>	USD 55	USD 60	USD 70	USD 80

## PAYMENT METHODS

**SCAN QR CODE**



[centreofbioinformaticsresearch@cnrb](mailto:centreofbioinformaticsresearch@cnrb)

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