

8. Animals Preserving Techniques
9. Cladistics
10. Systematics and Phenetics
11. 16S ribosomal RNA in phylogenetics
12. Phylogenetic comparative methods – use of evolutionary trees in other studies-comparative biology, adaptation or evolutionary mechanisms.

BSP561 BIOSTATISTICS AND BIOINFORMATICS LAB

Course Outcomes:

After successful completion of the course, students will be able to :

- CO 1. Perform suitable statistical tests for evaluation of data
- CO 2. Make suitable graphical representations of data
- CO 3. Perform statistical tests - t test, F-test, ANOVA
- CO 4. Develop the skill to use search engines, internet tools and databases.
- CO 5. Gain the practical knowledge of restriction mapping and microarray techniques.

Biostatistics

1. Measurement of Central tendencies, mean, median, mode
2. Measures of dispersion range SD, CV&SE
3. Scatter plot, Simple Correlation & Regression, Multiple Correlations
4. Construction of frequency table
5. Theoretical distribution, Binomial, Poisson & normal
6. Statistical inference, normal, t test, chi-square & F test
7. Analysis of Variance

Bioinformatics

1. Introduction to bioinformatics
2. Basic feature of computers; flow charts and problems.
3. Search engines and internet tools.
4. Biological databases
5. Use of databases (e.g. BLAST, FASTA)
6. Restriction mapping
7. Micro array techniques
8. Search engines
9. Web lab viewer and Ras mols