

INTRODUCTION TO PHYLOGENETICS

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This course will introduce the theory behind different phylogenetic tree reconstruction approaches and Hands on sessions using MEGA software. The target audience is students and researchers with no experience in Bioinformatics. Knowledge of computer programming is not required to attend this course.

- 1** Basics of Phylogenetic Tree Reconstruction Approaches
- 2** Different Phylogenetic Tree Reconstruction Methods & Quality Assessment using MEGA
- 3** Maximum Likelihood Phylogeny Reconstruction Using an Example Dataset (RAxML & MEGA)
- 4** Interactive session: Open Questions & Discussion

09:30 AM

Lecture: Basics of Phylogenetic Tree Reconstruction Approaches

10:30 AM

Demo:
Different Phylogenetic Tree Reconstruction Methods & Quality Assessment using MEGA

12:00 PM

Hands-on:
Maximum Likelihood Phylogeny Reconstruction Using an Example Dataset (RAxML & MEGA)

01:30 PM - 02:00 PM

Interactive session: Open Questions & Discussion

FEES

500 INR for Students and Academicians
2500 INR for Non-Academics
100 \$ for Foreign Participants

[More Info](#)

[Register](#)



Bioinformatics Facility

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