

Genome Annotation Winter Internships

Participate in Genome Annotation winter internships to explore genome annotation in coldenvironment species, focusing on identifying cold-tolerant genes, regulatory elements under cold stress, and functional regions in cold-adapted genomes.

Focussed Areas under Genome Annotation Winter Internship

- 1. Genome annotation in cold-tolerant species
- 2. Gene prediction under cold stress conditions
- 3. Annotation of cold-resistant regulatory elements
- 4. Cold-stress epigenetic marker identification
- 5. Comparative genomics in cold-adapted organisms
- 6. Cold-environment multi-omics data for genome annotation
- 7. Transcriptomics of cold-tolerant organisms
- 8. Functional annotation of cold-tolerant genes
- 9. Annotation of alternative splicing events in cold environments
- 10. Genome-wide association studies for cold-resistance
- 11. Epigenetic modification annotation in cold-stressed genomes
- 12. Cold-environment genome annotation for disease research
- 13. Genome annotation of microbial communities in cold environments
- 14. Gene function prediction in cold-adapted species
- 15. Annotation of cold-induced gene expression changes
- 16. Genome annotation of cold-resistant plant and animal systems
- 17. Annotation of genetic variants related to cold stress
- 18. Bioinformatics tools for cold-environment genome curation
- 19. Cold-stress genome annotation for evolutionary studies
- 20. High-throughput sequencing for cold-environment genome annotation

Protocols Covered across various focussed areas under Genome Annotation Winter Internship

- 1. Cold-environment gene prediction and annotation workflows
- 2. RNA sequencing for cold-stress transcript annotation
- 3. Functional annotation using cold-environment bioinformatics tools
- 4. Comparative genomics techniques for cold-adapted species
- 5. Genome-wide association study (GWAS) for cold-resistance genes
- 6. Cold-stress epigenetic marker identification protocols

- 7. Alternative splicing event annotation in cold environments
- 8. Multi-omics data integration for cold-tolerant species
- 9. Gene function prediction workflows for cold-environment genomes
- 10. Genome annotation in microbial communities in cold climates

Duration: 5, 10, 15, 20, and 30 Days

Note: Please cross confirm whether internship slots for this field are available before joining.

Click Here for Genome Annotation Winter Internship Fees

Application Process and Other info