Roberto Biello

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Research on

Demographic and evolutionary processes affecting populations.

I am particularly interested in genetic and genomic approaches to reconstruct population-level patterns and infer processes to study questions associated with local adaptation, genome evolution and wildlife conservation.

Keywords

Population genetics, genomics, molecular ecology, conservation, phylogenetics, comparative genomics, landscape genetics, genome assembly and annotation, bioinformatics.

I joined the course because

I want to improve my phylogenetic knowledge.

I am working on large-scale genome projects and focusing on comparative genomics analyses, so I want to learn novel approaches to properly use genomic resources for genome-scale phylogenetic inference.



