

# ComGenR: Community Genetics Analyses in R

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January 7, 2014

Community Genetics is a field of biology that works at the interface between ecological genetics and community ecology. Being inherently multi-disciplinary, the analytics involved have developed in separate fields. **ComGenR** is intended to synthesize these analytical techniques and facilitate new analytically and computationally driven research tools. Here, we present an introduction to the package, broken into five main sections:

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