**ComGenR: A package for community genetics analyses**

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* Community genetics works at the interface between community ecology and ecological genetics.
* As this field is inherently multi-disciplinary, it utilizes analytics from different fields of study in statistics, ecology and genetics.
* Here we present a package that brings together functions for community genetics analyses with the intent of merging and adding new analyses that are available to researchers in this field.
* The package is available as a stable release on CRAN and as a development version on GitHub