



Donnerstag: 31.01.19 16:00 s.t.

Zentrum für Bioinformatik, Bundesstraße 43, Raum 16

Biotite: A comprehensive framework for computational molecular biology in Python

Analyzing biological sequence and structure data can be tedious work. This talk presents Biotite, a Python package for computational molecular biology, that provides easy-to-use but still efficient methods for analyzing, modifying and visualizing such data.

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Everyone is welcome!