

Biological Sequence Analysis Using The SeqAn C++ Library (Chapman & Hall/CRC Mathematical And Computational Biology) By Andreas Gogol-Döring

By Andreas Gogol-Döring

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Web pages of the Chair of Genome Informatics at the Faculty of Biological Sequence Analysis Using the SeqAn C++ Library, CRC Mathematical & Computational Biology.
<http://www.rahmannlab.de/people/rahmann/publications>
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<http://www.slideshare.net/binal000/rna-seq-data-analysis-a-practical-approach-chapman-hallcrc-mathematical-and-computational-biology-1-e-2015-pdf-unitedvrg>

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William C. Hall, Adonis Moschovakis analysis, occurrence, and biological effects Peter M. Chapman, William J. Adams,

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http://en.wikipedia.org/wiki/Sequence_analysis

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