

RNA Lecture Series Part II: Modern methods to analyze features of ribonucleo-protein particles



H 53 | 17.00 h

Rigel *
0,5 CP
Methode
course

April 15, 2015

The more, the better: Why hidden Markov models are superior to a pairwise sequence alignment

Prof. Dr. Rainer Merkl, Biochemistry II,
Universität Regensburg

April 29, 2015

Computational modeling of RNAi data

Prof. Dr. Rainer Spang, Institute for Functional Genomics,
Universität Regensburg

May 13, 2015

Chipe chipe chip chip - High throughput analyses of immunoprecipitated chromatin

Prof. Dr. Michael Rehli, Internal Medicine III,
Universitätsklinikum Regensburg

May 27, 2015

High throughput methods to explore the architecture and function of mitochondrial RNPs

Dr. Johannes Popow, EMBL Heidelberg

June 10, 2015

Mass Spectrometry in Proteomics: Part II - Applications

Prof. Dr. Rainer Deutzmann, Biochemistry III,
Universität Regensburg

June 24, 2015

NMR-based structural analysis of biomolecules at atomic resolution

Prof. Dr. Werner Kremer, Biophysics,
Universität Regensburg

July 08, 2015

Structure probing of RNAs and RNPs

Dr. Philipp Milkereit, Biochemistry III,
Universität Regensburg

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