

Training Course



A Primer for RNA-Seq Processing, Interpreting and Visualization

RNA-Seq data analysis
Workflow development

Introduction to Galaxy RNA-workbench

Functional RNA analyses
Visualization concepts

Fundamental statistics
Basic file formats

Practical examples
Unix how-to

4th – 6th October 2017

Where?
Who should come?
How to apply?
What are the costs?
Contact:

University of Freiburg (RBC)
Highly motivated experimentalists
destair.bioinf.uni-leipzig.de
120€, incl. materials and meals
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The training is provided and supported by:











