

Upcoming Events

There are some events which could be of interest for you:

❑ **Production Pipelines for Virus Sequencing Data,**

12th–15th September 2017 in Basel.

This tutorial is addressed to practitioners who analyze virus sequencing data including staff of bioinformatics core units as well as developers who want to test their own tools based on the V-pipe open platform.

Deadline: 12th of August, details at <https://www.bc2.ch/2017/program/tutorials/t3/>

❑ **bionection – Partnering Conference for Technology Transfer in Life Sciences,**

17th – 18th October , 2017 in Jena.

This event is aimed at researchers, technology transfer experts, innovative growth companies and small and medium-sized enterprises (SMEs).

Deadline: 31st of August, details at <http://www.bionection.com/program>

*** Manja Marz is invited to this event to introduce the EVBC**

Tools

We have extended in our webpage the list of virus-bioinformatics tools and databases that could be of interest for you. The new tools added:

LRIsScan is a tool for the prediction of long-range interactions in full viral genomes based on a multiple genome alignment. It is able to find interactions spanning thousands of nucleotides.

SPAdes is a tool for assembling genomes and mini-metagenomes from highly chimeric reads.

For details, please have a look at <http://evbc.uni-jena.de/tools/>

News and Announcements

We are planning to apply for an Innovative Training Networks (ITN) - European Commission fund https://ec.europa.eu/research/mariecurieactions/about/innovative-training-networks_en

the guide from 2016 (new call will be opened on the coming September 2017)

http://ec.europa.eu/research/participants/data/ref/h2020/other/guides_for_applicants/h2020-guide-appl16-msca-itn_en.pdf

We would appreciate to learn who is interested to apply with us. Additionally, you propose to meet two days (e.g October) to discuss the idea and might also draft the proposal which has deadline on 10 January 2018. Please, let me know if you are interested until end of August.

Please help us to spread your news by sending us information and news which should be spread.