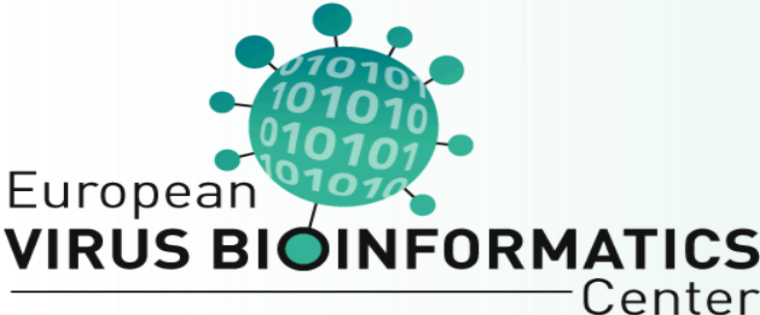


Theoretical and practical metagenomic approaches to viral discovery

Prof. Arthur Gruber - Institute of Biomedical Sciences - University of São Paulo - Brazil
Prof. Manja Marz - Friedrich Schiller University Jena - Germany

FRIEDRICH-SCHILLER-
UNIVERSITÄT
JENA

| Time | Monday - Oct 21 | Tuesday - Oct 22 | Wednesday - Oct 23 | Thursday - Oct 24 | Friday - Oct 25 |
|---------------------------------|--------------------------------------------------------------------------------------------------------------|-------------------------------------------------------------------------------------------------------|--------------------------------------------------------------------|-----------------------------------------------------------------------------------------------|--------------------------------------------------------------------------------------------------|
| 08:30-09:00 | Welcome announcements | Open time for discussion with instructors | Open time for discussion with instructors | Open time for discussion with instructors | Open time for discussion with instructors |
| 09:00-10:15 | Metagenomics - challenges for viral detection and discovery - Prof. Arthur Gruber | Viral genome reconstruction and provirus finding with profile HMMs - Prof. Arthur Gruber | SVM (miRNAs and virus) Prof. Manja Marz | Introduction into RNA world Prof. Manja Marz | LRIsScan/Circos (long-range Interactions of segmented viruses) - Prof. Manja Marz |
| 10:15-10:45 | Coffee break, open time for discussion with instructors | Coffee break, open time for discussion with instructors | Coffee break, open time for discussion with instructors | | Coffee break, open time for discussion with instructors |
| 10:45-12:15 | Profile HMMs of viral markers Prof. Arthur Gruber | Progressive assembly using profile HMMs as seeds and GenSeed-HMM program - Prof. Arthur Gruber | Random Forrest (to detect viral miRNAs) Prof. Manja Marz | Folding algorithm: MacCaskill and Partition functions Prof. Manja Marz | Covariance models Prof. Manja Marz |
| 12:15-13:30 | Lunch time | Lunch time | Lunch time | Lunch time | Lunch time |
| 13:30-15:30 | Retrieving public sequences, multiple sequence alignment - Prof. Arthur Gruber | Finding proviruses in bacterial genomes with e-Finder program - Prof. Arthur Gruber | PCA (for viral host classification) Prof. Manja Marz | RNAfold (to determine the secondary structures of RNA viruses) - Prof. Manja Marz | Infernal to detect viral elements from (meta-)genomic samples - Prof. Manja Marz |
| 15:30-16:00 | Coffee break, open time for discussion with instructors | Coffee break, open time for discussion with instructors | Coffee break, open time for discussion with instructors | Coffee break, open time for discussion with instructors | Coffee break, open time for discussion with instructors |
| 16:30-18:00 | Profile HMM construction with TABAJARA / screening datasets with HMM-Prospector - Prof. Arthur Gruber | Introduction to machine learning methods Prof. Manja Marz | CNN (for viral host classification) Prof. Manja Marz | Vienna RNA Package (for virus host interactions) Prof. Manja Marz | Final discussion and certificate distribution Prof. Arthur Gruber and Prof. Manja Marz |
| 19:00-21:00 | | | | Dinner and social event with all attendees Prof. Arthur Gruber and Prof. Manja Marz | |
| | | | | | |
| Activity: | | Place: | |  | |
| Discussion | | ICB2 - Auditorium | | | |
| Theoretical lecture | | ICB2 - Auditorium | | | |
| Short lecture+Practical session | | ICB4 - Room 5 | | | |
| Practical session | | ICB4 - Room 5 | | | |
| Social event | | To be defined | | | |
| Short lecture+Practical session | | ICB4 - Room 6 | | | |
| Theoretical lecture | | ICB4 - Room 3 | | | |