

BIBI/ZB MED-Workshop
24.10.2019
Uni Bielefeld, CeBiTec, Raum G2-104

Format: 20 (15+5) Minuten je Vortrag

10:00 Ankunft, Begrüßung

10:20 Dietrich Rebholz-Schuhmann (ZB MED - Information Centre for Life Sciences): Using Semantics in Data Analytics, e.g., for the identification of user communities

10:40 Konrad Förstner (ZB MED - Information Centre for Life Sciences): Information Services at ZB MED - Data science for the life science

11:00 Pause

11:30 Sebastian Jünemann (Uni Bielefeld - Genominformatik, de.NBI/BiGi, BIBI/DILS): Service Bioinformatics: Analysis of applications in the field of 16S rRNA amplicon-based phylomics

11:50 Eva Seidlmayer (ZB MED - Information Centre for Life Sciences): Data enrichment with Wikidata for infometrical applications

12:10 Bernd Müller (ZB MED - Information Centre for Life Sciences): Using Pre-Trained Paragraph Vector Models for the Recommendation of Contextual Terms

12:30 Mittagspause

13:30 Alexander Sczyrba (Uni Bielefeld - Computational Metagenomics, de.NBI-Cloud, BIBI): Large scale metagenome analyses in the cloud

13:50 Liren Huang (Uni Bielefeld - Genominformatik, Computational Metagenomics): Cloud-based bioinformatics frameworks for next-generation sequencing data

14:10 Klaus Lippert (ZB MED - Information Centre for Life Sciences): The QuaMedFo Project: patents as external indicator for the evaluation of biomedical research

14:30 Pause

15:00 Silvia Di Giorgio (ZB MED - Information Centre for Life Sciences): GRADitude: a computational tool for the analysis of Grad-seq data

15:20 Nils Kleinbölting (Uni Bielefeld - Genominformatik, de.NBI/BiGi): Analysis of T-DNA insertions by Nanopore sequencing

15:40 Tizian Schulz (Uni Bielefeld - Genominformatik, BIBI/DILS): PLAST: Querying a pan-genome

16:00 Abschluss, Ende der Veranstaltung