## Program for the 5<sup>th</sup> Symposium on Biological Data Visualization

## Friday, July 10

| Triday, July 10 |       |  |  |  |
|-----------------|-------|--|--|--|
| 8:30            | 8:45  | BioVis Welcome, Kay Nieselt  |  |  |
| 8:45            | 9:45  | Keynote Lecture: Visual data science - Advancing science through visual reasoning Speaker: Torsten Möller, University of Vienna, Austria                           |  |  |
| 9:45            | 10:30 | Primer: Four Levels of Visualization, Session Chair: Sean O'Donoghue Speaker: <i>Min Chen, Oxford e-Research Centre, University of Oxford, UK</i>                  |  |  |
| 10:30           | 11:00 | Coffee/Tea Break   |  |  |
| 11:00           | 12:40 | BioVis Papers: Omics Data Visualization, Session Chair: Michel Westenberg  |  |  |
| 11:00           | 11:25 | PathwayMatrix: Visualizing Binary Relationships between Proteins in Biological Pathways, <i>T. N. Dang, P. Murray, <u>A. Forbes</u></i>                            |  |  |
| 11:25           | 11:50 | Extended LineSets: A Visualization Technique for the Interactive Inspection of Biological Pathways, <i>F. Paduano, T. N. Dang, P. Murray, A. Forbes</i>            |  |  |
| 11:50           | 12:15 | Pan-Tetris: an interactive visualisation for Pan-genomes, <u>A. Hennig</u> , J. Bernhardt, K. Nieselt  |  |  |
| 12:15           | 12:40 | BactoGeNIE: A Large-Scale Comparative Genome Visualization for Big Displays,<br>J. Aurisano, K. Reda, J. Leigh, G.E. Marai, A. Johnson                             |  |  |
| 12:45           | 14:15 | Lunch Break with Lunch Box and Beverages   |  |  |
| 14:15           | 15:00 | Primer: <b>How to Visualize Data,</b> Session Chair: Sean O'Donoghue Speaker: <i>Robert Kosara, Tableau Software</i>   |  |  |
| 15:00           | 15:30 | Short Poster Talks Session, Session Chairs: Alex Lex and Michael Westenberg  |  |  |
| 15:30           | 16:00 | Coffee/Tea Break   |  |  |
| 16:00           | 17:15 | BioVis Papers: Visualizing Imaging Data, Session Chair: Jos Roerdink   |  |  |
| 16:00           | 16:25 | Physically-based In Silico Light Sheet Microscopy for Visualizing Fluorescent Brain Models, <u>M. Abdellah</u> , A. Bilgili, S. Eilemann, H. Markram, F. Schürmann |  |  |
| 16:25           | 16:50 | Visual parameter optimisation for biomedical image processing,<br>A.J. Pretorius, Y. Zhou, R.A. Ruddle   |  |  |
| 16:50           | 17:15 | GRAPHIE: Graph based Histology Image Explorer, H. Ding, C. Wang, K. Huang, R. Machiraju  |  |  |
| 17:15           | 17:30 | Poster Fast Forward, Chairs: Alexander Lex, Michel Westenberg  |  |  |
| 17:30           | 19:30 | Poster Presentation and Reception with Drinks and Finger Food  |  |  |
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## Saturday, July 11

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| 8:30   | 9:30    | Keynote Lecture: <b>The challenges of biological information visualization</b> Speaker: <i>Larry Hunter</i> , <i>University of Colorado</i> , <i>USA</i>                          |
| 9:30   | 10:45   | BioVis Papers: Proteins and Structures, Session Chair: Jan Aerts  |
| 9:30   | 9:55    | MoFlow: Visualizing Conformational Changes as Molecular Flow Improves Understanding, W. Rumpf, S. Dabdoub, A. Shindhelm, W. Ray   |
| 9:55   | 10:20   | ReactionFlow: Visualizing Relationships between Proteins and Complexes in Biological Pathways, <i>T.N. Dang, P. Murray, <u>A. Forbes</u>, J. Aurisano</i>                         |
| 10:20  | 10:45   | Integrated visual analysis of protein structures, sequences, and feature data, C. Stolte, K. S. Sabir, J. Heinrich, C. J. Hammang, A. Schafferhans, S. I. O'Donoghue              |
| 10:45  | 11:15   | Coffee/Tea Break  |
| 11:15  | 12:15   | Challenges Session, Session Chair: Kay Nieselt, Sean O'Donoghue   |
| 11:15  | 11:45   | New Advanced Solutions for Genomic Big Data Analysis and Visualization, Speaker: <i>Nacho Medina</i> , Computational Biology Lab, University of Cambridge, UK                     |
| 11:45  | 12:15   | Visualizing 3D Genomes, Speaker: <i>Marc Marti-Renom</i> , Genome Biology Group, Barcelona, Spain   |
| 12:15  | 14:00   | Lunch Break with Lunch Box and Beverages  |
| 14:00  | 15:40   | BioVis Papers: Omics Data Visualization, Session Chair: Nils Gehlenborg   |
| 14:00  | 14:25   | miRTarVis: An Interactive Visual Analysis Tool for microRNA-mRNA Expression Profile Data, <u>D. Jung</u> , B. Kim, R. Freishtat, M. Giri, E. Hoffman, J. Seo                      |
| 14:25  | 14:50   | VisRseq: R-based visual framework for analysis of sequencing data, <u>H. Younesy</u> , T. Möller, M. C. Lorincz, M. M. Karimi, S. J. M. Jones                                     |
| 14:50  | 15:15   | Epiviz: a view inside the design of an integrated visual analysis software for genomics, <i>F. Chelaru</i> , <i>H. Corrada Bravo</i>  |
| 15:15  | 15:40   | XCluSim: A visual analytics tool for interactively comparing multiple clustering results of bioinformatics data, <u>S. L'Yi</u> , B. Ko, D. Shin, YJ. Cho, J. Lee, B. Kim, J. Seo |
| 15:40  | 16:00   | Coffee/Tea Break  |
| 16:00  | 17:00   | Design Contest Presentations, Session Chairs: Eamonn Maguire and Ryo Sakai  |
| 17:00  | 17:30   | Awards Ceremony and Closing Remarks, Liz Marai  |
| 17:30  | 18:30   | Bring your own problem  |
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