

2nd Systems Microscopy Annual Consortium Meeting

23-24th of April, 2013, Leuven

Poster presentations

(venue: Van Hamaele)

No	Title and presenter
1.	“Tracking by structured learning: application to the identification of migratory patterns in cellular screening data”. Alice Schoenauer Sebag , ARMINES
2.	“New CellProfiler Modules for intelligent automated microscopy” Volker Hilsenstein , EMBL - Advanced Light Microscopy Facility, Heidelberg, Germany
3.	“HT-technology for Microscopy-based screening at FIMM” Carina von Schantz-Fant , FIMM, Helsinki, Finland
4.	“Role of vinculin-mediated force transmission in adhesion regulation” Pablo Hernandez Varas , Center for Biosciences, Department of Biosciences and Nutrition, Karolinska Institutet, Sweden
5.	“Automated High Content Imaging of Cell Organelle Morphometry and Function in Cellular Stress Responses” Steven Wink , Division of Toxicology, LACDR, Leiden University, the Netherlands
6.	“Discovering the genetic signal underlying cancer cellular heterogeneity in drug repositioning strategy” Ian Morilla, Aurelio Moya-Garcia and Anibal Bueno Amorós , UMA
7.	“Automated fluorescence correlation spectroscopy to study interactions in living cells” Malte Wachsmuth , European Molecular Biology Laboratory, Heidelberg, Germany
8.	“Integration of systems microscopy and gene expression profiling identifies genes that correlate with breast cancer cell migration and invasion capacity.” Vasiliki-Maria Rogkoti , Division of Toxicology, Leiden Academic Center for Drug Research, Leiden University, Leiden,
9.	“Curvature analysis of clonal boundaries implicates Fascillin III as a regulator of intercellular adhesion in the wing imaginal disc” Joseph Barry , European Molecular Biology Laboratory, Heidelberg, Germany
10.	“Connection of Screen Data with Expression Analysis Data” Sandra Scharaw , European Molecular Biology Laboratory, Heidelberg, Germany
11.	“ Microscopy-based RNAi screen to identify novel candidate metastasis genes” Michiel Fokkelman , Leiden Academic Centre for Drug Research, Leiden University
12.	"Automatic classification of adhesion complexes in migratory cells into protrusion/retraction domains" Jakob Kowalewsky , Center for Biosciences, Department of Biosciences and Nutrition, Karolinska Institutet, Sweden
13.	“WiScan and WiSoft - High-Content Screening at High-Throughput Rate” Yael Paran , IDEA-Bio, Israel

14.	<p>“Characterization of early secretory pathway regulators “ Fatima Verissimo, EMBL, Heidelberg, Germany</p>
15.	<p>"Analysis of tumor tissue microenvironment" Riku Turkki, University of Helsinki, Helsinki, Finland</p>
16.	<p>“Fully automated FRAP of ER-Exit sites” Christian Tischler Advanced Light Microscopy Facility, EMBL, Heidelberg, Germany</p>
17.	<p>“CellH5: A format for data exchange in high-content screening” Christoph Sommer, IMBA</p>
18.	<p>“The role of CMAC component expression levels in cell motility and CMAC organization” Xiaowei Gong, Center for Biosciences, Department of Biosciences and Nutrition, Karolinska Institutet, Sweden</p>
19.	<p>“Multivariate classification of Cell-Matrix Adhesion Complexes” Alexa Kiss, Center for Biosciences, Department of Biosciences and Nutrition, Karolinska Institutet</p>
20.	<p>“The analysis of Golgi <i>de novo</i> biogenesis reveals self-organizing principles of the early secretory pathway” Paolo Ronchi, Cell Biology and Biophysics Unit, EMBL Heidelberg, Germany</p>
21.	<p>“Towards a quantitative life-FRET set-up for Systems Microscopy” Ulrich Berge, Center for Biosciences, Department of Biosciences and Nutrition, Karolinska Institutet, Sweden</p>
22.	<p>“Towards a canonical four dimensional computational model of the dividing human cell to integrate and predict the spatio-temporal dynamics of mitotic proteins” Julius Hossain, EMBL Heidelberg, Germany</p>
23.	<p>“Analysis of Granger causality in Cell Migration based on Systems Microscopy” John Lock, Center for Biosciences, Department of Biosciences and Nutrition, Karolinska Institutet, Sweden</p>