



Oncode - CGC Annual Conference 2019

## Fundamental Cancer Biology

*KIT Royal Tropical Institute,  
Mauritskade 64, Amsterdam*

### Thursday November 7, 2019

8.30 - 9.30	<b>Registration &amp; welcome with coffee</b>	
9.30 - 10.05	Proteome-wide characterization of the targets and mechanisms of action of cancer drugs	<b>Bernhard Küster</b> <i>Technical University of Munich</i>
10.05 - 10.40	The Modular Proteome and its Clinical Significance	<b>Ruedi Aebersold</b> <i>EHT Zurich</i>
10.40 - 11.15	<i>TBA</i>	<b>Bernd Bodenmiller</b> <i>University of Zurich</i>
11.15 - 11.50	<b>Break</b>	
11.50 - 12.25	Multiscale cellular states shape the signaling landscape of cell populations	<b>Lukas Pelkmans</b> <i>University of Zurich</i>
12.25 - 13.00	Regenerative landscape of intestinal organoids	<b>Prisca Liberali</b> <i>Friedrich Miescher Institute for Biomedical Research, Basel</i>
13.00 - 14.10	<b>Lunch and Poster presentation</b>	
14:10 - 14.45	The power of ONE: Immunology in the age of single cell genomics	<b>Ido Amit</b> <i>Weizman Institute of Science, Tel Aviv</i>
14.45 - 15:20	Harnessing innate immunity in cancer therapy	<b>Eric Vivier</b> <i>Marseille-Luminy Immunology Center</i>
15:20 - 15:55	<b>Break</b>	
15:55 - 16:30	Epigenetics and inheritance	<b>Amanda Fisher</b> <i>Imperial College London</i>
16:30 - 17:05	Chromosomal instability as a source of chronic inflammation in cancer	<b>Samuel Bakhom</b> <i>Memorial Sloan Kettering Cancer Center, New York</i>
17.05-17.10	<b>Closing remarks, day 1</b>	

## Friday November 8, 2019

9.00 - 9.30	<b>Welcome with coffee</b>	
9.30 - 10.05	<i>TBA</i>	<b>Karen Knudsen</b> <i>Sidney Kimmel Cancer Center, Thomas Jefferson University, Philadelphia</i>
10.05 - 10.40	Machine learning for cancer diagnostics – finding patterns in Big Data	<b>Jeroen de Ridder</b> <i>Center for Molecular Medicine, University Medical Center Utrecht</i>
10.40 - 11.15	The I-SPY 2 trial, adaptive randomization of targeted drugs optimizes complete response and survival for high risk breast cancer	<b>Laura van 't Veer</b> <i>University of California, San Francisco</i>
11.15 - 11.45	<b>Break</b>	
11.45 - 12.20	Lessons from studies of tumour heterogeneity	<b>Greg Hannon</b> <i>Cancer Research UK Cambridge Institute, Cambridge</i>
12.20 - 12.55	Dissecting the bone marrow microenvironment in malignant hematopoiesis	<b>Rebekka Schneider</b> <i>Erasmus Medical Center, Rotterdam</i>
12.55 - 14.00	<b>Lunch and Poster presentation</b>	
14:00 - 14.35	The Comparative Genomics of Cancer	<b>Duncan Odom</b> <i>German Cancer Research Center, Heidelberg</i>
14.35 - 15:10	Lineage tracing on the organism-wide level	<b>Jan-Philipp Junker</b> <i>Max Delbrück Center for Molecular Medicine, Berlin</i>
15:10 - 15:40	<b>Break</b>	
15:40 - 16:15	Understanding the causes and consequences of polyploidy	<b>Matilde Galli</b> <i>Hubrecht Institute, Utrecht</i>
16:15 - 17.00	How T cells fine-tune their response to tumors	<b>Monika Wolkers</b> <i>Sanquin, Amsterdam</i>
17.00 - 17.10	<b>Poster prize award and closing remarks</b>	
17.10 - 18.00	<b>Drinks</b>	