

Annual Scientific meeting 8 January 2016 Conference Centre De Driehoek, Willemsplantsoen 1c, 3511 LA Utrecht

- 08.30-09.00 Registration and coffee
- 09.00-09.05 Opening remarks (session chair: Edwin Cuppen)
- 09:05-09.20 Liqin Wang (Bernards): A therapeutic intervention based on a vulnerability of drug resistant BRAF mutant melanoma
- 09:20-09:35 Robin van Geel (Schellens) Treatment strategies in advanced BRAFm and KRASm colorectal cancer
- 09:35-09:50 Sasha de Henau (Burgering): Cellular polarization as a model to study spatiotemporal redox regulation
- 09:50-10:05 Jarno Drost (Clevers): Modelling colorectal cancer in human intestinal organoids
- 10:05-10:20 Francis Blokzijl (Cuppen): Mutational patterns in human adult stem cells
- 10:20-10:35 Anna Ressa (Heck): High-throughput (phospho-) proteomics and transcriptomics in colon cancer: an integrated approach to new therapy
- 10:35-10:50 Evert Bosdriesz (Wessels): Unraveling signaling crosstalk in colorectal cancer, a multi-omics approach
- 10:50-11:10 Coffee break

session chair: Puck Knipscheer

- 11:10-11:25 Marcel Smid (Martens) Specific mutational signatures impact the immune response in luminal breast cancer
- 11:25-11:40 Jan Paul Medema: Subtyping colon cancer PDXs
- 11:40-11:55 Hugo Snippert (Bos): Life and death in patient-derived cancer organoids
- 11:55-12.10 Martijn Koppens (Van Lohuizen): Great variety between human colorectal cancer organoids in response to EZH2 inhibition
- 12:10-12:25 Nick Beije (Sleijfer) Optimal treatment sequence in castration-resistant prostate cancer driven by AR-V7 determination in circulating tumor cells

12:25-13:30 Lunch

session chair: Michiel Vermeulen

- 13:30-13:45 Peter ten Dijke Regulation of TGF-beta receptor signaling in breast cancer
- 13:45-14:00 Dylan Mooijman (Van Oudenaarden): Single Cell Hydroxymethylation Reveals Extensive Chromosome-wide Heterogeneity and allows for Lineage Reconstruction
- 14:00-14:15 Arianna Fumagalli (Van Rheenen): Orthotopic transplantation of tumor organoids as new tool to study colorectal cancer progression in vivo
- 14:15-14:30 Hannes Lans (Wim Vermeulen): Oxaliplatin-induced DNA damage response
- 14:30-14:45 Stefano Annunziato (Jonkers): Rapid in vivo testing of tumor suppressors in ILC by CRISPR-Cas9 mediated somatic gene editing
- 14:45-15:00 Nannette Jelluma (Kops): The impact of different levels of chromosomal instability on cancer
- 15:00-15:20 Tea Break

session chair: Lodewyk Wessels

- 15:20-15:35 Joppe Nieuwenhuis (Brummelkamp): Regulators of Histone Acylation
- 15:35-15:50 Jonne Raaijmakers (René Medema): Dissecting the spindle assembly checkpoint using haploid genetics in human cells
- 15:50-16:05 Rik Lindeboom (M. Vermeulen): A systems biology approach to investigate cell fate in intestinal organoids
- 16:05-16.20 Krijn Dijkstra (Voest): Exploiting tumor organoids for immunotherapy in colorectal cancer
- 16:20-16:35 Pavlo Grytsenko (Friedl): p120-catenin dependent glioma-cell networks sustain diffuse brain infiltration
- 16:35-16:45 Bastiaan Evers (Bernards): Comparison of CRISPR, CRISPRi and shRNA technologies for high throughput screens

16:45 Drinks

17.30-18.30 CGC group leader meeting (academiegebouw).

AIO's and Postdocs of the participating groups are particularly invited to join.

The capacity of the room is limited to 120 participants, so please register online on the CGC.nl website <u>www.cancergenomics.nl</u>