ONCODE INSTITUTE TECHNOLOGY SYMPOSIUM	
09.00 - 09.30	**Walk in, registration & coffee/tea**
09.30 - 09.40	**Welcome by Marvin Tanenbaum**
09.40 - 11.10	**Session I:**
09.40 - 10.10	*AI for molecular diagnostics* **Keynote speaker 1 - Jeroen de Ridder; Jeroen de Ridder Group, UMC Utrecht**
10.10 - 10.30	*In vitro dissection of enhancer-promoter compatibilities* Pia Mach, Bas van Steensel Group, NKI
10.30 - 10.50	*Better late than never: embracing appropriate models for proteomics data from complex experimental designs* Koos Rooijers, Monika Wolkers Group
10.50 - 11.10	*Measuring changes in translation efficiency with single-cell ribosome profiling* Michael van Insberghe; Alexander van Oudenaarden Group, Hubrecht Institute
11.10 - 11.40	**Coffee break (30min)**
	Session II:
11.40 - 12.00	*Phosphoproteomics to investigate resistance mechanisms and differences in drug sensitivity between chemonaive and pretreated metastatic colorectal cancer patients* Marin Bont, Jeanine Roodhart group, UMC Utrecht
12.00 - 12.20	*FLAGx – a multiplex imaging approach to spatially resolve transcription of extrachromosomal DNA in single cancer cells* Lukas Frank; René Medema Group, Prinses Maxima Centrum
12.20 - 12.40	*primeTF, a versatile technology for sensitive, multiplexed measurement of TF activity in cultured cells* Max Trauernicht, Bas van Steensel Group, NKI
12.40 - 13.40	**Lunch & networking (1 hour)**
13.40 - 15.20	**Session III:**
13.40 - 14.00	*Distinguishing combinatorial protein post-translational modifications with single-molecule nanopore measurement* Xiuqi Chen, Cees Dekker group at TU Delft
14.00 - 14.20	*Dissecting Transcription Dynamics with Single-Molecule Imaging* Charis Fountas; Tineke Lenstra Group, NKI
14.20 - 14.40	*Development of an in situ functional sequencing method with single-cell spatial resolution* Lars van Roemburg, Miao-Ping Chien Group, Erasmus MC
14.40 - 15.00	*Decoding the sequence requirements for translation initiation* Bram Verhagen; Marvin Tanenbaum Group, Hubrecht Institute
15.00 - 15.20	*Employing scRibo-seq to probe ribosomal states* Kseniia Sarieva; Alexander van Oudenaarden Group, Hubrecht Institute
15.20 - 15.45	**Afternoon break (25min)**
	Session IV:
15.45 - 16.05	*Interaction of transcription and chromatin* Wim Pomp; Tineke Lenstra group, NKI
16.05 - 16.25	*Proteomics across the Central Dogma: A glance into the Mass Spectrometry Toolbox.* Dick Zijlmans, Michiel Vermeulen Group, Radboud University Nijmegen
16.25 - 16.55	*High-throughput multiplexed genome-wide profiling of many chromatin states in single cells* **Keynote speaker 2 - Jop Kind; Jop Kind Group, Hubrecht Institute**
16.55 - 17.00	**Summary and closing by Marvin Tanenbaum**
17.00 - 18.00	**Networking drinks**