

CONTRA Innovative Training Network Workshop
Handling Single Cell Data
 Józefów near Warsaw, Poland, September 9-15, 2018
[Holiday Inn Józefów - Aquila](#)

Sunday, September 9, 2018	
14:00	Hotel Check-in starts
18:00	Conference check-in starts
19:00	Common dinner

PART I: WORKSHOP

Monday, September 10, 2018		
9:00-9:05	Introduction	
9:05-9:50	<i>Single cell RNA sequencing technologies</i> doi.org/10.1016/j.molcel.2015.04.005	Aleksandra Kołodziejczyk
9:50-10:35	<i>Analysis of transcriptional dynamics using single-cell RNA-seq data</i> doi.org/10.1038/s41586-018-0414-6	Peter Kharchenko
10:35-11:05	Coffee break	
11:05-11:50	<i>Spatial Transcriptomics - Bridging histology and RNA sequencing</i> doi.org/10.1126/science.aaf2403	Michaela Asp
11:50-12:35	<i>Charting Tissue Expression Anatomy by Spatial Transcriptome Deconvolution</i> doi.org/10.1101/362624	Jonas Maaskola
12:35-14:00	Lunch break	
14:00-14:45	<i>Unraveling new insights into human immune system landscape one cell at the time</i> doi.org/10.1126/science.aah4573	Alexandra Chloe-Villani
14:45-15:30	<i>Disease-Associated Microglia: A Universal Immune Sensor of Neurodegeneration</i> doi.org/10.1016/j.cell.2018.05.003	Aleksandra Deczkowska
15:30-16:00	Coffee break	
16:00-16:45	<i>Methods and principles for interpreting single-cell variation</i> doi.org/10.1038/nrg3833	Oliver Stegle
19:00	Common dinner	

Tuesday, September 11, 2018		
9:00-9:45	<i>Identification of somatic mutations with single cell DNA sequencing</i> doi.org/10.1016/j.tig.2018.04.003	Peter Park
9:45-10:30	<i>Efficient identification of somatic variants from bulk and single cell experiments</i> dx.doi.org/10.1101/121954	Alexander Schoenhuth
10:30-11:00	Coffee break	
11:00-11:45	<i>Genetic analysis of differentiating cells using single-cell RNA-seq</i> dx.doi.org/10.1098/rstb.2012.0362	Anna Cuomo
11:45-12:30	<i>Probabilistic inference of copy number clonal architecture from low-depth single-cell genomes</i> doi.org/10.1038/nmeth.4140	Adi Steif
12:30-14:00	Lunch break	
14:00-14:45	<i>Single cell analysis of circulating tumor cell clusters</i> doi.org/10.1016/j.cell.2014.07.013	Nicola Aceto
14:45-15:30	Special lecture: <i>introduction to drug resistance mechanisms and metastatic disease, part 2 (continuation from the first training event)</i>	Eike Staub
15:30-16:00	Coffee break	
16:00-16:45	<i>Single cell & big data approaches to understanding genome variation in normal individuals & disease</i> doi.org/10.1101/gr.201160.115	Jan Korbel
16:45-17:15	break	
17:15-18:30	Installation party: <i>the Tutors help the ESRs with problems with installing software necessary for the practical tutorial part.</i>	Anna Cuomo and Ximena Ibarra Soria
19:00	Conference dinner	

PART II: PRACTICAL TUTORIAL
Tutors: Anna Cuomo and Ximena Ibarra Soria

Wednesday, September 12, 2018	
9:00-10:30	DNA sequencing and experimental design.
10:30-11:00	Coffee break
11:00-12:30	Next-generation sequencing data processing.
12:30-14:00	Lunch break
14:00-15:30	Bulk and single-cell RNA-seq experiments and batch correction methods.
15:30-16:00	Coffee break
16:00-17:00	Single-cell RNA-seq data analysis: QC and technical variance.

Thursday, September 13, 2018	
9:00-10:30	Single-cell RNA-seq data analysis: clustering and dimensionality reduction.
10:30-11:00	Coffee break
11:00-12:30	Single-cell RNA-seq data analysis: differential expression and dynamic processes.
12:30-14:00	Lunch break
14:00-15:30	Other types of genomic data
15:30-16:00	Coffee break
16:00-17:00	Projects in groups

Friday, September 14, 2018	
9:00-10:30	Projects in groups
10:30-11:00	Coffee break
11:00-12:30	Projects in groups
12:30-14:00	Lunch break
14:00-15:30	Projects in groups
15:30-16:00	Coffee break
16:00-17:00	Projects in groups

Saturday, September 15, 2018	
9:00-10:30	Presentations of projects
10:30-11:00	Coffee break
11:00-12:30	Presentations of projects
12:30-14:00	Lunch and farewell