## New single cell sequencing methods and applications

## 6th Meeting Single Cell Center Heidelberg (SCC HD)

https://single-cell-center-hd.de

## Jun 13, 2018 14:30-17:40, BioQuant, SR 41, ground floor

Organizers: Eileen Furlong, Karsten Rippe, Josephine Bageritz, Charles Girardot Venue: BioQuant Center, Im Neuenheimer Feld 267, for directions see:

http://malone.bioquant.uni-heidelberg.de/contact/Campus Heidelberg BQ.jpg

Discussion with beer & pizza

Please send an e-mail to Sofie Steinfest (secretary B066@dkfz.de) if you want to attend.

14:30-15:45	Session I (15 min talks + 10 min discussion), chair Karsten
14:30-14:40	Qin Zhang (DKFZ): Full length single cell RNA-seq on different platforms
14:40-15:05	<b>Stefano Secchia</b> (EMBL): Dissecting cell fate determination during Drosophila development by single cell ATAC-seq
15:05-15:30	Carl Herrmann (BioQuant): Exploring cellular variability from multi-omics single-cell data
15:45-16:15	Coffee break with informal discussions
16:15-17:30	Session II (15 min talks + 10 min discussion), chair Eileen
16:15-16:40	Kati Lappalainen (DKFZ): Transcriptome analysis by sequencing single nuclei
16:40-17:05	<b>Josephine Bageritz</b> (DKFZ): How to make a gene expression atlas of a developing tissue
17:05-17:30	<b>Christoph Merten</b> (EMBL): Droplet microfluidics for single cell genomics, antibody screening and personalized medicine
17:30-17:40	Wrap up, announcements topics of next meeting on 12 <sup>th</sup> Sept 2019 (EMBL), 12 <sup>th</sup> Dec 2019 (BioQuant) and 5 <sup>th</sup> Mar 2020 (EMBL).