

Single-Cell and Spalial Biology

Umeå

April 20th Umen Universitet

Eukaryotic Single Cell Analysis Microbial Single Cell Analysis Spatial Transcriptomics Spatial Proteomics

A DAY FULL OF SINGLE CELLS

- From 9:00 to 16:00
- Fika provided
- UB333, Floor 3
- Universitetsbiblioteket

REGISTRATION FORM



bit.ly/SC_Spatial_Seminar_2023





MORE INFO: support@ngisweden.se

Single-Cell and Spatial Biology Seminar

Thursday, April 20th 2023, Umeå

Session 1: Single-Cell Analysis Methods and Facilities

9:00 - 9:05 Introduction to the event

Kristina Benevides, Project Coordinator - NGI

9:05 - 9:35 Overview of Eukaryotic Single Cell RNAseq

Anja Mezger, Head of Applications Development - NGI

10X Genomics protocols

- Protocols offered and workflow
- o Sample requirements and tips

SmartSeq3

- o Protocols offered and workflow
- o Sample requirements and tips

9:35 - 10:00 Microbial Single-Cell Facility

Henrik Gezelius, Acting Head at NGI-Single Cell Genomics - Uppsala

- Protocols offered
- o Sample requirements and tips

Break

10:20 - 10:40 Data analysis

Anastasios Glaros, Bioinformatician - NGI

10:40 - 11:30 Q&A

11:30 - 13:00 - Lunch break

Session 2: Spatial Analysis at the SB platform

13:00 - 13:05	Introduction to the session Renuka Kudva, Project Coordinator - NGI	
13:05 - 13:15	Spatial Biology Platform overview Anja Mezger, Head of Applications Development - NGI	
13:15 - 13:30	FF - Spatially Resolved Transcriptomics Xesús Abalo, Project Coordinator - NGI	
	0 0 0	Tissue Optimisation Sample requirements and tips H&E or IF + ST
13:30 - 13:50	FFPE - Spatially Resolved Transcriptomics Jörg Bachmann, Research Engineer - NGI	
	0 0 0	Tissue Adhesion CytAssist Sample requirements and tips; 11x11 mm
Break		
14:10 - 14:30	Spatial Proteomics Carolina Oses, Researcher, Spatial Proteomic Unit - SciLifeLab	
	0 0	Protocols offered Sample requirements and tips
14:30 - 14:45	Data anal TBA , NBIS	ysis
14:45 - 15:30	Q&A	

Concluding remarks