





## Join the IMC Summit on October 21, 2021

Learn how you can gain research insights by deciphering and uncovering the complex secrets of spatial biology using Imaging Mass Cytometry.

Whether you are just starting to think about using proteomic approaches to deeply interrogate the tissue microenvironment, are performing your first high-multiplex experiments or are a veteran user of Imaging Mass Cytometry $^{\text{M}}$  (IMC), this event has something for you. We are excited to share the agenda of what is sure to be an inspiring and thought-provoking half a day.

October is Breast Cancer Awareness Month, and we'll kick off with two excellent presentations on the use of IMC in research related to breast cancer. To support work aimed at combating this disease, Fluidigm will make a \$5 donation for every IMC Summit attendee,\* with the proceeds going to charities selected by our two speakers.

Here's the exciting agenda! Register today at fluidigm.com/imcsummit2021

## Speakers and Topics



Welcome to the IMC Summit 2021

Welcome Address
Chris Linthwaite
President and CEO, Fluidigm

Moderator: Michelle Macpherson

Director of Product Management, Imaging Mass Cytometry, Fluidigm

## Making an impact in breast cancer research



## Bernd Bodenmiller, PhD

Dual Professor for Quantitative Medicine, University of Zurich and Institute for Molecular Health Sciences, ETH Zurich



Capturing diversity in tumor microenvironments: Biological determinants of breast cancer disparities

#### Melissa Davis, PhD

Assistant Professor of Cell and Developmental Biology, Scientific Director of the International Center for the Study of Breast Cancer Subtypes, Weill Cornell Medicine

Obesity, inflammation and cancer

Daniela Quail, PhD

Assistant Professor, Department of Physiology, Rosalind and Morris Goodman Cancer Research Centre, McGill University Forum and Time

Auditorium 6:00 PT | 9:00 ET 14:00 BST | 15:00 CET

Auditorium 6:10 PT | 9:10 ET 14:10 BST | 15:10 CET

Speakers and Topics		Forum and Time
	IMC Summit Keynote Presentation Using IMC to understand the neuroinflammation of severe COVID19 infection Bertram Bengsch, MD, PhD Group Leader, Department of Internal Medicine II, University Medical Center Freiburg	Auditorium 7:10 PT   10:10 ET 15:10 BST   16:10 CET
	Break Visit The Promise of IMC and Product Kiosks	Kiosks 7:50 PT   10:50 ET 15:50 BST   16:50 CET
	Ask an FAS: Sample preparation and panel design  Leigh-Anne McDuffus  Field Applications Specialist, Proteomics, Fluidigm	Ask an FAS 7:50 PT   10:50 ET 15:50 BST   16:50 CET
	Expert User Insights: Roundtable Discussion  Moderator: Noah Saederup, PhD  Senior Product Manager, Mass Cytometry Reagents, Fluidigm  Panelists:  Dan Traum  Research Specialist, University of Pennsylvania  Marieke E. Ijsselsteijn  PhD Candidate, Leiden University Medical Center  Anthony Colombo  PhD Student in Biostatistics at University of Southern California, in collaboration with Cedars-Sinai Graduate School of Biomedical Sciences	Auditorium 8:05 PT   11:05 ET 16:05 BST   17:05 CET
	Ask an FAS: ROI selection and considerations in selecting a data analysis pipeline  Eric Swanson, PhD  Senior Field Applications Specialist, Proteomics, Fluidigm	Ask an FAS 8:15 PT   11:15 ET 16:15 BST   17:15 CET
	Introducing Maxpar® OnDemand  Clinton Hupple  Product Manager, Proteomics, Fluidigm	Auditorium 8:35 PT   11:35 ET 16:35 BST   17:35 CET
	2021 Publication Spotlights  The spatial landscape of lung pathology during COVID-19 progression  Andre Rendeiro, PhD  Postdoctoral Associate in Computational Biomedicine, Institute for Computational Biomedicine and Englander Institute	Auditorium 8:50 PT   11:50 ET 16:50 BST   17:50 CET

for Precision Medicine, Weill Cornell Medicine

## Joint Speakers:



Immunogenomics of colorectal cancer response to checkpoint blockade: Analysis of the KEYNOTE 177 trial and validation cohorts

#### Michele Bortolomeazzi

PhD Student, Cancer Research UK King's Health Partners Centre, Cancer Systems Biology Group, King's College London, Francis Crick Institute

## Lucia Montorsi, PhD

Postdoc in Peter Gorer Department of Immunobiology, King's College London, Francis Crick Institute

Speakers and Topics		Forum and Time
	Ask an FAS: Getting the most out of your data analysis  Wendell Smith, PhD  Field Applications Specialist II, Proteomics, Fluidigm	Ask an FAS 9:20 PT   12:20 ET 17:20 BST   18:20 CET
	Getting Started: Panel Discussion  Moderator: Alex Cherkassky, MS, MBA  VP, Franchise Head, Spatial Tissue Imaging and Mass Cytometry, Fluidigm  Panelists:  Hiran Ravichandran, MS  Research Associate, Weill Cornell Medicine	Auditorium 9:20 PT   12:20 ET 17:20 BST   18:20 CET
	Aida Meghraoui-Kheddar, PhD, PharmD Senior Postdoc, Institute of Molecular and Cellular Pharmacology Corrine Ramos, MBA, PhD Innovation & Marketing Director, ImaBiotech	
	Ask an FAS: Dual-mode system use, calibration and other instrument operation  Ron Cohn, MS  Senior Field Applications Specialist, Proteomics, Fluidigm	Ask an FAS 9:55 PT   12:55 ET 17:55 BST   18:55 CET
	Break	Auditorium 10:00 PT   13:00 ET 18:00 BST   19:00 CET
	Data Analysis Session I: The importance of cell segmentation in the data analysis pipeline	Data Analysis Sessions 10:15 PT   13:15 ET 18:15 BST   19:15 CET
	Notes from the bench: IMC data analysis from a biologist's perspective  Hartland Jackson, PhD  Investigator, Lunenfeld-Tanenbaum Research Institute	
	Joint Speakers:  Mass Cytometry and single cell segmentation of lung	



tissue samples

Jay Hirota, PhD

Canada Research Chair in Respiratory Mucosal Immunology, Assistant Professor of Medicine, McMaster University

Kjetil Ask, PhD

Associate Professor, Firestone Institute for Respiratory Health, McMaster University



Data Analysis Session II: Visiopharm® high-multiplex software and a new import function

Build a strong foundation: Powerful cell segmentation with Visiopharm's advanced phenotyping software

Brit Boehmer, PhD

Technical Sales Specialist, Visiopharm

Data Analysis Sessions 10:55 PT | 13:55 ET 18:55 BST | 19:55 CET

## **Speakers and Topics**



# Data Analysis Session III: An operationalized pipeline with IF data merging, HALO® and ImaCytE

Integrating IMC into an initial QC and deep multi-omic phenotyping workflow: A tissue-sparing approach to maximize the amount of remaining clinically collected tissues for other spatial technologies

M. Caleb Marlin, PhD

Project Manager, Arthritis and Clinical Immunology, Oklahoma Medical Research Foundation



19:15 BST I 20:15 CET

Forum and Time



Data Analysis Session IV: Data analysis methods in emerging applications

Barcoded IMC assays for an efficient spatial profiling of 3D spheroids

Vito Zanotelli, PhD

Recently from the Bodenmiller Lab, Department of Quantitative Biomedicine, University of Zurich

Data Analysis Sessions 11:35 PT | 14:35 ET 19:35 BST | 20:35 CET



Come celebrate 100-plus Imaging Mass Cytometry publications and hear how IMC is advancing our understanding of health and disease.

Learn more and register today at fluidigm.com/imcsummit2021

\*Donation of up to \$3,000 based on \$5 for each attendee who attends the summit for at least 1 hour

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