FLOW CAPS

U.S. NIH Campus Natcher Auditorium Bethesda, MD. 21 - 22 September 2010



Flow Cytometry:

Critical Assessment of Population Identification Methods (FlowCAP)

To advance the development of computational methods for the identification of cell populations of interest in flow cytometry data

http://flowcap.flowsite.org

Flow CAP Organizing Committee

- Ryan Brinkman, British Columbia Cancer Agency
- Raphael Gottardo, Fred Hutchison Cancer Research Center
- Richard Scheuermann, University of Texas Southwestern Medical Center
- Jill Schoenfeld, Tree Star, Inc.



Sponsored by the National Institute of Allergy and Infectious Diseases

FlowCAP Summit 2010

Date: September 21st & 22nd, 2010

Time: 8:30AM-5:50 PM (Tuesday) and 9:00AM-3:30 PM (Wednesday)

Location: Natcher Conference Center, NIH Campus

45 Center Drive

Bethesda, Maryland 20892

AGENDA

DAY ONE (Tuesday):

8:30-9:00AM	Registration
9:00-9:10AM	Importance of FlowCAP from the DAIT Perspective Dan Rotrosen Division of Allergy Immunology and Transplantation, NIAID
9:10-9:40AM	FlowCAP: history, goals and design of FlowCAP-I Richard H. Scheuermann U.T. Southwestern Medical Center
9:40-10:10AM	Rapid Cell Population Identification in Flow Cytometry Data Nima Aghaeepour British Columbia Cancer Agency
10:10-10:40AM	Automatic determination of the number of mixture components in Flow Cytometry with Variational Bayes Hannes Bretschneider University of British Columbia
10:40-11:00AM	BREAK
11:00-11:30AM	GPU accelerated Bayesian mixture models for FCM analysis Cliburn Chan Duke University
11:30-12:00PM	flowMerge: Merging mixture components for automated gating of flow cytometry data Greg Finak Clinical Research Institute of Montreal
12:00-12:30PM	Flow Cytometry Data Assessment with L2 Discrepancy Learning Process: Analysis and Visualization Faysal Khettabi British Columbia Cancer Agency
12:30-1:30PM	LUNCH

1:30-2:00PM	On the use of NMF and curvHDR to cluster flow cytometry data Joe Maisog Medical Numerics, Inc. / Georgetown University MedicalCenter
2:00-2:30PM	TBD (FLAME) Geoff McLauchlan University of Queensland
2:30-3:00PM	SWIFT: Scalable Weighted Iterative Flow-clustering Technique Iftekhar Naim University of Rochester
3:00-3:30PM	Self-organizing Maps for Flow Cytometry Data Analysis Radina Nikolic British Columbia Institute of Technology (BCIT)
3:30-3:50PM	BREAK
3:50-4:20PM	FLOCK: a density-based clustering method for automated identification and comparison of cell populations in high-dimensional flow cytometry data Yu Qian U.T. Southwestern Medical Center
4:20-4:50PM	Support Vector Machines for classification of flow data John Quinn TreeStar, Inc
4:50-5:20PM	Misty Mountain - A Parallel Clustering Method. Application to Fast Unsupervised Flow Cytometry Gating Istvan Sugar Mt. Sinai School of Medicine
5:20-5:50PM	SamSPECTRAL: Efficient spectral clustering on flow cytometry data Habil Zare British Columbia Cancer Agency
5:50PM	ADJOURN

DAY TWO (Wednesday):

9:00-10:00AM **KEYNOTE ADDRESS**

Coherent Single Cell Analysis in the 21st Century: ROFLMAO

Mario Roederer

National Institutes of Health

10:00-10:20AM *BREAK*

10:20-11:20AM FlowCAP-I Results

Ryan Brinkman

11:20-12:00PM FlowCAP-I Debrief: What worked and what didn't work

Richard H. Scheuermann

U.T. Southwestern Medical Center

12:00-1:00PM **LUNCH**

1:00-1:30PM Comparative Metrics: Measuring the quality of a classification method

without a known ground truth

Adam Triestar TreeStar, Inc

1:30-1:50PM FlowCAP-II: comparative metrics

Jill Schoenfeld TreeStar, Inc

1:50-2:20PM FlowCAP-II: Dataset classification and identification of important gaps in

datasets used for FlowCAP-I Richard H. Scheuermann

U.T. Southwestern Medical Center

2:20-2:40PM FlowCAP-II: Beyond classification – other opportunities for FCM data

analysis

Ryan Brinkman

British Columbia Cancer Agency

2:40-3:00PM FlowCAP-II: design, funding

Ryan Brinkman

British Columbia Cancer Agency

3:00-3:30PM **Wrap up**

Richard H. Scheuermann

U.T. Southwestern Medical Center

3:30PM ADJOURN