

ABRF 2024

Annual Meeting

*Preparing today's cores
for tomorrow's needs*



Minneapolis, MN
April 21-24

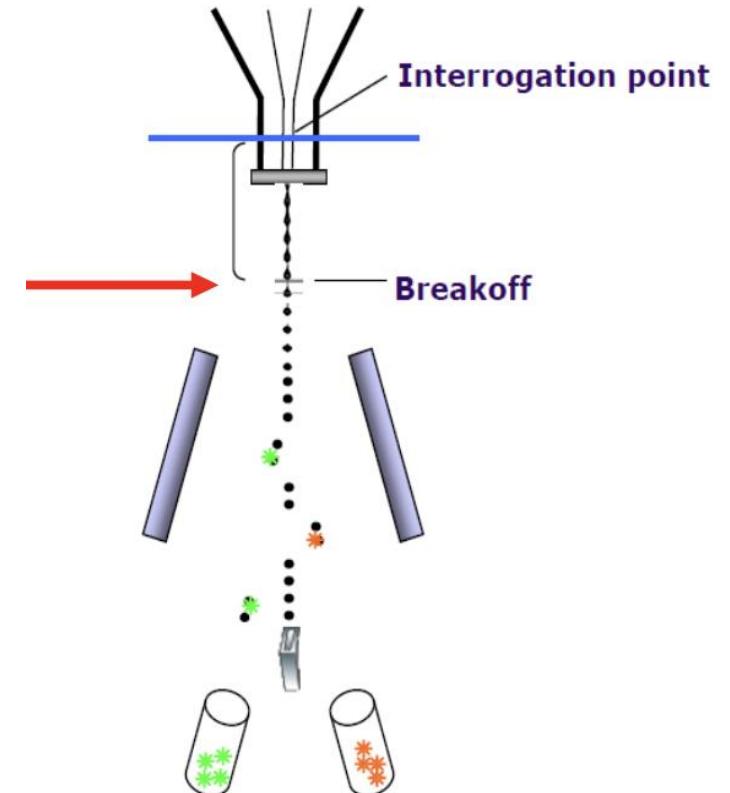
ABRF FCRG Drop Delay Study Year 2

ABRF Meeting – Monday April 22, 2024



Drop Delay

- Drop delay is the time needed by an event to travel from the interrogation point to the droplet breakoff point
- If the delay is measured incorrectly, the particle of interest will not be contained in the sorted drop, reducing the recovery of the target particles



Evolution of Drop Delay

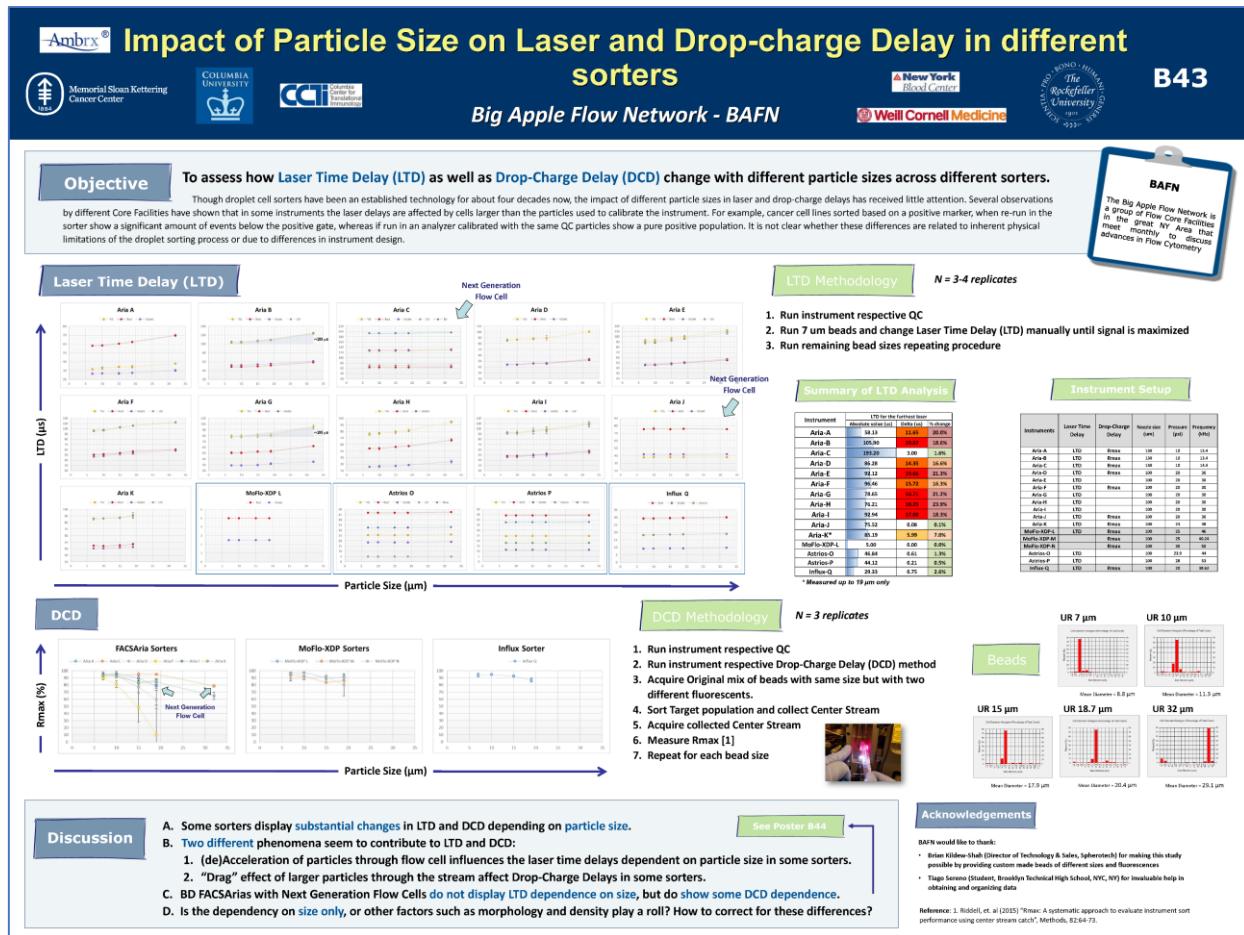
- Early cell sorters required a manual drop delay process = time consuming
- This evolved to an automated drop delay calibration:
 - Standardized beads
- Now - fully automated wizard driven instrument sort set up
 - Includes drop delay calibration
- New fully automated systems:
 - Streamline sort set up and drop delay calibration
 - **BUT** block the ability of the sort operator from adjusting the delay
- Why is changing drop delay necessary?
 - Allow sorting of a wider variety of sample types
 - Increases ability to sort larger cells on instruments with limited nozzle diameter options



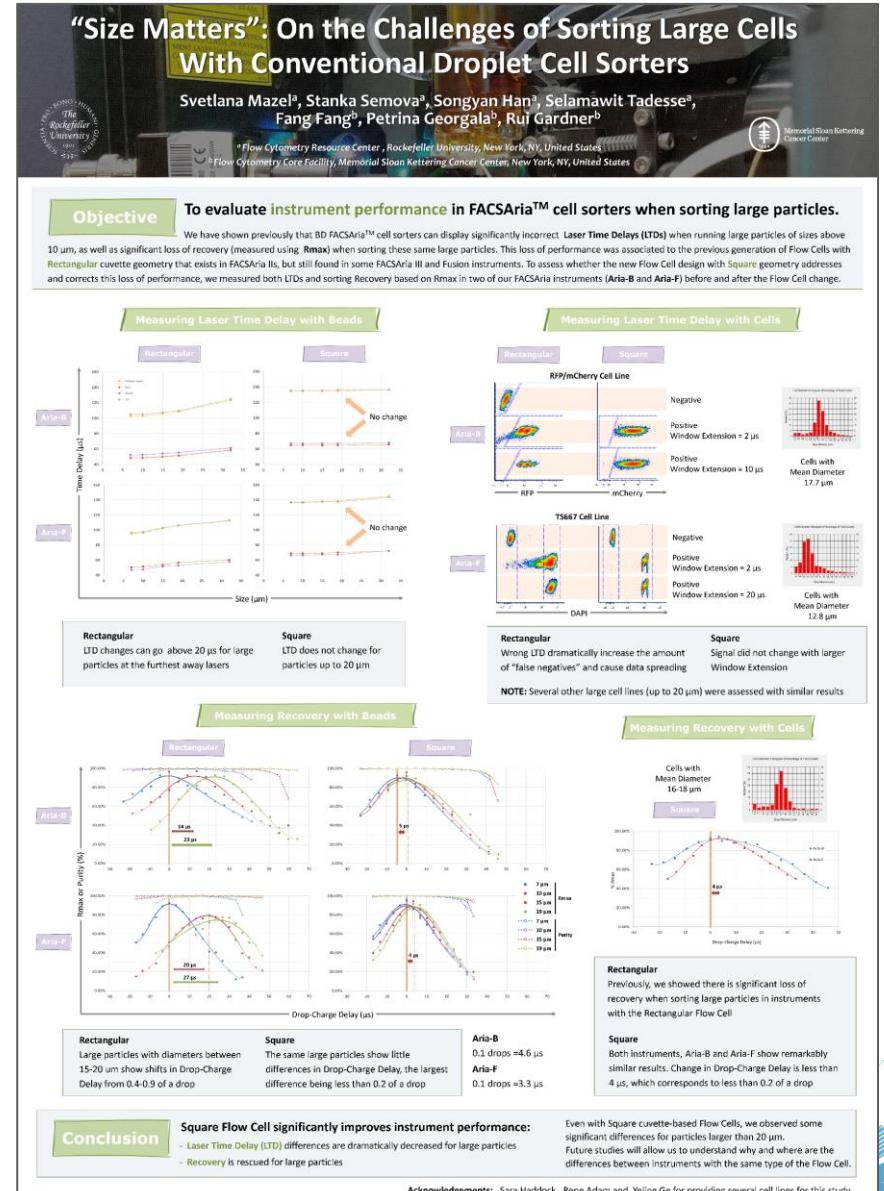
What factors influence Drop Delay?

- Drop delay is influenced by several parameters including:
 - ✓ Temperature
 - ✓ Sheath Pressure
 - ✓ Drop drive settings
 - ✓ Fluidics design
 - ✓ Particle size
- Of these, particle size is the most variable among sorts on a given instrument
- Increased size can subtly alter drop delay before causing noticeable deterioration of sort streams

Previous drop delay testing



Rockefeller University, MSKCC, and the BAFN group

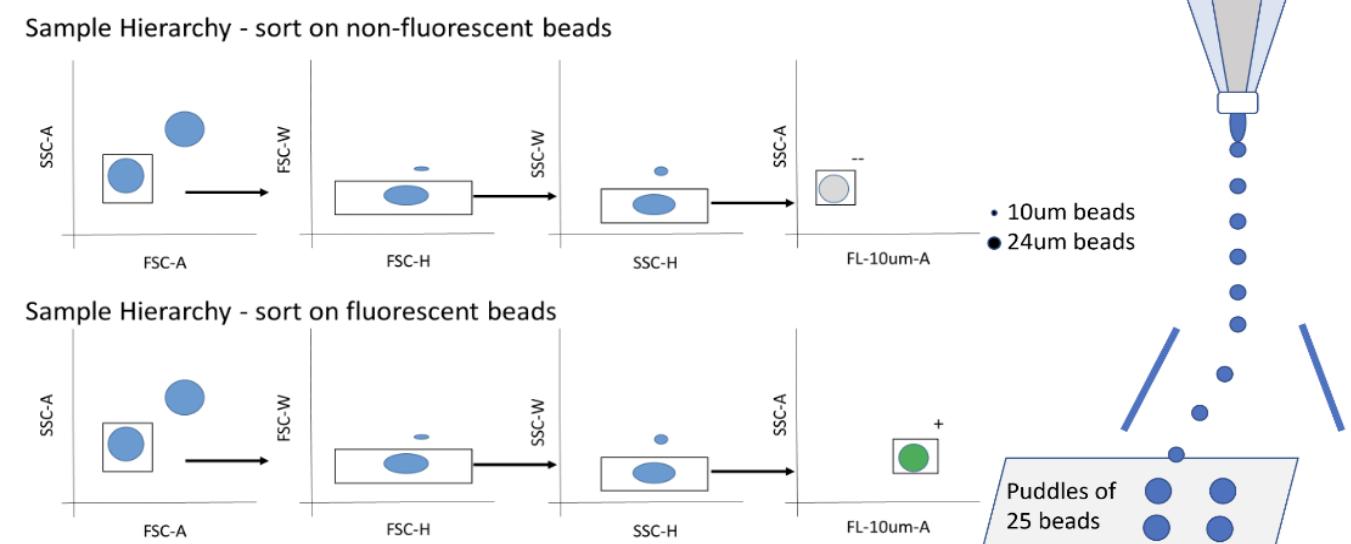


ABRF FCRG Drop Delay Study – Year 1 recap

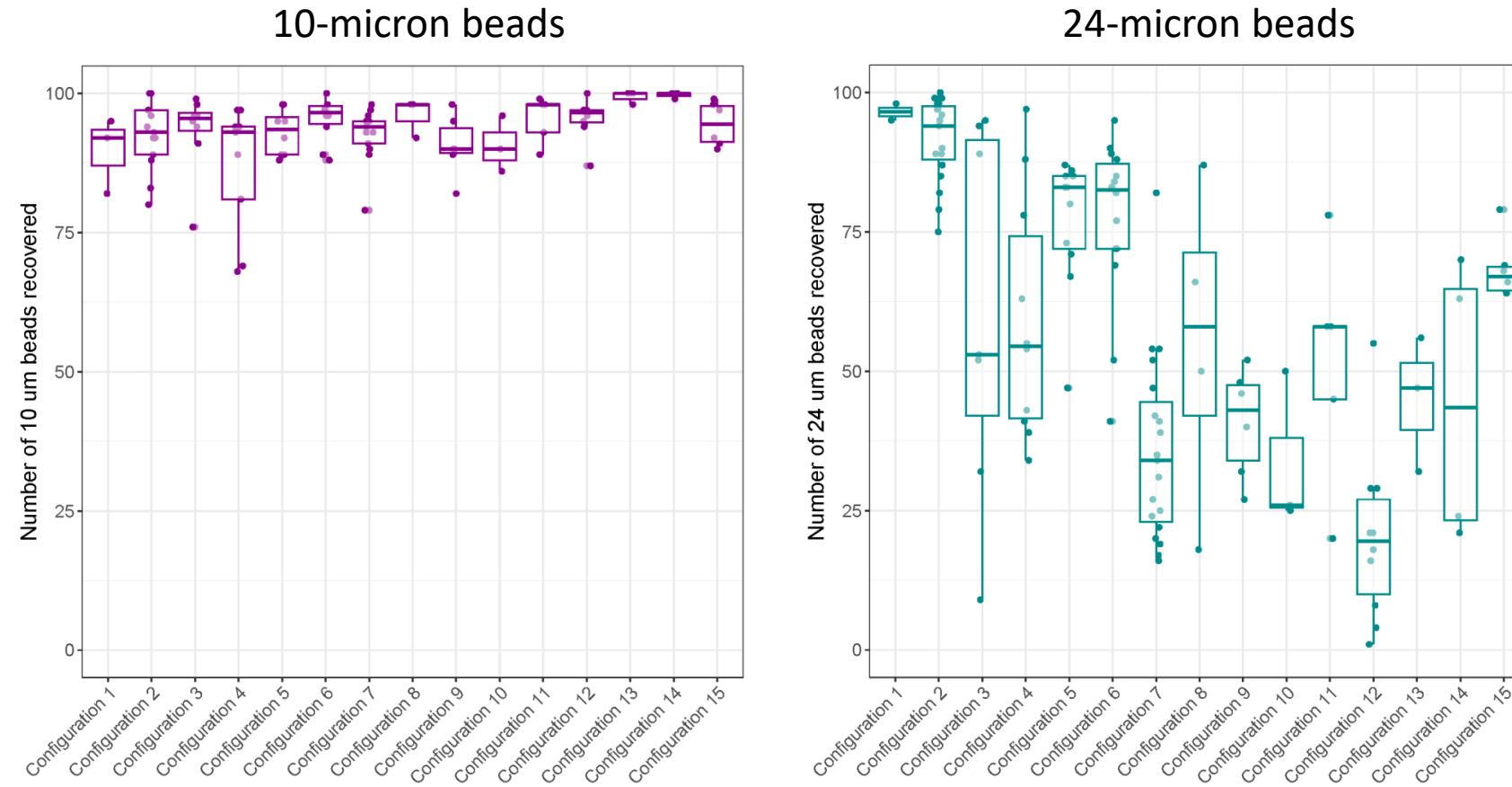
- First Year: Measure automated drop delay accuracy with different particle sizes across commercially available sorters
 - ✓ 10- and 24-micron beads
 - ✓ Beads decrease variability introduced when working with cells
 - ✓ 11 institutions and 10 cell sorter models, 15 different configurations
 - ✓ For consistency, a 100-micron orifice was used for each configuration

Materials & Methods

- 10 cell sorter models, using a 100-micron orifice for sorting
- 10- and 24-micron beads:
 - ACURFP2.5-250-5, ACURFP20-100-1, PPX-100-10, PPX-200-10, Spherotech, Inc.
- Drop delay was set using the automated drop delay for each configuration with a 100-micron orifice
- 10- or 24-micron beads were sorted in four puddles of 25 beads each, using a one-drop envelope and set to maintain sort purity
- Sorting was repeated on two additional days for three total replicates



ABRF FCRG Drop Delay Study – Year 1 Results



ABRF FCRG Drop Delay Study – Year 2

Second year: increase robustness of study

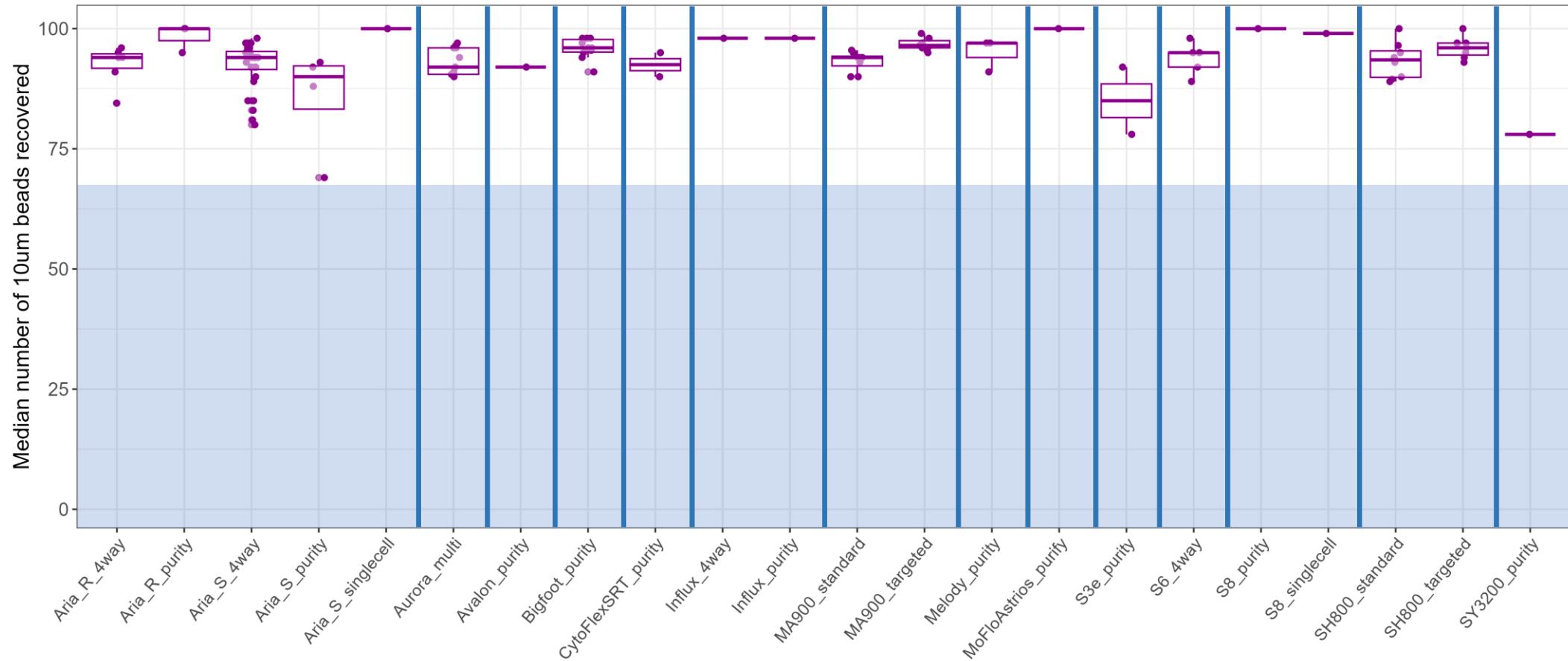
- Include more sorters
 - Sorters and configurations are named
- Similar materials and methods
 - Use fluorescent beads only - ACURFP2.5-250-5, ACURFP20-100-1, Spherotech Inc. – larger beads were closer to 25-micron diameter

ABRF FCRG Drop Delay Study – Year 2 Results

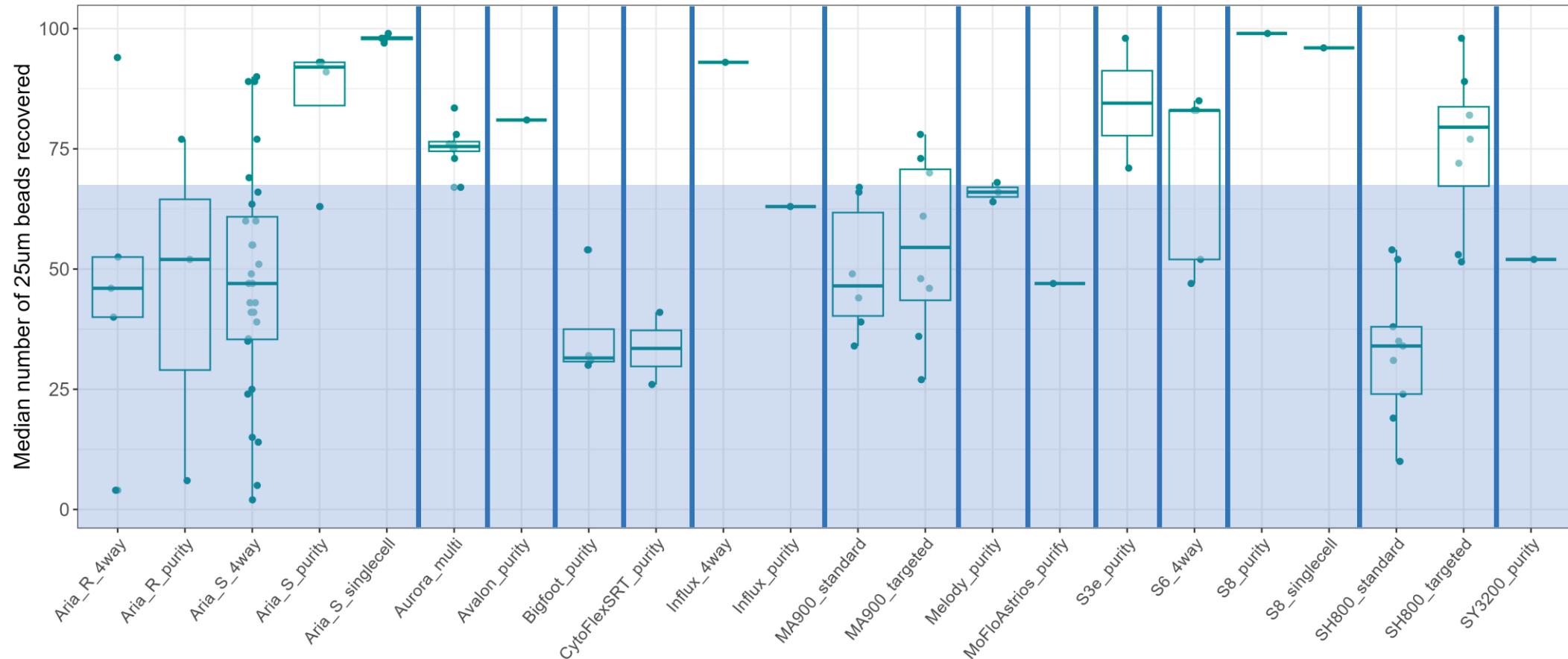
Before the big reveal

- Wanted to include at least 3 sorters for each model
 - In some cases, this was not possible
- To account for this, the FCRG is interested in conducting a continuation of the study for Year 3
 - Increase robustness of data for sorters of $n \leq 3$

ABRF FCRG Drop Delay Study – Year 2 results – 10-micron beads

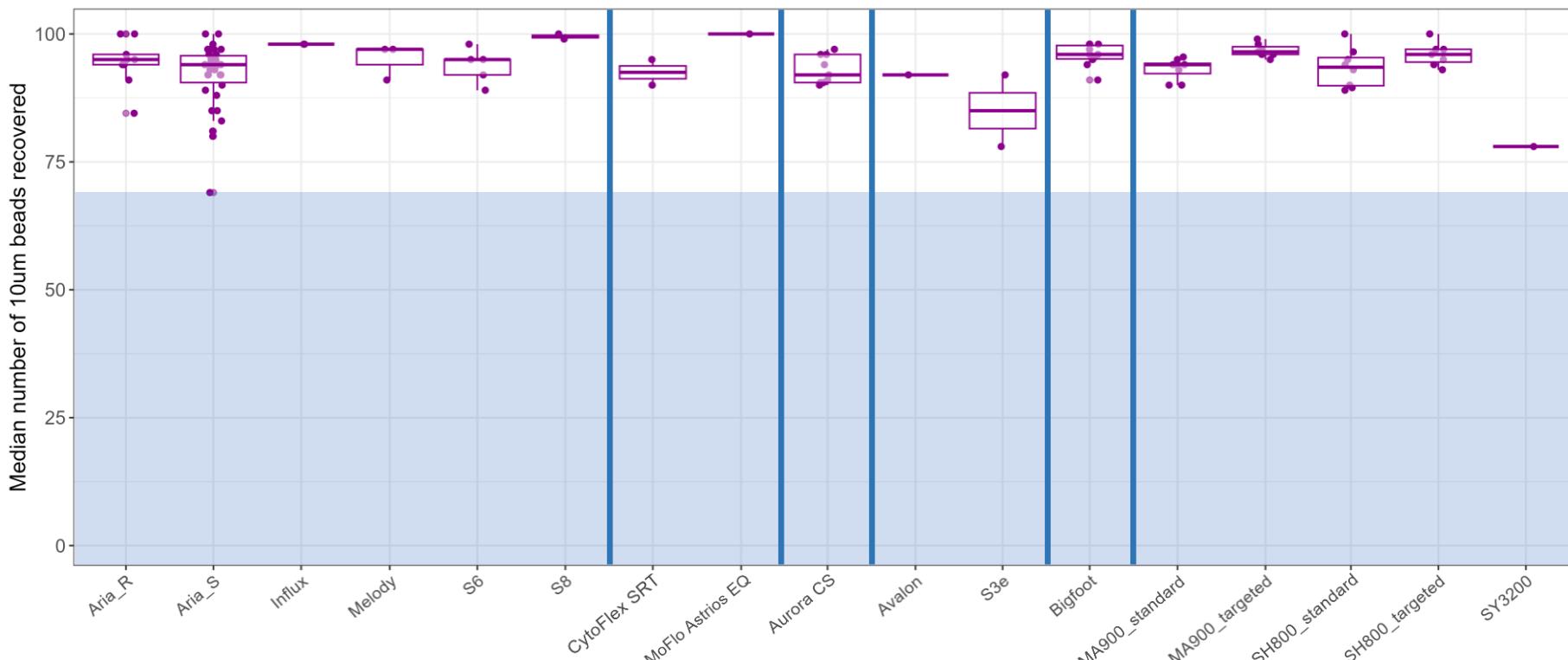


ABRF FCRG Drop Delay Study – Year 2 results – 25-micron beads



Data consolidated by instrument

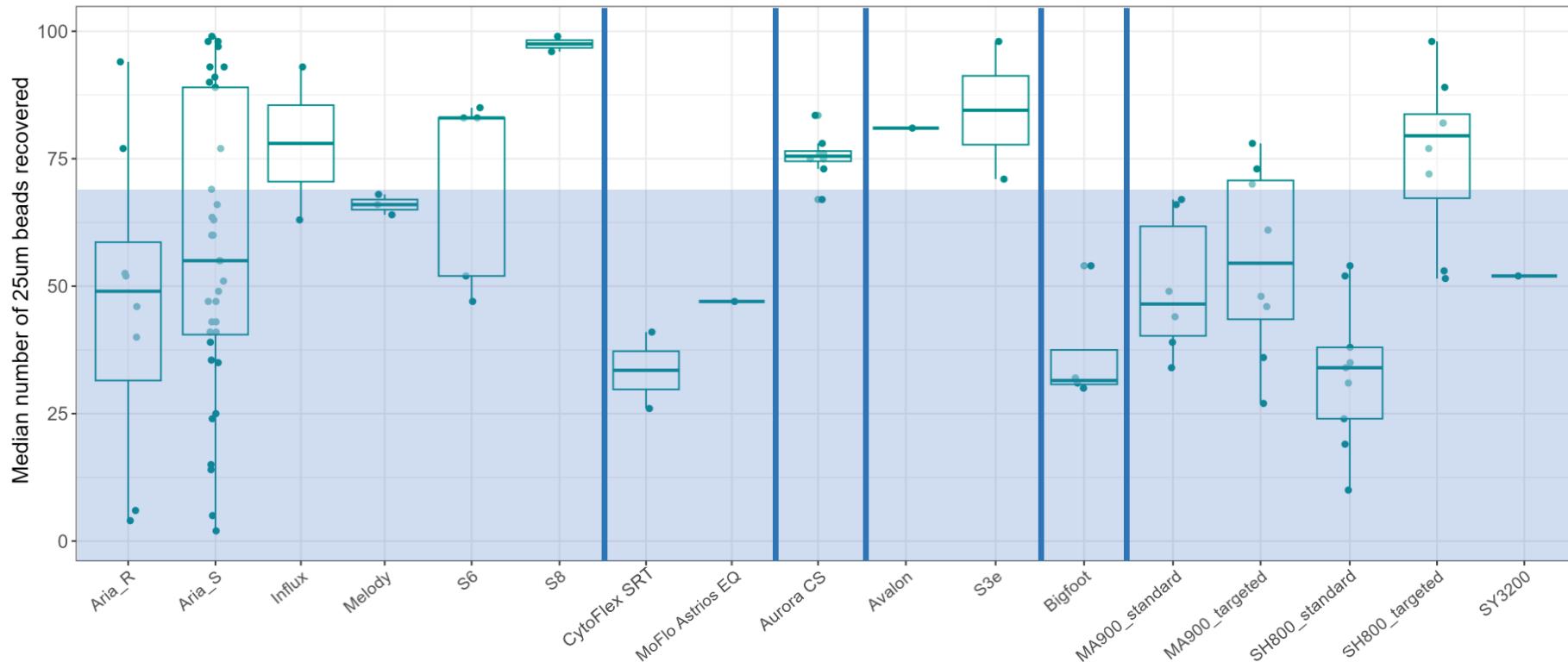
Consolidated 10-micron data



Number of each instrument:

Aria-R - 10
Aria-S - 27
Influx - 2
Melody - 2
S6 - 3
S8 - 1
CytoFLEX SRT - 2
MoFlo Astrios EQ - 1
Aurora CS - 7
Avalon - 1
S3e - 1
Bigfoot - 4
MA900-S - 7
MA900-T - 7
SH800-S - 7
SH800-T - 7
SY3200 - 1

Consolidated 25-micron data

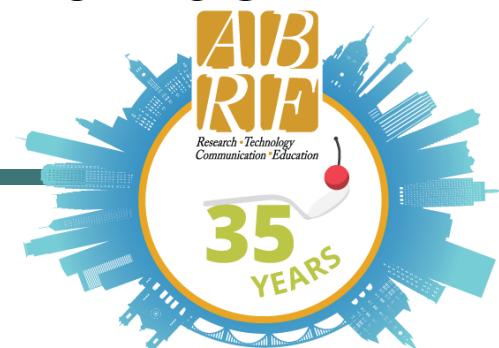


Number of each instrument:

Aria-R - 10
Aria-S - 27
Influx - 2
Melody - 2
S6 - 3
S8 - 1
CytoFLEX SRT - 2
MoFlo Astrios EQ - 1
Aurora CS - 7
Avalon - 1
S3e - 1
Bigfoot - 4
MA900-S - 7
MA900-T - 7
SH800-S - 7
SH800-T - 7
SY3200 - 1

Conclusions – Year 2

- Like Year 1, 10-micron beads exhibited good recovery and 25-micron exhibited poor recovery across instruments
 - More variation in 25-micron data in year 2; number of instruments increased
- Automated drop delay settings were accurate for 10-micron but not 25-micron bead sorting across most sorters
- Some variability may be due to a 1-drop vs. 2-drop sort envelope
 - Study participants were instructed to use a 1-drop envelope, but this did not always occur
- Need additional data for sorters where the sorter number was $n \leq 3$



Future Directions – Year 3

- Test additional sorters where $n \leq 3$
- On sorters where changing drop delay is possible, test whether incremental changes improve recovery of 25-micron particles
 - This has been shown previously for Aria instruments by the BAFN group: https://cdn.fccf.aws.mskcc.org/CYTO_2017_B43_impact_of_size_on_LTD_and_DCD_12_049764b2fa.jpg
- Test sort precisions

Acknowledgments

Amsterdam UMC - Microscopy and Cytometry Core Facility

Sara Garcia-Garcia

Tanja Konijn

Baylor College of Medicine - Cytometry and Cell Sorting Core

Amanda White

Madhavi Chintalapati

Claude Chew

Courtney Harvey

Mason Perry

Paul Porter

Columbia University - Columbia Stem Cell Initiative

Michael Kissner

Cornell Institute of Biotechnology - Flow Cytometry Facility

Amanda Ferguson

Lydia Tesfa

Emory University - Emory + Pediatric's/Winship Flow Cytometry Core

Aaron Rae

Prasanthi Chappa

Houston Methodist Research Institute

David Haviland

Ellie Jardinella

Johns Hopkins University - Integrated Imaging Center

Hanhvy Bui

Erin Pryce

Ian Dobbie

Massachusetts General Hospital

Daire Daly

Mayo Clinic - Cytometry and Cell Imaging Lab (CCIL)

Laura Lewis-Tuffin

Margaret Ushman

Michigan State University - MSU Flow Cytometry Core

Daniel Vocelle

Rice University - Shared Equipment Authority (SEA)

Harshavardhan Deshmukh

University of Auckland - Auckland Cytometry

Thaize Chometon

Tanvi Damani

Sandy Chen

University of Iowa - University of Iowa Flow Cytometry Facility

Heath Vignes

Mike Shey

Tom Kaufman

University of Nebraska Medical Center - Flow Cytometry Research Facility

Craig L. Semerad

University of Utah - Flow Cytometry Core Facility

James Marvin

Eduardo Salustiano Jesus dos Santos, Ph.D.

University of Wisconsin - Madison - UWCCC Flow Cytometry Laboratory

Kathryn Fox

Dagna Sheerar

West Virginia University - WVU Flow Cytometry & Single Cell Core Facility

Kathleen Brundage

Raven Forshee

ABRF 2024 Annual Meeting | Minneapolis, MN | April 21-24

Preparing today's cores for tomorrow's needs



ABRF FCRG

(includes current and former study participants)

Mehrnoosh Abshari, NIH/NIDCR

Roxann Ashworth (Executive Board liaison), Johns Hopkins University, School of Medicine

Claudia Bispo, University of California San Francisco

Sara Bowen, Dignity Health St. Joseph's Hospital and Medical Center (former)

Ching-Yuan (Steve) Chen, Columbia University Irving Medical Center

Xiaoxuan Fan, University of Maryland School of Medicine

Kevin Ferro, Stowers Institute for Medical Research (former) – **statistical analysis year 1**

Claire Fraser, Barrow Neurological Institute – **organized data and ran statistical analysis year 2**

Christiane Hassel, Indiana University – **study co-author**

Celine Lages, Cincinnati Children's Hospital Medical Center

Pam Moody, Cold Spring Harbor Laboratory – **study co-author**

Steven Polter, University of Rochester Medical Center

Kenneth Quayle, Cincinnati Children's Hospital Medical Center – **organized data year 2**

Kathy Schaefer, HHMI Janelia Research Campus (former)

Rachael Sheridan (Co-Chair), Van Andel Research Institute Flow Cytometry Core

Jane Srivastava (Chair), Gladstone Institutes

John Tigges, Beth Israel Deaconess Medical Center (former)

Eric Wieder, University of Miami Miller School of Medicine

Thank you to the ABRF for funding the study!

ABRF 2024 Annual Meeting | Minneapolis, MN | April 21-24

Preparing today's cores for tomorrow's needs



Do you have these instruments?

- Beckman Coulter – CytoFLEX SRT & MoFlo Astrios EQ
- BD Biosciences – Influx, FACSMelody & FACSDiscover S8
- Bio-Rad - S3e
- Cytek Biosciences – Aurora CS
- ThermoFisher – Bigfoot

Join Year 3 of the ABRF DD Study! →



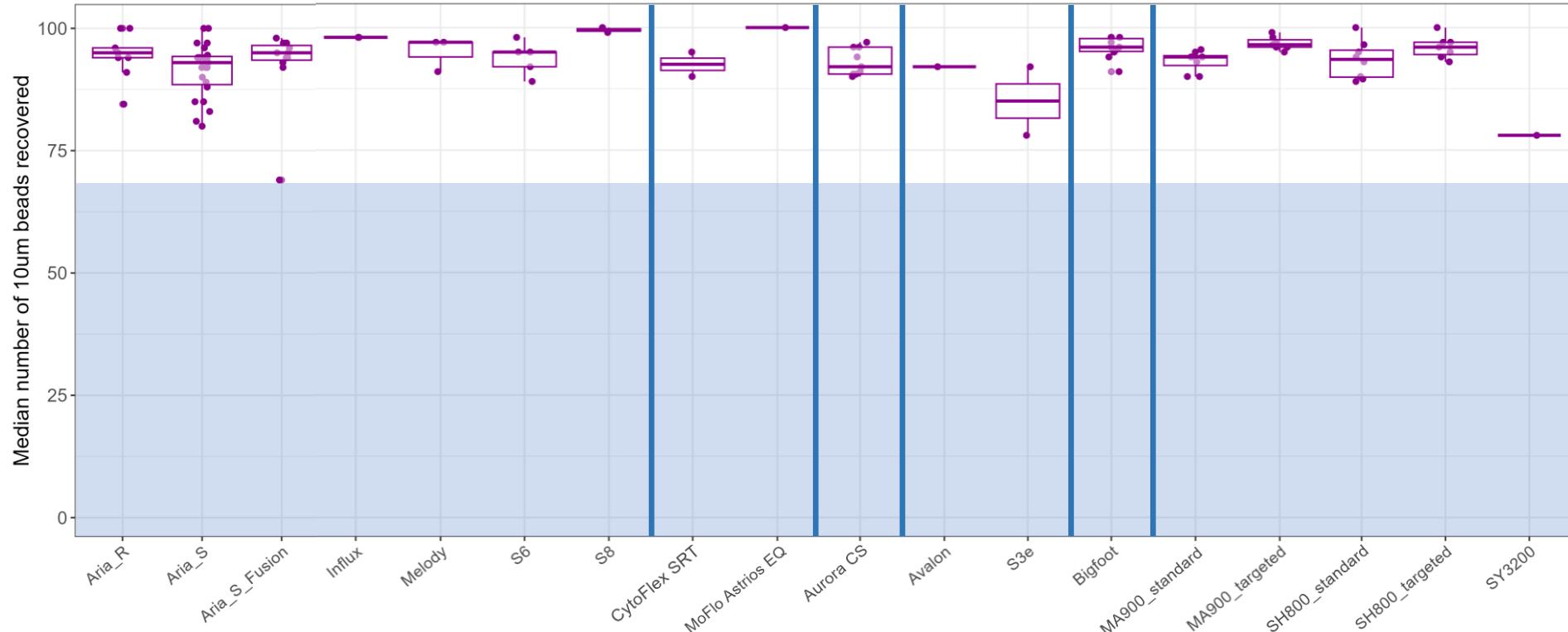
Thank you!

ABRF 2024 Annual Meeting | Minneapolis, MN | April 21-24
Preparing today's cores for tomorrow's needs



Extra slides

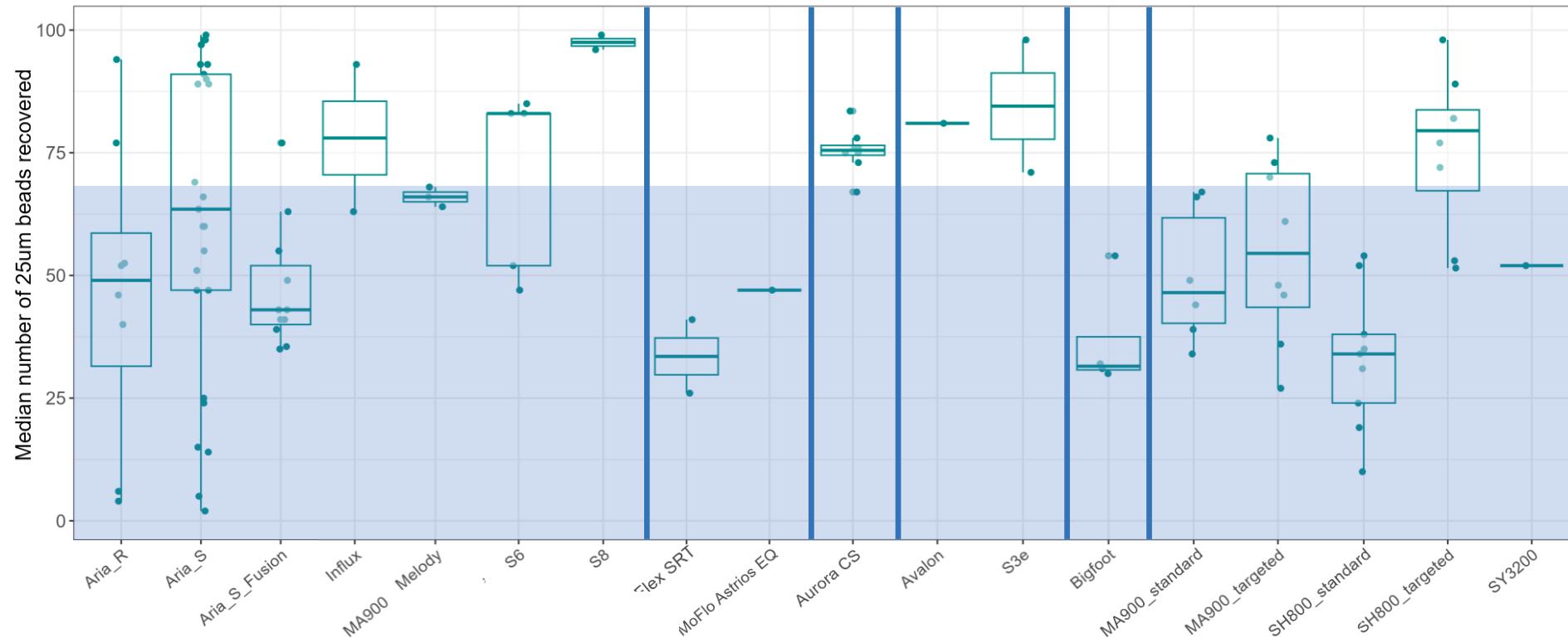
Consolidated 10-micron data – Fusion separate



Number of each instrument:

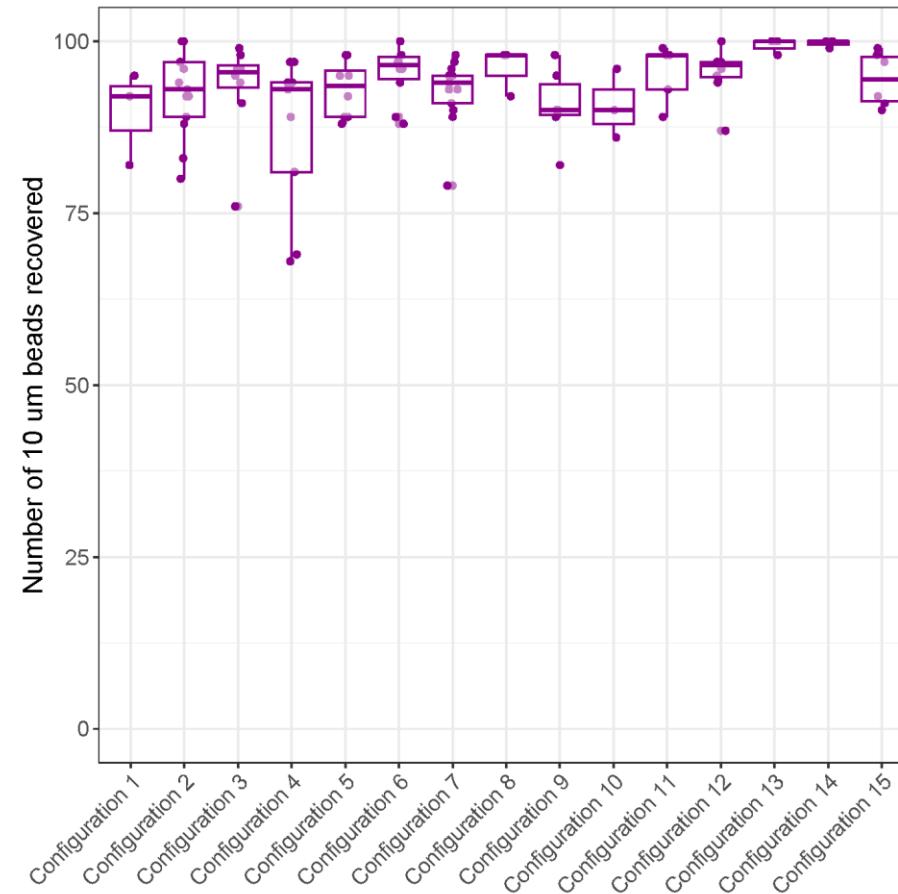
Aria-R - 10
Aria-S - 19
Aria Fusion - 8
Influx - 2
Melody - 2
S6 - 3
S8 - 1
CytoFLEX SRT - 2
MoFlo Astrios EQ - 1
Aurora CS - 7
Avalon - 1
S3e - 1
Bigfoot - 4
MA900-S - 7
MA900-T - 7
SH800-S - 7
SH800-T - 7
SY3200 - 1

Consolidated 25-micron data – Fusion separate



RESULTS – year 1

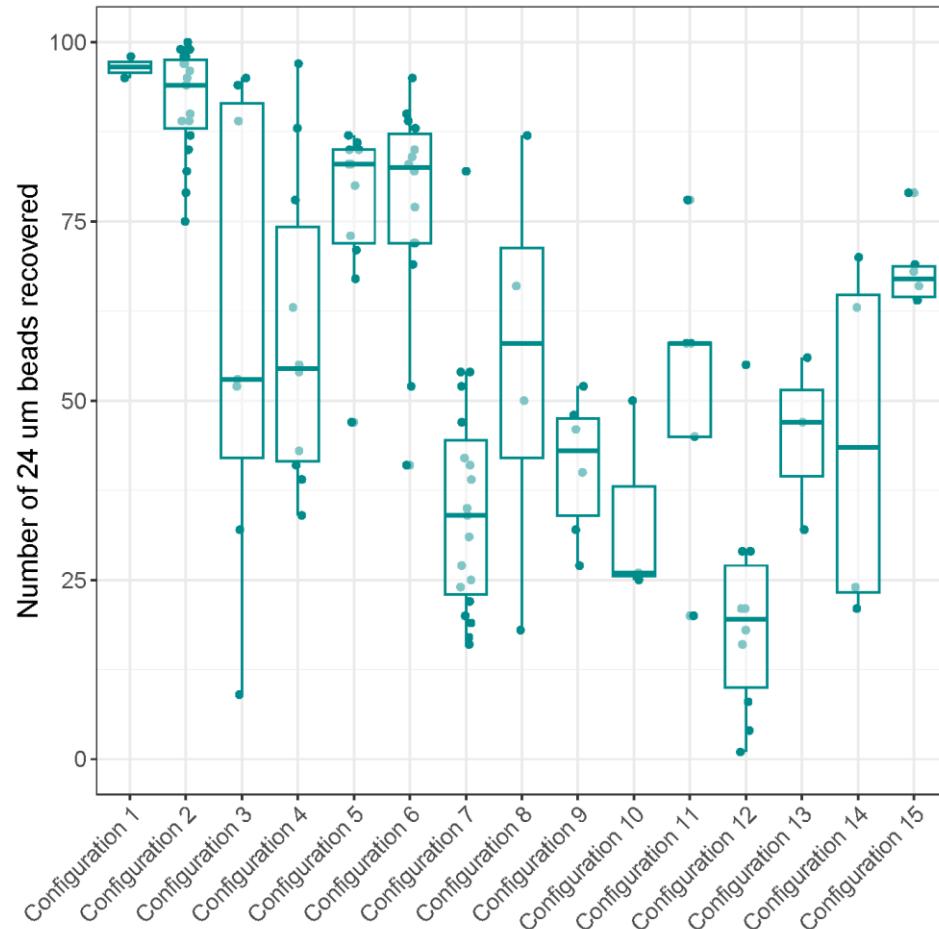
There was minimal variability in 10-micron bead recovery across all configurations using the Automated Drop Delay Setting.

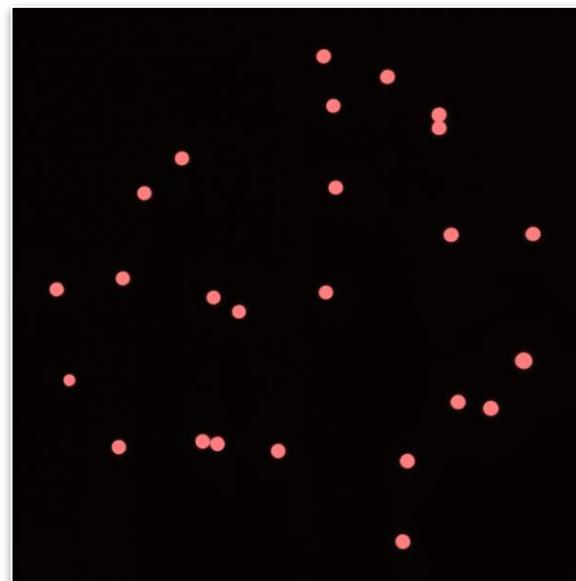
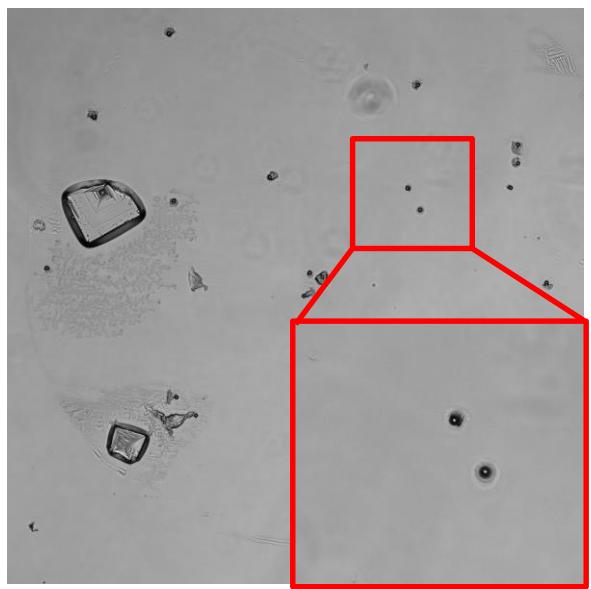


There was significant variability among 24-micron bead recovery across most configurations.

Configurations 1 and 2 show almost no change in the number of beads that were recovered using 10- or 24-micron beads

All other configurations show a 17 to 84 percent decrease in the recovery of 24-micron beads compared to 10-micron





Beads were sorted on to a slide and counted manually using a fluorescence or light microscope.

Side stream fanning of varying degree was observed on some models of cell sorter. This fanning was seen predominantly when sorting the 24-micron beads, although minor fanning was experienced while sorting the 10-micron beads with at least one model of sorter.

Left: Example of side stream fanning of 24-micron beads.

Right: Example of side stream fanning of 10-micron beads.

