The ImageStream[®] System for Imaging Cells in Flow: A New Tool for New Applications



www.amnis.com

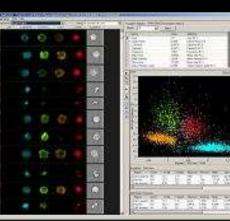
The ImageStream[®] System

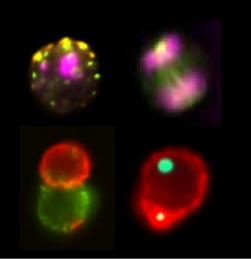


- ImageStream 100 Flow Imaging Instrument Brightfield, darkfield, and 4 fluorescent images at 5000 cells per minute
- IDEAS[®] Statistical Image Analysis Software Quantitative cellular image analysis and population statistics
- Novel Applications

Translocation, co-localization, cell classification, cell cycle, FISH-IS, etc.



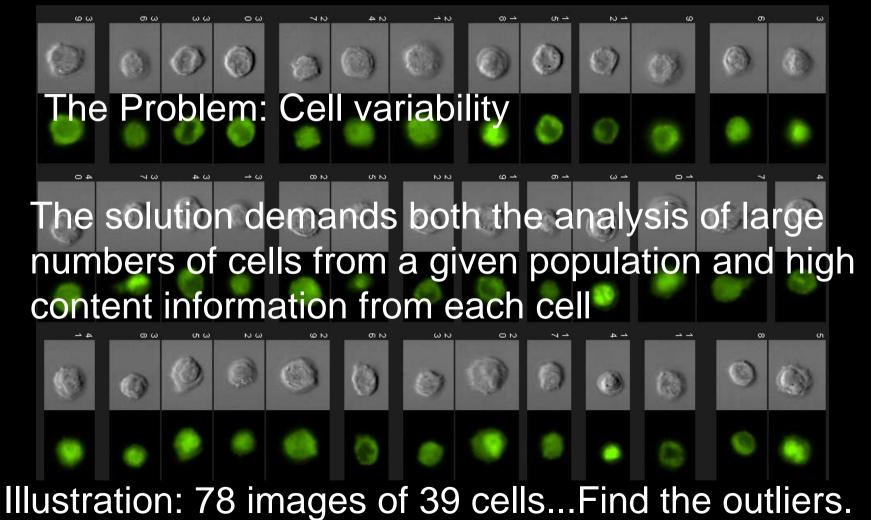




Issues in Quantitative Cell Biology



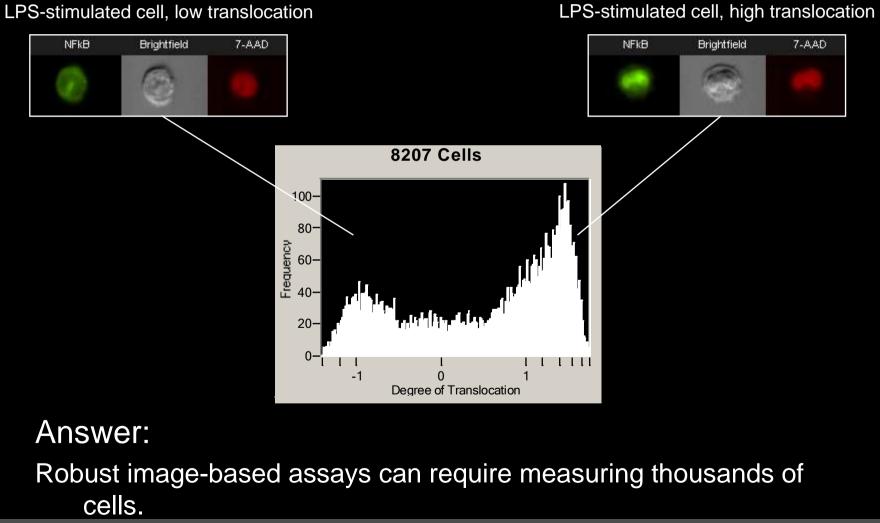
The Goal: Statistically robust assays with predictive power



Question: How Many Cells Do You Need?



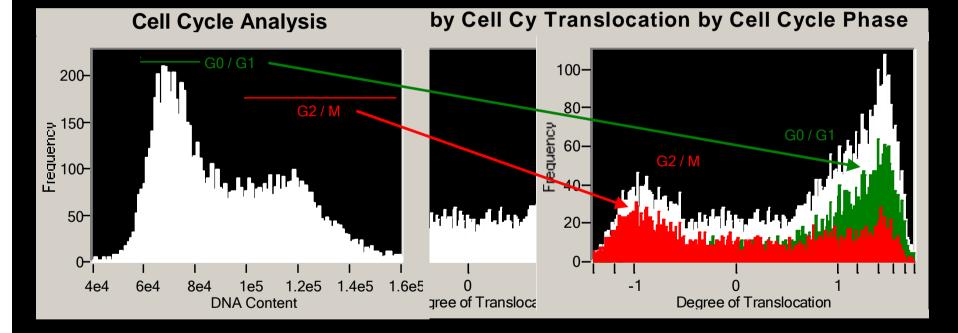
Example: NF-kB translocation assay in monocytes



How Much Information per Cell is Required? amnis

Example:

Incomplete NF-κB translocation in LPS-stimulated monocytes

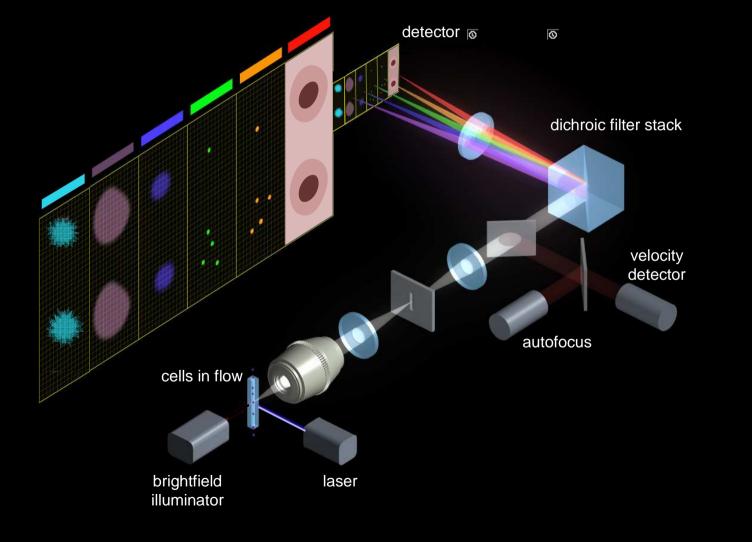


Conclusion:

The more you measure, the more you can understand.

ImageStream 100 Optical Layout



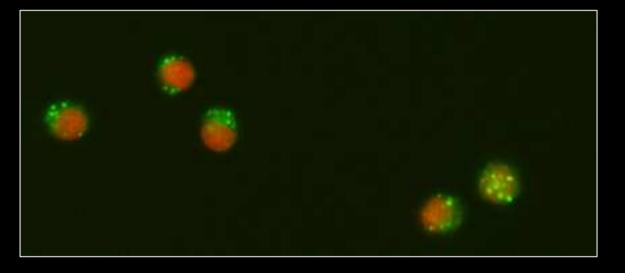


Resolution Comparable to Microscopy

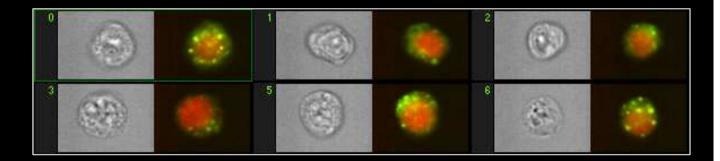


Intracellular Identification of Early Endosomes: EEA1-Alexa Fluor[®] 488 + 7-AAD

fluorescence microscopy: (40X mag)



ImageStream 100: (36X mag)



IDEAS[™] Software



| Population Statistics Object Data Co | mpensation Matrix | | |
|--------------------------------------|-------------------|---------------------|--|
| Object: 492 - | Select | | |
| Feature | Value | Definition | |
| ▶ 1_Area | 3445 | Area M1 0 | |
| 1_Aspect Ratio | 0.7613 | Aspect Ratio M1 0 | |
| 1_CentroidX | 39.5376 | CentroidX M1 0 | |
| 1_CentroidY | 55.1402 | CentroidY M1 0 | |
| 1_Frequency | 3.9042 | Frequency M1 0 | |
| 1_Major Axis | 35.5175 | Major Axis M1 0 | |
| 1_Mean Intensity | 7.683 | Mean Intensity M1 0 | |
| 1_Minor Axis | 30.9907 | Minor Axis M1 0 | |

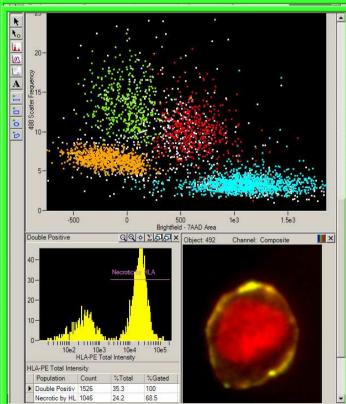




Image Gallery

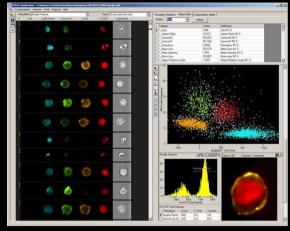
see every cell flexible viewing enhance & color tag populations virtual cell sort

Tabular Data 200+ params/cell population statistics object values

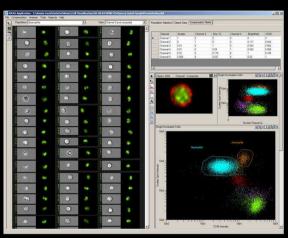
Workspace uni + bivariates flexible gating click dot to view cell custom parameters

Example Applications

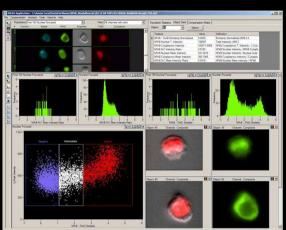




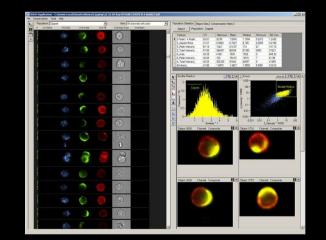
Apoptosis / Necrosis



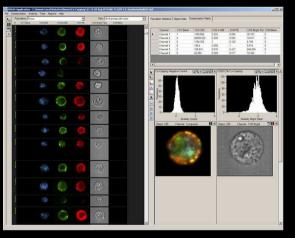
Cell Classification



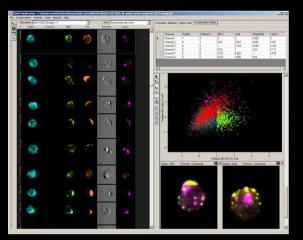
NF-κB Translocation



Marker Cap Quantitation



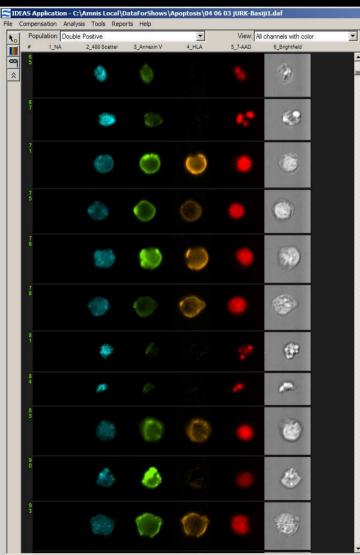
Marker Co-Localization



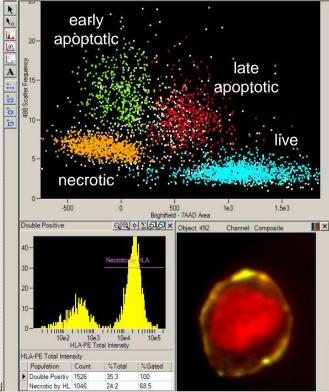
Molecular Trafficking

Mechanisms and Stages of Cell Death





| Object: 492 💌 | Select | |
|-------------------------|---------|----------------------------|
| Feature | Value | Definition |
| ▶ 1_Area | 3445 | Area M1 0 |
| 1_Aspect Ratio | 0.7613 | Aspect Ratio M1 0 |
| 1_CentroidX | 39.5376 | CentroidX M1 0 |
| 1_CentroidY | 55.1402 | CentroidY M1 0 |
| 1_Frequency | 3.9042 | Frequency M1 0 |
| 1_Major Axis | 35.5175 | Major Axis M1 0 |
| 1_Mean Intensity | 7.683 | Mean Intensity M1 0 |
| 1_Minor Axis | 30.9907 | Minor Axis M1 0 |
| 1 Object Rotation Angle | 1,3001 | Object Rotation Angle M1 0 |



Jurkat cells

- 0 ×

-

Treat with peroxide or camptothecin

Imagery (L-R): darkfield Annexin V-AF488 α-HLA PE nucleus (7-AAD) brightfield

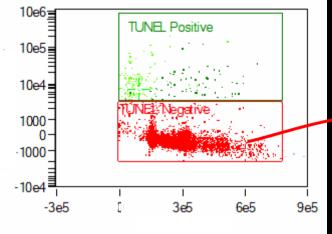
Use morphology to discriminate live cells from early apoptotic, late apoptotic, and necrotic cells.

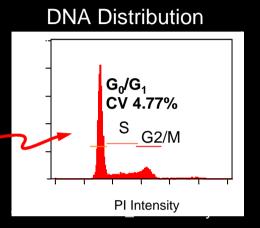
Apoptosis: Accurate cell death measurements



Negative control

| Population Statistics | | | | |
|------------------------------|-------|--------|--|--|
| Population | Count | %Gated | | |
| Single | 19155 | 100 | | |
| TUNEL Positive & Single | 180 | 0.94 | | |
| TUNEL Negative & Single | 18969 | 99 | | |
| TUNEL True Positive & Single | 113 | 0.59 | | |

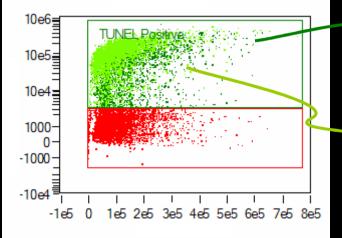


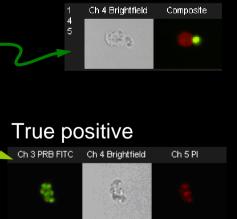


False positive**33%**

Apoptosis-induced

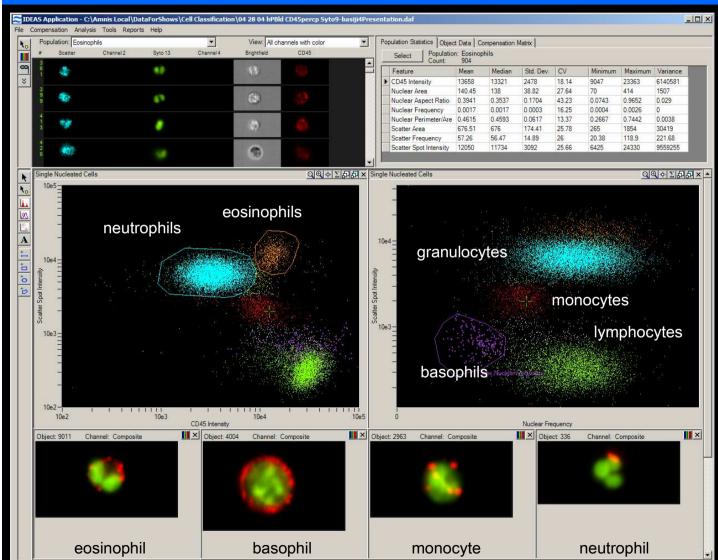
| Population Statistics | | | | |
|------------------------------|-------|--------|--|--|
| Population | Count | %Gated | | |
| Single | 20613 | 100 | | |
| TUNEL Positive & Single | 6521 | 31.6 | | |
| TUNEL Negative & Single | 14092 | 68.4 | | |
| TUNEL True Positive & Single | 4310 | 20.9 | | |





Morphologic Cell Classification





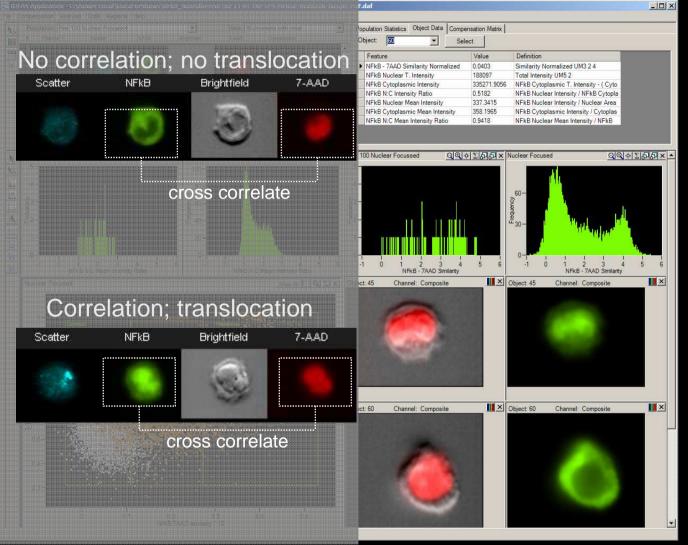
Lyse erythrocytes

Human blood

Imagery: darkfield SYTO-DNA α-CD45-PerCP brightfield

Build classifiers using correlation of stained populations with morphologic parameters.

NF-kB Translocation in Monocytes



THP-1 cells

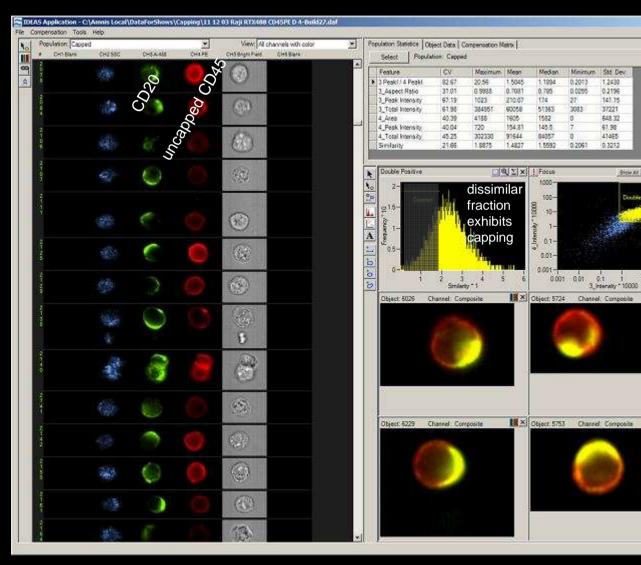
Treat with LPS

ar

Images (L-R) darkfield α-NFκB-FITC brightfield nucleus (7-AAD)

Measure NFκB to 7-AAD similarity to quantify degree of translocation

Quantitation of Marker Capping am



Raji cells

_IDIX

1.2433

0.2196

141.15

648.32

61.98

41455

0.3212

Show All Q X X

N X

X

37221

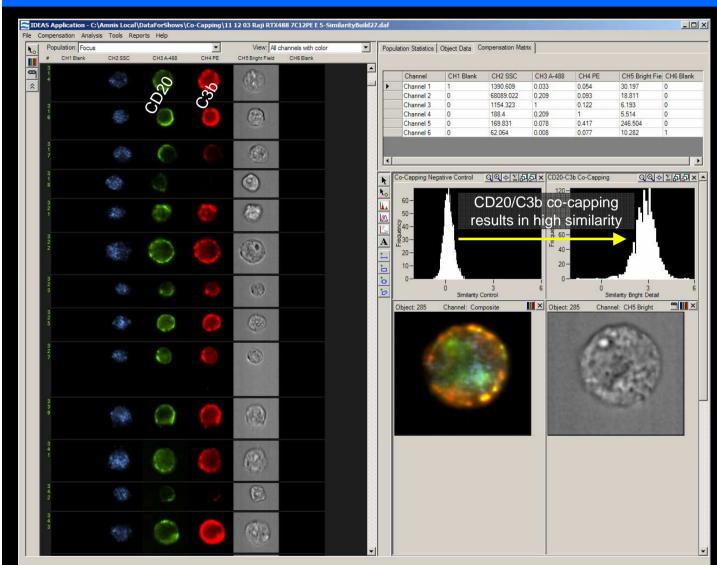
Imagery (L-R): darkfield rituximab - AF488 α-CD45 - PE brightfield

Measure similarity between fluorescent images

Data produced in collaboration with Dr. Paul Beum and Dr. Ronald Taylor, University of Virginia School of Medicine

Marker Co-Localization





Raji cells

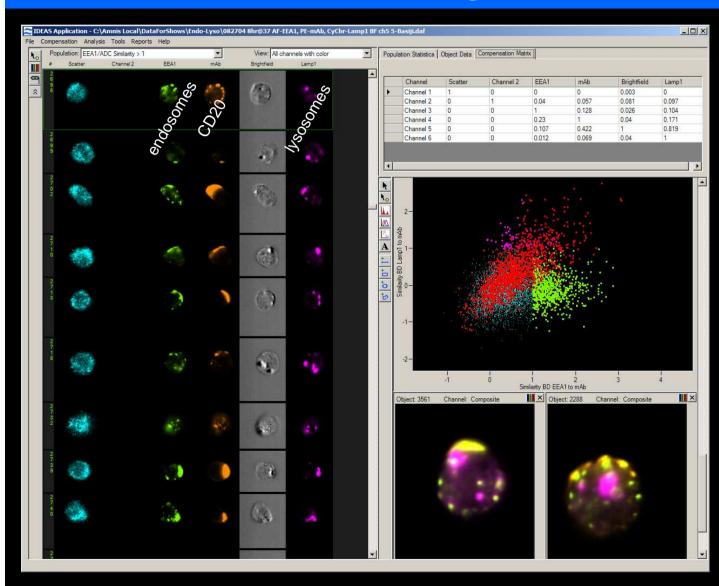
Imagery (L-R): darkfield rituximab - AF488 α-C3b(i) - PE brightfield

Measure similarity between fluorescent images and compare to control.

Data produced in collaboration with Dr. Paul Beum and Dr. Ronald Taylor, University of Virginia School of Medicine

Molecular Trafficking





Ramos cells

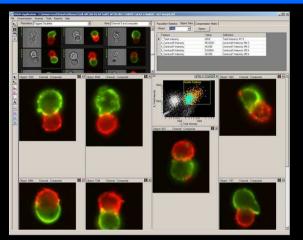
Imagery (L-R): darkfield EEA1 - AF488 α-CD20 - PE brightfield Lamp1 - CyChrome

Incubate at 37 °C for 0.5, 1, 2, 4, 8 hours. Measure similarity between mAb and endo / lyso images.

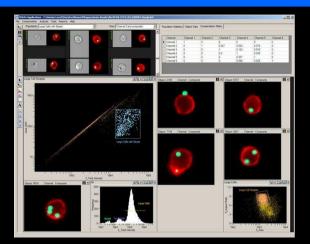
Monitor time-course of mAb-endo and mAb-lyso association.

Additional Applications

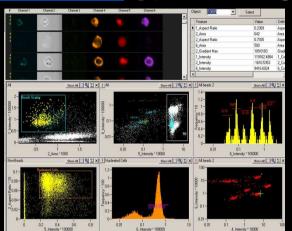
amnis



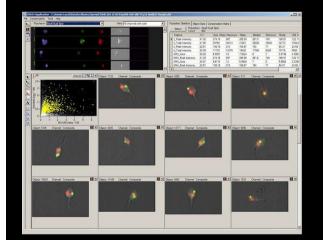
T Cell / APC Conjugates



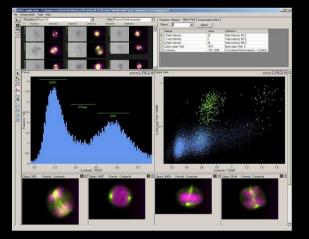
Phagocytosis



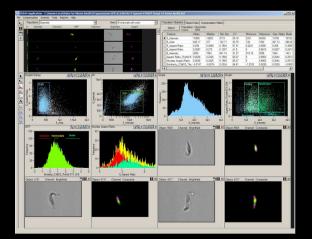
Multiplexing



High Throughput FISH-IS



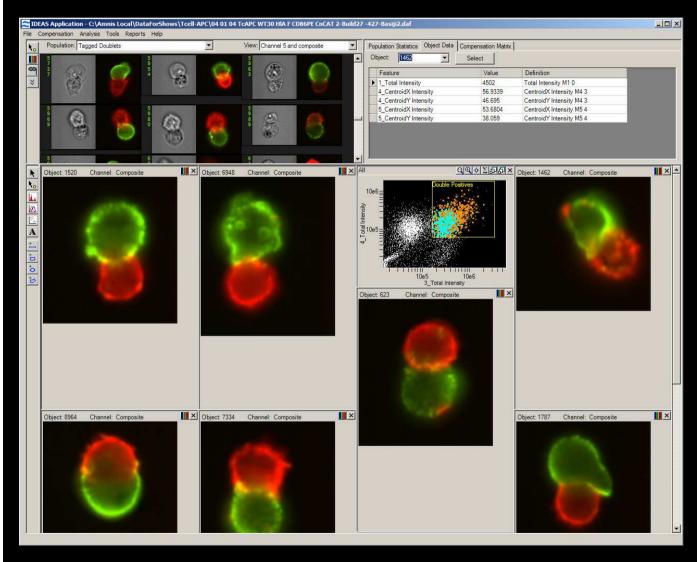
Cell Cycle and Mitosis



Infectious disease

T-cell / APC Interactions





Murine cells

Imagery (L-R): brightfield HLA-FITC + CD86-PE

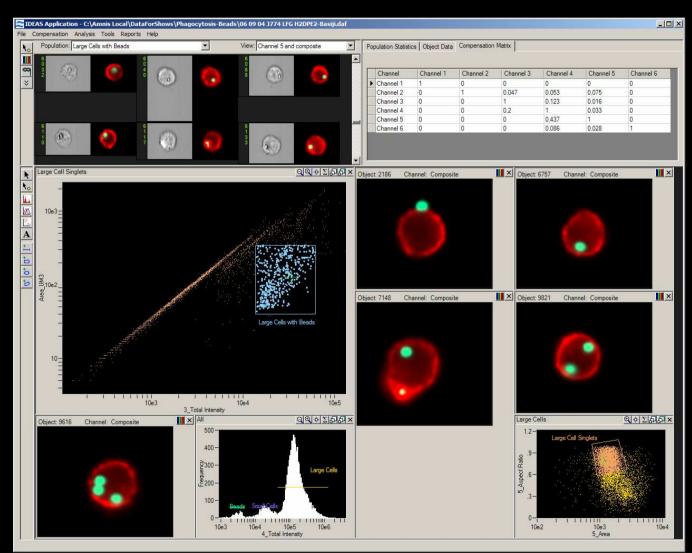
Define contact area using logical AND of FITC and PE masks.

Measure mean CD86 intensity at synapse vs. remaining APC area.

Data produced in collaboration with Dr. Rafick-Pierre Sekaly, University of Montreal

Phagocytosis





Murine J774a cells

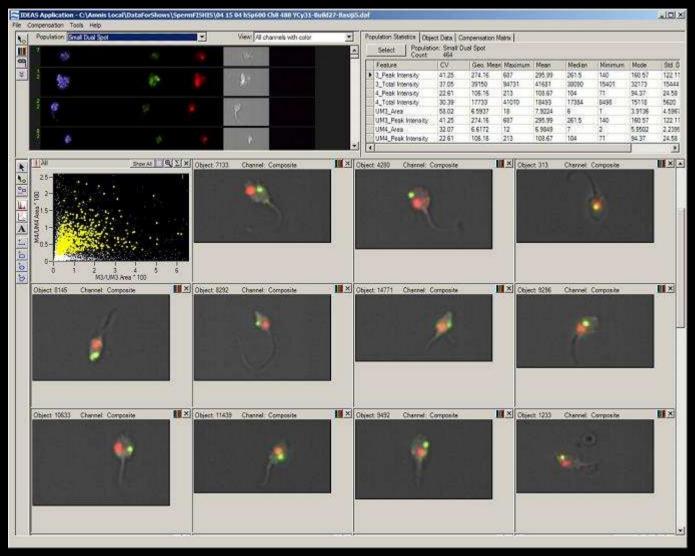
Mix with 2.5um fluorescent beads at equal concentration, shake 2 hr @ 37° C Wash, fix, run.

Imagery (L-R): brightfield FITC beads + H-2D-PE

Find H-2D positives, define cell singlets using brightfield area and aspect ratio, find phagocytic 5% using FITC intensity.

High Throughput FISH-IS[™]





FISH-IS with Chr. 8-FITC probe Chr. Y-Cy3 probe

Human Sperm

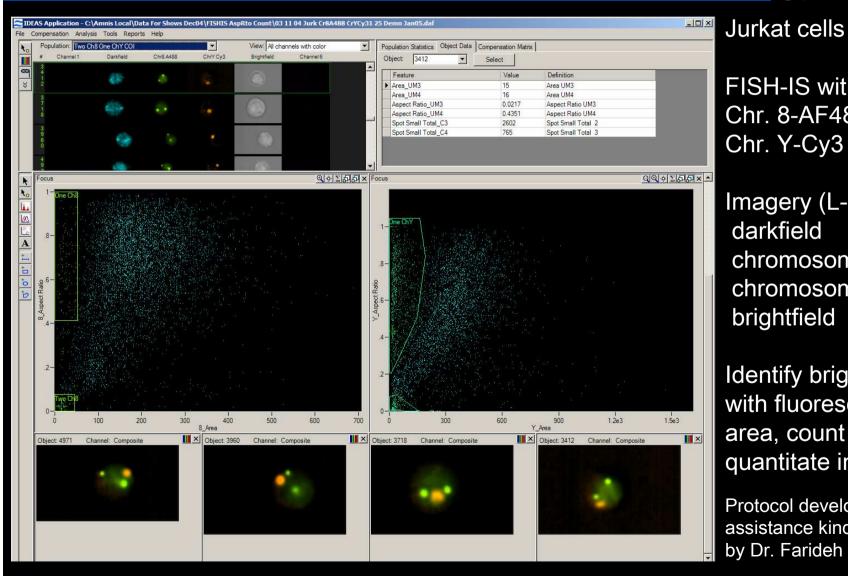
Imagery (L-R): darkfield chromosome 8-FITC chromosome Y-Cy3 brightfield

Identify bright FISH with fluorescent mask area, count spots, quantitate intensity

Work supported in part by NIEHS SBIR N43-ES-35507.

High Throughput FISH-IS[™]





FISH-IS with Chr. 8-AF488 probe Chr. Y-Cy3 probe

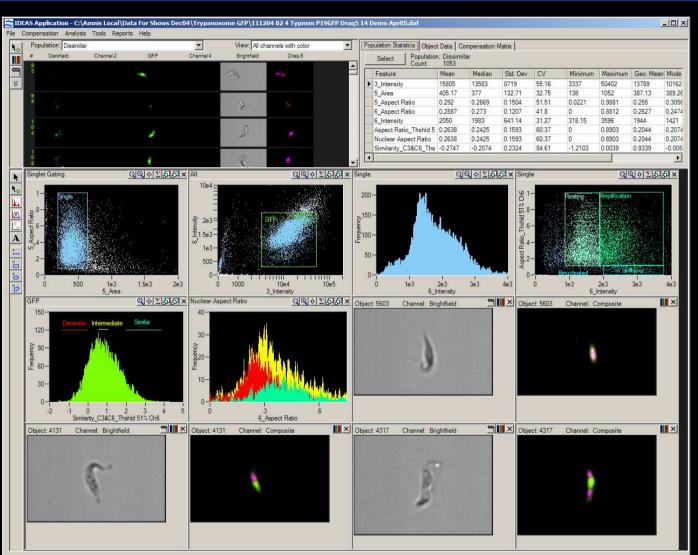
Imagery (L-R): darkfield chromosome 8 chromosome Y brightfield

Identify bright FISH with fluorescent mask area, count spots, quantitate intensity

Protocol development assistance kindly provided by Dr. Farideh Bischoff.

Infectious Disease





GFP-transfected *Trypanosome brucei*

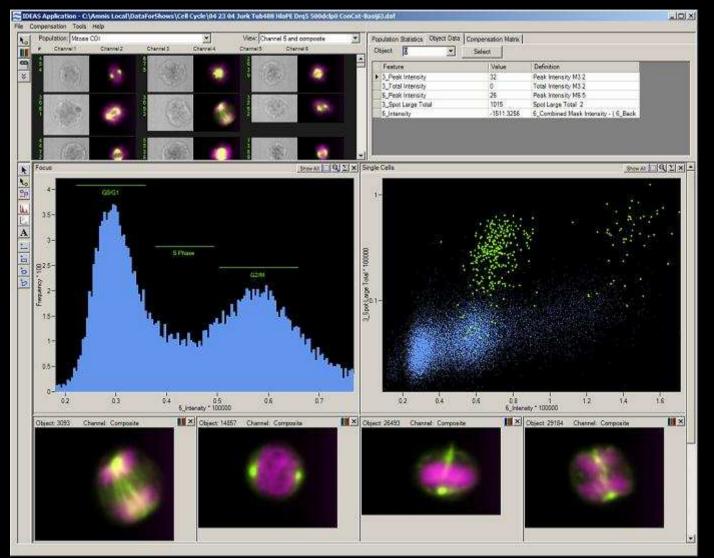
Imagery (L-R): darkfield GFP-NP19 brightfield DRAQ5[™] (DNA)

Measure similarity of GFP to DRAQ5 to quantify degree of translocation

Data produced in collaboration with Dr. Marilyn Parsons, Seattle Biomed. Res. Inst.

Cell Cycle and Mitosis Analysis





Jurkat cells

Stain with DRAQ5[™] α-tubulin-AF488, α-HLA-PE

Imagery: brightfield DRAQ5[™] + tubulin

Quantitate cell cycle using total intensity, identify mitotic cells using peak intensity of tubulin and/or DRAQ5[™].

Multiplexed Cytokine/Hematology



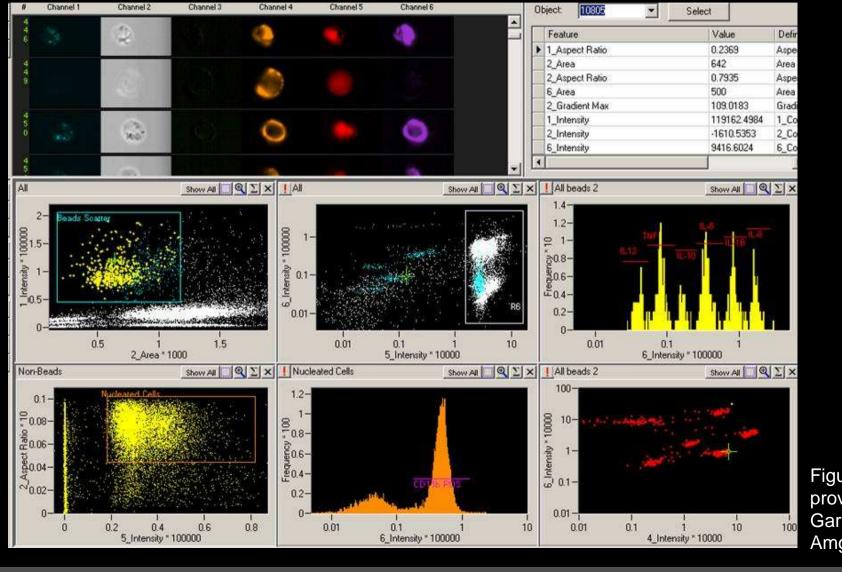


Figure kindly provided by Gary Elliott, Ph.D. Amgen, Inc.

Customer Driven Applications



- Human bone marrow
- Nuclear translocation in multiple cell types
- Phagocytosis in rare human blood cells
- DNA repair
- Shape change assays
- Bead Cell conjugates
- Biomarker identification

Summary



ImageStream System Delivers Quantitative Cell Biology:

- Easily measure many cells to describe population structures
- Replace biomarkers with quantitative morphology
- Apply rigorous assays with analytical flexibility
- Get results that are objective and verifiable



Clarity from Complexity