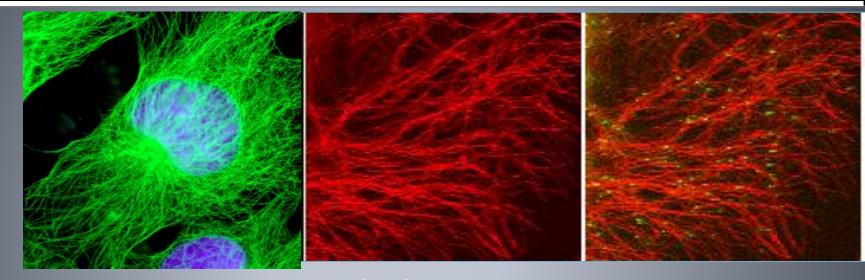
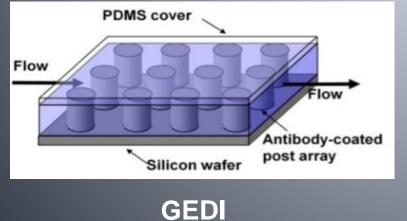
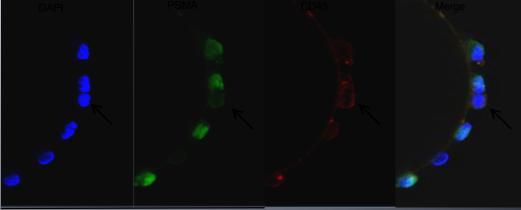
Using Circulating Tumor Cells for Precision Medicine



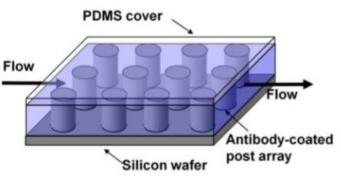
Biology

CTCs: Portal to individual patient Tumor





The Solution: Circulating Tumor Cells

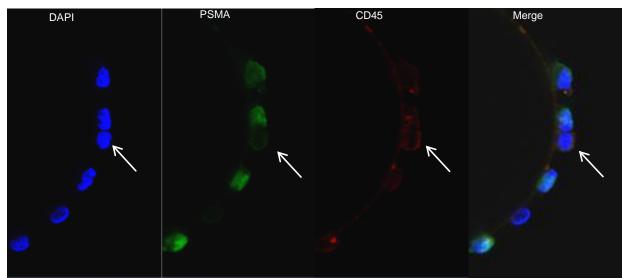


CTCs: Portal to individual patient Tumor

Brian Kirby, Engineering, Cornell, Ithaca CTSC Pilot Funding, 2008

GEDI

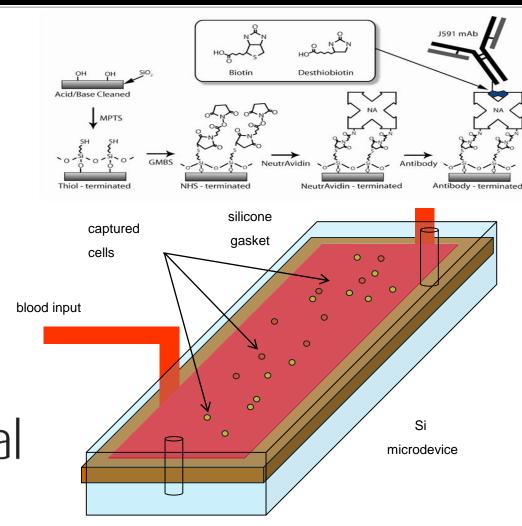




Finding a needle in a haystack: Geometrically enhanced differential immunocapture (GEDI)

- 1 mL of whole blood is injected into the device over 1 hour (flow rate at 1ml/h)
- GEDI: geometrically enhanced differential immunocapture

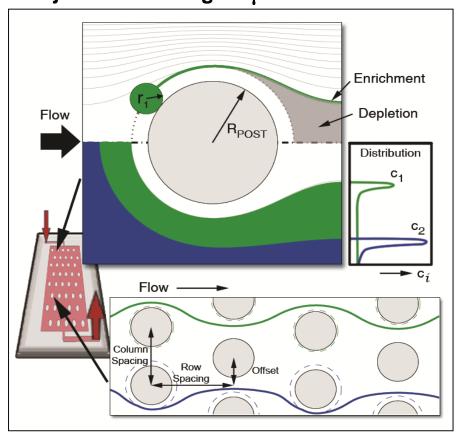
Weill Cornell Medical College
Clinical & Translational

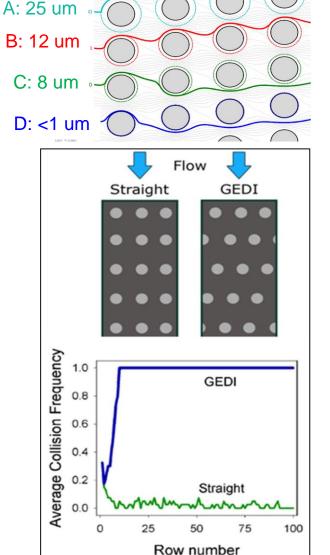


Capturing the elusive cancer cell: Fluid mechanical trickery and the development of GEDI microdevice.

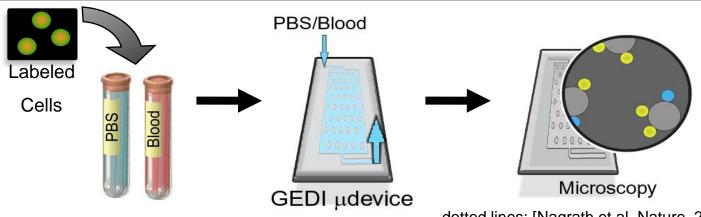
 GEDI technique combines size and surface specificity to exploit differences between cancer cells and blood cells

Particle trajectories through a µfluidic obstacle array.



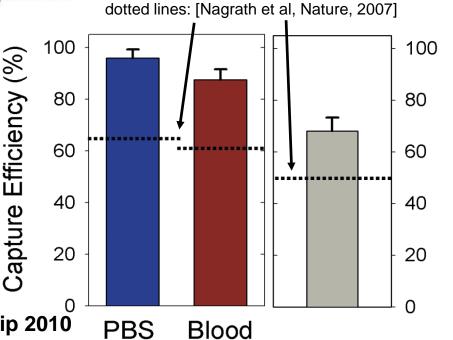


concentrates cancer cells by a factor of 10-100 million



 150-220 LNCaP cells/mL spiked in healthy whole blood or PBS+1% BSA

Cell type	#/mL	# captured
LNCaP	200	170
Blood	5 billion	91

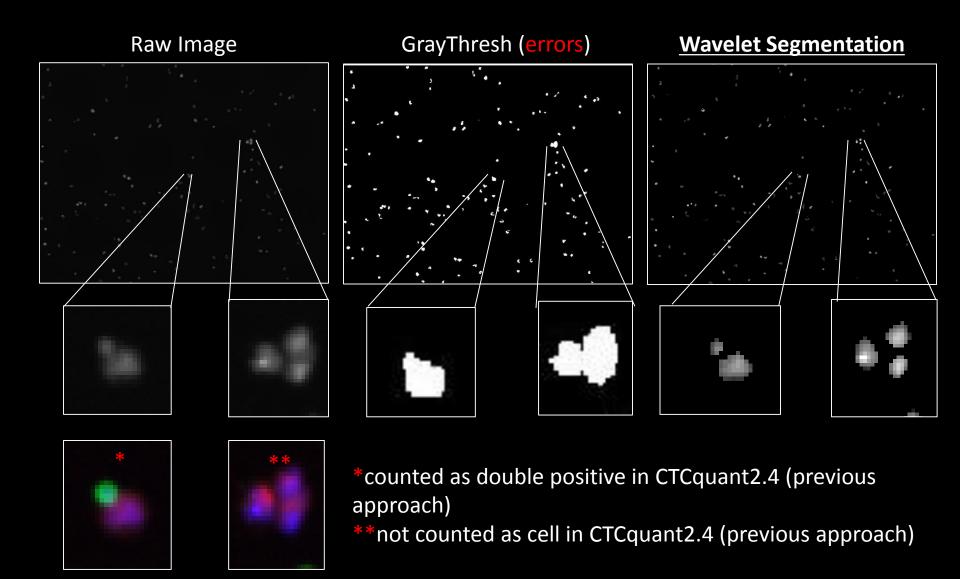


Gleghorn, Giannakakou, Kirby et al Lab on a Chip 2010

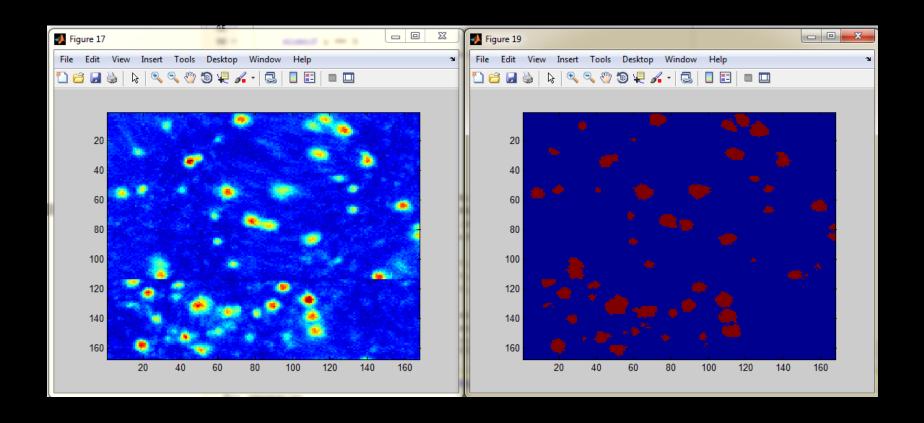
Timeline of GEDI Funding

- Partnership with Brian Kirby, Engineering, Cornell, Ithaca
- 2008, CTSC Pilot Funding
- 2009, NIH Ro1 (Giannakakou; Nanus)
- 2009, NYSTAR Designated Center for Advanced Technology (Kirby; Giannakakou)
- 2009, PCF Creativity Award (Nanus; Giannakakou; Kirby)
- 2011, NIH Center on the Microenvironment and Metastasis
 (Giannakakou; Project 2 co-leader; Kirby GEDI nanofabrifaction, Nanus, Project 4 co-leader)
- Sanofi, Prospective Phase II Clinical Trial in CRPC, 2013
- 2012, Development of HER2-GEDI for Breast and Gastric Cancer
- 2012, MWCBC, Breast Cancer Grant (Giannakakou)
- 2012, Eisai Pharma Breast cancer study (Giannakakou)
- 2013, Alex Matov, NRSA postdoctoral fellowship
- 2013, Alex Matov, CTSC Pilot grant

Wavelet-based DAPI Segmentation (Incubation chip pos350) cells merged with earlier algorithm, which was erroneous

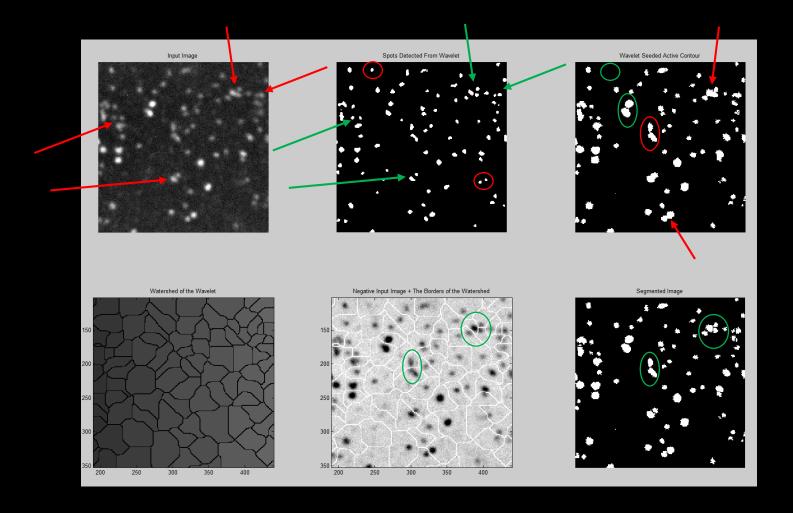


New segmentation method based on wavelet seeding and active contour using watershed limits of the area



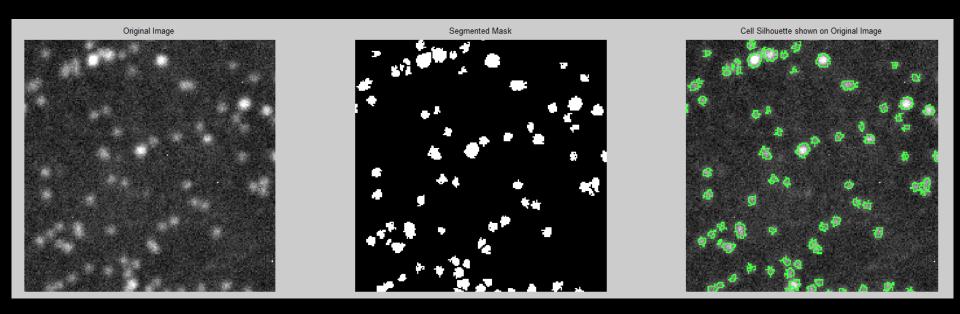
Algorithm designed and initiated by Alex Matov, and implemented in Matlab by Shayan Modiri (a student hired by A. Matov to work on his NCATS-NIH grant)

Novel PSMA Segmentation Method (Matov)



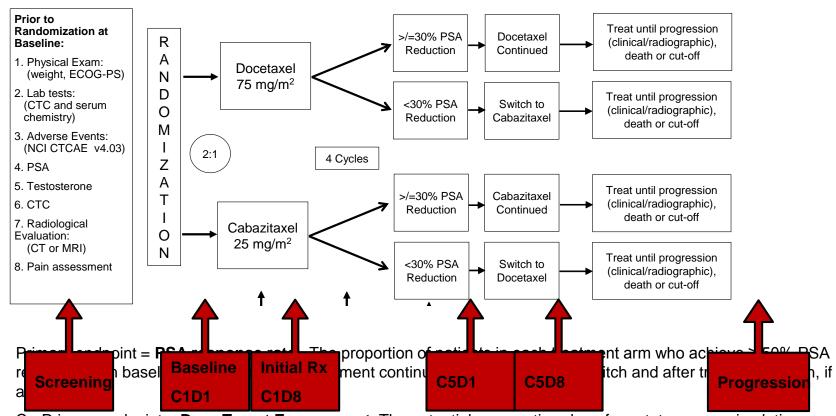
The areas of adjacent cells are correctly segmented even if the cells partially overlap

Novel PSMA Segmentation Method (Matov)



- The initial image segmentation is accomplished by stationary wavelet transform, which identifies bright pixel clusters in noisy images (seeding step)
- Active contour, next, identifies precisely the edges of the image features based on the seeds
- Watershed transformation of the seeding step image is overlaid, with reversed intensities, on the
 active contour image; logical conjunction ('and') of the active contour image and the watershed image
 identifies the bright areas and their exact borders

TAXYNERGY: Phase II Trial to Evaluate Benefit of Early Switch from first-Line Docetaxel/Prednisone to Cabazitaxel/Prednisone and the opposite sequence, exploring molecular markers and mechanisms of taxane resistance in men with Metastatic Castration-Resistant Prostate Cancer (mCRPC) who have not received prior chemotherapy



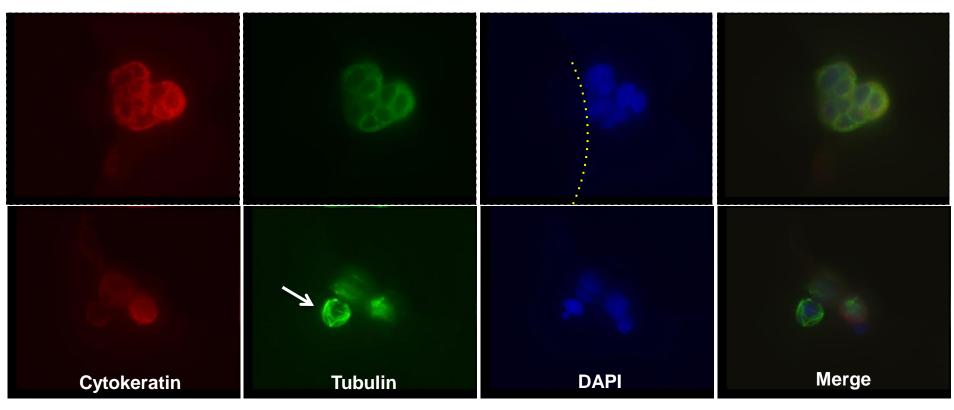
Co-Primary endpoint = **Drug Target Engagement**: The potential prognostic value of prostate cancer circulating tumor cells will be determined by using CTCs isolated at baseline and at defined time-points using the geometrically enhanced differential immunocapture (GEDI) method; by performing CTC enumeration, AR localization (cytoplasmic and nuclear) and tubulin bundling. The findings will be correlated with the clinical response/resistance to docetaxel/cabazitaxel for all patients

Clinical co-chairs: Antonarkis, Tagawa

Biomarker chair: Giannakakou

Assays on the Chip 4. Ex Vivo Treatment of LIVE CTCs to Predict Chemotherapy Response

Captured CTCs Ex Vivo treatment with Different Taxanes: Can CTC Response Predict Clinical Response?



Dotted Line: micropost margins

Arrow head: intricate microtubule network

Arrow: bundled microtubule (DTE effect)

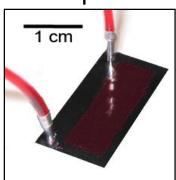
Customize Chemotherapy for the Individual

We aim to facilitate personalized treatment via capture and analysis of rare cells

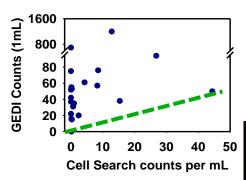
Patient Sample



Cancer cell Capture



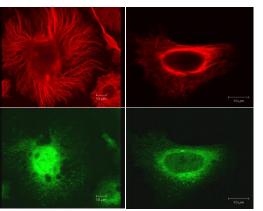
- our unique contribution: viable CTCs on chip for FUNCTIONAL ANALYSIS
 - Enumeration/
 - Genomics



Functional

Response

- Can we predict the response of prostate cancer patients to chemotherapy?
- Can we Tailor Therapy in Real Time?



Acknowledgements

Giannakakou Lab

- Ada Gjyrezi
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- Maria Thadani-Mulero, PhD
- Peppe Galletti, MD/PhD
- Luigi Portella, PhD
- Nancy Chan, MD
- Sidharta Shen, PhD
- Marisa Carbonaro, PhD
- Alex Matov, PhD

Collaborators

- David Nanus, MD (Weill Cornell)
- Scott Tagawa, MD (Weill Cornell)
- Neil Bander, MD (Weill Cornell)
- Brian Kirby, PhD (Cornell University)
- Stephen Plymate, MD (University of Washington, Seattle)

Collaborator Group Members

- He Liu PhD
 - Vincent Navarro, PhD
- Guang Lee
- Ganjun Gakhar, PhD
- Shihua Sun, PhD
 - Jason Gleghorn, Phd
 - Erica Pratt, PhD candidate
 - Yusef Sved
 - Erica Pratt
 - Steven Santana
 - Tim Lannir
 - Tim Smith
 - Charlie Huang