



SUMMARY OF MY MAIN ACTIVITIES

FILIPPO PICCININI

May 10, 2011



Who am I?



f.piccinini@unibo.it

| | |
|---------------------|-------------------------------------|
| First Name, Surname | Filippo Piccinini |
| Place of birth | Forlimpopoli, FC, Italy |
| Date of birth | April 20, 1985 |
| Title | Biomedical Engineer |
| Email | f.piccinini@unibo.it |
| Web site | www.filippopiccinini.altervista.org |

PH.D STUDENT

European Doctorate in Information Technology by

ARCES

Advanced Research Centre on Electronic Systems,
University of Bologna

SUPERVISOR

Prof. Alessandro Bevilacqua

CO-TUTOR

Prof. Mauro Ursino

MAIN RESEARCH TOPIC

Biomedical Image Elaboration





ARCES



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My office



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My Professor's office



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Biomedical engineering



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WEB SITE: <http://www.ing2.unibo.it>



Hospital partners



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IOR - BOLOGNA



**BONE
REGENERATION
LABORATORY**



IRST - MELDOLA



**OPTICAL
MICROSCOPE
LABORATORY**

**CELL
CULTURE
LABORATORY**



Your contact



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SUMMER SCHOOLS

- **ICVSS 2010 Computer Vision Summer School**

Scicli (Ragusa), Italy, July 12-17, 2010



F. Piccinini, A. Bevilacqua, A. Gherardi, L. Carozza, "*Accurate Mosaicing of Cell Images Acquired with Non Automated Equipment*". Poster to: ICVSS 2010 International Computer Vision Summer School, Scicli (Ragusa), Italy, July 12-17, 2010

- **CIMST 2010 Interdisciplinary Summer School on Bio-medical Imaging**

ETH Zurich, Switzerland, September 6-17, 2010



F. Piccinini, A. Bevilacqua, A. Gherardi, L. Carozza, E. Lucarelli, B. Dozza, "*Illumination Field Detection in Microscopic Images*". Poster to: 4th CIMST Interdisciplinary Summer School on Bio-medical Imaging, ETH Zurich, Switzerland, September 6-17, 2010.



Outline



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- Real time mosaicing
- Vignetting correction
- Microscope calibration
- Depth from focus
- Publications
- Contacts



Outline



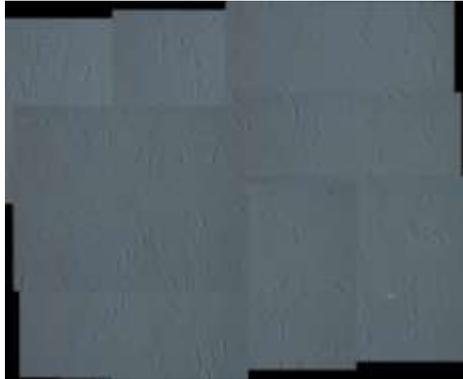
f.piccinini@unibo.it

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MOSAIC

A large image obtained stitching together two or many more images. A mosaic allows to extend the field of view of a camera, without losing resolution.



WHY MOSAICING IS IMPORTANT?



MOTIVATIONS OF MOSAICING

- To see a global condition of objects
- To store an instantaneous state of objects in contest
- To capture large objects of interest with a high resolution
- To perform time lapse experiments
- To perform high content analysis

MOTIVATIONS OF REAL TIME MOSAICING

- To acquire information of unexpected (tissue, cells) objects during a normal inspection
- To guide the operator during the navigation
- To get an immediate feedback when material is at one's disposal



BIOLOGICAL APPLICATIVE CONTEST

PROJECT ADVANCE

automatic non-invasive system based on high content analysis to detect and characterize vital mesenchymal stem cells in a spatio-temporal context

Collaboration with the Istituto Ortopedico Rizzoli (Bologna, Italy)

The Bone Regeneration Lab is the first in Italy to conduct a clinical trial of level 2 for utilization of Mesenchymal Stem Cells in regeneration medicine

GOAL

To study Mesenchymal Stem Cell's properties and behaviors to build biological pharmaceuticals to regenerate bones affected by tumors and other pathologies

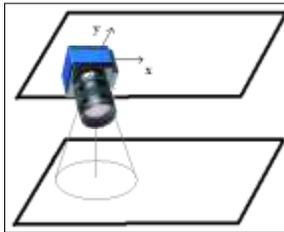


CONSTRAINTS

- Live cells for clinical uses → cells cannot be manipulated
- Non automated light microscope

PROBLEMS

- Very low contrast typical of mesenchymal stem cells observed in phase contrast mode
- Holder is not perfectly perpendicular to the optical axis
- Not uniform illumination field distribution in the microscopic field of view





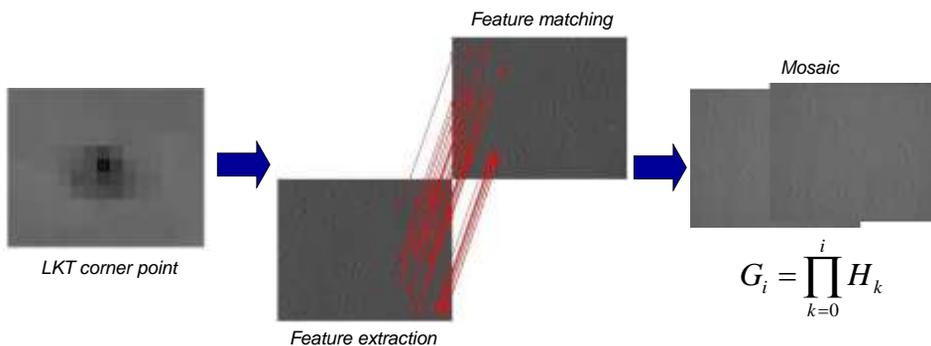
MATERIALS

- Microscope Nikon Eclipse TE2000-U equipped with Camera Nikon DXM1200F
- Cell cultures and histological samples



GEOMETRIC REGISTRATION

- Feature choice
- Lukas Kanade Tracker (LKT): corner point extraction and matching
- Transform matrix estimation





GEOMETRIC REGISTRATION

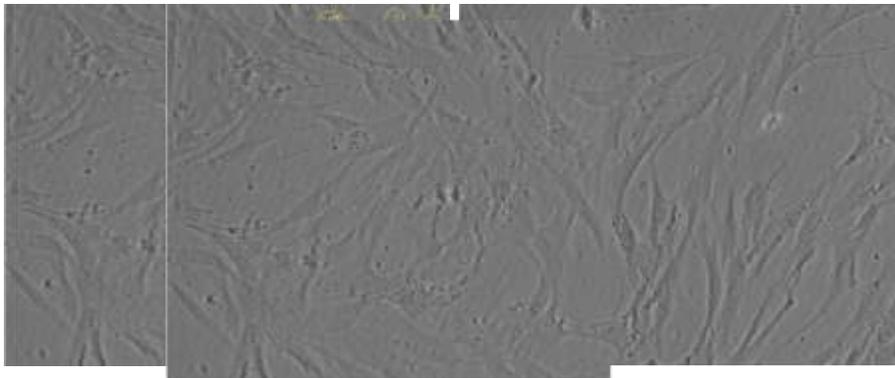
- Feature choice
- Lukas Kanade Tracker (LKT): corner point extraction and matching
- **Transform matrix estimation**

| | TRANSLATIVE | AFFINE | PROJECTIVE |
|---|---|---|---|
| | $H = \begin{bmatrix} 1 & 0 & t_1 \\ 0 & 1 & t_2 \\ 0 & 0 & 1 \end{bmatrix}$ | $H = \begin{bmatrix} a_{11} & a_{12} & t_1 \\ a_{21} & a_{22} & t_2 \\ 0 & 0 & 1 \end{bmatrix}$ | $H = \begin{bmatrix} a_{11} & a_{12} & t_1 \\ a_{21} & a_{22} & t_2 \\ v_1^T & v_2^T & 1 \end{bmatrix}$ |
|  |  |  |  |
| Original image | One corresponding point | Three corresponding points | Four corresponding points |



GEOMETRIC REGISTRATION

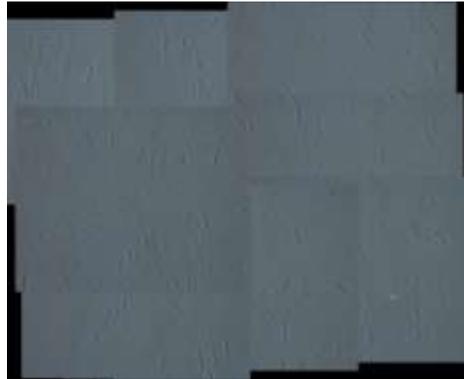
SUMMARY





PROBLEM

Seams between images can be markedly visible in stitching zone



PROBLEM

Seams between images can be markedly visible in stitching zone



In a first step **blending** can resolve problem of seams, but it can bring to color inhomogeneity, ghost effect, movement effect, etc.



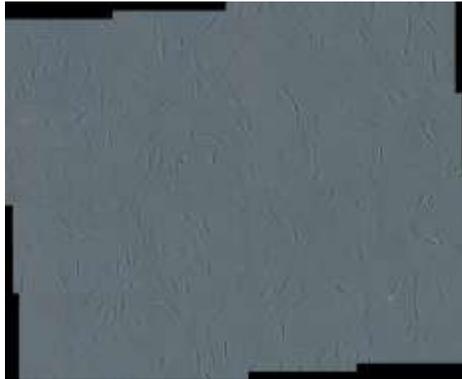
Real time mosaicing



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PROBLEM

Seams between images can be markedly visible in stitching zone



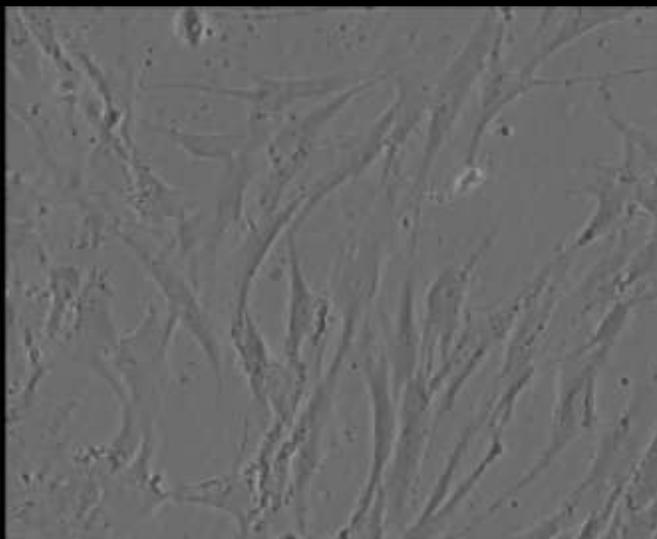
To completely resolve the problem the **vignetting function** due to uneven illumination field, lens, etc, must be studied...



Cell culture



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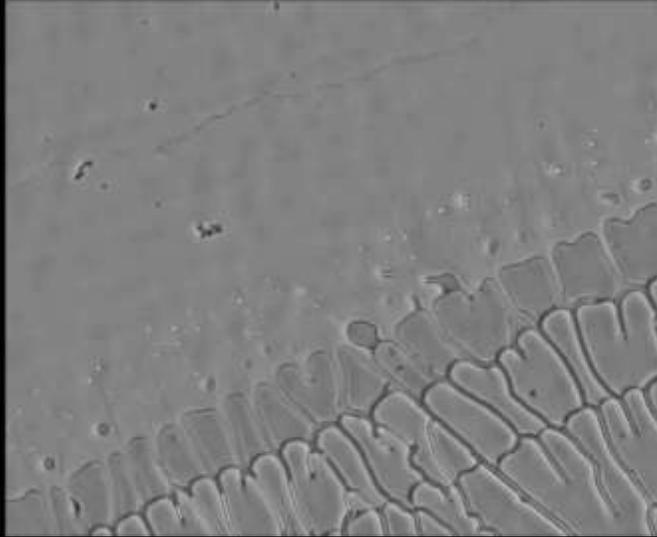
Courtesy of Dr. Ludovico Carozza



Histological sample



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Courtesy of Dr. Ludovico Carozza



Outline



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- Real time mosaicing
- **Vignetting correction**
- Microscope calibration
- Depth from focus
- Publications
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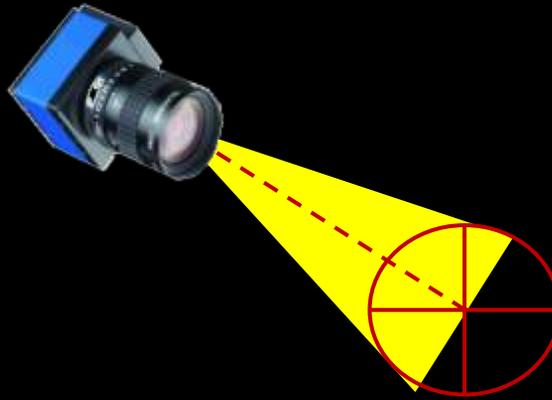


Vignetting



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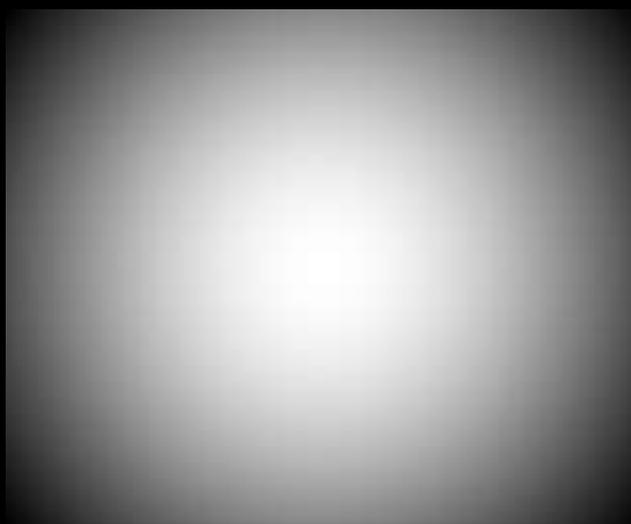
An effect of brightness attenuation away from the centre to the optical axis due to an uneven illumination field, lens, etc



Vignetting: a synthetic image



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Super-resolution Enhancement of T

David Capel and Andrew Z
Robotics Research Gr
Department of Engineering
University of Oxfor
Oxford OX1 3PJ, U

Abstract

The objective of this work is the super-resolution enhancement of image sequences. We consider in particular images of scenes for which the point-to-point image transformation is a plane projective transformation.

We first describe the imaging model, and a maximum

comput
tradition
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MAP es
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variation



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Vignetting: ideal case



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$$I = (F + B) IF$$

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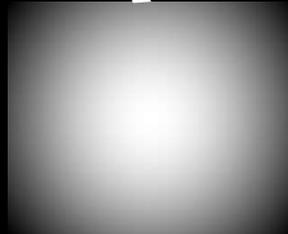
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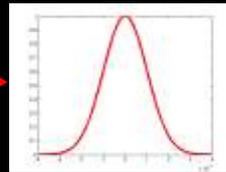


Vignetting: real case



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$$I = (F + B) IF + n \rightarrow$$



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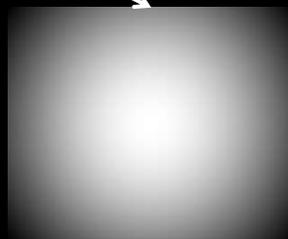
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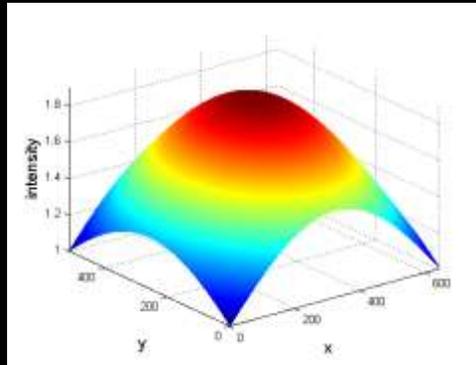
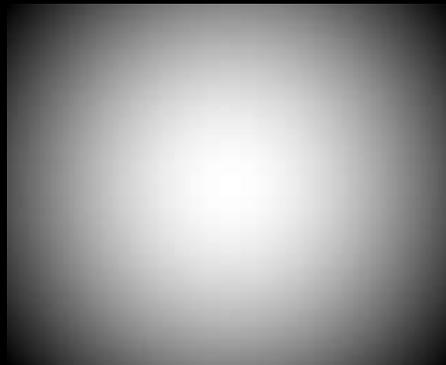
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THEORETIC ILLUMINATION FIELD



FLAT FIELD CORRECTION

1) $I = (F + B) IF$

2) $I_{ff} = I / IF = (F + B)$

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Vignetting: real case



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FLAT FIELD CORRECTION

$$1) I = (F + B) IF + n$$

$$2) I_{ff} = I / IF = (F + B) + n'$$

Super-resolution Enhancement of Text Image Sequences

David Capel and Andrew Zisserman
Robotics Research Group
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Abstract

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Vignetting



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computing a maximum a posteriori (MAP) estimate. The traditional approach is to model the texture as a first-order, stationary Markov Random Field (MRF). We propose a MAP estimator which uses the Huber edge-penalty function, and compare this with regularization based on the total variation norm.

In this work our target images are of text, and the image to image transformation is a planar projective transformation. This is the most general transformation required to



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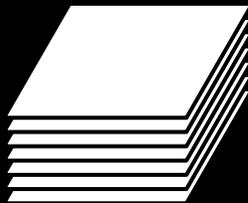


FLAT FIELD CORRECTION

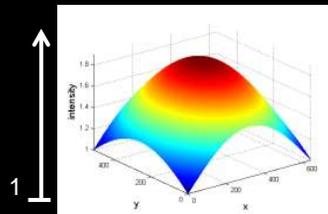
$$I_{ff} = I / IF$$

Typically, the illumination field is detected by an empty field (for example a white paper) acquired in advance and often temporally averaged (Light Field, LF)

I) LF = temporal mean (EF_i)



II) IF = LF / min (LF)



FLAT FIELD CORRECTION

$$I_{ff} = I / IF$$

Typically, the illumination field is detected by an empty field (for example a white paper) acquired in advance and often temporally averaged (Light Field, LF)

PROBLEM:

To know the illumination field is not always possible, for examples in real time applications or when the acquisition device is not available

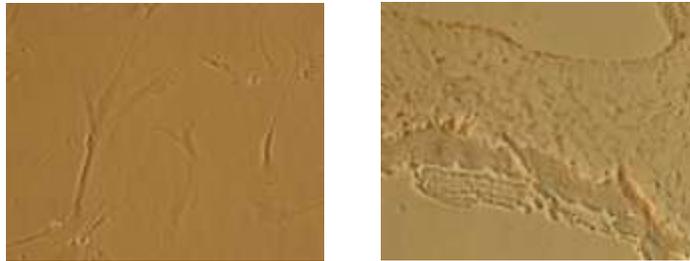
SOLUTION?

VIGNETTING FUNCTION FROM IMAGES' BACKGROUND REGIONS



IN OPTICAL MICROSCOPY

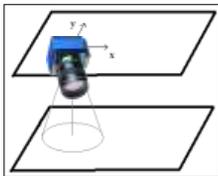
Cell cultures and histological examinations cover the most relevant part of the biological routine examinations



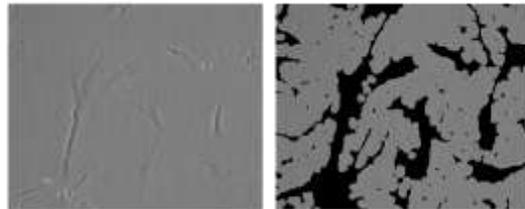
In this kind of images the background (the culture's medium and the specimen support, respectively) can be considered as being uniform



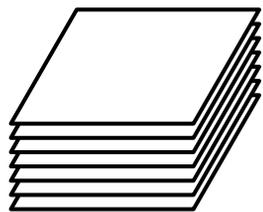
STAGE 1
image acquisition
and gray conversion



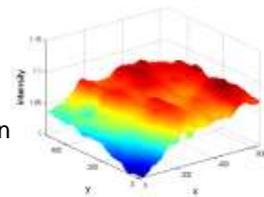
STAGE 2
background detection



STAGE 3
temporal
median filter

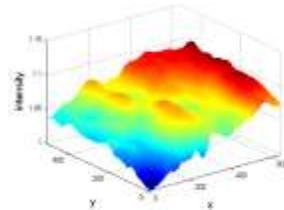
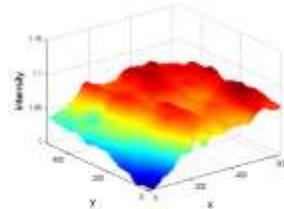


STAGE 4
illumination
field estimation

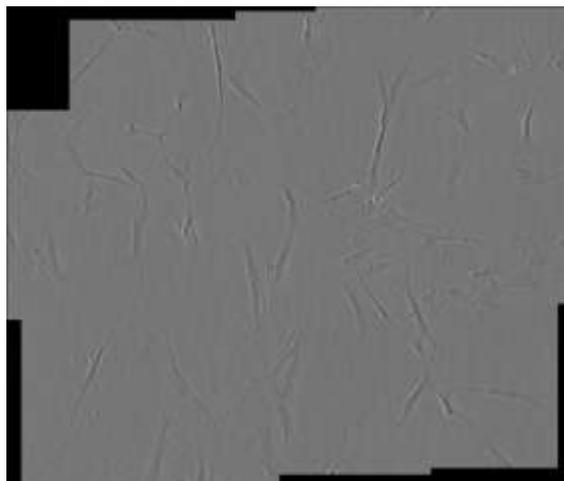
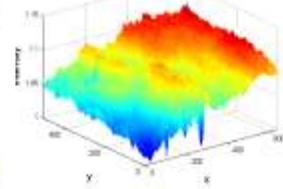




STAGE 4: illumination field estimation



Ground Truth



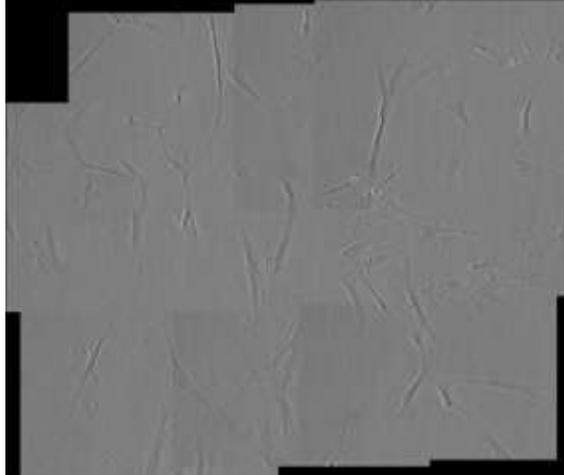
Mosaic with flat field correction



Flat field correction



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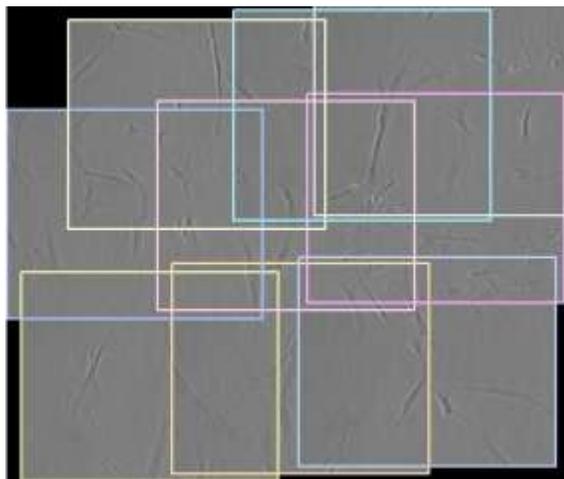
INPUT: 9 images (512 x 640). OUTPUT: mosaic (1162 x 1407)



Flat field correction



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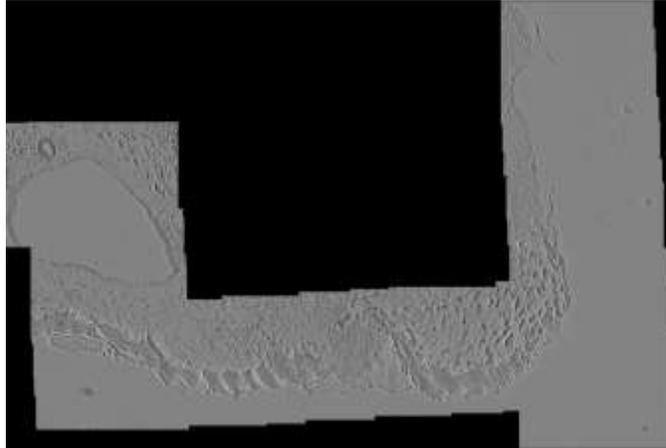
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Flat field correction



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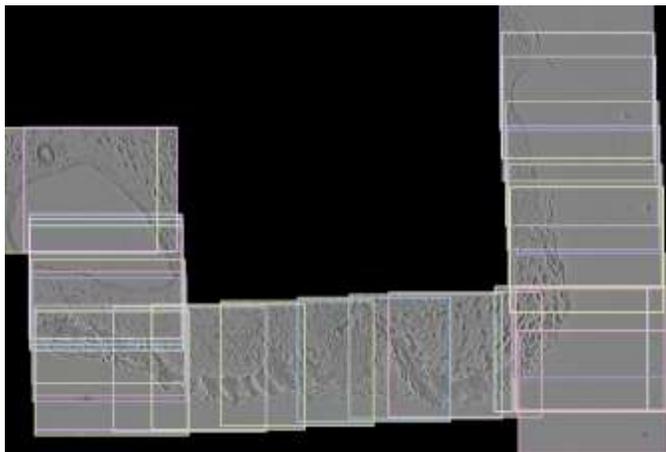
Mosaic with flat field correction



Flat field correction



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INPUT: 38 images (512 x 640). OUTPUT: mosaic (1847 x 2787)



Flat field correction – Error metrics



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- Root Mean Squared Error – RMSE

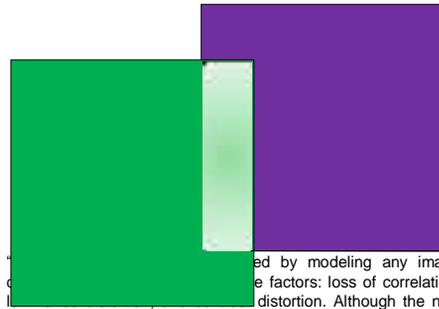
$$RMSE = \sqrt{\frac{\sum_x \sum_y (I(x,y) - R(x,y))^2}{N}}$$

- Signal to Noise Ratio – SNR

$$SNR = 10 * \log_{10} \frac{\sum_x \sum_y R(x,y)^2}{\sum_x \sum_y (I(x,y) - R(x,y))^2}$$

- Universal Quality Index – UQI (*)

$$UQI = \frac{4 * \sigma_{IR} * \bar{I} * \bar{R}}{(\sigma_I^2 + \sigma_R^2) * (\bar{I}^2 + \bar{R}^2)}$$



...ed by modeling any image
... factors: loss of correlation,
... distortion. Although the new
index is mathematically defined and no human visual system
model is explicitly employed, our experiments on various image
distortion types indicate that it performs significantly better than
the widely used distortion metrics..."

(*) Zhou Wang and Alan C. Bovik, "A Universal Image Quality Index",
IEEE Signal Processing Letters, Vol. 9, No 3, pp. 81-84, March 2002.



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| | RMSE | SNR | UQI |
|--------------------------------|------|-------|------|
| CELL w/o Flat Field | 3.76 | 30.45 | 0.76 |
| CELL Flat Field | 2.08 | 35.10 | 0.90 |
| HISTOLOGICAL w/o Flat Field | 4.75 | 29.29 | 0.93 |
| HISTOLOGICAL Flat Field | 3.25 | 32.02 | 0.96 |

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R G B images





Color vignetting correction



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Input: RGB image not flat field \longrightarrow Image_{RGB}IN = $f(R, G, B)$

RGB to GRAY conversion \longrightarrow GRAY = $0.299 * R + 0.587 * G + 0.114 * B$

Flat field correction \longrightarrow $\frac{GRAY}{IF} = 0.299 * \frac{R}{IF} + 0.587 * \frac{G}{IF} + 0.114 * \frac{B}{IF}$

Output: RGB image flat field \longrightarrow Image_{RGB}OUT = $f(\frac{R}{IF}, \frac{G}{IF}, \frac{B}{IF})$

ALGORITHM

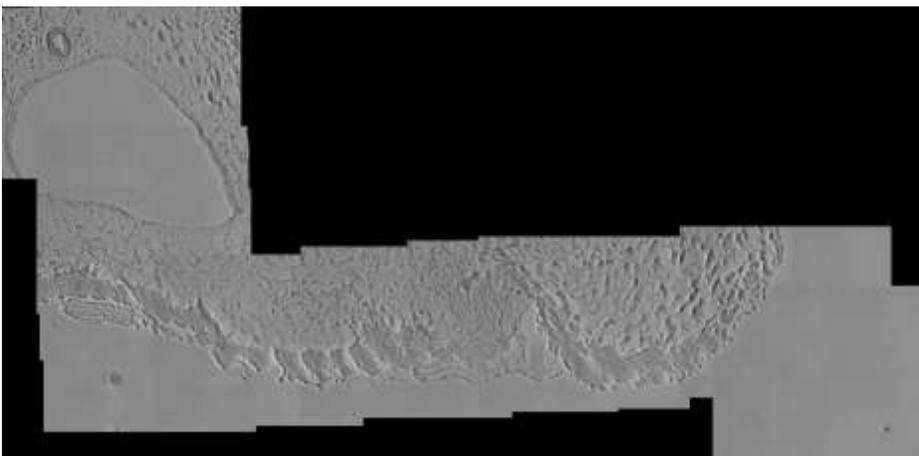
```
INPUT: RGB image not flat field
for each RGB channels {
    New channel = Old channel / Illumination Field
}
OUTPUT: RGB image flat field
```



Color vignetting correction



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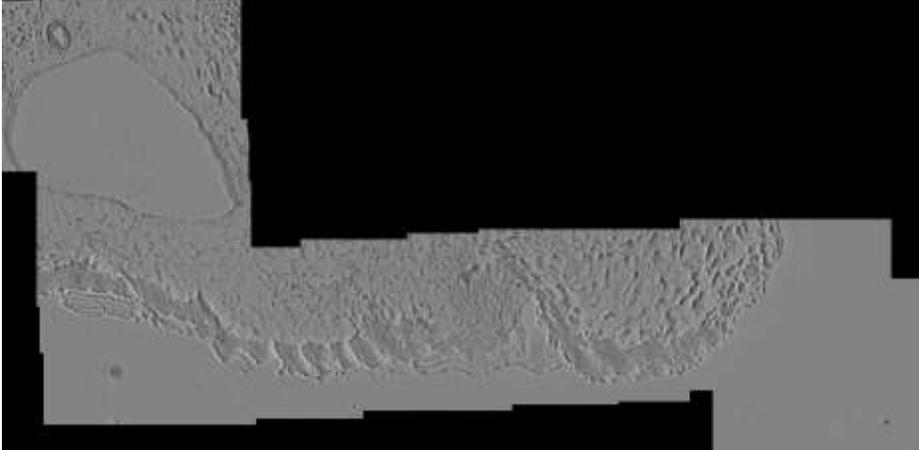
GRAY mosaic without flat field correction



Color vignetting correction



f.piccinini@unibo.it



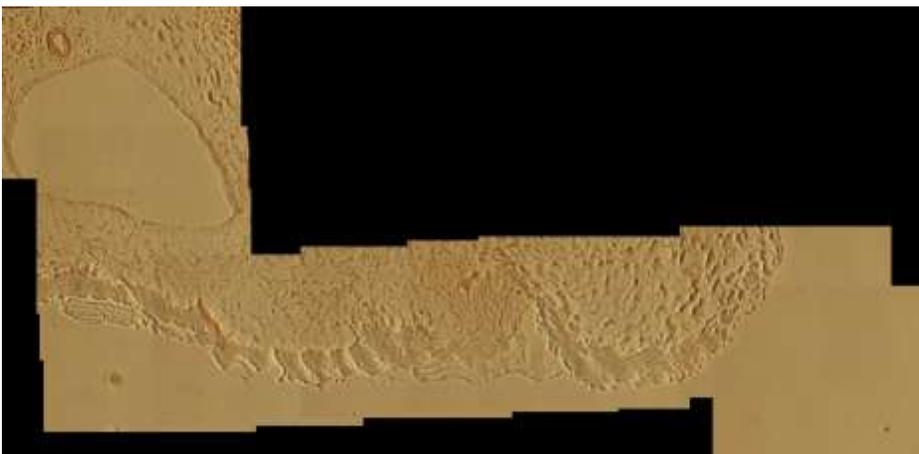
GRAY mosaic with flat field correction



Color vignetting correction



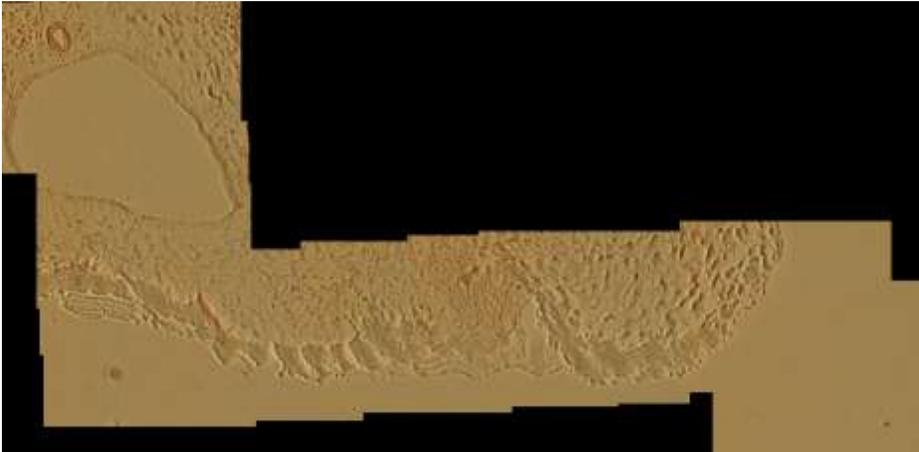
f.piccinini@unibo.it



RGB mosaic without flat field correction



Color vignetting correction



RGB mosaic with flat field correction



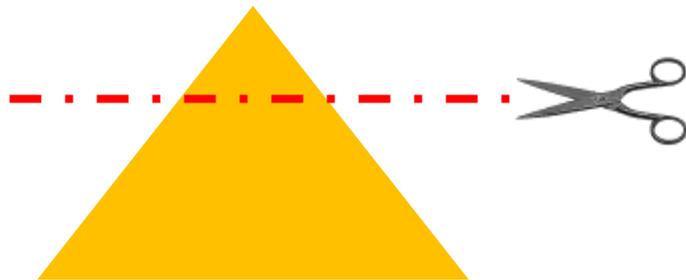
Saturation

DEFINITION: image in saturation

Image with some regions in which the intensity values are all at the maximum value

PROBLEM

- Where image is in saturation you cannot understand the original curve





Saturation

DEFINITION: image in saturation

Image with some regions in which the intensity values are all at the maximum value

PROBLEM

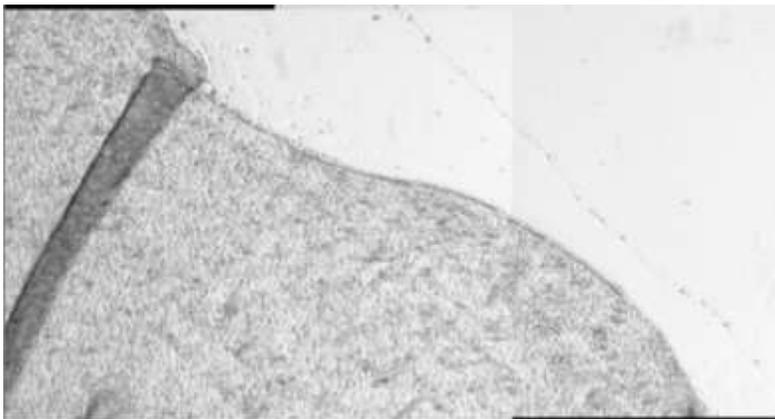
- Where image is in saturation you cannot understand the original curve
- When one or several channels are in saturation the color flat field correction generates incoherent colors

IMPORTANT NOTE

If there is even a channel not in saturation, the RGB to GRAY conversion generates a output image not in saturation and, in this case, there aren't any problems with the GRAY flat field correction.



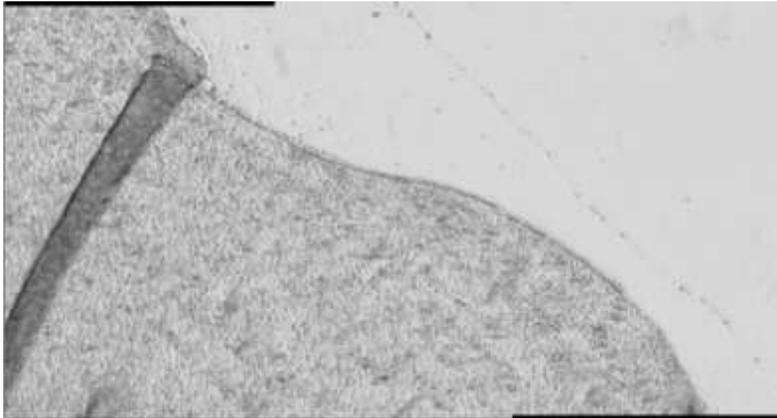
Correction with saturation



GRAY mosaic without flat field correction



Correction with saturation



GRAY mosaic with flat field correction



Correction with saturation



RGB mosaic without flat field correction



Correction with saturation



RGB mosaic with flat field correction



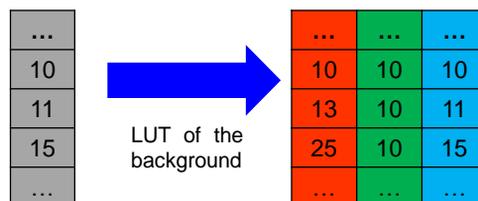
Correction with saturation

IDEA

Use a dynamic remapping color LUT (Look Up Table) to force that for each gray level there is only a RGB conversion color.

PRACTICAL SOLUTION

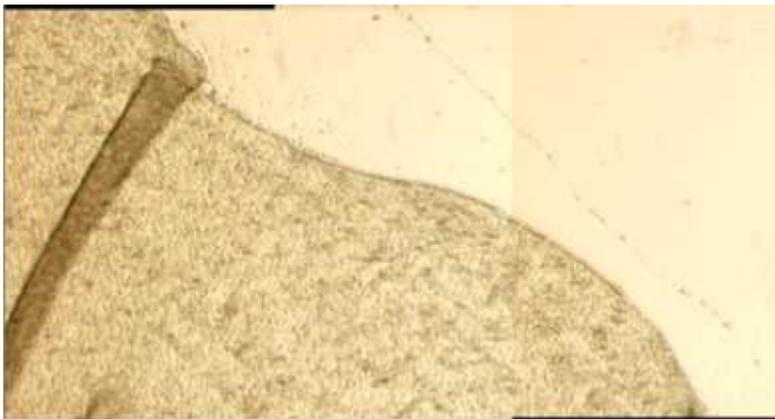
During mosaicing operation, when is detected a image in saturation, LUTs are calculated by comparing flat field mosaic in gray and in RGB (with incoherent colors due to the flat field correction applied on images in saturation). One LUT is referred to background regions and the other LUT is referred to foreground region. For each GRAY level in each region, one and only one RGB conversion is permitted. To select the conversion, a mean value is calculated for each channel, of the RGB mosaic, with the same GRAY conversion. LUT are dynamical improved with the new images added to the mosaic .





ALGORITHM

```
when RGB image in saturation is added to mosaic {  
  for each intensity levels in GRAY mosaic {  
    LUTfg = a look up table from foreground is calculated  
    LUTbg = a look up table from background is calculated  
  }  
  RGB mosaic is remapped using the two LUT  
}  
after for each new RGB image added to mosaic {  
  if there are new intensity levels in the flat field GRAY image {  
    LUTfg is improved  
    LUTbg is improved  
  }  
  RGB image is remapped using the two LUT  
}
```



RGB mosaic without flat field correction



Correction with saturation



RGB mosaic with flat field correction and without saturation correction



Correction with saturation



RGB mosaic with flat field correction and saturation correction



- Real time mosaicing
- Vignetting correction
- **Microscope calibration**
- Depth from focus
- Publications
- Contacts



MOTIVATIONS:

- To know the real size of the observed objects
- To understand if something in microscope it is wrong

NORMAL PROCEDURE:

Typically, when you buy a microscope with a camera, you could ask to the selling company to calibrate the system to understand the conversion factor " $\mu\text{m}/\text{pixel}$ ". It is commonly done using a set of reference gold standard grids of which they know very accurately the size of each different square.

PROBLEMS:

- Every time you change something in the setup (e.g.: move the camera) of the system the calibration should be redone
- High cost
- Time to call the technician of the selling company with the grid

OUR IDEA:

To use a secondary reference object to calibrate ourselves systems composed by microscope and camera. It permits to save money and time.



HYCOR KOVA GLASSTIC SLIDE



Disposable microscope slide made of optically clear plastic. Designed for microscopic examination of urine and other body fluids. Commonly used to count cells thanks to the printed counting grid.

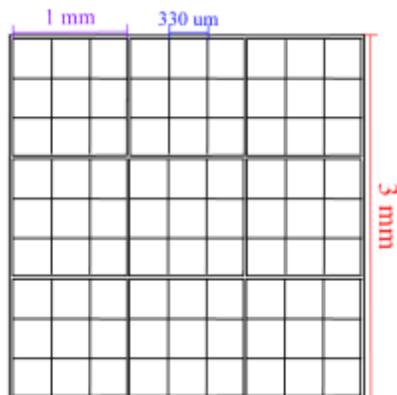
UNIT PRICE: only 1 dollar!!!



SIZE DIMENSIONS:

After a correspondence of about 7 email and several telephone calling with the selling company, and a verification test conducted with a microscope calibrated with a gold standard, we have prove that:

- smallest squared grid dimensions between line centroids: $330 \mu\text{m} \pm 6\%$
- lines dimension: $25 \mu\text{m} \pm 10\%$

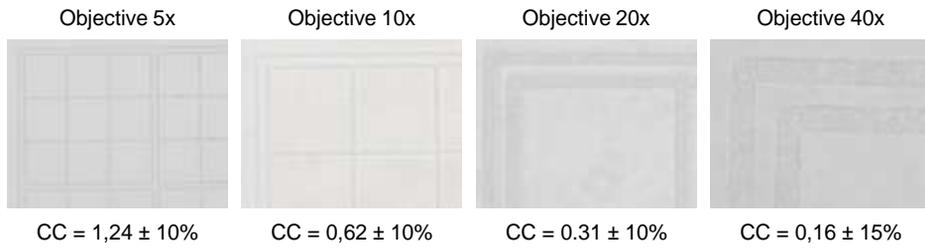
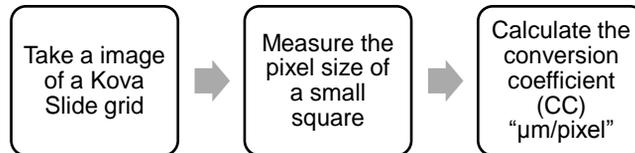




Microscope calibration

METHOD

- For each microscope's objective
- Whenever microscope's setup change

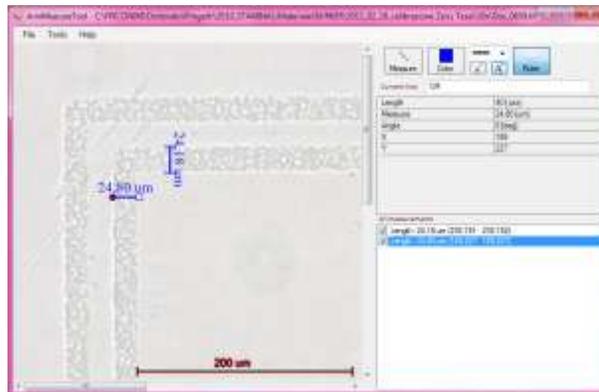


Microscope calibration

C# SOFTWARE: ANN MEASURE TOOL

User friendly image editor used to:

- Calibrate microscope system through a reference object
- Measure and label image object



Courtesy of Dr. Alessandro Gherardi



Outline



- Real time mosaicing
- Vignetting correction
- Microscope calibration
- **Depth from focus**
- Publications
- Contacts



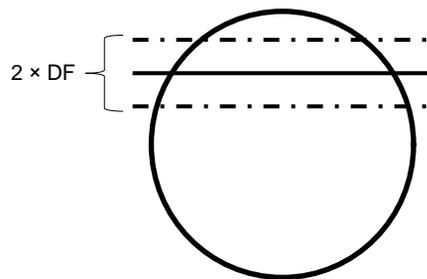
Depth from focus



DEPTH OF FOCUS (DF)

the distance over which the image plane can be displaced while a single object plane remains in acceptably sharp focus. Knowing several characteristics of camera and lenses is theoretically possible to calculate the DF.

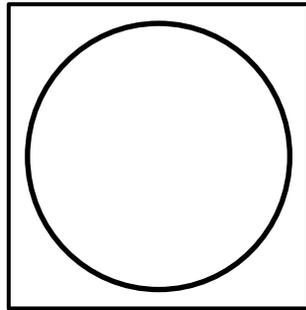
<http://www.microscopyu.com/tutorials/java/depthoffield/index.html>



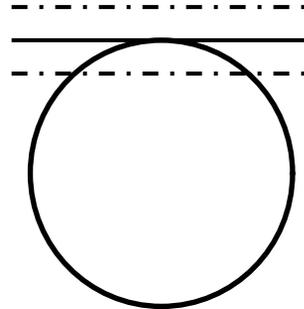


OBJECT RECONSTRUCTION

Starting from a images' stack of the same object, where images are acquired at different depth of plane (Z-plane), if the Z distance between two subsequent image is less that DF, is possible use the stack to reconstruct the single 2D image completely sharp of the object.

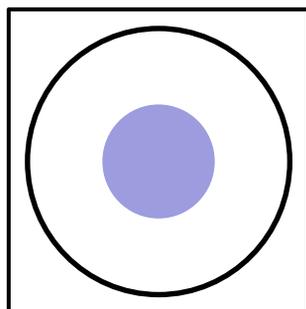


2D RECONSTRUCTED IMAGE

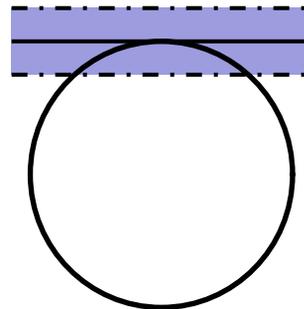


OBJECT RECONSTRUCTION

Starting from a images' stack of the same object, where images are acquired at different depth of plane (Z-plane), if the Z distance between two subsequent image is less that DF, is possible use the stack to reconstruct the single 2D image completely sharp of the object.



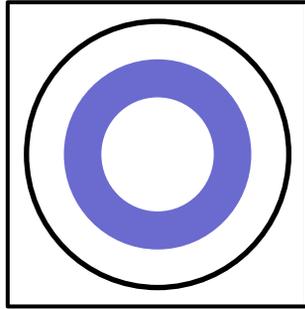
2D RECONSTRUCTED IMAGE



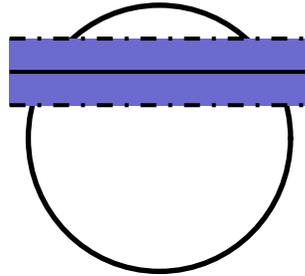


OBJECT RECONSTRUCTION

Starting from a images' stack of the same object, where images are acquired at different depth of plane (Z-plane), if the Z distance between two subsequent image is less that DF, is possible use the stack to reconstruct the single 2D image completely sharp of the object.

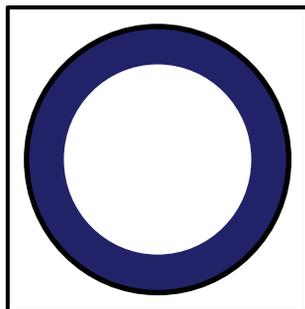


2D RECONSTRUCTED IMAGE

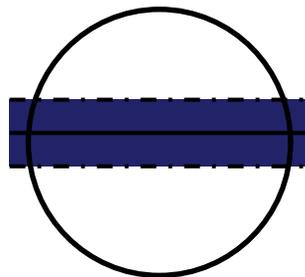


OBJECT RECONSTRUCTION

Starting from a images' stack of the same object, where images are acquired at different depth of plane (Z-plane), if the Z distance between two subsequent image is less that DF, is possible use the stack to reconstruct the single 2D image completely sharp of the object.



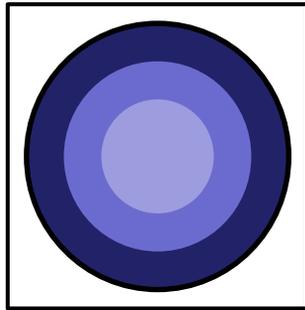
2D RECONSTRUCTED IMAGE



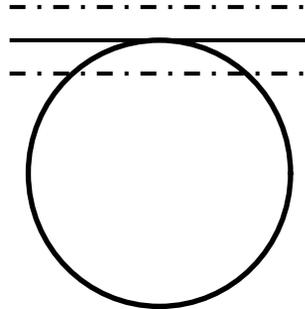


OBJECT RECONSTRUCTION

Starting from a images' stack of the same object, where images are acquired at different depth of plane (Z-plane), if the Z distance between two subsequent image is less that DF, is possible use the stack to reconstruct the single 2D image completely sharp of the object.



2D RECONSTRUCTED IMAGE



BIOLOGICAL APPLICATIVE CONTEST

PROJECT STAMINAL

Characterization of stem cells through automatic analysis of microscopic images in preclinical therapy.

Collaboration with the Istituto Scientifico Romagnolo per lo Studio e la Cura dei Tumori, IRST (Meldola, FC, Italy)

The IRST Pre-Clinical Pharmacology Lab is the first to isolate of stem cells from normal lung tissue of adult humans and to obtain spheroids.

GOAL

To study spheroids properties and behaviors to build solid biological model, useful to study treatment and drugs' effects against cancer diseases.



PROBLEM

The obtained spheroids have a diameter much greater of the depth of focus. A single 2D image acquired with optical microscopy is not sufficient to observe a spheroid with all boundary regions sharp.

SOLUTION

2D plane spheroid reconstruction starting from a stack of images in which depth difference between Z-plane of two subsequent images are less that value of depth from focus of the system.

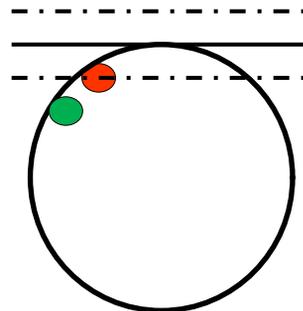
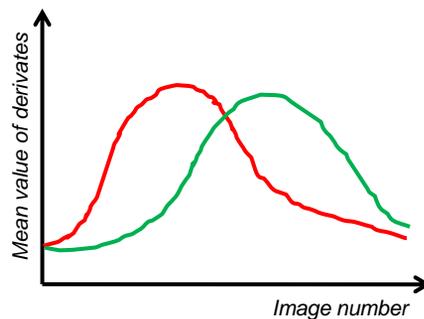
MATERIALS

Spheroid cultures and microscope system compose by non-automated inverted Zeiss Axiovert 200 microscope and Zeiss AxioCAM MRC camera. The micrometric microscope holder's shift is of $1\ \mu\text{m}$ (from data sheet). For each observed spheroid we have acquired a sequence of several images in which depth between Z-planes is $5\ \mu\text{m}$.



ALGORITHM

While an object's region is in focus, it is sharp and well defined. When the focus is lost the region became smooth. Starting from a stack of images acquired at different Z-plane positions, to understand the image in which an object's region is at the best focus, we can search in the stack the maximum of derivatives mean value of the small region.

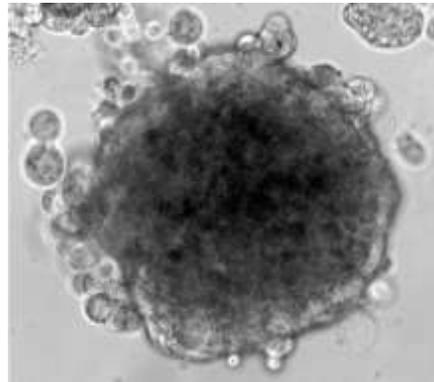




Depth from focus

REAL CASE

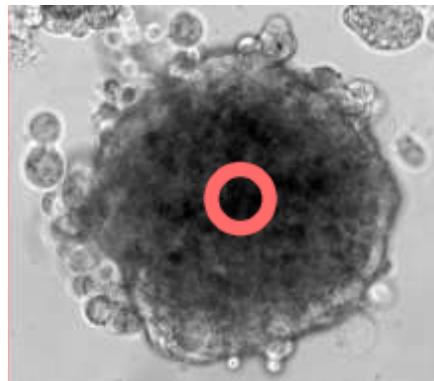
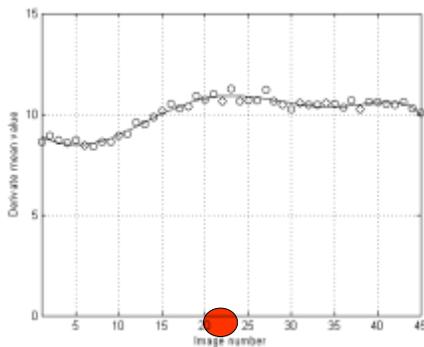
- Sequence of 45 images relative to a spheroid of lung cancer cells.
- Each images is acquired moving holder of step of $5\ \mu\text{m}$ in Z.



Depth from focus

REAL CASE

- Sequence of 45 images relative to a spheroid of lung cancer cells.
- Each images is acquired moving holder of step of $5\ \mu\text{m}$ in Z.

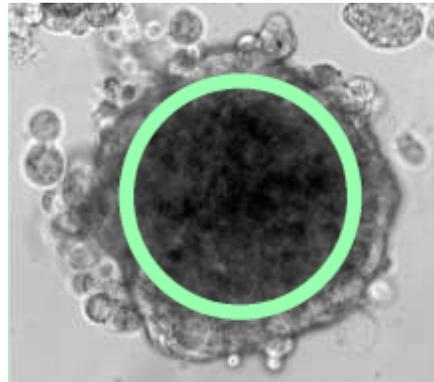
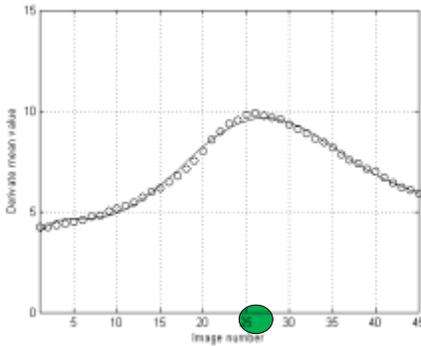




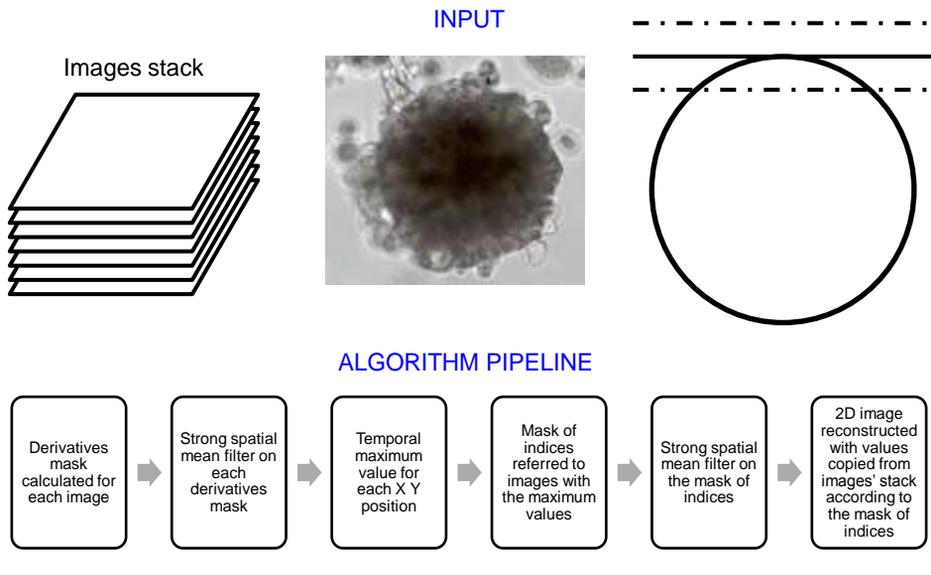
Depth from focus

REAL CASE

- Sequence of 45 images relative to a spheroid of lung cancer cells.
- Each images is acquired moving holder of step of 5 μm in Z.



Depth from focus



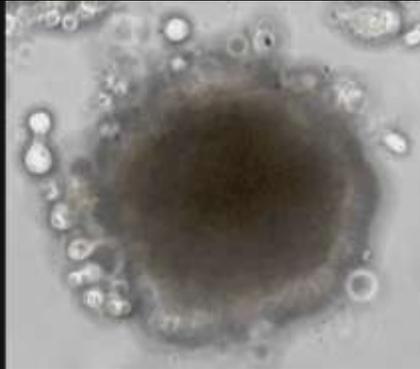


Depth from focus

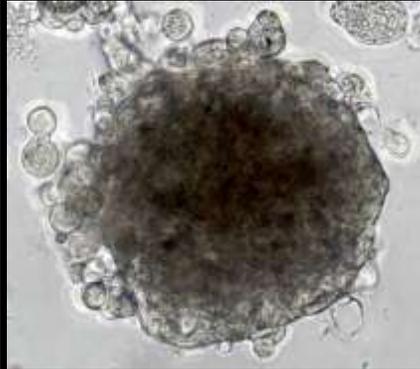


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INPUT:
IMAGES STACK



OUTPUT:
OUR METHODS

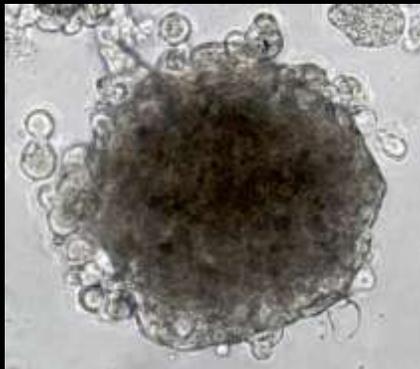


Depth from focus

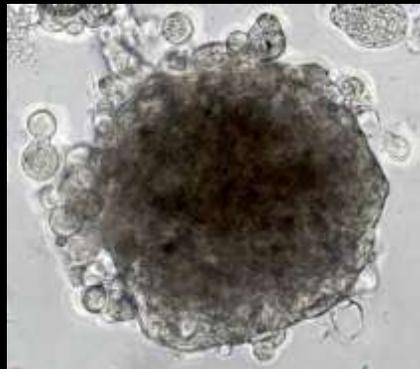


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OUTPUT:
OTHER METHODS



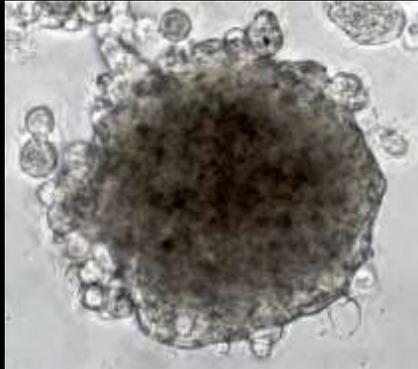
OUTPUT:
OUR METHODS



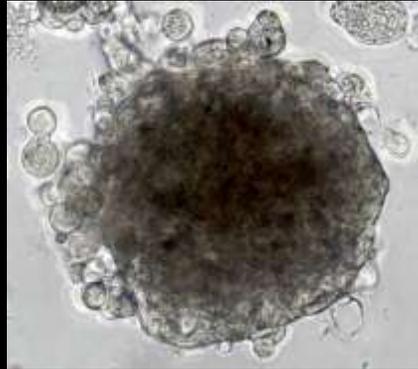
MICHAEL UMORIN IMAGEJ PLUGIN
<http://imagej.nih.gov/ij/plugins/stack-focuser.html>



OUTPUT: OTHER METHODS



OUTPUT: OUR METHODS



MICHAEL UNSER (EPFL) IMAGEJ PLUGIN
<http://bigwww.epfl.ch/demo/edi/index.html>



- Real time mosaicing
- Vignetting correction
- Microscope calibration
- Depth from focus
- **Publications**
- Contacts



Publications



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- A. Bevilacqua, F. Piccinini and A. Gherardi, "**Vignetting correction by exploiting an optical microscopy image sequence**", submitted to the 33rd Annual International Conference of the IEEE Engineering in Medicine and Biology Society (EMBS), Boston, USA, August 30 – September 3, 2011.
- A. Bevilacqua and F. Piccinini, "**Is an empty field the best reference to correct vignetting in microscopy?**", submitted to the 33rd Annual International Conference of the IEEE Engineering in Medicine and Biology Society (EMBS), Boston, USA, August 30 – September 3, 2011.
- A. Gherardi, A. Bevilacqua, and F. Piccinini, "**Illumination field estimation through background detection in optical microscopy**", IEEE Symposium on Computational Intelligence in Bioinformatics and Computational Biology (CIBCB), Paris, France, April 11-15, 2011, pp. 49-54.
- L. Carozza, A. Bevilacqua, and F. Piccinini, "**An Incremental Method for Mosaicing of Optical Microscope Imagery**", IEEE Symposium on Computational Intelligence in Bioinformatics and Computational Biology (CIBCB), Paris, France, April 11-15, 2011, pp. 55-60.

IN PREPARATION FOR SUBMISSION TO JOURNAL OF MICROSCOPY

A. Bevilacqua, F. Piccinini and A. Gherardi, "**Vignetting correction by exploiting an optical microscopy image sequence**", extension of article submitted to the 33rd Annual International Conference of the IEEE Engineering in Medicine and Biology Society (EMBS).



Outline



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- Real time mosaicing
- Vignetting correction
- Microscope calibration
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- Publications
- **Contacts**



Contacts



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- Filippo Piccinini, web site: www.filippopiccinini.altervista.org
- Prof. Alessandro Bevilacqua, email: alessandro.bevilacqua@unibo.it
- Computer Vision Group, University of Bologna: lab and web



<http://www.ing2.unibo.it/Ingegneria+Cesena/Studenti/Laboratori>



<http://cvg.deis.unibo.it/>

THANK YOU!