



FACULTAD DE MEDICINA
UNIVERSIDAD DE CHILE



LA SERENA SCHOOL
FOR DATA SCIENCE
Applied Tools for Data-driven Sciences

• AURA Campus
La Serena - Chile •

MAURICIO CERDA

LAB "BIO-RELATED": IMAGE PROCESSING METHODS FOR MICROSCOPY IMAGING

- La Serena, 8/24/2017 -

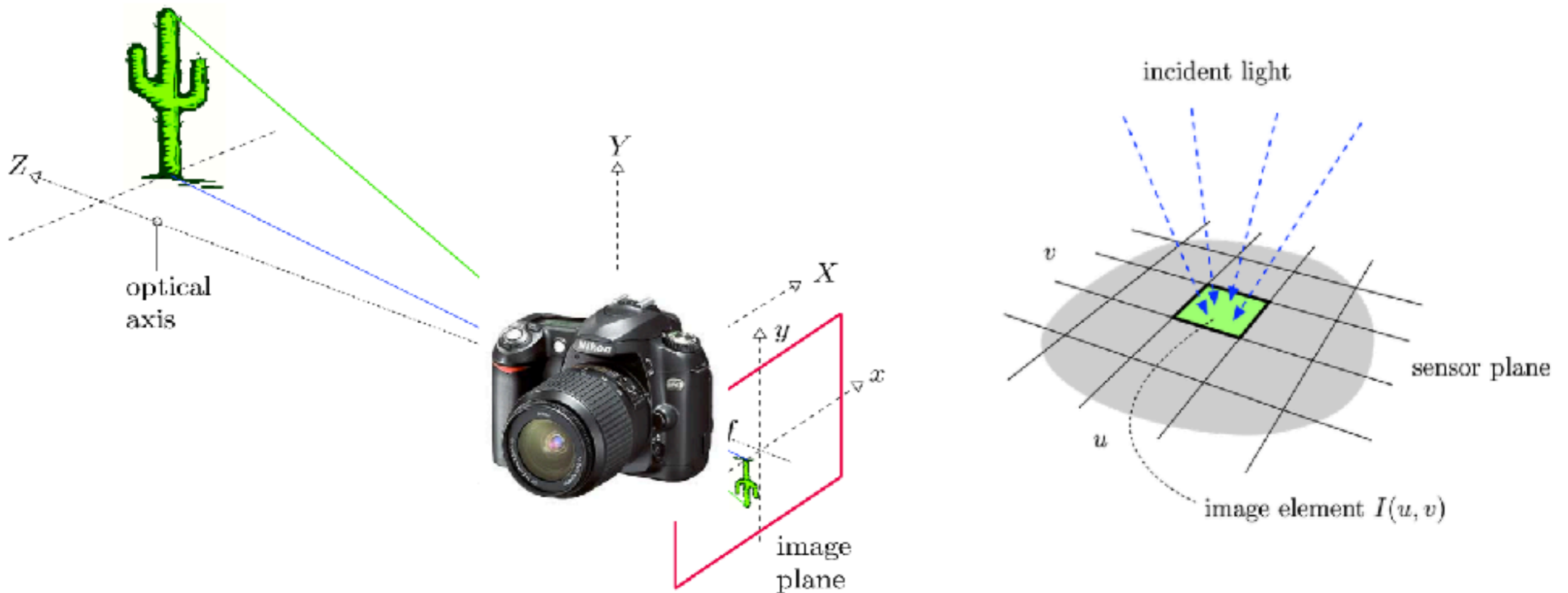


OUTLINE

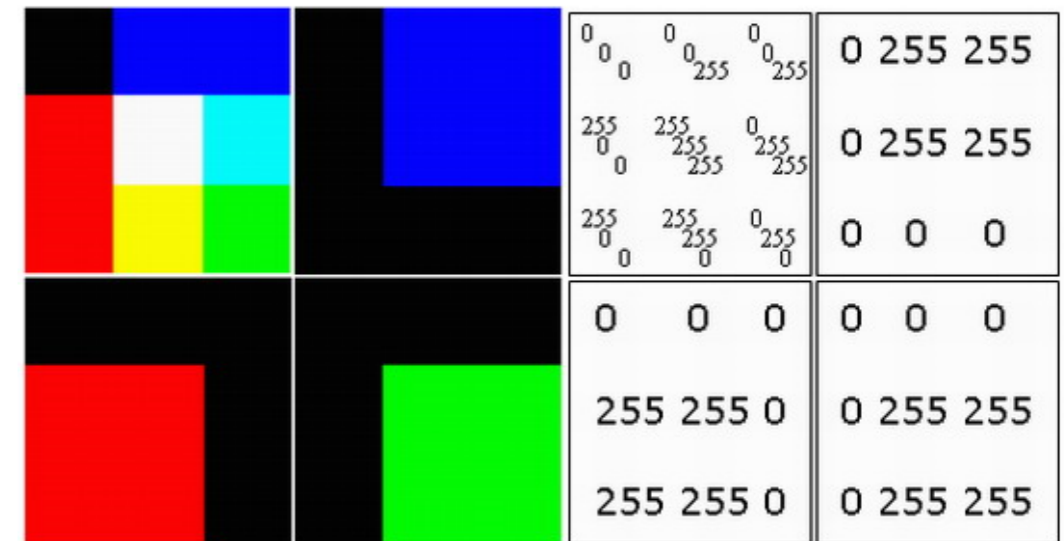
- ▶ Image processing?
- ▶ Segmentation (clustering)
- ▶ Shape description (PCA)
- ▶ Lab: challenge !

IMAGE PROCESSING: IMAGE

- ▶ **Image:** is an artifact that depicts visual perception, for example, a photo or a two-dimensional picture, that has a similar appearance to some subject—usually a physical object or a person, thus providing a depiction of it [wikipedia, 2017].



- ▶ Raster image:
 - ▶ Array or matrix of pixels with spatial coordinates $I(x,y)$.
 - ▶ A numerical value or color per location.



- ▶ Vectorial image:
 - ▶ Defined by basic functions (points, lines), instead of pixels.
 - ▶ To be displayed on screen needs to be rasterized (transforms to a raster image).
 - ▶ Raster or Vectorial in a paper?

Scalable Vector Graphics

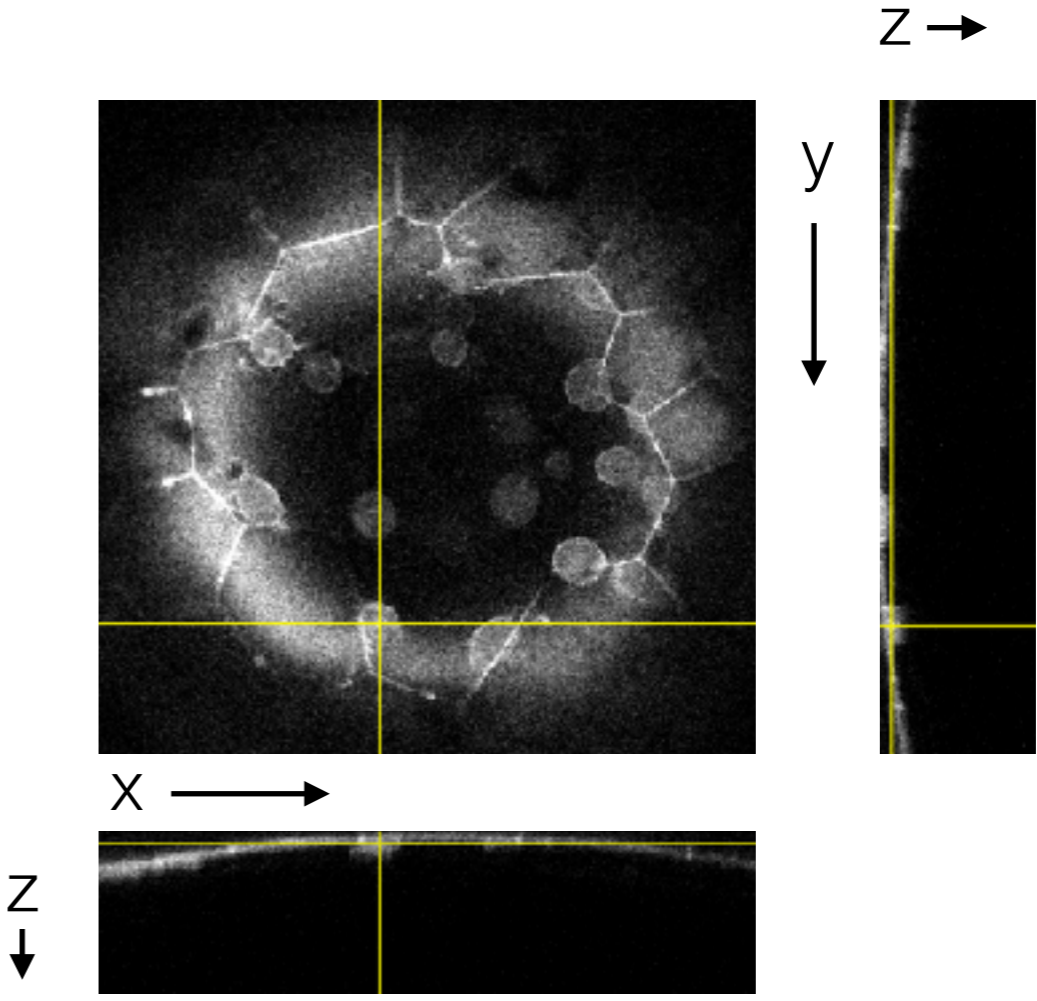
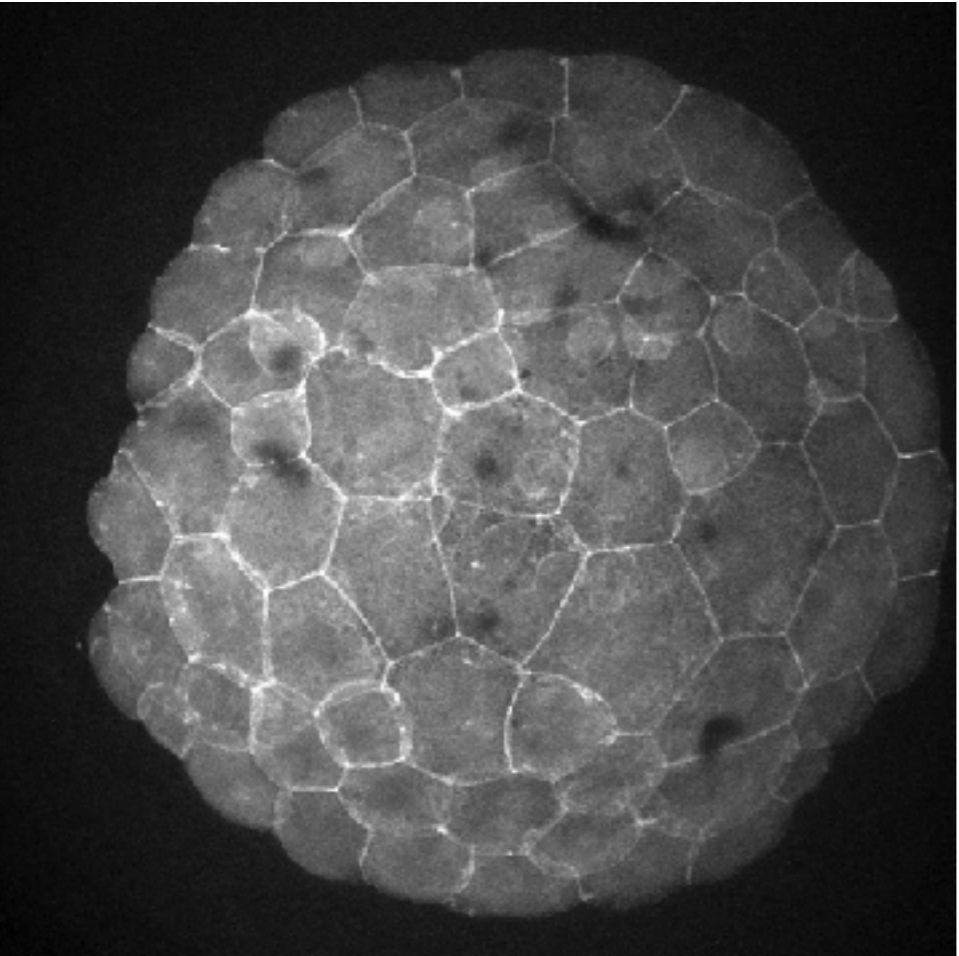
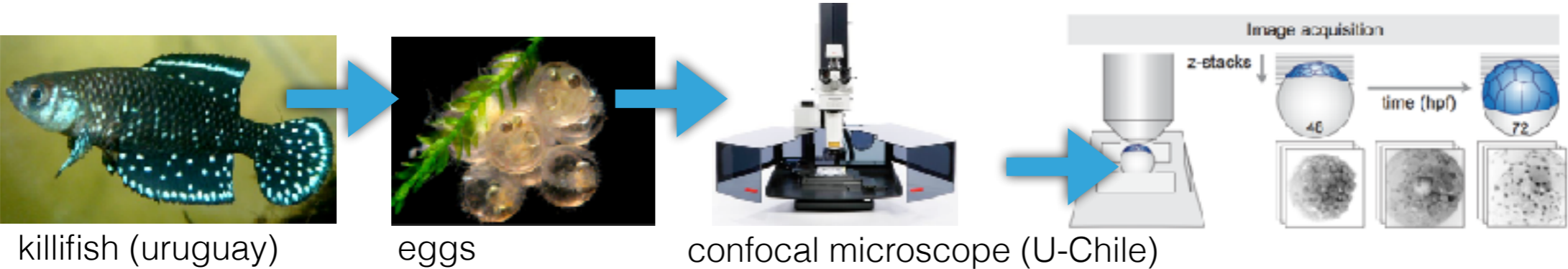


```
<?xml version="1.0" encoding="UTF-8" standalone="no" ?>
<svg version="1.0" xmlns="http://www.w3.org/2000/svg" ?>
  <defs>
    <linearGradient x1="99.7" x2="0" y1="0" y2="0" ?>
    </defs>
    <use xlink:href="#box_gr" ?>
    <use xlink:href="#circle" ?>
    <use xlink:href="#circle" ?>
    <line x1="100" y1="300" ?>
    <!--add more content here-->
    <circle cx="90" cy="300" r="10" ?>
  </svg>
```



- ▶ We want to process images to understand biological systems (models):
 - ▶ Tissue development
 - ▶ Medical imaging (parasites counting)

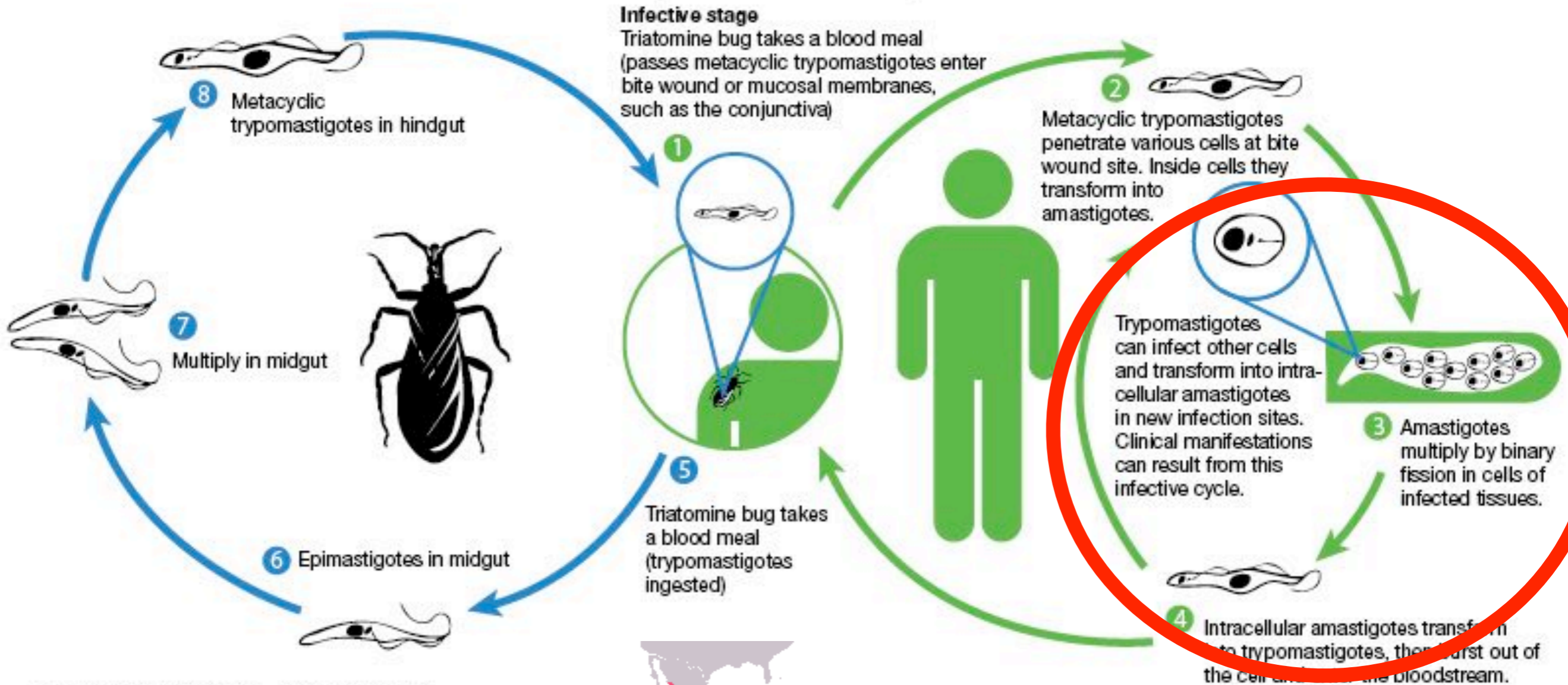
IMAGE PROCESSING: TISSUE DEVELOPMENT






Austrolebias nigripinnis 48-72 HPF, membrane marked with actine EGFP.

▶ How cells migrate over other cells?

Infection cycles of Chagas disease



-  Trypomastigoten = mobile pathogen
-  Amastigoten = immobile pathogen
-  Epimastigoten = divisible pathogen



Source: www.dpd-cdc.gov/dpdx



IMAGE PROCESSING: PARASITES

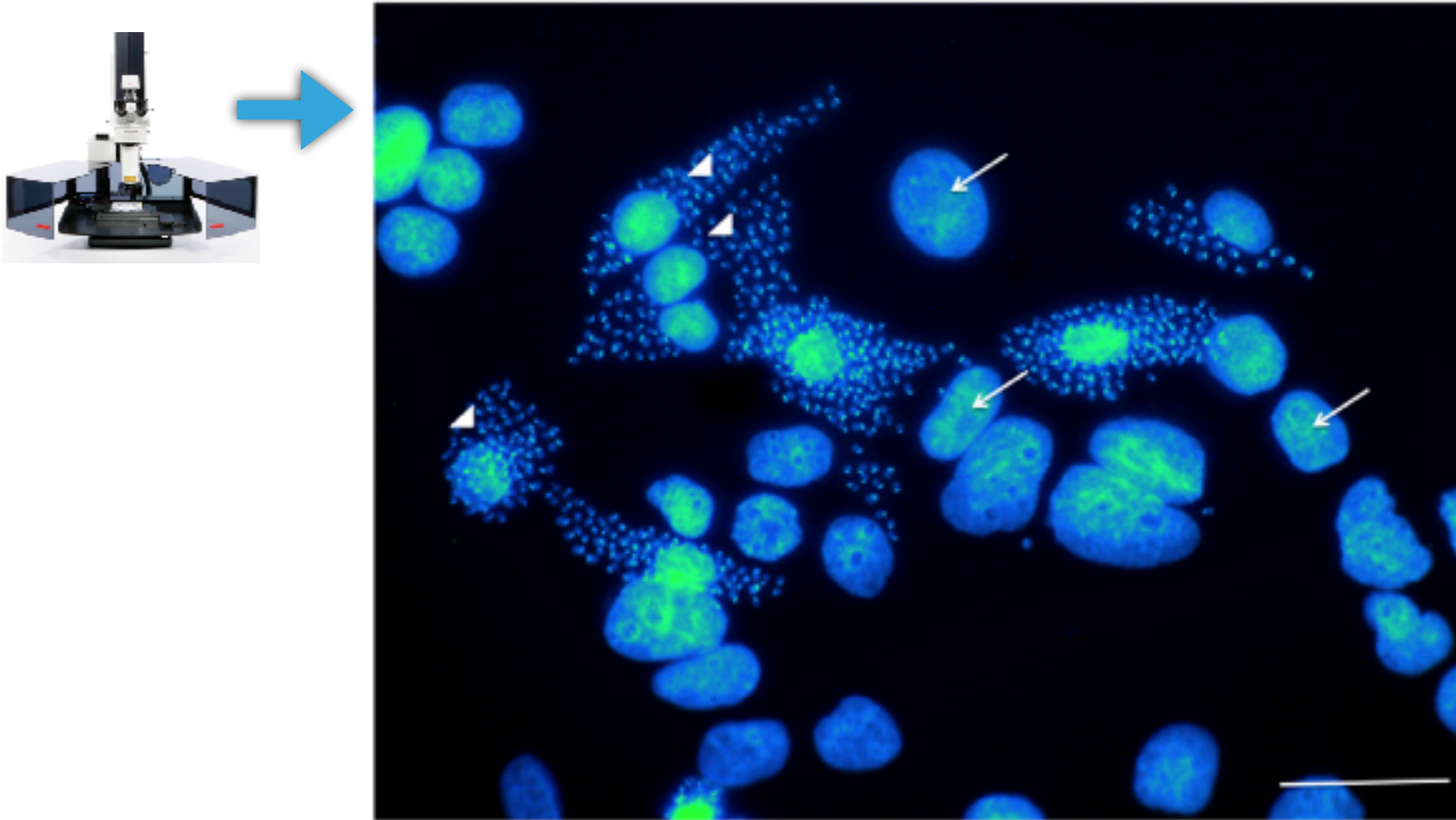
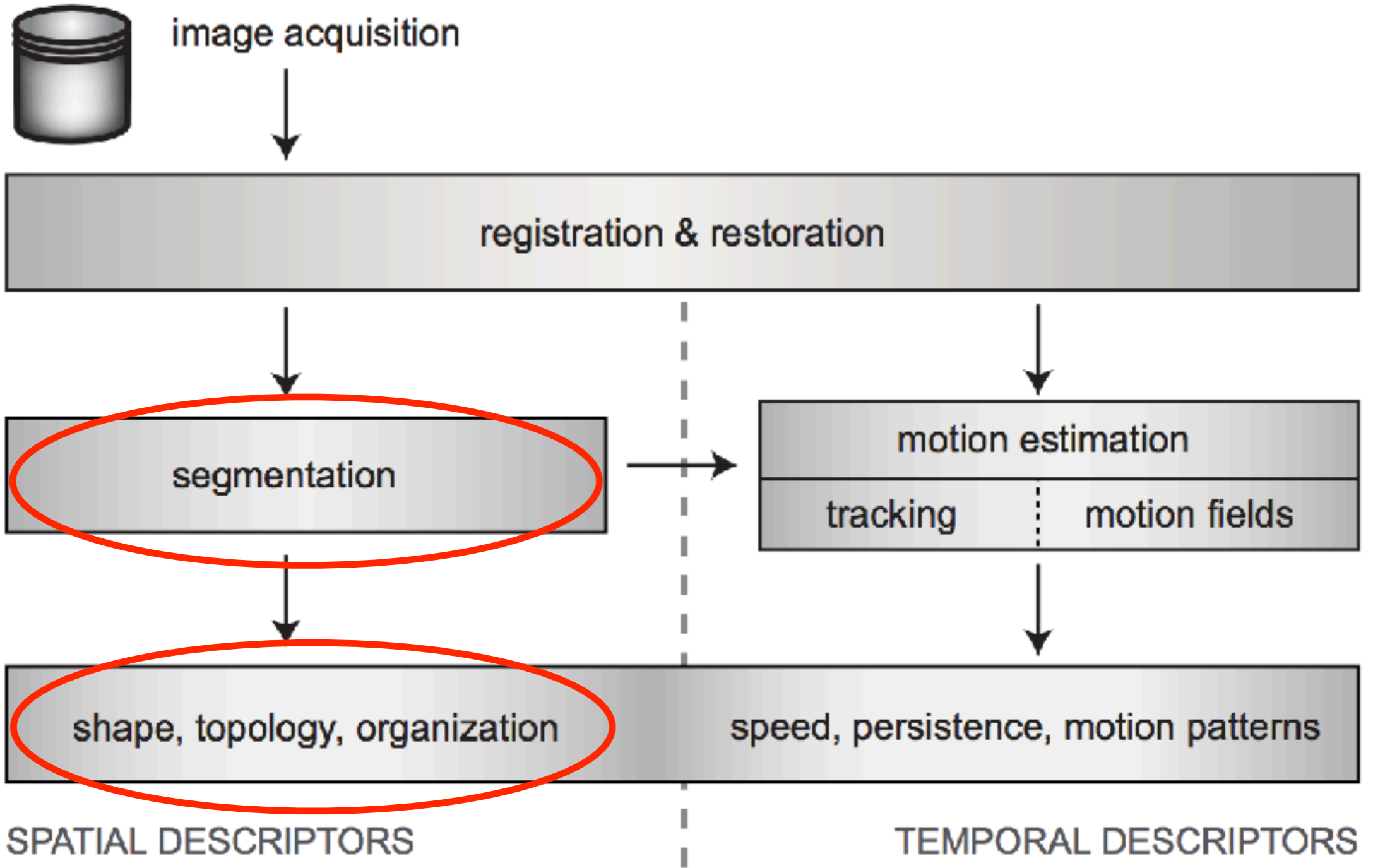


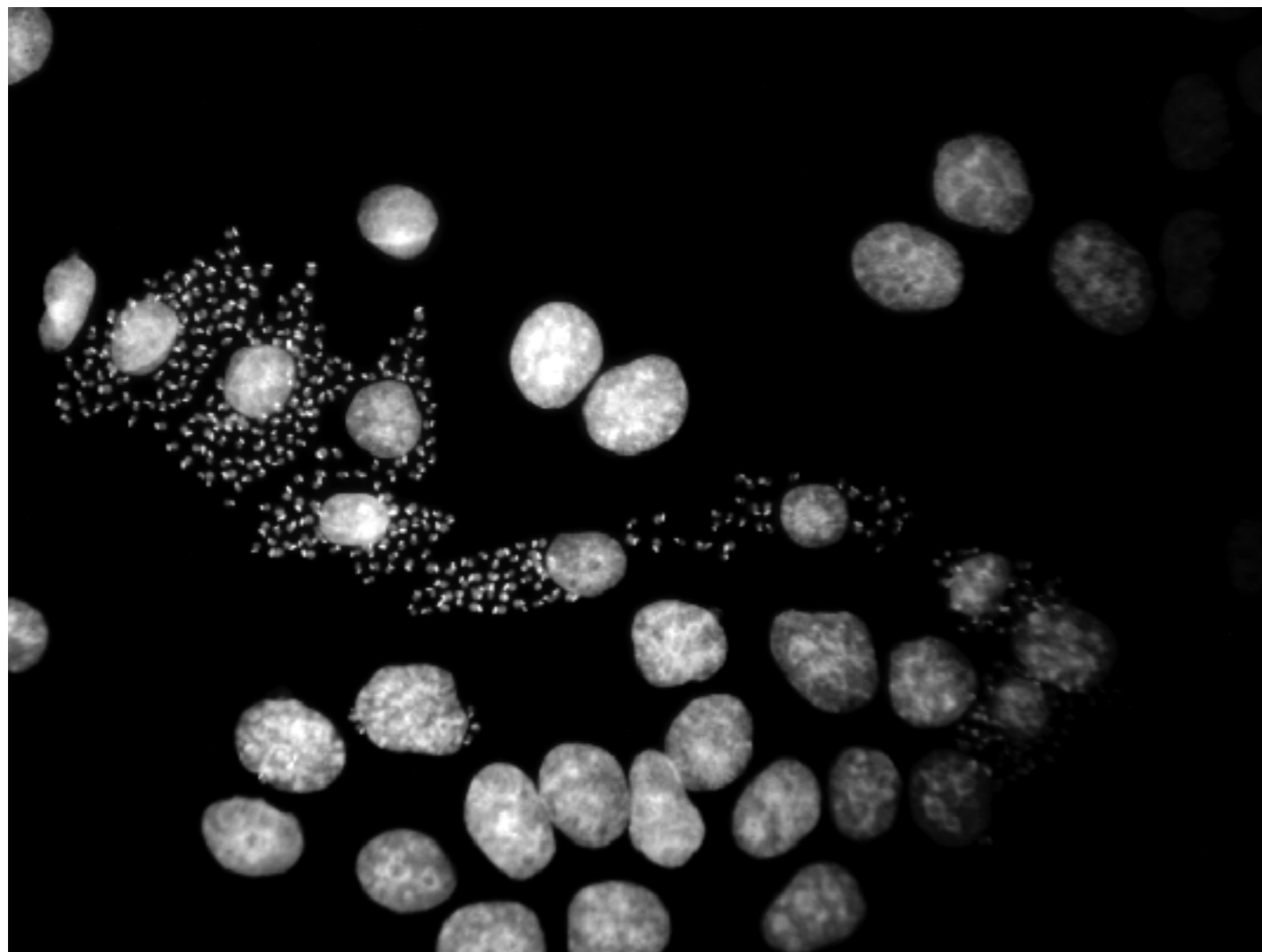
Fig. 1. Infection of BeWo cells with *T. cruzi* amastigotes. BeWo cells were challenged with *T. cruzi* Ypsilon strain trypomastigotes at a parasite:cell ratio of 1:1 for 24 h and were processed for DAPI staining after 48 h. The arrows show BeWo cell nuclei, and the arrowheads show intracellular amastigotes. Scale bar: 10 μm .

► Pregnancy?

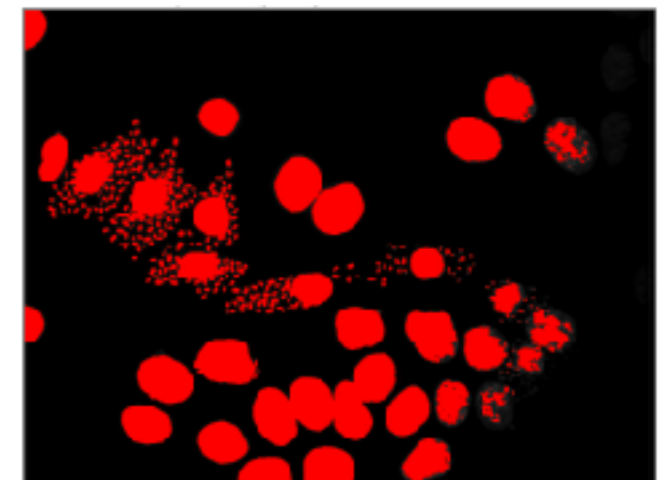
IMAGE PROCESSING: PIPELINE



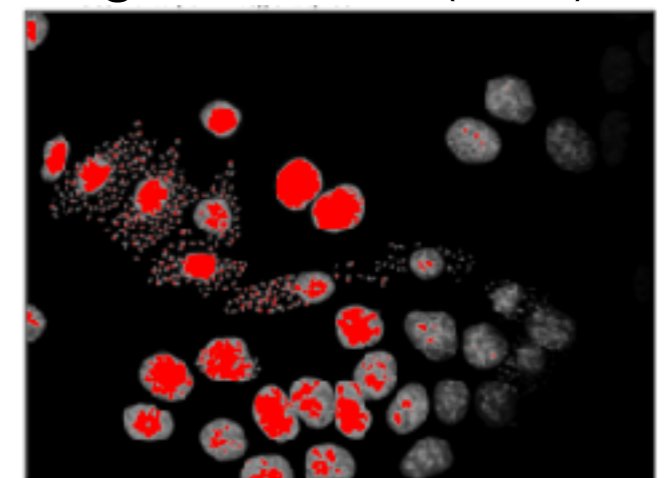
- ▶ The simplest segmentation... a manual global threshold [demo FIJI].



raw image



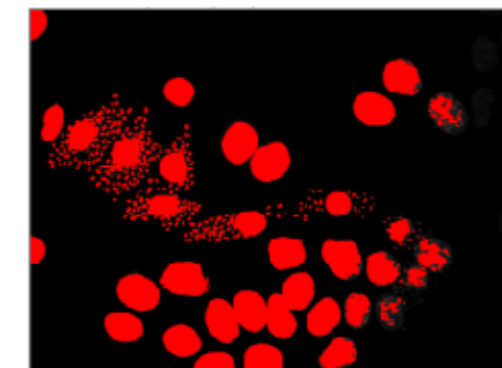
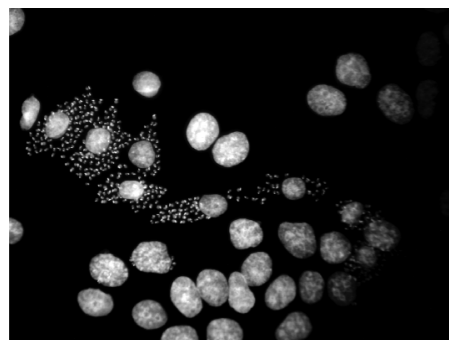
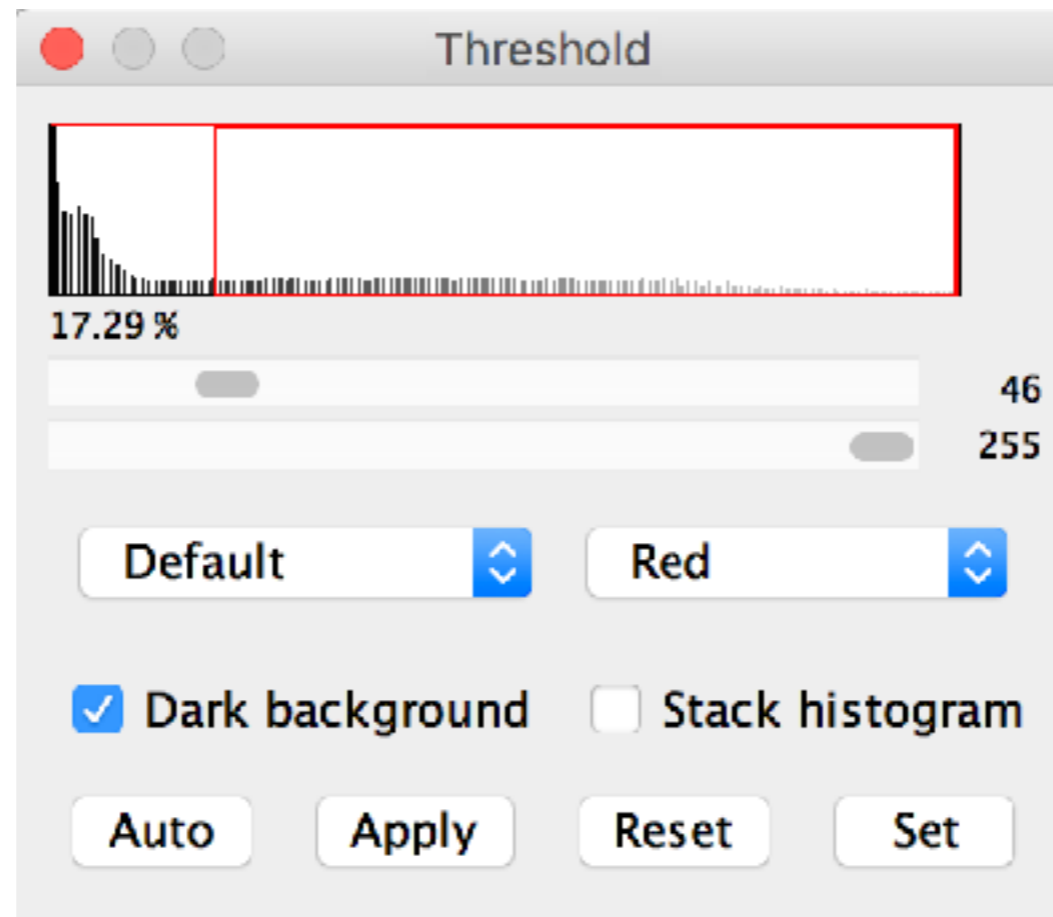
segmentation (>46)



segmentation (>158)

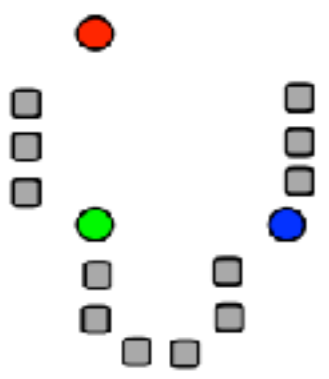
IMAGE SEGMENTATION

- ▶ But, it looks like ? ...

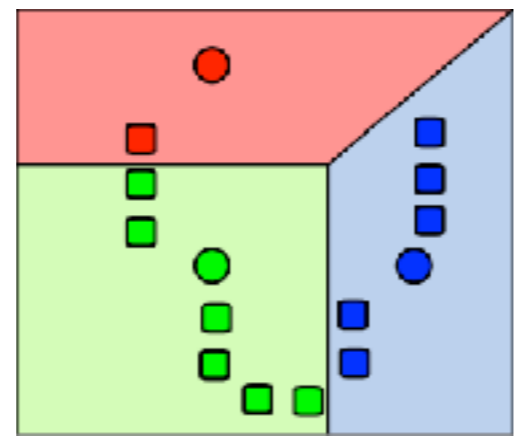


- ▶ We will start with two objects: cells, and background.
- ▶ We don't have examples (!)
- ▶ This is another kind of learning problem:
 - ▶ Supervised: regression, classification
 - ▶ Unsupervised: **clustering**

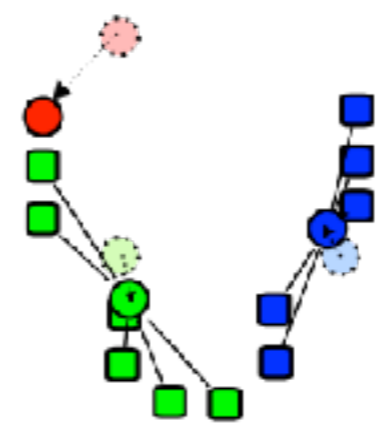
- ▶ We can model it as how to discover the best k groups or clusters at a pixel level.
- ▶ K-means clustering ($k=3$):



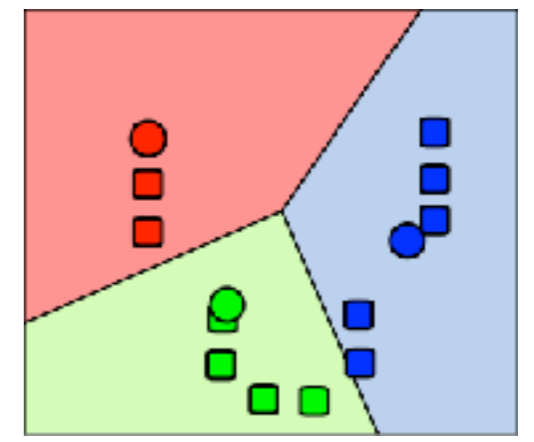
Random centroids



clusters assignation + voronoi diagram

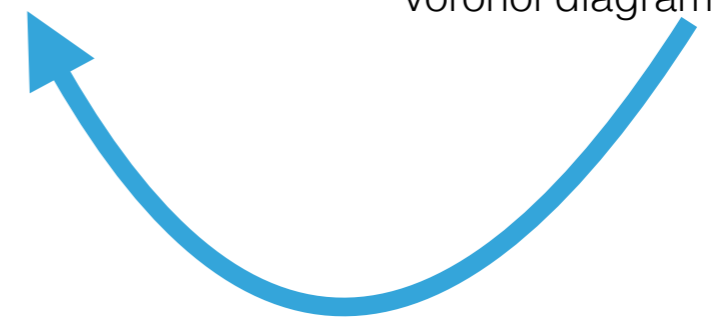


centroids re-computation



cluster assignation + voronoi diagram

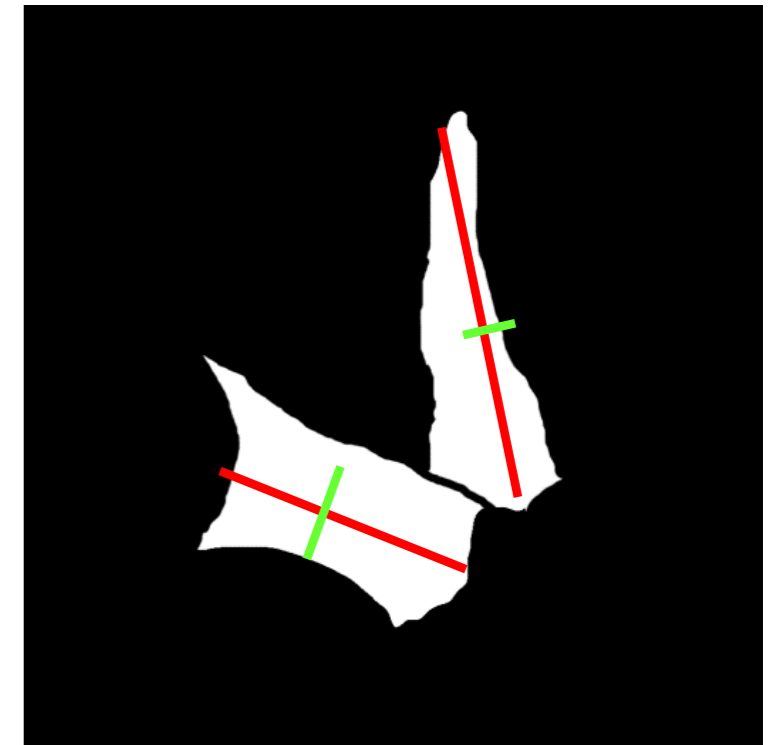
- ▶ K-Mean pseudocode [demo FIJI]:



- ▶ But we can also make examples! (label data)
- ▶ In that case, segmentation may be a supervised problem. Let's try to solve it as a Random Forest problem [demo FIJI].
- ▶ How can we decide?

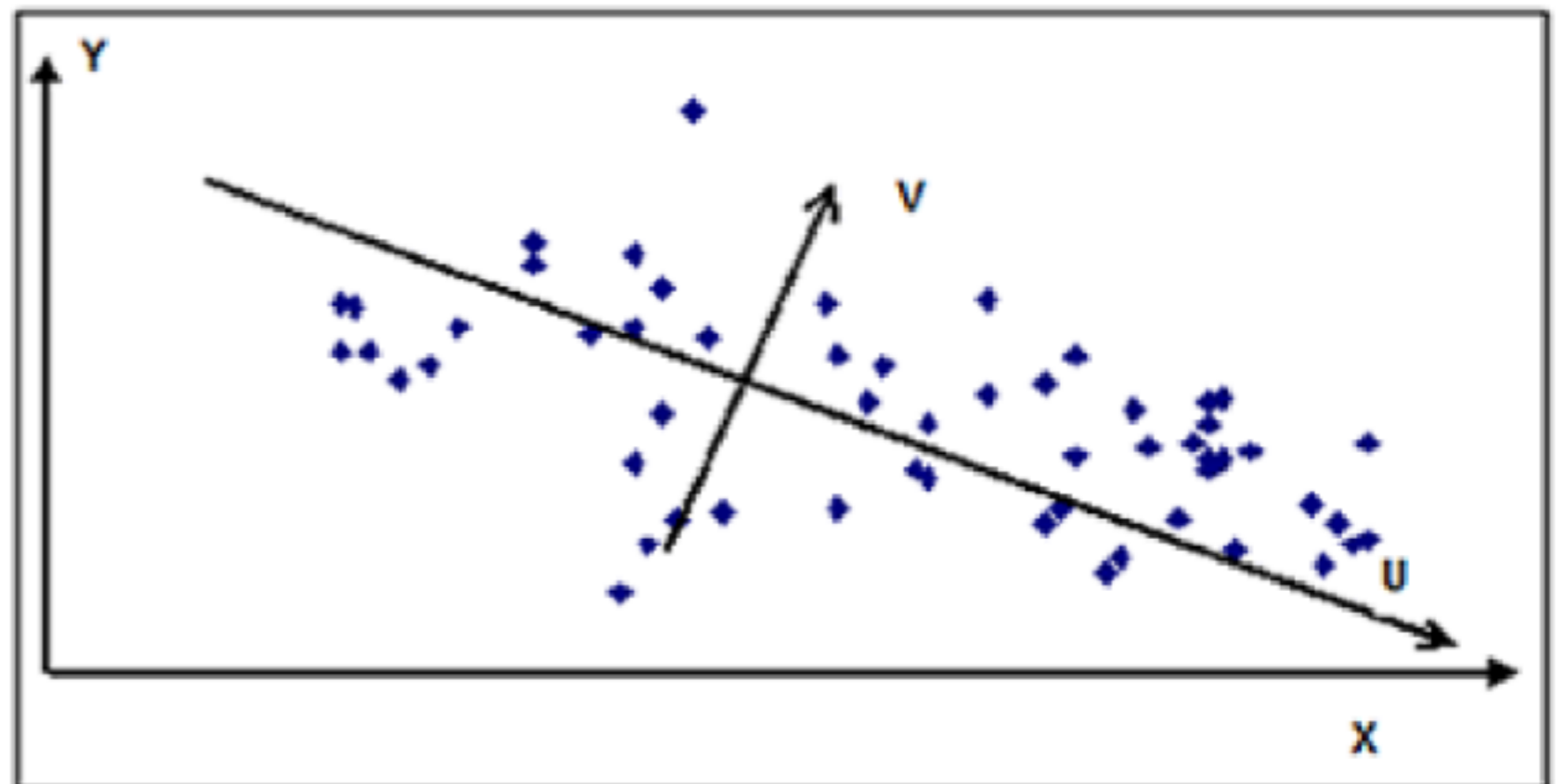
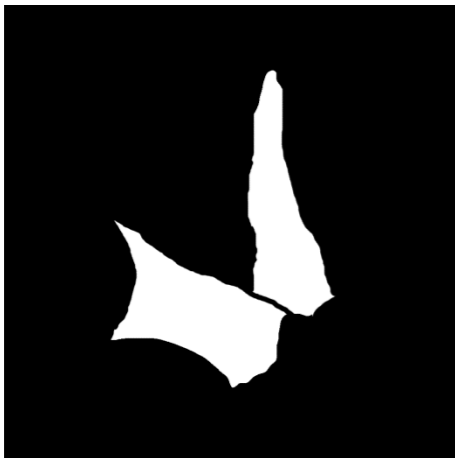
SHAPE DESCRIPTION: MOTIVATION

- ▶ If images are segmented, we can easily count objects.
- ▶ But, we cannot tell the difference between small and big or circle-like vs elongated cells.



SHAPE DESCRIPTION: IMAGE AS DISTRIBUTION

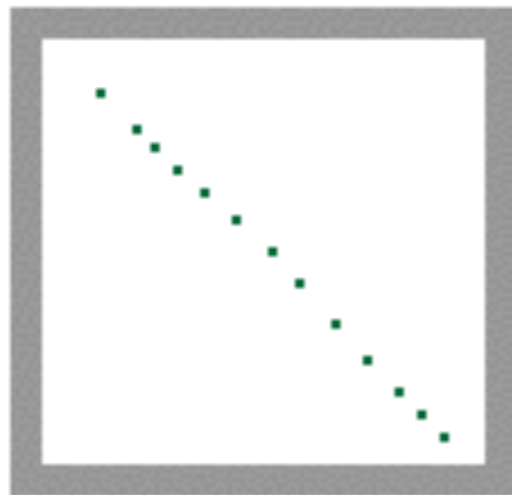
- ▶ The image as a set of pixels
- ▶ We can always find a direction to maximize variance



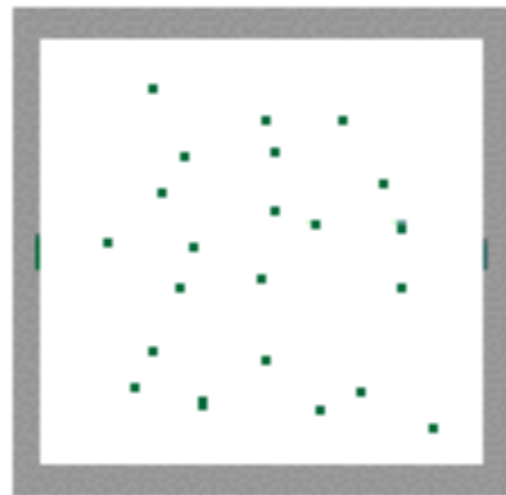
SHAPE DESCRIPTION: IMAGE AS DISTRIBUTION

- ▶ We can always find a direction to maximize variance
- ▶ Equivalent to diagonalize covariance matrix

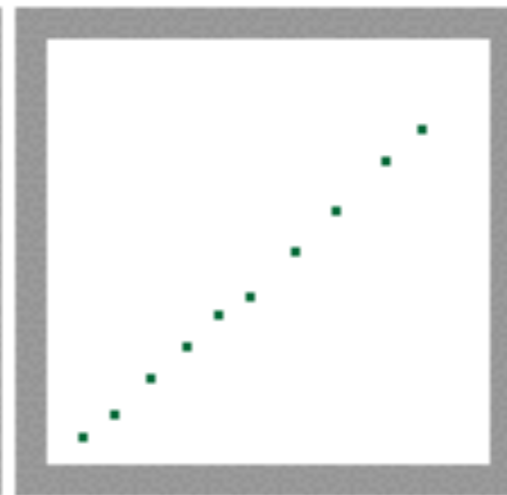
COVARIANCE



**Large Negative
Covariance**



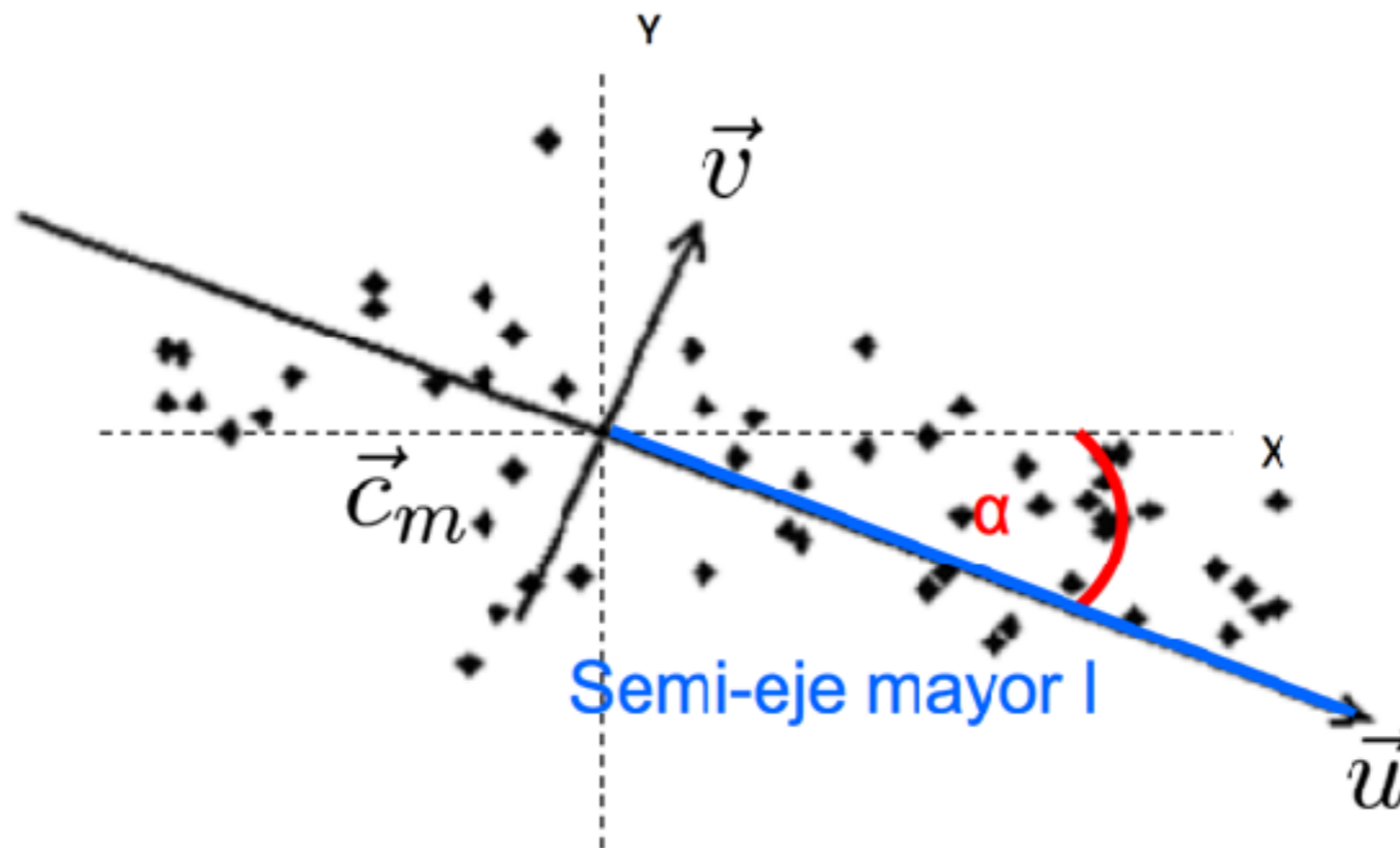
**Near Zero
Covariance**



**Large Positive
Covariance**

SHAPE DESCRIPTION: GEOMETRICAL VIEW

- ▶ We look for a rotation where covariance matrix is diagonal.



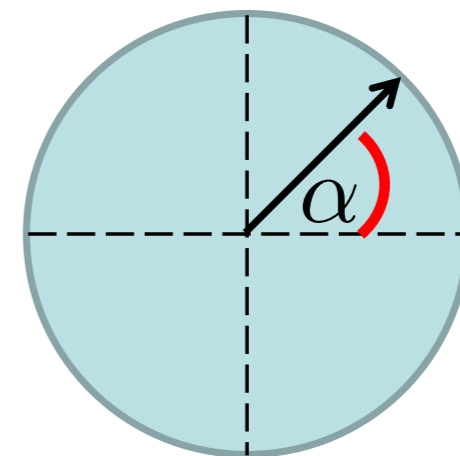
- ▶ If we call the rotation α
- ▶ Covariance matrix is diagonal (eigenvectors):

$$\sigma^2 \vec{u} = \lambda \vec{u} \quad \sigma^2 : \text{covariance matrix}$$

- ▶ If we assume a size 1 vector

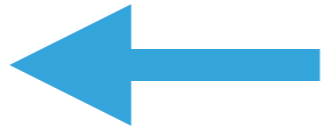
$$\hat{u} = \begin{pmatrix} a \\ b \end{pmatrix} = \begin{pmatrix} \cos(\alpha) \\ \sin(\alpha) \end{pmatrix}$$

$$\cos(\alpha)^2 + \sin(\alpha)^2 = 1$$



SHAPE DESCRIPTION: GEOMETRICAL VIEW

$$\sigma^2 \vec{u} = \lambda \vec{u}$$



$$\vec{u} = \begin{pmatrix} a \\ b \end{pmatrix} = \begin{pmatrix} \cos(\alpha) \\ \sin(\alpha) \end{pmatrix}$$



$$\begin{pmatrix} (\sigma_{xx}^2 - \lambda) \cos(\alpha) + \sigma_{xy}^2 \sin(\alpha) \\ (\sigma_{xy}^2 \cos(\alpha) + (\sigma_{yy}^2 - \lambda) \sin(\alpha) \end{pmatrix} = \begin{pmatrix} 0 \\ 0 \end{pmatrix}$$



$$\alpha = \frac{1}{2} \arctan \left(\frac{2\mu_{1,1}}{\mu_{2,0} - \mu_{0,2}} \right)$$

- ▶ With 1st eigenvalue we can measure the “length” (l) of the object in its intrinsic shape.

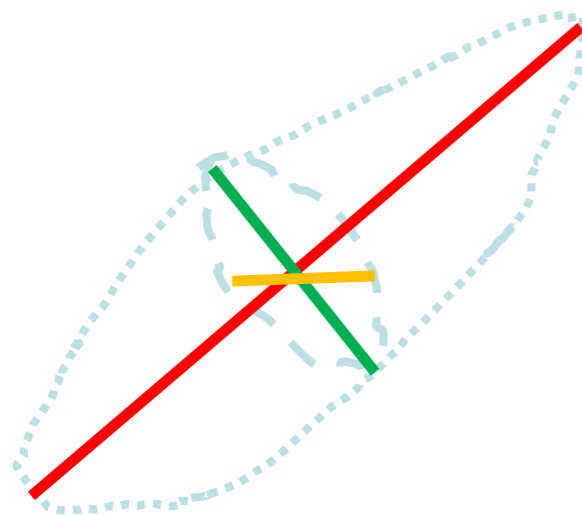
$$l^2 = \lambda = \frac{\text{Tr}(\sigma^2)}{2} + \sqrt{\frac{\text{Tr}^2}{4} - \det(\sigma^2)}$$

$$l^2 = \lambda = \frac{1}{2} \left(\sigma_{xx}^2 + \sigma_{yy}^2 + \sqrt{(\sigma_{xx}^2 - \sigma_{yy}^2)^2 + 4(\sigma_{xy}^2)^2} \right)$$

SHAPE DESCRIPTION: ECCENTRICITY

- ▶ We can now define eccentricity as:

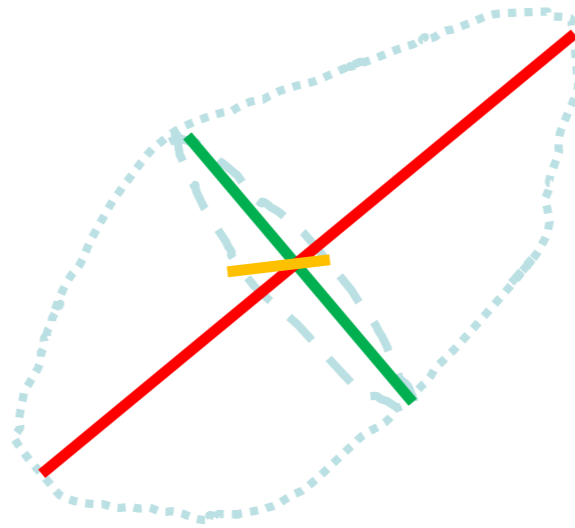
$$Elong = 1 - \frac{\lambda_2}{\lambda_1}$$



$$\lambda_1 \gg \lambda_2$$

Elong. ~ 1

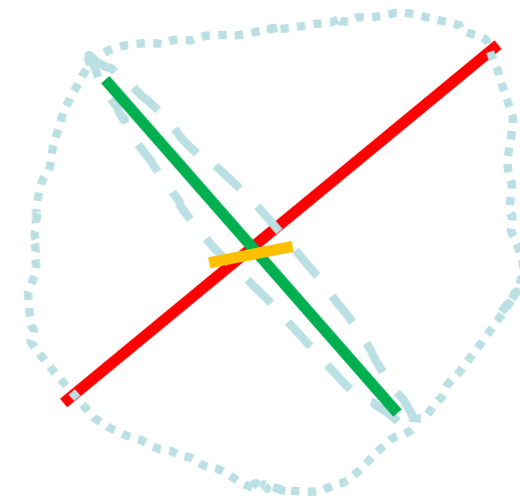
$$R.Elong = 1 - \frac{\lambda_3}{\lambda_1}$$



$$\lambda_1 \gg \lambda_3$$

R. Elong. ~ 1

$$Flatness = 1 - \frac{\lambda_3}{\lambda_2}$$



$$\lambda_2 \gg \lambda_3$$

Flatn. ~ 1

- ▶ We can compute it fast in binary images with image moments.

$$m_{p,q} = \sum x^p y^q I(x, y)$$

$$\mu_{p,q} = \sum (x - \bar{x})^p (y - \bar{y})^q I(x, y)$$

Order 0

$$m_{0,0} = \sum I(x, y)$$

$$area = m_{0,0}$$

Order 1

$$m_{1,0} = \sum xI(x, y)$$

$$m_{0,1} = \sum yI(x, y)$$

$$\vec{c}_m = \left(\frac{m_{1,0}}{m_{0,0}}, \frac{m_{0,1}}{m_{0,0}} \right)$$

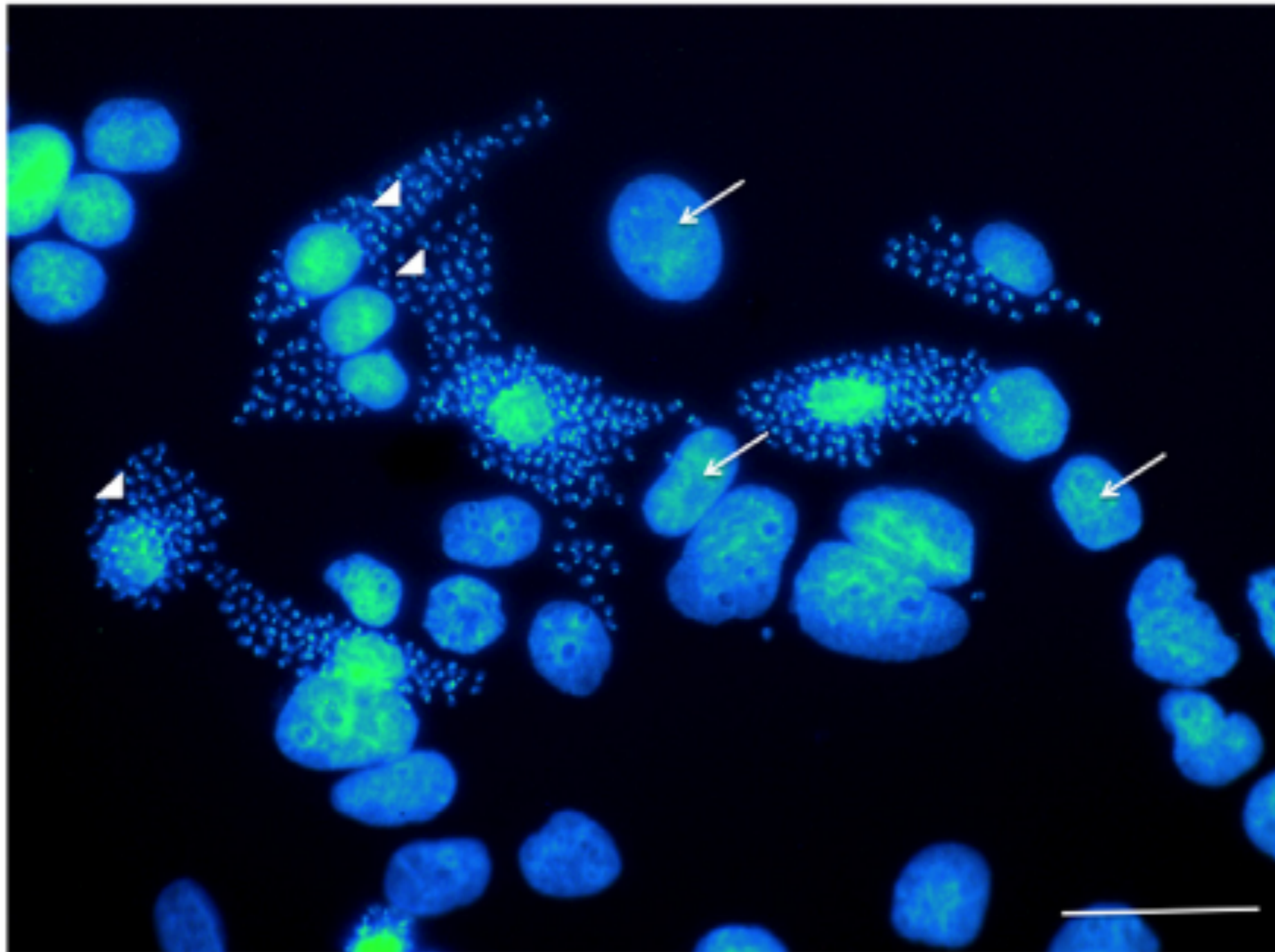
- ▶ With 2nd order moments covariance matrix is:

$$\sigma_x^2 = \frac{\sum (x - \bar{x})^2}{N} = \frac{\mu_{2,0}}{\mu_{0,0}} \quad \sigma_{xy} = \frac{\sum (x - \bar{x})(y - \bar{y})}{N} = \frac{\mu_{1,1}}{\mu_{0,0}}$$

$$\sigma^2 = \frac{1}{\mu_{0,0}} \begin{bmatrix} \mu_{2,0} & \mu_{1,1} \\ \mu_{1,1} & \mu_{0,2} \end{bmatrix}$$

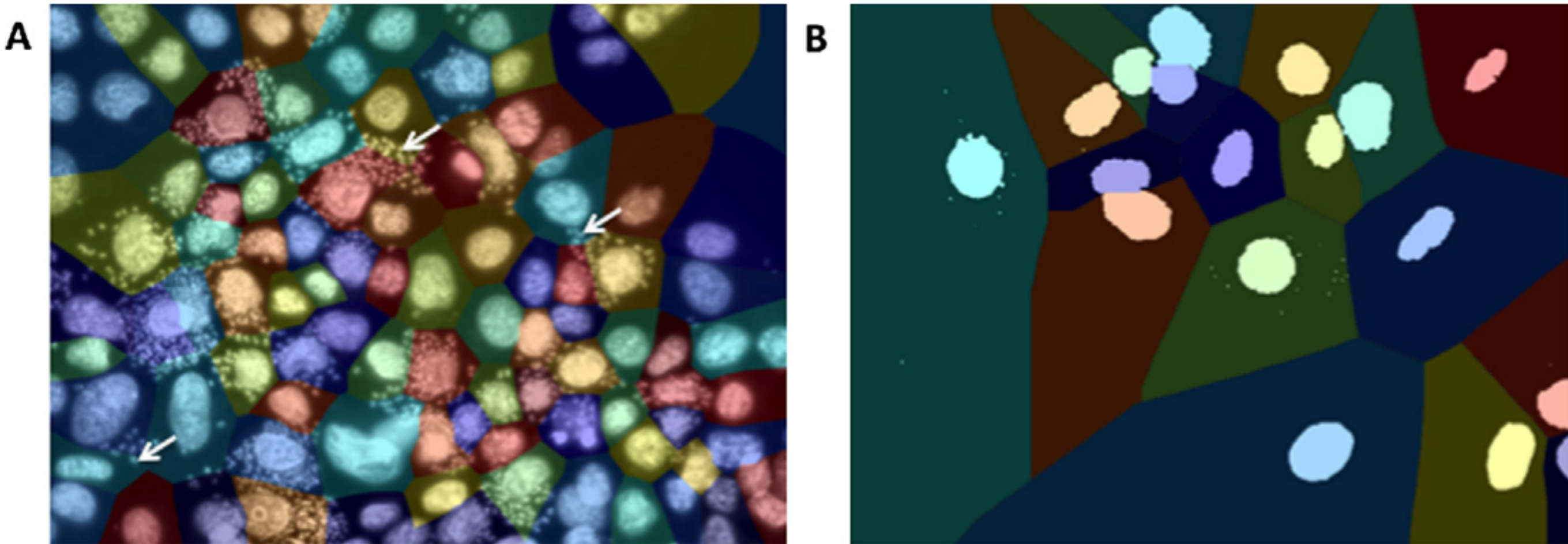
- ▶ To find the rotation is known as *Principal Component Analysis* (PCA).
- ▶ Multiple applications, eg. select eigenvectors to simplify distribution (objects) to a few numbers.
- ▶ PCA in FIJI [demo].

- ▶ Challenge 1: to estimate the number of parasites, BeWo cells, and their eccentricity.



LAB: CHALLENGE

- ▶ Count the number of infected cells by assigning (eg Voronoi) parasites to BeWo cells.



THANKS

F-Medicine

SCIAN-Lab

Steffen Härtel

Mauricio Cerda

Victor Castañeda

Jorge Jara

Marilyn Gatica

Paula Llanos

Alejandro Lavados

Juan Jose Alegría

Jimena López

Jorge Mansilla

Lucas Ale

Jorge Toledo

LEO-Lab

Miguel Concha

German Reig

Loreto López

Eduardo Pulgar

Lab Chagas

Ulrike Kämmerling

Anita Liempi

Hospital Clínico

U-Chile

F-Engineering

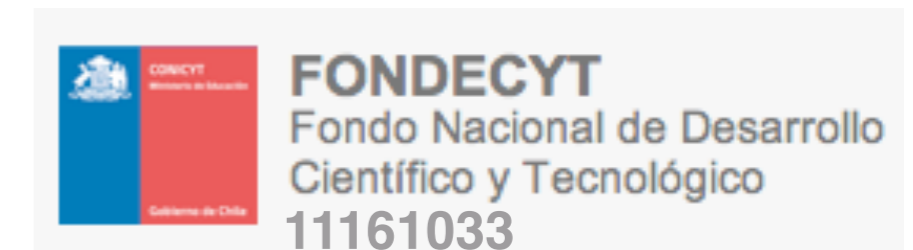
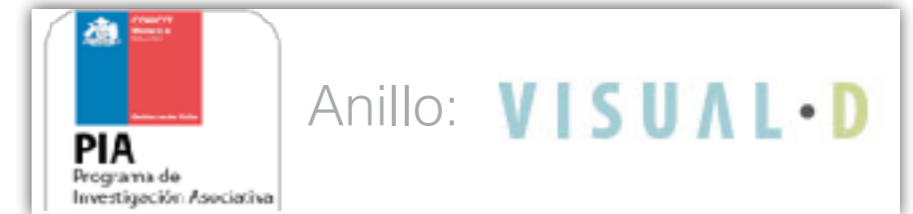
Computer Science

Nancy Hitschfeld

Physics

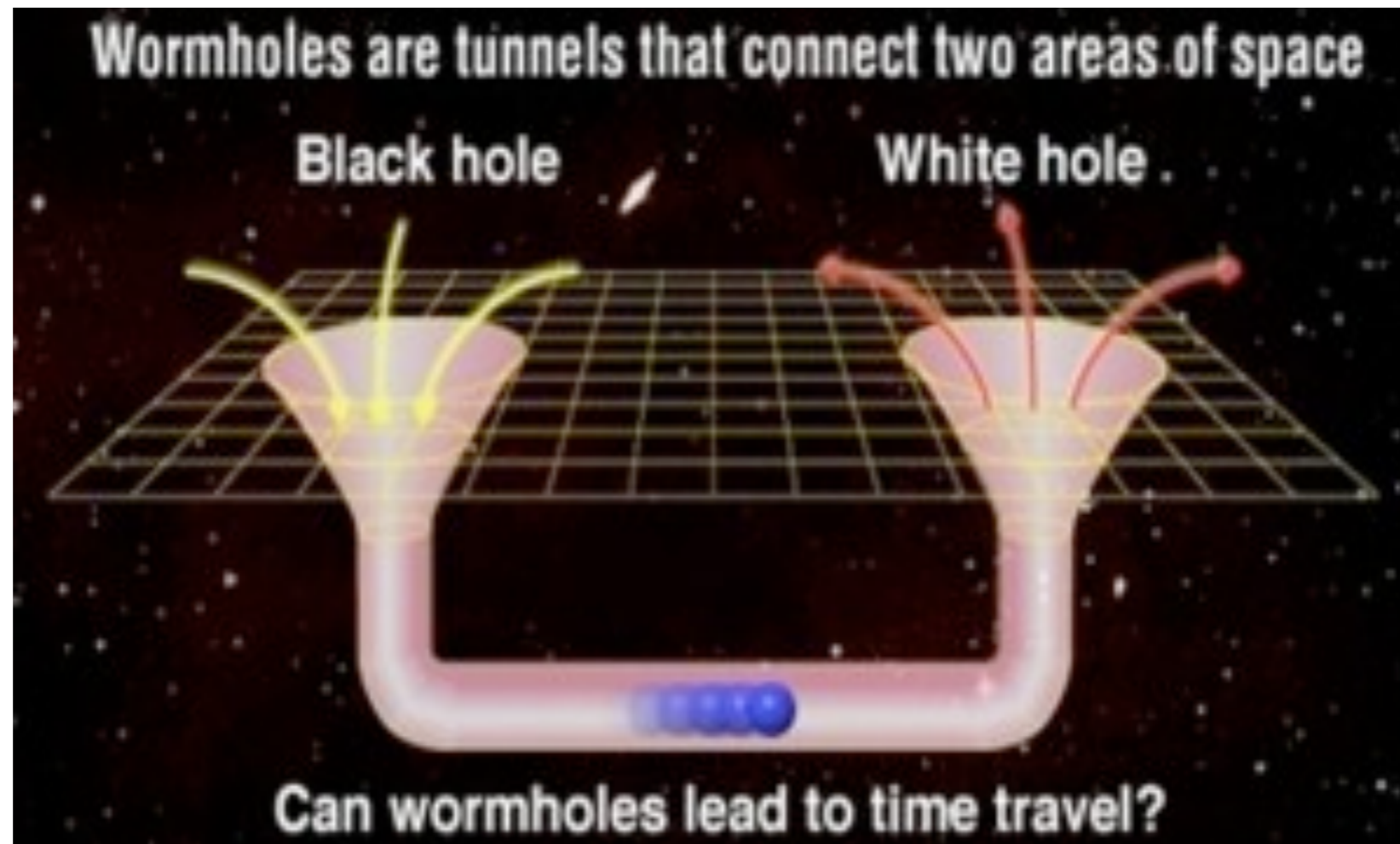
Rodrigo Soto

Nestor Sepúlveda



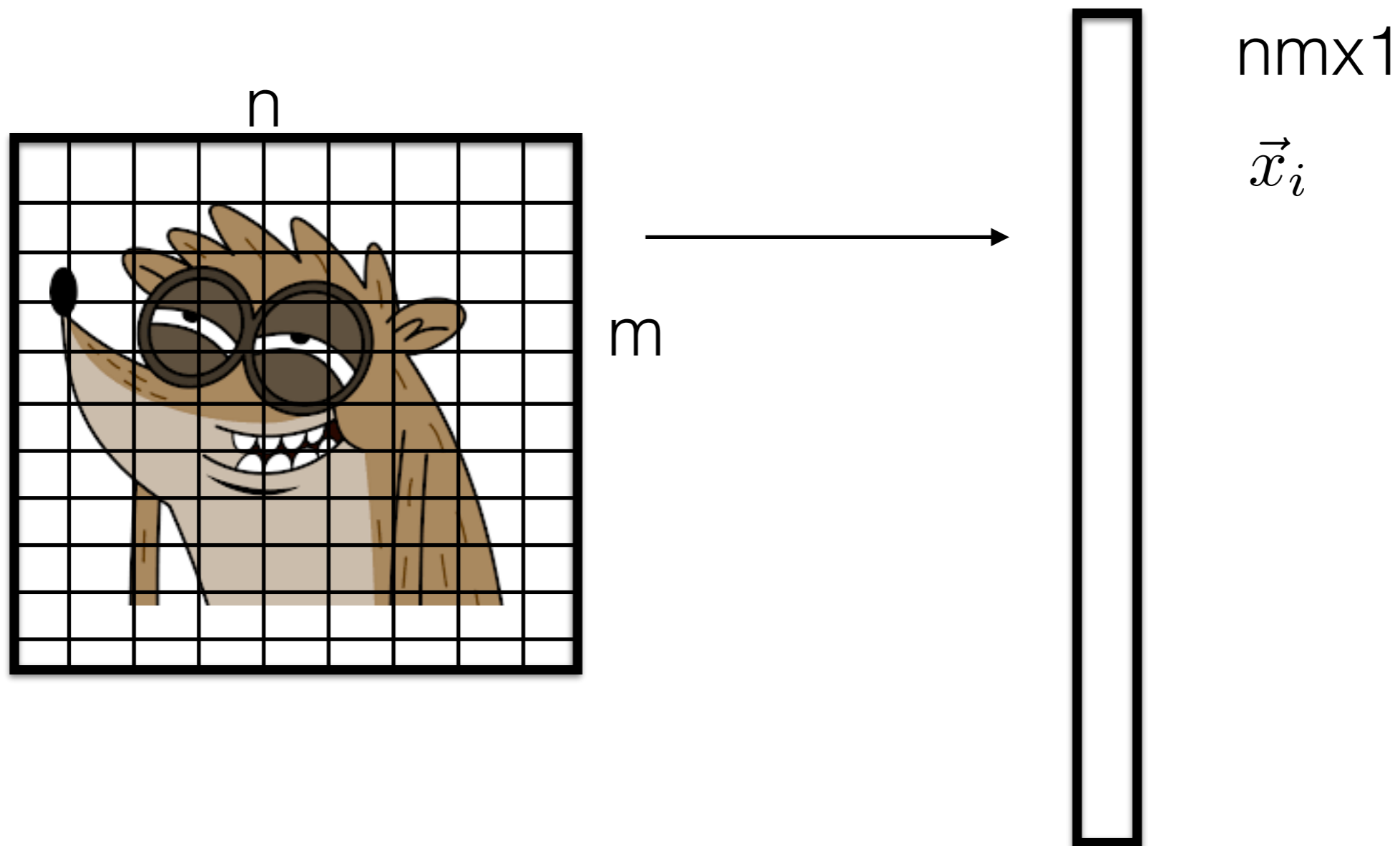
MAURICIO CERDA
MAURICIO CERDA@MED.UCHILE.CL
[HTTP://WWW.SCIAN.CL](http://www.scian.cl)
[HTTP://WWW.BNI.CL](http://www.bni.cl)

SHAPE DESCRIPTION: HIGH DIMENSION?



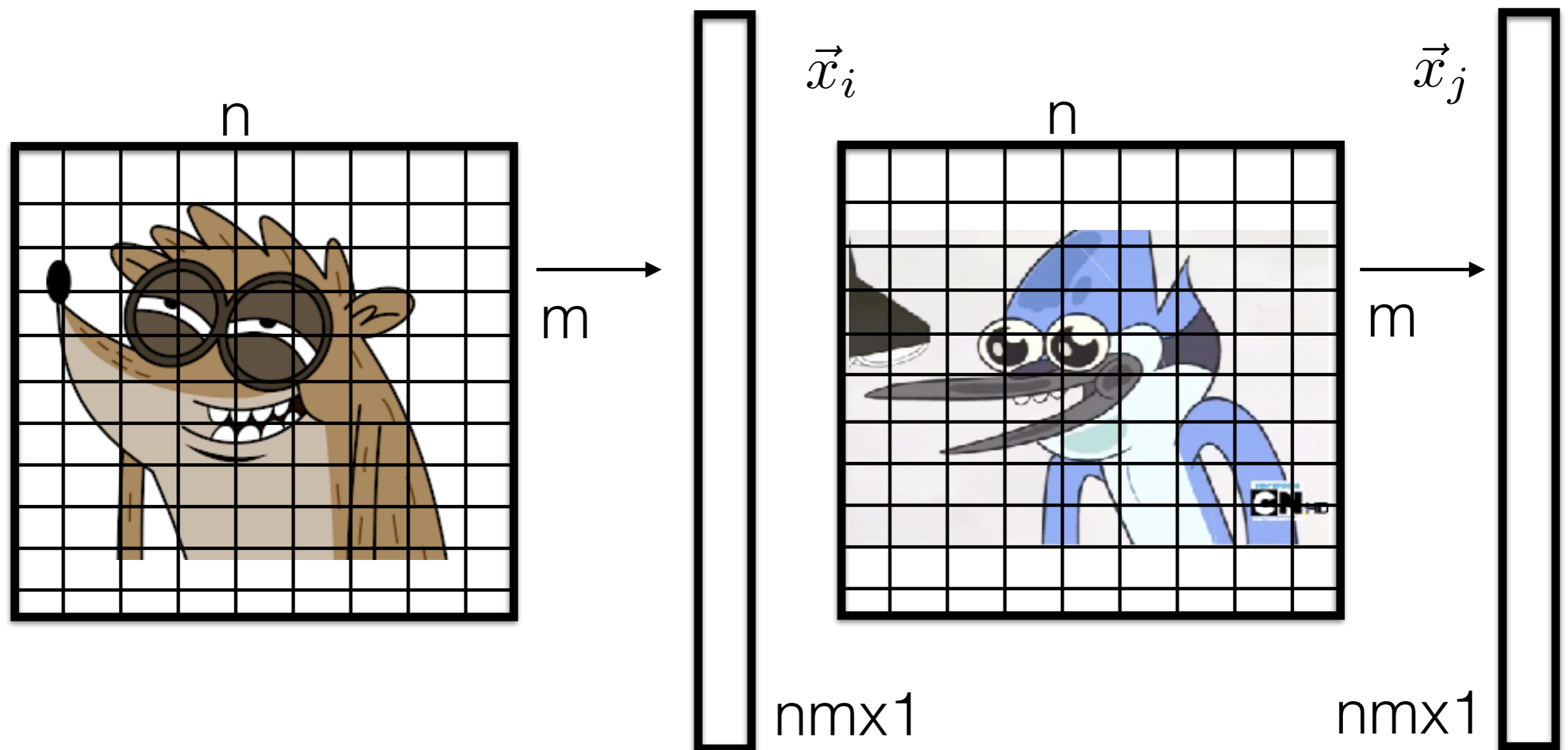
SHAPE DESCRIPTION: HIGH DIMENSION?

- ▶ How to understand an image in high dimension?



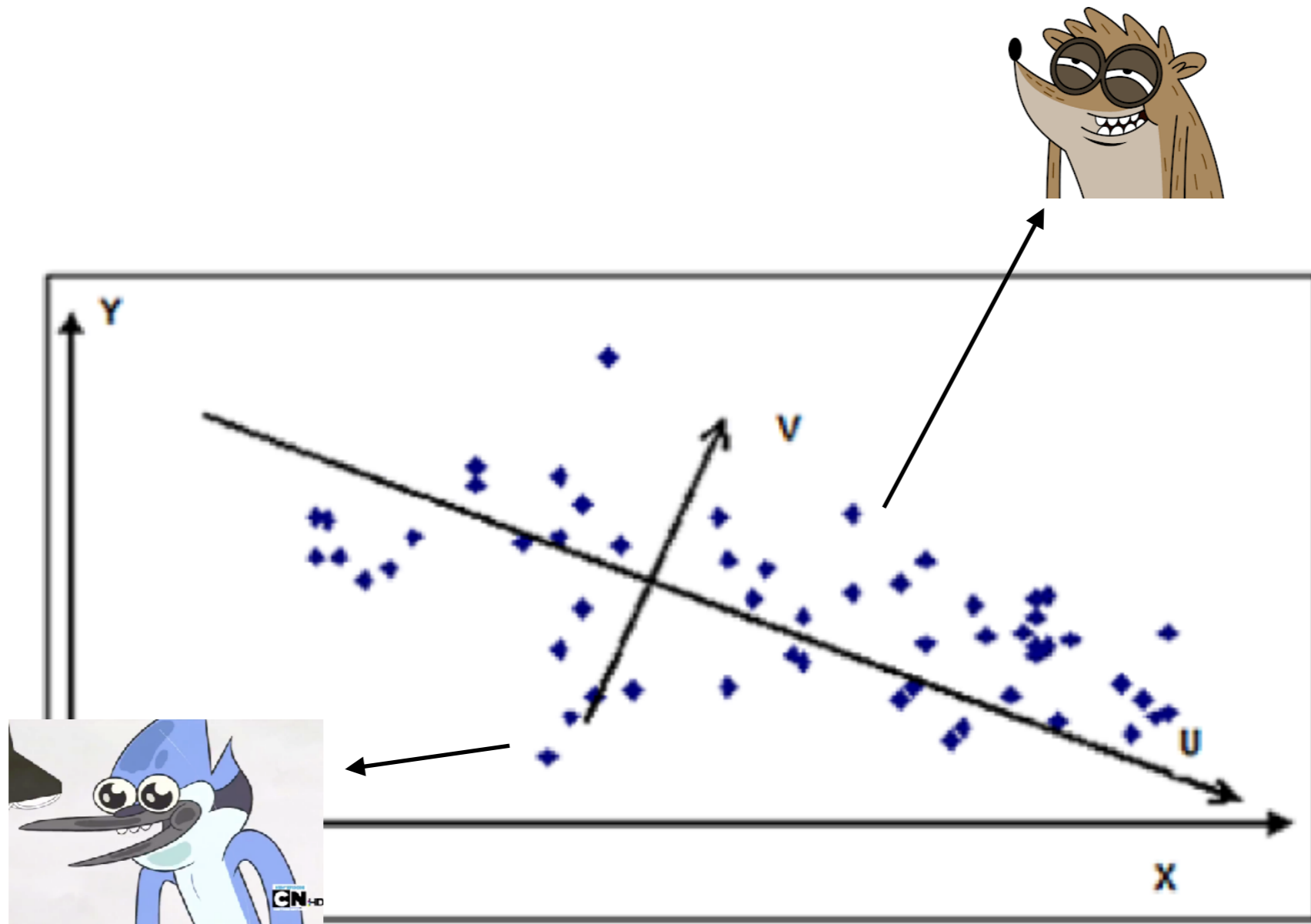
SHAPE DESCRIPTION: HIGH DIMENSION?

- ▶ For 2D images, we now have a nm size vector per image



SHAPE DESCRIPTION: HIGH DIMENSION?

- ▶ Now, each image is a point in your feature space.



SHAPE DESCRIPTION: COVARIANCE MATRIX

- ▶ Now, each image is a point in our feature space.

$$\mathbf{X} = [\vec{x}_1 \dots \vec{x}_i \dots \vec{x}_k] \quad (\text{column vector})$$

$$\mu_i = E(\vec{x}_{:,i}) \quad (\text{mean of row } i)$$

$$\Sigma_{ij} = \text{cov}(\vec{x}_{:,i}, \vec{x}_{:,j}) = E[(\vec{x}_{:,i} - \mu_i)(\vec{x}_{:,j} - \mu_j)]$$

$$\Sigma = \begin{bmatrix} E[(\vec{x}_{:,1} - \mu_1)(\vec{x}_{:,1} - \mu_1)] & E[(\vec{x}_{:,1} - \mu_1)(\vec{x}_{:,2} - \mu_2)] & \cdots & E[(\vec{x}_{:,1} - \mu_1)(\vec{x}_{:,k} - \mu_k)] \\ \vdots & \ddots & \vdots & \vdots \\ E[(\vec{x}_{:,k} - \mu_k)(\vec{x}_{:,1} - \mu_1)] & E[(\vec{x}_{:,k} - \mu_k)(\vec{x}_{:,2} - \mu_2)] & \cdots & E[(\vec{x}_{:,k} - \mu_k)(\vec{x}_{:,k} - \mu_k)] \end{bmatrix}$$

- ▶ If $\mu_i = 0$ (centered data)

$$\Sigma = \frac{1}{k} \mathbf{X}^T \mathbf{X}$$

- ▶ We can diagonalize the matrix (eg. using SVD)

$$\Sigma = \mathbf{U} \mathbf{D} \mathbf{V}^T$$

Eigenvectors

Eigenvalues (diagonal matrix)

- ▶ If eigenvalues are sorted from higher to lower.
- ▶ The first eigenvector will indicate the direction that maximizes variance.
- ▶ If the input vector are size nm , how many eigenvector are in the base?

SHAPE DESCRIPTION: PCA AS DESCRIPTOR

- ▶ Example. Face representation (*eigenfaces*) from a set of k photos form the same person.



1



2

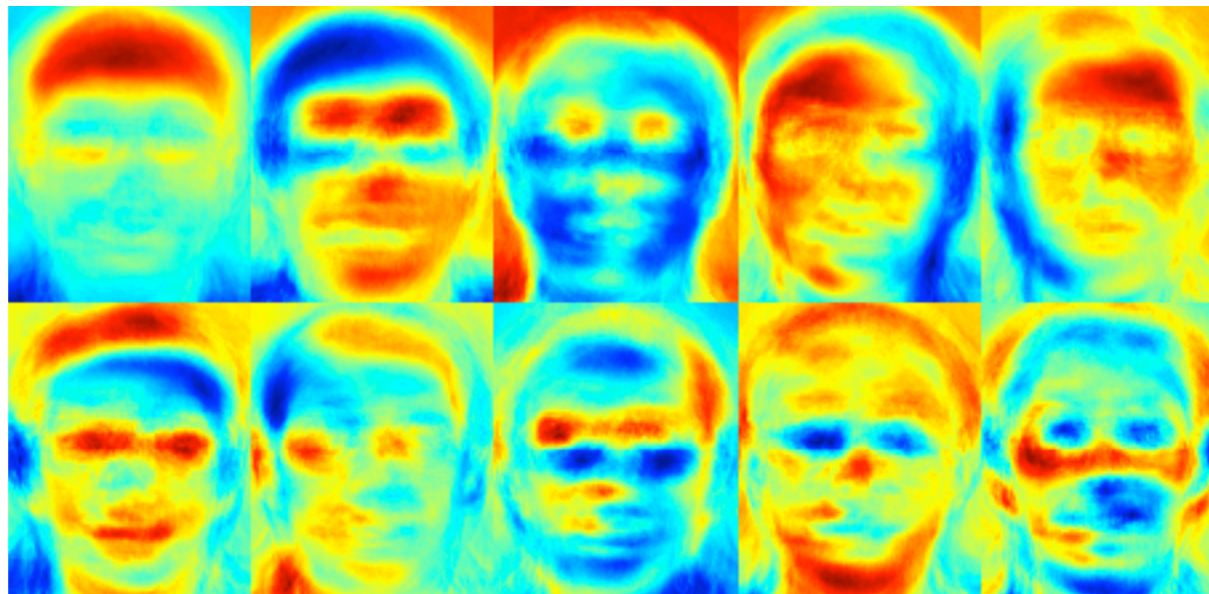
...



k

SHAPE DESCRIPTION: PCA AS DESCRIPTOR

- ▶ Example. Face representation (*eigenfaces*). The first 10 eigenvectors are:



- ▶ Example. Face representation (*eigenfaces*). We can use it as a way to reduce dimensionality:

