# Yeast Cells Classification

Machine Learning Approach to Discriminate Saccharomyces cerevisiae Yeast Cells Using Sophisticated Image Features.

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#### Table of Contents

- Introdution
- 2 Features
- 3 Classification
- 4 Results
- 6 Conclusion





Introdution

#### Section 1

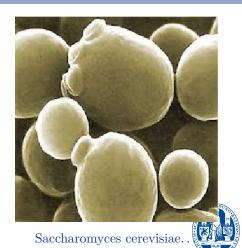
#### Introdution



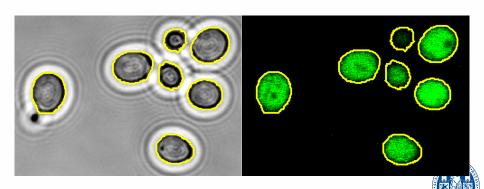


### Saccharomyces cerevisiae

- Originally isolated from grapes skin.
- Intensively studied eukaryotic model.
- Used to understand gene behaviour, under stress response.

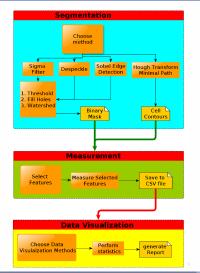


### BMH1-GFP under high stress 50mM NaCl media



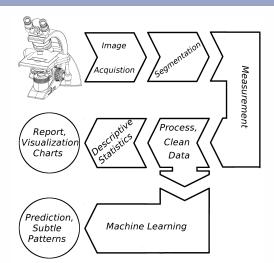
### Image Analysis Platform

- Has the following components:
  - Segmentation Module.
  - Measurement Module.
  - Statistics and Data Visualization Module.
  - GUI.





### Image Analysis Workflow

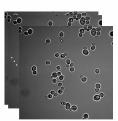


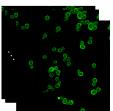




## Yeast Cells Image Modality

- Images Acquired by Zeiss LSM5 Exciter.
  - 2-Channels
    - Bright-Field
    - GFP- Protein





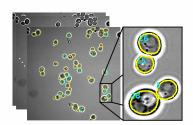


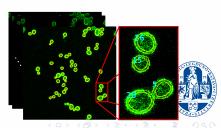


### Yeast Cells Segmentation

 Segmentation on Bright-Field Channels.

Resulted masks used to measure all channels.





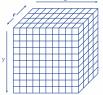
### Hough Transform To Detect Circles

- Detect Geometrical Circles.
  - Using 3D cube-like Accumulator.
  - Threshold to estimate cell locations.

$$T = 2\pi r - \{2\pi r \times \alpha + p\}$$
$$p = \beta \times (r_{max} - r_{min}) - r_{index}$$

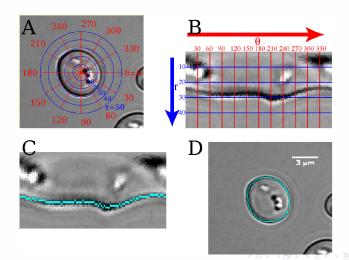






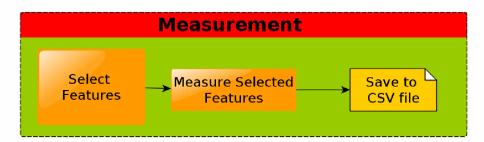


Introdution





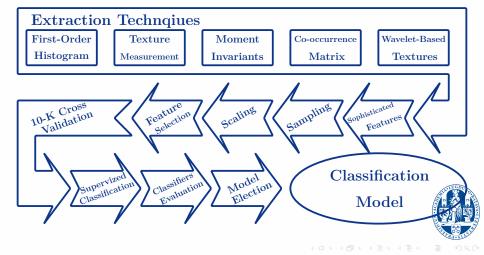
#### Measurement Module



- Subtle patterns not easy to be extracted.
- Sometimes it's not possible to see differences in cell groups.
- We need an automatic system to extract hidden features.



### Machine Learning



#### Section 2

#### **Features**





- A feature is a representation/attribute of an image.
- Texture:
  - is the visual effect produced by spatial distribution of variations.
  - is a rich Source of visual Information.
- Feature Extraction is locating pixels with distinctive characteristics.





### First-Order Histogram

• An Image as a function f(x, y).

$$h(i) = \sum_{x=0}^{N-1} \sum_{y=0}^{M-1} \delta(f(x, y), i), \qquad (1)$$

$$\delta(j,i) = \begin{cases} 1, j = i \\ 0, j \neq i \end{cases}$$
 (2)





Features	Description
Size	The number of pixels occupied by the cell.
Total Intensity	Sum of intensity values of pixels occupied by the cell.
Intensity Stan-	The standard deviation from the mean (intensity/pixel)
dard Deviation	of the intensity values at each pixel.
Perimeter	Cell perimeter.
Circularity	The circularity of detected shapes.
	$Circularity = \frac{4\pi Size}{perimeter^2}.$ (3)
Vacuole Size	Estimation of the vacuole size.
Membrane Fea-	Size, total Intensity, Intensity standard deviation.
tures	





### Textures based on First-Order Histogram

Textures	Description
Variance	Measure of intensity contrast.
	$\mu_2(z) = \sum_{i=0}^{L-1} (z_i - m)^2 . P(z_i)$
Relative	Zero for constant intensities.
Smoothness	$R(z) = 1 - \frac{1}{1 + \sigma^2(z)}$
Skewness	Indication of the skewness of the histogram. $\mu_3(z) = \sum_{i=0}^{L-1} (z_i - m)^3 . P(z_i)$
Uniformity	Has a maximum value when intensity levels are equal. $U(z) = \sum_{i=0}^{L-1} P^2(z_i)$
Entropy	A measure of variability, is zero for constant images. $e(z) = -\sum_{i=0}^{L-1} P(z_i) . log_2 P(z_i)$





Features

#### **Moment Invariants**

- An image moment:
  - Is a certain particular weighted average, i.e. moment of pixel intensities.
  - Computed based on the information from shape and interior region.
  - Useful descriptor after segmentation.
- Simple Properties from low order moments.
- Invariant to translation, scale and rotation.
- frequently used as features for:
  - Image Processing.
  - Remote sensing.
  - Shape recognition.
  - Classification.





#### Hu's Set of Moment Invariants

$$hu = \{\Phi_1, \Phi_2, \Phi_3, \Phi_4, \Phi_5, \Phi_6, \Phi_7\}.$$

- Are widely Known set of seven invariants.
- $\Phi_1$  &  $\Phi_2$  are based on second order moments.
- $\bullet$   $\Phi_3$  ...  $\Phi_7$  are based on third order moments.
- More effective when fused with other techniques.





#### Co-occurrence Matrix

- Simple texture attributes can not characterize cells.
- Similar texutres agree in their second-order statistics.
- Second Order statistics:
  - Are given by pairs of pixels.
  - Have good discrimination rates.
  - Important in automated image analysis.
  - Features derived from co-occurrence matrix.
- Co-occurrence Matrix:
  - For Image f(x,y) with L discrete levels (Dimension L x L).
  - The  $(i, j)^{th}$  is # of times that f(x1,y1) = i and f(x2,y2) = j.
  - where  $(x2,y2) = (x1,y1) + (d \cdot \cos \theta, d \cdot \sin \theta)$ .



#### Co-occurrence Matrix Derived Features

- Features used for texture discrimination.
  - Angular Second Moment.
  - Correlation.
  - Intertia.
  - Absolute Value.
  - Inverse Difference.
  - Entropy.
  - Maximum Probability.





#### Multi-Scale Features

- Methods to calculate multi-scale features:
  - Wigner Ditributions.
    - Has interference terms between components.
  - Gabor Transform.
    - Non-orthogonal -> Redundant features.
  - Wavelet Transforms.





#### Section 3

#### Classification





- Dataset of 1440 yeast cell instances.
- 14-3-3 proteins with GFP in 50mM vs. 0mM NaCl
- Measure all features per individual cell instance
- Construct a contigency table to represent dispositions of the set of instances.
- Evaluate 23 different linear and non-linear classifiers.
  - ... including: decision trees, naive Bayes, least-square linear preictors, SVM, etc...





### Imbalanced Dataset & Sampling Techniques

- Unequal distribution between classes.
- Sampling Techniques improves classifier accuracy.
  - UnderSampling.
  - OverSampling.
  - SMOTE.





### Data Scaling, i.e. Normalization

- Applied at data pre-processing.
- Some Algorithms will not work without Normalization.
- Normalization Techniques:

• UL. 
$$x_i^* = \frac{x_i}{||x||}, i = 1, 2, ...d,$$

• MV. 
$$x_i^* = \frac{(x_i - \mu)}{\sigma}, i = 1, 2, .., d,$$





### Feature (Attribute) Selection

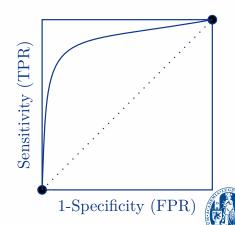
- To optimally reduce feature space.
- Advantages:
  - Improves the prediction performance.
  - Provides faster and more cost effective classifiers.
  - Provides a better understanding of the underlying process that generated the data.
  - Reduces overfitting.
  - Reduces training time.
- Avoid selecting redundant and irrelevant features.
- Selected Algorithms:
  - Information Gain (IG).
  - Correlation Feature Selection (CFS).
  - Principal Component Analysis (PCA).





### Evaluation metrics, ROC and AUC

- ROC curve is a 2D graphical plot.
  - $\bullet$  **AUC** = **1**, Perfect
  - 1 > AUC > 0.9, Excellent.
  - 0.9 > AUC > 0.8, Good.
  - 0.8 > AUC > 0.7, Fair.
  - 0.7 > AUC > 0.6, Poor.



#### Classifiers Evaluated

- 23 different classifiers.
- $\bullet$  Using Weka, R and rWeka





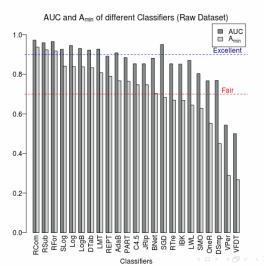


Results



Results •000000

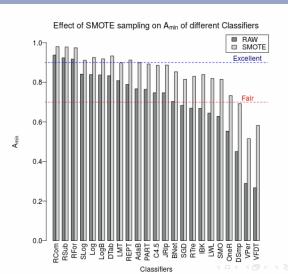
### AUC vs. Amin







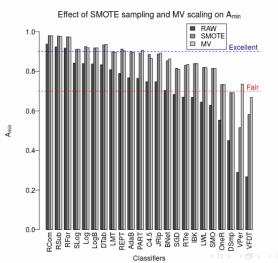
### Power of Sampling







#### Normalization and Feature Selection

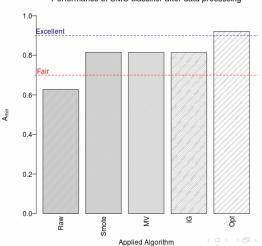






### $\overline{\text{SVM} : \text{SMO}}$

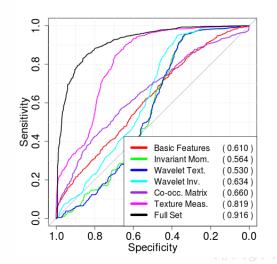
#### Performance of SMO classifier after data processing







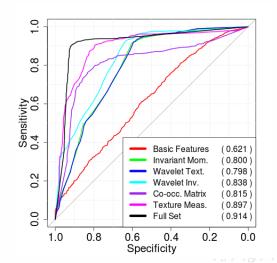
### Analysis and AUC value of Logistic Classifier







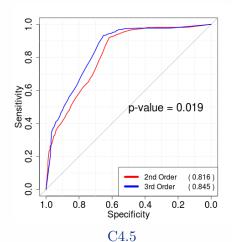
### Analysis and AUC value of C4.5 Classifier

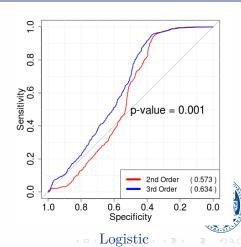






### Performance of Classifiers using second and up-to third order invariant moment features





#### Section 5

#### Conclusion





- A machine learning approach can discriminate yeast cells cultivated under different stress levels.
- A feature set is powerful in predicting cell groups, combined features from 1st-order histogram, moment invariants, Co-occurrence matrix and Wavelet-based texture features.
- Using SMOTE for data sampling, MV for data normalization and IG for feature selection.
- As future work:
  - Classify different cell strains and conditions in a high-volume HTS studies.
  - $\bullet~$  Use developmental techniques to create optimal classifier.



### Acknowledgement

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- Supervisor : Dr. Ir. Fons J. Verbeek (section Imaging & BioInformatics, LIACS)
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