

High-speed Recognition of Micro-array Genomic Images Using Multi-scale Representations



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INTRODUCTION

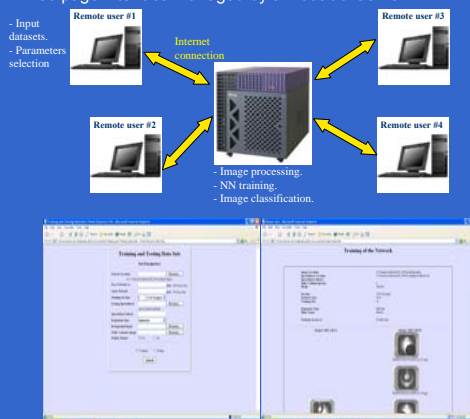
- This project is part of the Northeast Structural Genomics Consortium (NESG). The goal of this consortium is to develop efficient and integrated technologies for high-throughput (HTP) protein production and 3D structure determination.
- This project focuses on the design of an image analysis system to classify protein crystal structures in a production oriented environment.
- The method performs classification of microscopic images as clear droplets versus non-clear droplets (precipitates and crystals).
- Using expert classification for ground truth, current results show high classification accuracy with a large image datasets.

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METHODOLOGY

5. Remote Application Server

Web page interface managed by a Matlab® Server.



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RESULTS

1. Classification with the Image Database

► Image Features for Neural Network

Quantitative shape descriptions of a first-order Laplacian pyramid coefficients histogram combined with the power spectrum and autocorrelation information.

8 statistics for each Laplacian coefficient subset with totally 5 subsets. ($8 \times 5 = 40$)

► Binary classification

0 = clear drop, 1 = not a clear drop

► Training data set

100, 500, 1000, 1500, 2000, 2500, 3000, 3500 images
1/2 clear drops, 1/2 drops with precipitates/crystals

► Testing experiments

200 images with precipitates/crystals: TP = 90%.
200 images with clear drops: TN = 92%.

► Definition of the classification system

Accuracy = $(TP+TN)/2 = 91\%$

TP = True Positive (Percentage) = Correctly classified images with precipitates or crystals / Total images with precipitates or crystals * 100%

TN = True Negative (Percentage) = Correctly classified images with clear drops / Total images with clear drops * 100%

2. Examples of Misclassified Images



Skin effects, complex boundaries, light reflection, etc.

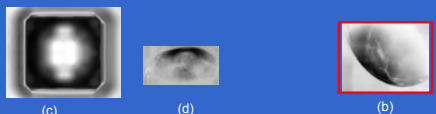
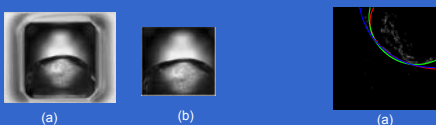


Small bubble size, no sharp gradient, background influence, etc.

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METHODOLOGY

1. Preprocessing of Microscopic Images



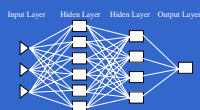
(a) Original image with an oil droplet containing precipitates. (b) Cropped image with Radon transform. (c) Background image cropped without drops. (d) Pre-processed image with Ellipsoidal Hough transform.

(a) Ellipsoidal Hough transform [1] to detect the three most probable ellipses, plotted over the edge map of the pre-filtered image. (b) Pre-processed image cropped with the minimal rectangular area encompassing the three ellipses.

2. Feature Extraction with Laplacian Pyramidal Expansion [2]



3. Classification with a Feed-Forward Neural Network



4. We used the quantitative shape descriptions of a first-order histogram combined with the power spectrum and autocorrelation information:

$$\begin{aligned} \text{Mean:} & S_0 = \sum_{i=1}^n b_i P(b_i) \\ \text{Standard Deviation:} & S_1 = \left[\sum_{i=1}^n (b_i - S_0)^2 P(b_i) \right]^{1/2} \\ \text{Skewness:} & S_2 = \frac{1}{S_1^3} \sum_{i=1}^n (b_i - S_0)^3 P(b_i) \\ \text{Kurtosis:} & S_3 = \frac{1}{S_1^4} \sum_{i=1}^n (b_i - S_0)^4 P(b_i) - 3 \\ \text{Energy:} & S_4 = \sum_{i=1}^n [P(b_i)]^2 \\ \text{Entropy:} & S_5 = - \sum_{i=1}^n P(b_i) \log_2 [P(b_i)] \\ \text{Power:} & S_6 = \sum_{i=1}^n [f(b_i)]^2 \\ \text{Autocorrelation:} & S_7 = \sum_{i=1}^n \sum_{j=1}^n \text{Im}(i) \cdot \text{Im}(j) \end{aligned}$$

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DATA

- Microscopic images were acquired with a CGD camera under robotic control.
- Gray scale 8-bits images saved in tiff format.
- Image Database of 5,000 manually classified images:
 - 2500 drops containing precipitates and/or crystals.
 - 2500 clear drops (clear, skin, etc)



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DISCUSSION

- Introduction of robotic manipulation of the crystals for HTP protein production requires the automation of image analysis of crystallization experiments for classification of solution content.
- The proposed feed forward neural network showed promising results in classifying microscopic images.
- Most features of representation were computed from Laplacian pyramid expansion histograms. The histogram made the features invariant to orientation which was a desirable feature in order to be able to characterize the diversity and complexity of precipitate appearances.
- The Laplacian expansion provided a representation of the image edge and texture patterns at different scales with extremely fast implementation.

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CONCLUSION

- We are working on further classification of the microscopic image database to separate crystals from precipitates.
- A parallel task of this project includes the creation of a web-based infrastructure for testing and development. An experimental test-bed for crystallization screening is currently under development.

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REFERENCES

- Angelini, E. D., Y. Wang, et al. (2004). Classification of Micro Array Genomic Images with Laplacian Pyramidal Filters and Neural Networks. Workshop on Genomic Signal Processing and Statistics (GENSIPS), Baltimore, Maryland, USA.
- D. H. Ballard, "Generalizing the Hough transform to detect arbitrary shapes," *Pattern Recognition*, vol. 13, pp. 111-122, 1981.
- P. J. Burt and E. H. Adelson, "The Laplacian pyramid as a compact image code," *IEEE Transactions on Communications*, vol. 31, pp. 532-540, 1983.
- C. A. Cumbaa, A. Lauricella, N. Fehman, C. Veatch, R. Collins, J. Luft, G. DeTitta, and I. Jurisica, "Automatic classification of sub-microlitre protein-crystallization trials in 1536-well plates," *Acta Crystallographica Section D-Biological Crystallography*, vol. 59, pp. 1619-1627, 2003.
- I. Jurisica, P. Rogers, J. I. Glasgow, R. J. Collins, J. R. Woffley, J. R. Luft, and G. T. DeTitta, "Improving objectivity and scalability in protein crystallization," *IEEE Intelligent Systems*, vol. 16, pp. 26-34, 2001.

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