Alpha Divergence for Feature Pruning and Indexing of Large Biological Databases

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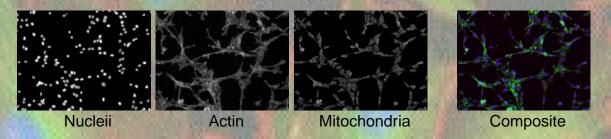
URSI 2002 National Radio Science Meeting, January 9-12, 2002, University of Colorado, Boulder, CO.

Outline

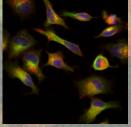
- 1. Indexing Application: Large-Scale Cellular Imagery Database
- 2. Statistical Framework: Alpha-Divergence for Indexing
- 3. Application Cellular Imagery
- 4. Summary, Open Issues

Cellular Imaging with Fluorescence Microscopy

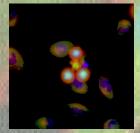
Can label and image several distinct cell components



Can measure biological effect of drugs by tracking changes vs.
 dose



Before



After

Large-Scale Cellular Imagery Database

- Combinatorial Biology: large biological effects database
 - \sim 30 cell lines (heart, liver, cervix, prostate, lung, etc.)
 - ~ 30 labels/markers (nucleus, membranes, cytoskeleton)
 - ~ 10000 compounds (pharmacopia)
 - ~ 100 images (4 replicates \times 5 doses \times 5 timepoints)
 - \Rightarrow 10¹⁰ images (300M multi-color image sets)
- Objectives:
 - Catalog biological effects of known compounds
 - Infer properties of novel compounds by comparison to known drugs

Methodology

- Combinatorial Biology ⇒ Imaging Experiments
- ◆ High-Throughput Cellular Imaging ⇒ Image Responses

$$\{(X^k, Y^k), k = 1, \dots, K\}$$

 X^k are images, $Y^k \in \mathcal{Y}$ are labels assigned by experiment protocol

Image Responses ⇒ Response Signatures (feature vectors)

$$Z^k = Z(X^k)$$

Response Signatures ⇒ Biological Effects

$$Z^k = Z(X^k) \sim f(z|Y^k)$$

f a feature density (likelihood) conditioned on label Y^k

 \Rightarrow Biological Similarity? $X^1 \sim X^2$ if $f(z^1|Y^1) \sim f(z^2|Y^2)$

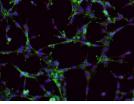
Comparison of Biological Effects

Define a Cell Imaging Experiment

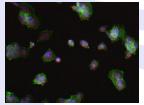
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\mathcal{E}_k = \{ Y^k \sim \text{treatment k} : \\ n \text{ cell lines} \times m \text{ labels} \times p \text{ time points} \times q \text{ replicates} \}
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Can we infer biological effects by comparing experiments?

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Y^k \in \mathcal{Y} \stackrel{\mathrm{def}}{=} \{ \text{ Apoptosic (cell death), Normal (blank)} \}
\mathcal{E}_{\mathsf{Blank}} = \{ \text{ "Blank" : 400 imgs of untreated 3T3 cells } \}
\mathcal{E}_{\mathsf{Apop}} = \{ \text{ "Pos. Control" : 90 imgs of 3T3 w/ Paclitaxel} \}
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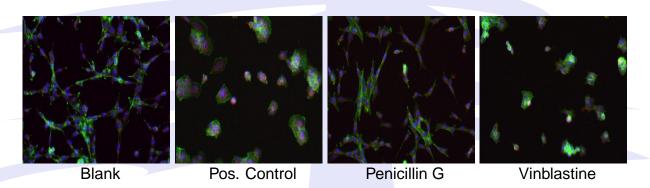
Blank



Pos. Control

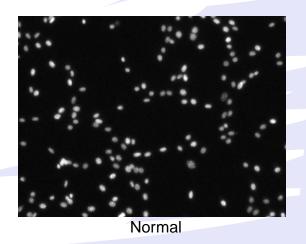
Apoptosis Experiment

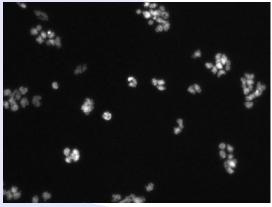
- Collected 1000 images of 10 different compounds, $\mathcal{E}_k, k = 1, \dots, 10$
 - 3 known apoptosis inducers, 7 known non-apoptotic agents
 - Imaged 3T3 cells treated with various compound doses
- Objective: Classify compounds according to effect (apoptosis inducer?) (classify: $\mathcal{E}_k \sim \mathcal{E}_{\mathsf{Blank}}$ or $\mathcal{E}_k \sim \mathcal{E}_{\mathsf{Apop}}$)



Apoptosis Effects: Nucleus

• Nuclear Fragmentation ("blebbing") and Condensation

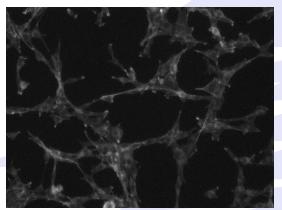




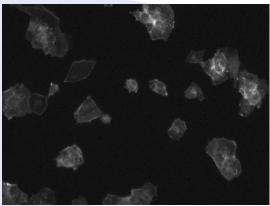
Apoptosis

Apoptosis Effects: Actin Filaments

Actin Reorganization



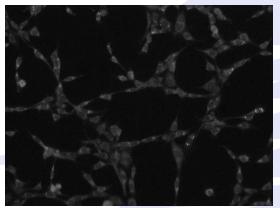
Normal



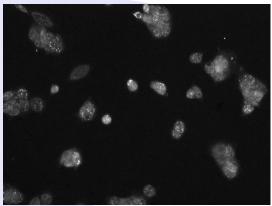
Apoptosis

Apoptosis Effects: Mitochondria

• Mitochondrial Disintegration



Normal



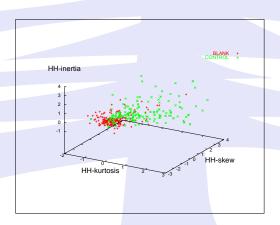
Apoptosis

Feature Extraction

Top 12 Features:

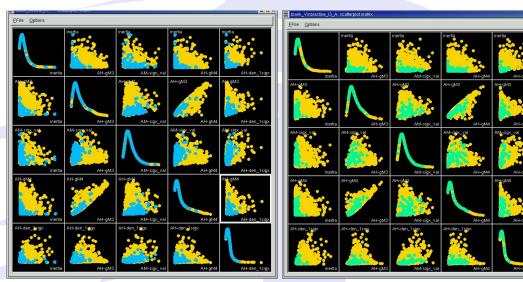
	Nucleus	Actin	Mitochondria
	H-Mass(Size in pix)	A-Mass(Size in pix)	HM-Coef.Var.
	H-Coef.Var.	A-Rot.Inert.	M-2nd X-Mom. Inert.
٠.	H-Kurt.	AH-3rd Jt. Cumulant	M-Rot.Inert.
	H-Skew	AH-4th Jt. Cumulant	
		AH-2nd Y-Mom. Inert.	

• Graylevel features from Nuclear channel:



Feature Extraction

- Lattice plot of five selected features
 - Blank vs. Control and Blank vs. Vinblastine scatterplots visually similar



Blank vs. Control

Blank vs. Vinblastine

Candidate Feature Similarity Metrics

- Univariate: Kolmogorov-Smirnov Distance:
 - Product density approximation: combine several 1D feature scores
- Multivariate: Rényi α-Divergence
 - Computation via second-order approximation of densities
 - Direct computation via minimal graphs

Univariate Distance Measures

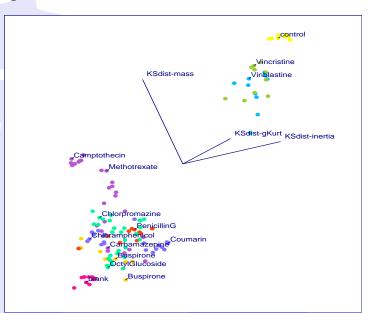
Score computed from 1D Kolmogoroff-Smirnoff distance

$$D_b(z) = D_{KS}(f_1(z)||f_{\mathsf{blank}}(z))$$

$$D_c(z) = D_{KS}(f_1(z)||f_{control}(z))$$

$$D_{bc}(z) = D_{KS}(f_{\mathsf{blank}}(z) || f_{\mathsf{control}}(z))$$

$$Score(z) = (D_b(z) - D_c(z))/D_{bc}(z)$$



Renyi α**-Divergence**

Define: $f_i = f(z|Y^i), f_0 = f(z|Y^0)$

The Rényi α -divergence of fractional order $\alpha \in [0,1]$ [Renyi:61, 70]

$$D_{\alpha}(f_i||f_0) = \frac{1}{\alpha - 1} \ln \int f_0 \left(\frac{f_i(z)}{f_0(z)}\right)^{\alpha} dz$$
$$= \frac{1}{\alpha - 1} \ln \int f_i^{\alpha}(z) f_0^{1 - \alpha}(z) dz$$

Note: $D_{\alpha}(f_i||f_0) = D_{\alpha}(Y^i||Y^0)$ is indexed by Y^i and Y^0 .

Renyi α-Divergence: Special Cases

α-Divergence vs. Batthacharrya-Hellinger distance

$$D_{\frac{1}{2}}(f_i \parallel f_0) = \ln \left(\int \sqrt{f_i(z) f_0(z)} dz \right)^2$$

$$D_{BH}(f_i \parallel f_0) = \int \left(\sqrt{f_i(z)} - \sqrt{f_0(z)} \right)^2 dz$$

$$= 2 \left(1 - \int \sqrt{f_i f_0} dz \right)$$

α-Divergence vs. Kullback-Leibler divergence

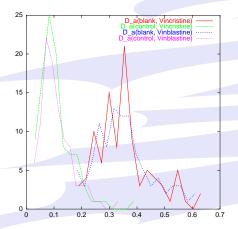
$$\lim_{\alpha \to 1} D_{\alpha}(f_i, f_0) = \int f_0(z) \ln \frac{f_0(z)}{f_i(z)} dz \tag{1}$$

Renyi α-**Divergence via Approximation**

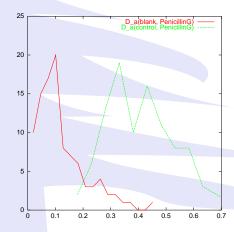
• Compute D_{α} using second-order approximation:

$$D^u_{\alpha}(f(x;\mu_1,\Lambda_1)\|f(x;\mu_0,\Lambda_0)) = \underbrace{-\frac{1}{2}\ln\frac{|\Lambda_0|^{\alpha}|\Lambda_1|^{1-\alpha}}{|\alpha\Lambda_0+(1-\alpha)\Lambda_1|}}_{\text{Term A}} + \underbrace{\frac{\alpha(1-\alpha)}{2}\Delta\mu^T(\alpha\Lambda_0+(1-\alpha)\Lambda_1)^{-1}\Delta\mu}_{\text{Term B}}$$

where $\Delta \mu = \mu_1 - \mu_0$.



 $D_{\alpha}(Vin*||Blank)$ vs. $D_{\alpha}(Vin*||Control)$



 $D_{\alpha}(\text{Penicillin G}||\text{Blank}) \text{ vs. } D_{\alpha}(\text{Penicillin G}||\text{Control})$

Experiment Results

Compounds used in Experiment

Training Data	Test Data	
Negative	Non-Apoptotic	
Blank, Penicillin G	Chloramphenicol, Buspirone, Sodium Azide,	
	Carbamezapine, Octyl Glucoside,	
	Carbon Tetrachloride	
Positive	Apoptotic	
Paxlitaxel (pos. control)	Methotrexate, Vincristine, Vinblastine	

- Correctly classified Vincristine and Vinblastine as apoptosis inducers
- Methotrexate did not penetrate the 3T3 cells with the experiment protocol we used → did not induce apoptosis.

Summary and Ongoing Issues

- Successful Comparison of Biological Experiments
 - Similarity of cell image changes
 similarity of biological effect
- Rényi Divergence via 2nd Order Approximation
 - Reasonable fit for nuclear features (unimodal, symmetric)
 - Will be an issue for other feature densities (skew, multimodal)
- Open Questions:
 - What is the appropriate feature set for cellular imagery analysis?
 - Does image similarity imply biological similarity?