

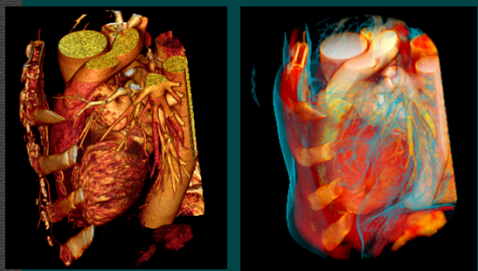
影像處理與統計應用

生物影像統計學

(BIS: Bio-Image Statistics)

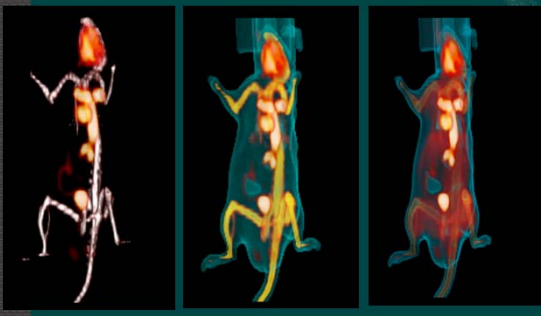
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2017/04/22

3D Cardiac Volume Rendering



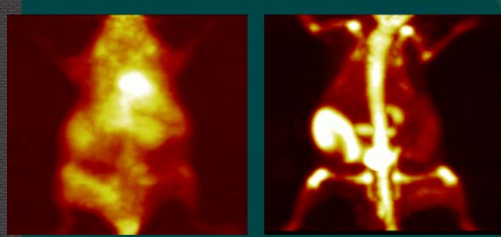
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3D Mice Volume Rendering



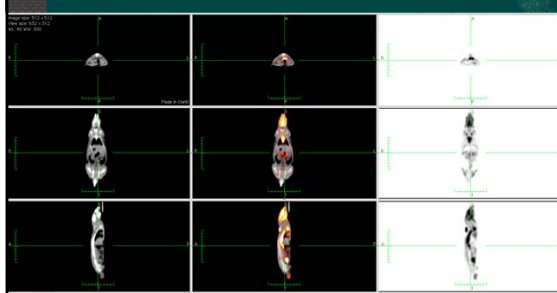
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3D Maximum Intensity Projection



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Mice PET+CT and Image Fusion



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3D Registered Image



3D registered and overlaid neuronal patterns of multiple fruit fly central complexes (top) and thoracic ganglia (bottom). Hanchuan Peng, 2008.

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Bioinformatics: MicroArray Image

Enlarge one block and spike spots index

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Segmentation a Spot of MicroArray Using Kernel Density

Estimated Density Curve

Estimated Density Curve

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Feature Calculation for MicroArray

GKDE KDE GMM

Ratio of mean=0.965 Ratio of mean=1.013 Ratio of mean=0.901

GenePix Irregular GenePix Circular GenePix Rectangular ScanAlyze

Ratio of mean=1.086 Ratio of mean=0.813 Ratio of mean=0.866 Ratio of mean=0.685

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Evaluation Accuracy of Segmentation Methods for Spike Spots

The comparisons of SSREs obtained by GMM, GKDE, KDE, ScanAlyze, and GenePix 6.0 for spike genes are listed

Array	Sum of square of relative errors							Relative performance		
	GKDE	KDE	GMM	ScanAlyze	GenePix irregular	GenePix circular	GenePix rectangular	GKDE	KDE	GMM
1	85.482	82.495	85.482	243.383	301.408	320.264	258.886	64.878	66.105	64.878
2	35.009	45.899	55.025	117.817	152.598	123.384	128.127	53.310	61.042	53.296
3	80.421	77.148	80.421	286.480	317.287	303.845	317.147	71.838	73.070	71.838
4	29.861	28.021	30.170	36.042	31.277	34.409	35.664	4.528	10.409	3.539
5	10.401	7.908	10.410	15.256	20.094	16.070	19.737	31.823	48.161	31.765
6	7.605	7.229	7.647	53.292	57.603	55.908	58.491	85.757	85.525	85.678
7	60.911	58.383	60.916	90.118	115.513	115.779	114.534	32.411	34.106	32.405
8	110.991	101.908	110.992	163.735	219.798	196.928	207.335	32.213	37.761	32.213
9	31.005	31.740	32.980	130.081	147.429	132.150	132.280	74.744	75.712	74.763
10	26.500	22.211	26.505	32.285	31.157	33.630	37.984	13.662	12.664	13.640
11	149.074	130.196	149.739	244.790	286.494	261.726	272.590	39.101	46.813	38.830
12	675.010	648.212	674.388	769.750	826.411	761.916	767.239	11.406	14.923	11.408
13	12.302	12.108	12.222	16.525	23.215	14.541	17.244	16.086	16.745	15.956
14	6.781	6.550	6.786	94.839	107.370	101.951	104.794	92.850	93.093	92.843
15	12.705	12.425	12.739	29.883	33.558	44.113	45.079	37.404	38.419	37.371
16	9.910	10.025	9.987	17.570	24.496	21.485	24.715	43.590	42.842	43.305
Average relative performance								45.283	48.593	45.231

Array 1 is obtained by swapping the dyes of Array 1. Relative improvement is specified by $(\text{Min}(\text{GenePix}, \text{ScanAlyze}) - \text{Method}) / \text{Min}(\text{GenePix}, \text{ScanAlyze})$ as a percentage.

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3D Image Segmentation

39th Slice Sagittal image Coronal image GMM

Rat Images

GMM

K-mean

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Image Reconstruction and Analysis

Target image 5% noise 10% noise 30% noise

Line profiles

POEM

FBP

Simulated prompt sinogram

OSEM

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What is bio-image statistics?

- ◆ Five image types of features:
 - ◆ Pixel statistics (max, min,...)
 - ◆ Textures (Entropy, histogram,...)
 - ◆ Polynomial decompositions
 - ◆ High contrast features (e.g. object number, spatial distribution, size, shape, ..., etc.)
 - ◆ Standard image transforms (Fourier, wavelet, Chebyshev)

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What is bio-image statistics?

- ◆ A good understanding of **mathematics and statistics** and an interest in mastering the biological/biochemical/bio-image/Optical images background is essential.

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Main Goal

- ◆ In recent years (after 2000 B.C.), new state-of-art scanning machines like nano-PET/CT, naon-SPECT/CT, micro-MR,...,and etc bring imaging science into investigation of super micro details inside live animal.
- ◆ This course is designed to understand how to analyze the feature and interesting of bio-image by using statistical tools.

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Tool

- ◆ Statistics
- ◆ Feature extraction and selection
- ◆ Segmentation
- ◆ Registration
- ◆ Clustering, classification and annotation
- ◆ Indexing and retrieval
- ◆ Visualization

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Resources

- ◆ **BioImage Suite** is an integrated image analysis software suite developed at Yale University. BioImage Suite has been extensively used at different labs at Yale since about 2001. The last stable version is 2.6.1 which was released on April 6th, 2009.
<http://www.bioimagesuite.org/>

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Resources

- ◆ **Center for Bio-Image Informatics**: The UCSB Bio-Segmentation Benchmark dataset consists of 2D/3D images and time-lapse sequences that can be used for evaluating the performance of novel state of the art computer vision algorithms. Tasks include segmentation, classification and tracking. For each class of problem, at least one ground truth dataset is available. We also provide performance metrics for comparing the results of the algorithms with the ground truth. Additional ground truth data will be posted as they become available. <http://www.bioimage.ucsb.edu/>

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Statistics

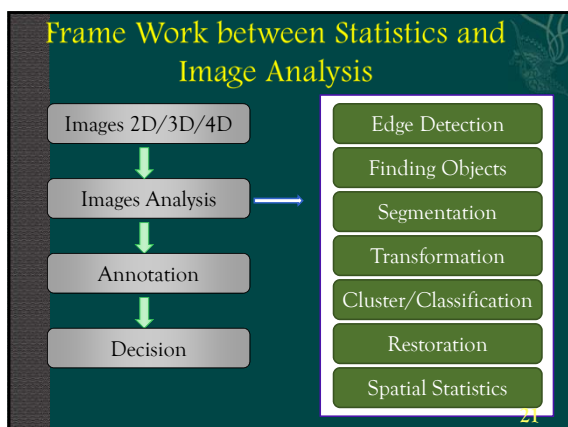
- ◆ **Smoothing:** Gaussian, Median, Mean,...
- ◆ **Classification:** Gaussian mixture model, Bayesian approach,...
- ◆ **Correlation:** Pearson, Kendall, Spearman
- ◆ **Relevant Test:** Chi-test, Kappa,...
- ◆ **Prediction:** Spatial statistics, Regression, General linear model,...

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Feature extraction and selection

- ◆ ROI-Based Analysis
- ◆ Feature extraction: Size, Shape, Geometric, Area, Volume, Distance,...
- ◆ Pixel intensity
- ◆ Clustering and Classification

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什麼是醫學影像定量?

影像定量分析: 透過數學或統計方法找出影像特徵並與疾病進行關連分析的一種技術。

目的: 建立客觀量化數據(資訊), 提升(輔助)定(質)性判讀結果。

重點: 如何從影像上找出影像特徵??

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Examples

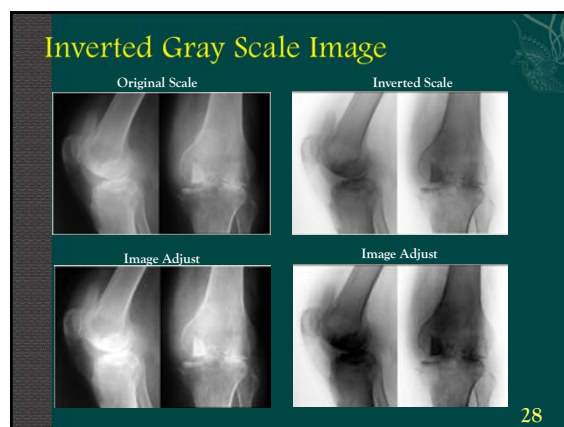
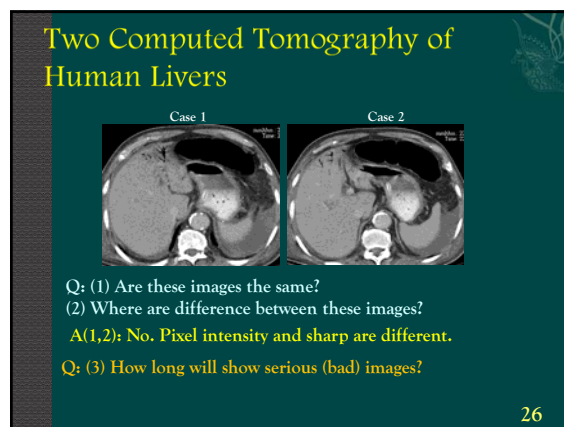
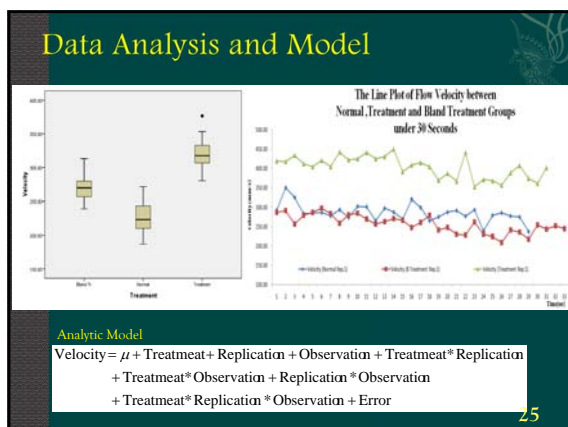
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Two Treatments of Doppler Ultrasound Images

Q: What are different features between these images?

A: (1). Time interval of Peaks (2). Maximum value of peaks (3). Distance between positive and negative peaks.

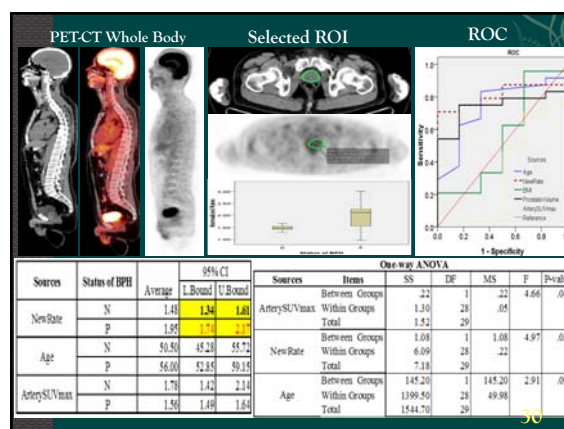
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PET-CT : Benign Prostate Hyperplasia

- ◆ PET: the metabolic information was estimated from PET images, including SUV of arteries, iliac arteries, kidney, skin, whole body, and etc.
- ◆ CT: The rate of prostate volume
- ◆ Analysis
 - ◆ ROC (Receiver Operating Characteristic), AUC (Area under ROC), Pearson's correlation, and ANOVA

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SPECT : The Parkinson Diseases

- ◆ Trodat-1 Imaging
 - ◆ The functional activity volume of corpus stratum was varying depend on the progress of PD
- ◆ MR/CT/SPECT (Normal):
 - ◆ Realignment, Normalization
- ◆ Evaluation risk of PD by statistical model
 - ◆ Naïve Bayes, Logistic Regression,.....etc.

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Co-registered with Trodat-1 brain images by using MR

Before After

MR template image

Abnormal Normal

■ E-cobaltame
■ E-cobaltame
■ E-cobaltame

■ E-cobaltame
■ E-cobaltame

Probability (Risk) of PD By Naive Bayse

Index	Values(%)
Accuracy	86
Specificity	67
Sensitivity	100
PPV	80
NPV	100

Risk

0.996 0.998 0.998 0.546 0.551 0.001 0.037

Abn Abn Abn Abn Nor Nor Nor

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Cardiac 64-Slice MDCT

- ◆ 64-Slice MDCT
 - ◆ Dedicate Anatomical Information of Cardiac Coronary Artery Disease (CAD)
- ◆ Extracted Physical Information
 - ◆ Long axis, Short Axis, Intensity Values, ..., etc.
- ◆ Statistical tools
 - ◆ ROC (Receiver Operating Characteristic), AUC (Area under ROC), Pearson's correlation, and ANOVA

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3D MPR of Cardiac 64-Slice MDCT

Extracted Features

Item	SS	df	MS	F	P-Value
Middle length rate of systolic long axis/short axis	0.18	1	0.18	5.024	0.020
Widest Crosses	1.46	47	0.03		
Total	1.64	48			
Lower length rate of systolic long axis/short axis	0.02	1	0.02	6.257	0.015
Widest Crosses	0.17	47	0.13		
Total	0.19	48			
Middle length rate of diastolic long axis/short axis	0.18	1	0.18	7.301	0.003
Widest Crosses	0.95	36	0.03		
Total	1.07	37			
Upper average of sagittal slice in cross-section average of length	0.22	1	0.22	8.756	0.005
Widest Crosses	0.28	47	0.06		
Total	0.30	48			
Middle average of sagittal slice in cross-section average of length	1.29	1	1.29	13.773	0.001
Widest Crosses	4.40	47	0.09		
Total	0.69	48			
Upper average of sagittal slice in diastolic average of length	0.60	1	0.60	15.399	0.000
Widest Crosses	1.41	36	0.04		
Total	0.03	37			

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Ultrasound: Elastogram

- ◆ Elastogram
 - ◆ Visualized the elasticity of Breast Cancer by RGB
 - ◆ Avoiding embarrassing and keeping privacy
- ◆ Extracted Color Information
 - ◆ Intensity of Red/Blue channel
 - ◆ Computing characteristic of ROI
- ◆ Statistical tools
 - ◆ ROC (Receiver Operating Characteristic), AUC (Area under ROC), Pearson's correlation, Logistic Regression, ANOVA,....., and etc.

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Normal Benign Malignant

Mean Plot between Treatments

ANOVA Test

处理组	数目	平均数	df	平均平方和	F	显著性
校正后的常数	6688	4.314	2	3349.215	50.514	0.000
校正后的因子	1417793	2.89	1	1417793.289	21341.125	0.000
校正后的残差	6688	4.30	2	3349.215	50.414	0.000
总数	1442006	5.40	99	10352.344		
校正后的总数	10352	3.44	57			

a. R 平方 = .647 (调整后模型的 R 平方 = .634)

预测数 Mean (red)

Cat	平均数	标准误差	95% 信頼区間	
			下界	上界
Malignant	171.434	2.105	167.216	175.651
Benign	160.969	1.870	157.222	164.716
Normal	145.316	1.684	141.982	148.650

The Future

- ◆ An integrated project union together with interdisciplinary
- ◆ Imaging/Images
- ◆ Information Science
- ◆ Bioinformatics
- ◆ Integrated Medical Information and Biological Images for Personal Medicine

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Acknowledgement

- ◆ Thank all students for their efforts in projects
- ◆ Thank E-DAH and VGHKS for providing images
- ◆ Thank Prof. Hung, Ding, Du, MD Lu, MD Chen, and other related experts/technological radiologist for their support and comments in projects



Thanks for Your Attention



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