



NTU Center of Genomic Medicine

臺大基因體醫學研究中心

主持人：楊泮池 教授
陳明豐 教授
陳為堅 教授
執行長：錢宗良 教授



NTU Center of Genomic Medicine (NTU-CGM)

- 2002/10/30 教育部正式通過計畫
- 2003/03/25 通過設置**臺大基因體醫學研究中心** (NTU-CGM)
- 2003/04/02 協助臺大SARS團隊相關研究支援
- 2004/04/01 臺大基因體醫學研究中心正式啟用
- 2005/12/01 更名為**臺大醫學卓越研究中心** (NTU-RCME)
- 2007/07/27 生醫分子影像核心實驗室正式落成啟用
- 2008/09/12 幹細胞核心實驗室正式啟用
- 2009/09/09 永齡基金會捐贈共軛焦顯微鏡等貴重儀器
- 2009/12/15 美商應用生命系統(ABI)捐贈新一代核酸定序儀
- 2010/03/19 代謝體核心實驗室成立
- 2010/08/01 校方通過變更回為「**臺大基因體醫學研究中心**」
- 2010/11/20 與臺灣大學/醫院/醫學院合作，推動轉譯醫學研究

NTU Center of Genomic Medicine

Mission

To establish state-of-the-art core facilities, support cutting edge genomic research for scientists in NTU

Our Focus

Disease-based translation research, emphasize on **cancer** and **infectious diseases**

Bench Mark

Top-ranking university in Asia-Pacific area
Osaka University, Melbourne University



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NTU Center of Genomic Medicine

主持人：楊泮池 院長 陳明豐 院長 陳為堅 院長

Genomic Medicine Cores

執行長：錢宗良教授

Cancer Division

楊泮池教授

Infectious Disease Division

陳培哲教授

Administration Office & Teaching Resource Center



Proteomics

周綠蘋

Microbial Genomic

陳培哲、
王錦堂等

Microarray

楊泮池、
俞松良等

Tissue Bank

林中梧

Transgenic & Knockout Mouse

蘇銘嘉、
林淑華

Genetic Epidemiology

陳為堅、
簡國龍等

Bioinformatics Biostatistics

莊曜宇、
蕭朱杏等

Stem Cell

何弘能
錢宗良

Metabolomics

郭明良、
曾宇鳳、
郭錦樺

任務目標

- 基因醫學研究領域設置10個設備完善之核心實驗室
整合跨領域、跨院校研究交流
- 每年支持計畫主持人發表50篇以上之國際學術論文
期刊，及超過2項專利申請案
- 每年舉辦20個以上之教育訓練課程
- 每年舉辦暑期研習營，培訓學生超過30人次
- 促進國際學術交流及合作

Milestone Achieved

- Fully function core facilities in genomic research
- Facilitate multidisciplinary collaborations in NTU
- Support NTU PI publish high ranking research papers > 50/year, patent > 2/year
- Educational programs and workshops > 20/year
- NTU students summer research program
- International credibility and collaboration

國際學術交流：諾貝爾獎得主參訪研究中心

Barry Marshall
Nobel Laureate Medicine 2005
At 112th Anniversary
Symposium
26.6.2007



Dr. Barry Marshall
The 2005 in Medicine

It has been a most interesting and impressive visit. Thank you for showing us NTU Medical Center.

Best wishes,

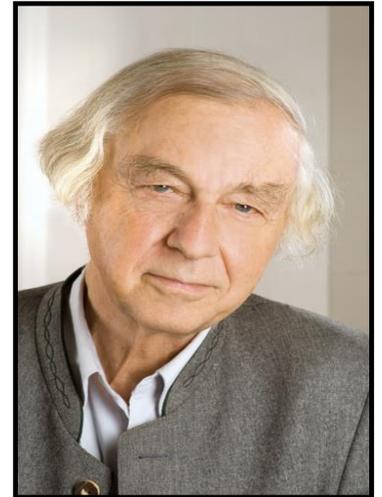
Joh. Deisenhofer
Aug. 15, 2008



Dr. Johann Deisenhofer
The 1988 in Chemistry

A most interesting tour showing the NTU Research Center of Medical excellence, thank you,
Robert Huber
(ROBERT HUBER)

10. Nov. 2008



Dr. Robert Huber
The 1988 in Chemistry

醫學研究中心暑期研習營 (2006-2011)

臺灣大學



量化研究成果 (2011/04-2011/12)

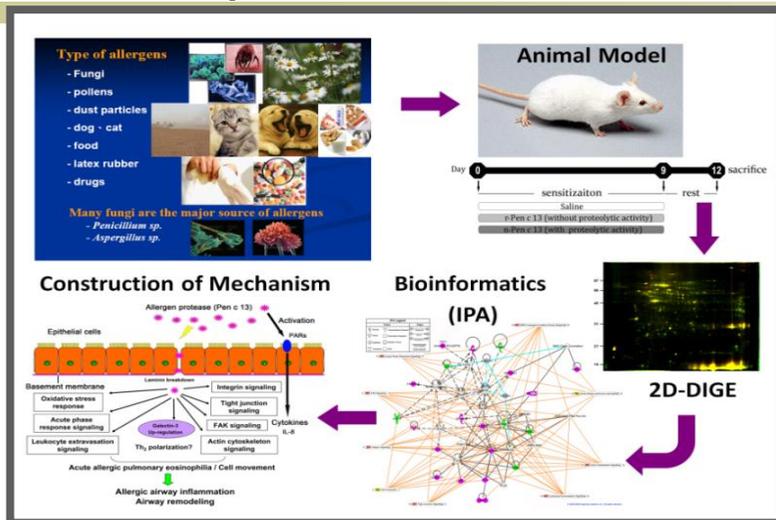
Listing		Total	Significant
Published Articles	Journals	196	96
	Conferences	61	27
	Technology Reports		
Patents	Pending	8	
	Granted	3	
Personal Achievements	Honors Awards	15	4
	Keynotes Given by PIs	24	4
	Editor for Journals	29	18

量化研究成果 (2011/04-2011/12)

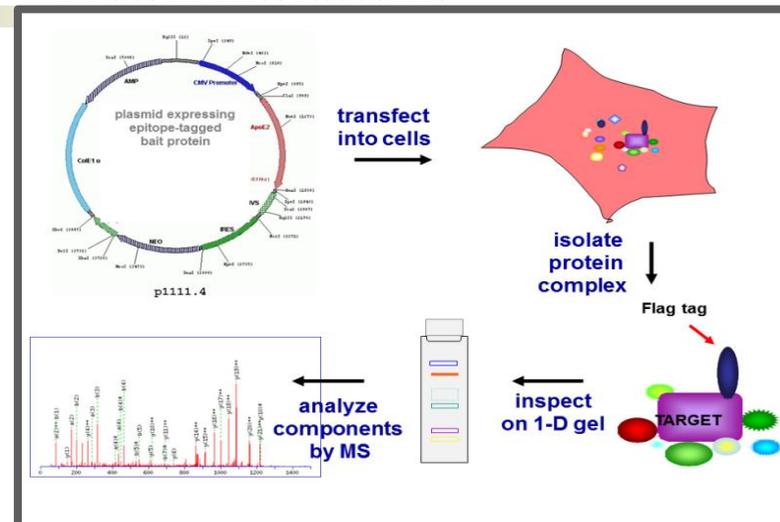
Listing		Total	Significant
Workshops Conferences	Item	20	4
	Participants	2679	1200
International Cooperation Activities	Item		
	Participants		
Training Courses	Hours	1453	120
	Participants	345	210
Technological Services	Item	2459	
	Service Fee	1,842,280	

Proteomics strategies for clinical research

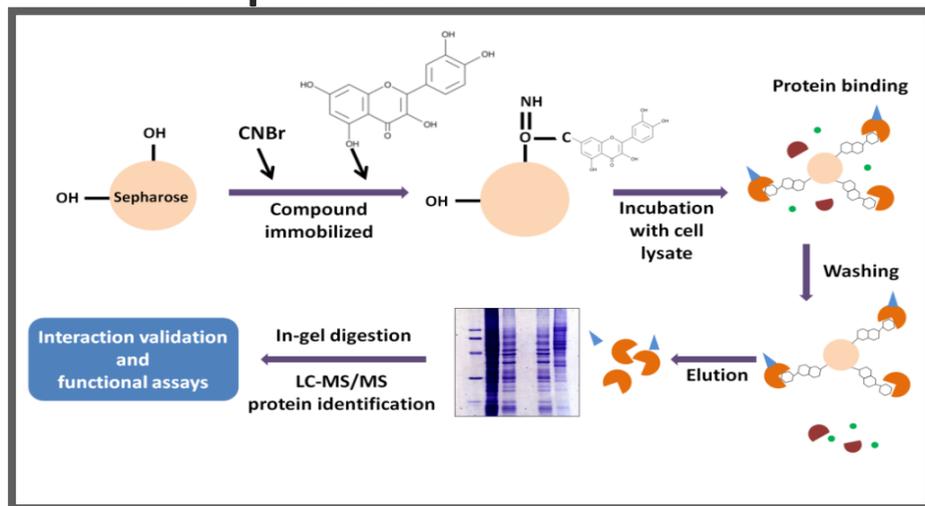
Differential proteome in murine model



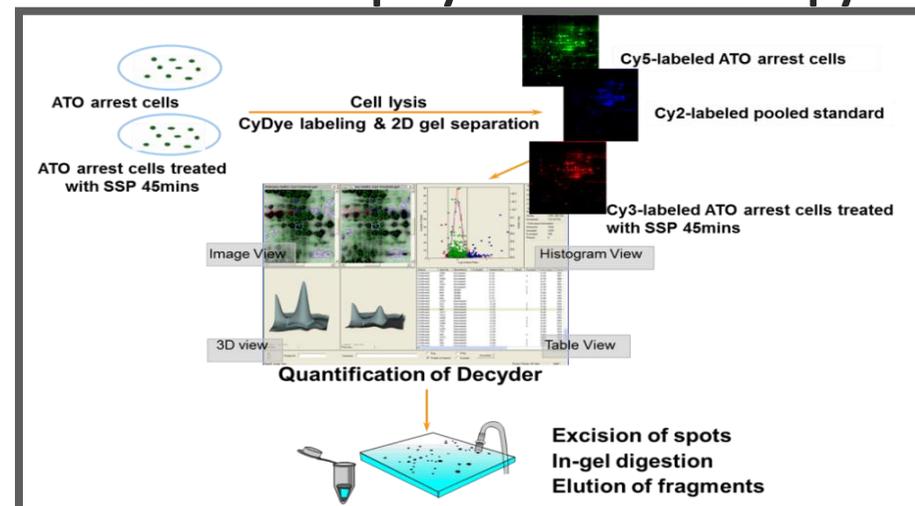
Protein interactome



Chemical proteomics



Differential display of cancer therapy



Associations Between Hepatitis B Virus Genotype and Mutants and the Risk of Hepatocellular Carcinoma

Hwai-I Yang, Shiou-Hwei Yeh, Pei-Jer Chen, Uchenna H. Iloeje, Chin-Lan Jen, Jun Su, Li-Yu Wang, Sheng-Nan Lu, San-Lin You, Ding-Shinn Chen, Yun-Fan Liaw, Chien-Jen Chen

For the REVEAL-HBV Study Group

(A successful collaboration between Microbial Genomics Core Laboratory and Academia Sinica)

HBV genotype	Precore 1896 variant	BCP 1762/1764 variant	No. of participants	Person-years of follow-up	No. of HCC cases	HCC incidence (per 100 000 person-years)	Adjusted HR† (95% CI)	P‡
B	Wild type	Wild type	198	2404.8	14	582.2	1.0 (referent)	
B	Wild type	A1762T/G1764A	82	950.1	9	947.3	1.35 (0.58 to 3.12)	.48
B	G1896A	Wild type	413	5175.0	9	173.9	0.20 (0.09 to 0.46)†	<.001
B	G1896A	A1762T/G1764A	110	1319.9	7	530.3	0.62 (0.25 to 1.54)	.30
C	Wild type	Wild type	129	1633.2	6	367.4	0.73 (0.28 to 1.89)	.51
C	Wild type	A1762T/G1764A	115	1242.1	28	2254.2	2.99 (1.57 to 5.70)†	<.001
C	G1896A	Wild type	41	497.3	4	804.3	1.38 (0.45 to 4.20)	.57
C	G1896A	A1762T/G1764A	73	911.2	2	219.5	0.36 (0.08 to 1.59)	.18

* HCC = hepatocellular carcinoma; HBV = hepatitis B virus; BCP = basal core promoter; HR = hazard ratio; CI = confidence interval.

† Adjusted for sex and age at recruitment.

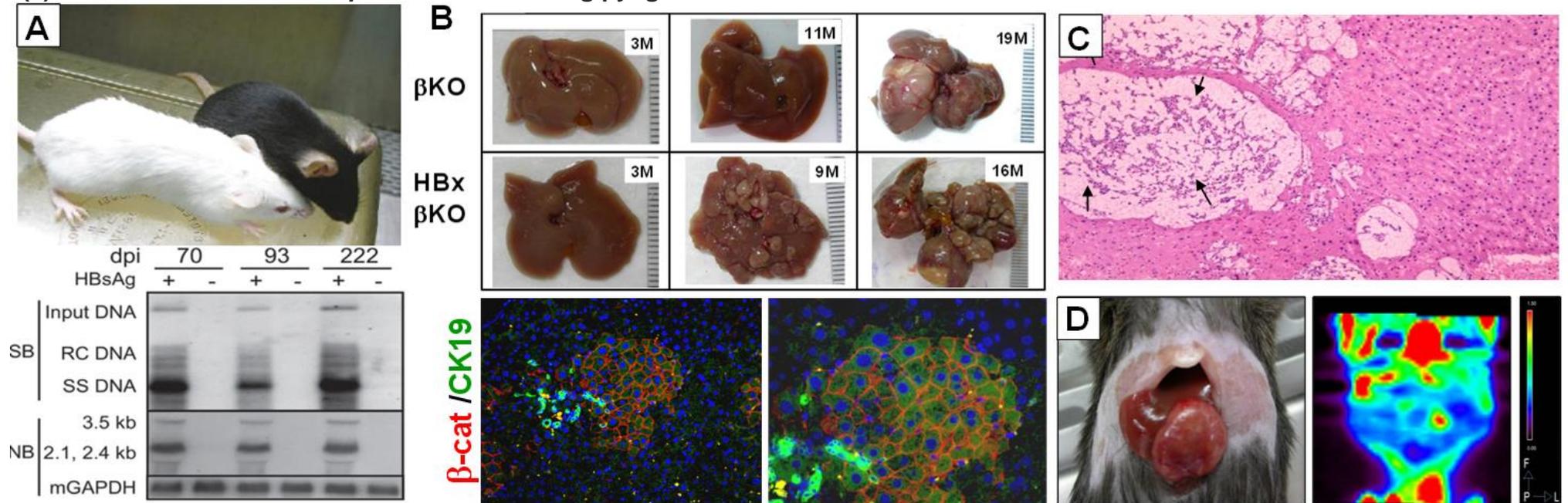
‡ *P* values (two-sided) were from Cox proportional hazards models.

(Yang HI, et al. *J Natl Cancer Inst.* 2008)

Establish the animal models for Infectious diseases



- (1) Immunocompetent mouse model for chronic HBV infection
- (2) Orthotopic HCC mouse model → Gene therapy and drug development
- (3) The knockout mouse model for viral protein induced HCC from progenitor cells
- (4) Woodchuck models for hepatitis virus infection and HCC
- (5) The mice model for *K. pneumoniae* causing pyogenic liver abscess

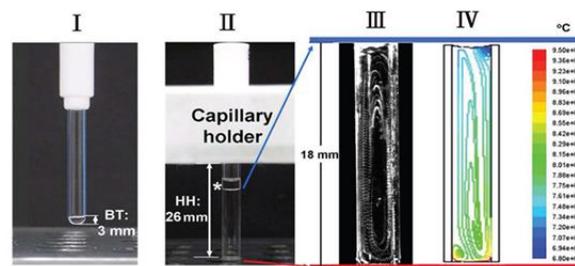
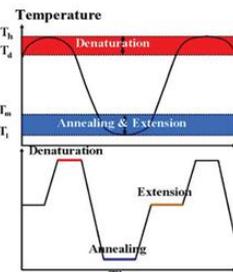
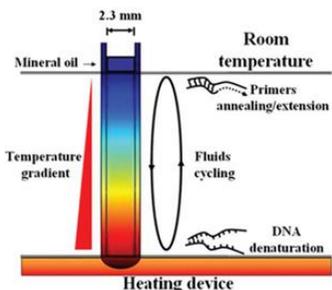
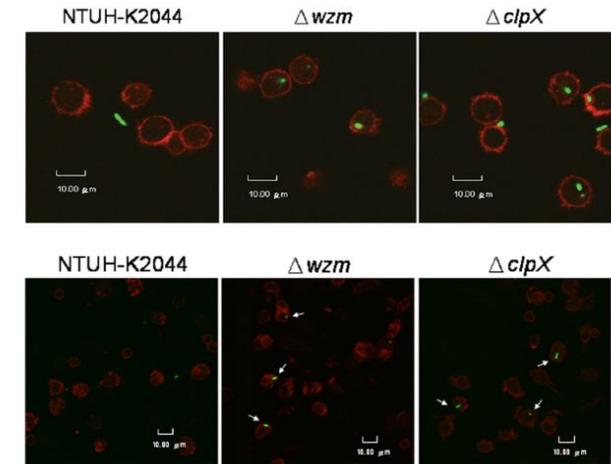
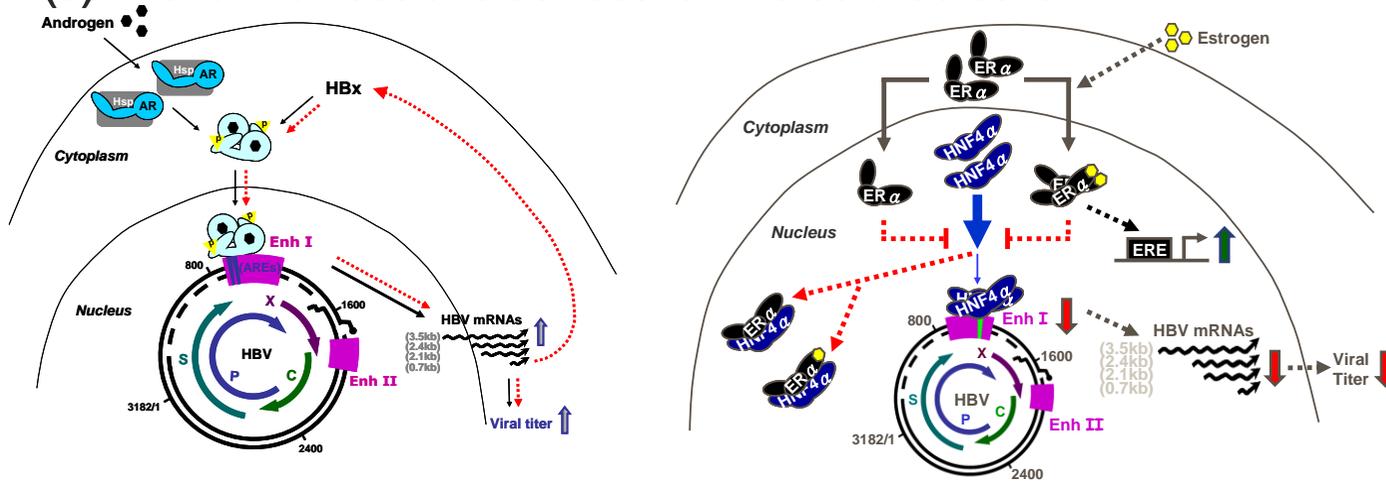


Lin YJ, et al. *Proc. Natl. Acad. Sci. USA.* 2010
 Fang CT, et al. *J. Exp. Med.* 2004
 Huang KW, et al. *Proc. Natl. Acad. Sci. USA.* 2010
 Wang EY, et al. *Proc. Natl. Acad. Sci. USA.* 2011

Identify novel mechanisms for infectious diseases



- (1) the sex steroids in regulating virus infection and liver carcinogenesis
- (2) the genetic loci for *K. pneumoniae* pathogenesis
- (3) the factors in determining the persistence of viral infection
- (4) the genome evolution in liver cancer development
- (5) novel methods and devices for microbial detection

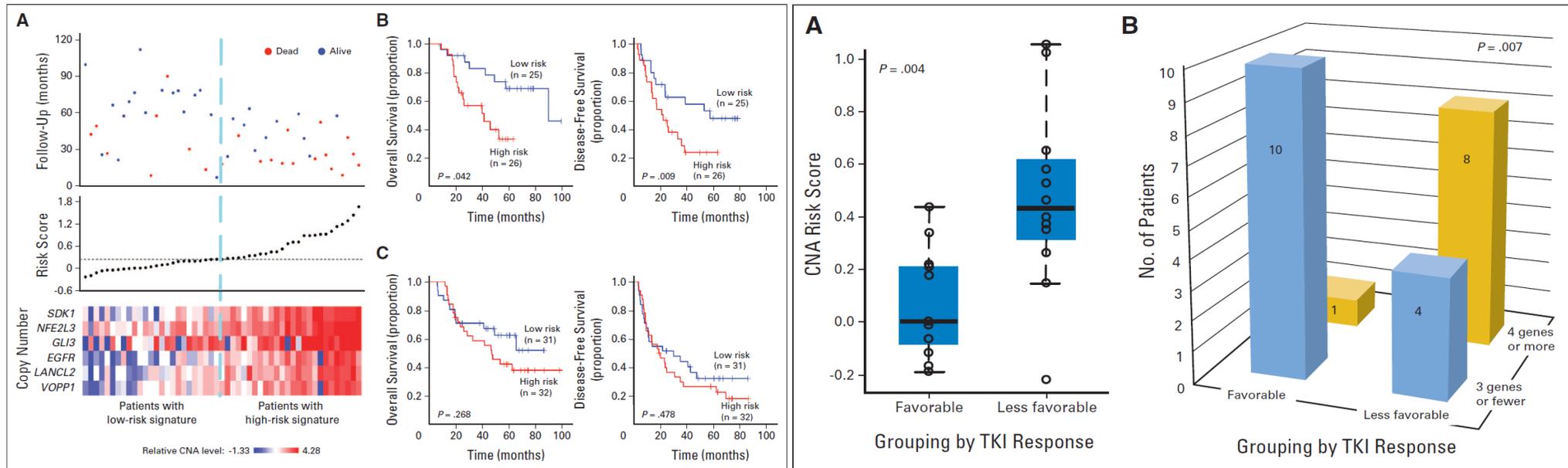


Wang SH, et al. *Gastroenterology*. 2012
 Tao Y, et al. *Proc. Natl. Acad. Sci. USA*. 2011
 Pan YJ, et al., *Infect Immun*. 2011
 Chou WP, et al. *Biotechniques*. 2011
 Wang SH, et al. *Hepatology*. 2009
 Liu WH, et al. *Gastroenterology*. 2009

Predicting Therapeutic Response of Lung Adenocarcinoma with EGFR Mutations



30% of lung adenocarcinoma patients with EGFR-activating mutation are non-responders in epidermal growth factor receptor tyrosine kinase inhibitors (EGFR-TKIs) treatment. We identify a prognostic signature unique to the EGFR-activating mutant patients. Our finding may lead to a better patient management for EGFR-mutant patients.



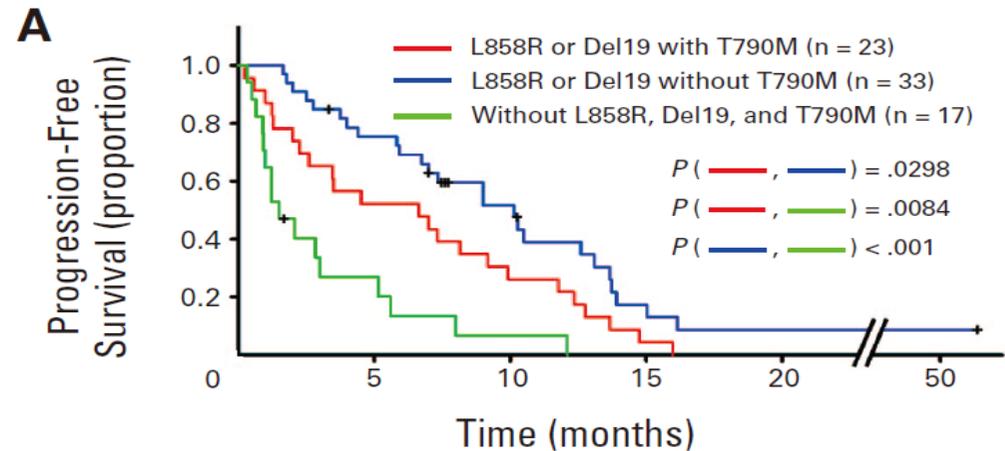
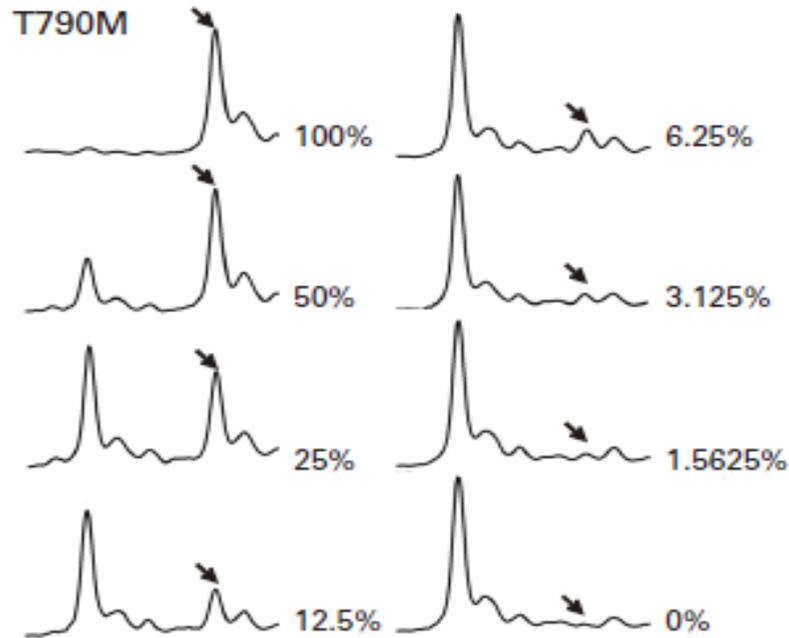
Microarray Core

Publications: [Journal of Clinical Oncology 2011, 29:3435-42.](#)
 Patent Application: [US US1975-2011.7.4, PCT](#)

Pre-treatment EGFR T790M Mutation Predicts Shorter EGFR-TKI Response in Lung Cancer



Non-small cell lung cancer (NSCLC) patients with EGFR-activating mutations have excellent response to EGFR-tyrosine-kinase-inhibitors (EGFR-TKIs) but T790M mutation accounts for most TKI drug resistance. This study used highly sensitive methods to detect T790M before and after TKI therapy and found the pre-treatment T790M mutation was associated with shorter PFS of EGFR TKI therapy in NSCLC patients.



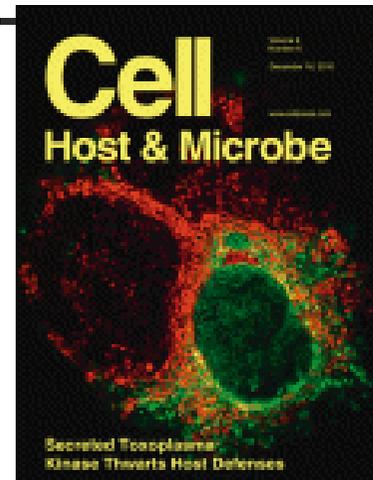
Cox Regression Model

Variable	Hazard Ratio	95% CI	P
L858R or Del19 without T790M	1.000		
L858R or Del19 with T790M	1.854	1.044 to 3.292	.035
Without L858R, Del19, and T790M	4.965	2.524 to 9.765	< .001

Enterovirus vs. microRNA



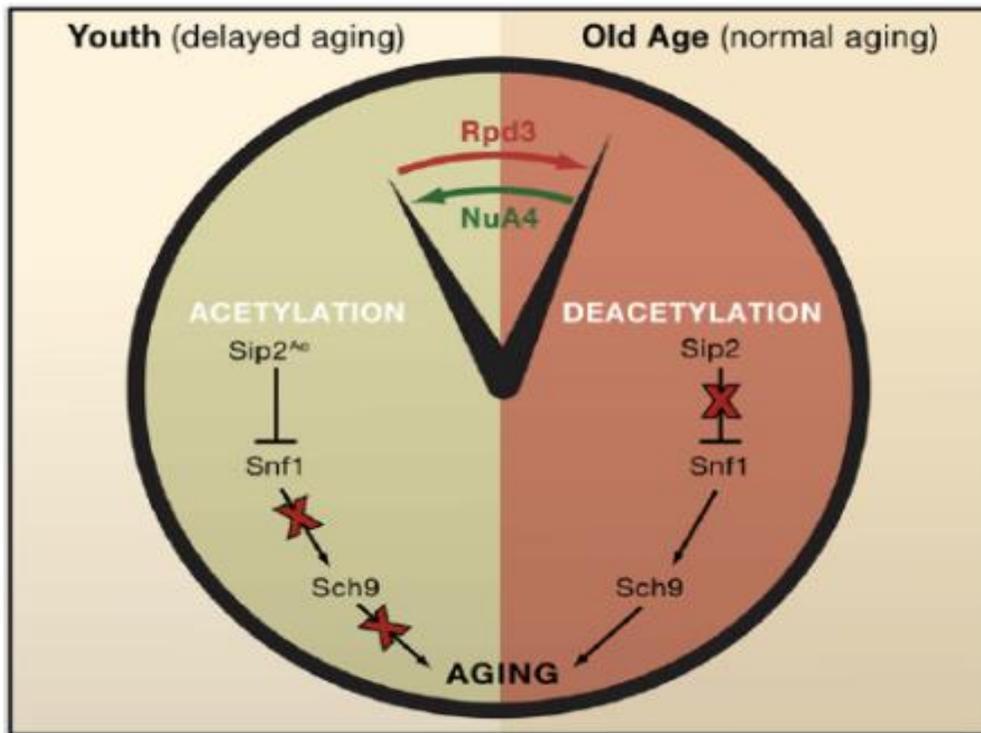
- Enterovirus can up-regulate microRNA-141 to shutdown host protein synthesis and start to make viral proteins and promote viral replication ([Cell Host & Microbe](#), January issue of 2011).
- This work demonstrated that microRNA-141 will be a novel therapeutic target.



Acetylation of Yeast AMPK Controls Intrinsic Aging Independently of Caloric Restriction

Cell

Jin-Ying Lu,^{1,2} Yu-Yi Lin,³ Jin-Chuan Sheu,⁴ June-Tai Wu,⁵ Fang-Jen Lee,⁵ Yue Chen,⁶ Min-I Lin,¹ Fu-Tien Chiang,¹ Tong-Yuan Tai,⁴ Shelley L. Berger,⁷ Yingming Zhao,⁶ Keh-Sung Tsai,^{1,2,4} Heng Zhu,^{8,10,*} Lee-Ming Chuang,^{2,4,*} and Jef D. Boeke^{9,10,*}



Acetylation and The Elixir of Life

在Yeast模式，限制卡路里可延緩老化，然而細胞內生性的老化仍然受到 *Acetylation* 調控！

¹Department of Laboratory Medicine, National Taiwan University Hospital

²Graduate Institute of Clinical Medicine, College of Medicine

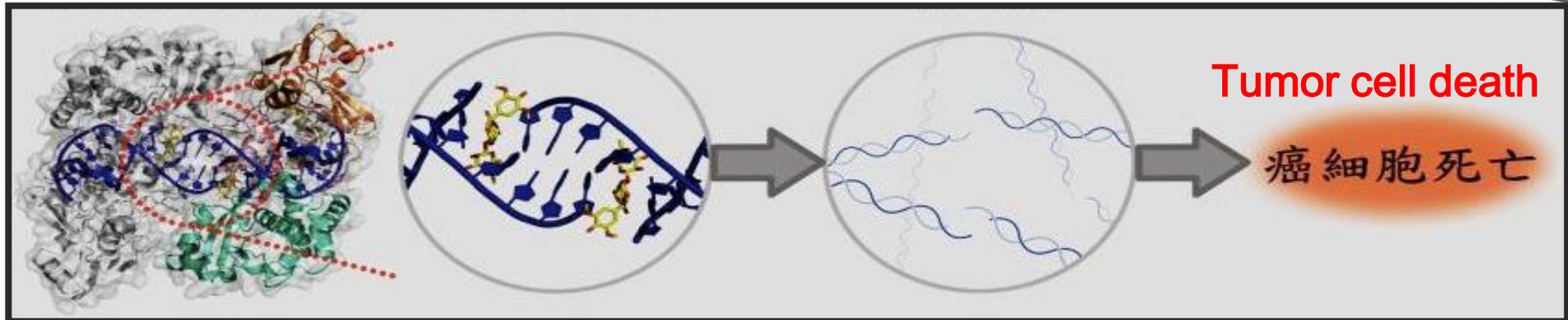
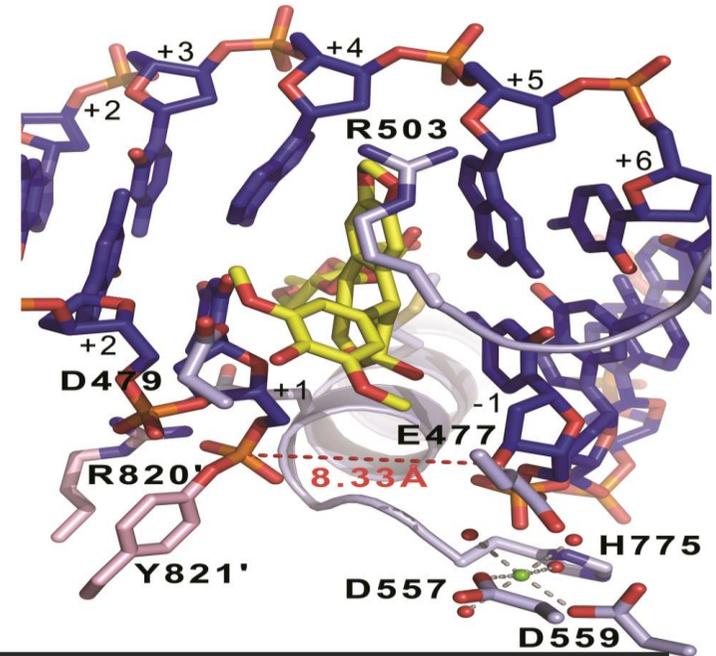
³Institute of Biochemistry and Molecular Biology, College of Medicine

Structural Basis of Type II Topoisomerase Inhibition by the Anticancer Drug Etoposide

Chyuan-Chuan Wu,¹ Tsai-Kun Li,^{2,3} Lynn Farh,⁴ Li-Ying Lin,^{1,5} Te-Sheng Lin,^{1,5} Yu-Jen Yu,¹ Tien-Jui Yen,¹ Chia-Wang Chiang,^{1,5} Nei-Li Chan^{1,5*}

¹台大醫學院生物化學暨分子生物學研究所

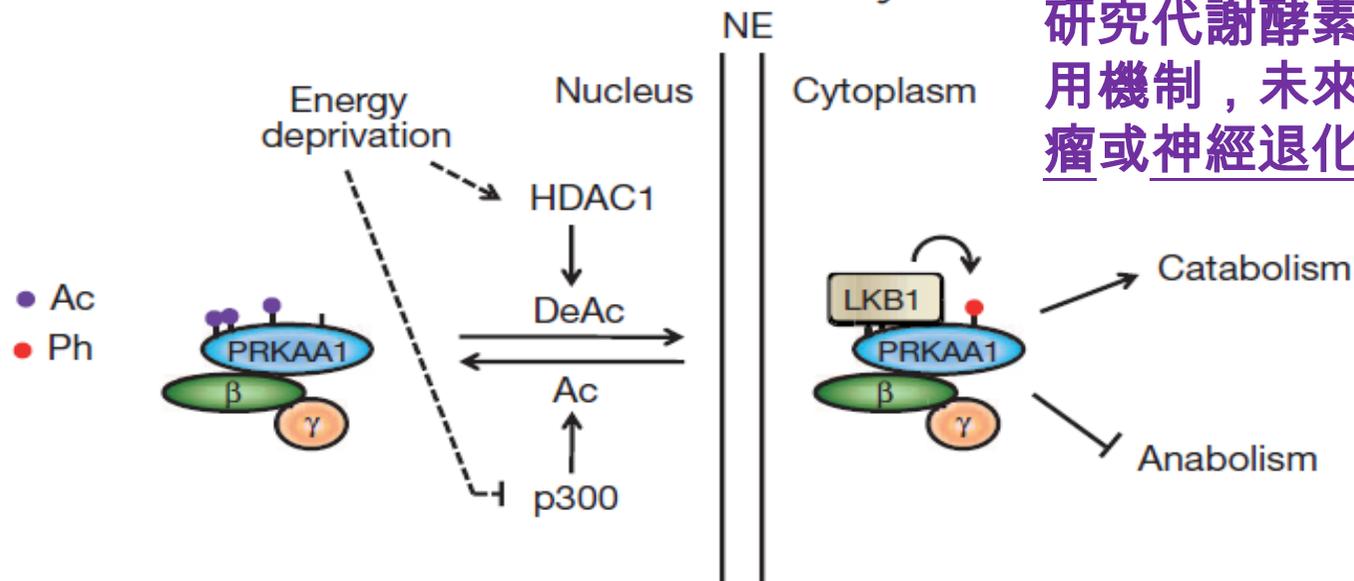
研究闡明抗癌藥etoposide引發DNA斷裂，導致癌細胞死亡結構基礎，可用於開發低副作用的新型抗癌藥物。



Functional dissection of lysine deacetylases reveals that HDAC1 and p300 regulate AMPK

Yu-yi Lin^{1,2,3}, Samara Kiihl^{4*}, Yasir Suhail^{5*}, Shang-Yun Liu¹, Yi-hsuan Chou¹, Zheng Kuang^{6,7}, Jin-ying Lu^{2,8}, Chin Ni Khor⁹, Chi-Long Lin⁹, Joel S. Bader⁵, Rafael Irizarry⁴ & Jef D. Boeke^{6,7}

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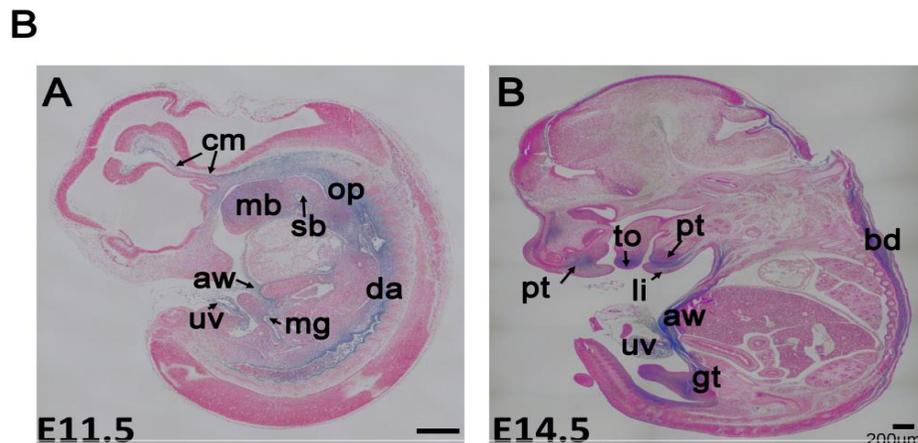
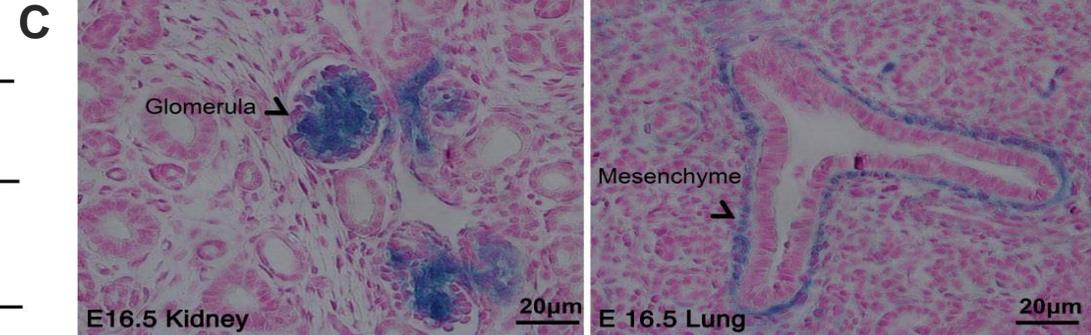
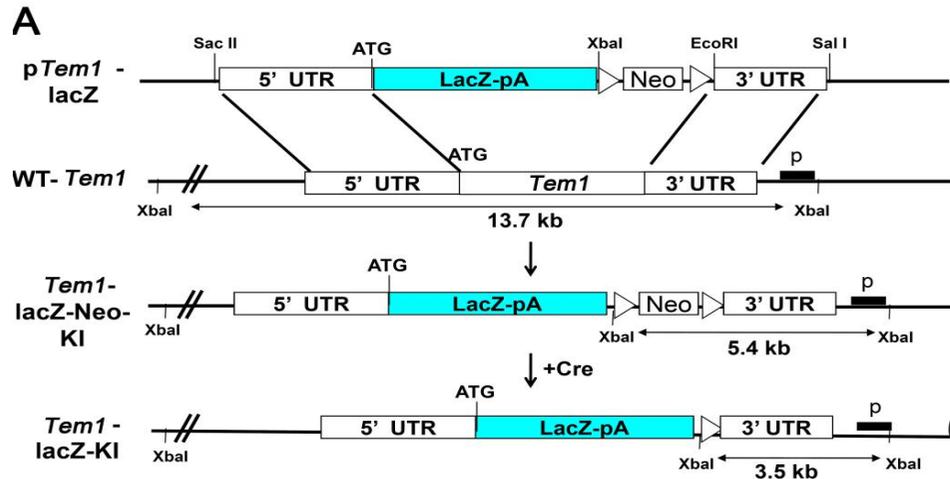
研究代謝酵素Lysine deacetylases 作用機制，未來將可能運用在治療新生腫瘤或神經退化等之疾病。

Figure 4 | Deacetylation of PRKAA1 specifically enhances its physical interaction with LKB1 kinase. a, Phosphorylation of endogenous PRKAA1 is reduced and unresponsive to acetylation status upon knockdown of *LKB1*.

Gene Knockout Mouse Core Laboratory



Gene targeting and expression analysis of mouse protein using the lacZ reporter



(A) Maps of wild-type *Tem1* allele, recombinant allele with the single exon of the *Tem1* gene replaced by the lacZ/neo gene, and final recombinant allele with lacZ but without neo.

(B) Whole-mount Bluo-Gal staining followed by sagittal sectioning showing *Tem1*-lacZ expression in E11.5 and E14.5 mouse embryos.

(C) At E16.5, strong *Tem1*-lacZ expression emerges in the glomeruli of the kidney and the mesenchymal cells or mesenchymal-like cells or around the bronchial epithelial cells of the lung.

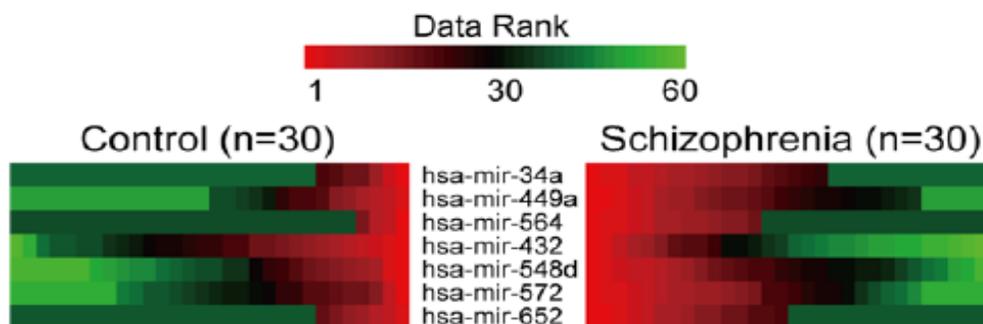
(ref. Huang et al., Gene Expression Patterns, 2011)

週邊血液之7種微小核糖核酸 (microRNA) 表現偏差 作為精神分裂症生物標記的可能性

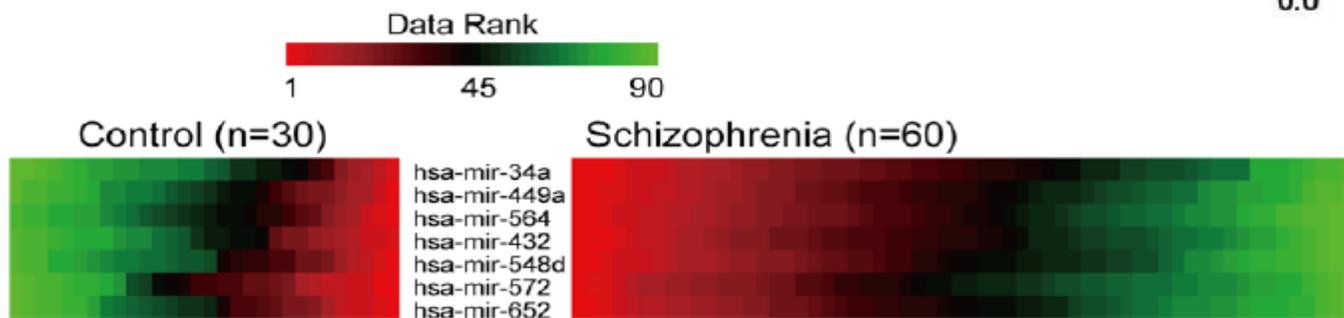
(Lai et al., 2011, *PLoS ONE* 6: e21635)



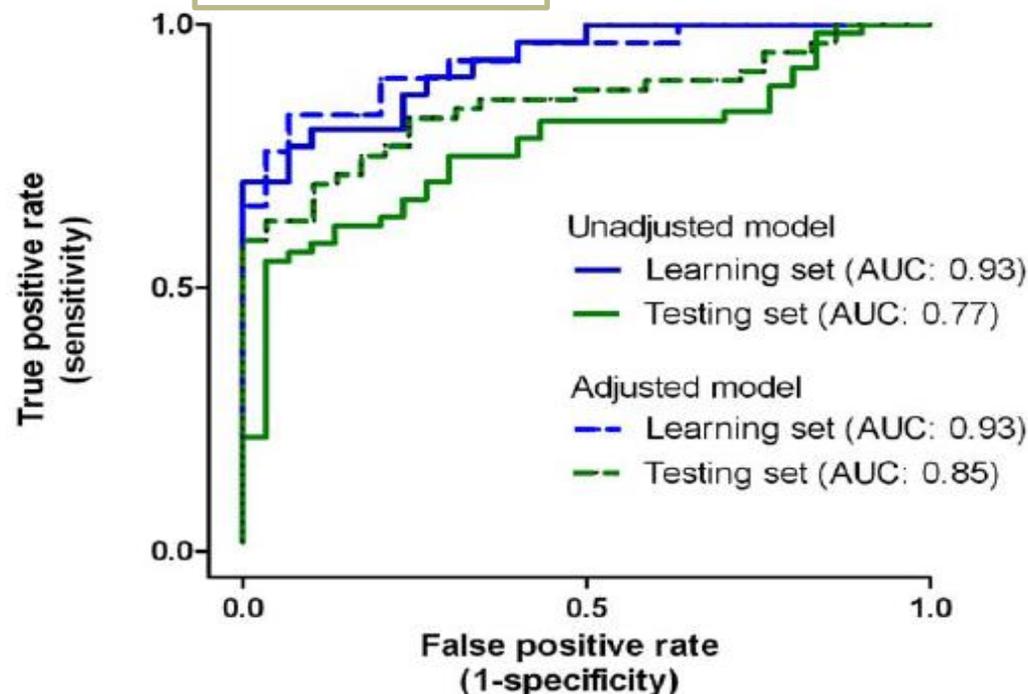
A Learning Set



B Testing Set



ROC curve



針對病人及其家屬之菸鹼酸貼片皮膚紅腫反應進行QTL 全基因體掃描，尋獲新的精神分裂症遺傳連鎖標誌14q32.12

(Lien et al., 2011, *Schizophrenia Bulletin*. doi: 10.1093/schbul/sbr054)

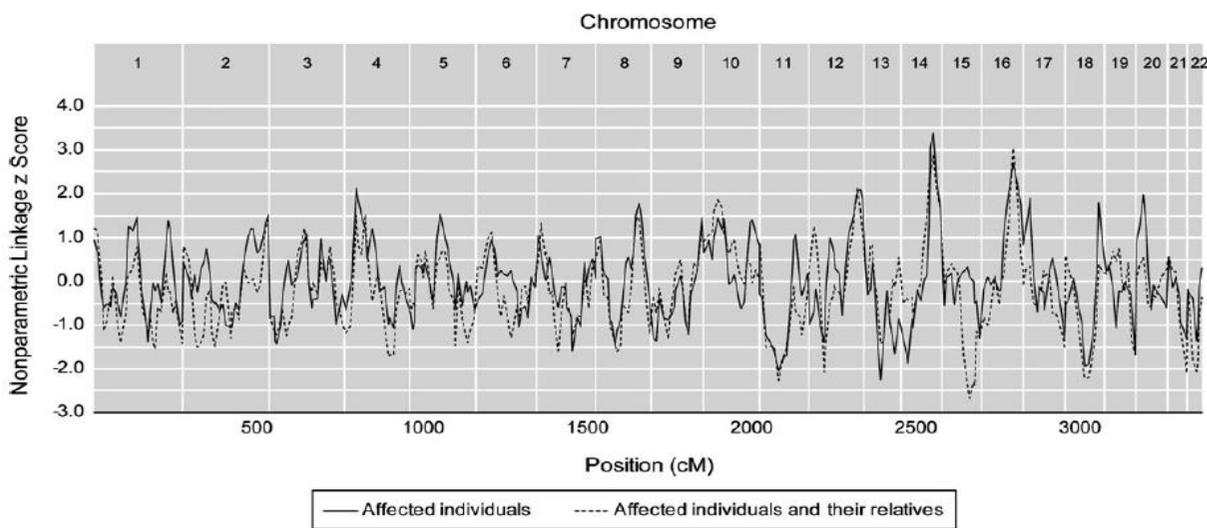
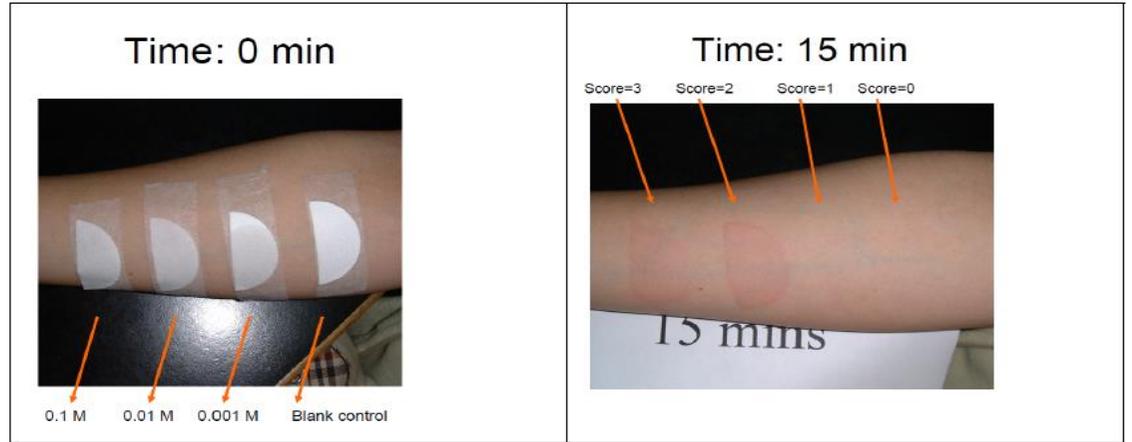
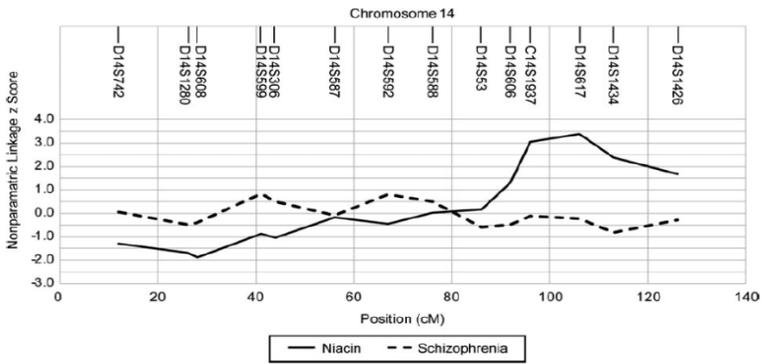
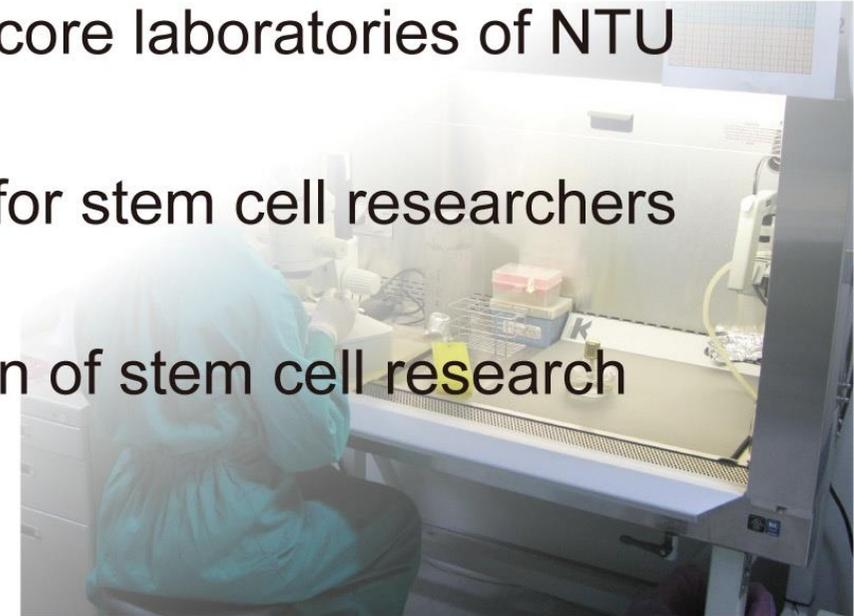


Figure S1. Implementation of the niacin skin patch and the scoring for the flush response.



Stem Cell Core

- To supply hESC for the researchers within NTU medical campus
- To provide the techniques and routines for characterization of hESC
- To integrate the service provided by the core laboratories of NTU **Center of Genomic Medicine**
- To supply hESC and technical supports for stem cell researchers nationwide
- To promote the international collaboration of stem cell research



Human ES cell and feeder cell supply

(2008/10/31-2011/06/15)

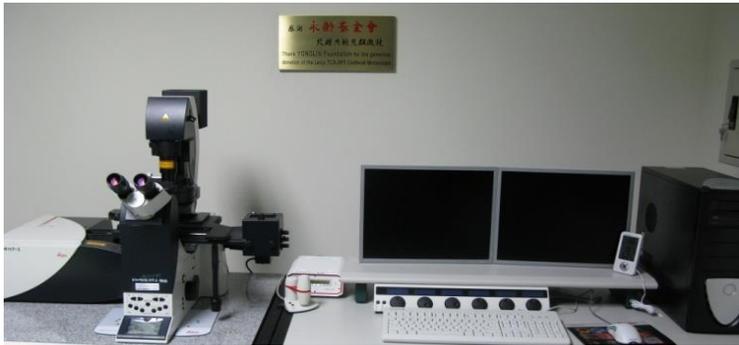


Applicant Units	PI	Dish	Research subject
NTU, IOB	李宣書	318	Generation of insulin-producing cells
NTU, Dept. of Food Science and Technology	蔣丙煌	61	The effect of Ginkgo extract on dopaminergic neuron differentiation
NTUH, Dept. of Surgery	吳耀銘	86	Hepatic differentiation of ES cell
NTU, Dept. of Pharmacology	郭明良	52	Cancer stem cell study
NTU, Institute of Microbiology	鄧述諄	275	Telomerase capacity of ES cell differentiation
NTU, Institute of Clinical Genomics	陳佑宗	60	Genetic manipulation of ES by PiggyBac transposon
NTU, Dept. of Pharmacology	陳青周	20	Hepatic differentiation of ES cell
NTU, Biomedical Engineering	林峰輝	192	Expansion ES cell by bioreactor
NTUH, Dept.of Obs/Gyn	何弘能	70	Establish new ES cell line
NTUH, Dept. of Ophthalmology	楊長豪	184	The oxidative stress of ES cells induced by paraquat

Core Facilities



Confocal (Leica TCS SP5)



Tissue FAXS



Laser-Enabled Analysis and Processing System (LEAP™)



Cell Sorter (BD FACS Aria II)



Stem Cell Core Workshop



Introducing Stem Cell Core service and its core facilities on 2010/05/26

NTU participants		Non-NTU participants	
Professors	11	Students	1
Students	27	Industrial	5
Research Assistants	22	YongLin Healthcare Foundation	2
Postdoctoral Research Fellow	8		
Medical Doctors	2		
Medical Laboratory Scientist	1		
Subtotal	71	Subtotal	8
Total		79	



Core PI, Prof. Ho gave a brief introduction.



Core Co-PI, Prof. Chien introduced core service and instruments



Instrument Demonstration: Confocal microscope



Instrument Demonstration: IVIS, Xonogen

Education



International Students

Year	Nationality	Subtotal
2008	Thailand、 Spain(2)、 Poland、 Slovenia、 Serbia(2)	7
2009	Spain、 Portugal、 Hungary、 Italy	4
2010	Thailand、 the Netherlands(2)、 Spain、 Italy、 Italy、 Austria、 Hungary	8
2011	Thailand、 Slovenia(2)、 Hungary、 Italy、 USA	6
Total		25

Summer Camp Students

Year	Grade	Subtotal
2008	3rd grade 、 4th grade	2
2009	1st grade 、 2nd grade 、 5th grade (2)	4
2010	2nd grade	3
2011	1st grade 、 2nd grade (2) 、 3rd grade	4
Total		13

- ❖ **代謝體研究服務**: 樣品處理、樣品分析、代謝體諮詢服務、生物標的確效(biomarker verification)、代謝圖譜分析
- ❖ **代謝體分析方法開發**: 開發並改良LC-MS與GC-MS分析方法
- ❖ **數值分析方法開發**: run order analysis, outlier detection, peak detection, baseline correction, alignment
- ❖ **代謝體研究**: 包括臨床及營養代謝體研究以達到轉譯醫學之個人化醫療。

代謝體核心實驗室 Metabolomics Core Lab.

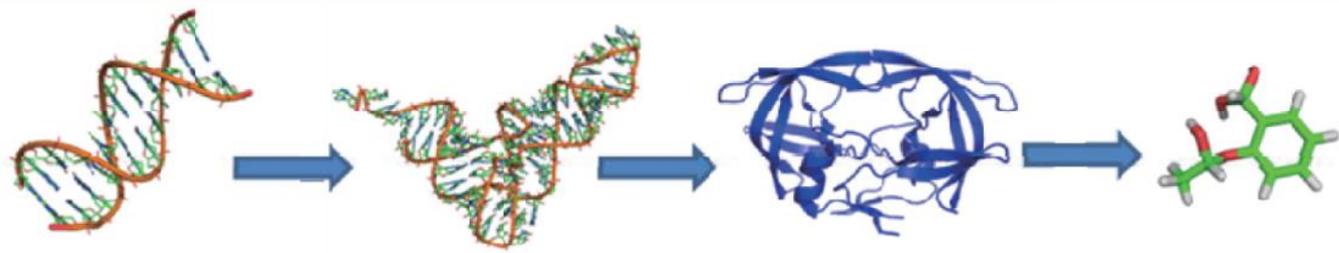


Figure 1. Central Dogma of Molecular Biology with Metabolome



Honors Awards

MetaPick, a denoising and peak picking algorithm for extracting single compound from complex mixtures in liquid chromatography/mass spectrometry (LC/MS) metabolomics data, Metabolomics 2011, Best informatics presentation award.

Services

- 代謝體服務包括代謝體研究與實驗設計諮詢、樣品處理、儀器分析質譜圖譜分析；服務申請者包括臨床醫師與基礎學術研究者，樣品種類包括血液、尿液、細胞萃取液及呼出氣體等。
- 100年度已完成之服務共計21案，總計樣品數量1124個，進樣針數4496針，品管針數540針。收樣中及預計進行之服務共計25案，總計樣品數量2380個。

Future Directions

- Backup new research Institutes:
Cancer, Genomics, IBMS (Neurosciences)
- Multidisciplinary translational research
- International collaborations (MD Anderson, Terry Fox, Peking University, etc.)
- New area and facility:
Metabolomics
High throughput sequencer
Novel drug design for target therapy
Molecular Imaging & Medical devices, etc.

校內跨中心合作

針對 系統生物學研究中心、分子生醫影像研究中心 兩大研究領域之合作，將以人才交流共同提升國際化為目標。

每年將規劃暑期研習營培訓大學部同學，栽培未來跨領域研究新秀。

共同舉辦國際學術研討會，為臺灣大學與英、美、日及歐盟等大學建構學術合作平臺。

2011 暑期研習營說明會

分子生醫影像中心
(Molecular imaging center)
生物醫學訊息分析
分子標記設計
基因組學
分子影像、顯微晶片

蛋白質學蛋白質質
分析核心
(Proteomics core)
蛋白質序、蛋白質互作用
蛋白質表現及活性分析
蛋白質核酸交互作用

遺傳流行病學核心
(Genetic Epidemiology core)
基因組學標記
遺傳學分析
遺傳學診斷

組織庫核心
(Tissue bank core)
組織收集保存
雷射微切、免疫染色
分子標記分析
基因組學表現分析

代謝物核心
(Metabolomics core)
作為基礎研究的橋梁與
臨床藥物代謝的潛在發展

時間：6月24日(四) 08:50-12:00

地點：醫學院基醫大樓 102 講堂(舊社)

08:50 - 09:05 報到(領取書面資料)

09:05 - 09:10 基因體醫學研究中心主任
楊院長 泮池 致詞

09:10 - 09:15 分子生醫影像中心主任
孫副院長 啓光 致詞

09:15 - 09:30 Molecular Imaging Center

09:30 - 09:45 Proteomics Core

09:45 - 10:00 Microbial Genomics Core

10:00 - 10:15 Microarray Core

10:15 - 10:30 Tissue Bank Core

10:30 - 10:45 Coffee Break

10:45 - 11:00 Transgenic Mouse and
Gene Knockout Core

11:00 - 11:15 Genetic Epidemiology Core

11:15 - 11:30 Bioinformatics and
Biostatistics

11:30 - 11:45 Stem Cell Core

11:45 - 12:00 Metabolomics Core

微生物基因组核心
(Microbial genomics core)
病原體基因组
病原體基因组
病原體基因组
病原體基因组

基因微陣列及晶片核心
(Microarray core)
基因篩選、晶片製備及服務
寡核苷酸及短干涉RNA
診斷晶片

基因轉殖與基因敲除核心
(Transgenic and
knockout mouse core)
基因轉殖、基因敲除
轉殖動物模型、基因敲除
動物模型

幹細胞核心
(Stem Cell core)
建立長期且穩定人
類胚胎幹細胞的技術以
及資料庫

生物資訊核心
(Bioinformatics and
biostatistic core)
生物資訊分析
生物統計
生物資訊諮詢
資料庫

幹細胞核心
(Stem Cell core)
建立長期且穩定人
類胚胎幹細胞的技術以
及資料庫

歡迎大學部同學們踴躍參加!

台大基因體醫學研究中心、分子生醫影像中心 敬邀

中心已獲得或申請中之其他計畫/經費說明

99年度教育部補助之邁向頂尖大學經費佔中心實際經費資源約30%，校務基金及中心核心服務收入約10%，向外爭取之經費包含貴重儀器與其他經常費用超過60%，教育部補助本中心之效益比超過兩倍以上。

100年度中心已獲得之其他計畫包括：

- 教育部顧問室轉譯醫學人才培育計畫
- 生技醫藥國家型科技計畫：通過多項計畫
- 國科會100年幹細胞及再生醫學前瞻計畫
- 跨國頂尖中心計畫 (MD Anderson)

中心100年度核定經費 (單位：千元)

補助項目	核定金額
研究人力費	21,040
業務費 (耗材、物品及雜項費用)	8,960
儀器設備費	10,000
合 計	40,000

中心101年度期經費規劃說明

年次	研究人力費	耗材及雜項費用	研究設備費	國外或大陸地區差旅費	出席國際學術會議差旅費	小計
101	42,417	12,598	20,455	1,280	3,327	80,077
102	43,536	12,602	21,623	1,280	3,327	82,368
103	44,422	12,600	20,513	1,280	3,327	83,142
104	45,173	12,604	20,149	1,280	3,327	82,533

中心5年期經費規劃說明

- 研究人力費：(核心實驗室基本服務人力; 規劃聘用10位專案助理教授，儲備具國際競爭力人才):
~ NT\$ 43,000,000/year
- 耗材及雜項費 (實驗室耗材及研究中心基本維護費):
~ NT\$ 13,000,000/year
- 研究設備費 (汰舊換新設備、大型儀器組件維護):
~ NT\$ 21,000,000/year
- 國際交流經費 (邀請國外學者及出席國際會議差旅費):
~ NT\$ 5,000,000/year

Total: ~ NT\$ 82,000,000/year



中華民國精彩一百

敦品勵學 賞國慶人

敬請 委員們支持與指正

