分子生物學應用

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主題介紹

I. 即時定量聚合酶連鎖反應 (Real-time quantitative PCR, RT-qPCR/Q-PCR)

II. 報告基因分析 (Reporter gene assay)

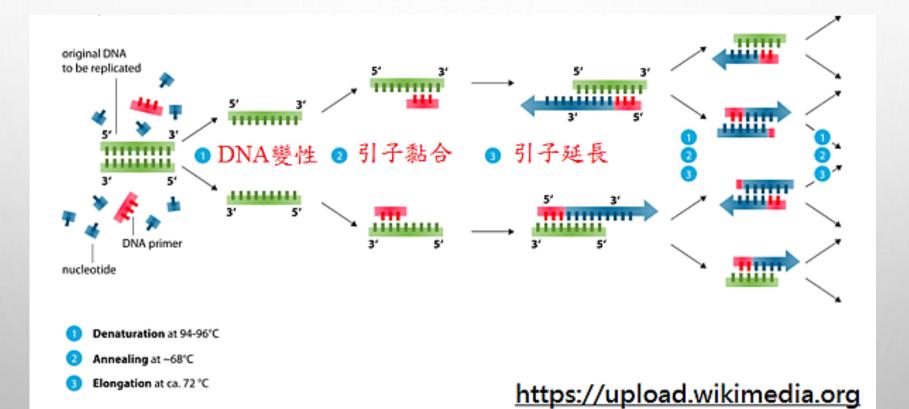
III. 核糖核酸干擾 (RNA interference, RNAi)

I. 即時定量聚合酶連鎖反應 (Q-PCR)

上課內容

- · 聚合酶連鎖反應(PCR)
- Q-PCR原理
- Q-PCR步驟
- Q-PCR反應混合物
- TaqMan Probe(探針)& SYBR Green(染料)
- 使用的儀器/系統
- · 如何避免Q-PCR錯誤
- Q-PCR應用

聚合酶連鎖反應 (Polymerase Chain Reaction, PCR)



- -以電泳確認PCR產物
- -以2ⁿ(n表示循環次數)的速度在複製DNA



The Nobel Prize in Chemistry 1993

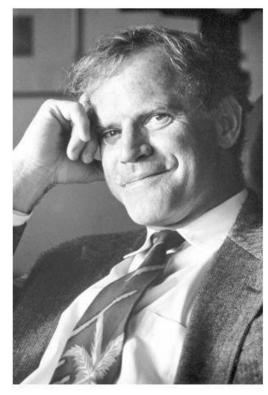


Photo from the Nobel Foundation archive.

Kary B. Mullis

Prize share: 1/2

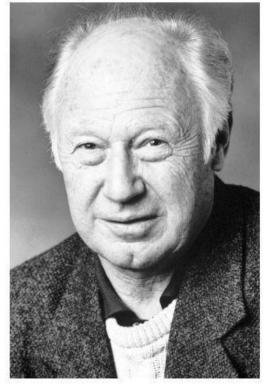


Photo from the Nobel Foundation archive.

Michael Smith

Prize share: 1/2

Kary B. Mullis: for his invention of the polymerase chain reaction (PCR) method.

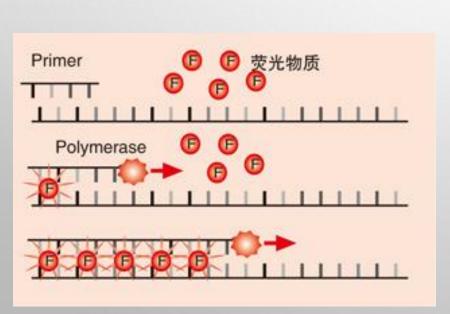
Michael Smith: for his fundamental contributions to the establishment of oligonucleotide-based, site-directed mutagenesis and its development for protein studies.

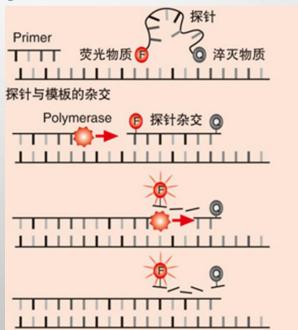
Q-PCR 原理

uses reverse transcription (反轉錄) to produce a DNA template from an RNA source that can then be amplified.



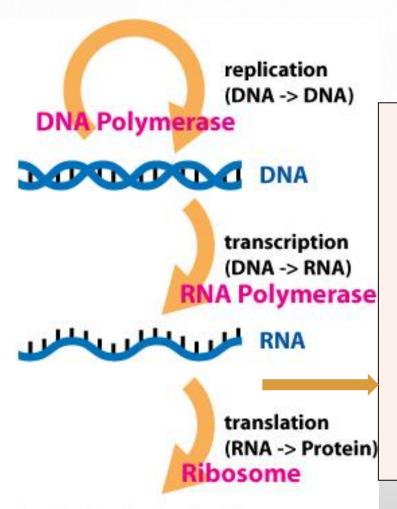
- 這是一種基於聚合酶鏈反應(PCR) 的技術。
- 將反應放入機器中,該機器使用相機或檢測器實時監測觀察 並螢光反應檢。
- 檢測分子產生的螢光,隨著反應的進行,該分子增加。
- 可以定量相對少量的PCR產物。







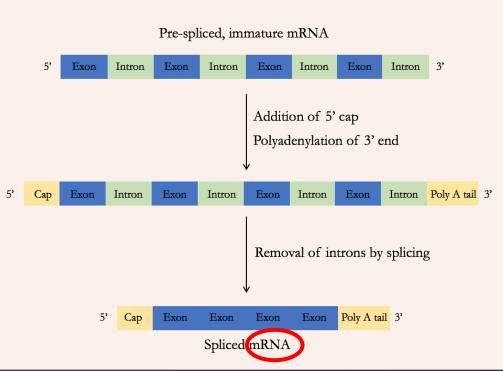
中心法則



0-0-0-0-0 Protein

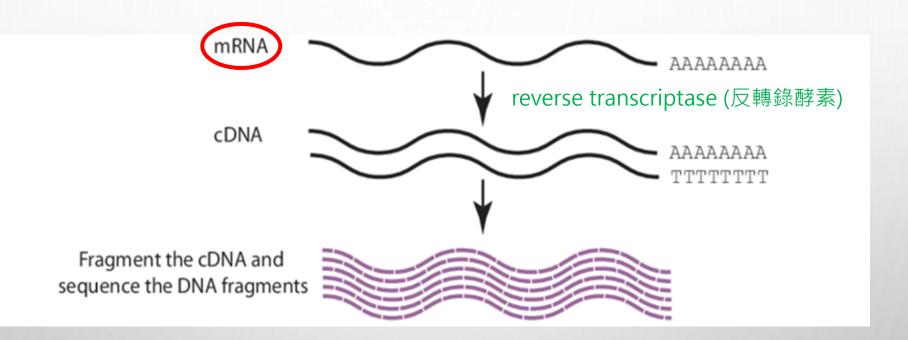
https://zh.wikipedia.org

成熟RNA修飾

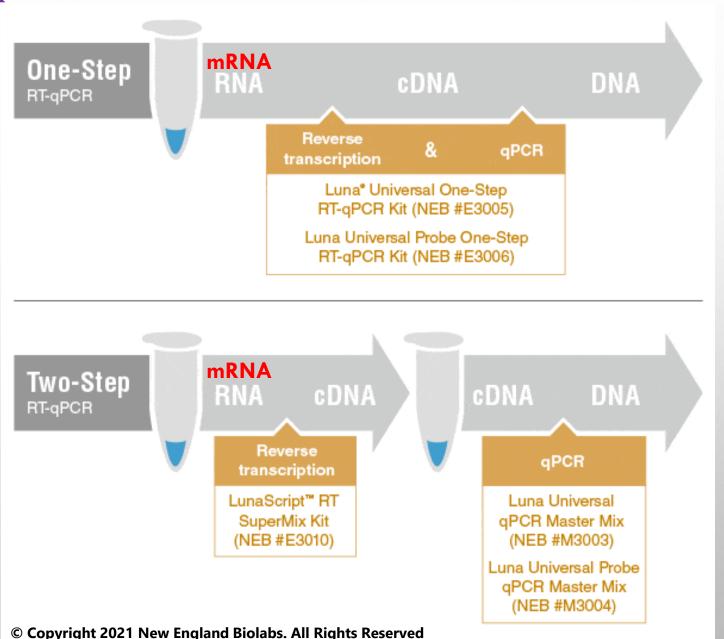


https://commons.wikimedia.org/

信使RNA (mRNA) 反轉錄成為互補 DNA (cDNA) 的 流程

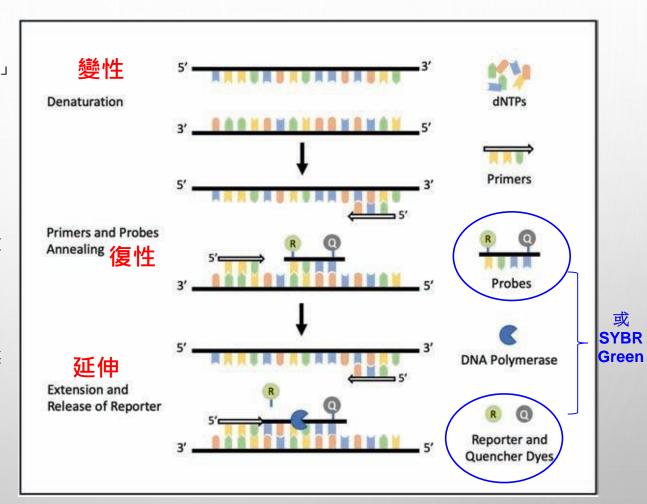


Q-PCR步驟





- a. 變性-高溫用於將雙鏈DNA「熔化」 成單鏈,並鬆動單鏈DNA中的二級 結構。通常使用DNA聚合酶可以承 受的最高溫度(通常為95°C)。如 果範本GC含量高,可以增加變性時 間。
- b.復性-在復性過程中,互補序列有機會雜交,因此使用基於引物的計算熔融溫度(Tm)的適當溫度(低於引物Tm的5°C)。
- c. 延伸-在70-72°C時,DNA聚合酶的活性是最佳的,引物延伸以高達每秒100個鹼基的速率發生。當即時螢光定量PCR中的擴增子很小時,該步驟通常與使用60°C作為溫度的復性步驟相結合。



Q-PCR反應混合物

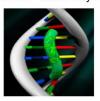




TaqMan Probe(探針)& SYBR Green(染料)

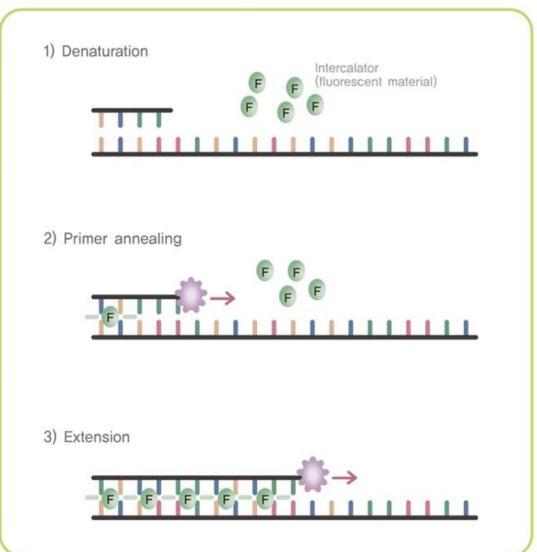
- 每個Q-PCR都包含一個螢光分子(例如 TAQMAN® 探針或 SYBR® GREEN染料),用於監測 PCR產物的積累。
- 隨著標靶基因擴增數量的增加,螢光團發出的螢光量也隨之 增加。
- 為使用Q-PCR 進行基因表達研究而開發的兩種類型的化學物質是:
- (1) SYBR Green 染料
- (2) TaqMan Probe 探針



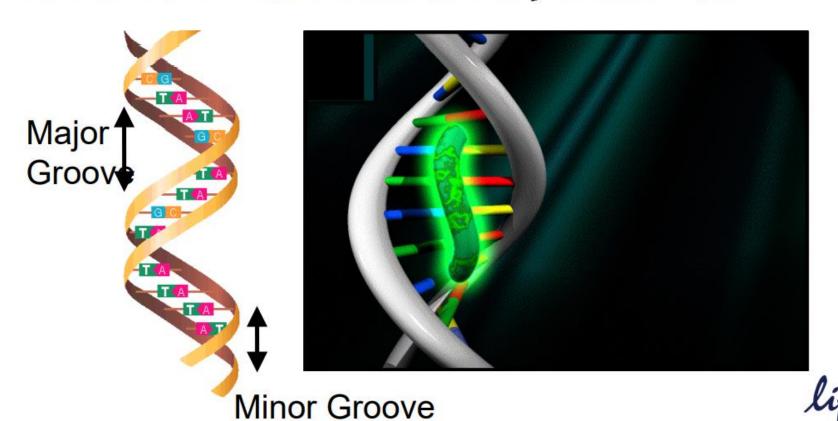


(1) SYBR GREEN 染料

- 這是一種染料,當它在 DNA的微小凹槽處非特異 性結合時,會發出突出的螢 光信號。
- 也可以使用其他螢光染料,如 溴化乙錠(EtBr)或吖啶橙 (Acridine Orange),但SYBR Green更適合用於其更高的信號 強度。
- SYBR Green比TaqMan Probe更受歡迎,因為它可 以提供有關每個擴增周期的 資訊以及有關TaqMan Probe無法獲得的熔化溫度 的資訊。
- 然而,與TaqMan Probe相 比,其缺點是缺乏特異性。

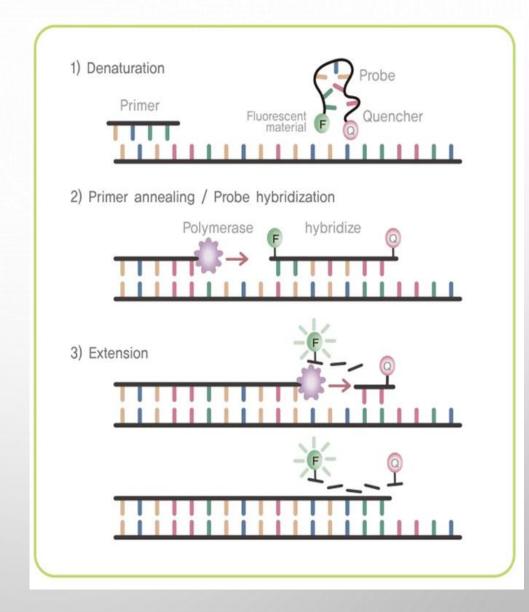


- A 'minor groove'-binding molecule specific to the minor groove of double-stranded DNA
- Fluoresces at an increased intensity when bound



(2) TaqMan Probe 探針

- 它是一種水解探針,帶有染料,通常在其5端有螢光素(FAM),並且在寡核苷酸的3端附著一個淬滅劑(non-fluorescent quencher)。
- 在正常情況下,探針保持盤繞在自身上,使螢光染料靠近淬滅劑,從 而抑制或淬滅染料的螢光信號。
- 當聚合酶在延伸階段開始合成新的 DNA鏈時,它通過5'末端核酸酶活 性導致探針降解,螢光素與淬滅劑 分離,從而產生螢光信號。
- 隨著該過程的繼續,在每個迴圈中, 信號分子的數量增加,導致螢光的 增加,這與靶標的擴增呈正相關。





TaqMan® Probe: TaqMan® MGB/NFQ Probes

- Minor Groove Binder (MGB)
 - Small molecule that fits snugly into minor groove of duplex DNA
 - Stabilizes probe annealing
- Non-fluorescent Quencher (NFQ)
 - "Dark" quencher acts as energy transfer acceptor that doesn't emit a detectable fluorescent signal
 - MGB probe design uses a special algorithm in Primer Express® Software

AGGCCTTGAGAGATAT

MGB (minor groove binder)





ThermoFish@CTACACAGTCCGGAACTCTCTATAGCATCACAC

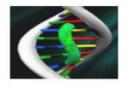


TaqMan® Probe

SYBR® Green 1 Dye

Less specific





Specificity

- Highly specific
- Probe Hybridization

Sensitivity

Very High

Flexibility

- Multiplex PCR
- SNP detection
- +/- application

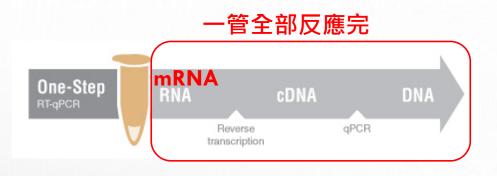
Very High

- No Probe is required
- Screening tool

Optimization

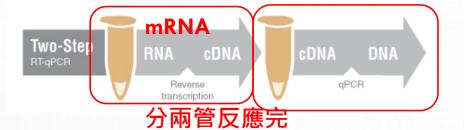
- Ready to use 20x primer/probe mix no need to optimize
- Gold standard for MAQC
- PCR efficiency 100% ±10%

- Need to optimize PCR program
- Need to check primer-dimer info
- Need to check PCR efficiency





Reaction mixture					
Master mix	Volume per one reaction				
2X SYBR Green RT-PCR reaction mix	$12.5\mu\mathrm{L}$				
Forward primer (10 μ M)	$1~\mu { m L}$				
Reverse primer $(10 \mu\text{M})$	$1~\mu { m L}$				
Nuclease-free H ₂ O	$9\mu ext{L}$				
RNA template (1 pg to 100 ng total RNA)	$1~\mu { m L}$				
iScript reverse transcriptase for one-step RT-PCR	$0.5\mu\mathrm{L}$				
Final reactions volume	25 μL				



Reagents for reverse transcription



Volume (μL)

16.9

25

1st

 Random primers
 3

 dNTP mix 10 mM
 3

 ddH2O
 36

 5X First strand buffer
 12

 DTT 0.1 M
 6

 RNase OUT 40 U/μL
 3

 RT enzyme
 1

 Total volume
 64

 Reagents for PCR
 Volume (μL)

 10X PCR buffer with MgCl2
 2.5

 dNTP mix 10 mM
 0.5

 Fwd primer 10 μM
 2

 Rev primer 10 μM
 2

 Taq DNA polimerase (5U/μL)
 0.5

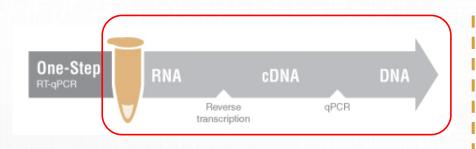
 Biotinylated dUTP
 0.6

2nd

Autoclaved ddH₂O

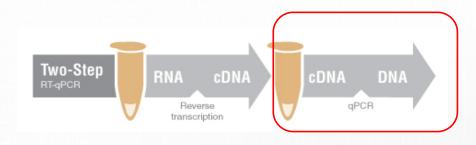
Total volume

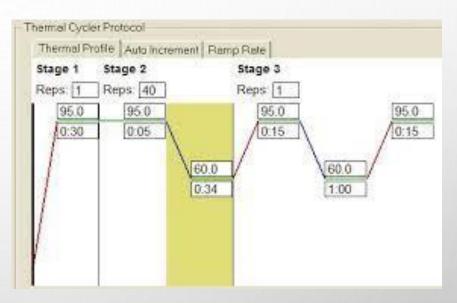
20



Protocol

Process	Duration/temperature			
cDNA synthesis	20 min at 50°C			
iScript reverse transcriptase inactivation	4 min at 95°C			
PCR cycling and detection (standard PCR cycle)				
Denaturation	10 sec at 95°C			
Annealing/extension	30 sec at 61°C (data collection step)			
Repeated for (30 to 45 cycles)				
	1 min at 95°C			
Melt curve analysis (optional)	1 min at 55°C			
	10 sec at 55°C (80 cycles) increasing each by 0.5°C each cycle)			





常用哪種儀器/系統

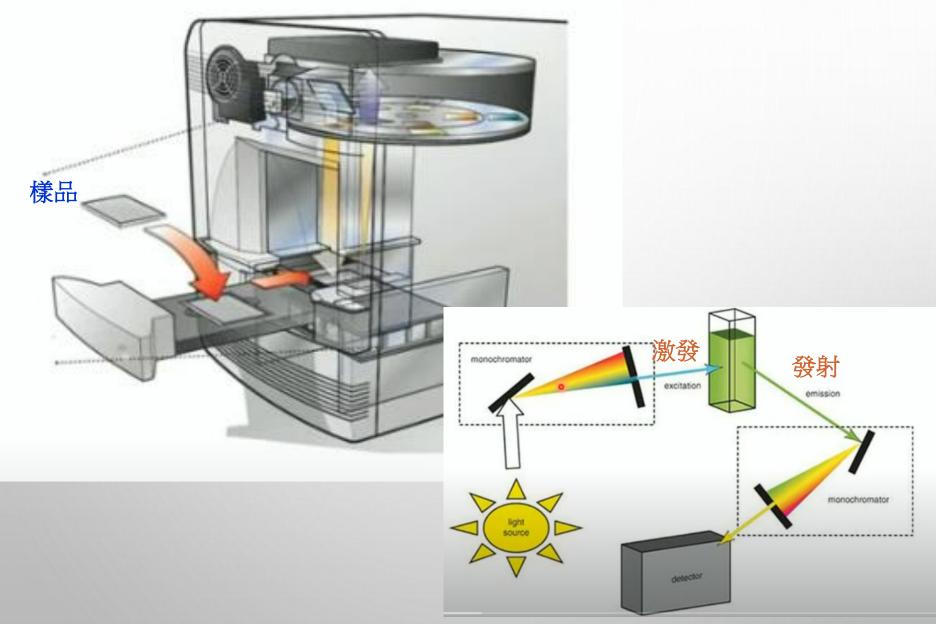
(1) ABI系統

Quantstudio systems 1,3,5,6,7





分光螢光計



實驗材料













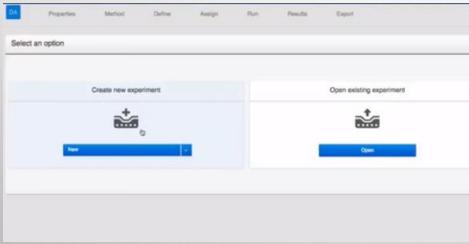


https://youtu.be/K5flNp46wxo

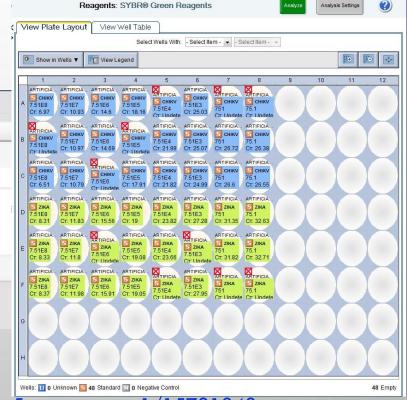
程式設置

Thermo Fisher







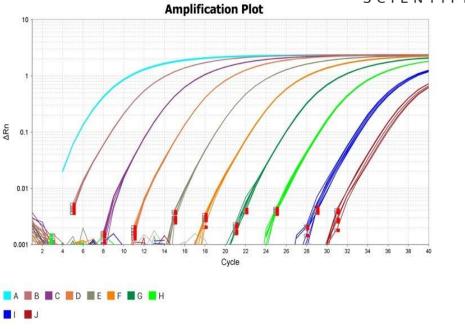


10-Log dynamic range sensitivity

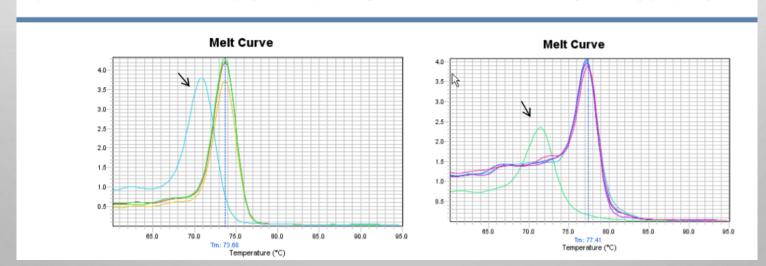




Amplification plots are displayed with the ability to drill down to a subset of sample wells.

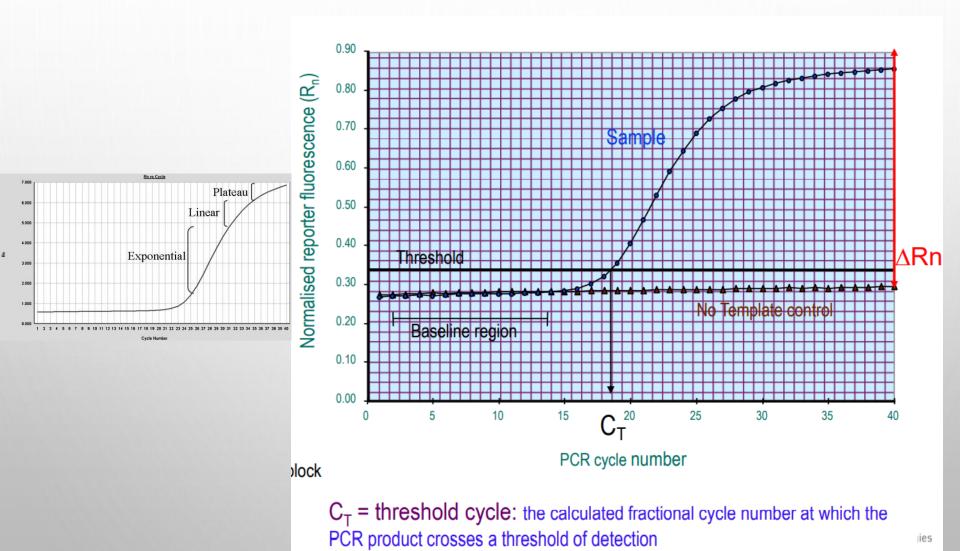


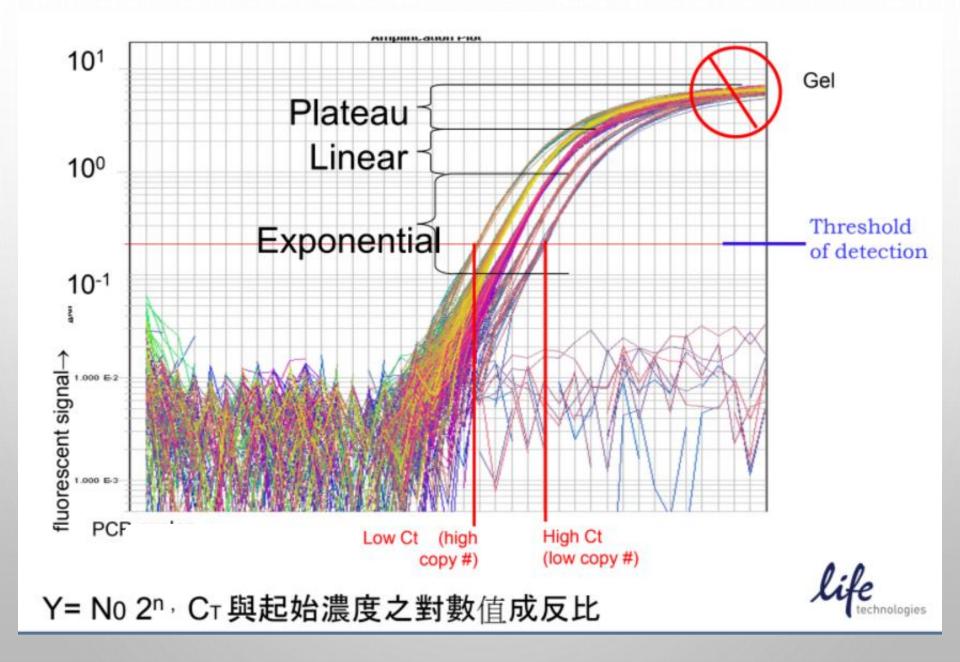
從Melt Curve確認是因為Primer Dimer或是有汙染



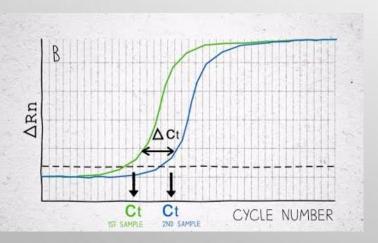
26

Cycle Threshold (Ct)





Raw Ct		Delta Ct	Delta Delta ct	2^delta d
GAPDH	p53			
21.00	23.00	2.00	-3.93	=2^-(F3)
20.50	22.00	1.50	-4.43	
20.60	22.50	1.90	-4.03	
20.00	26.00	6.00	0.07	
20.50	26.20	5.70	-0.23	
20.30	26.40	6.10	0.17	
		5.93		
	GAPDH 21.00 20.50 20.60 20.00 20.50	GAPDH p53 21.00 23.00 20.50 22.00 20.60 22.50 20.00 26.00 20.50 26.20	GAPDH p53 21.00 23.00 20.50 22.00 20.60 22.50 20.00 26.00 20.50 26.20 20.30 26.40 6.10	GAPDH p53 21.00 23.00 20.50 22.00 1.50 -4.43 20.60 22.50 1.90 -4.03 20.00 26.00 6.00 0.07 20.50 26.20 5.70 -0.23



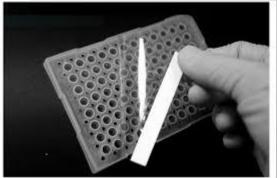
$$\begin{split} \Delta Ct &= Ct_{target} - Ct_{18SrRNA} \\ \Delta \Delta Ct &= \Delta Ct_{(siRNA\ treated)} - \Delta Ct_{(siRNA\ nontreated)} \\ Relative\ expression\ level &= 2^{-\Delta\Delta Ct} \\ \% KD &= 100\ \times\ \left(1-2^{-\Delta\Delta Ct}\right) \end{split}$$

(2) Bio-Rad系統

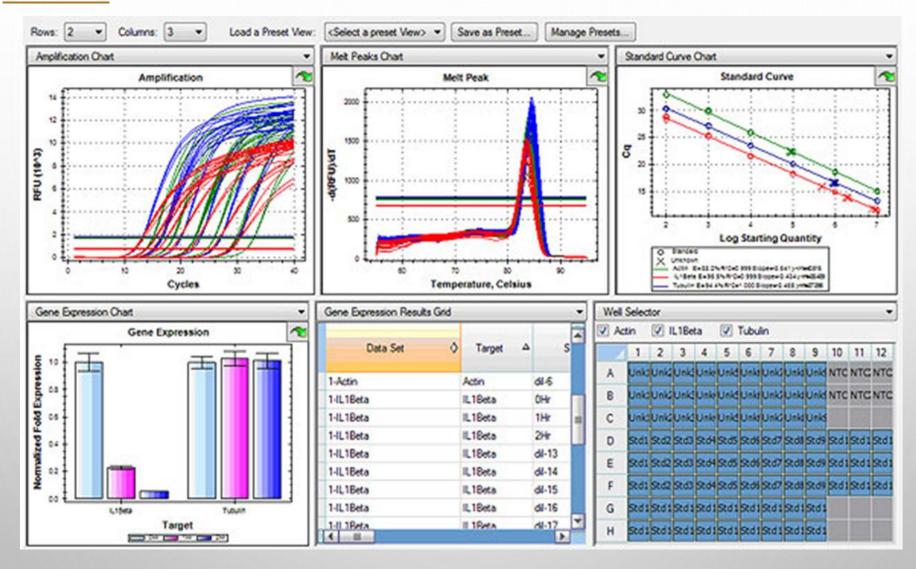








程式設置



如何避免Q-PCR錯誤



(1) 核糖核酸品質

- 是成功製備cDNA並執行高效且可重複的Q-PCR的最關鍵 步驟。
- RNA對RNA酶的降解非常敏感,在核酸分離步驟中必須小心處理。降解或污染的RNA會對Q-PCR實驗的效率和產量產生負面影響。
- 您的實驗室工作台、移液器和吸頭必須不含 RNase,提取的 RNA 必須儲存在不含 RNase 的溶液中。
- 當您使用分光光度計評估RNA純度時,260和280nm (A260/280)處的吸光度比應在1.8-2.0的範圍內。如果 較低,則可能表明樣品被苯酚或蛋白質污染。

(2) 預混液與表格使用



使用預混液可大幅減少實驗變異性,從而減少孔間和樣品間 差異來提高再現性。

在執行Q-PCR實驗時最好事先製作一張準備要加入每項試劑 的清單與樣品相對位置表格。

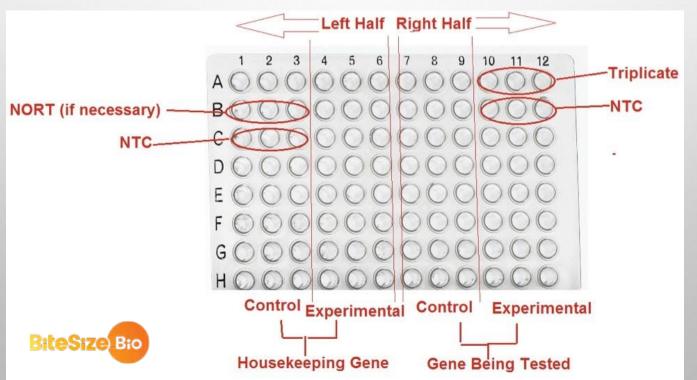


Component	Volume/ reaction
Master mix	20 μl
Nuclease free water	12 μl
5x Qiagen OneStep RT-PCR buffer	5 μl
dNTP Mix (10 mM of each dNTP)	1 μl
Primer Forward (100 pmol/ul)	0.5 μl
Primer Reverse (100 pmol/ul)	0.5 μl
Qiagen OneStep RT-PCR Enzyme Mi	x 1.0 μl
Template RNA	5 μl
Total	25 μl

(3) 控制組



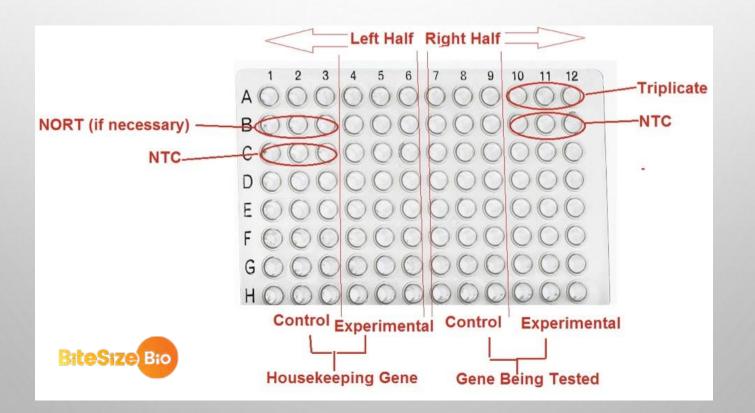
- Q-PCR測定時,適當的控制組是很重要的。
- 應該包括一個缺乏cDNA的陰性對照組,可以檢測表面或試劑的 交叉污染。
- 也需要一組反轉錄的陰性對照組,其中沒有逆轉錄酶,如果觀察到產品,表明您的樣品受到DNA污染。



(4) 參考基因

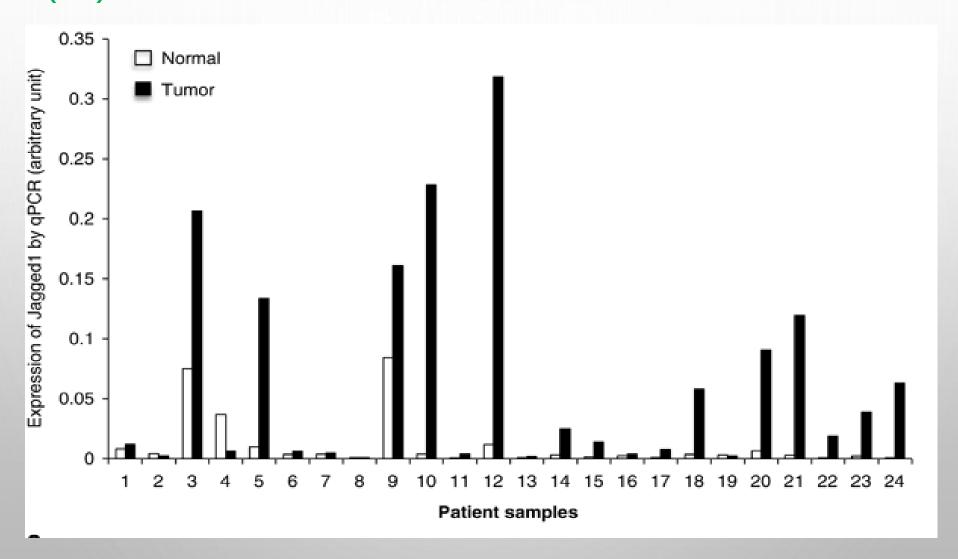


- 內源性對照基因的擴增可以解釋起始cDNA的數量或質量的差異,以及RNA製備方法或cDNA合成的差異。
- 可靠的參考基因是指其表達量不受實驗變數的影響,並且 在樣品條件的相關生理狀態之間沒有差異的基因。

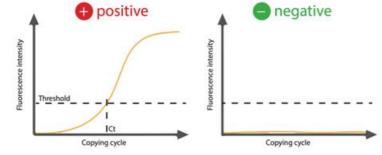


Q-PCR 應用 - Gene expression Disease diagnosis More Specificity Cancer genetics Infectious studies microRNA analysis More Sensitivity Microbial load testing Read time Low template **PCR** Genotyping cfDNA detection Cost effective SNP analysis Rapid Microarray validation

(1) 基因表達分析-例如病人正常細胞與癌細胞基因表現差異



(2)微生物檢測量 a) sample collection b) RNA extraction c) Reverse transcription d) RT-PCR amplification DNA polymerase primer probe e) Results positive negative





一次認識三種檢驗方式

健康醫療網





抗原檢測 抗原快篩)

♥♥ PCR檢測 【(RT-PCR)

高風險地區,快速 找出感染者的利器

檢測

檢體中是否含有 病毒的抗原

優點

檢驗時間短 快速得到結果

缺 準確率較PCR低,容點 易產生偽陽、偽陰性

資料來源:疾管署

全球判斷染疫的 標準檢測

檢測

檢體中是否含有 病毒的遺傳物質

慢點

準確度高,病毒 量低也可檢驗出

缺 耗時、成本高,需要點 專業設備及人員執行



抗體檢測 (抗體快篩)

後續瞭解病毒的 盛行率、研究用

檢測

血清中是否含有 病毒的抗體

慢點

可找出曾經感染過或 打疫苗者是否有抗體

缺 感染後期才能驗出,也 點 可能會有偽陰性產生

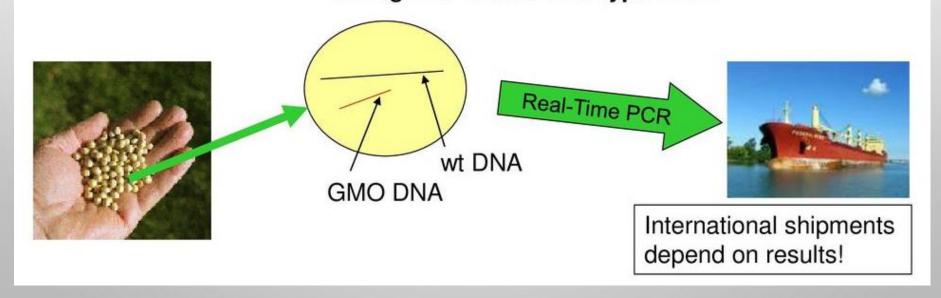
CONS MADE BY FLATICON

(3)基因食品檢測

Example: Determining percentage of GMO food content

Determination of percent GMO food content important for import / export regulations.

Labs use Real-Time PCR to measure amount of transgenic versus wild-type DNA.







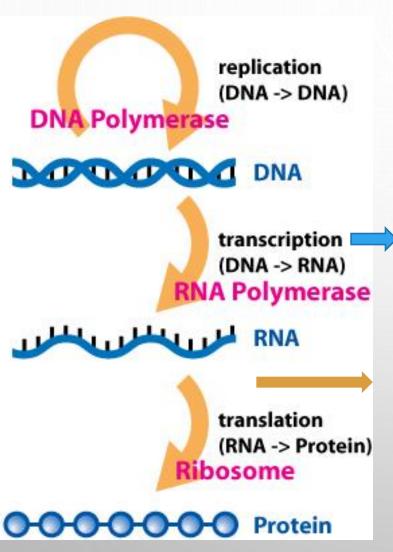
II. 報告基因分析 (Reporter gene assay)

上課內容

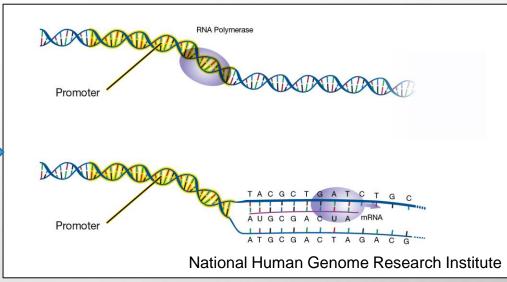
- 基因表現調節
- 什麼是報告基因分析
- 常見的報告基因
- 報告基因分析的應用

基因表現調節

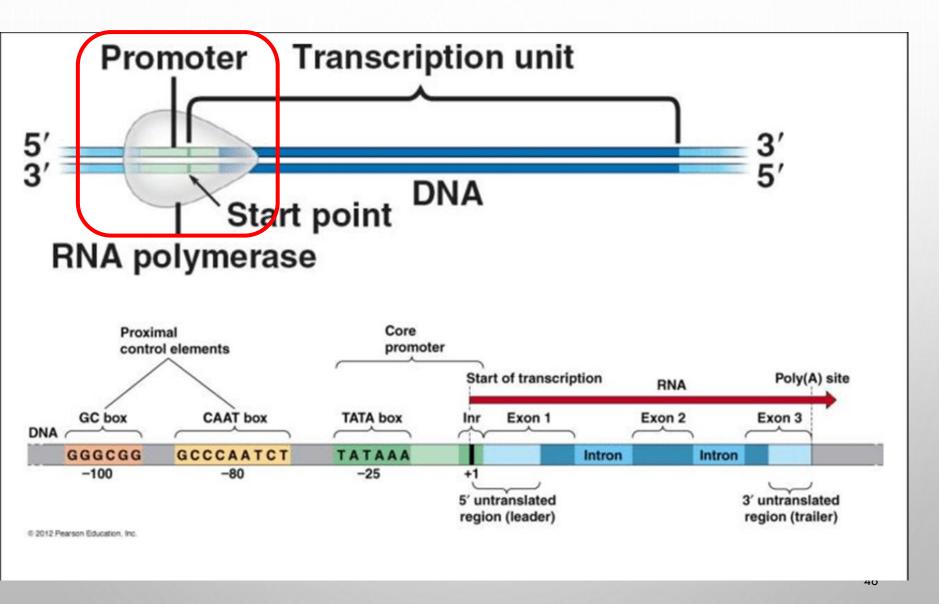
中心法則



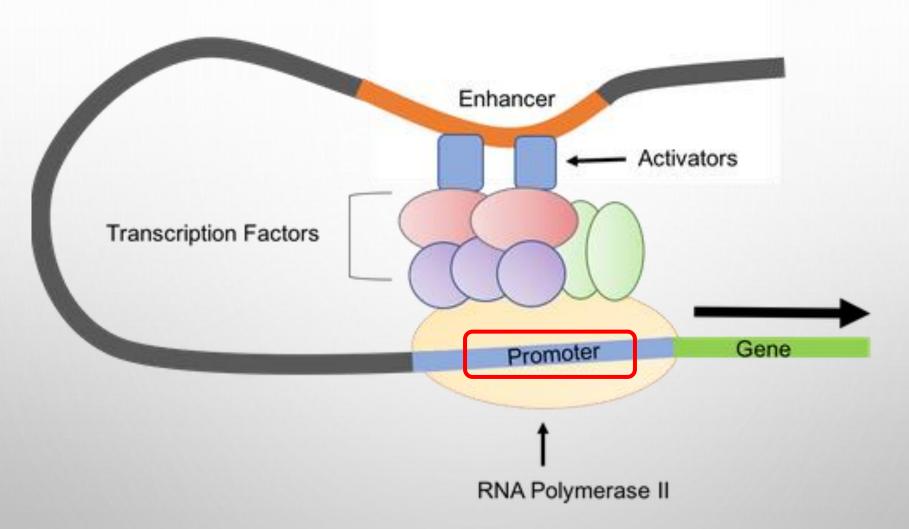
DNA > RNA



啟動子 (Promoter)



→ addgene

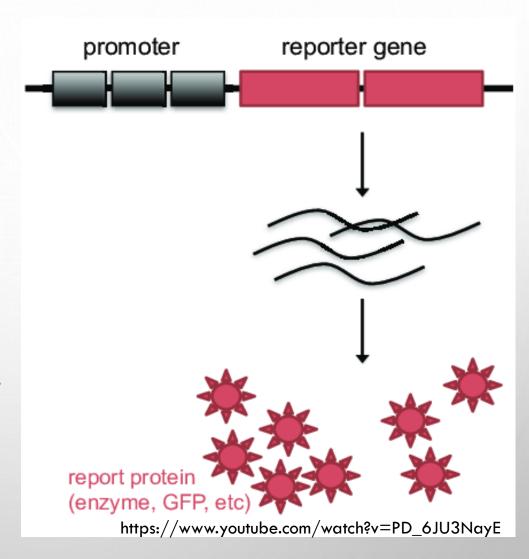


什麼是報告基因分析

報告基因分析主要為報告基因 的啟動子活性的測量。

 報告基因是把啟動子DNA序列 和報告基因DNA序列的融合。

• 報告基因可以為酵素或螢光物。





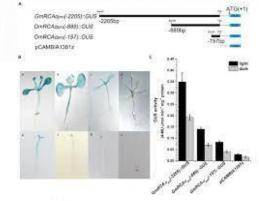
如何才是好的報告基因?

- 1. 易於檢測其功能。
- 2. 其表達產物能進行定量測定。
- 3. 不會影響實驗材料本身。
- 4. 表達產物在受體細胞中原本就不存在。

常見的報告基因

- β-葡萄糖醛酸酶 (β-glucuronidase, GUS)
- β-半乳糖苷酶(β-galactosidase, LacZ)
- 鹼性磷酸酶 (alkaline phosphatase, ALP)
- 氯黴素乙醯轉移酶 (Chloramphenicol acetyltransferase, CAT)
- 綠色螢光蛋白 (Green fluorescent protein, GFP)
- 螢光素酶 (luciferase)

(1)β-葡萄糖醛酸酶(GUS)



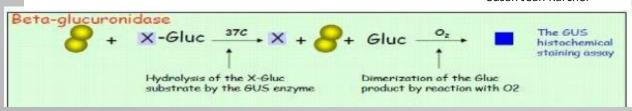
B r o-glucuronic acid

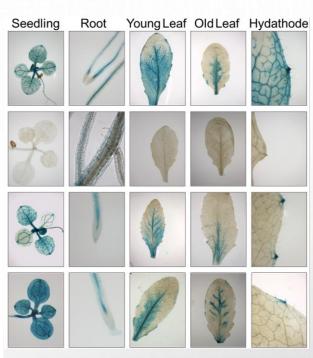
X-Gluc (5-bromo-4-chloro-3-indolyl-β-D-glucuronide)

oxidation dimerization

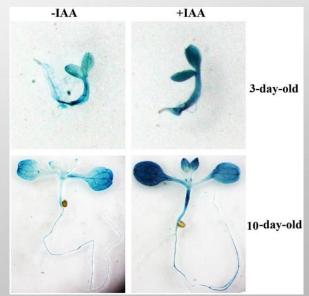
+ glucuronic acid

Susan Jean Karcher



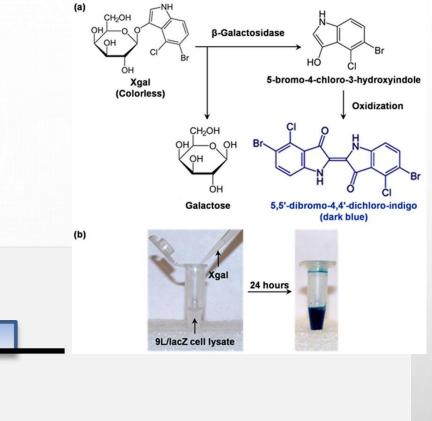


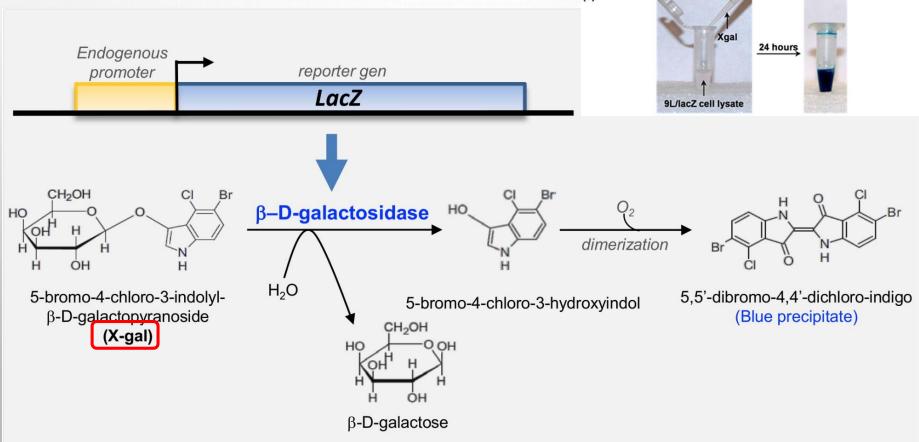
doi: https://doi.org/10.1371/journal.pone.0159875.g004



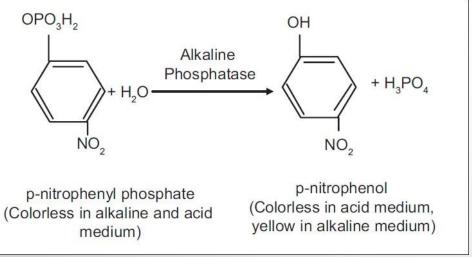
Frontiers in Plant Science 6(295) · May 2015

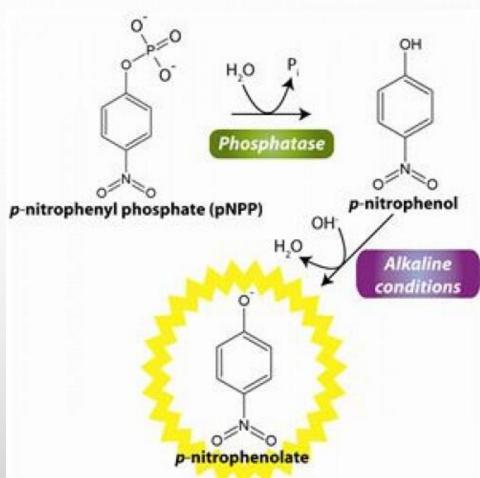
(2) β-半乳糖苷酶(LacZ)



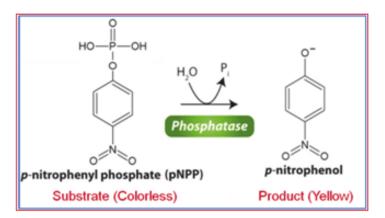


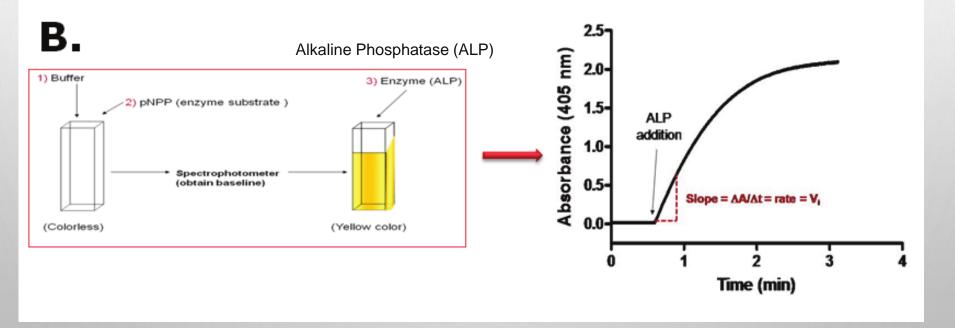
(3) 鹼性磷酸酶 (ALP)





Α.





(4) 氯黴素乙醯轉移酶(CAT)

CAT: Chloramphenicol (CAM) acetyl transferase

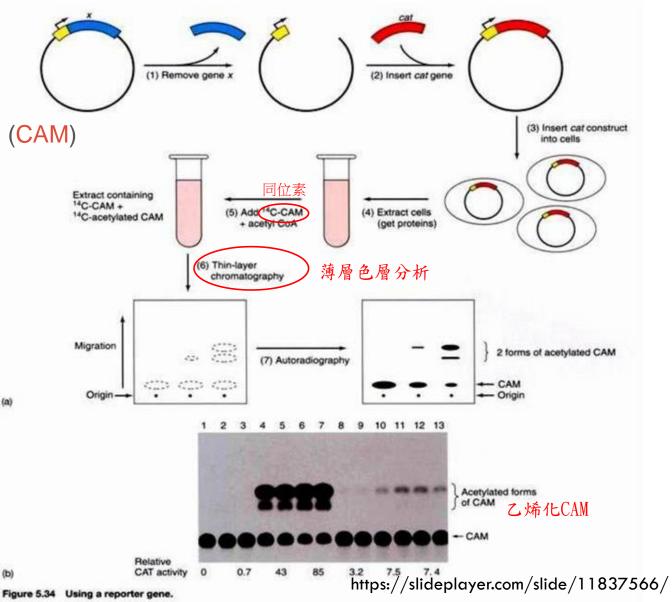
CAM: chloramphenicol (CAM)

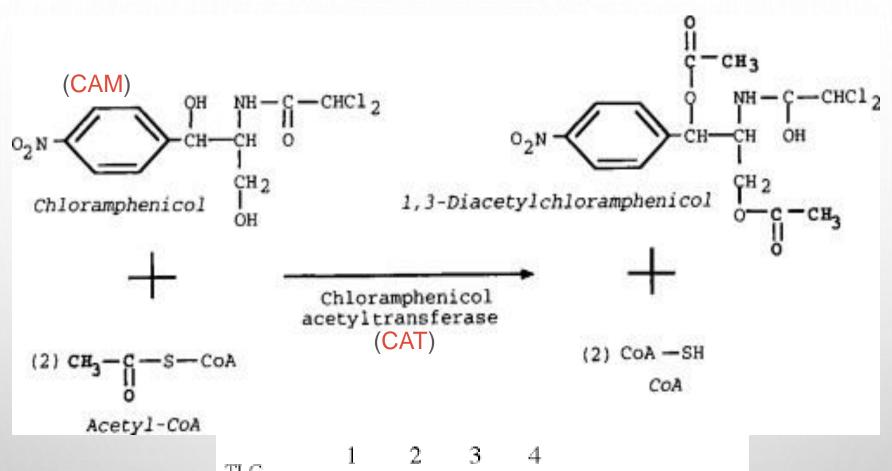
Protein synthesis inhibitor

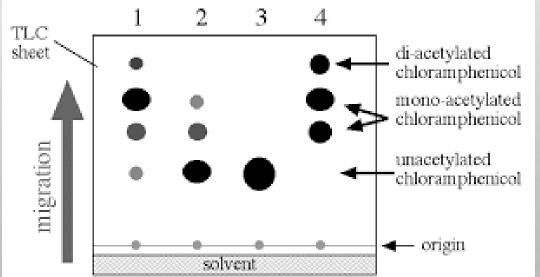
↓ acetylation by CAT

Loss inhibitor activity

Other reporter enzymes: β-galatosidase luciferase







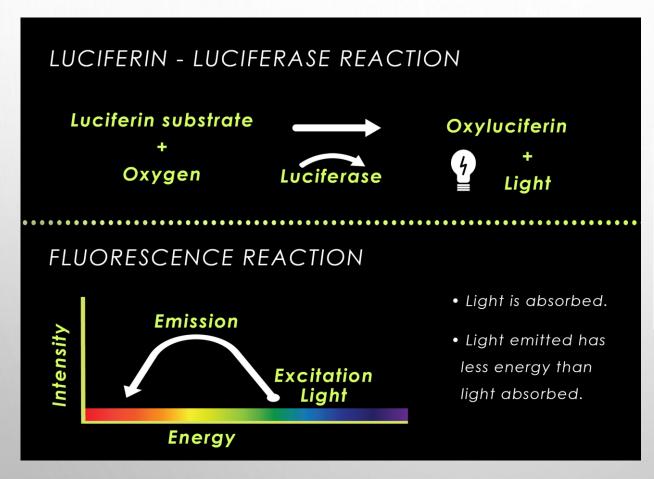
冷光與螢光反應

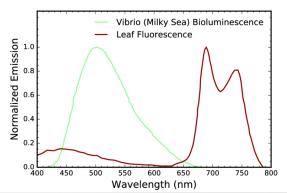
從微孔盤底部會不斷產生<u>冷光</u>自發光,這時後選擇**反光 效果較佳的白盤** 可有效檢測樣品的冷光訊號值



微孔盤底部會會有一道散射光, 為了有效阻隔鄰近樣品間的干擾, 螢光實驗建議選用**遮蔽力較佳的黑盤**來進行

https://blog.xuite.net/clare690725/twblog/111356956





DISCOVER BIOLUMINESCENT ORGANISMS



Renilla reniformis

Subsrate: Coelenterazine

Wavelength: 480 nm

Cypridina noctiluca

Subsrate: Vargulin Wavelength: 465 nm

Gaussia princeps

Subsrate: Coelenterazine

Wavelength: 460 nm

Photinus pyralis

Subsrate: D-Luciferin

Wavelength: 560 nm

Pyrophorus

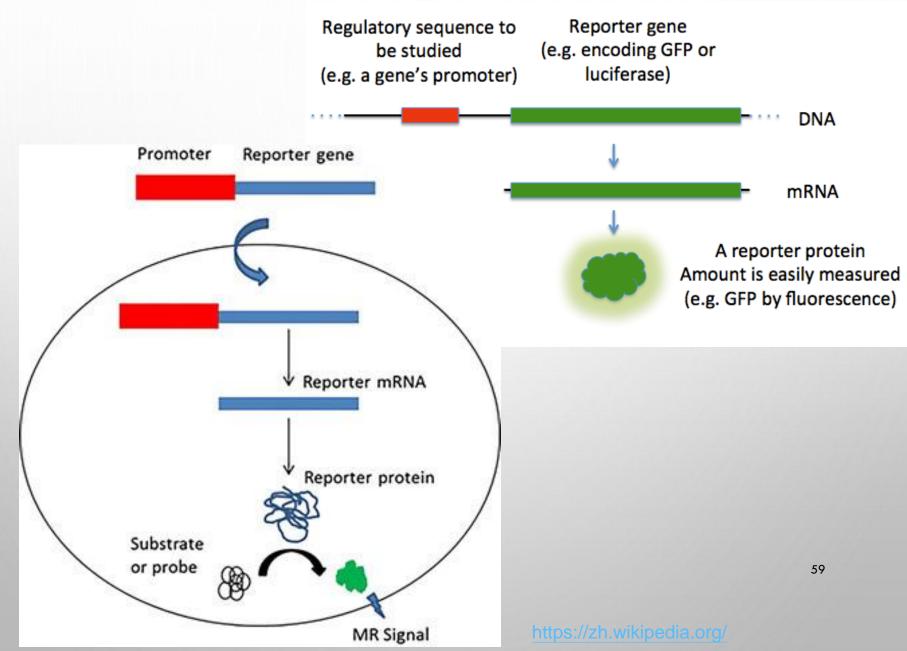
plagiophthalamus

Subsrate: D-Luciferin

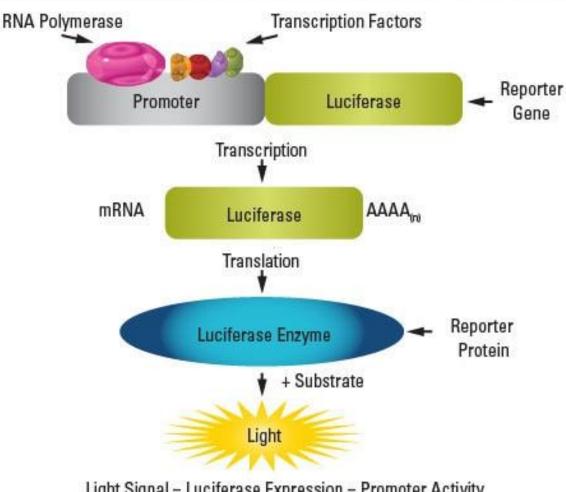
Wavelength: 613 nm (red)

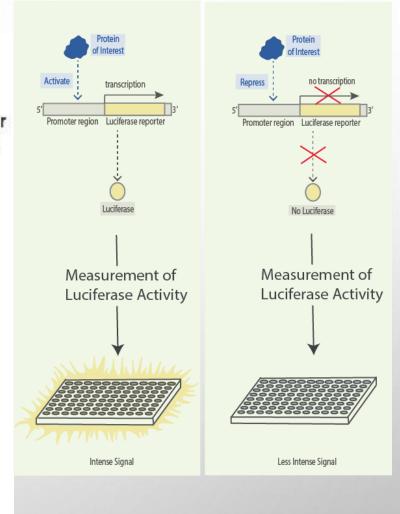
400 500 600

(5)綠色螢光蛋白(GFP)



(6)冷光素酶 (Luciferase)





Light Signal = Luciferase Expression = Promoter Activity

SCIENTIFIC



The Nobel Prize in Chemistry 2008

"for the discovery and development of the green fluorescent protein, GFP"

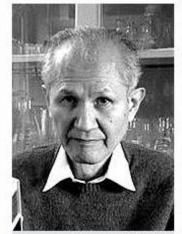


Photo: J. Henriksson/SCANPIX

Osamu Shimomura

O 1/3 of the prize

USA

Marine Biological Laboratory (MBL) Woods Hole, MA, USA; Boston University Medical School Massachusetts, MA, USA



Photo: J. Henriksson/SCANPIX

Martin Chalfie

O 1/3 of the prize

USA

Columbia University New York, NY, USA



Photo: UCSD

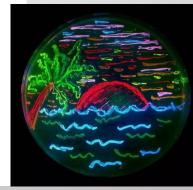
Roger Y. Tsien

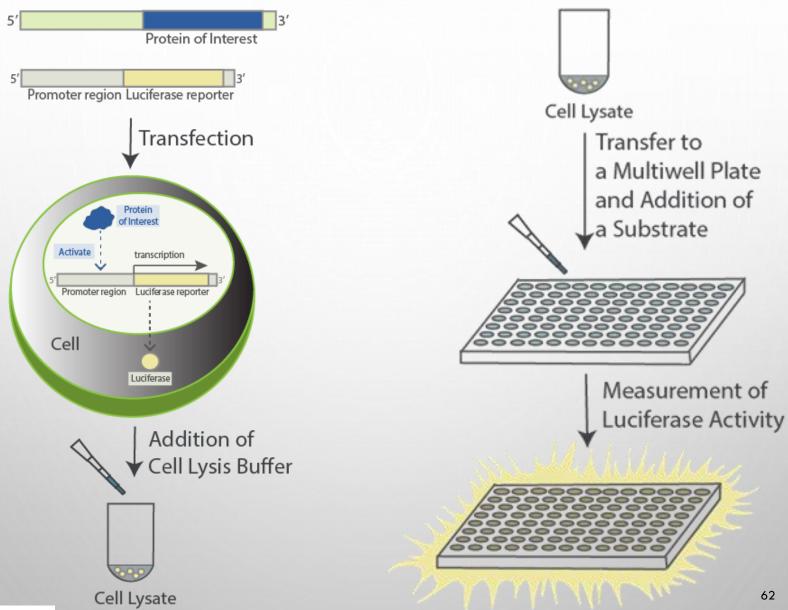
O 1/3 of the prize

USA

University of California San Diego, CA, USA; Howard Hughes Medical Institute



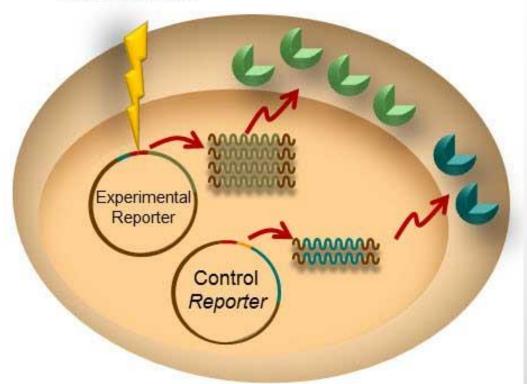


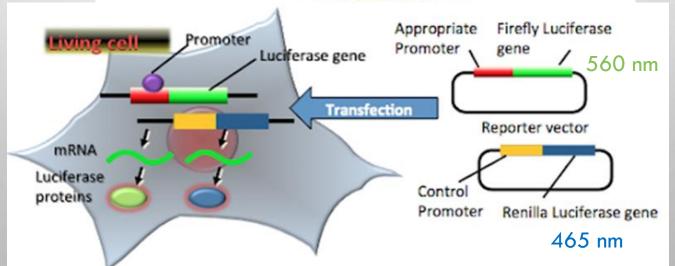




雙報告書檢測

TREATMENT

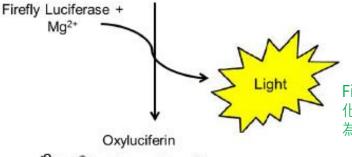






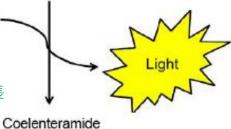


Renilla luciferase作用為校正轉染效率的內參,以消除孔間細胞數量和轉染效率的差異。

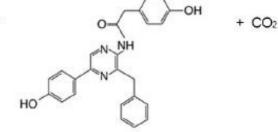


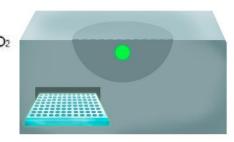
Renilla Luciferase -

Firefly luciferase催 化Luciferin發光波長 為560 nm。



Renilla luciferase催化 Coelenterazine發光波 長為465 nm





Read results on any luminometer



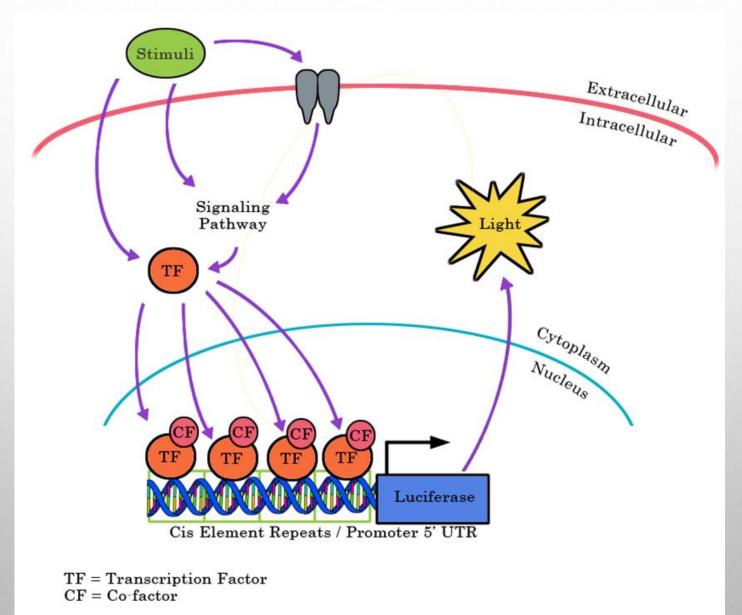
cell extracts Firefly luciferase Renilla luciferase

Promoter activity = Firefly luciferase activity / Renilla luciferase activity

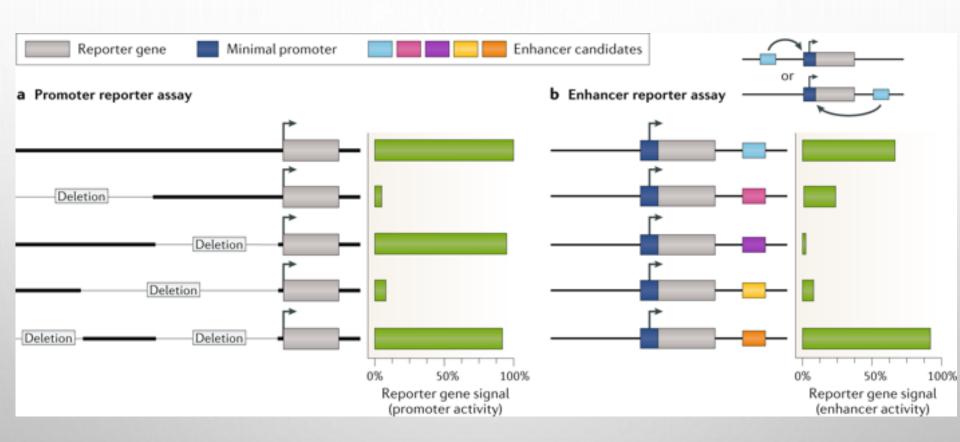
報告基因分析的應用

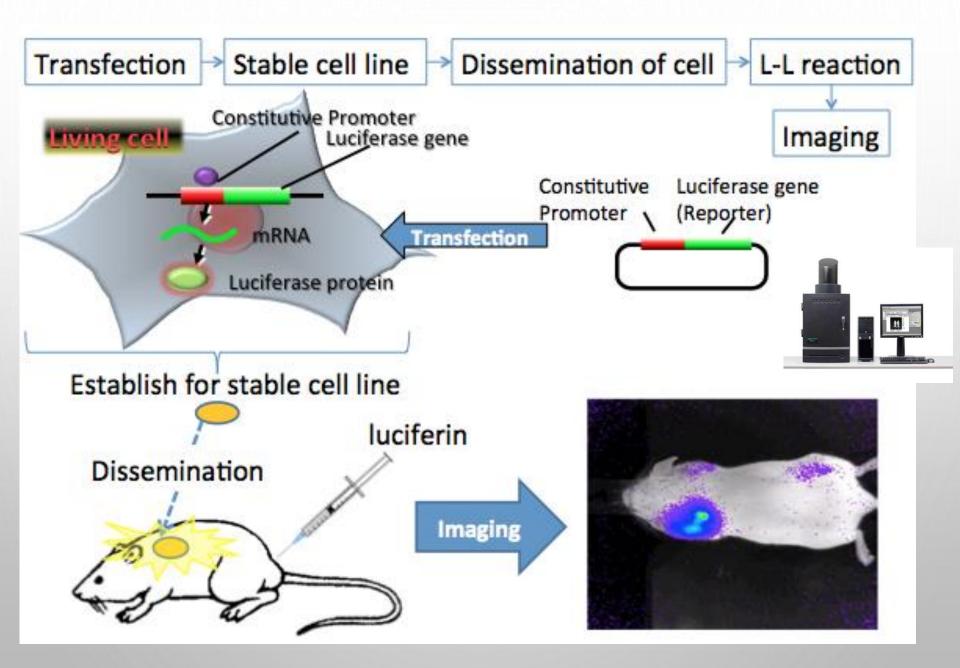
- 分析基因調控
- 分析miRNA靶標
- 分析核受體信號傳導
- 研究蛋白質相互作用
- 高通量篩選

(1)分析基因調控



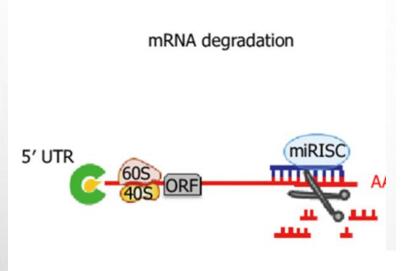




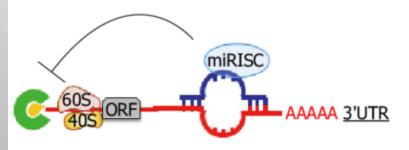


(2) 分析MiRNA靶標

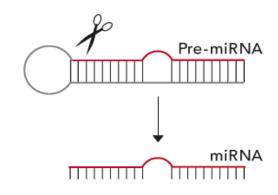
MicroRNA (miRNA) 是短RNA, 與轉錄本 3¹UTR中的靶標相互作用,導致mRNA降解或 抑制翻譯。



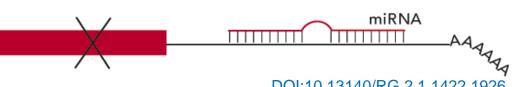
Translation inhibition



DOI:10.4331/wjbc.v8.i1.45

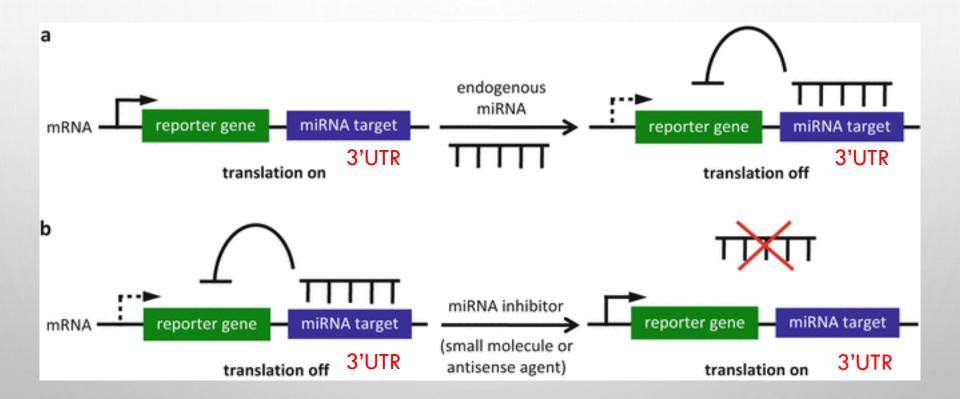


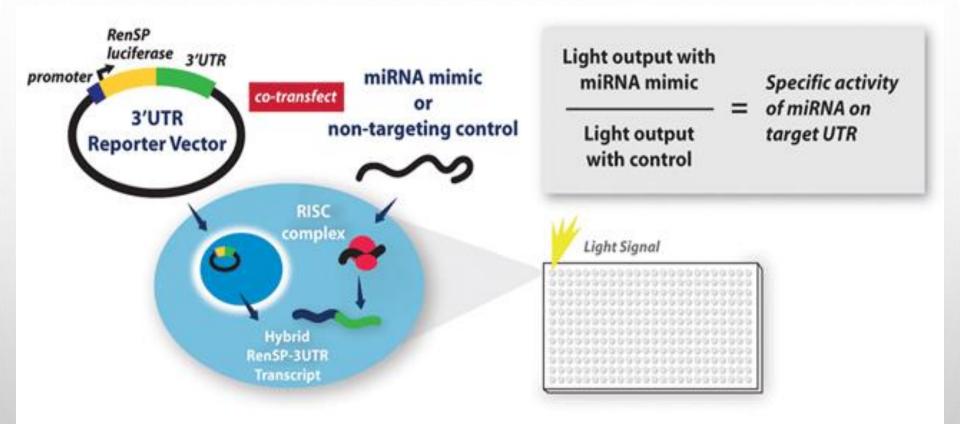




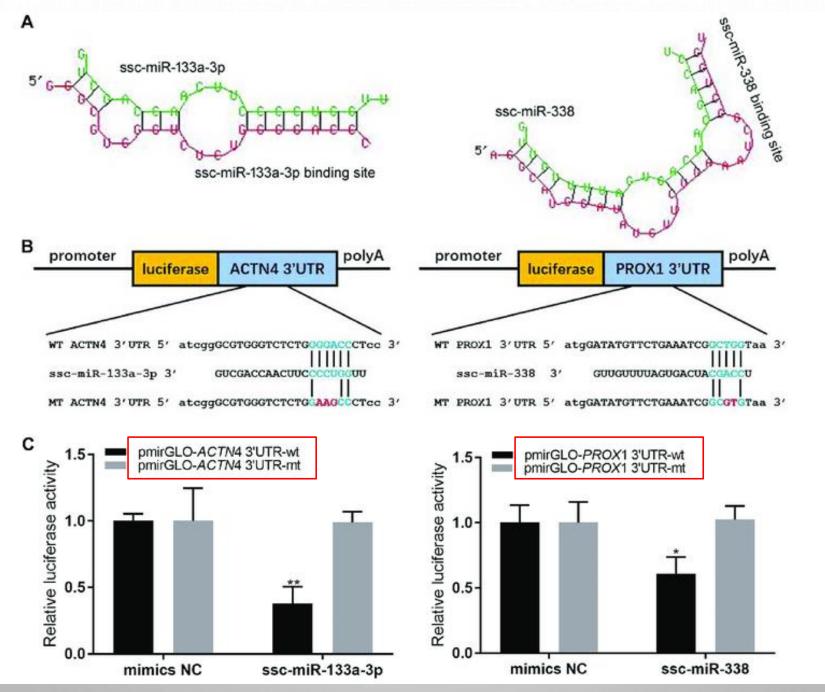
DOI:10.13140/RG.2.1.1422.1926

觀察MiRNA介導的效應需要一個在較弱的啟動子的控制下的報告者,以便可以觀察到基因表達的細微變化。

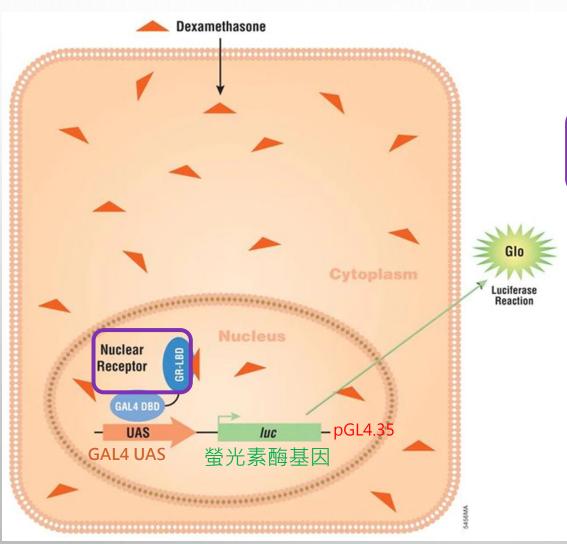


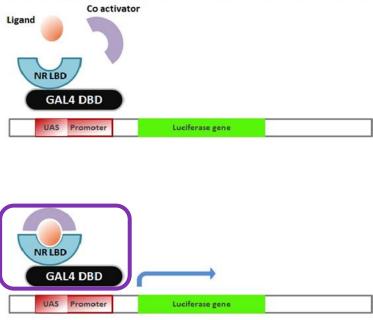


- 1. Transfect the GoClone 3'UTR construct into your cell line
- 2. A constitutive promoter drives production of a hybrid RenSP-human 3'UTR transcript
- 3. Total luciferase output depends on the effect of the human 3'UTR on the hybrid transcripts stability and/or translation efficiency
- 4. Include a co-transfected synthetic miRNA or miRNA inhibitor to measure its impact on transcript stability and/or translation efficiency
- 5. Use optimized LightSwitch Luciferase Reagents to obtain maximun sensitivity and dynamic range



(3)分析核受體信號傳導





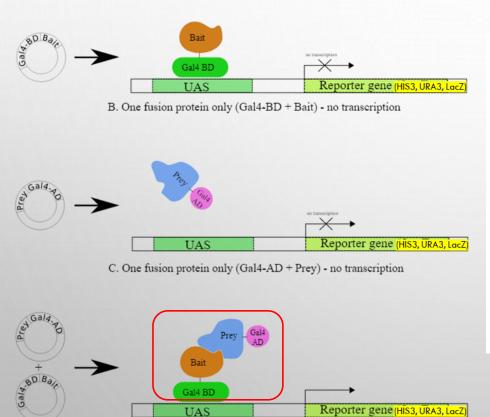
- 關鍵是一個優化的PGL4載體,其中包含9個重複的GAL4 UAS, 一個最小的啟動子和LUC2P螢光 素酶基因(PGL4.35)。
- 製備兩個含有雌激素受體配體結 合域(PBIND-ER∝)或糖皮質激 素受體配體結合域(PBIND-GR) 的PFN26A載體。
- 用載體、培養和檢測轉染細胞。



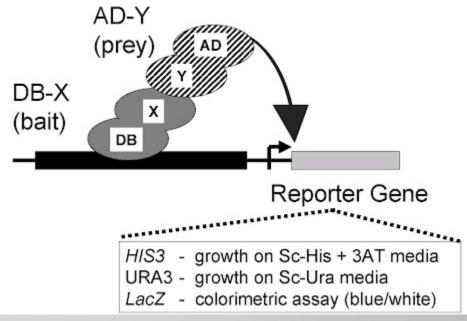
(4)研究蛋白質相互作用



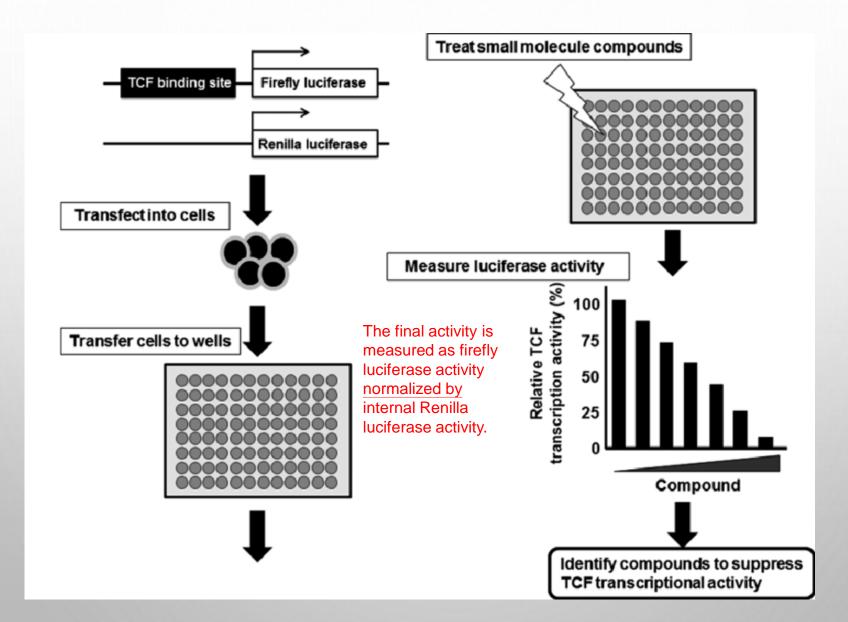
A. Regular transcription of the reporter gene



D. Two fusion proteins with interacting Bait and Prey



(5)高通量篩選







III. 核糖核酸干擾 (RNAi)

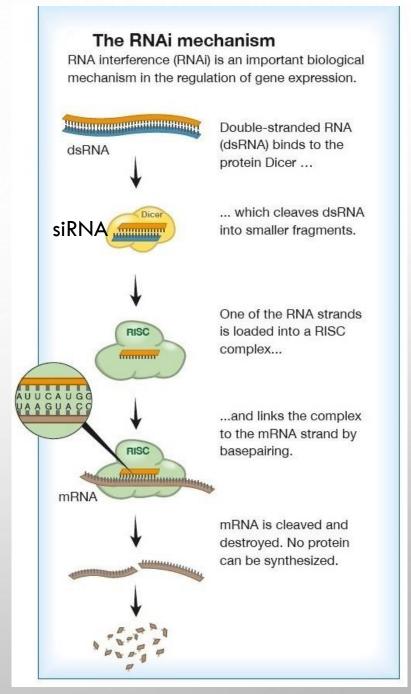
上課內容

- RNAi原理
- 誰發現了RNAi
- RNAi技術
- RNAi的應用

RNAi原理

- 雙鏈RNA(dsRNA)由RNase III家族 成員<u>Dicer</u>處理,產生21-23nt小干擾 RNA。
- siRNA由稱為RNA誘導的沉默複合物 (RISC)的多組分核酸酶操縱。

siRNA: small interfering RNA



誰發現了RNAi

2006 Nobel Prize in Medicine or Physiology RNA Interference

NATURE | VOL 391 | 19 FEBRUARY 1998

Potent and specific genetic interference by double-stranded RNA in Caenorhabditis elegans

Andrew Fire*, SiQun Xu*, Mary K. Montgomery*, Steven A. Kostas*†, Samuel E. Driver‡ & Craig C. Mello‡

A naturally occurring mechanism

Destruction of mRNA results in the post transcriptional inhibition of gene expression and the prevention of protein synthesis.

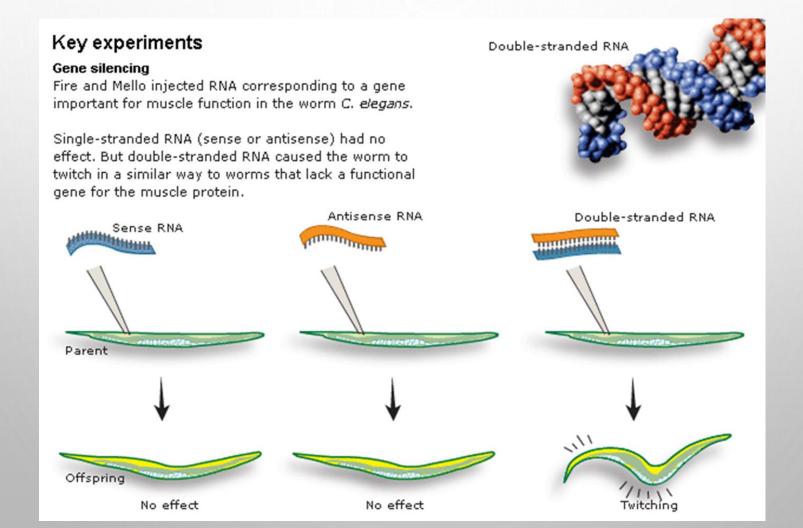
克雷格·梅洛

安德魯·菲爾



動物模型-線蟲

- RNAi最初被發現於秀麗隱桿線蟲 (*Caenorhabditis elegans, C. elegans*)。
- 觀察到雙鏈RNA (dsRNA) 在沉默標靶基因表達方面比單獨使用反股 RNA (anti-sense RNA)有效10倍。

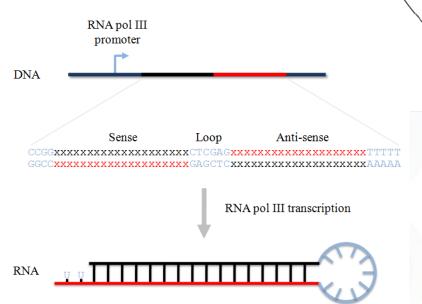


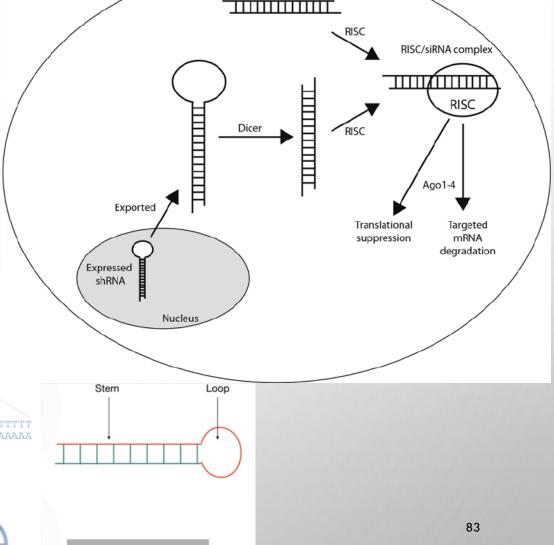


小髮夾RNA (short hairpin RNA, shRNA)

19-29 個核甘酸

短髮夾 RNA (shRNA) 包含一個環結構,該環結構被加工成 siRNA,並且還導致 mRNA 以依賴於靶 mRNA 的互補結合的序列特異性方式降解。

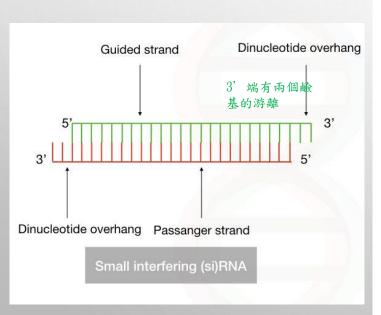


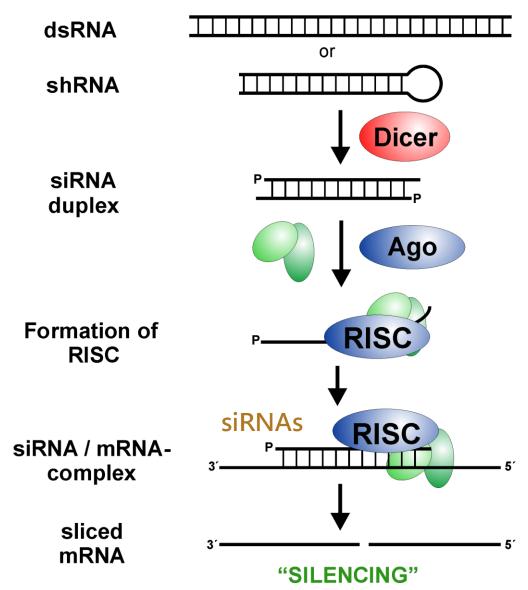


Transfected siRNA

小干擾 RNA (small interfering RNAs, siRNAs)

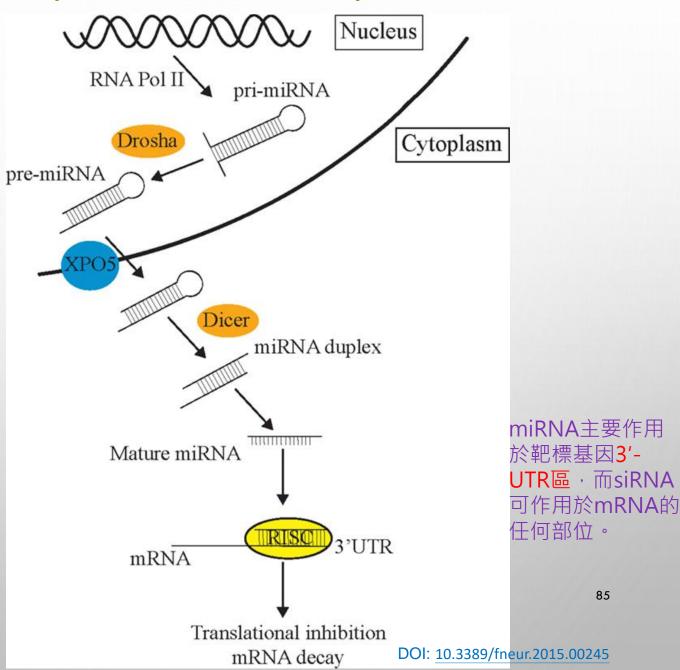
小干擾 RNA (siRNA)。 具有 20-25 個核甘酸, 3'末端突出的 siRNA 可激活 RNAi,導致 mRNA 以依賴於靶 mRNA 的互補結合的序列特異性方式降解。





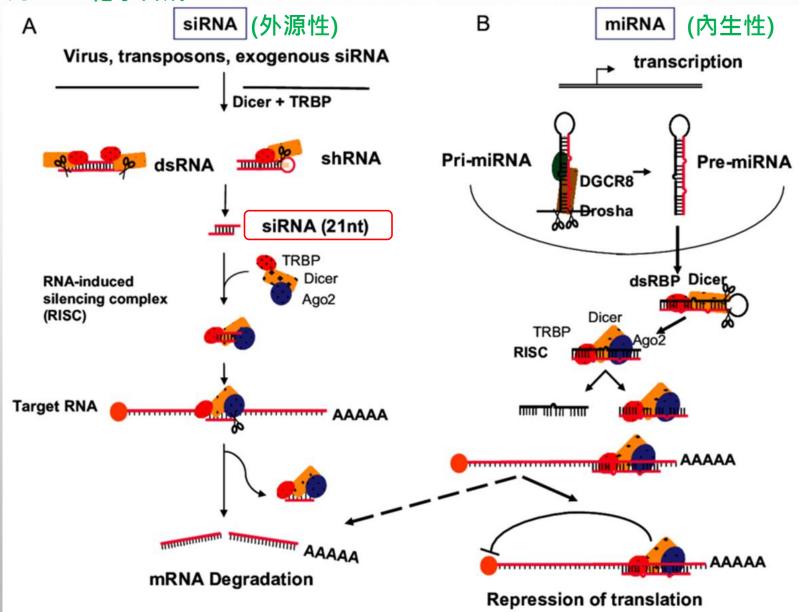
小分子核糖核酸 (microRNA, miRNA)

小分子核糖核酸 (miRNA),是真核 生物中廣泛存在的一 種長約19-25個核苷 酸的RNA分子,可調 節其他基因的表現

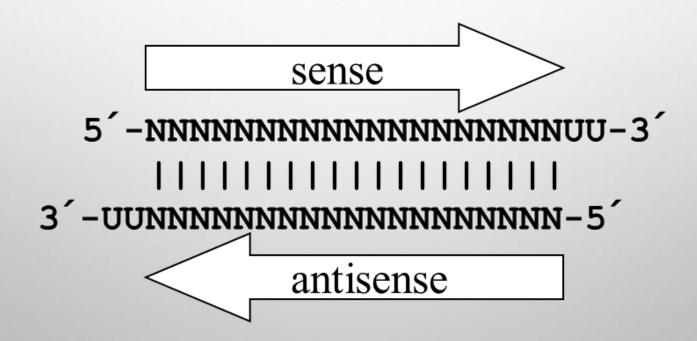


RNAi相關技術

1. siRNA化學合成

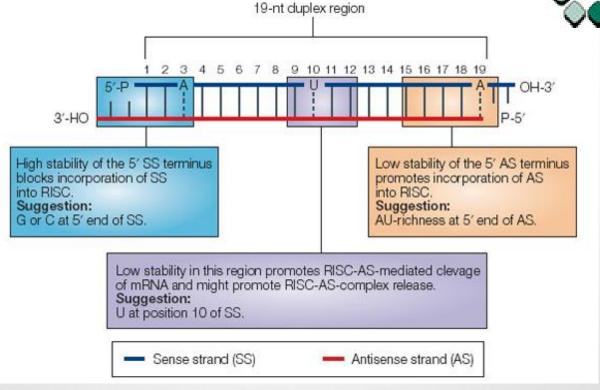


- 靶向任何序列的合成siRNA可以通過化學合成製備
- 在哺乳動物細胞中, siRNA在敲低靶基因表達方面的有效性範圍(50-95%)
- siRNA的有效性取決於標靶序列



siRNA 設計



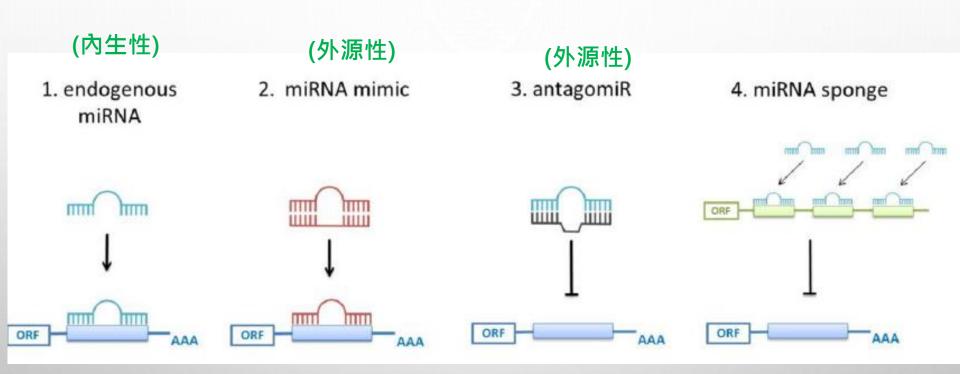


- 1.低至中等GC含量(30-50%)。
- 2. 無內部重複或回文。
- 3. 在感應鏈的位置 3 處存在 A。
- 4. 在感應鏈的位置 19 處存在 A。
- 5. 在感測鏈的位置 **19** 處沒有 **G** 或 **C**。
- 6. 在感應線的位置 **10** 處存在 U。
- 7. 感應鏈位置 **13** 處沒有 **G**。
- 8. 在感應鏈的 15-19 位置至少有 3 個 A/U。

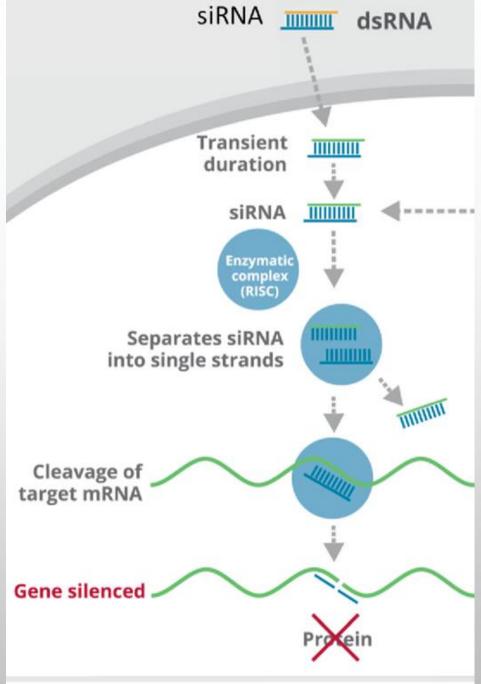
Table 2. Names and addresses of siRNA design computational soft wares

Soft wares	Address	
siDirect	http://genomics.jp/sidirect/index.php?type=fc	
siDESIGN Center	dharmacon.gelifesciences.com/design-center	
siRNA Design Software	http://www.genscript.com/ssl-bin/app/rnai	
Block-iT RNAi Designer	https://rnaidesigner.invitrogen.com	
siRNA Target Finder	http://www.ambion.com/techlib/misc/siRNA_finder.html	
RNAi explorer	www.genelink.com/sirna/siRNAorder.asp	

靶向miRNA活性的不同方法

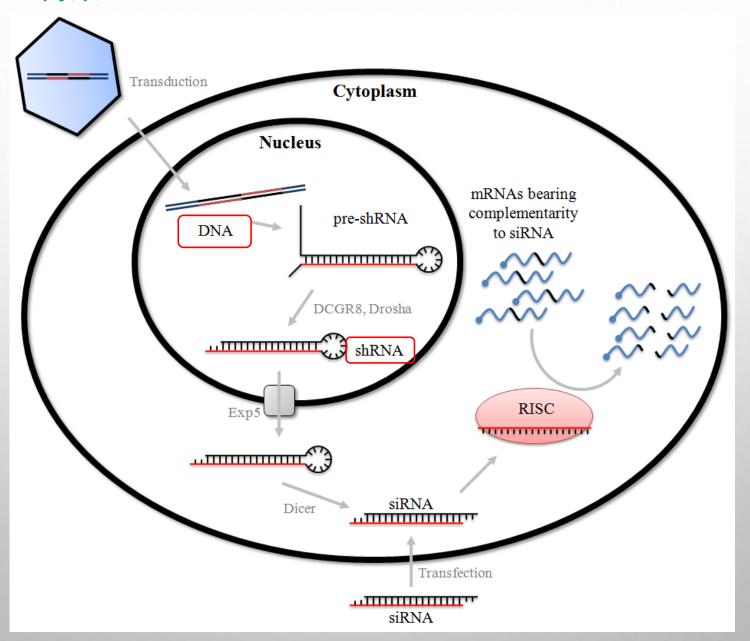


2. siRNA轉染



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3. shRNA轉染





關於 RNAi▼ | 公告訊息▼ | 服務說明▼ | C6 RNAi Citations▼ | 資料庫 | 檔案下載▼ | 軟體工具▼ | 常見問題▼ |



線上回報 knockdown feedback information



Login

免費獲贈 shRNA glycerol stock



C6-1 shRNA質體菌株分讓

⇒訂購control shRNA質體菌株

C6-2 盤式VSV-G pseudotyped lentivirus

C6-3 混合型VSV-G pseudotyped lentivirus

C6-4 單株VSV-G pseudotyped lentivirus

C6-9 客製化VSV-G pseudotyped lentivirus

C6-10 Genome-wide pooled lentivirus

C6-13 shRNA細菌株到病毒或DNA製備服務

Bacteria





Virus

會員登入 Password Account

◆服務項目清單及收費準則

帳號申請

核心公告 000

C6-22 AAV 客製化服務說明+new+

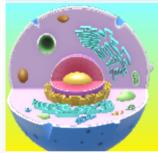
2019/11/26

108年12月停止取件服務

2019/11/1

RC 3女 T古 口 ≐T R世

93



新服務其他客製化服務

服務項目訂購(核心服務項目&收費準則查詢) Login 會員登入: 帳號 密碼 申請會員 C6-1 C6-2 C6-5 C6-6 C6-7 C6-8 C6-10 C6-11 C6-12 C6-13 C6-14 C6-9 C6-15 C6-16 C6-17 C6-18 C6-21 shRNA 質體細菌株分讓(點選可展開) [產品腦介] [使用須知] [技術支援] shRNA 關鍵字查詢 INeedControlshRNA Search

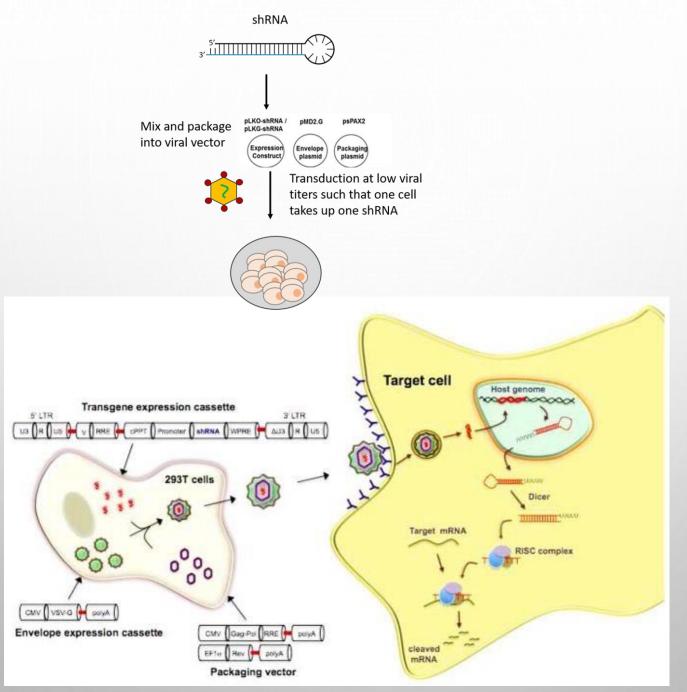
可利用 Cloneld / Symbol / targetSeq 作查詢

Search Entrez Gene to find official symbols.



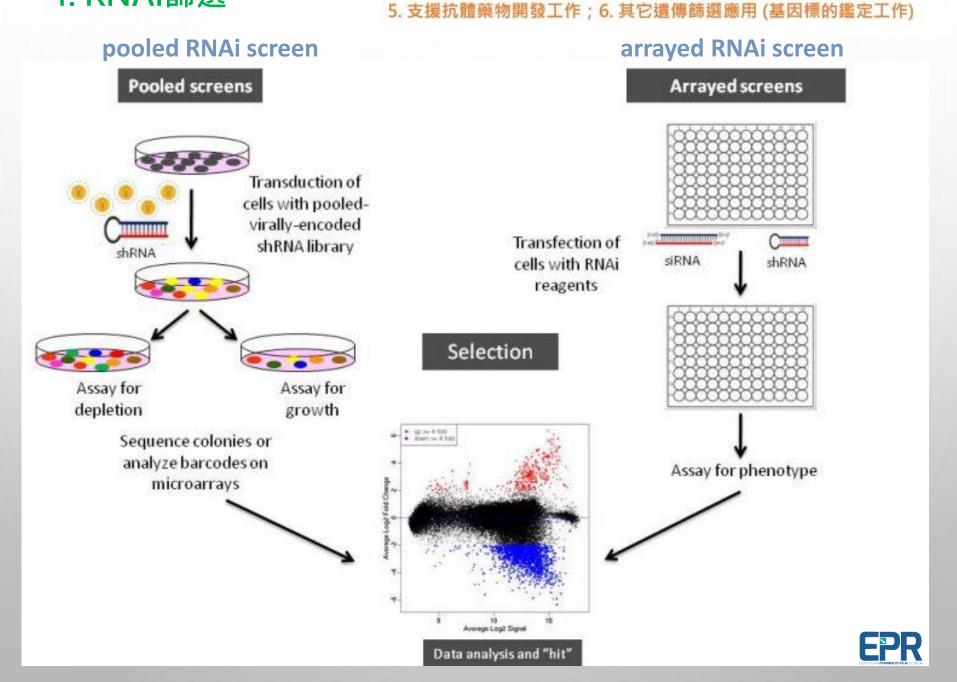
訂單取貨方式	收單(繳費完成)時間	出貨時間說明
核心取貨	※ 每週一及三早上9:00前	收單當日製作·3-4個工作天後取貨。
宅配寄送	※ 每週一及三早上9:00前	收單當日製作,每週四及隔週一出貨,預計5-8個工作天到貨。

Showing 1 to 10 of 10



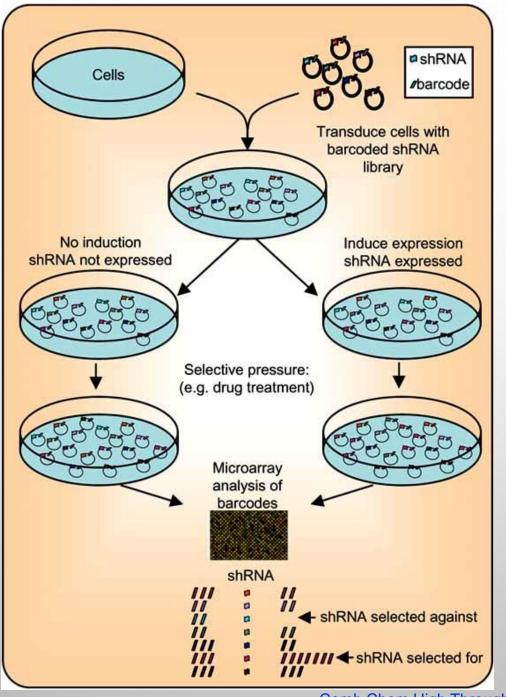
4. RNAi篩選

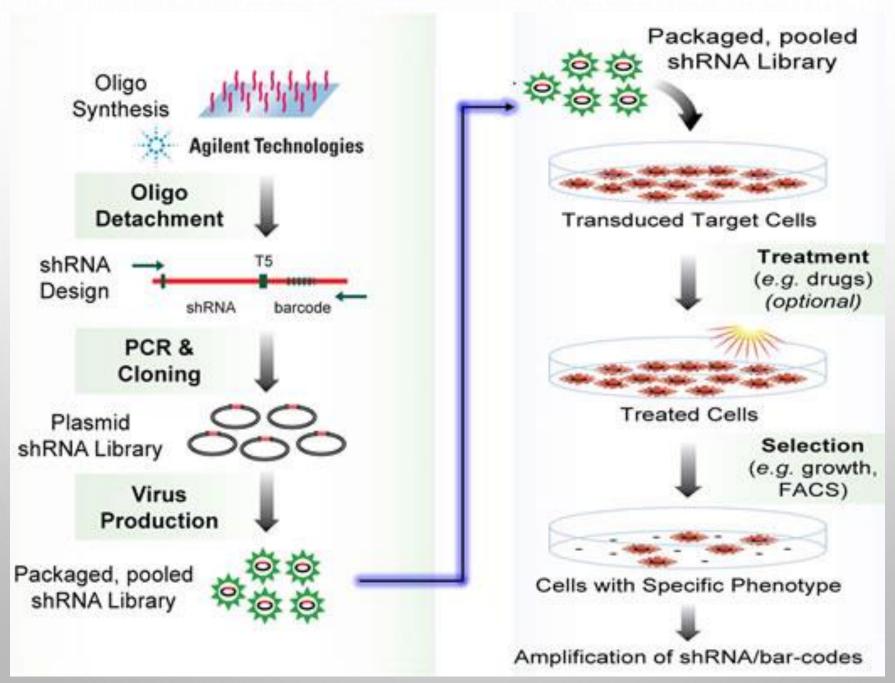
產業應用: 1. 鑑定藥物作用機轉; 2. 抗藥性基因; 3. 抗癌基因標的; 4. 抗病毒基因



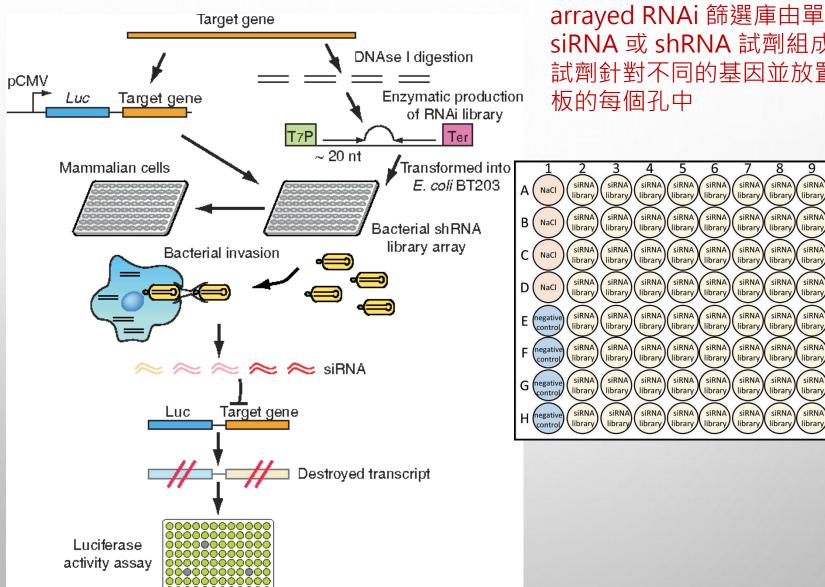
shRNA (a) Pooled RNAi 篩選庫 Mix and package into viral vector Transduction at low viral titers such that one cell takes up one shRNA Negative selection Positive selection Reference control

97





(b) arrayed RNAi 篩選庫

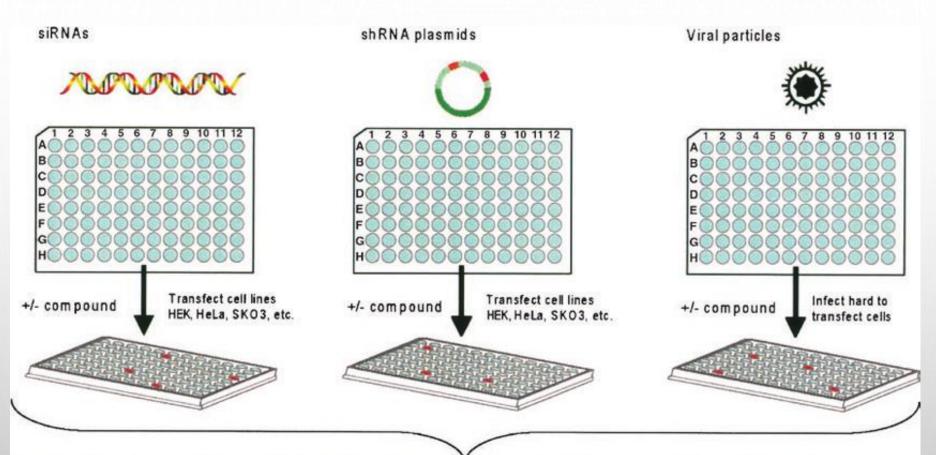


arrayed RNAi 篩選庫由單獨的 siRNA 或 shRNA 試劑組成,這些 試劑針對不同的基因並放置在多孔

negative

library

library



Score phenotypes from cell based assays

- · Luminescent reporter
- Cell viability
- High content imaging
- Fluorescent reporter
- ELISA
- FRET
- · Barcode PCR

Oncology specific cell based assays

- · Wound healing
- Proliferation
- Caspase activation
- · Colony formation
- 3D Coculture
- Invasion assay
- Angiogenesis assay



Research Update

TRENDS in Immunology Vol.23 No.12 December 2002

Research News

RNA interference of HIV replication

Miguel Angel Martínez, Bonaventura Clotet and José A. Esté

Double-stranded RNA-mediated interference (RNAi) induces sequence-specific post-transcriptional gene silencing and has emerged as a powerful tool to silence gene expression in multiple organisms. In mammalian cells, duplexes of 21 nucleotide RNAs, known as short-interfering RNAs (siRNAs), efficiently inhibit gene expression, Recent research demonstrates the general use of siRNAs to specifically inhibit HIV-1 replication by targeting viral or cellular genes. Importantly, RNAi opens a new avenue for gene-based therapeutics.

siRNA可以將HIV水準 降低30-50倍!!!

Inhibition of Retroviral Pathogenesis by RNA Interference

Short interfering RNA confers intracellular antiviral immunity in human cells

Leonid Gittin**; Sveta Karelsky* & Raul Andino*

A lentivirus-based system to <u>functionally silence genes</u> in primary mammalian cells, stem cells and transgenic mice by RNA interference

Douglas A. Rubinson^{1*}, Christopher P. Dillon^{1,2*}, Adam V. Kwiatkowski¹, Claudia Sievers^{1,2,3}, Lili Yang⁴, Johnny Kopinja⁵, Mingdi Zhang⁵, Michael T. McManus^{1,2}, Frank B. Gertler¹, Martin L. Scott⁵ & Luk Van Parijs^{1,2}
*These authors contributed equally to this work.

Blocking oncogenes in malignant cells by RNA interference— New hope for a highly specific cancer treatment?

A little more than one year after the first demonstration that silencing of endogenous human genes is possible in cell culture, the new tool of RNA interference (RNAi) enters the field of tumor therapy.

Advantages of co-delivery of siRNA and small molecule anticancer drug

Overcome multidrug resistance

Generate synergistic apoptotic effect

Reduce toxicity

