

# Bi/Ch 113: Biochemistry of the Cell, Spring 2017

- protein and membrane trafficking

[www.its.caltech.edu/~bich113](http://www.its.caltech.edu/~bich113)

## Goals:

- To present our current understanding of these topics, with an emphasis on the underlying molecular mechanisms.
- To illustrate the experimental strategies biologists use to analyze cell biological processes.
- To read research papers in detail and to discuss them critically.

# Bi/Ch 113: Biochemistry of the Cell, Spring 2017

Lectures: Tuesday and Thursday 11-11:55 am, Broad 100

*Section attendance is mandatory*

## **Teaching assistants**

Amanda Mock, [jmock@caltech.edu](mailto:jmock@caltech.edu)

Greg Varuzhanyan, [gvaruzhanyan@caltech.edu](mailto:gvaruzhanyan@caltech.edu)

Ruohan Wang, [rwwang@caltech.edu](mailto:rwwang@caltech.edu)

Section 01: Wednesday 2-3:55 pm, Broad 200 (Ruohan)

Section 02: Thursday 7-8:55 pm, Kerckhoff 101 (Greg)

## **Grading**

Midterm: due at the beginning of class, May 9.

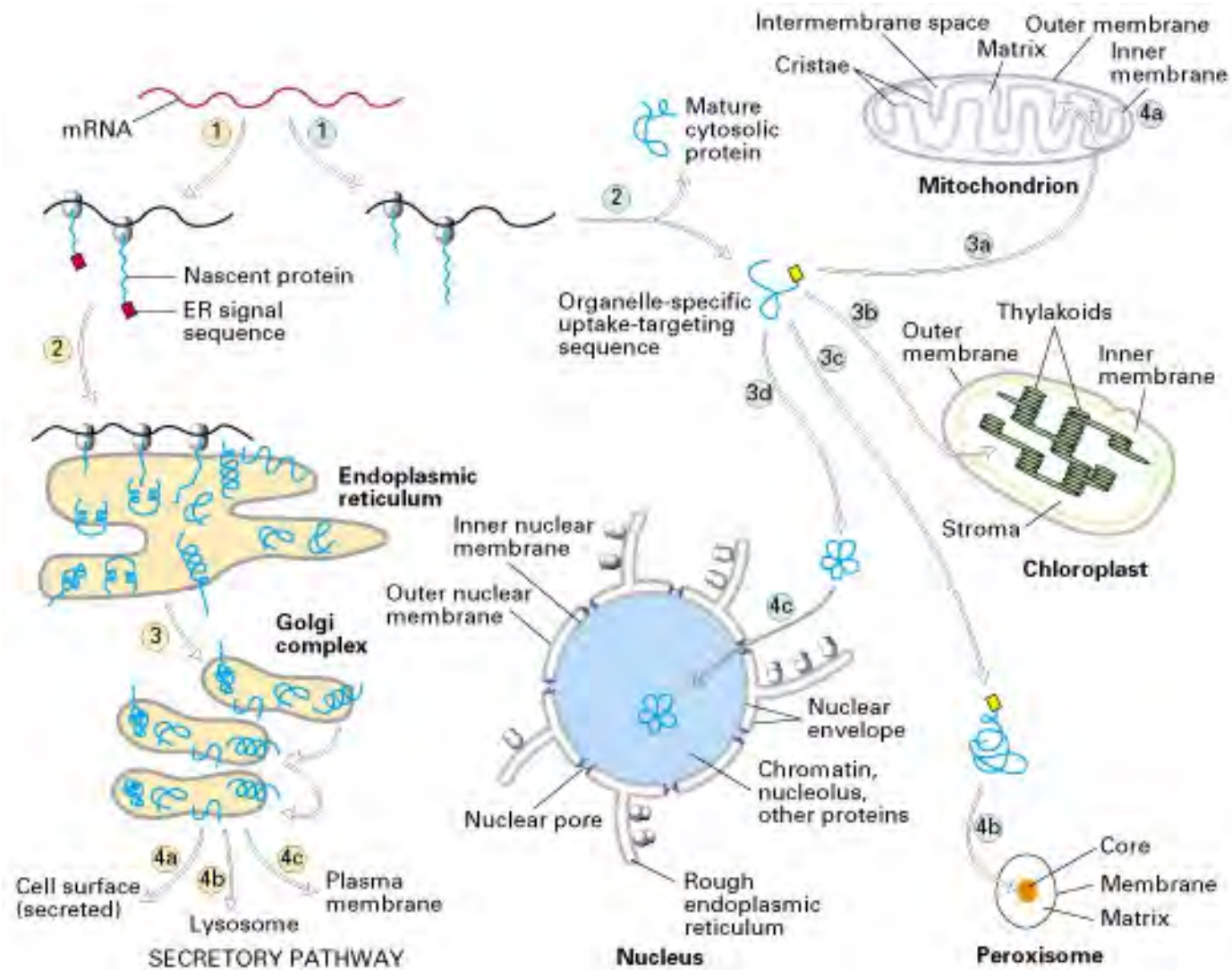
Final: due June 8 for seniors and graduate students; due June 15 for others.

Course grade is determined by: section performance, 1/3; midterm, 1/3; final exam 1/3.

No extensions are allowed for paper summaries, the midterm, or the final.

# Protein and membrane trafficking

- How do proteins get to their destination?



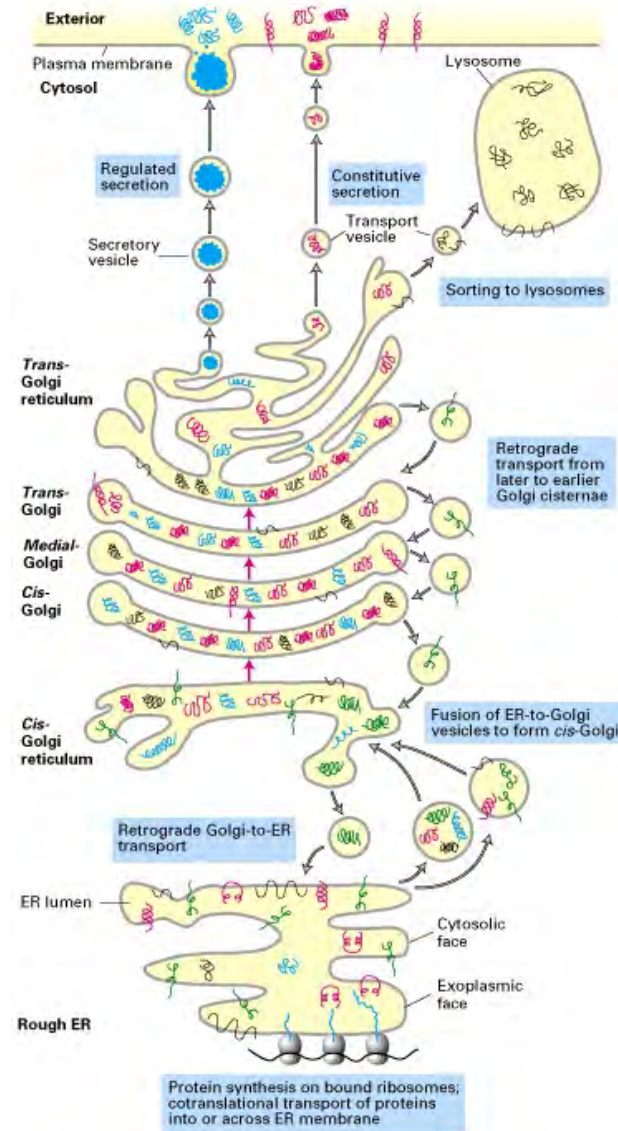
# Protein and membrane trafficking

Endocytosis

Secretion

Vesicle fusion

Protein translocation



How are proteins, lipids, and organelles sorted to the correct compartment?

Model systems:

ER: translocation, secretion, morphology

Viruses

Endocytosis

Nucleus structure and import

Autophagy

Apoptosis

Mitochondria

Relation to disease

Oncogenesis

Lodish et al, 2000

# Guidelines for reading and discussing papers

- 8 research papers will be presented and critiqued in section.

## Sections of paper

Title

Abstract

Introduction

Materials and methods

Results (Figures/Tables)

Discussion

# Reading and discussing research papers

1. What is the overall question the authors are addressing?  
Is it an important question; does it address a gap in knowledge?
2. What experimental strategy do the authors use to answer this question?
3. Are the experiments appropriate for the question, and are they well-done? Are the data convincing?
4. Are the experiments properly controlled?
5. How strongly do the data support the conclusions? That is, do you believe their conclusions?
6. Are the findings interesting and important?
7. Where does one go from here?
  - What additional studies can be done to test the conclusions?
  - Is a new line of research suggested by these findings?

# Presenting research papers in section

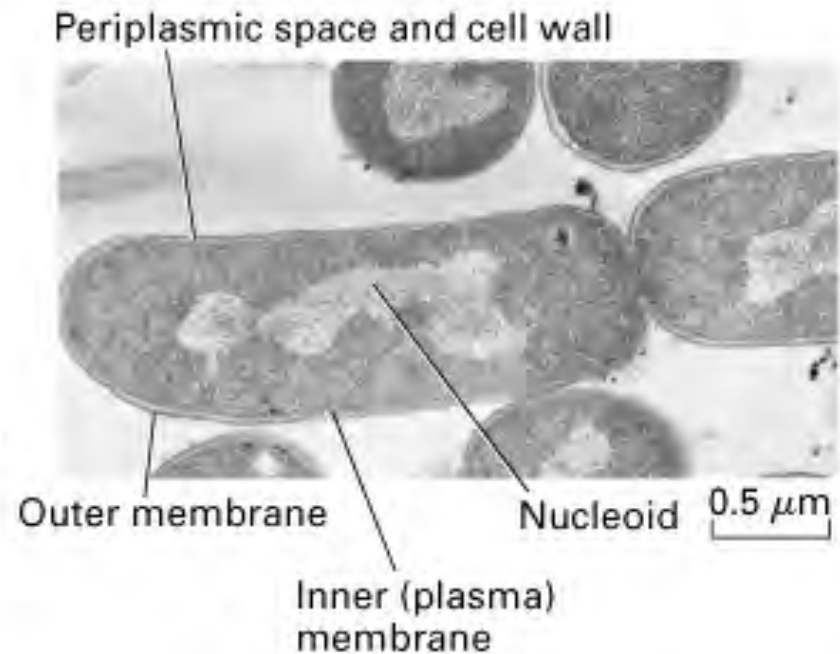
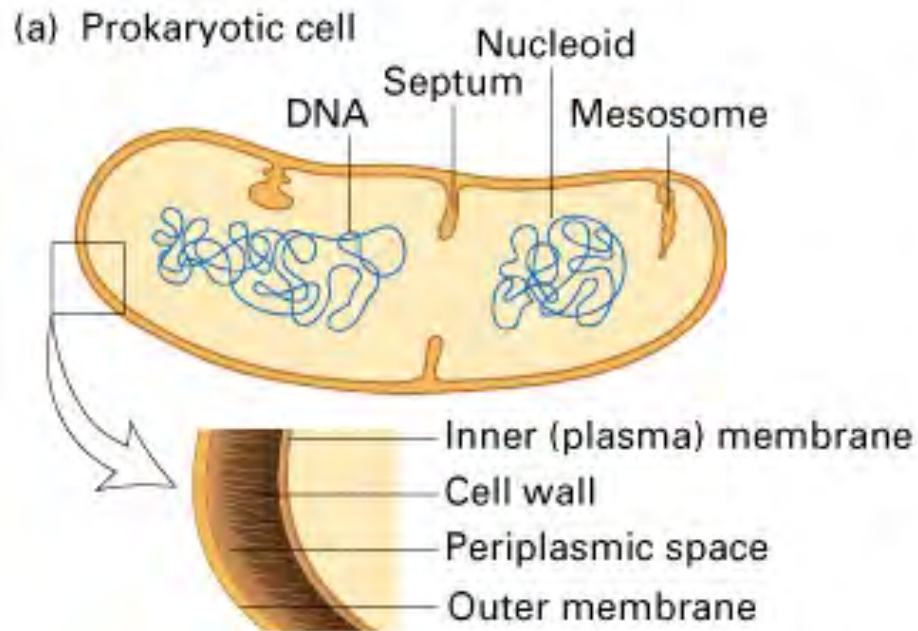
TAs will provide guidance on presentations

1) Provide **introduction** giving background material and describing the state of knowledge at the time the experiment was done. How does this work fit into the overall state of the field?

2) Analyze the **individual figures/tables** in the paper. What is the figure purporting to demonstrate, and what method is used? Are the data convincing? Which are the most critical pieces of data?

3) **Overall evaluation/conclusions**: does the paper accomplish what it claims to do? Are the results important? What are the directions for future work?

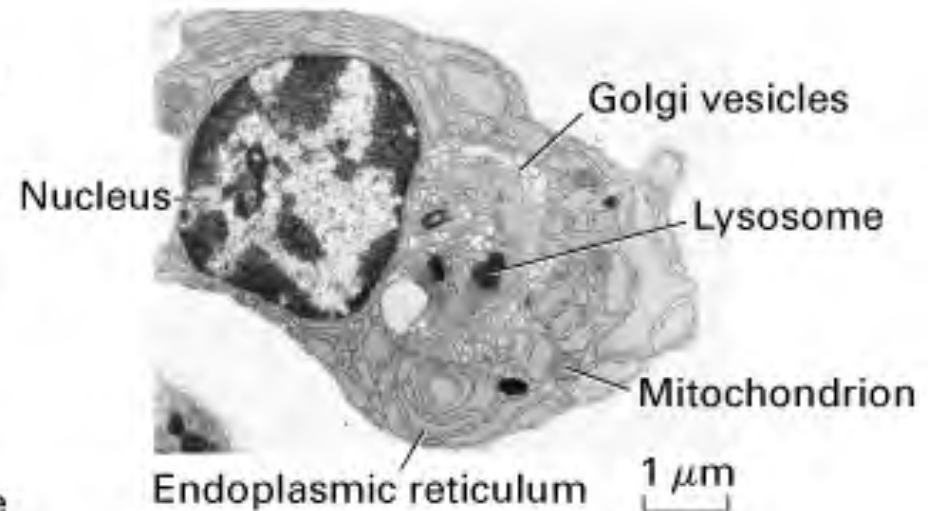
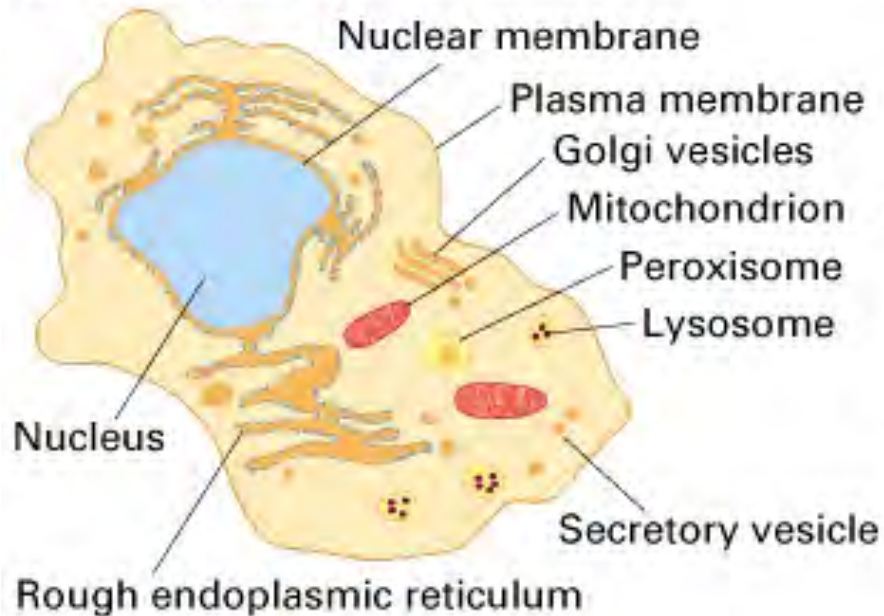
# Cellular dimensions- prokaryotes



Bacteria: 0.5  $\mu\text{m}$  diameter

# Cellular dimensions- eukaryotes

(b) Eukaryotic cell



Mammalian cell: 10-30  $\mu\text{m}$

Nucleus: 3-10  $\mu\text{m}$

Lipid bilayer:  $\sim 4$  nm

Vesicles:  $>50$  nm

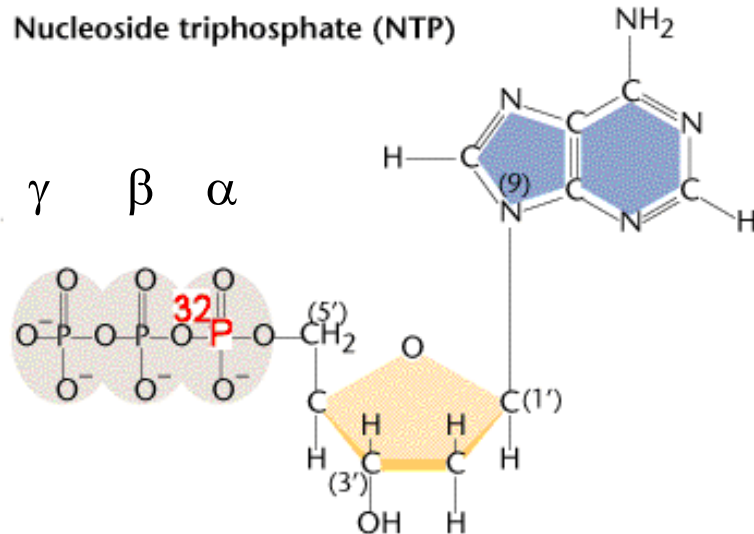
Mitochondria: 0.4  $\mu\text{m}$  diameter

$\alpha$ -helix: 11 Å diameter; 1.5 Å translation per residue

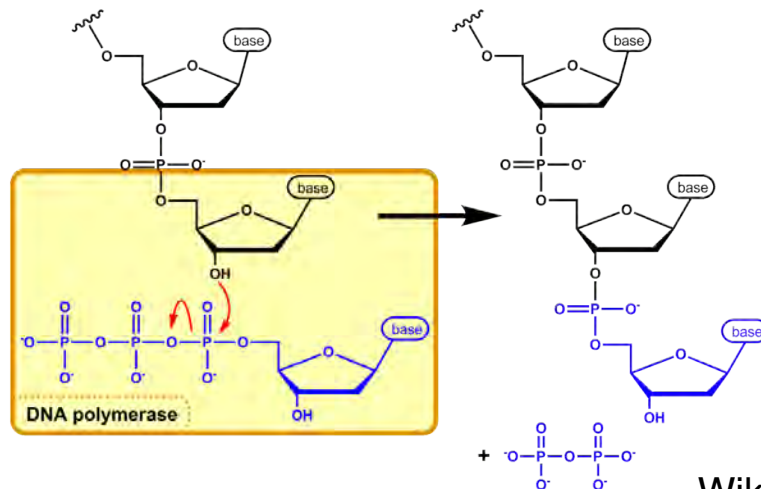
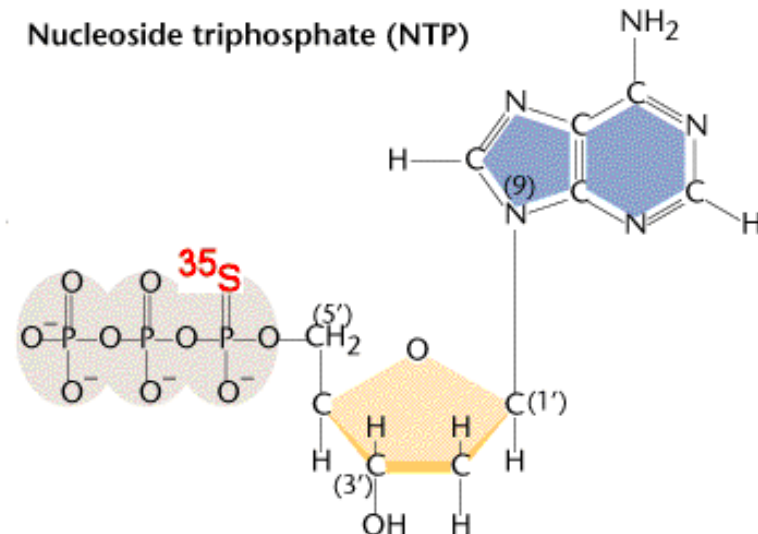
**Lodish et al, 2000**

# Labeling nucleic acids with radioactive isotopes

Nucleoside triphosphate (NTP)

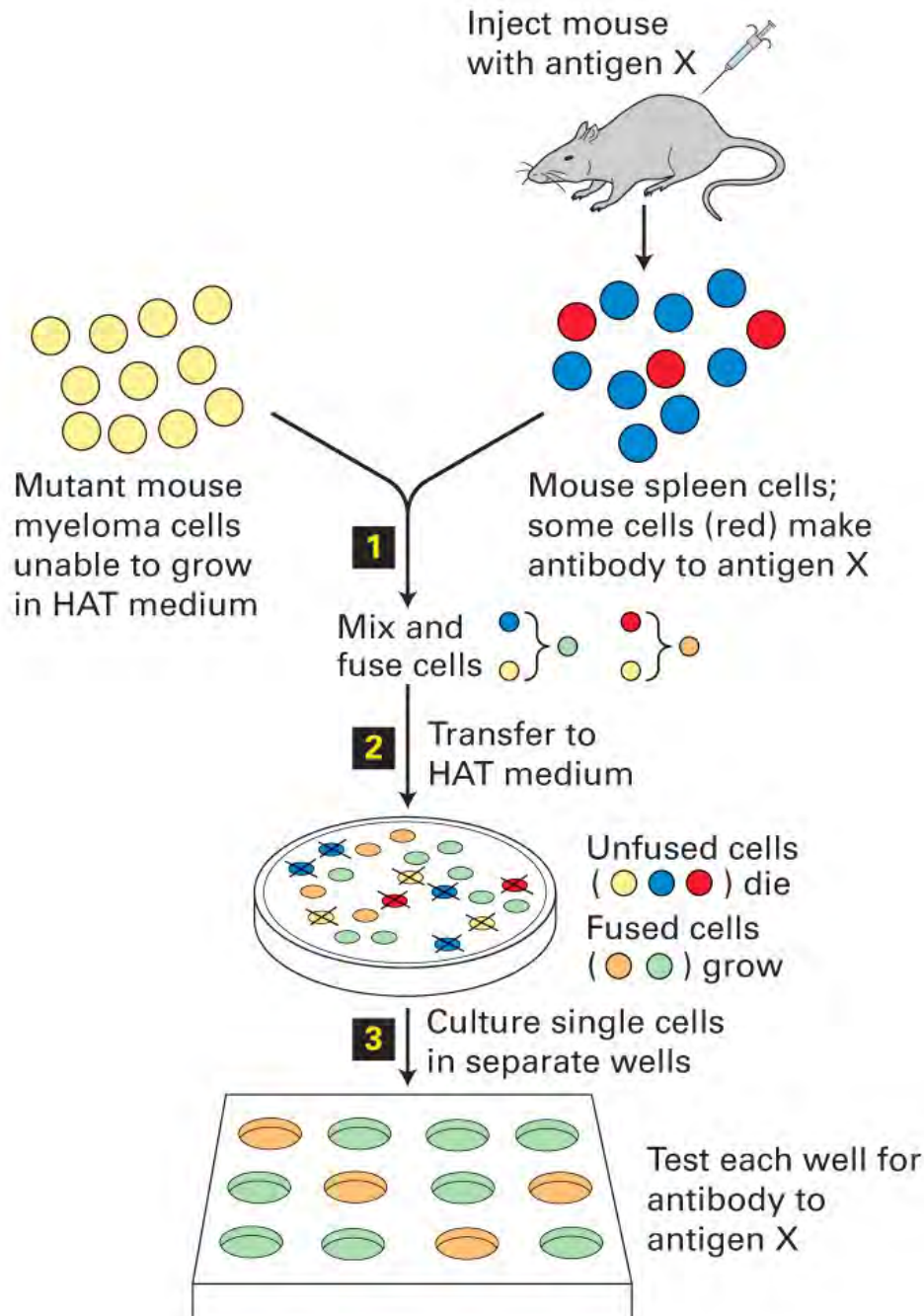


Nucleoside triphosphate (NTP)



During DNA synthesis the  $\alpha$ -phosphate group of the free nucleotide is added to the 3' OH of the growing chain.

# Monoclonal antibodies



- Polyclonal antibodies recognize multiple epitopes.
- A monoclonal antibody recognizes a single epitope.
- Monoclonal antibody is a defined, renewable reagent.
- Specificity versus sensitivity.

# Assigning gene function

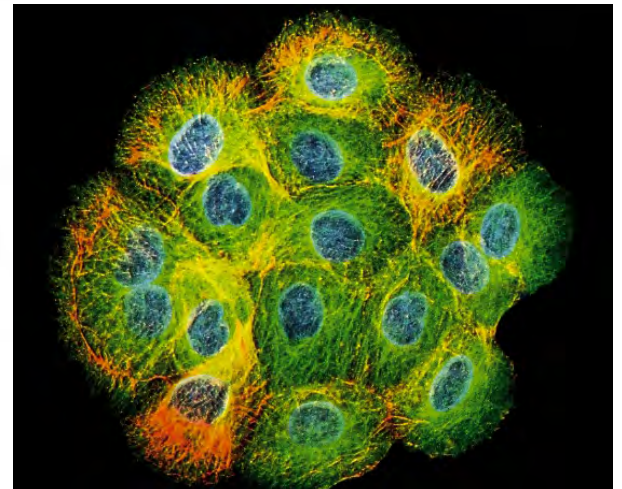
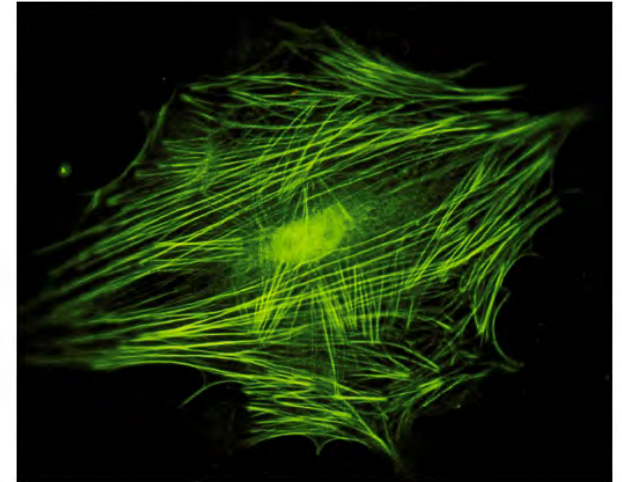
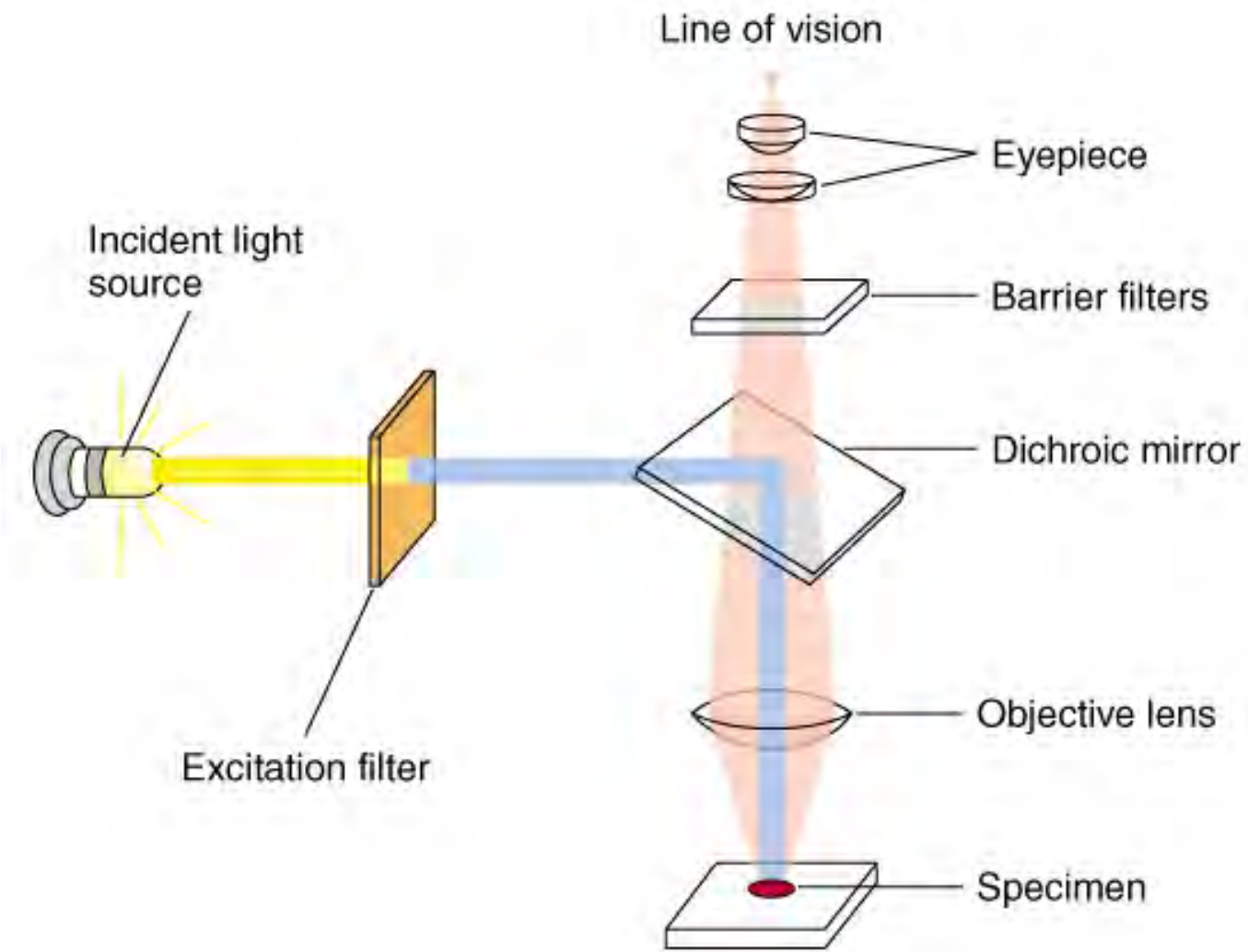
## A. Temporal and spatial correlations-- limitations

1. RNA localization- Northern blot analysis, RT-PCR, in situ hybridization
2. Protein localization- Western blot analysis, immunofluorescence, epitope tagging, chimeric GFP

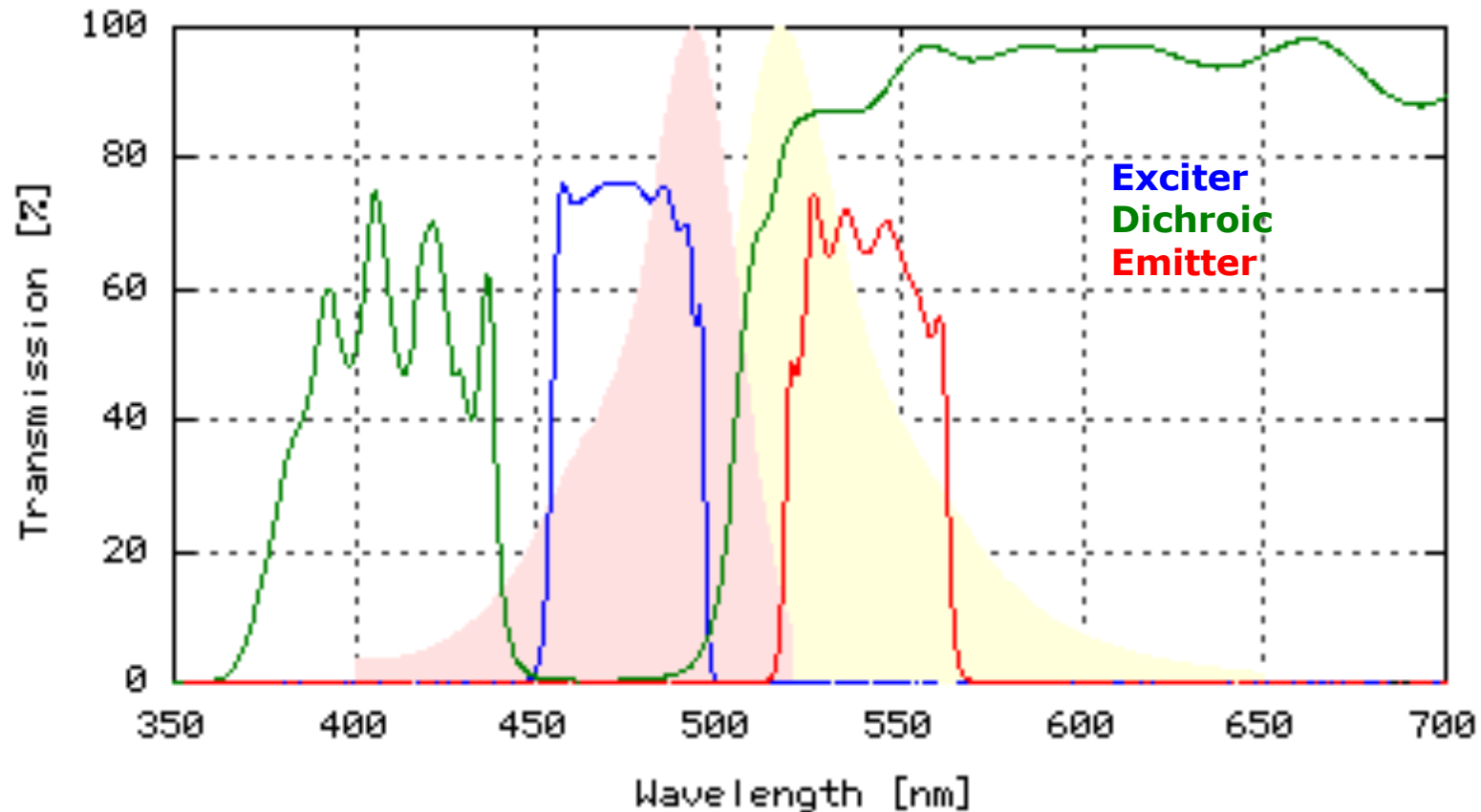
## B. Functional approaches

1. Overexpression- transfection, viral infection-- limitations
2. Misexpression- viral infection, oocyte injection
3. Gene disruption
  - Yeast
  - homologous recombination in mice- standard and conditional
  - RNA interference
  - dominant negative alleles
4. Mutational studies

# The fluorescent microscope



# Fluorescence filter set for fluorescein (FITC)



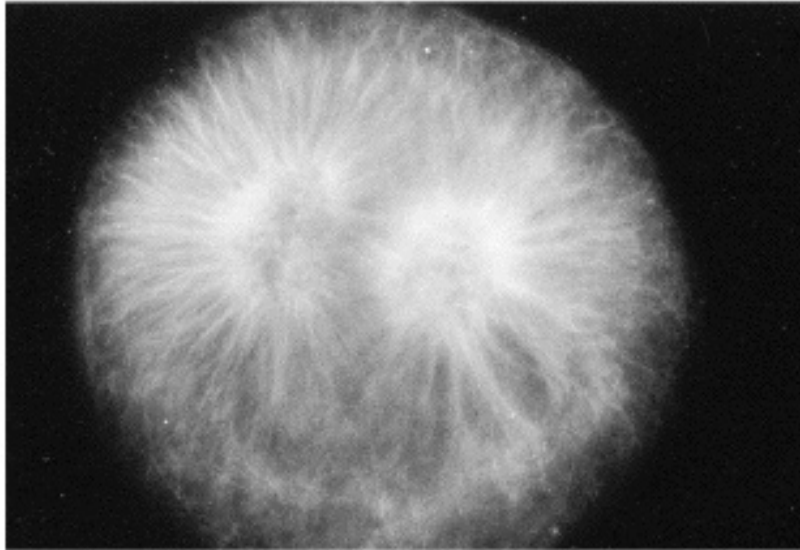
**Exciter filter:** allows excitation (shorter wavelength) light to pass

**Dichroic filter:** deflects excitation light and allows emitted (longer wavelength) light to pass

**Emitter filter:** allows emitted light to pass

# Confocal and deconvolution microscopy create sharper images

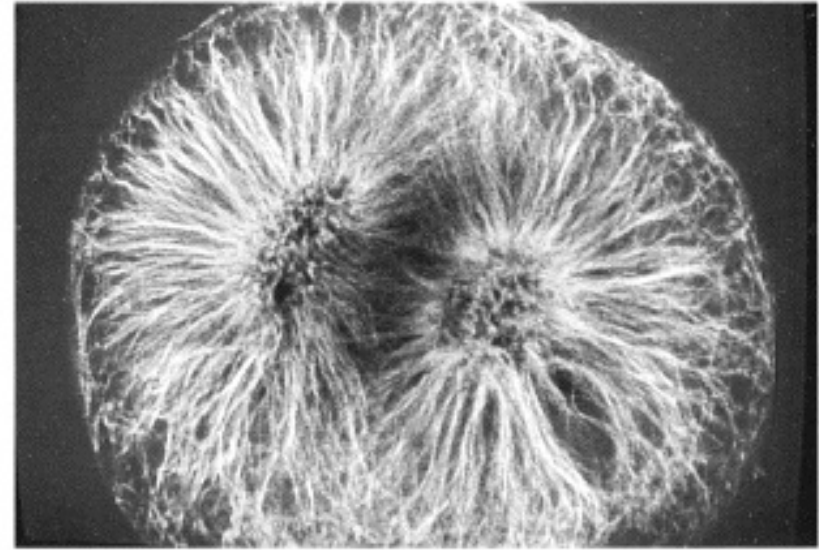
(a)



conventional

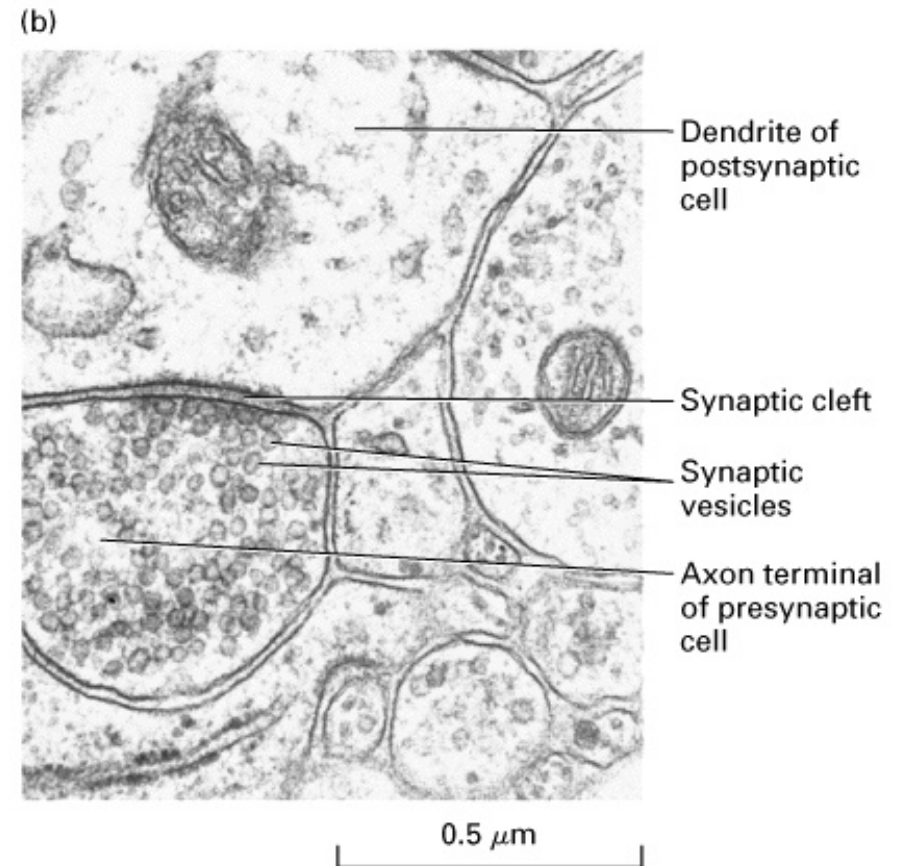
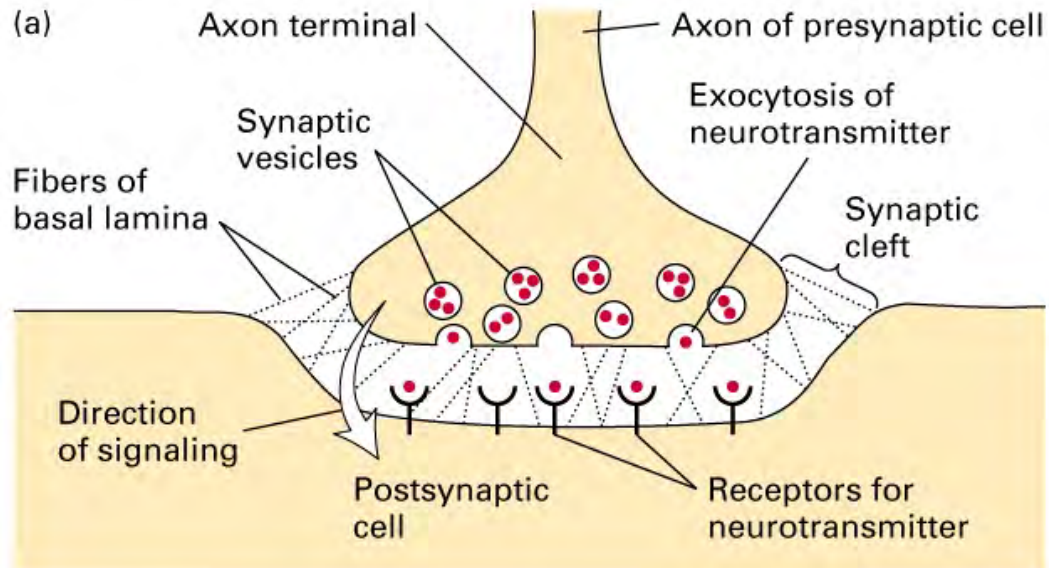
(b)

Confocal microscopy:  
optical section

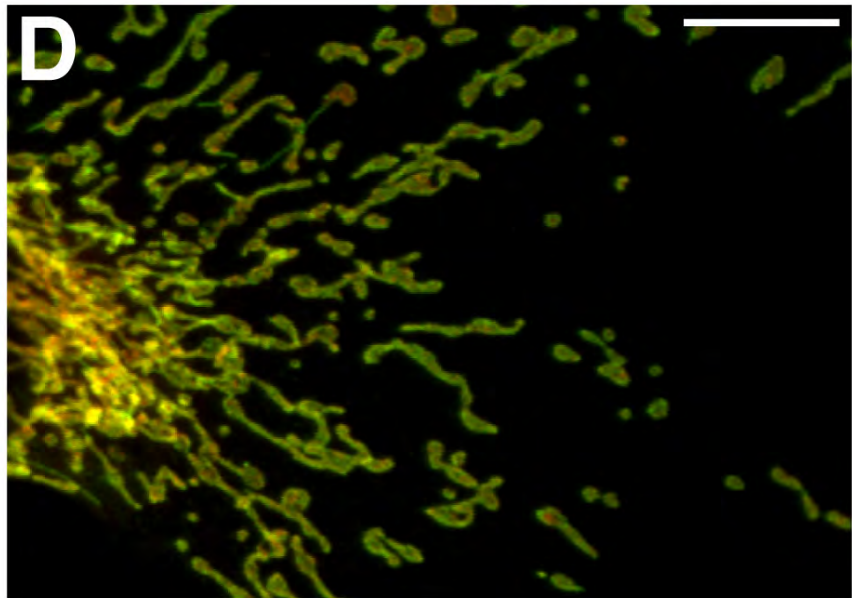
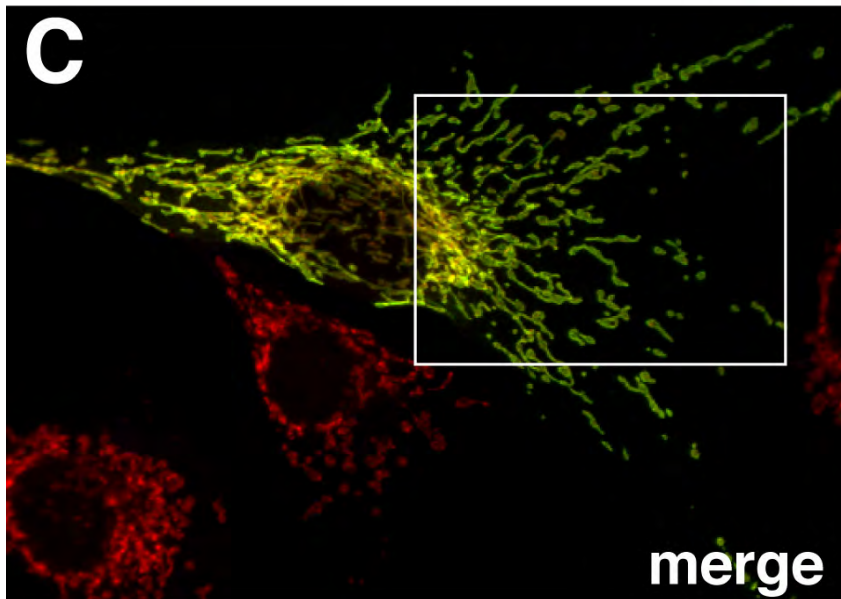
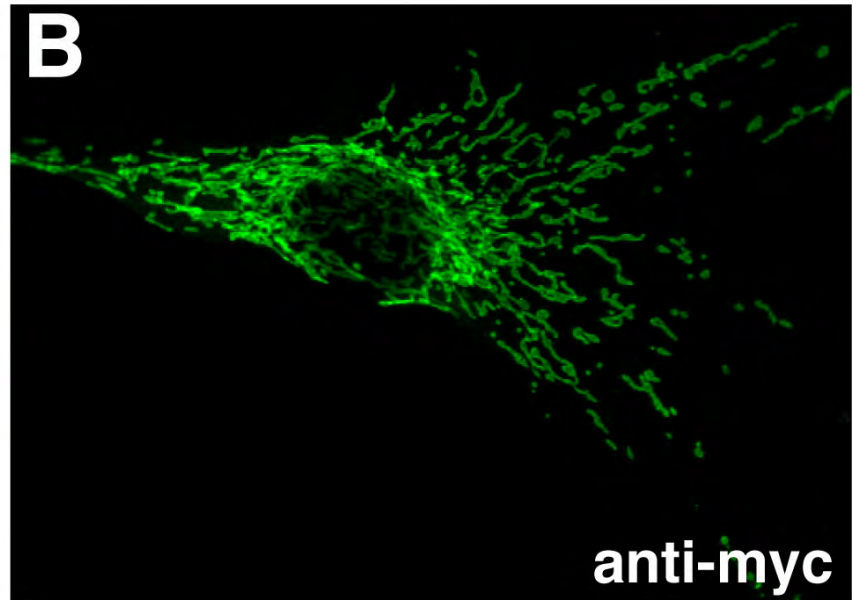
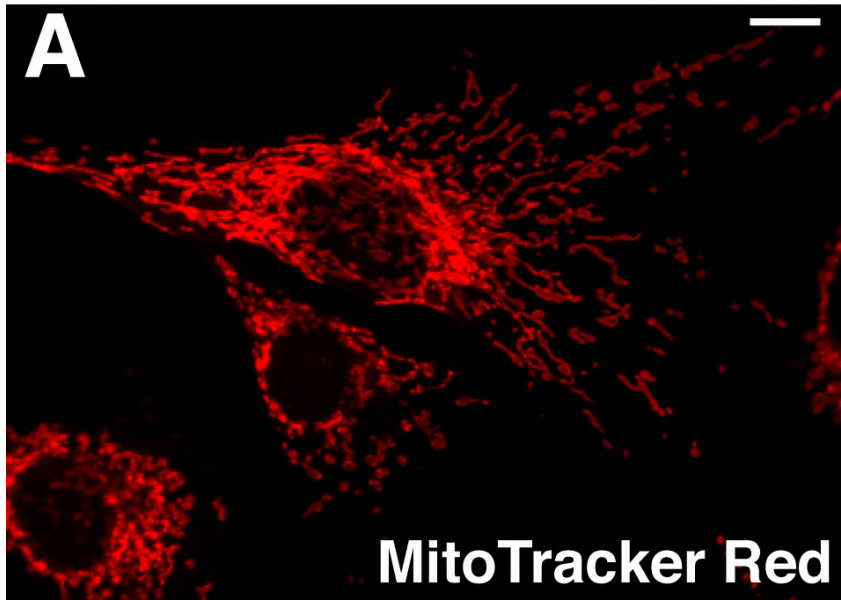


confocal

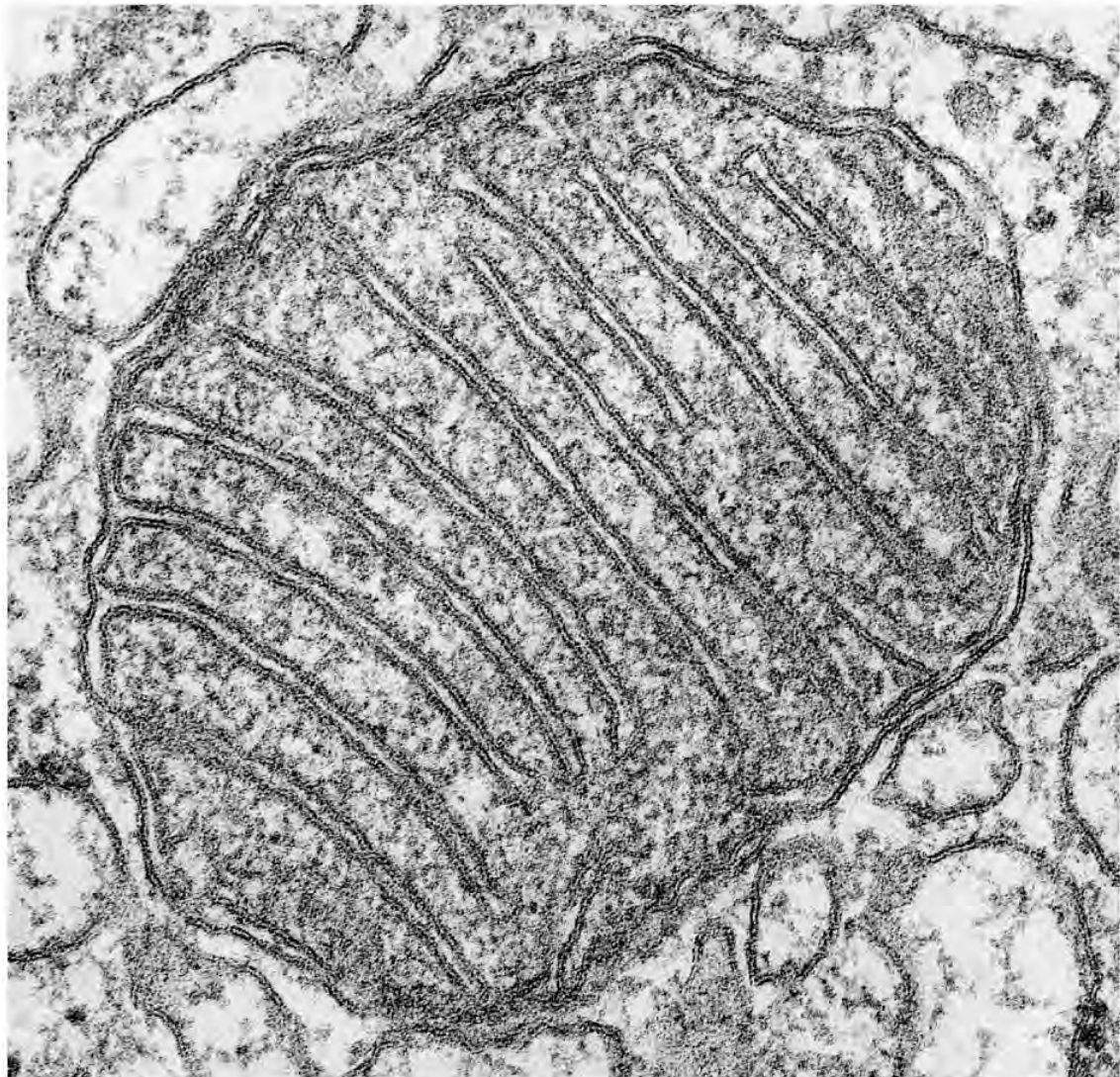
# Electron microscopy



## Mitofusins localize to mitochondria

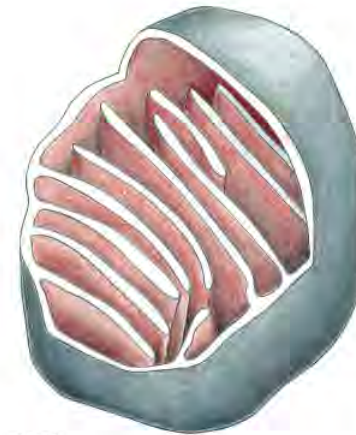


# Structure of mitochondria



(A)

100 nm

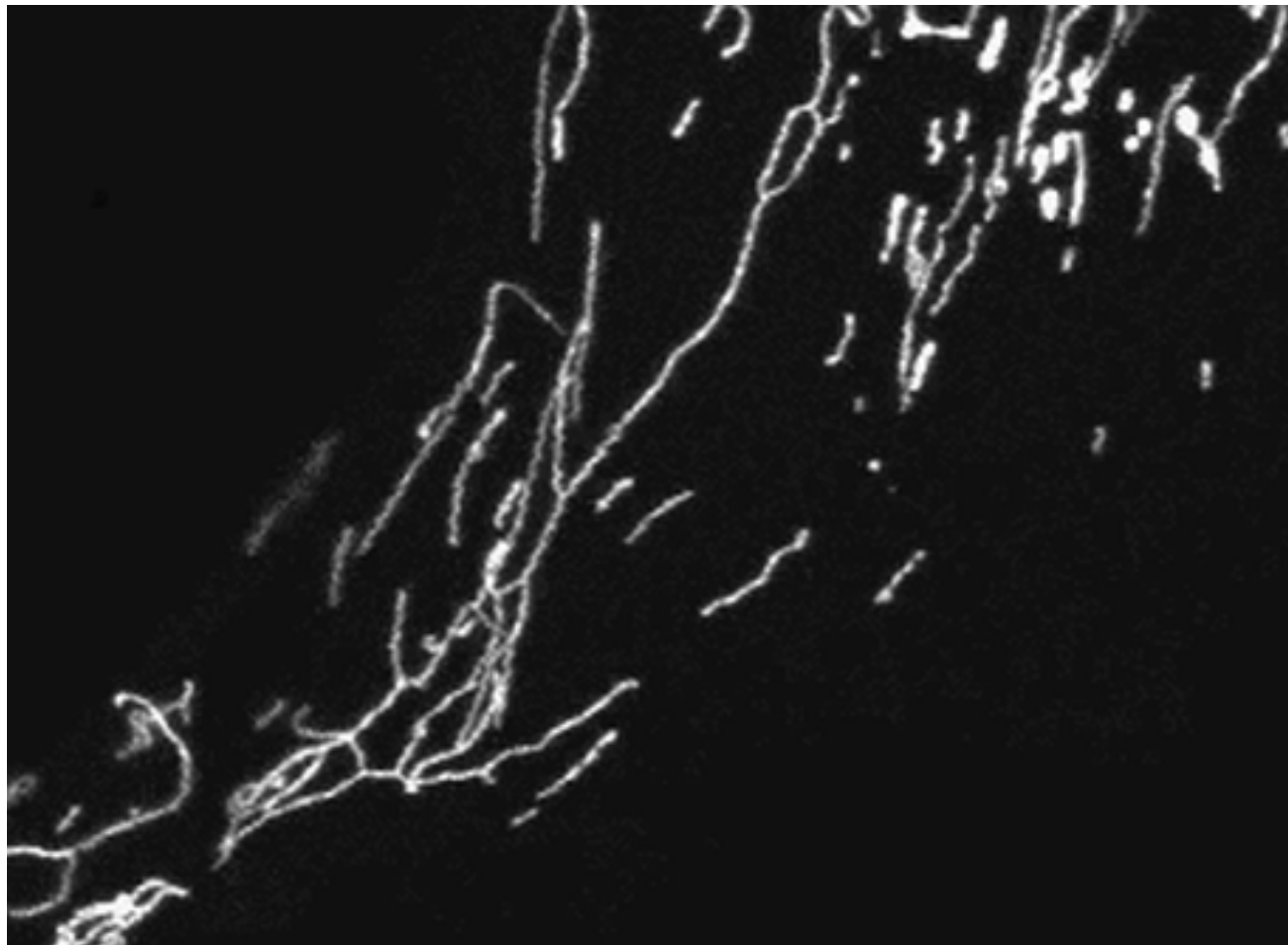
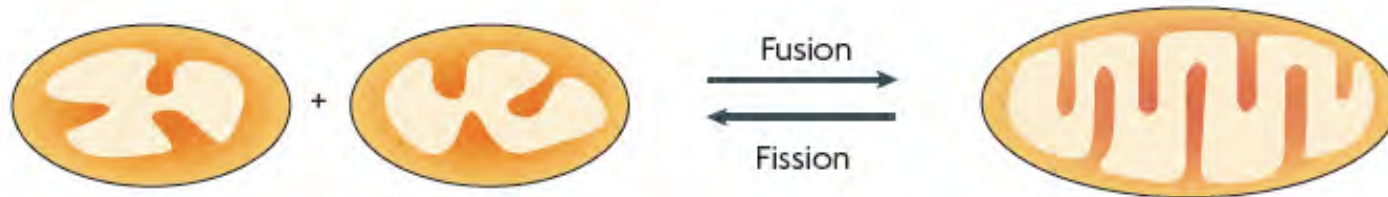


(B)

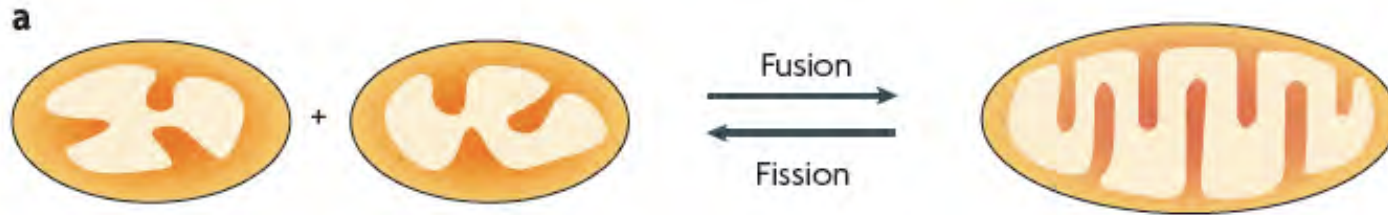


(C)

# Use of fluorescent proteins to track organelle dynamics in live cells

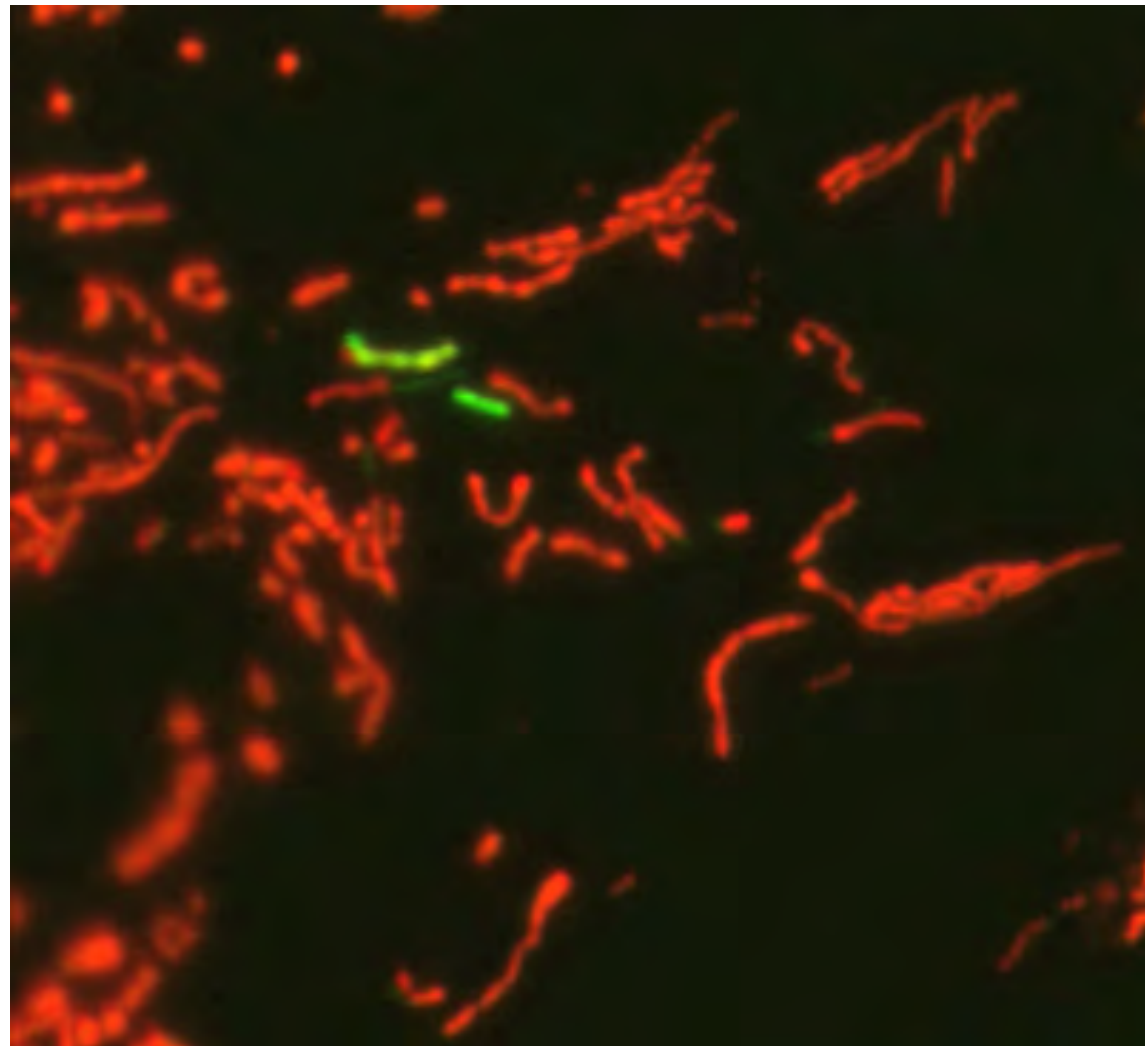


# Tracking individual mitochondria by photoactivation

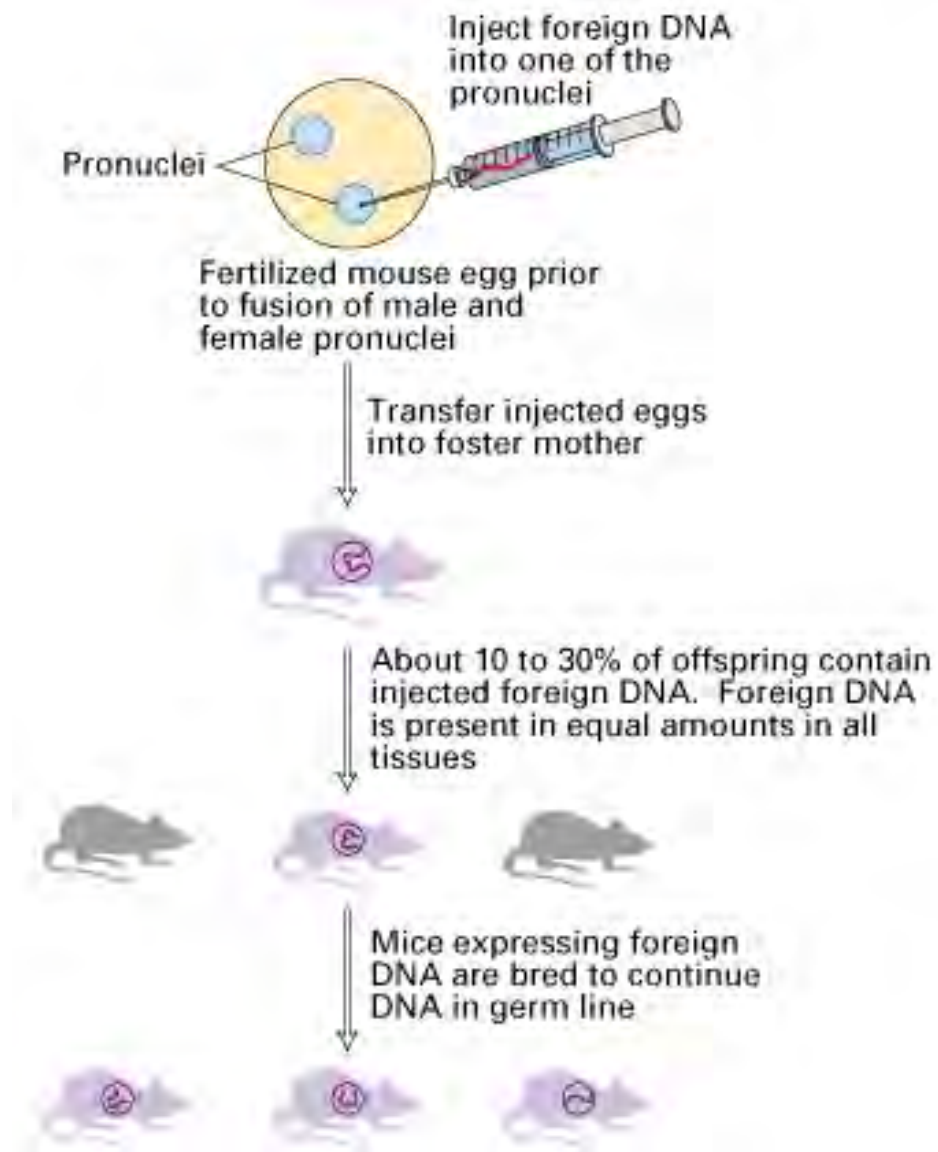


Mito-DsRed

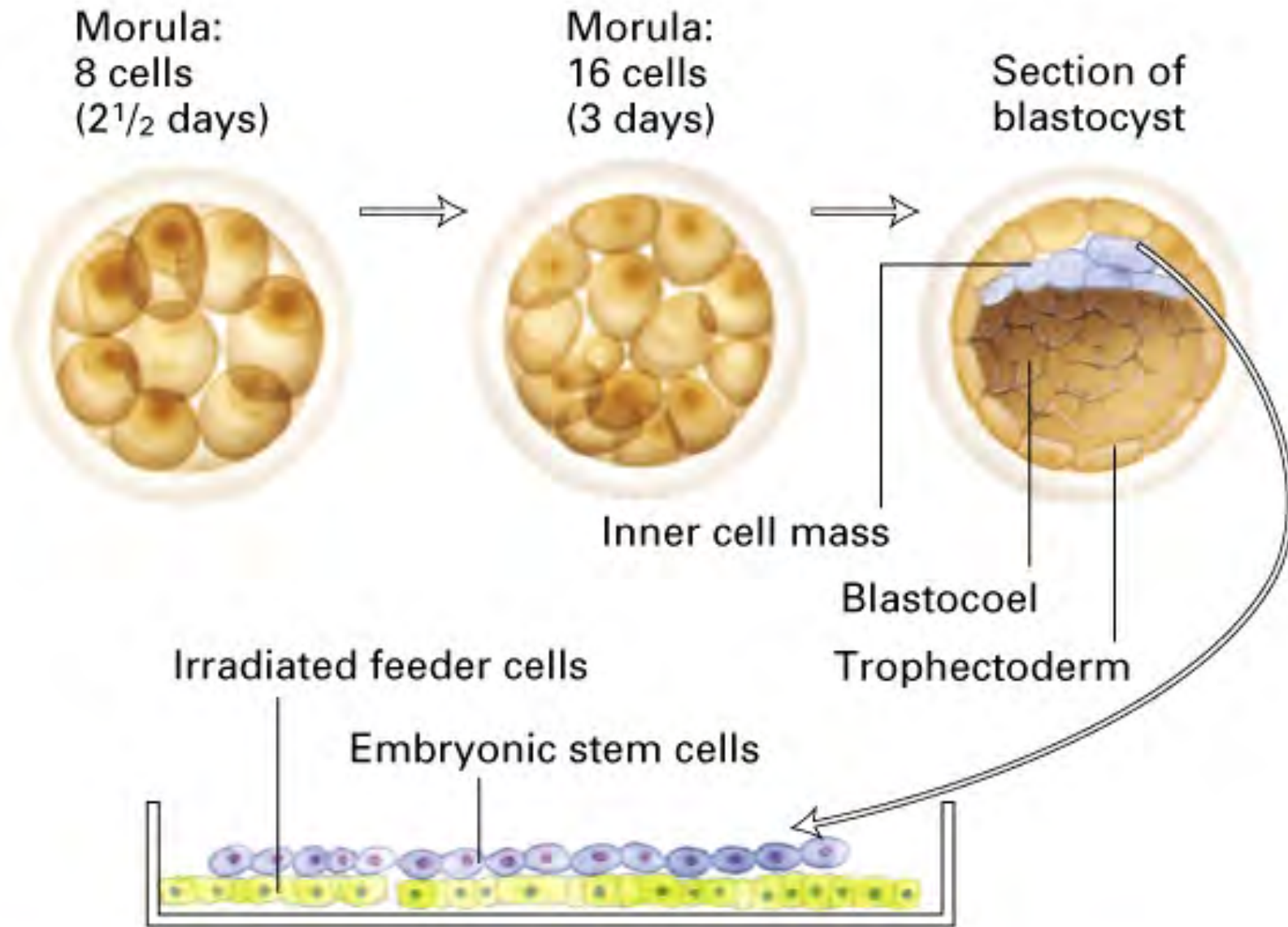
Mito-PA-GFP



# Production of transgenic mice

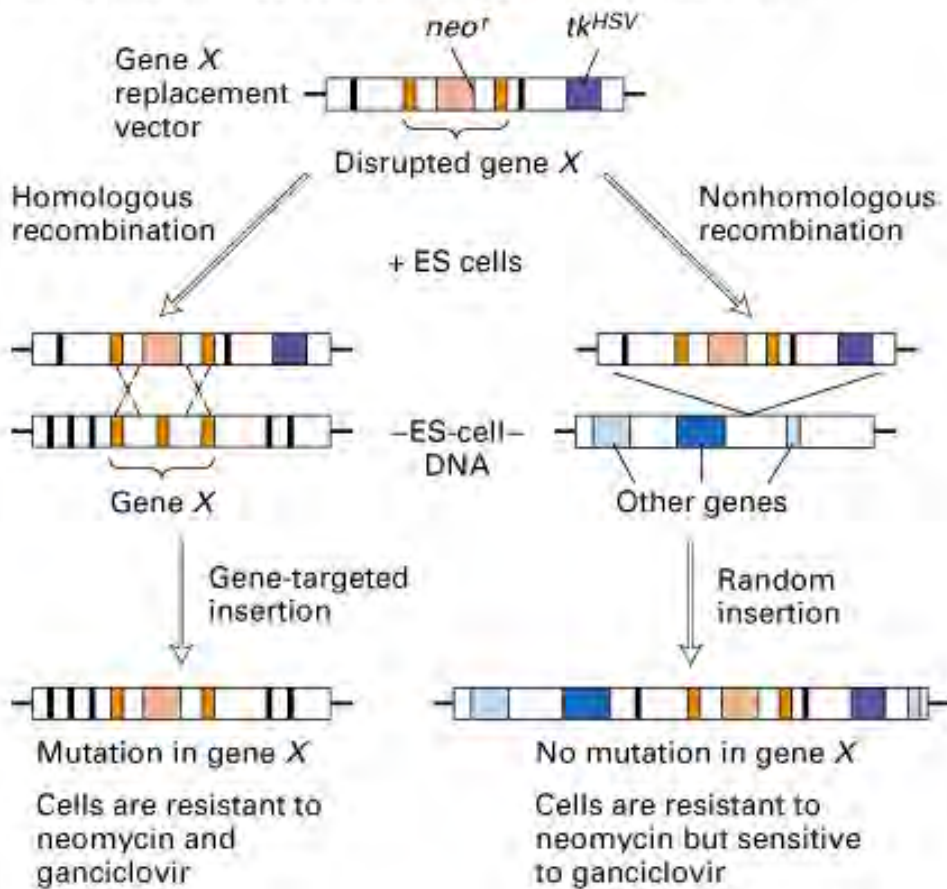


# Embryonic stem cells are totipotent

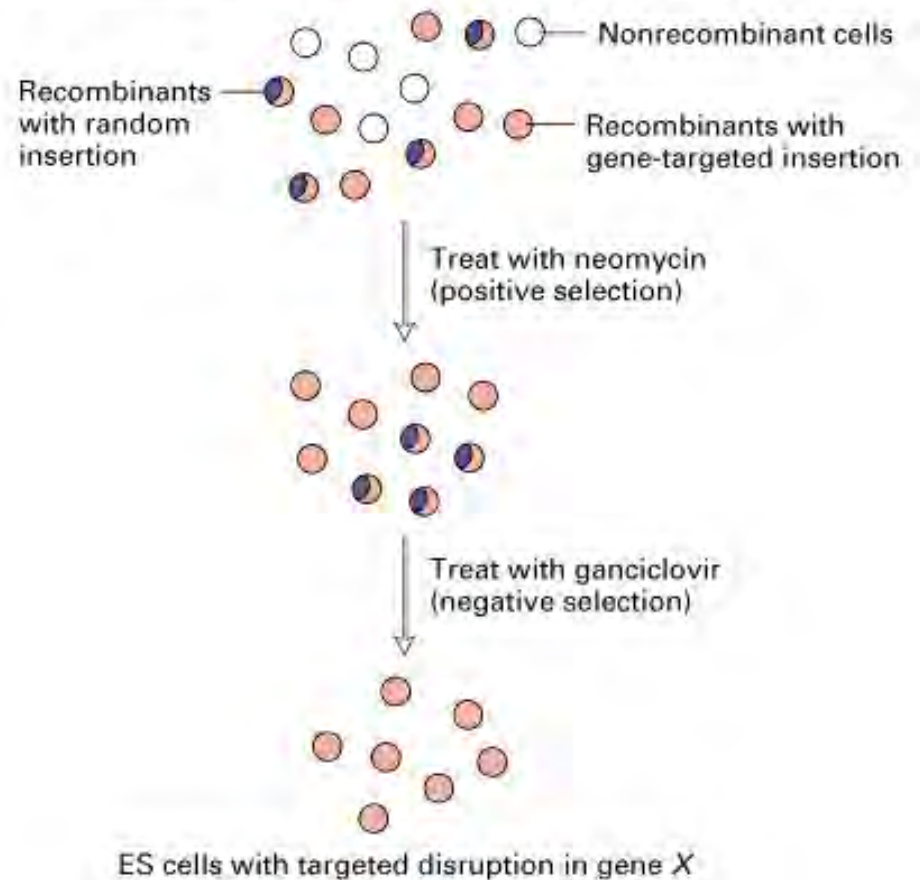


# Creation of mice ES cells carrying a knockout mutation

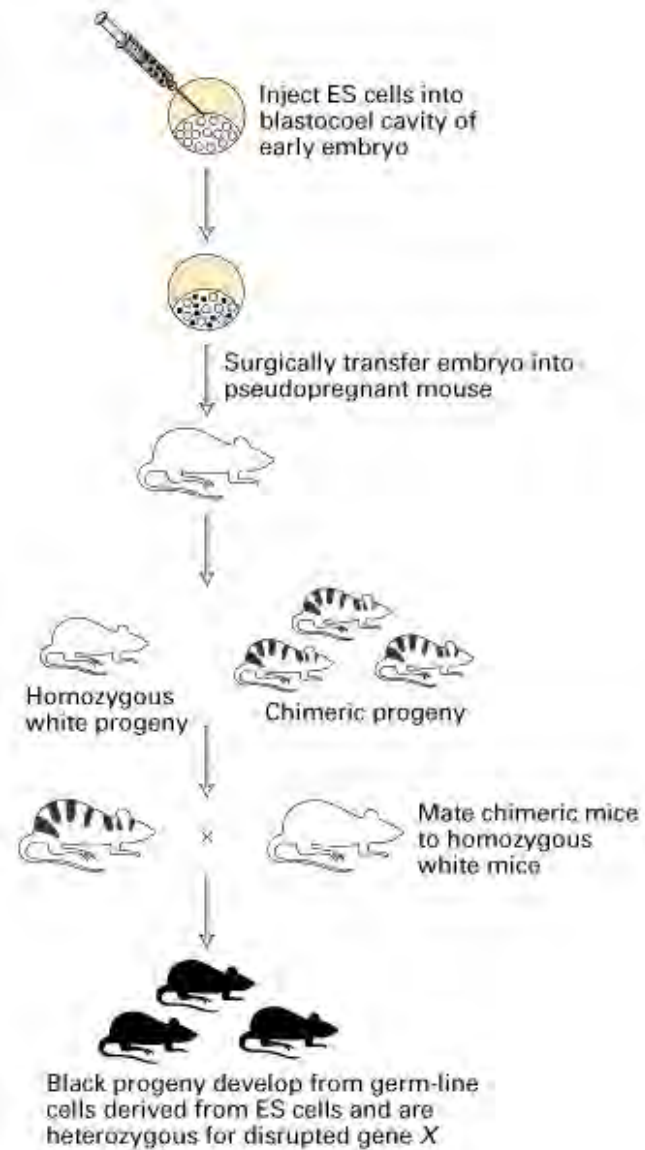
(a) Formation of ES cells carrying a knockout mutation



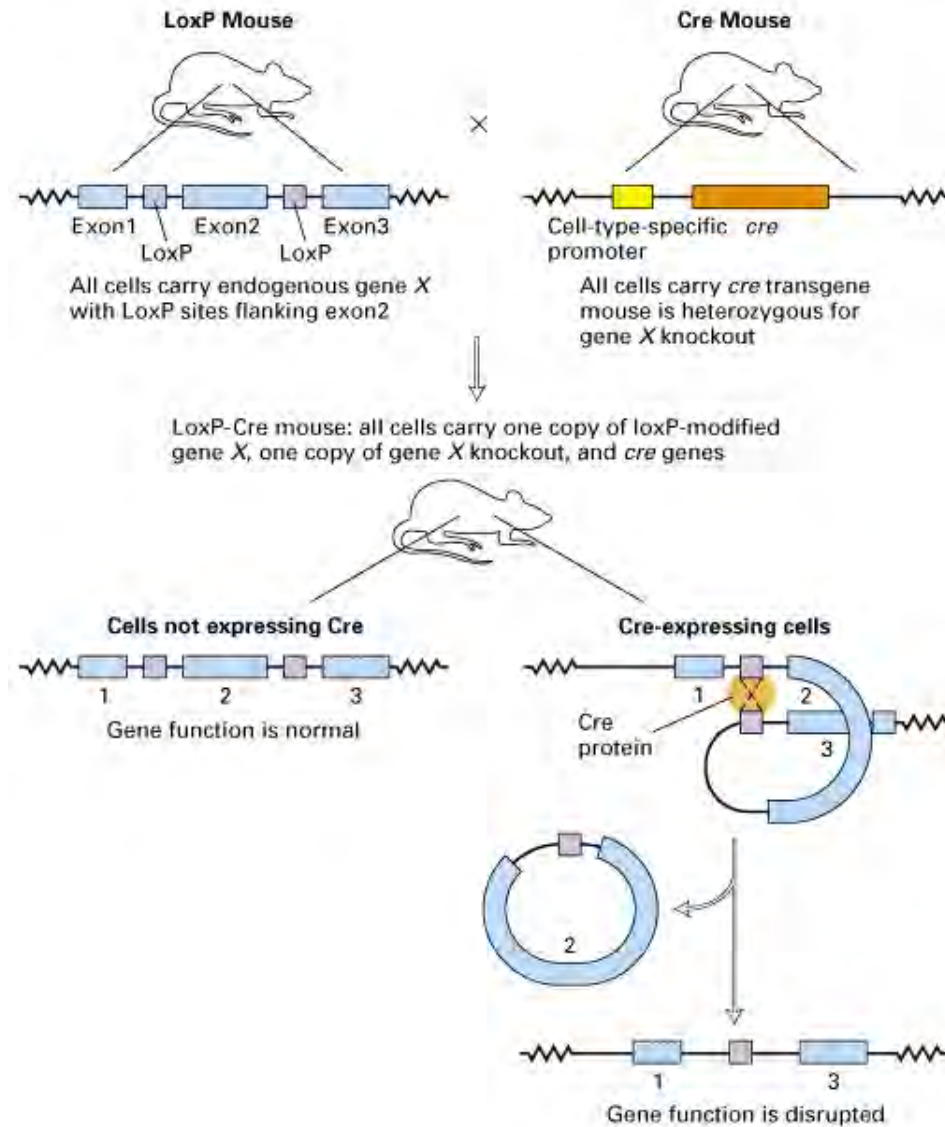
(b) Positive and negative selection of recombinant ES cells



# Gene knockout in mice

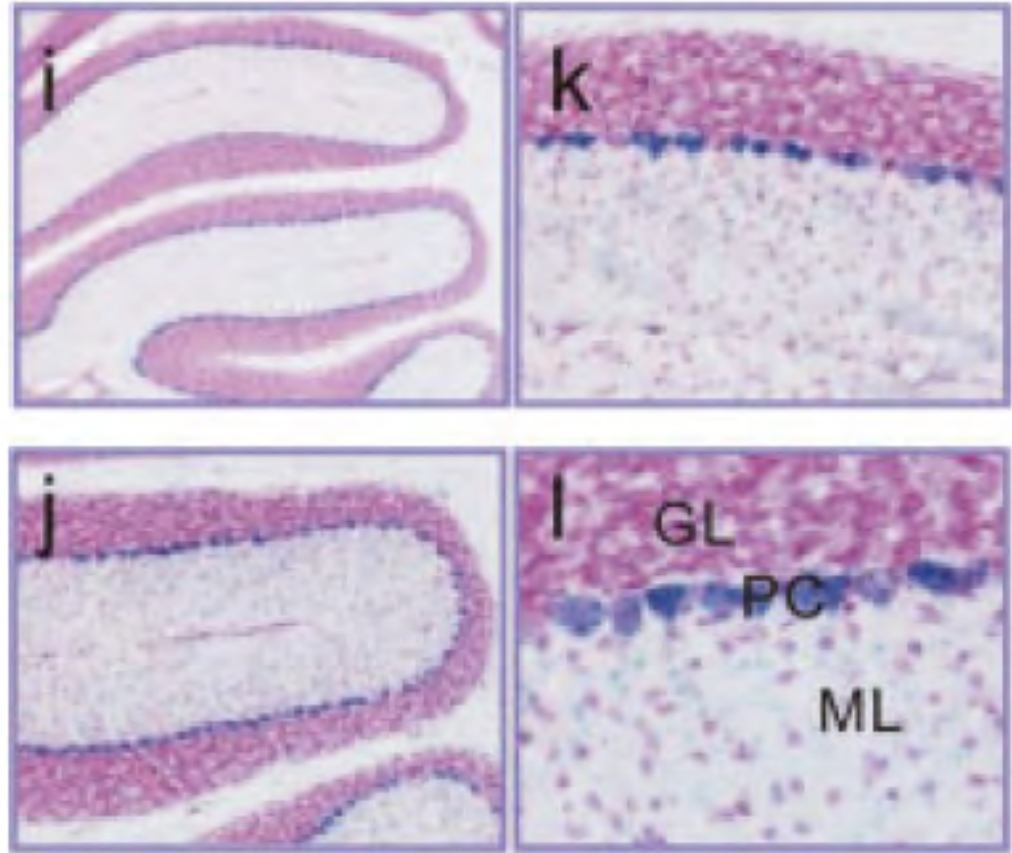
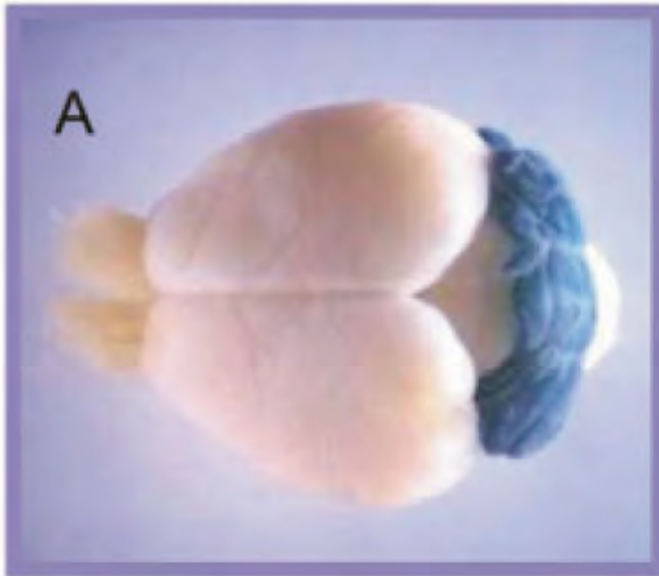


# Cell-type-specific (conditional) gene knockouts in mice



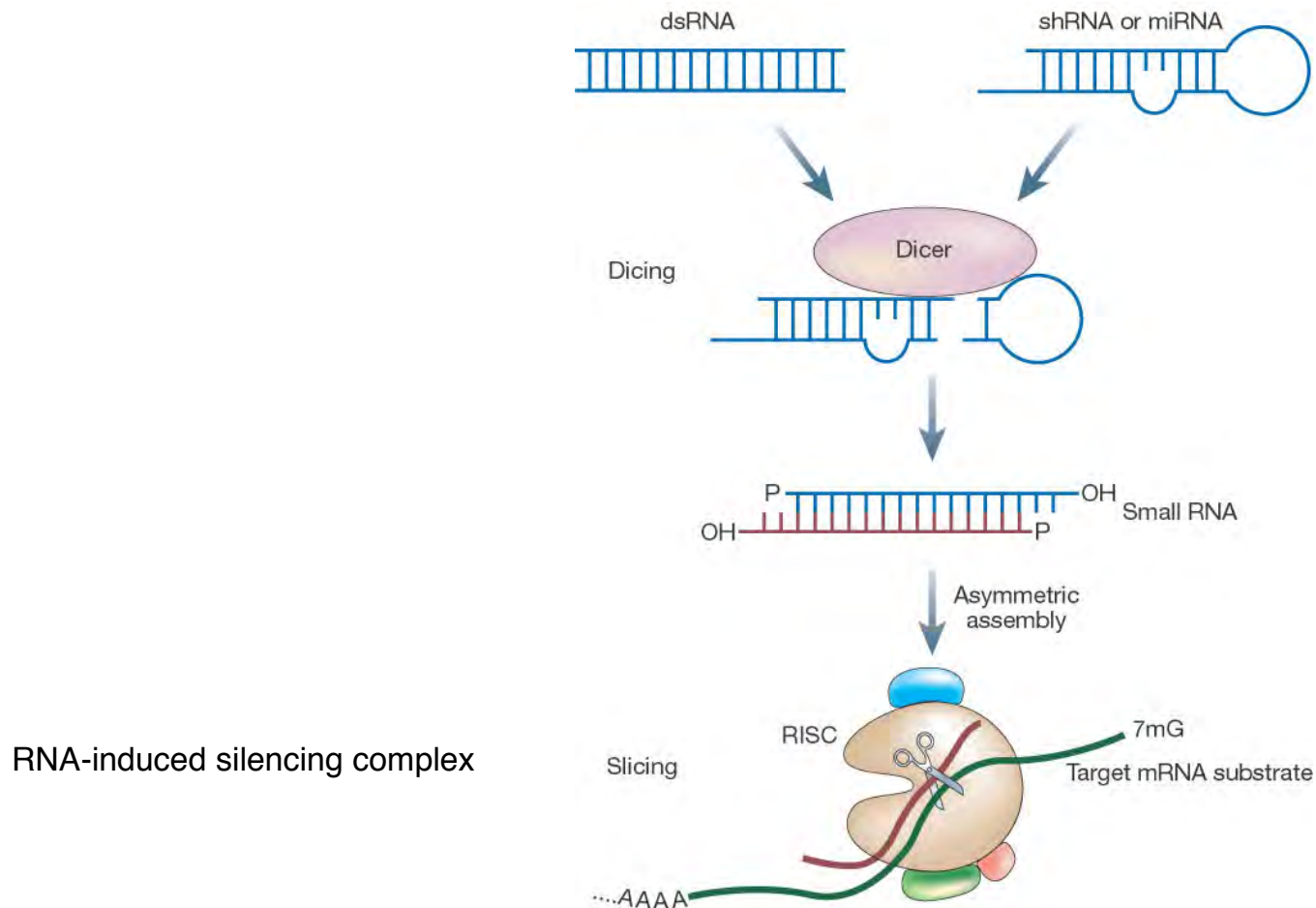
# The *cre-loxP* system allows temporal and spatial control of gene disruption

BAC-Pcp2-IRES-Cre(+/-)

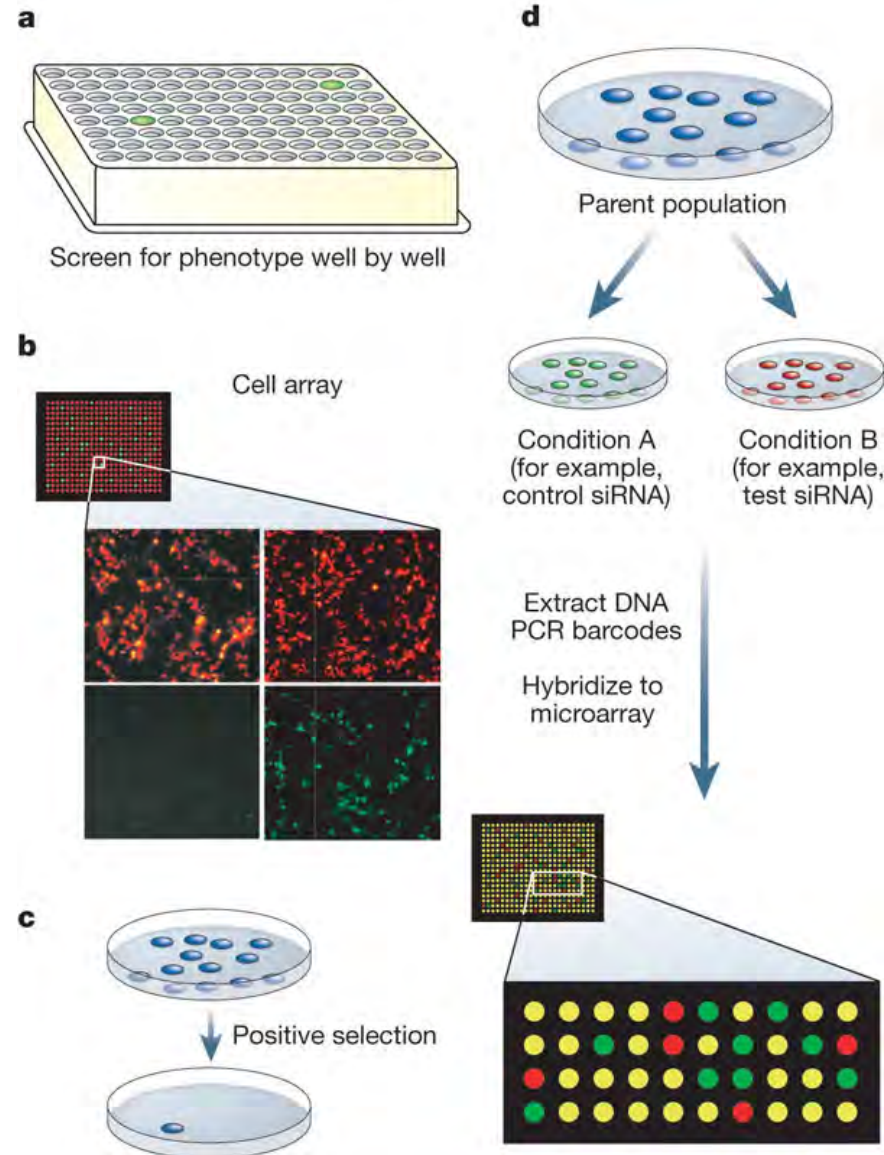


specific expression of Cre recombinase  
in Purkinje cells of cerebellum

# Using RNA interference to disrupt gene function

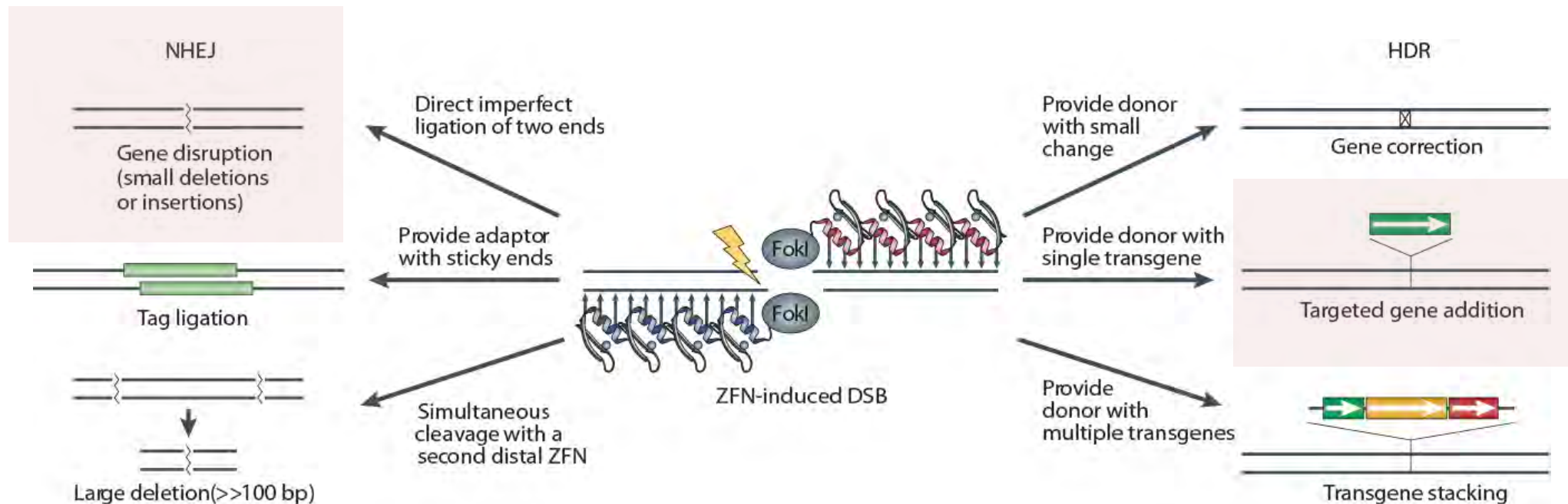


# RNAi screens to discover new genes

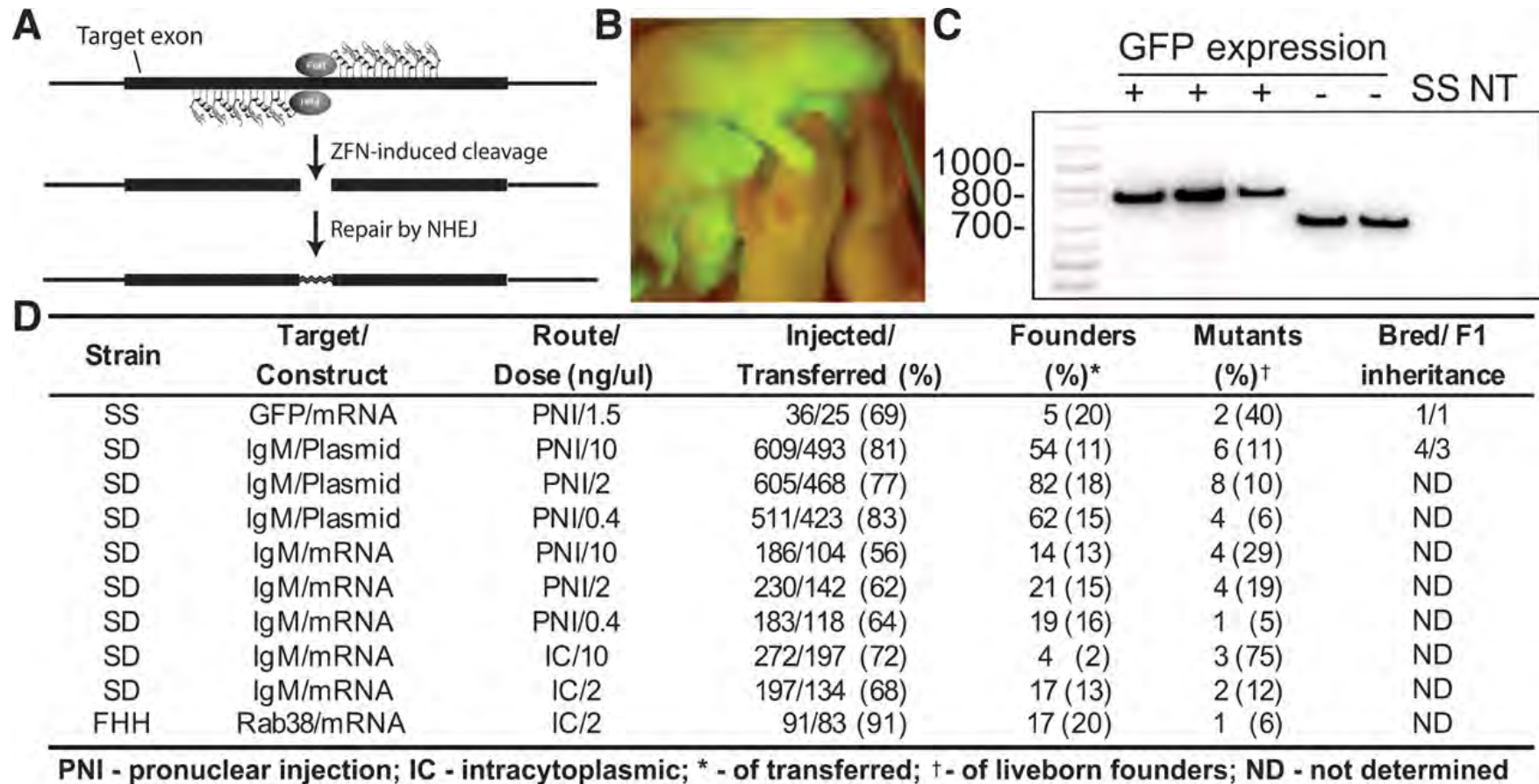


# Genome editing with engineered zinc finger nucleases

*Fyodor D. Urnov, Edward J. Rebar, Michael C. Holmes, H. Steve Zhang and Philip D. Gregory*



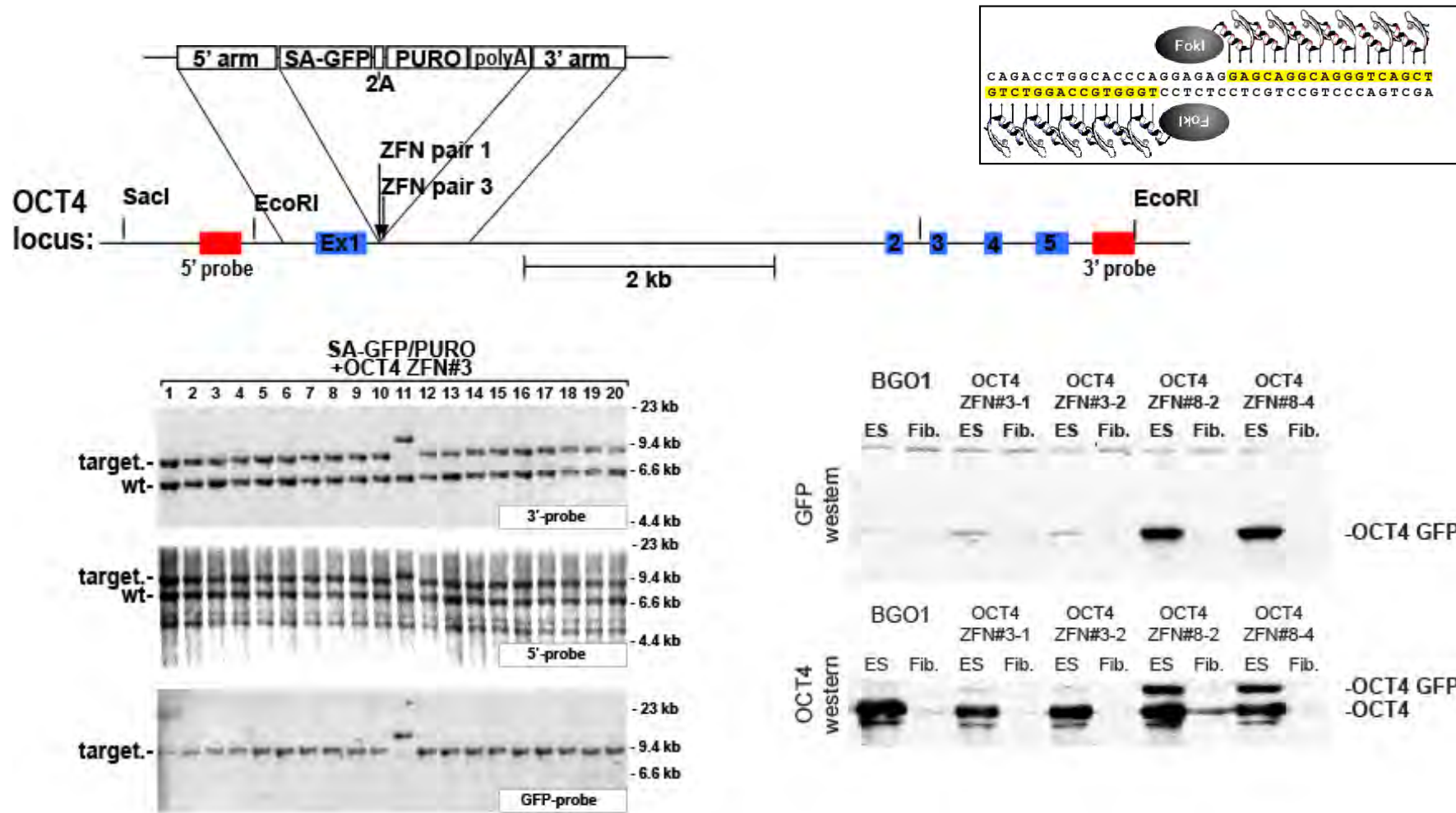
# ZFN-mediated gene disruption in rat embryos



A M Geurts et al. Science 2009;325:433-433

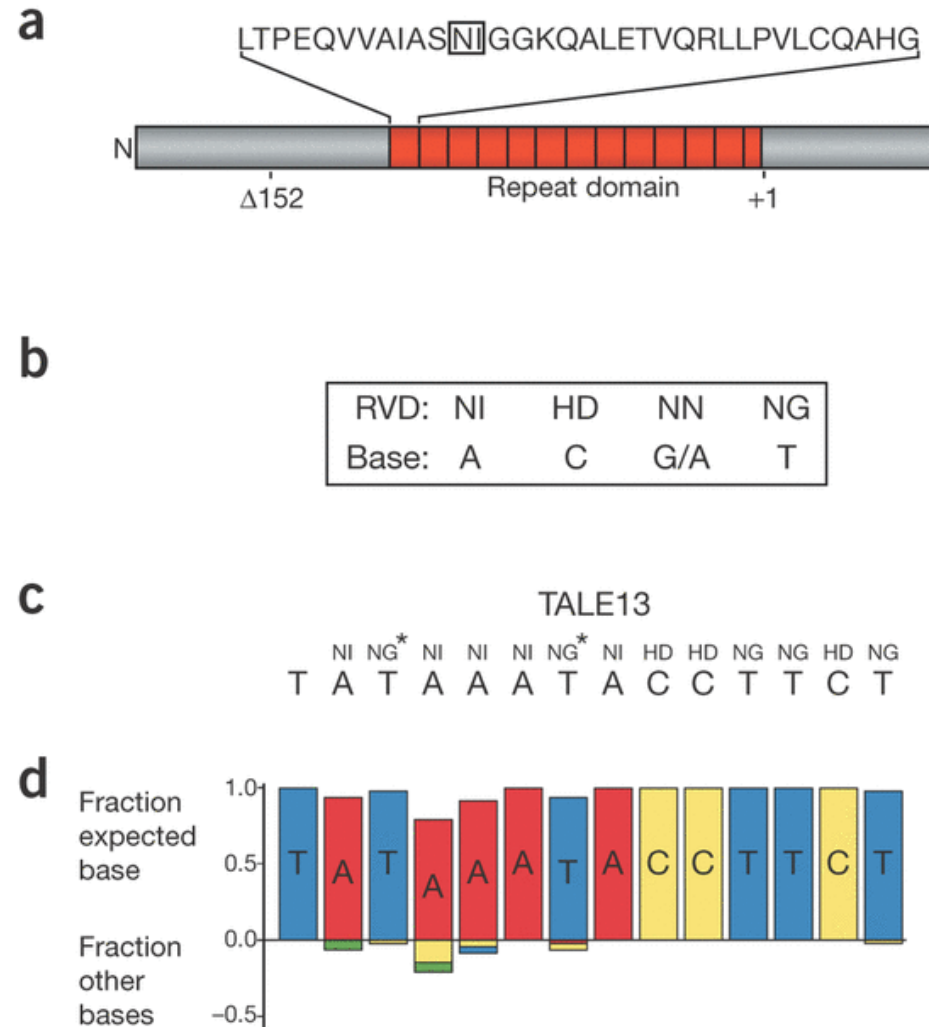
ZFN targeted against GFP transgene in transgenic rats

## Human ES cells and iPS cells: Endogenous Gene → Reporter at 95% Efficiency



# TALE: transcription activator–like effector proteins from *Xanthomonas*

- **TAL effectors:** virulence factors produced by a genus of plant pathogens, *Xanthomonas* spp.
- bind to specific host promoter sequences
- contain 17–18 repeats of 34 amino acids.
- DNA binding specificity determined by positions 12 and 13



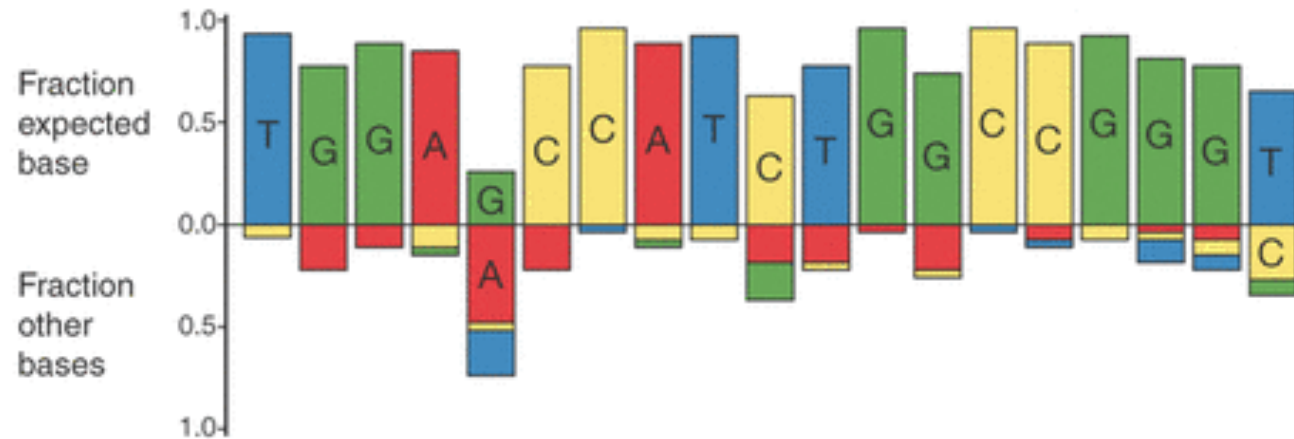
# Engineering TALEs using a simple code

d

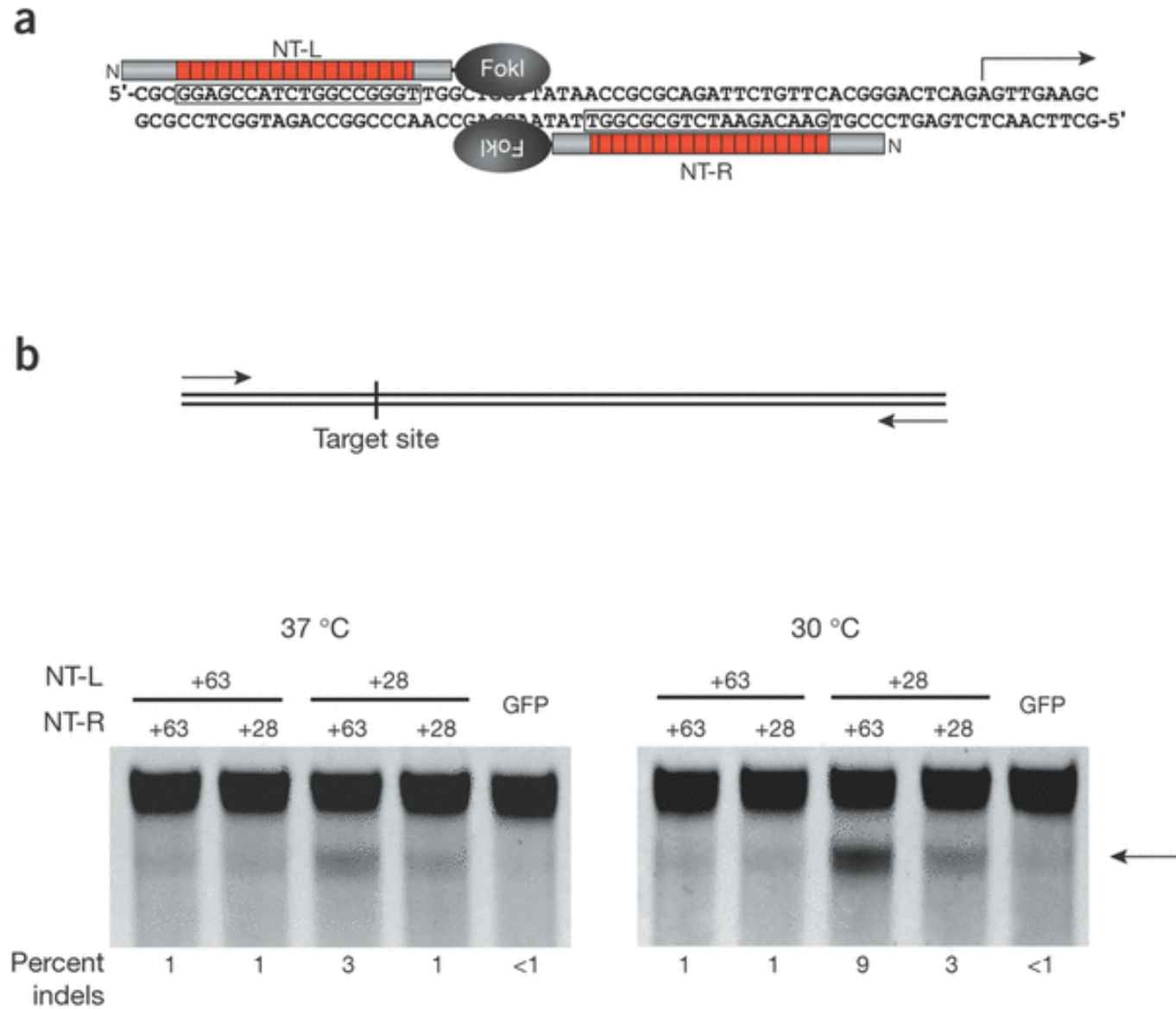
NT-L

	NN	NN	NN	NN	CC	CC	NN	NG	CC	NG	NN	NK	CC	CC	NN	NK	NN	NG
T	G	G	A	G	C	C	A	T	C	T	G	G	C	C	G	G	G	T

e



# TAL effector nucleases (TALENs)



- FokI endonuclease is an obligate dimer

# TAL effector nucleases (TALENs) induce indels

*hey2*:

#1297/ #1257 Mutations in 12 of 110 sequences: ~11%

GC TCTTCCGTTTCCACATCC ACCACATCCCAACAGAGC AGCGGGAGCAGCAGTAAACC WT

<-----GAGCAGCAGTAAACC Δ142

GCTCTTCCGTTTCCACATCCACC-----CAGCGGGAGCAGCAGTAAACC Δ14

GCTCTTCCGTTTCCACATCCACC-----ACAGCGGGAGCAACAGTAAACC Δ13

GCTCTTCCGTTTCCACATCCAC-----AGAGCAGCGGGAGCAGCAGTAAACC Δ11 [2x]

GCTCTTCCGTTTCCACATCCACCAC-----AGAGCAGCGGGAGCAGCAGTAAACC Δ8 [3x]

GCTCTTCCGTTTCCACATCCACCACAT-----tGAGCAGCGGGAGCAACAGTAAACC Δ6 (Δ7 and +1)

GCTCTTCCGTTTCCACATCCACCACATC--AACAGAGCAGCGGGAGCAGCAGTAAACC Δ2

GCTCTTCCGTTTCCACATCCACCACATaaaccaccacACAGAGCAGCGGGAGCAGCAG +6 (Δ4 and +10)

ACCTTCCCTCTATCATT<----- / /----->TCTGGAAGAAAAGAAA Δ303

*gria3a*:

#1258/ #1260 Mutations in 13 of 89 sequences: ~15%

GGAG TCGTCCAATAGCTTCT CAGTCACGCACGCCTGT GAGTTTCTGCTCTTTA TCTT WT

GGAGTCGTCCAATAGCTTC-----GCCTGTGAGTTTCTGCTCTTTATCTT Δ12

GGAGTCGTCCAATAGCTTCTCA-----GCCTGTGAGTTTCTGCTCTTTATCTT Δ9

GGAGTCGTCCAATAGCTTCTCAGT-----CTGTGAGTTTCTGCTCTTTATCTT Δ9

GGAGTCGTCCAATAGCTTCTCAG-----CCTGTGAGTTTCTGCTCTTTATCTT Δ9 [2x]

GGAGTCGTCCAATAGCTTCTCAGTCA-----GCCTGTGAGTTTCTGCTCTTTATCTT Δ5

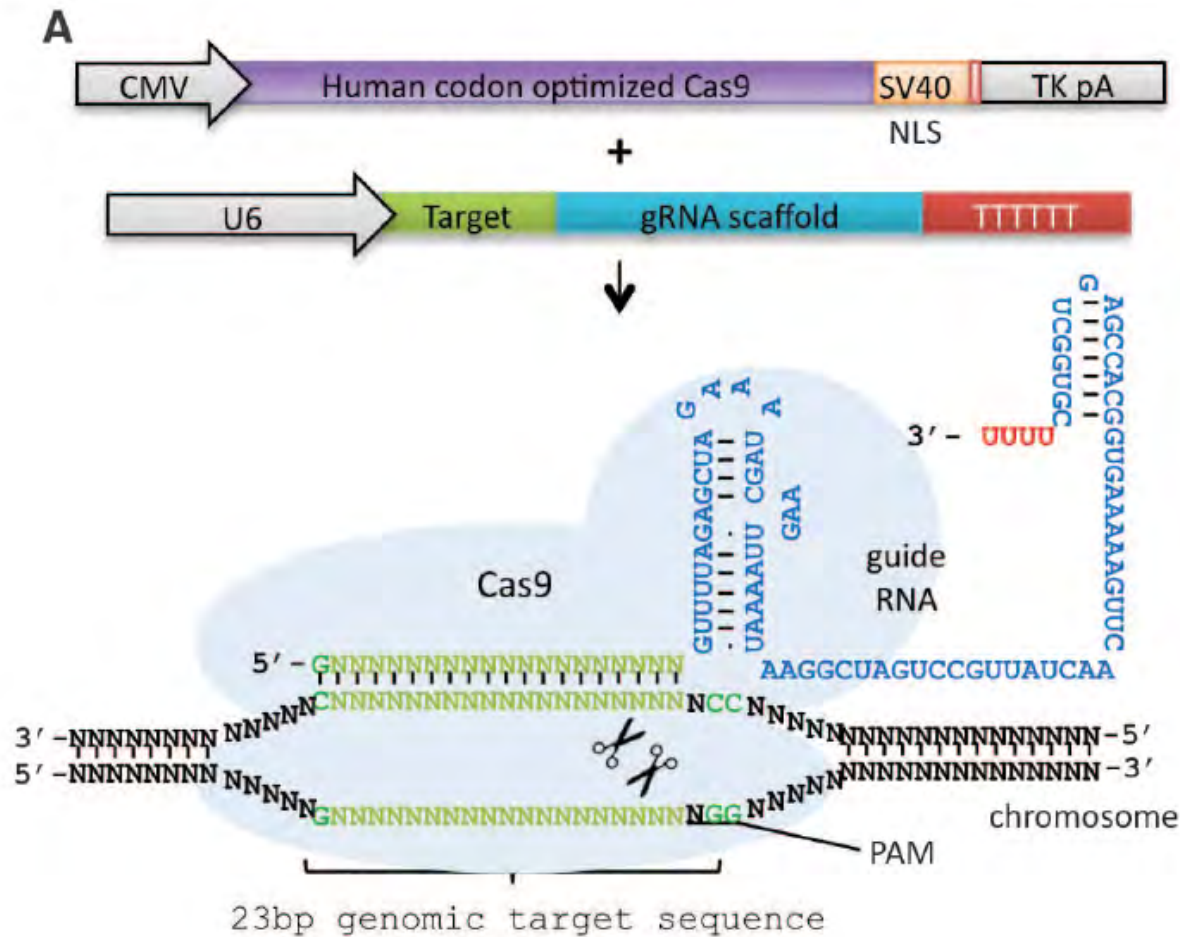
GGAGTCGTCCAATAGCTTCTCAGTCAgaaa--CCTGTGAGTTTCTGCTCTTTATCTT Δ2 (Δ6 and +4)

GGAGTCGTCCAATAGCTTCTCAGTctcagtcGCCTGTGAGTTTCTGCTCTTTATCTT +0 (Δ5 and +5)

GGAGTCGTCCAATAGCTTCTCAGTcacgcACGCACGCCTGTGAGTTTCTGCTCTTTA +4 [3x]

GGAGTCGTCCAATAGCTTCTCAGctgtgcctgtaACGCCTGTGAGTTTCTGCTCTTT +5 (Δ6 and +11) [2x]

# CRISPR-mediated genome editing



- CRISPR: clustered regularly spaced short palindromic repeat system
- Found in archaea and bacteria
- Key components consists of Cas9 nuclease and guide RNA
- By engineering the guide RNA, Cas9 cleavage can be targeted to any desired site.

# Assigning gene function

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## B. Functional approaches

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