

# How mammals sense infection: from endotoxin to the Toll-like receptors

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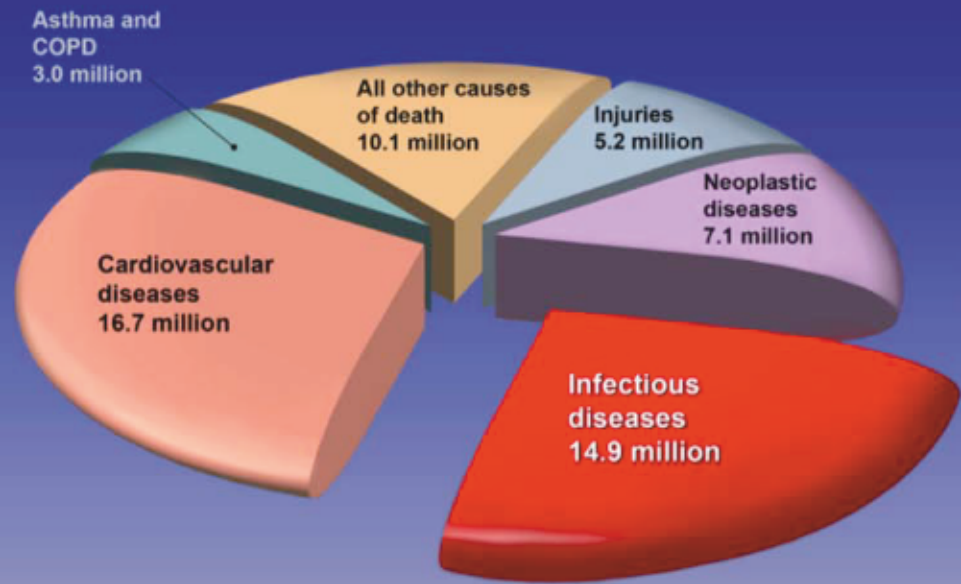
Mutagenetix

*A database of mutations and phenotypes induced by ENU*





Ernest Beutler, M.D., 1928-2008



Infectious Diseases	Annual deaths (millions)
Respiratory infections	3.96
HIV/AIDS	2.77
Diarrhoeal diseases	1.80
Tuberculosis	1.56
Vaccine-preventable childhood diseases	1.12
Malaria	1.27
STD's (other than HIV)	0.18
Meningitis	0.17
Hepatitis B and C	0.16
Tropical parasitic diseases	0.13
Dengue	0.02
Other infectious disease	1.76

Based on: The challenge of emerging and re-emerging infectious diseases. D.M. Morens, G.K. Folkers, and A. S. Fauci, Nature 463, 122 (7 January 2010)

# Infections and their transmissible character were known in antiquity...



460 - 370, BCE

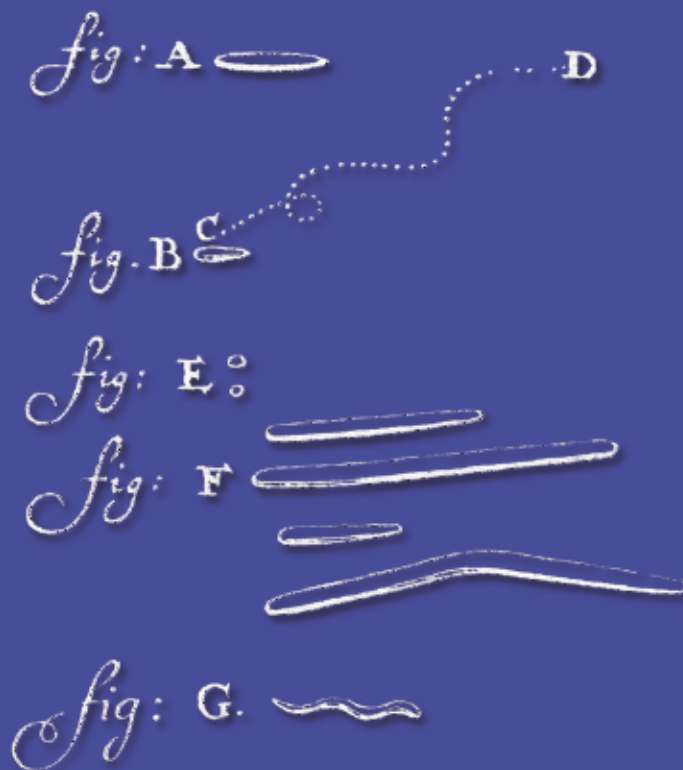
129 - 199, AD

1135-1204, AD

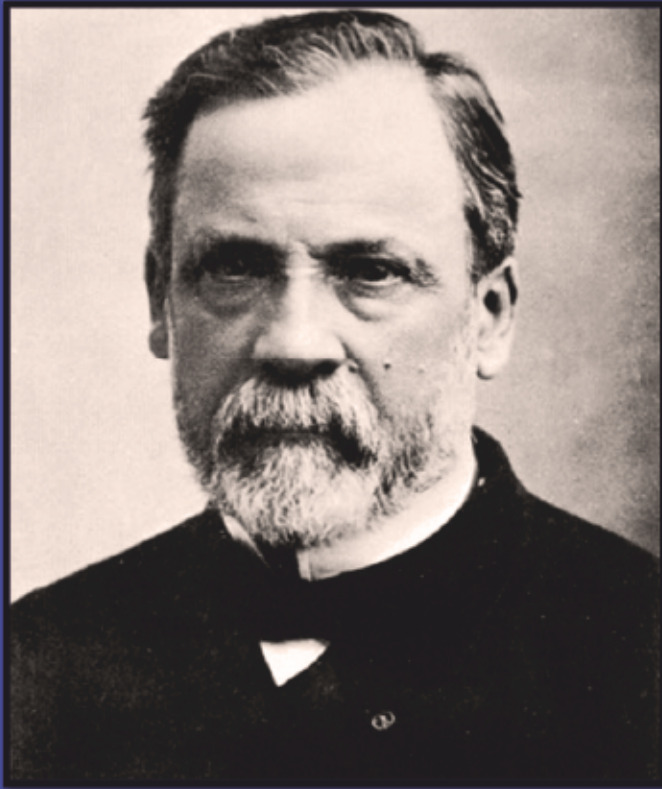
...while microbes were only discovered in  
the 17<sup>th</sup> century...



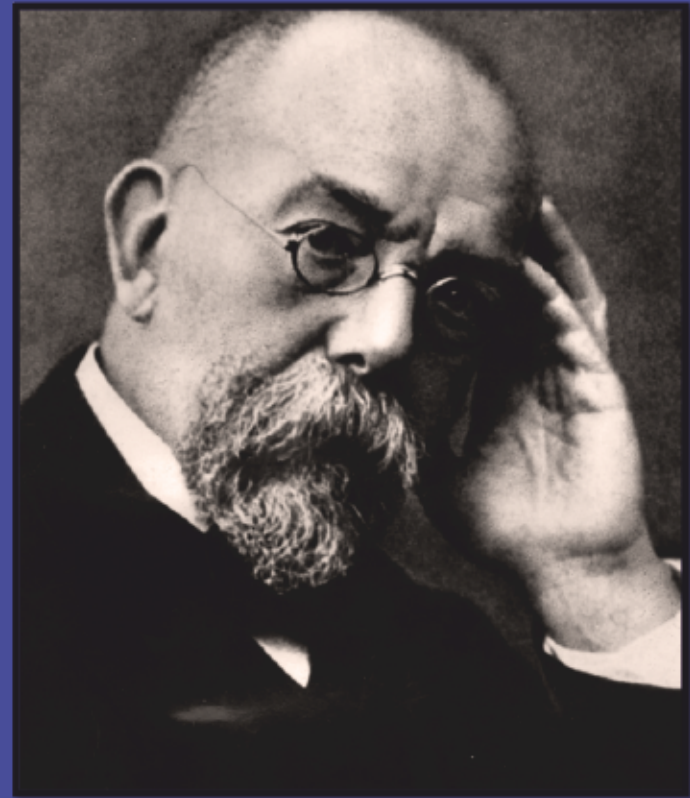
Antonie van Leeuwenhoek  
1632 - 1723



...and the association between microbes and infection was only discovered in the 19<sup>th</sup> century.

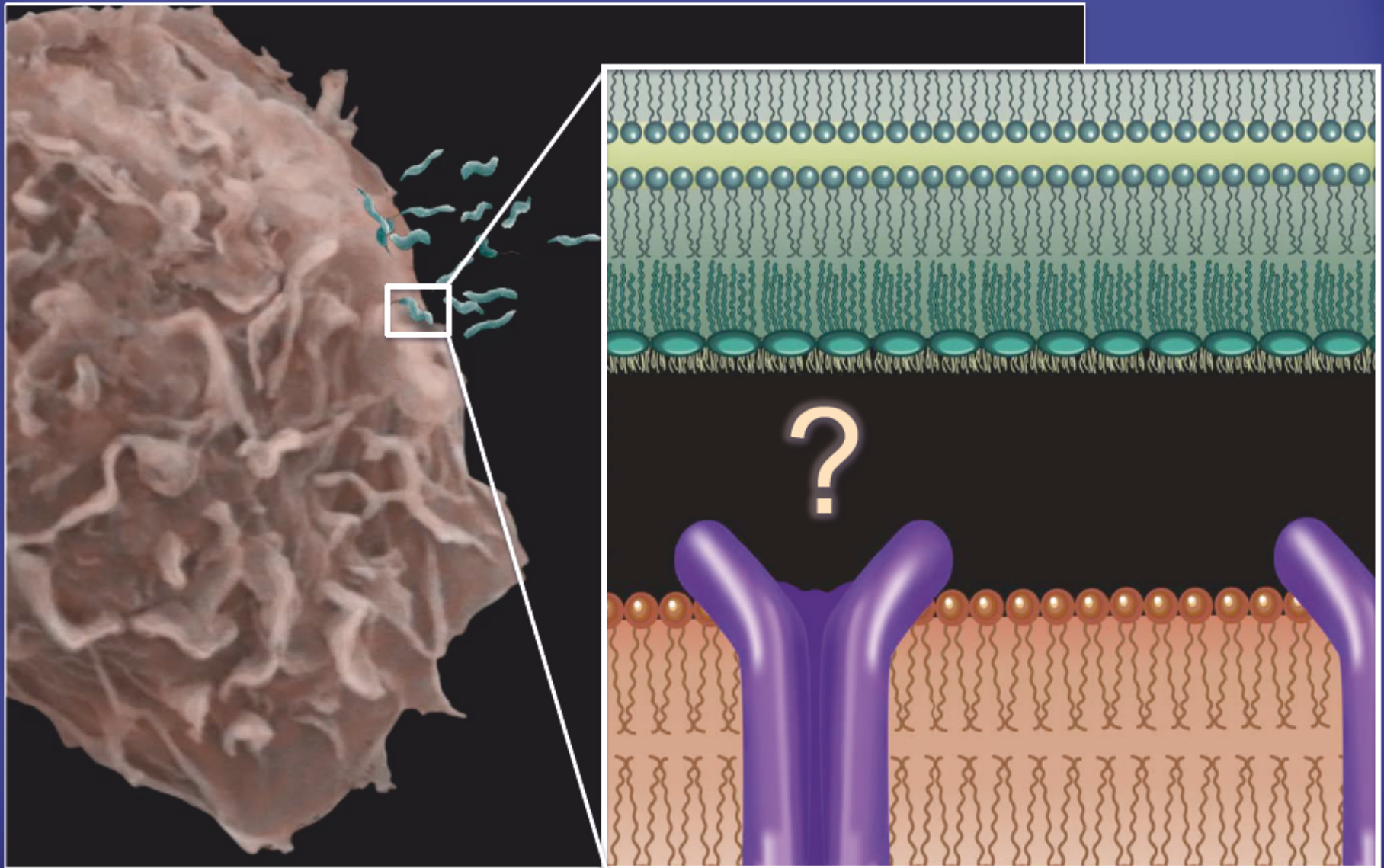


Louis Pasteur  
1822-1895

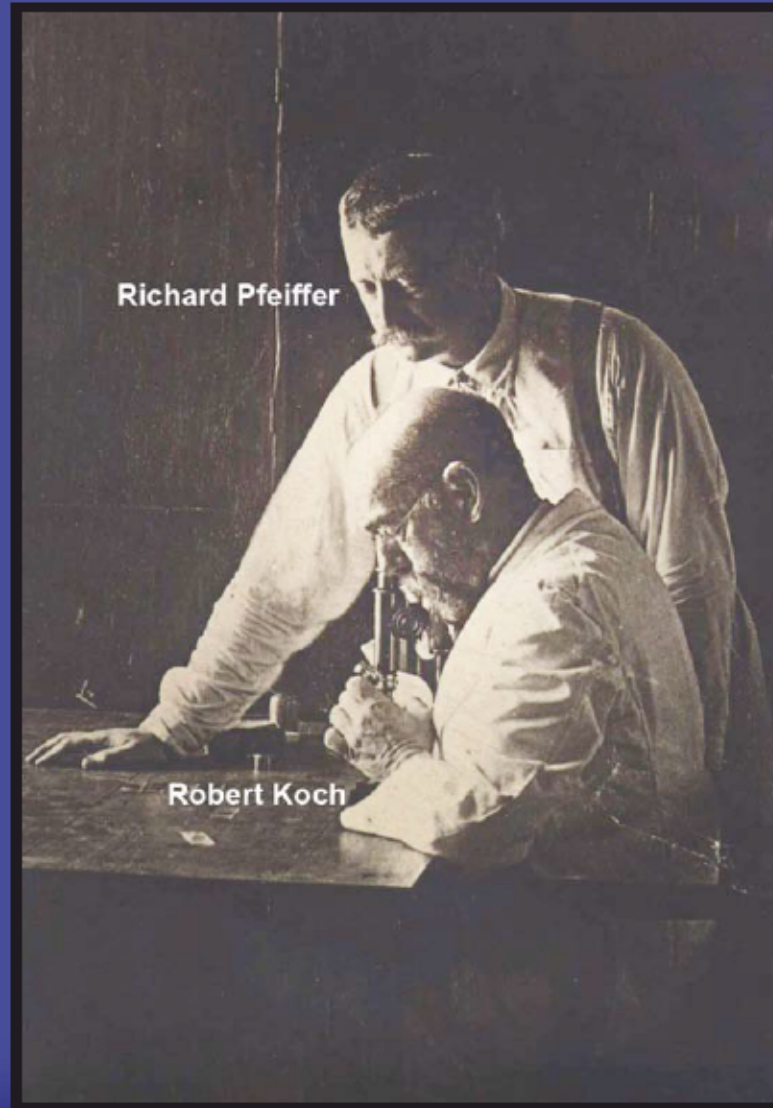


Robert Koch  
1843-1910

What might be the nature of contact between microbe and host?

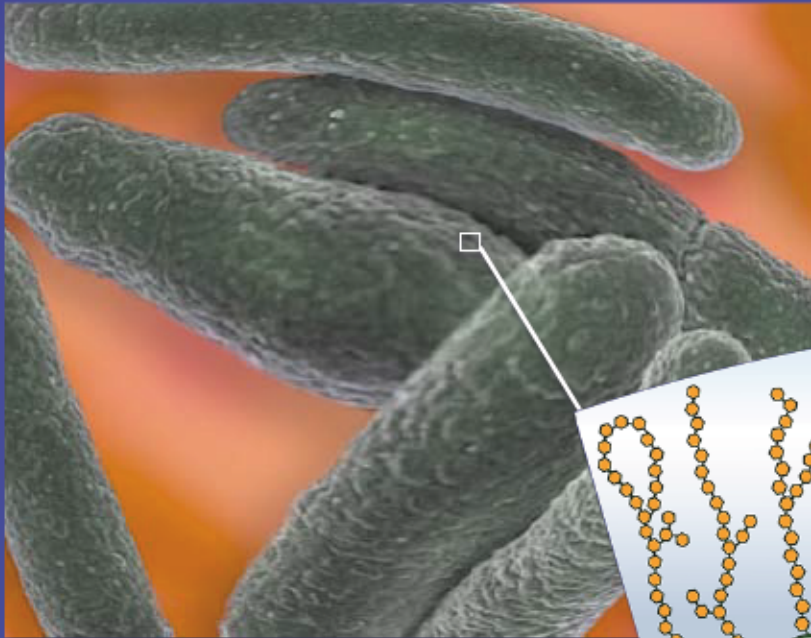


Soon after microbes were discovered, it was appreciated that mammals recognize them as foreign and mount an intense inflammatory response

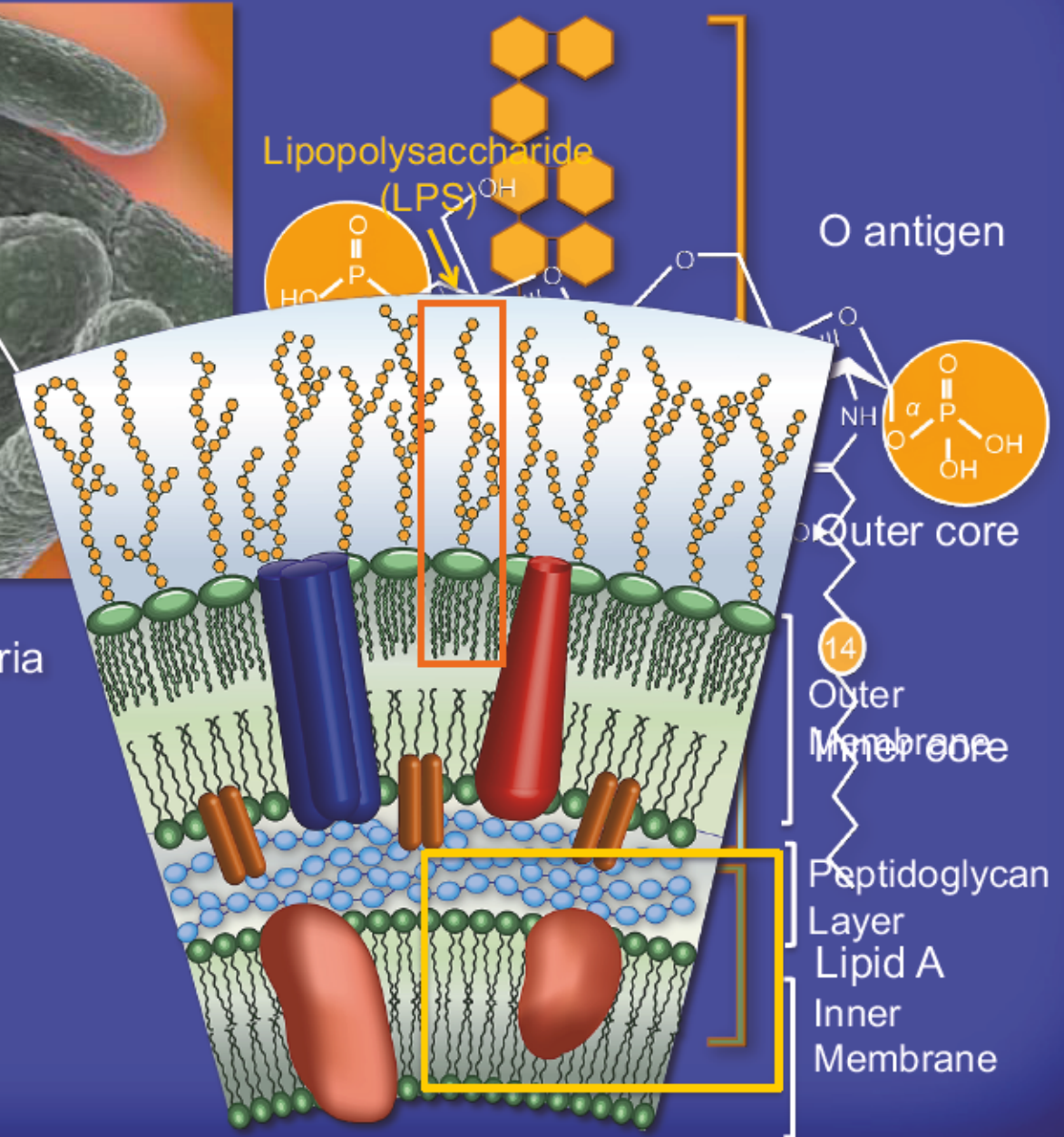


Richard Pfeiffer, a  
Robert Koch, noted  
microbes  
reaction in  
on after they  
into these  
oined the term  
describe the  
st-stable  
iated with  
nsible for fever,  
hock and  
th.

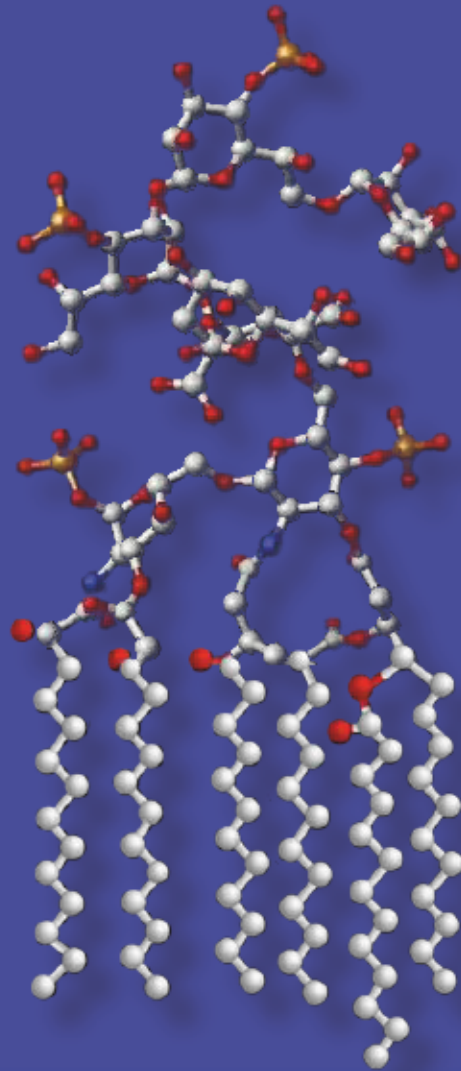




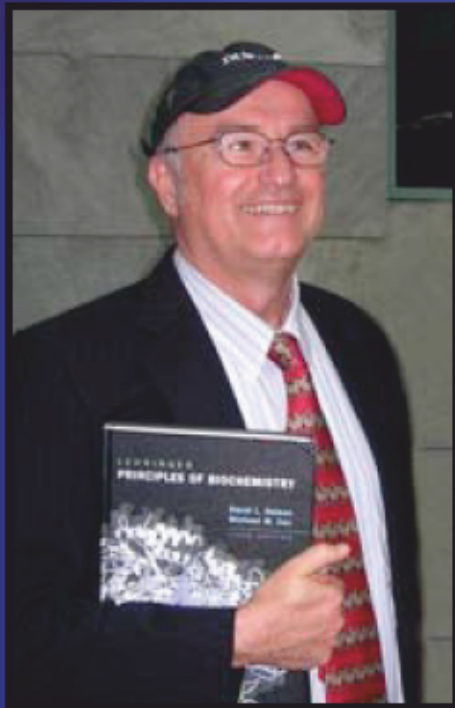
Gram negative bacteria  
*Escherichia coli*



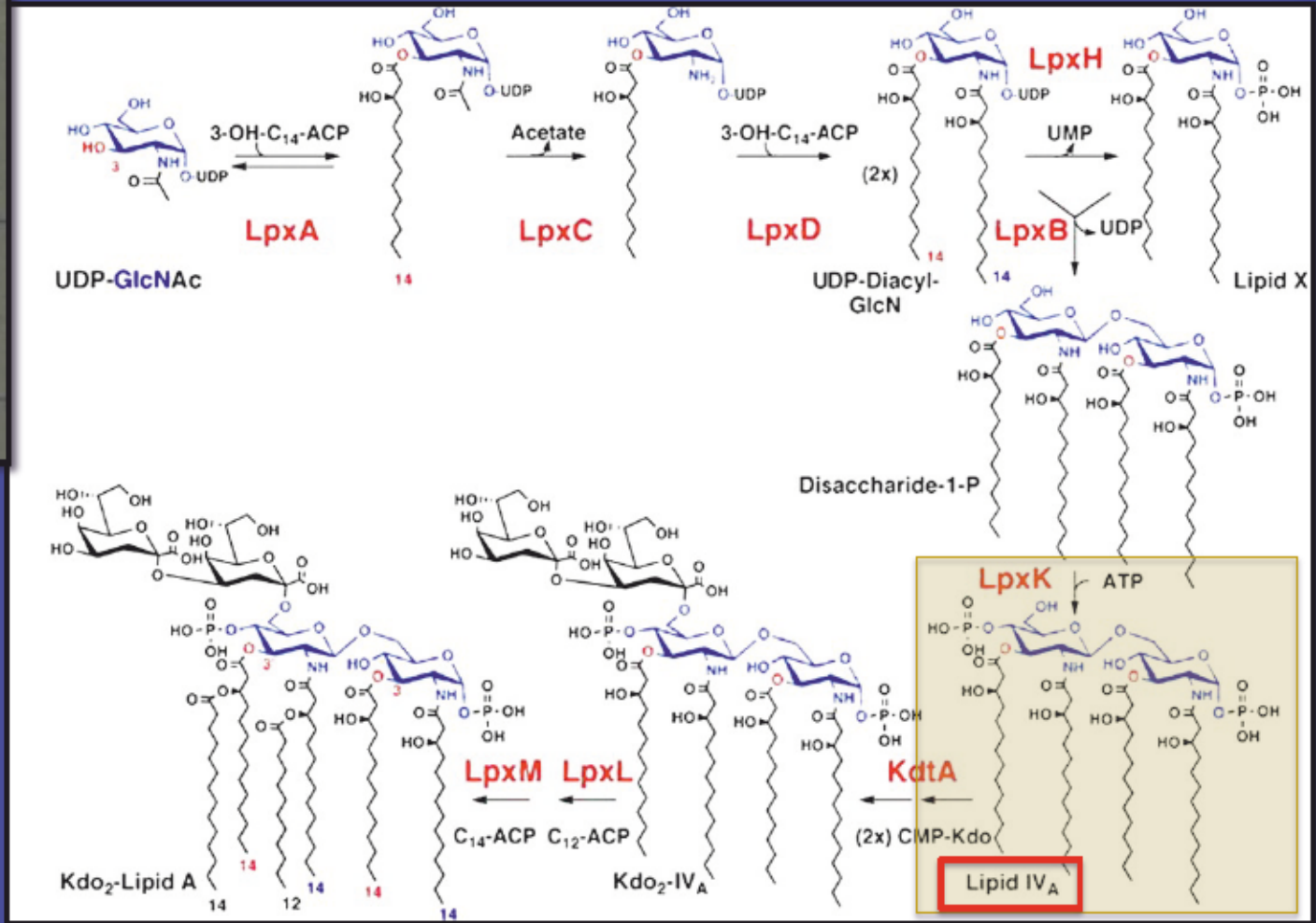
# LPS



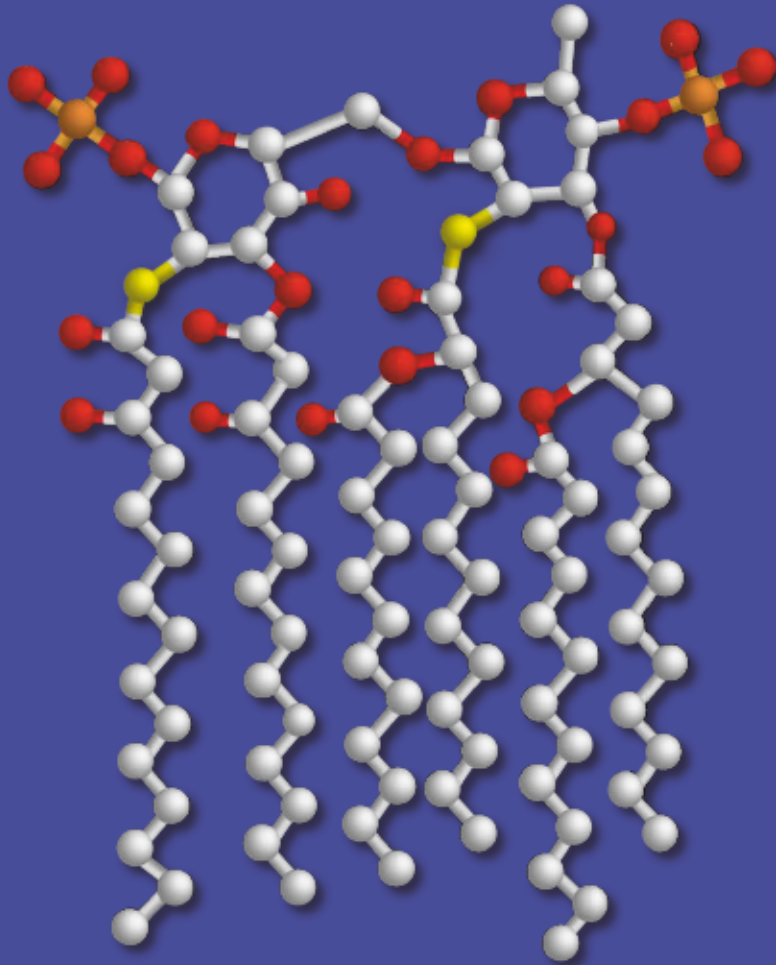
# Biosynthesis of Lipid A



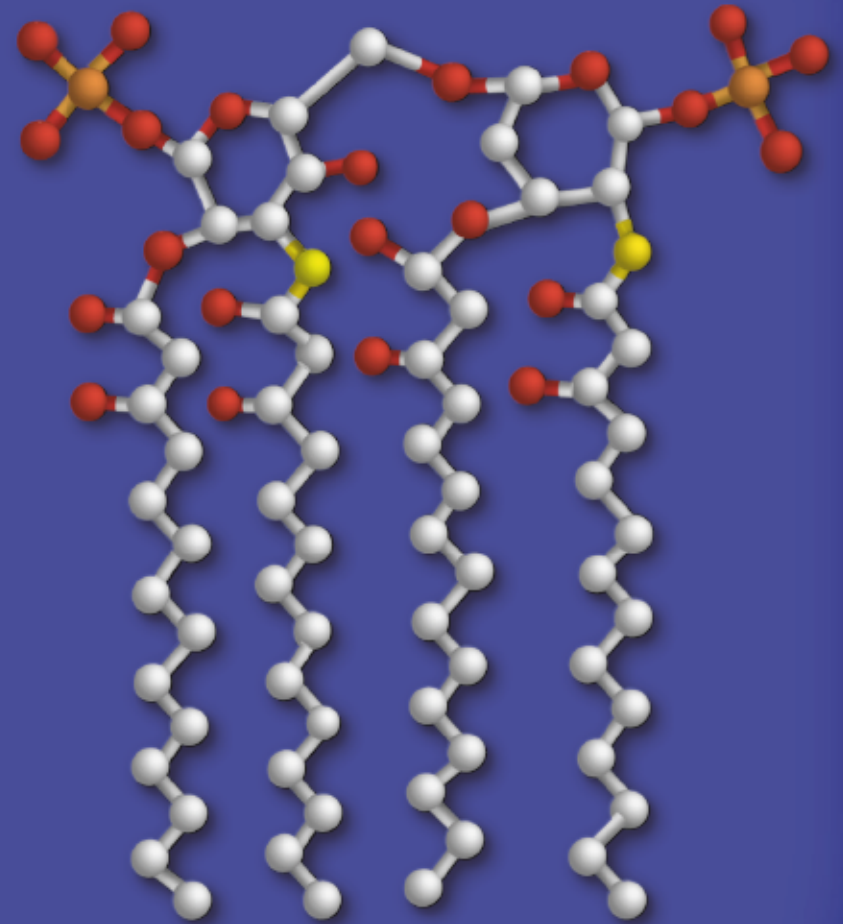
Christian R.H. Raetz  
1946 - 2011

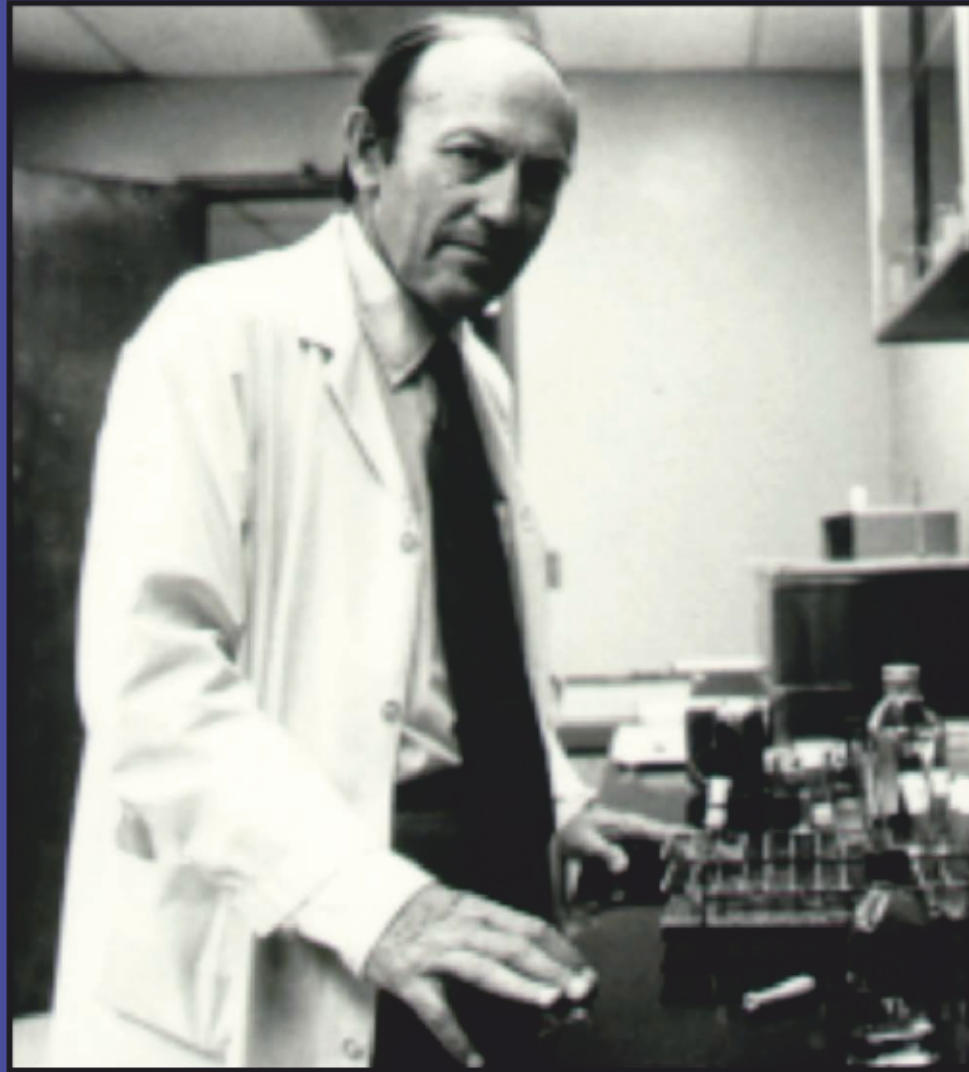


Lipid A



Lipid IVa

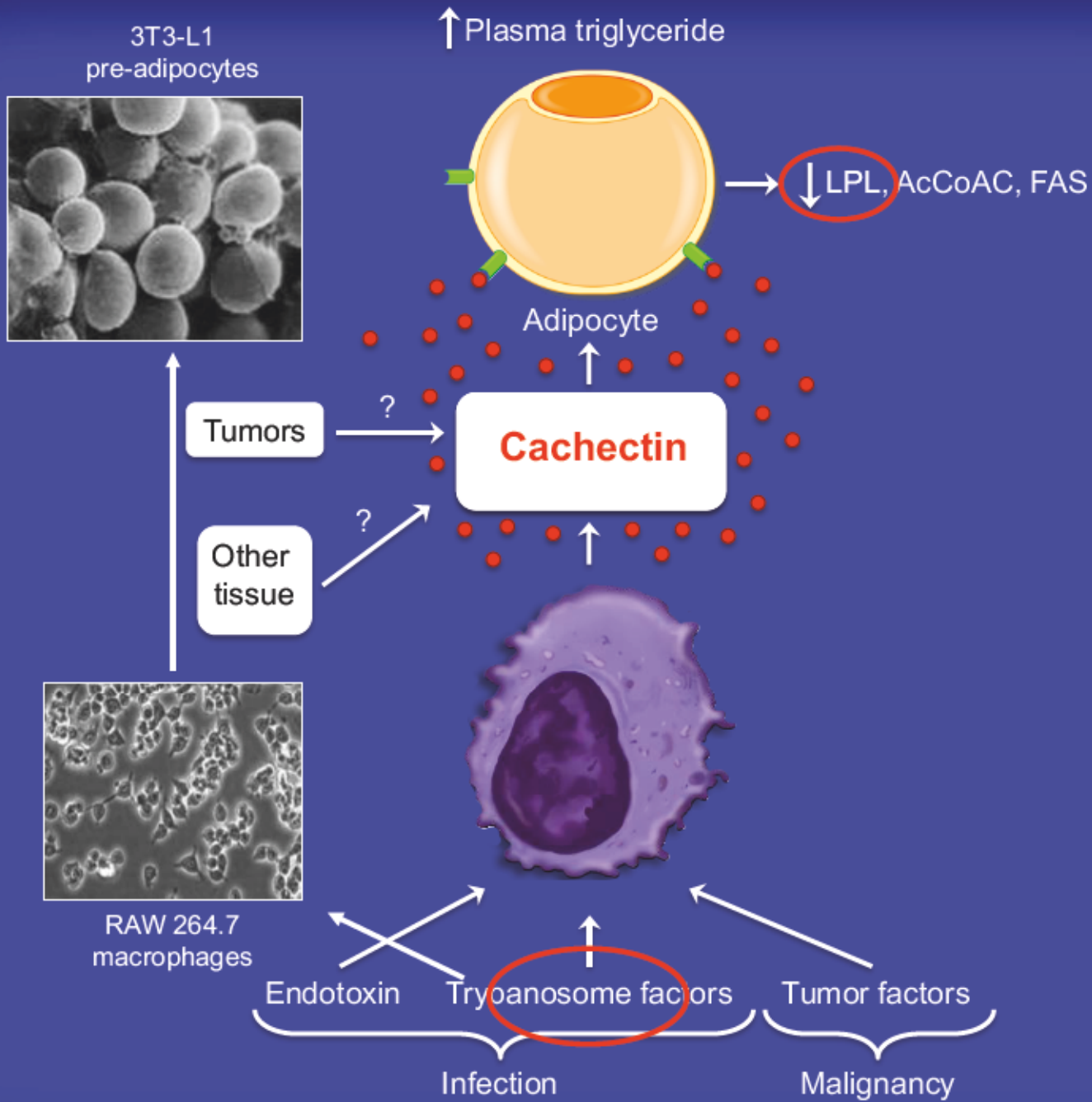




Abraham Braude, 1917-1984

Wasting disease (cachexia) in a cow with African trypanosomiasis



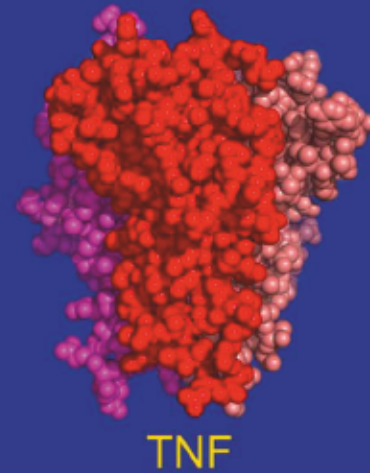
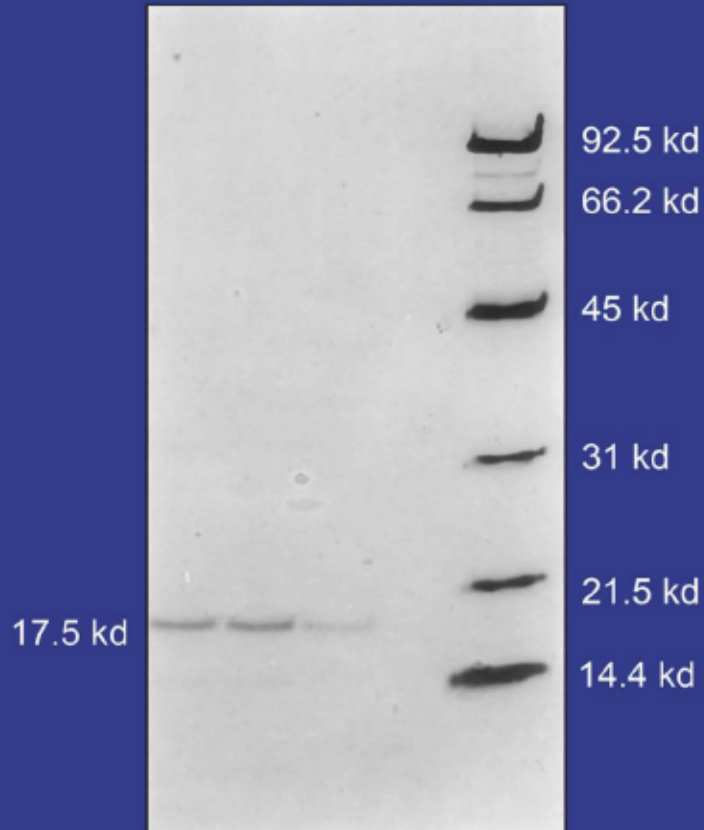


1983

# Isolation of mouse cachectin

- Pressure dialysis of medium from ~500 10 cm plates of LPS-activated RAW 264.7 cells (early harvest)
- ConA sepharose chromatography
- Isoelectric focusing in a glycerol gradient
- Preparative native gel electrophoresis
- Preparative SDS gel electrophoresis
- Yielded microgram quantities of an apparently pure 17.5 kD protein with approximately 2% yield of initial biological activity (prior to denaturing gel electrophoresis).
- Cachectin comprised 1-2% of the protein secreted by RAW 264.7 cells during the first two hours following LPS activation.





## Cachectin = Mouse tumor necrosis factor

(mouse CACH)

H<sub>2</sub>N LEU-ARG-SER-SER-SER-GLU-ASN-SER-SER-ASP-PRO-PRO-VAL-ALA- ? -VAL-VAL-ALA-ASN...

H<sub>2</sub>N VAL-ARG-SER-SER-SER-ARG-THR-PRO-SER-ASP-LYS-PRO-VAL-ALA-HIS-VAL-VAL-ALA-ASN...

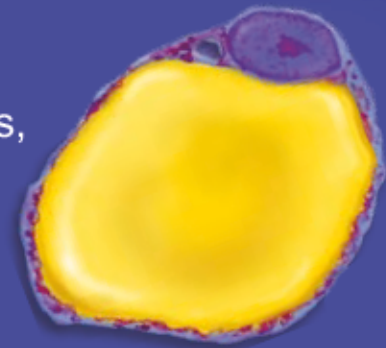
(human TNF)

1 µg of cachectin had 10<sup>8</sup> U of TNF activity

This raised the question:  
might TNF mediate *all*  
effects of LPS, including  
the lethal effect?

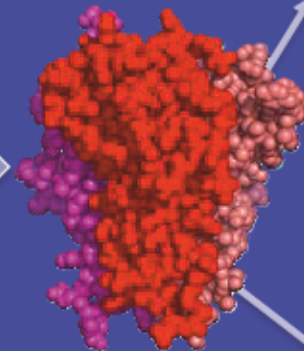
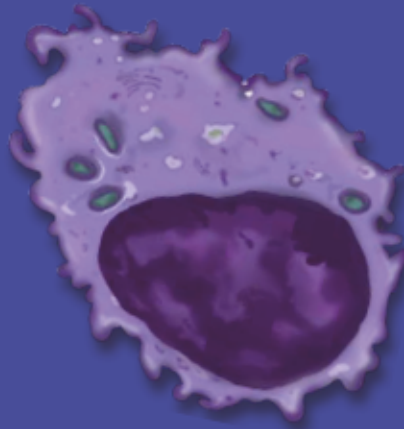
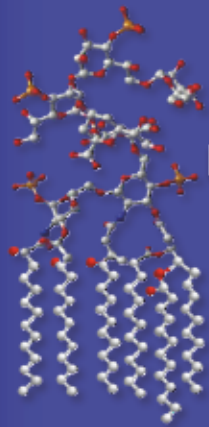
↓ Triglyceride synthesis,  
LPL, FAS

↑ AcCoA carboxylase,  
glycerol release



LPS

MACROPHAGE

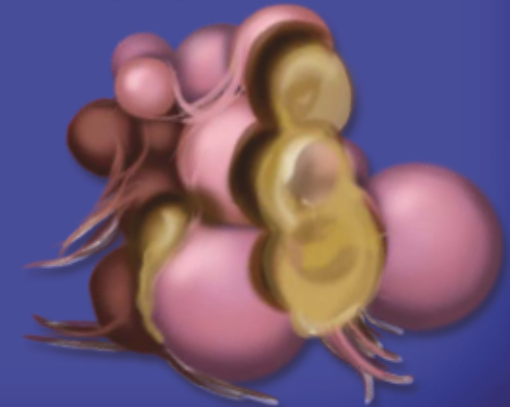


FAT

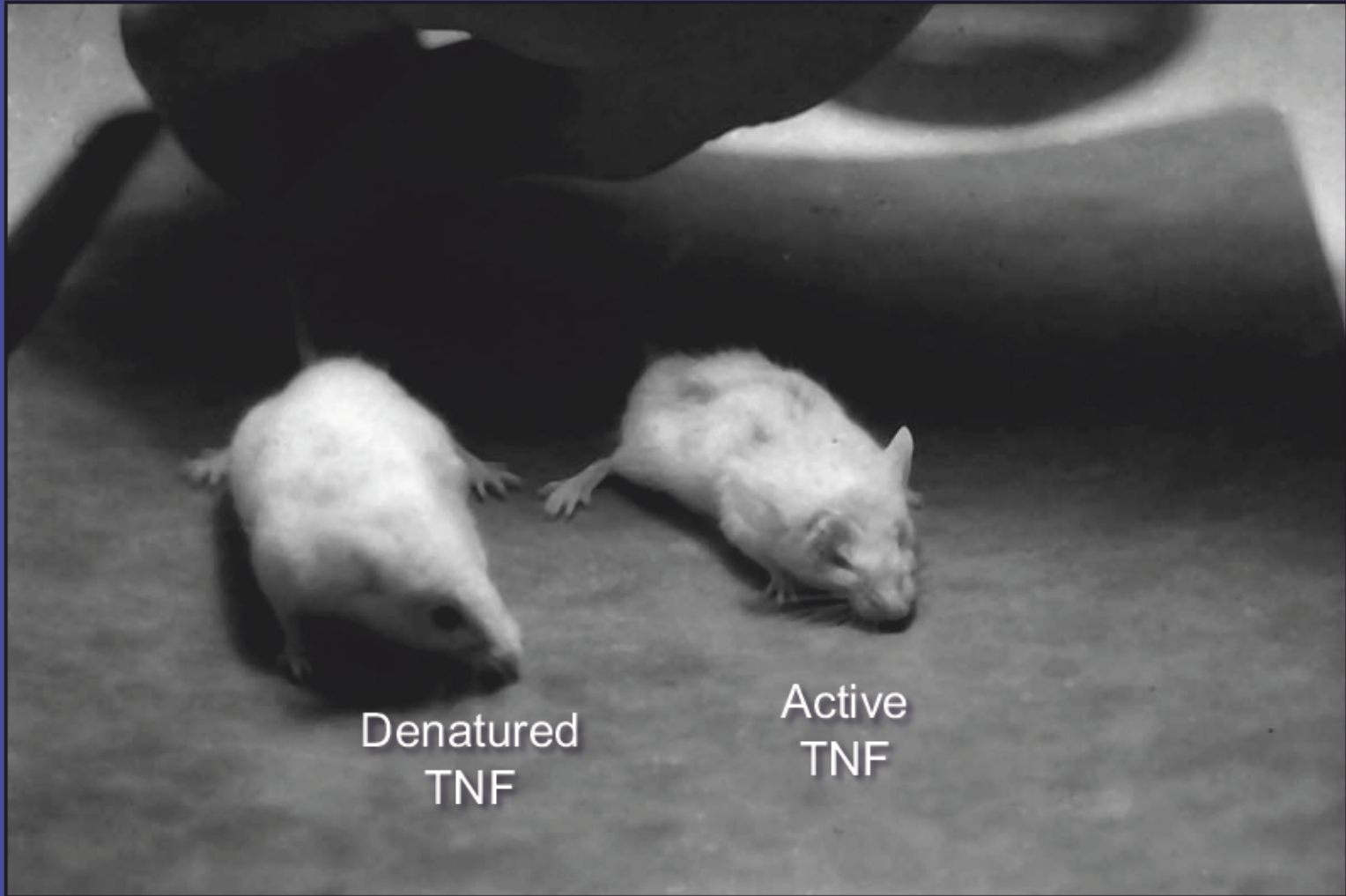
TNF

TUMOR

↑ Tumor necrosis  
and cytolysis

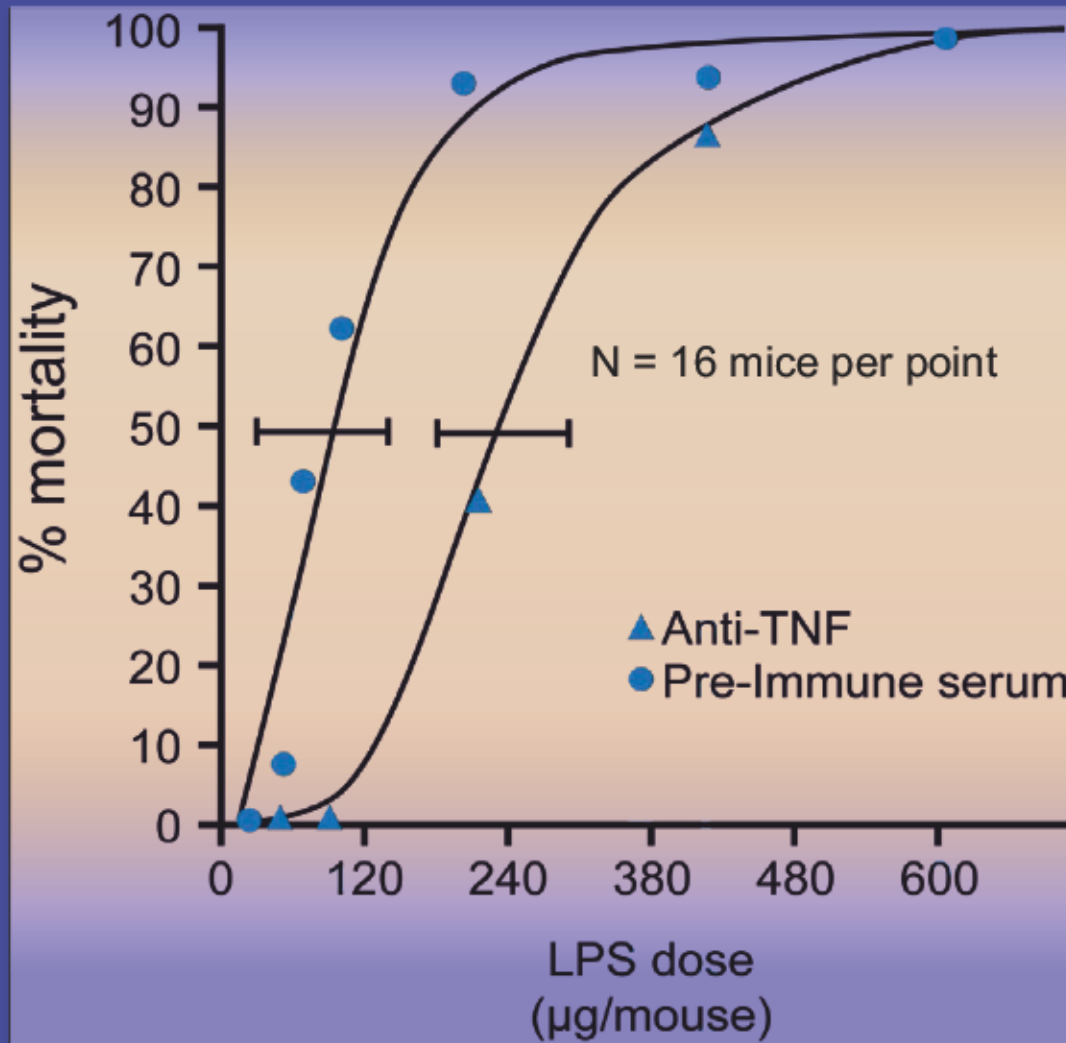


# Purified TNF mimics LPS toxicity

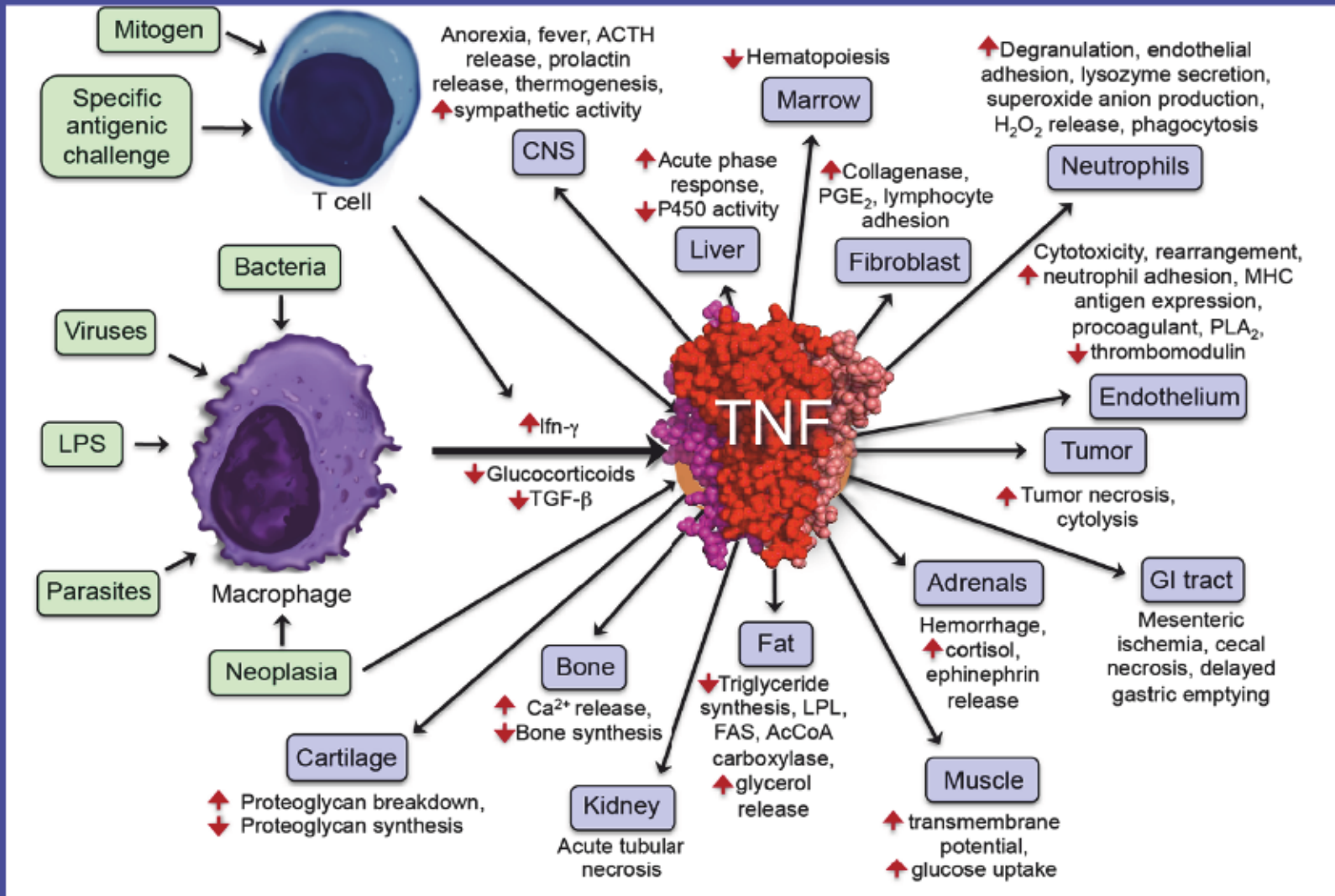


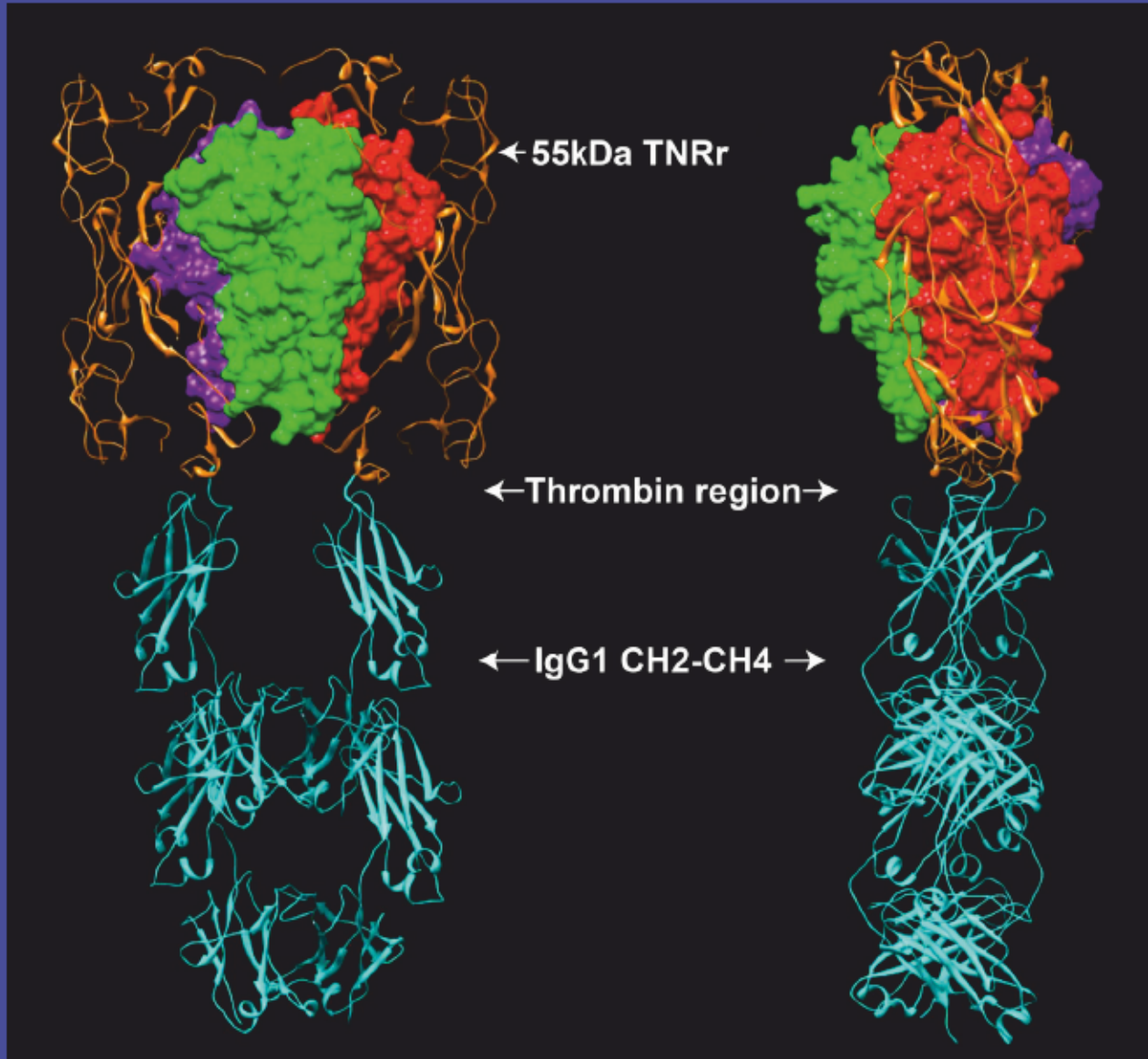
1984

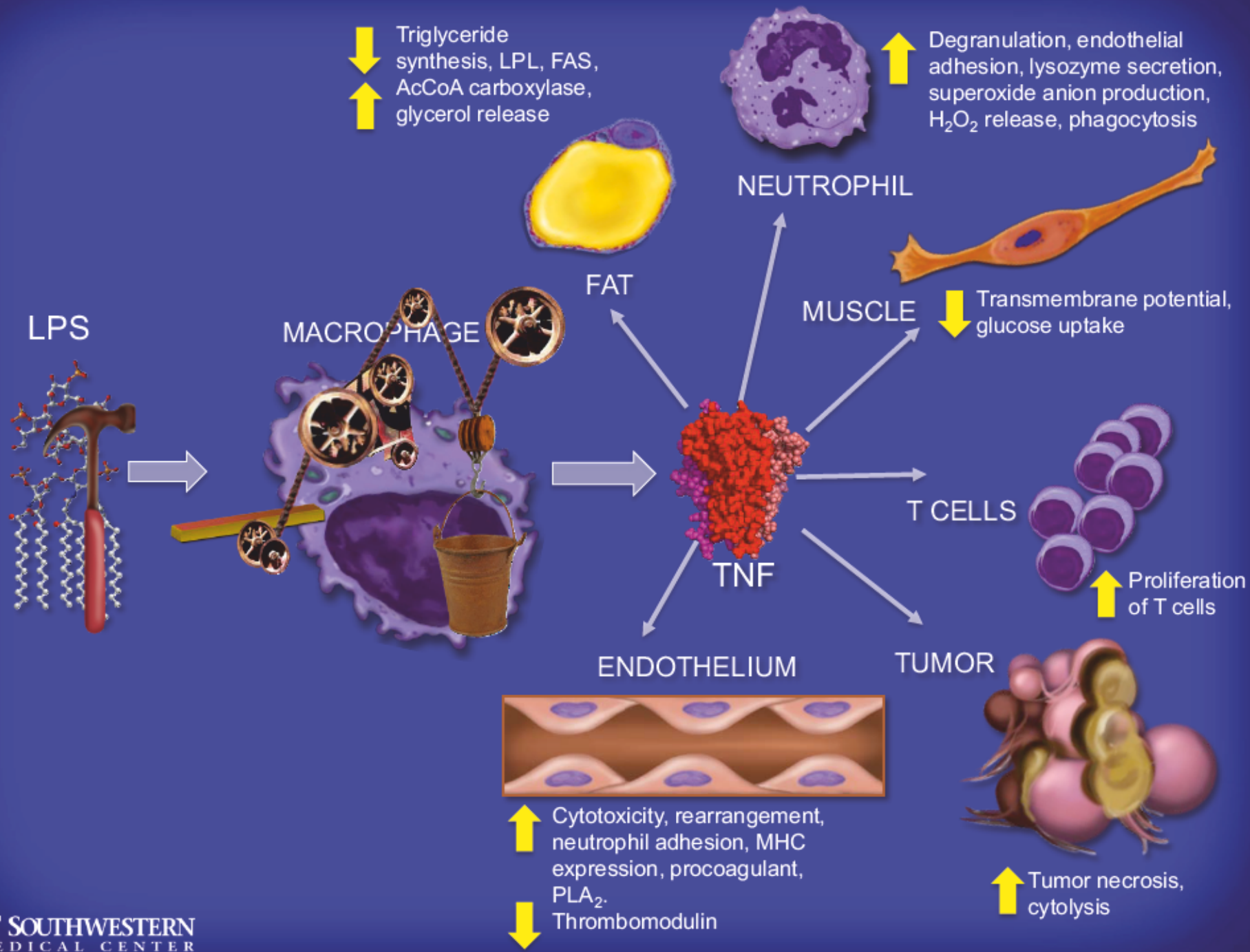
# The lethal effect of LPS is attenuated by passive immunization against TNF



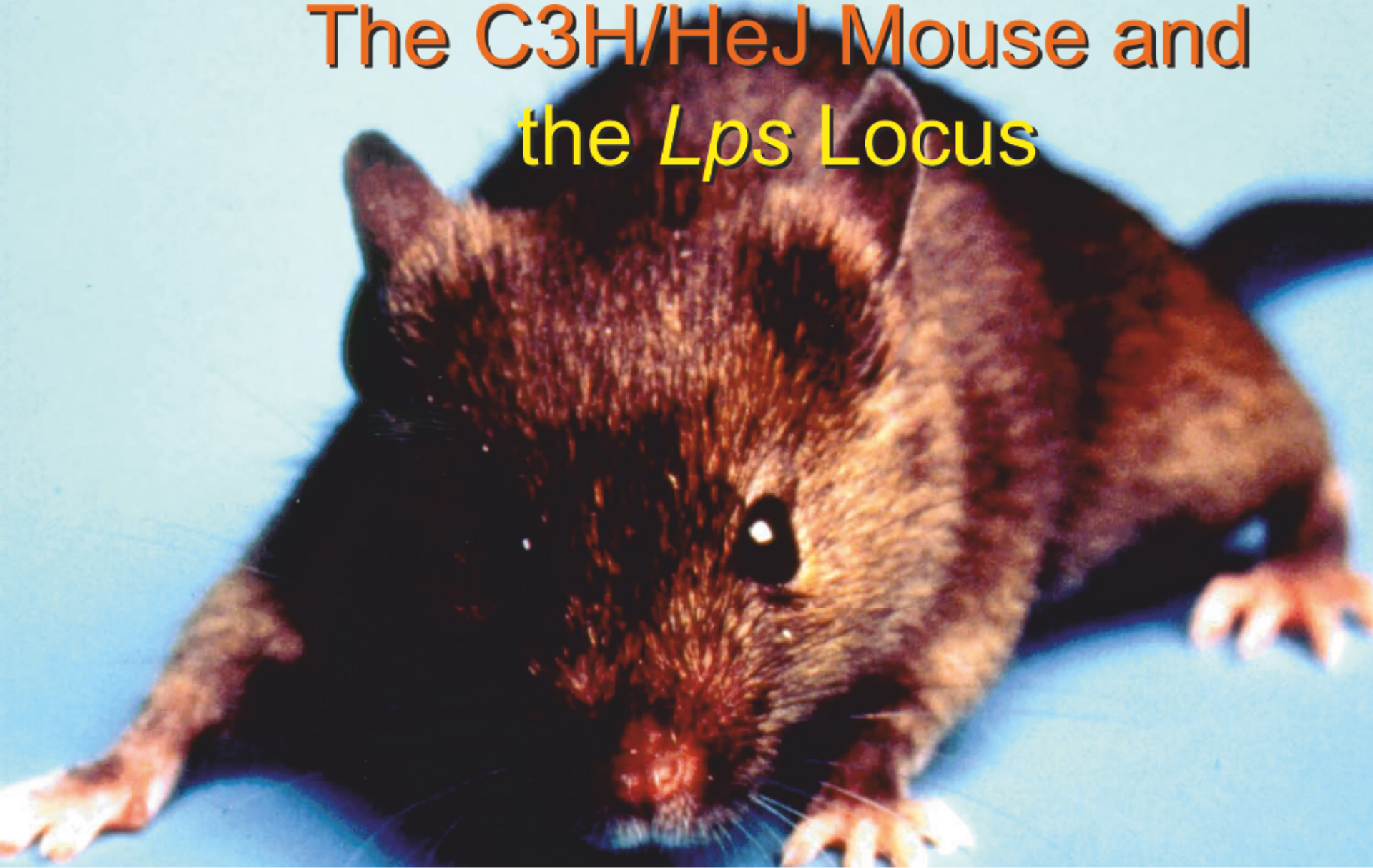
1985







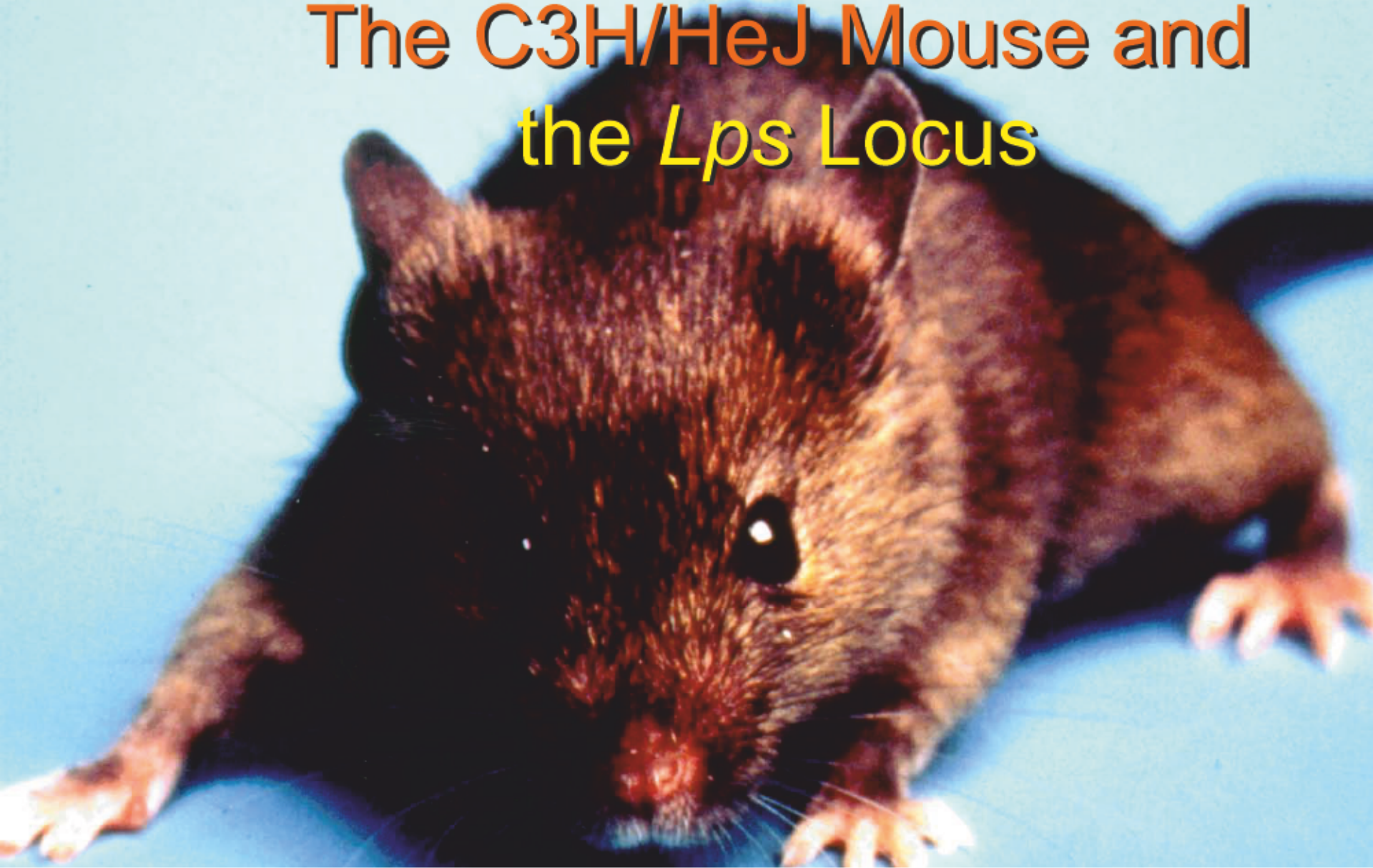
# The C3H/HeJ Mouse and the *Lps* Locus



- Resistant to LPS (Heppner and Weiss, 1965)

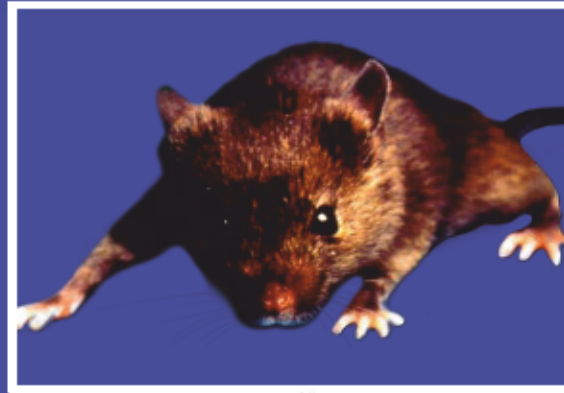


# The C3H/HeJ Mouse and the *Lps* Locus

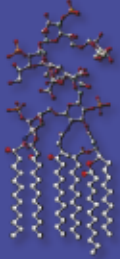


- Fail to make a cytokine response to LPS (for example, no TNF), suggesting a proximal defect.

C3H/HeJ mice:  
resistant to LPS  
(and *only* LPS)



LPS



Gram negative bacteria

Nucleic Acids



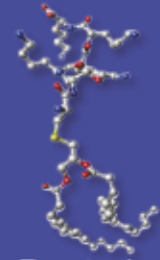
Viruses

Flagellin

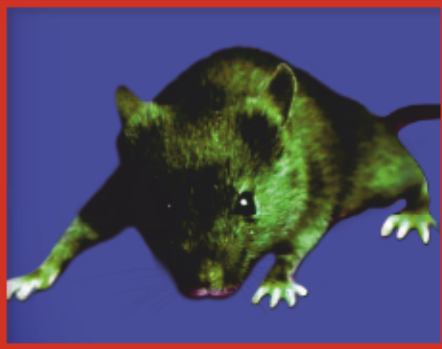


Bacteria

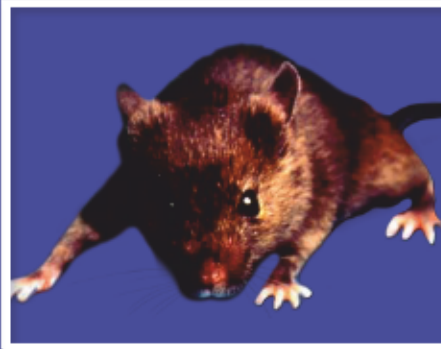
Lipoproteins



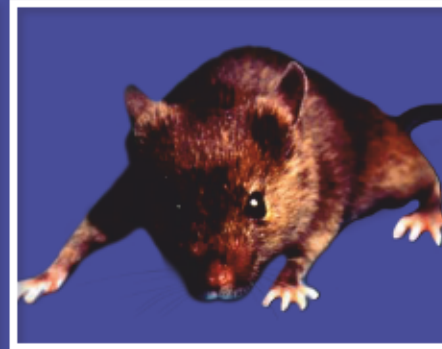
Bacteria



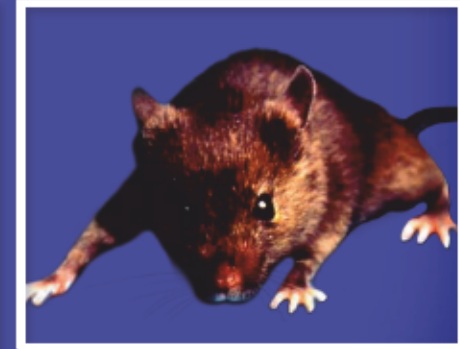
Unresponsive



Responsive

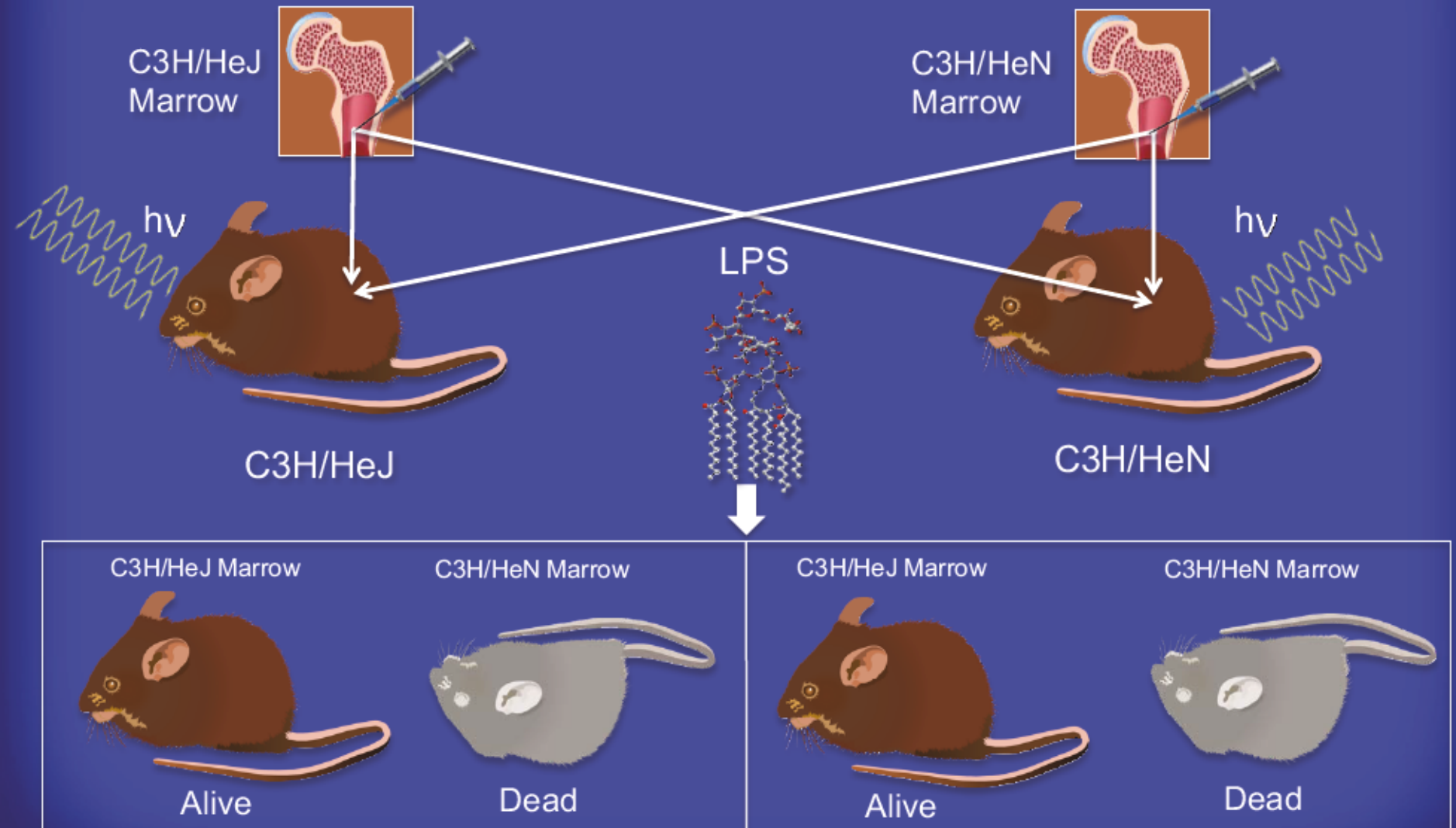


Responsive

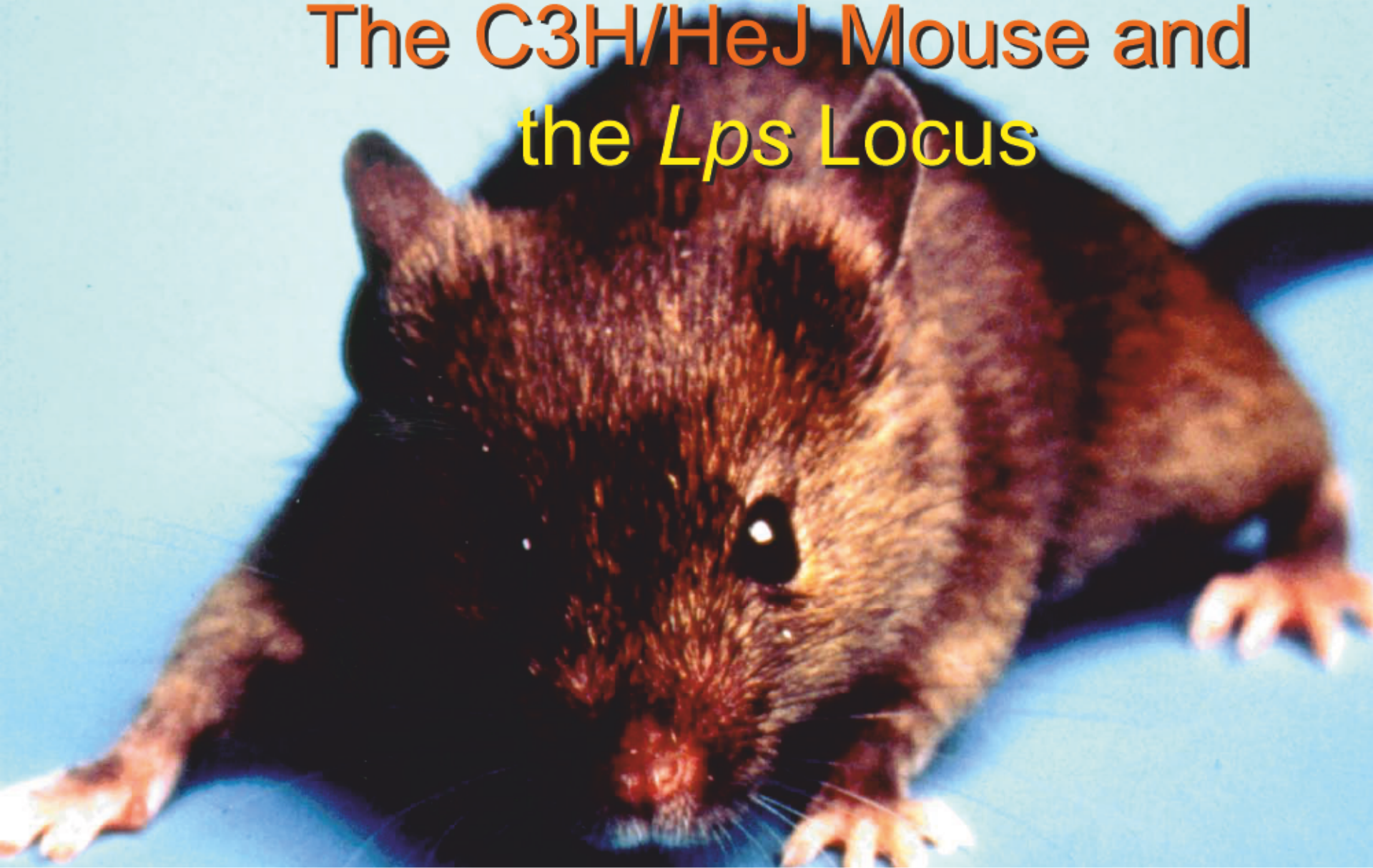


Responsive

# C3H/HeJ bone marrow transfer to C3H/HeN and vice versa: susceptibility to LPS-induced lethality is determined by the donor

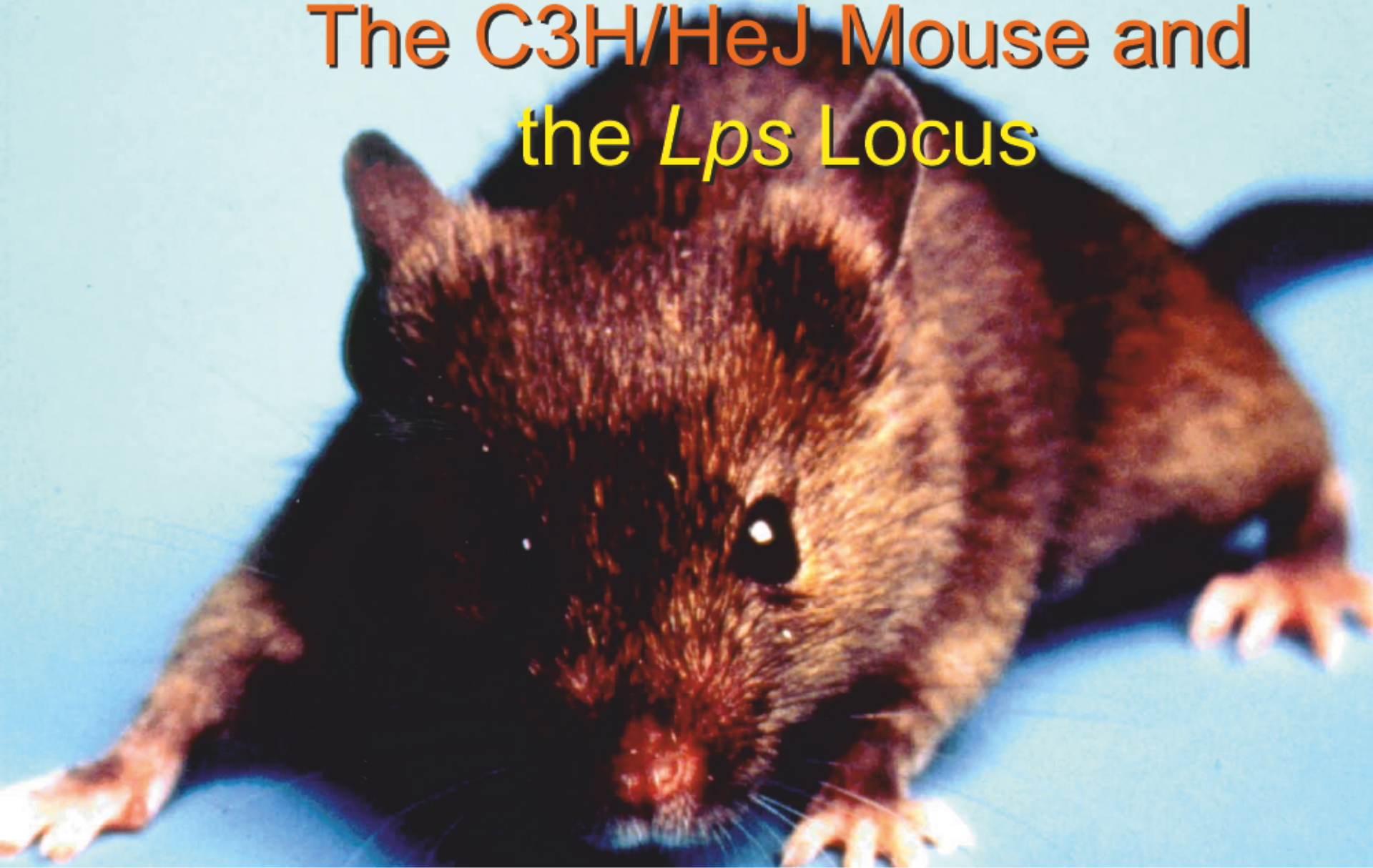


# The C3H/HeJ Mouse and the *Lps* Locus



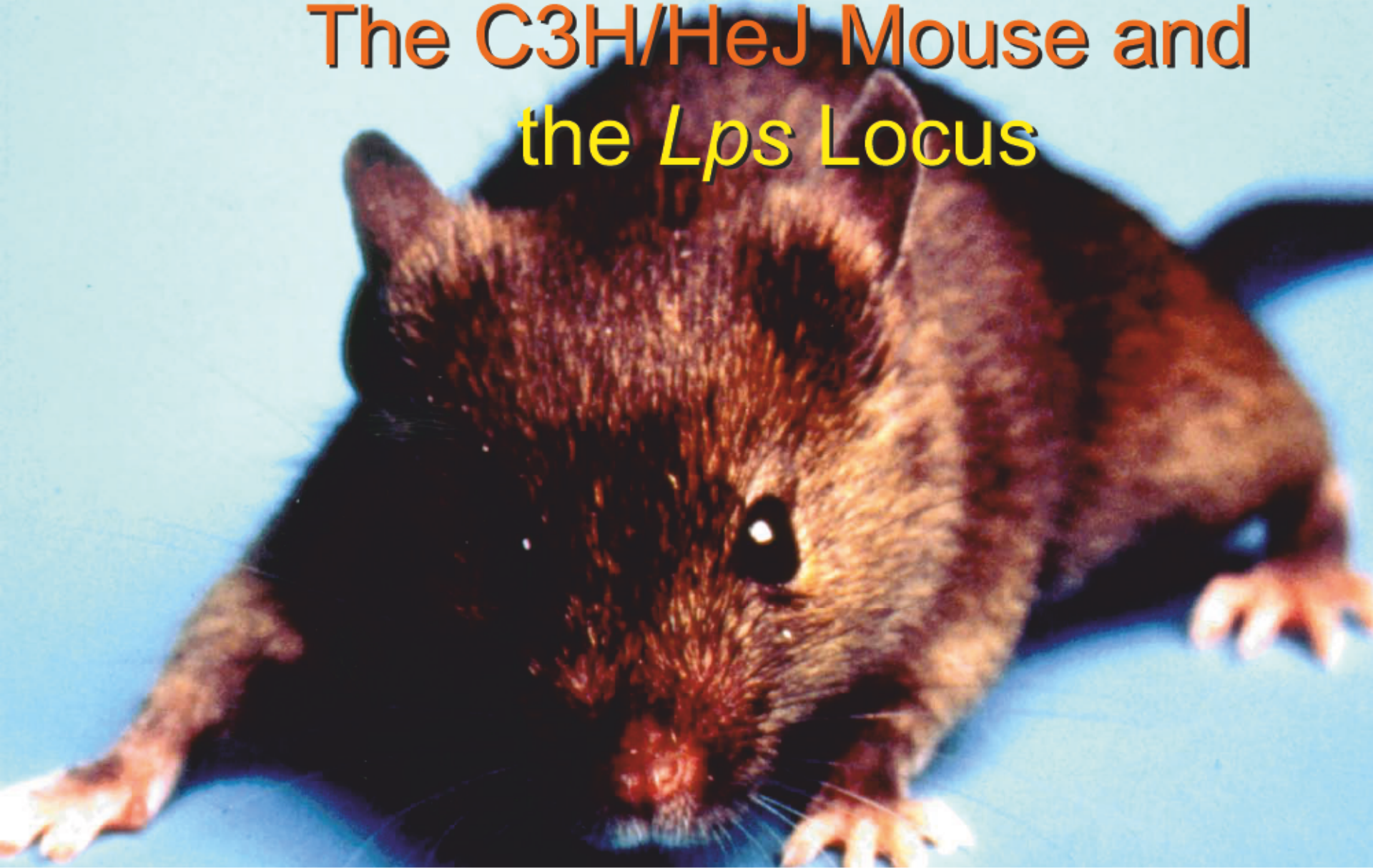
- Hypersusceptible to authentic G(-) infections (O'brien, et al., 1980; Svanborg-Eden, et al., 1983)

# The C3H/HeJ Mouse and the *Lps* Locus



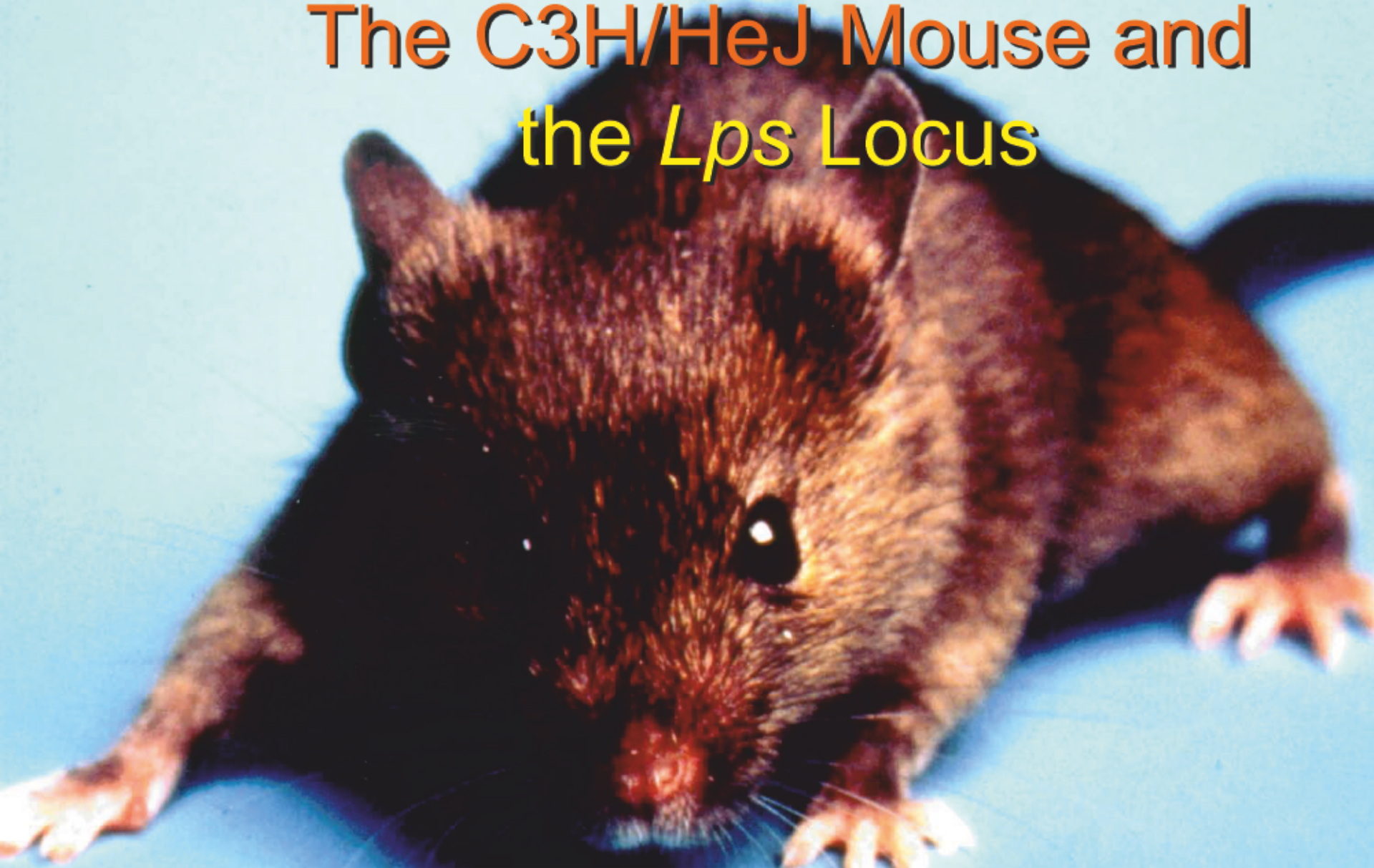
- LPS does not work as an adjuvant in C3H/HeJ mice (B.J. Skidmore et al, 1976)

# The C3H/HeJ Mouse and the *Lps* Locus

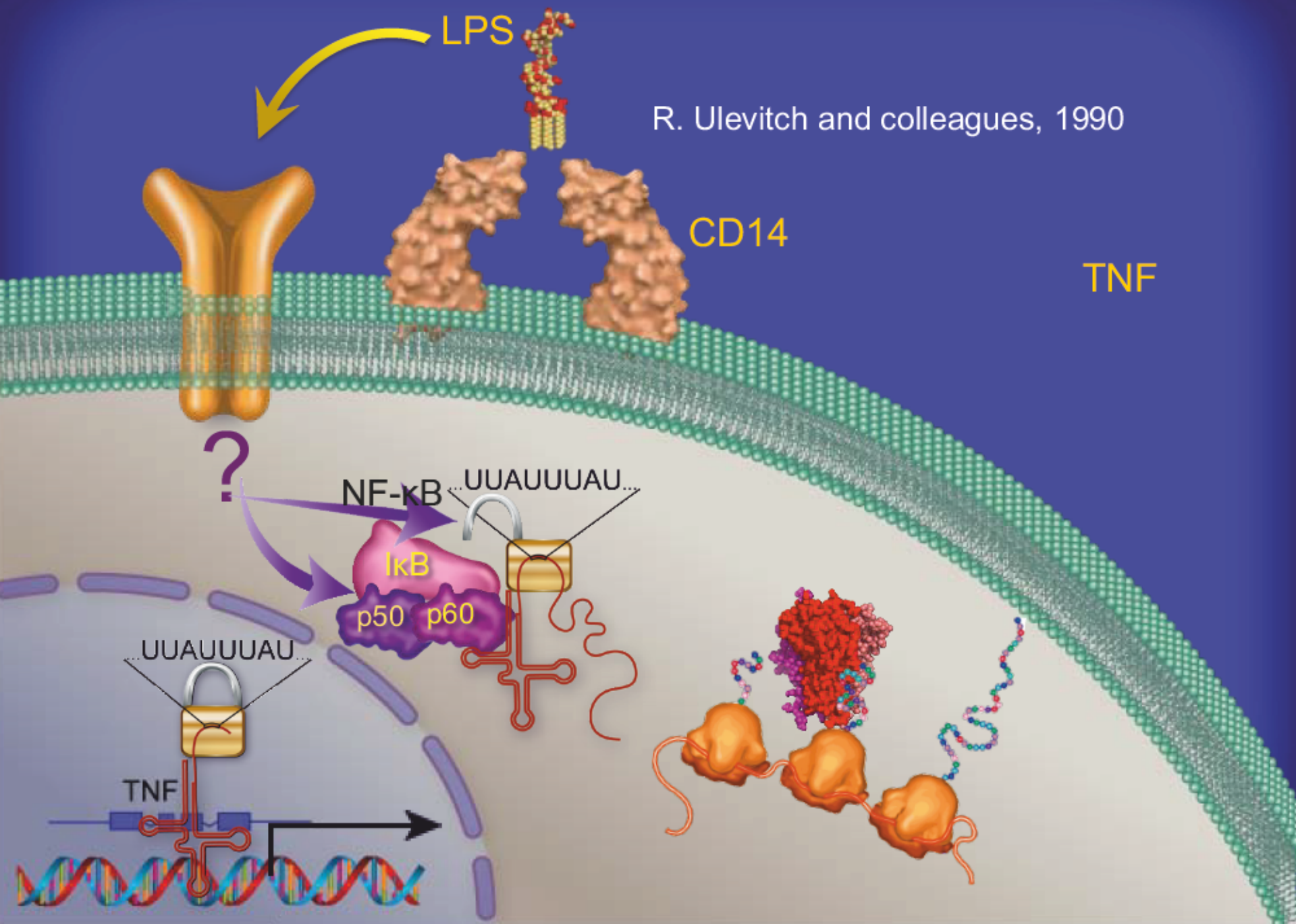


- Single locus (*Lps*); allelic to a mutation in the LPS-refractory C57BL/10ScCr strain (Coutinho and Meo, 1978).

# The C3H/HeJ Mouse and the *Lps* Locus



- *Lps* mapped to Chr. 4 between *Mup1* and *Polysyndactyly* loci by Watson et al. in 1978.



LPS

R. Ulevitch and colleagues, 1990

CD14

TNF

?

NF-κB

IκB

p50

p60

UUAUUUUAU...

UUAUUUUAU...

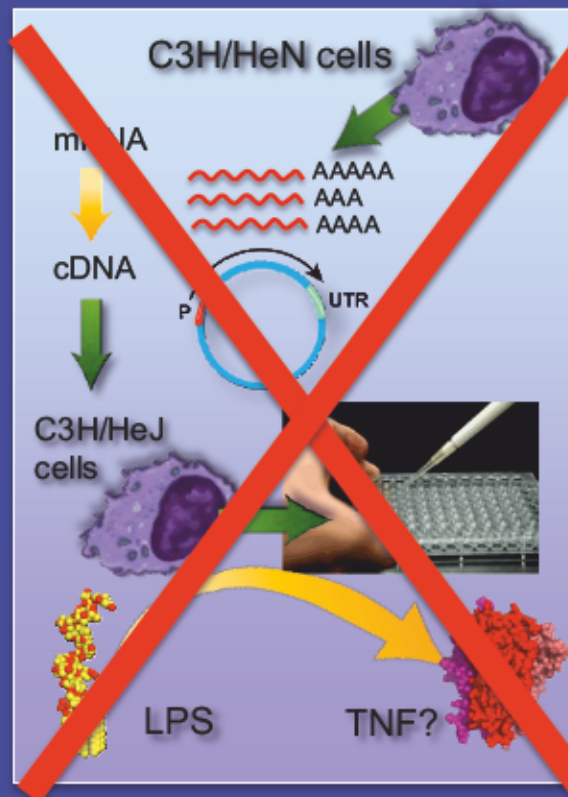
TNF



# Conventional searches for a difference between C3H/HeJ and C3H/HeN



1. Cross immunization of C3H/HeJ and C3H/HeN mice.



2. Transfect cDNA from C3H/HeN to C3H/HeJ cells.



3. Comparisons at the protein level.

Betsy  
Layton



Alexander  
Poltorak



Christophe  
Van Huffel



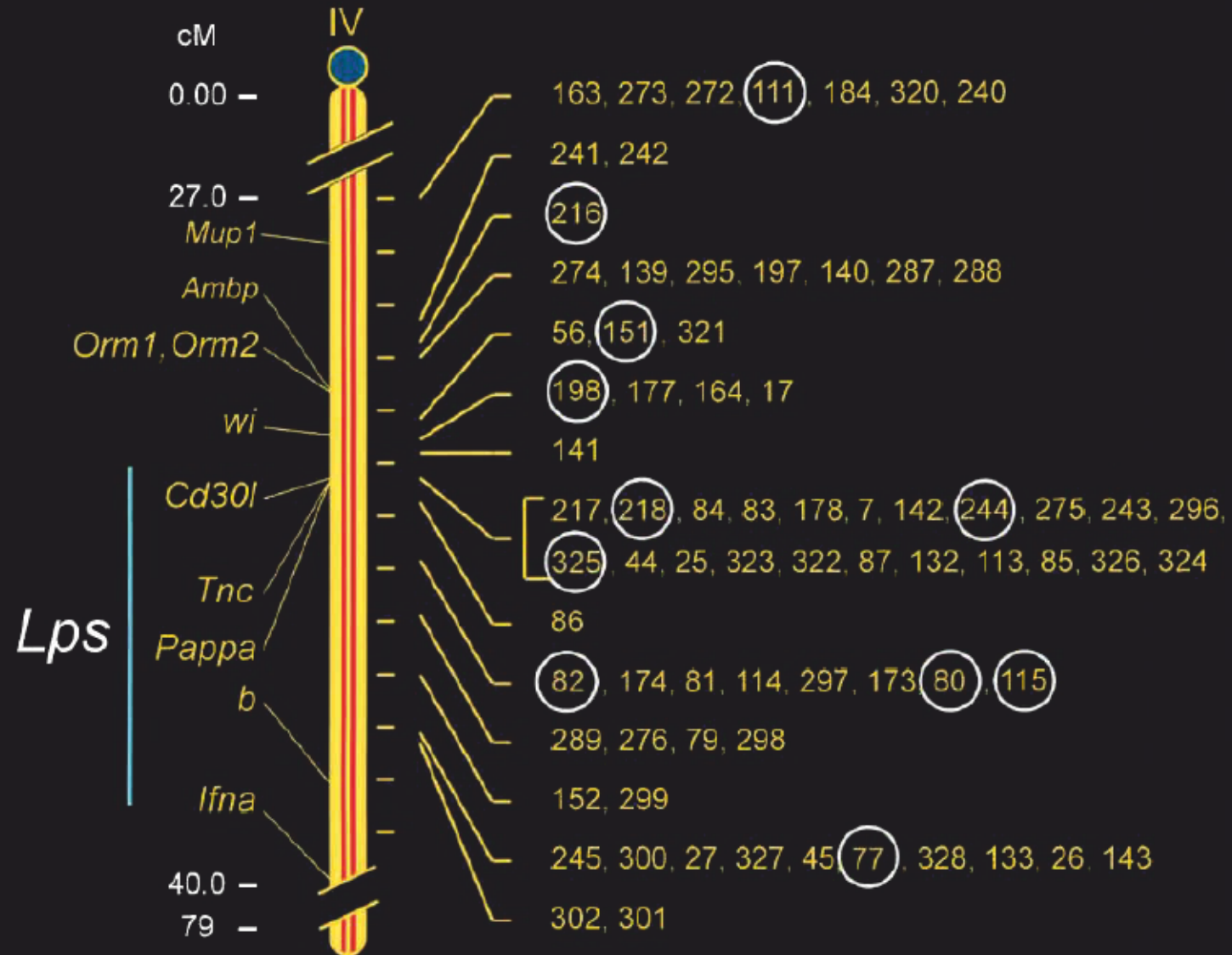
Irina  
Smirnova



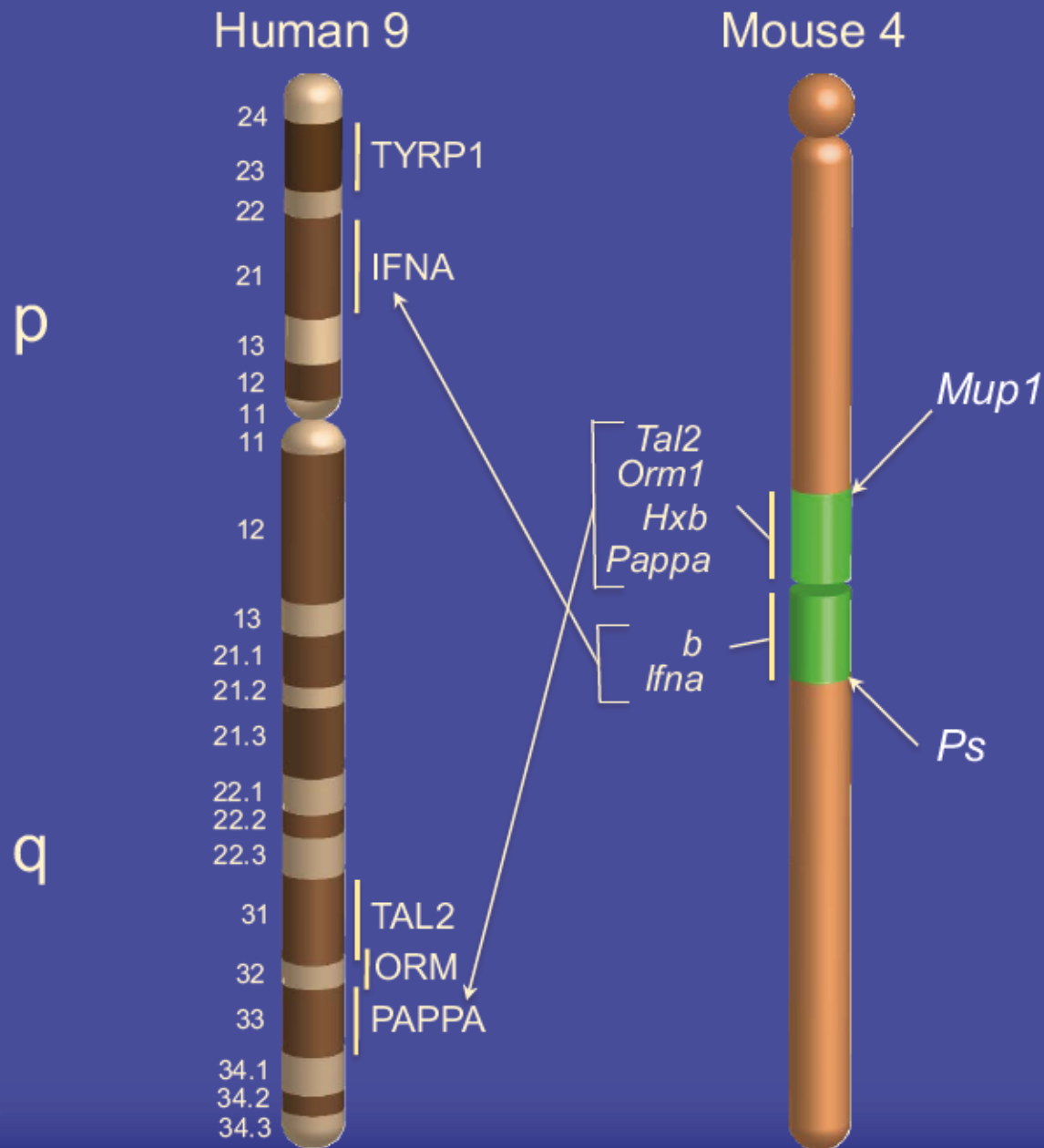
# Positional cloning entails...

- *Genetic mapping* (in our case, on 2093 meioses)
- *Physical mapping* (in our case, entire interval cloned in 66 BACs and 2 YACs)
- *Exploration for genes* (in our case, 1 authentic genes and 7 pseudogenes)
- *Mutation identification* (find the one and only genetic change responsible for the phenotype.)

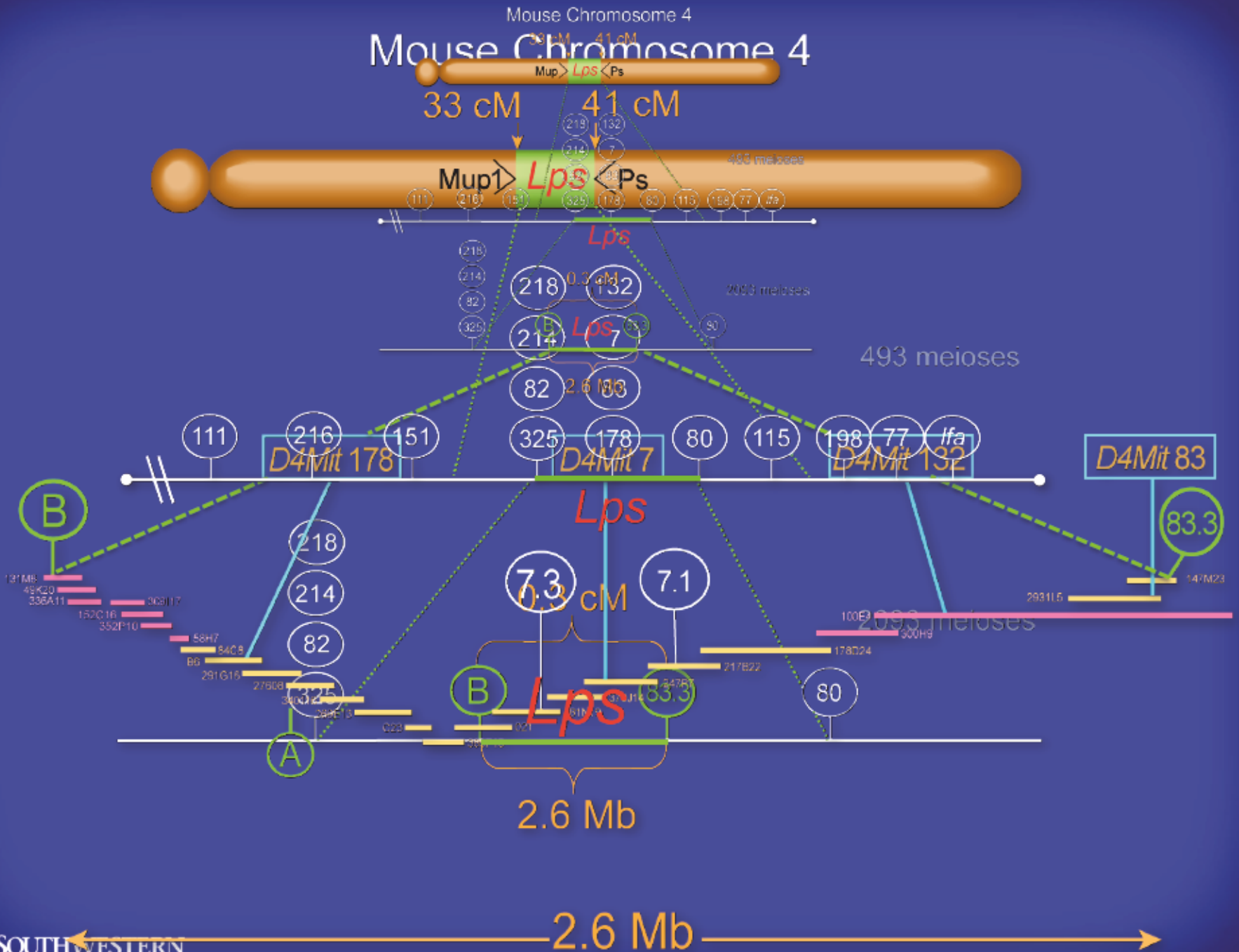
# MAPPING *Lps* WITH RESPECT TO D4MIT MARKERS



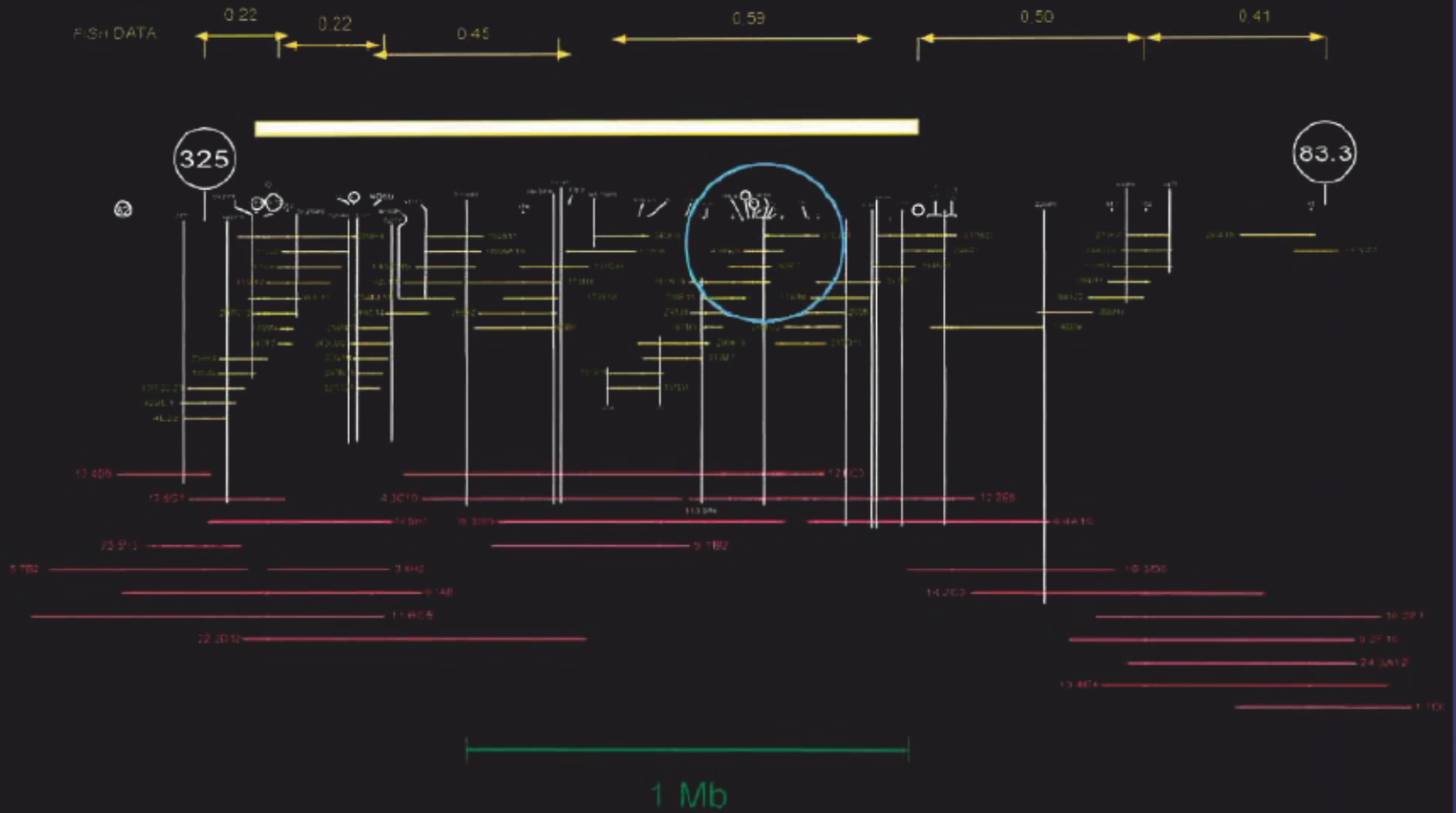
# Comparison of Mouse and Human LPS Gene Locus



# Mouse Chromosome 4



# CONTIG: D4MIT325 THRU 83.3



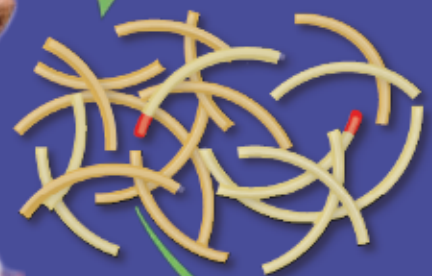
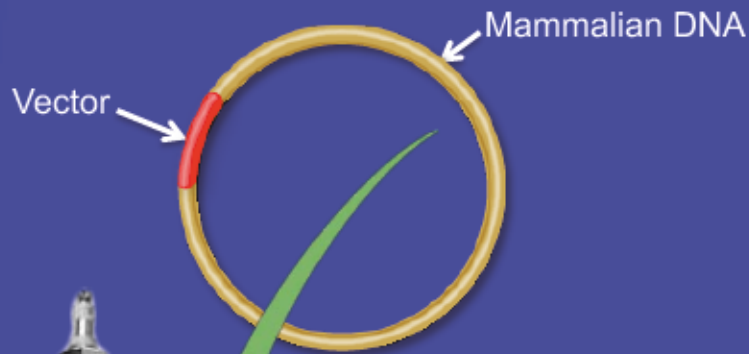




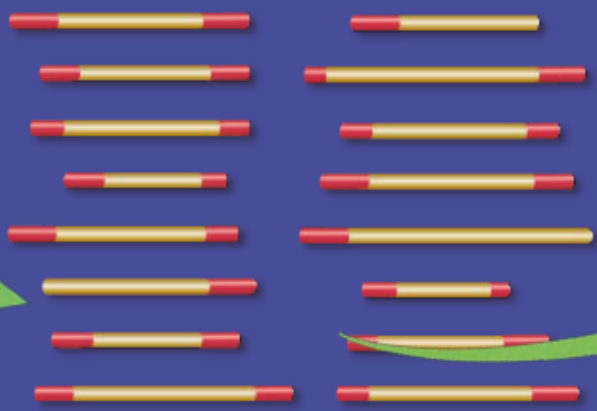
# Methods for finding genes within BACs

- Exon trapping
- Hybridization selection
- Computational prediction (GRAIL)
- Shotgun sequencing and EST database searching

# Creation of BACs

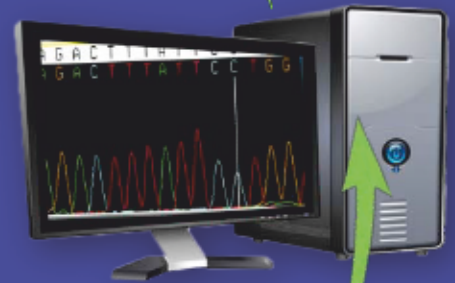


# Fragmentation of BACs



# Cloning of Fragments

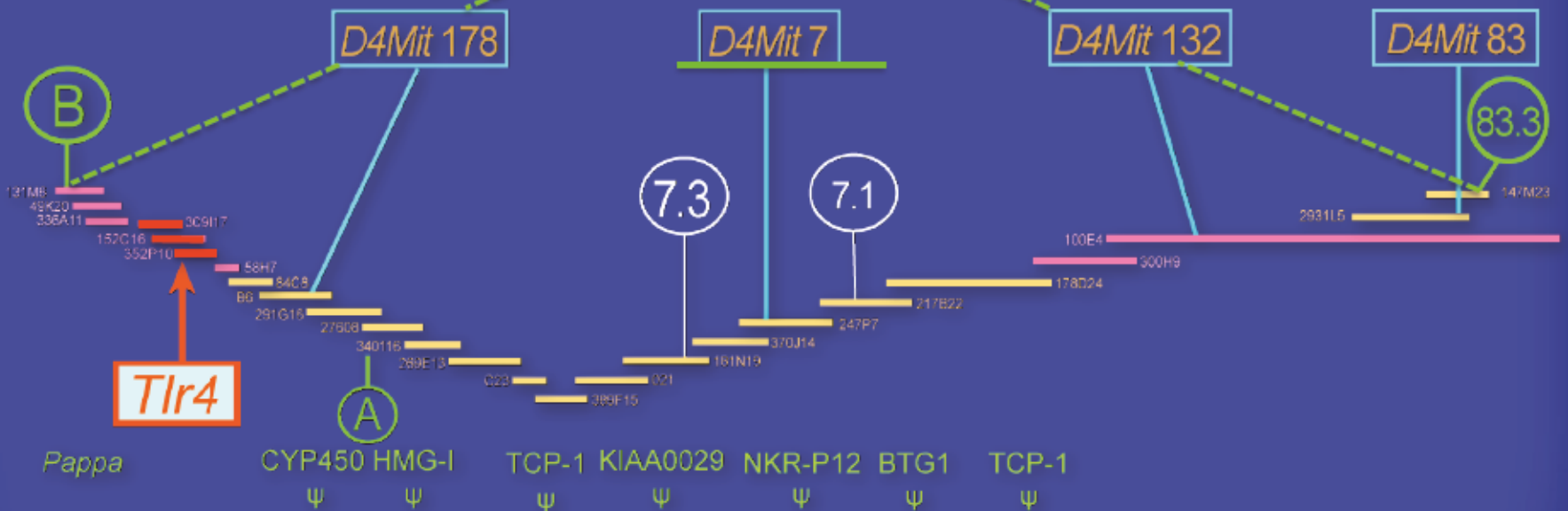
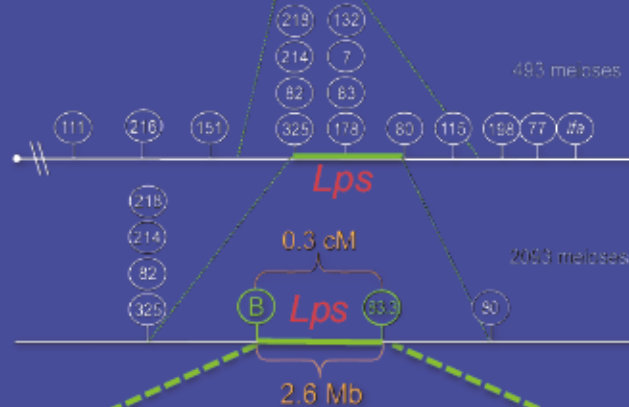
# Remote EST database



# BLASTing

Mouse Chromosome 4

33 cM 41 cM



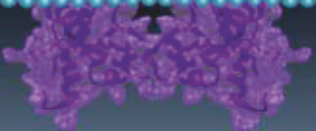
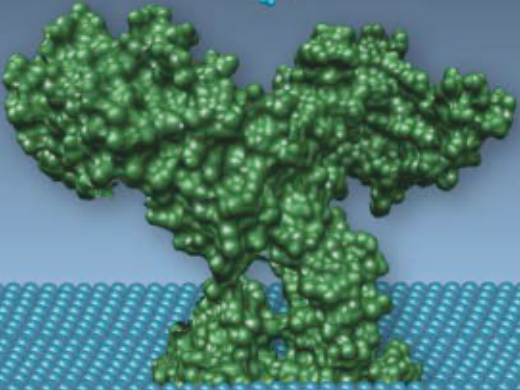
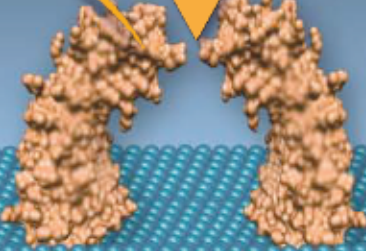
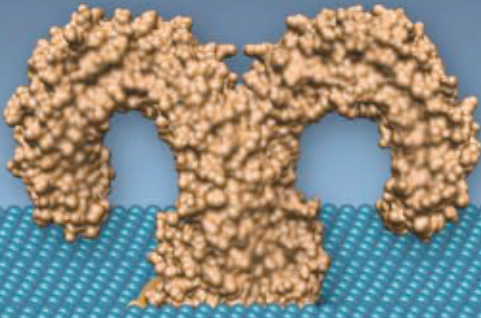
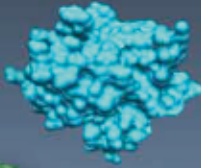
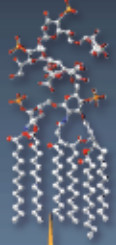
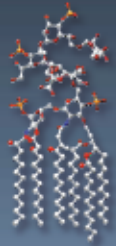
2.6 Mb



LPS

LPS

IL-1



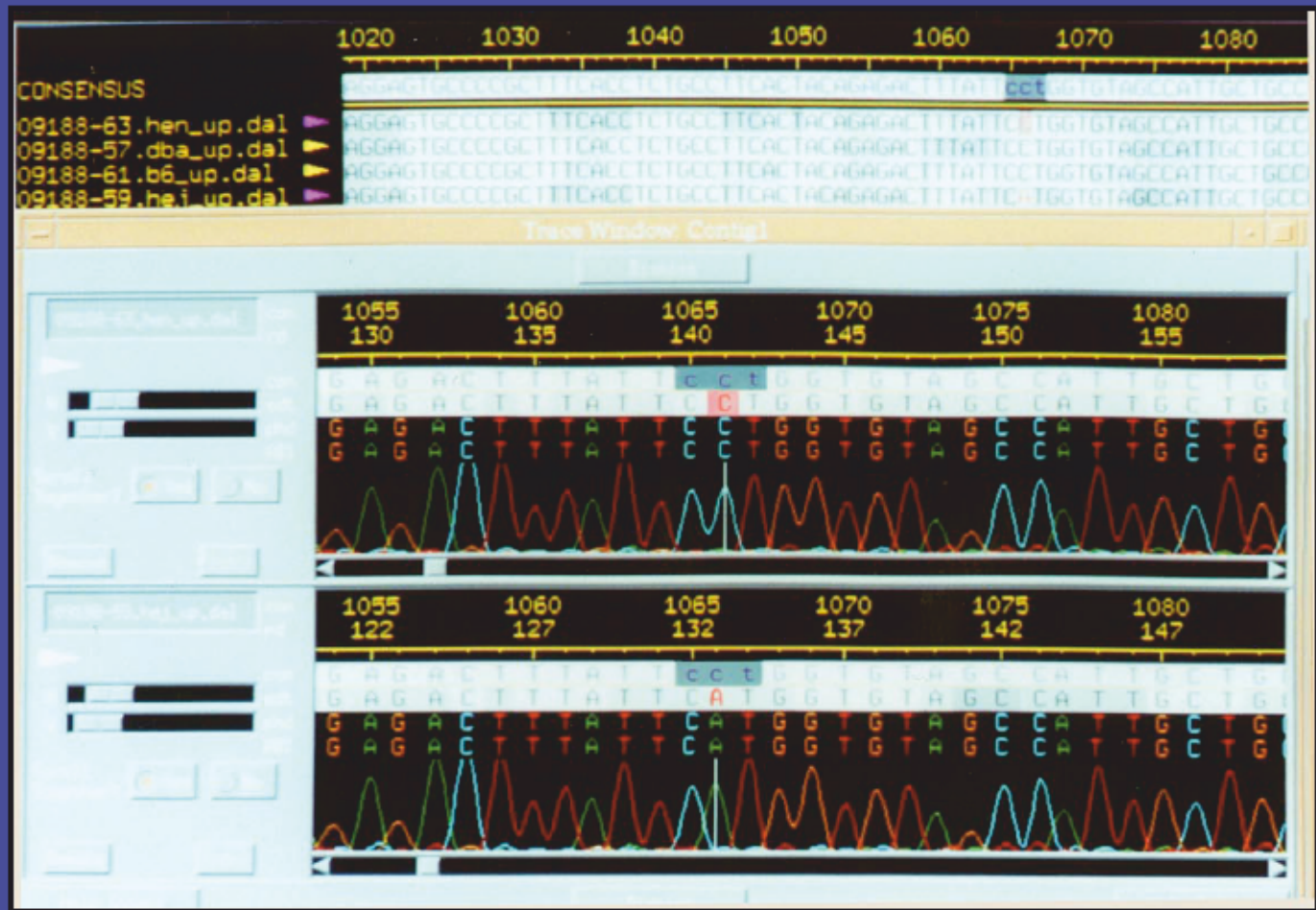
TLR4

CD14

IL-1R

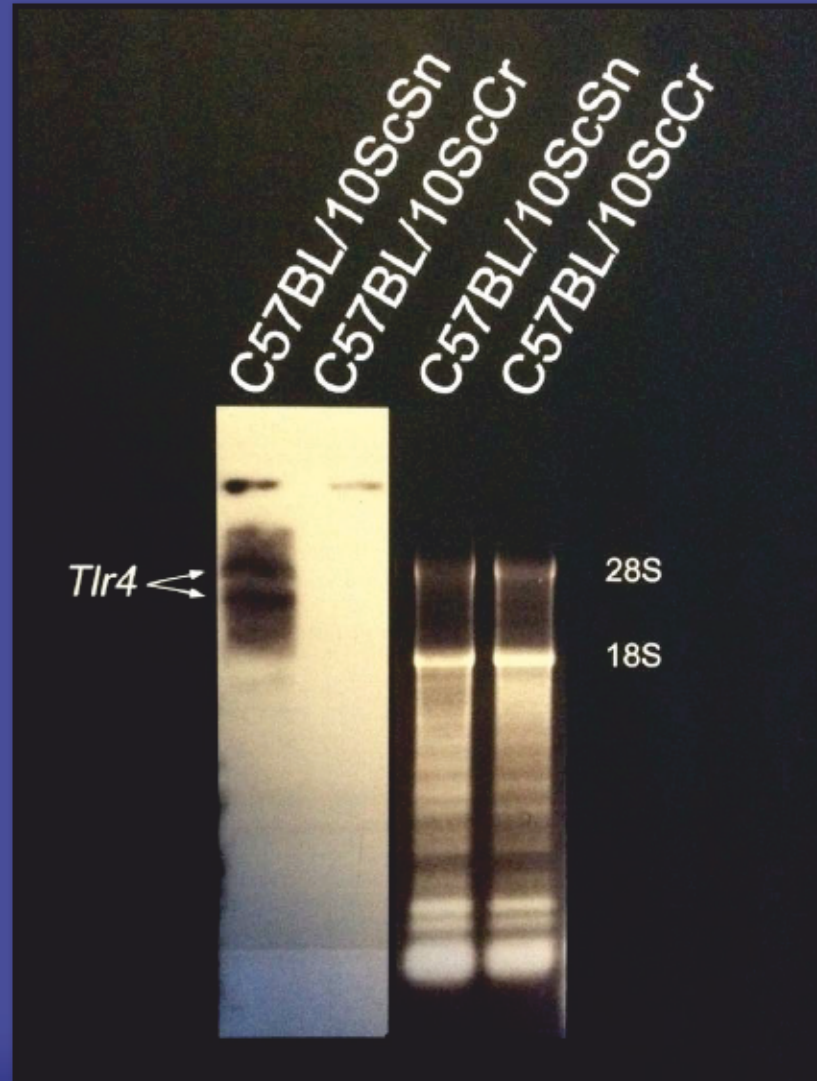


# Screen shot of a mutation in TLR4 distinguishing C3H/HeJ from C3H/HeN mice

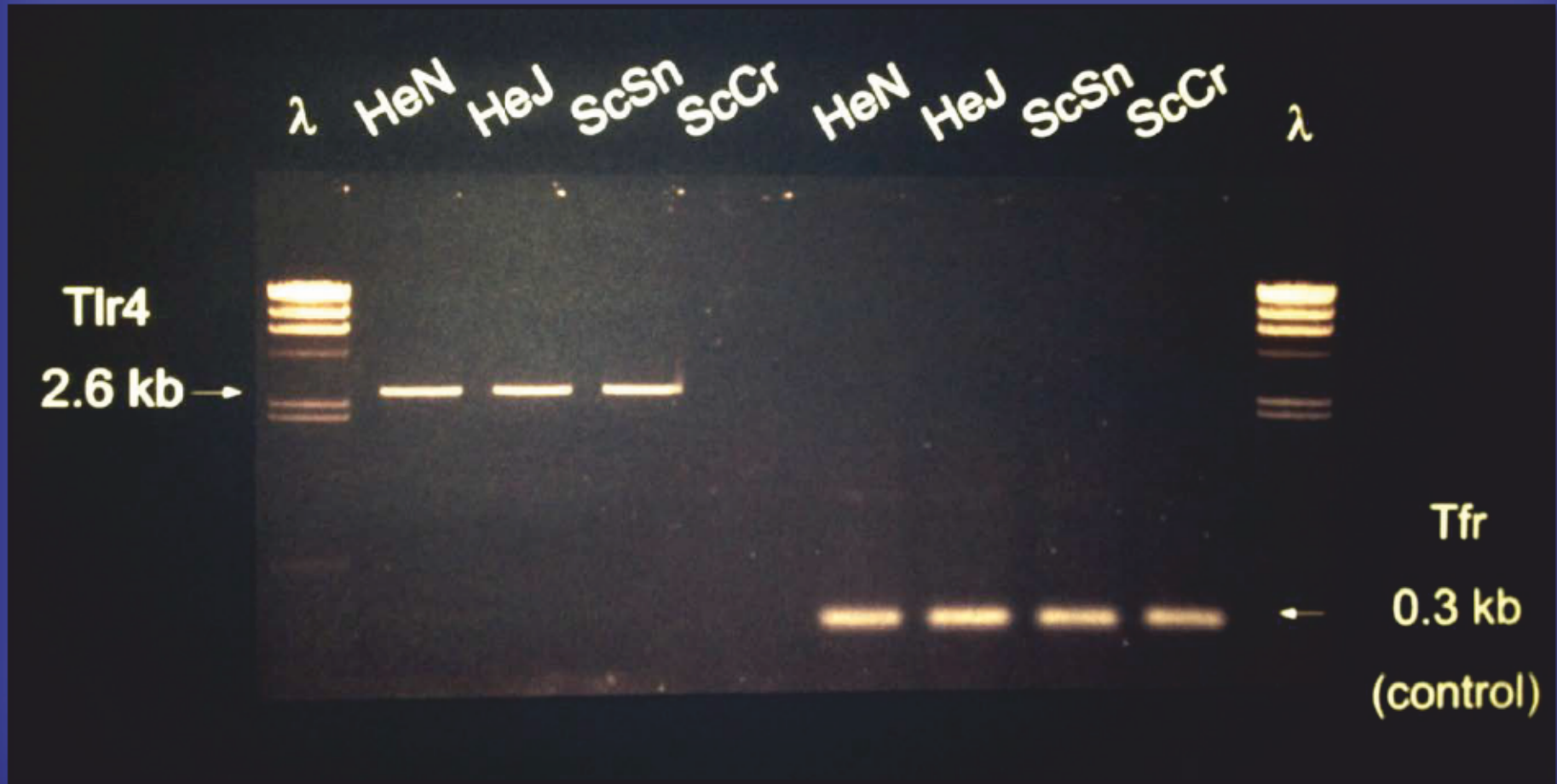


1998

On Northern blot analysis, C57BL/10ScCr mice appear not to express *Tlr4*



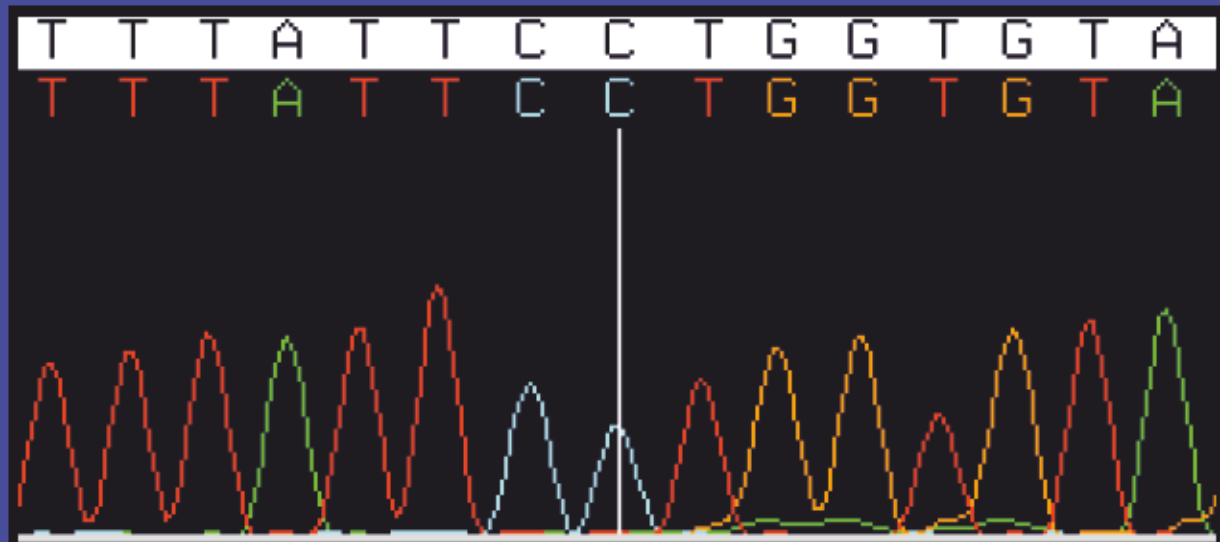
# RT-PCR also shows non-expression of Tlr4 in C57BL/10ScSr mice



1998

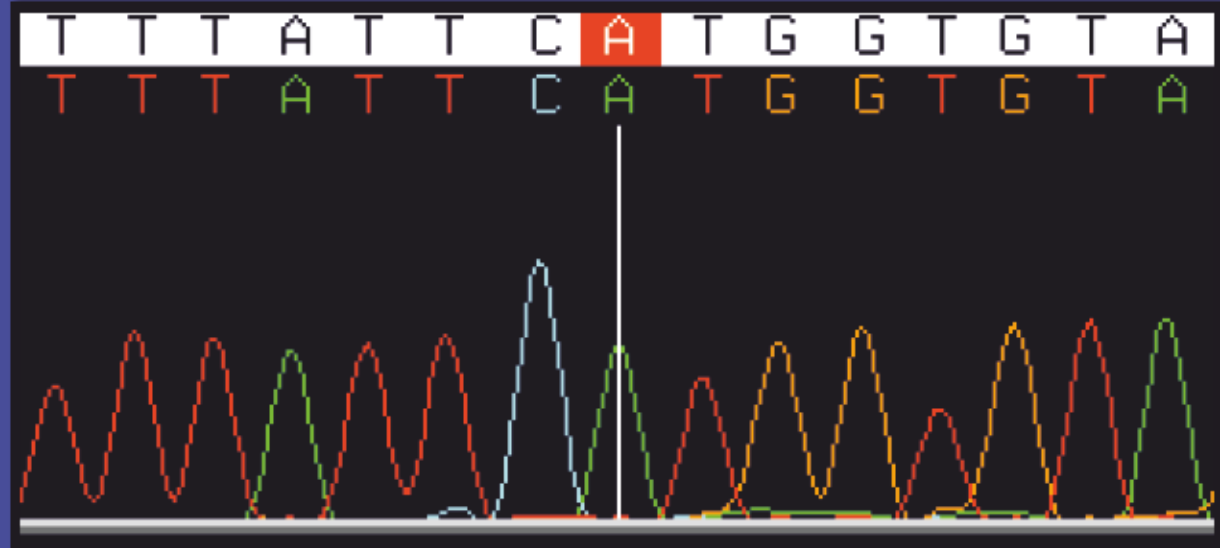
712

... F I P G V ...



C3H/HeN

*Tlr4*<sup>lps-n</sup>



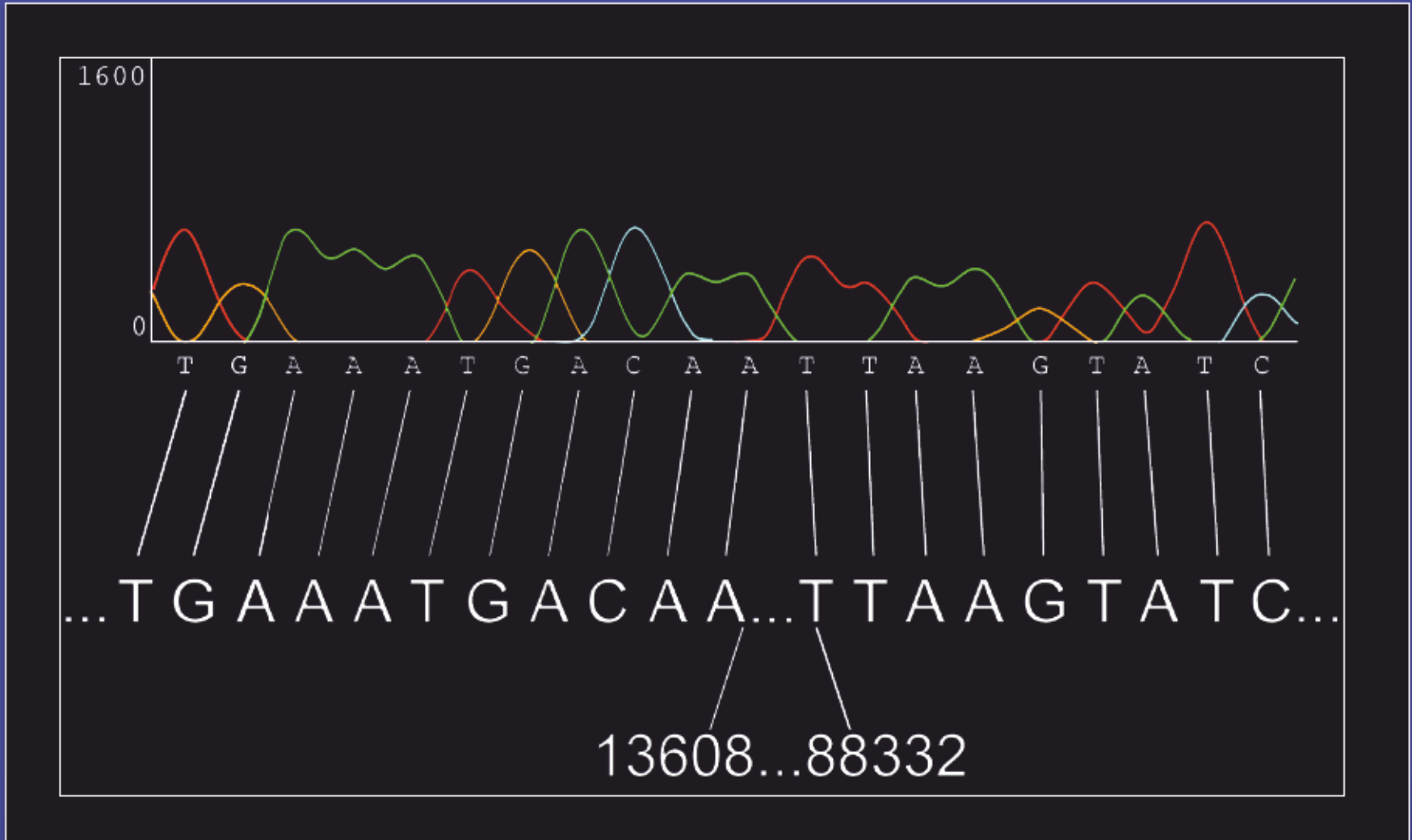
C3H/HeJ

*Tlr4*<sup>lps-d</sup>

... F I H G V ...



# Deletion of 74K in the C57BL/10ScCr mouse



obtained with  
d RNA analyses  
esis was in 8%  
× TEB (45 mM  
els. For analyses  
ed at pH 5.0 and  
cribed (24), and  
acrylamide, 8 M  
ete oxidation of  
d J. E. Dahlberg  
aminoacylation  
ected with <sup>35</sup>S-  
omigration of  
ermined by RNA  
dried gels (18).  
rotein synthesis,  
ncentration of  
s added to 200  
n synthesis was  
nine (50 μCi/ml

quences at the  
fingerprinting  
nces (13).  
ortase, *Nature*  
an, B. D. Holl,  
82).  
ev. *Biol.* 8, 65

results.  
*Science* 276,

*Cell*, 9, 3041  
elidis, *Cell* 14,  
b, L. M. Spitz,  
2929 (1993).  
enzymes with  
st to their nu-  
cytes (3), may  
GTP in splicing  
e inhibition of  
dional Ran-GTP  
on by accumu-  
chanism would  
unts of nuclear  
rsors accumu-  
Ran system in

26. F. Muller, S. G. Clarkson, D. J. Galas, *Nucleic Acids Res.* 15, 7191 (1987); A. Kressmann, S. G. Clarkson, V. Pirota, M. L. Birnstiel, *Proc. Natl. Acad. Sci. U.S.A.* 75, 1176 (1978).

the aminoacyl-AMS compounds. Supported by NIH grant GM30220.

9 September 1998; accepted 2 November 1998

## Defective LPS Signaling in C3H/HeJ and C57BL/10ScCr Mice: Mutations in *Tlr4* Gene

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Christophe Van Huffel,‡ Xin Du, Dale Birdwell, Erica Alejos,  
Maria Silva, Chris Galanos, Marina Freudenberg,  
Paola Ricciardi-Castagnoli, Betsy Layton, Bruce Beutler§

Mutations of the gene *Lps* selectively impede lipopolysaccharide (LPS) signal transduction in C3H/HeJ and C57BL/10ScCr mice, rendering them resistant to endotoxin yet highly susceptible to Gram-negative infection. The codominant *Lps<sup>d</sup>* allele of C3H/HeJ mice was shown to correspond to a missense mutation in the third exon of the Toll-like receptor-4 gene (*Tlr4*), predicted to replace proline with histidine at position 712 of the polypeptide chain. C57BL/10ScCr mice are homozygous for a null mutation of *Tlr4*. Thus, the mammalian Tlr4 protein has been adapted primarily to subservise the recognition of LPS and presumably transduces the LPS signal across the plasma membrane. Destructive mutations of *Tlr4* predispose to the development of Gram-negative sepsis, leaving most aspects of immune function intact.

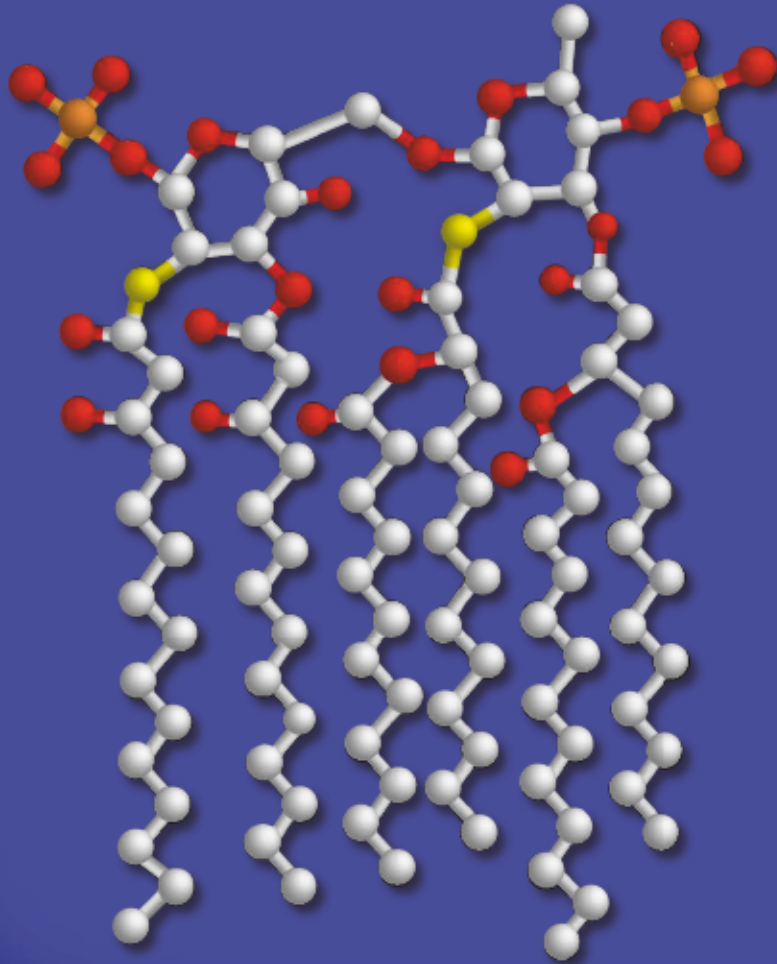
Conservative estimates hold that in the United States alone, 20,000 people die each year as a result of septic shock brought on by Gram-negative infection (1). The lethal effect of a Gram-negative infection is linked, in part, to the biological effects of bacterial

lipopolysaccharide (endotoxin), which is produced by all Gram-negative organisms. A powerful activator of host mononuclear cells, LPS prompts the synthesis and release of tumor necrosis factor (TNF) and other toxic cytokines that ultimately lead to shock in

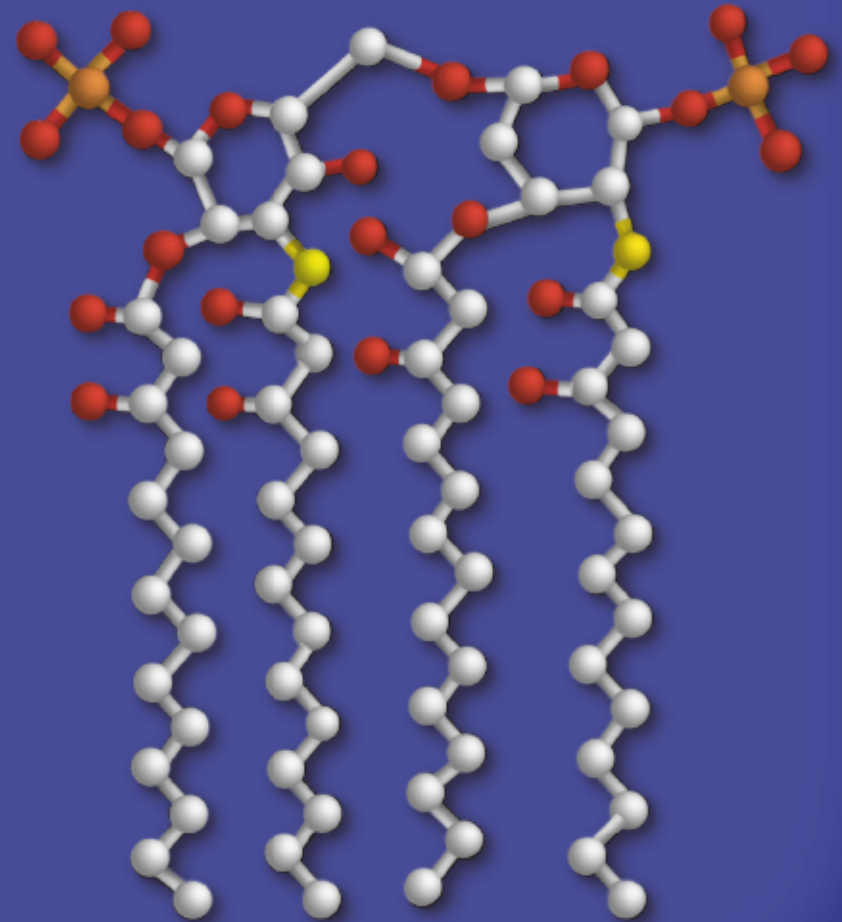
But the question remained:

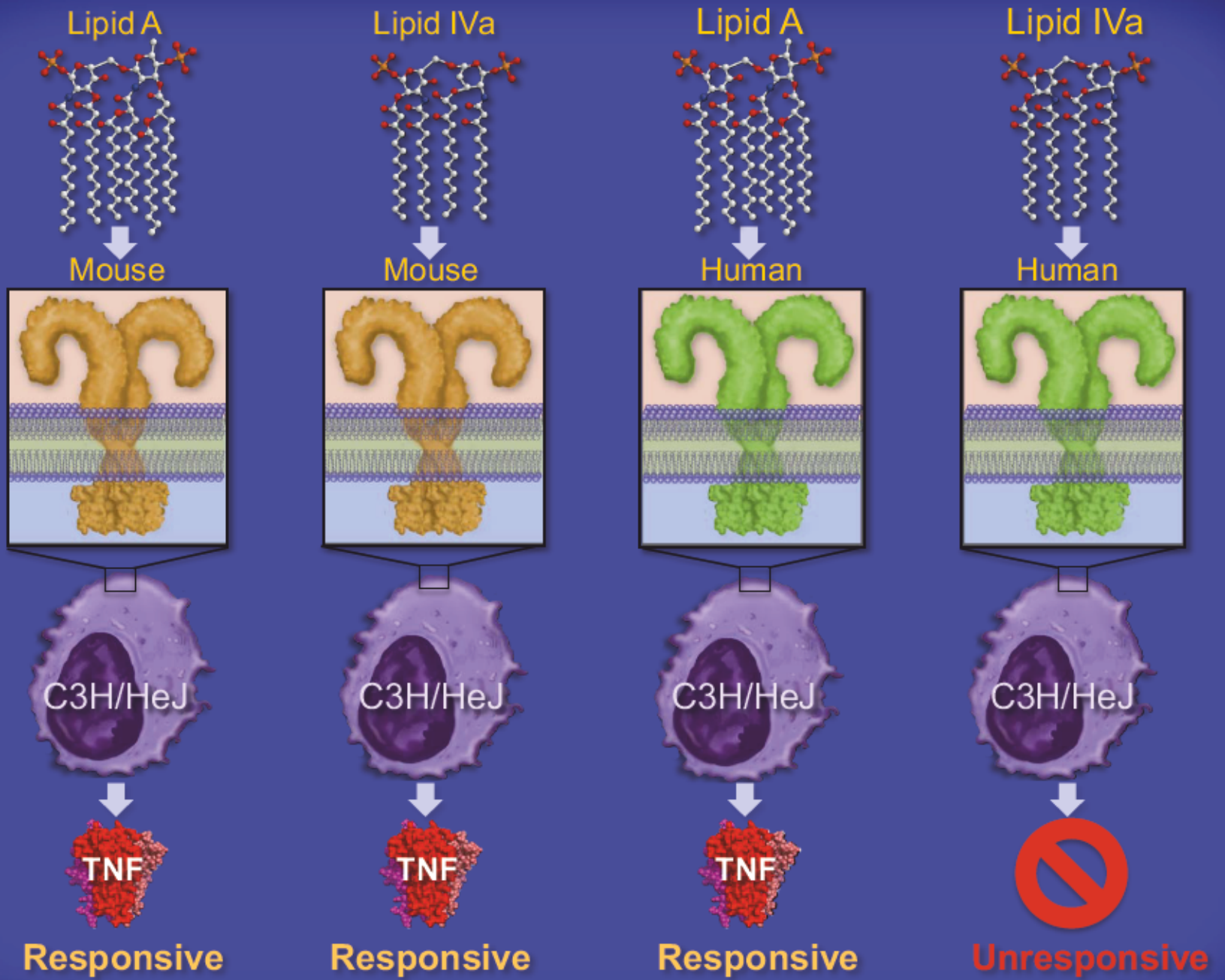
Was there direct contact between  
TLR and LPS?

Lipid A:  
Agonist for both  
mouse and human



Lipid IVa:  
Agonist for mouse;  
antagonist for human





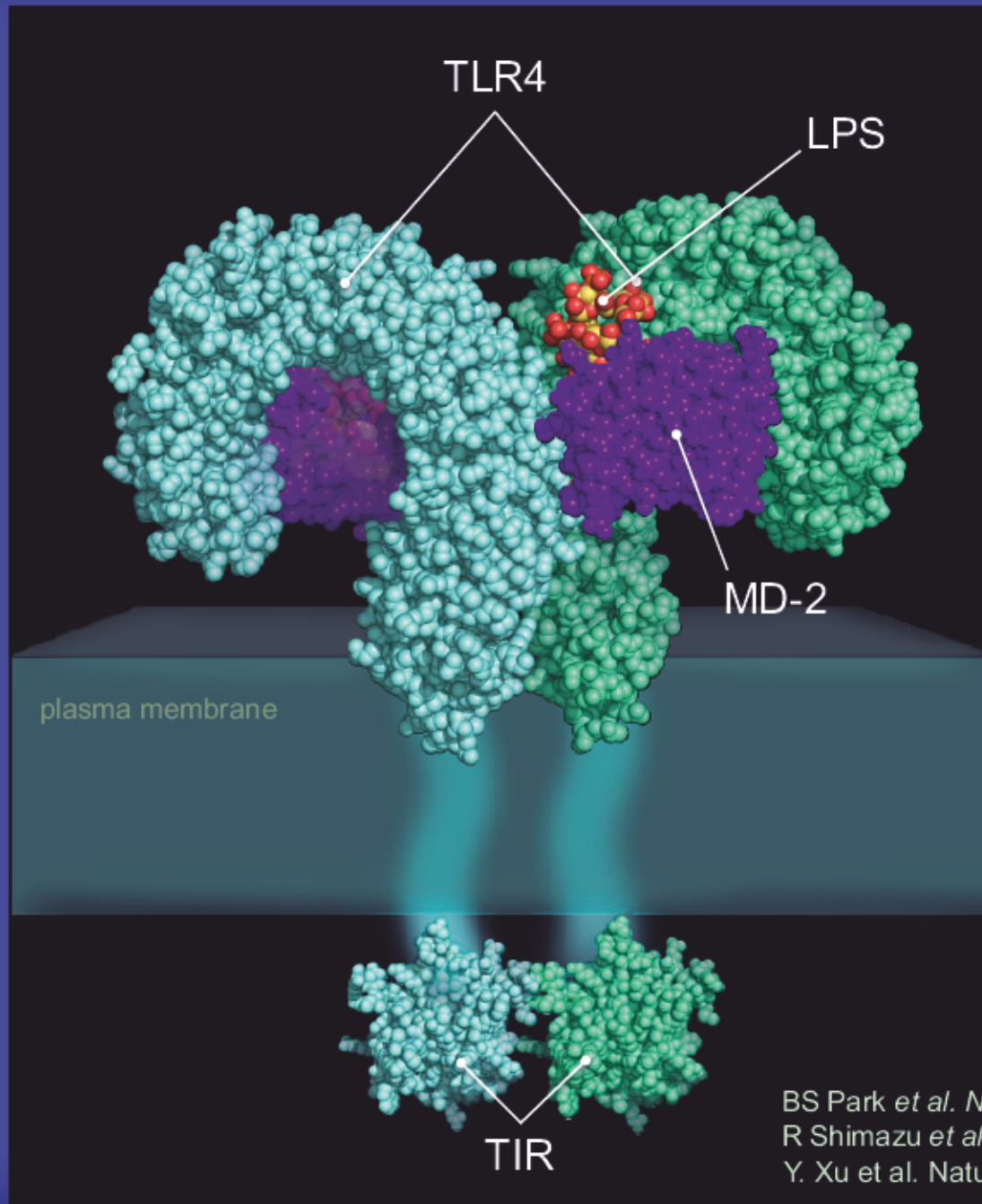
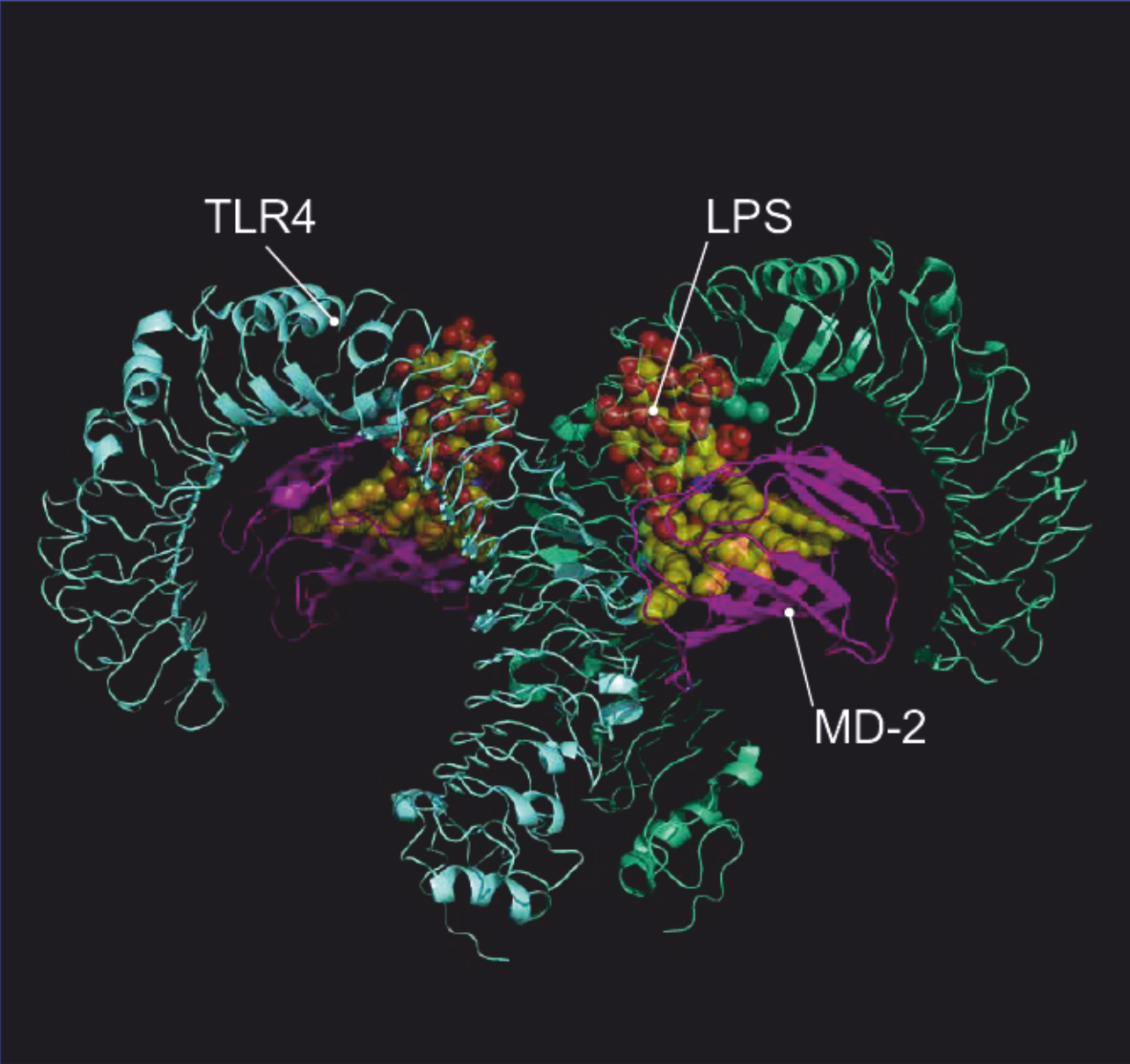
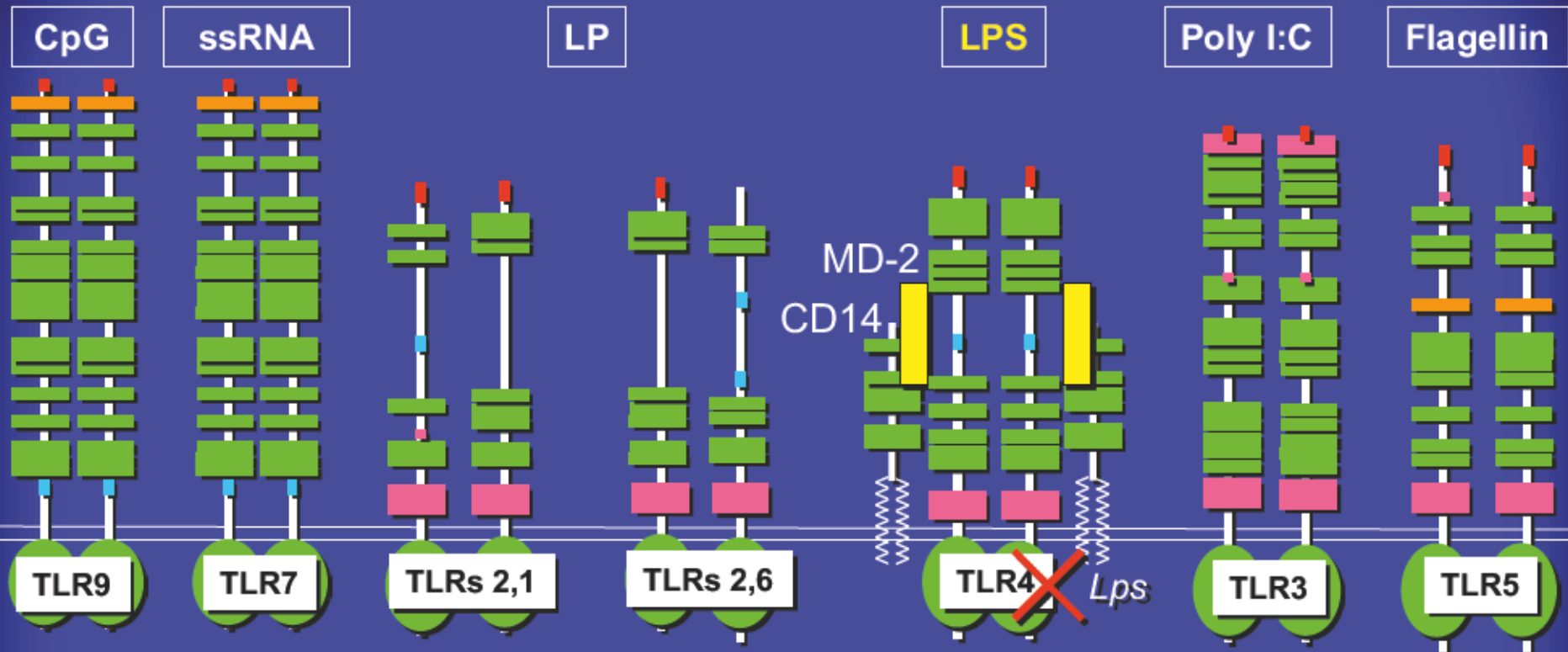


Image created using PyMol

BS Park *et al.* *Nature* **458**, 7242 (2009)  
R Shimazu *et al.*, *J Exp Med* **189**, 11 (1999)  
Y. Xu *et al.* *Nature* **408**, 6808 (2000)



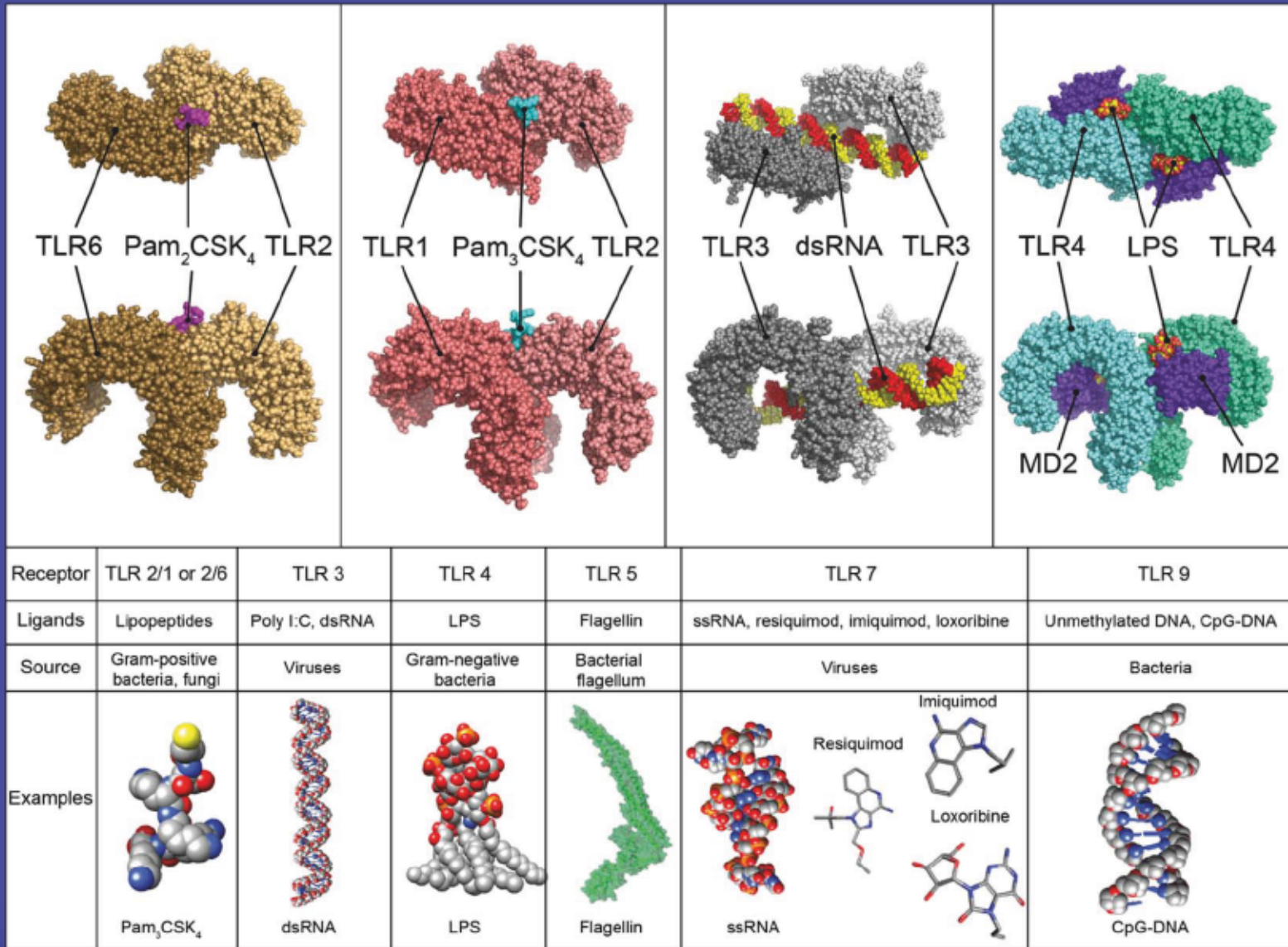


The role of TLR4 as a sensor of LPS suggested that other microbial ligands were sensed by other TLRs, and the specificity of most TLRs was revealed by knocking the genes out one at a time, largely in the Akira lab.

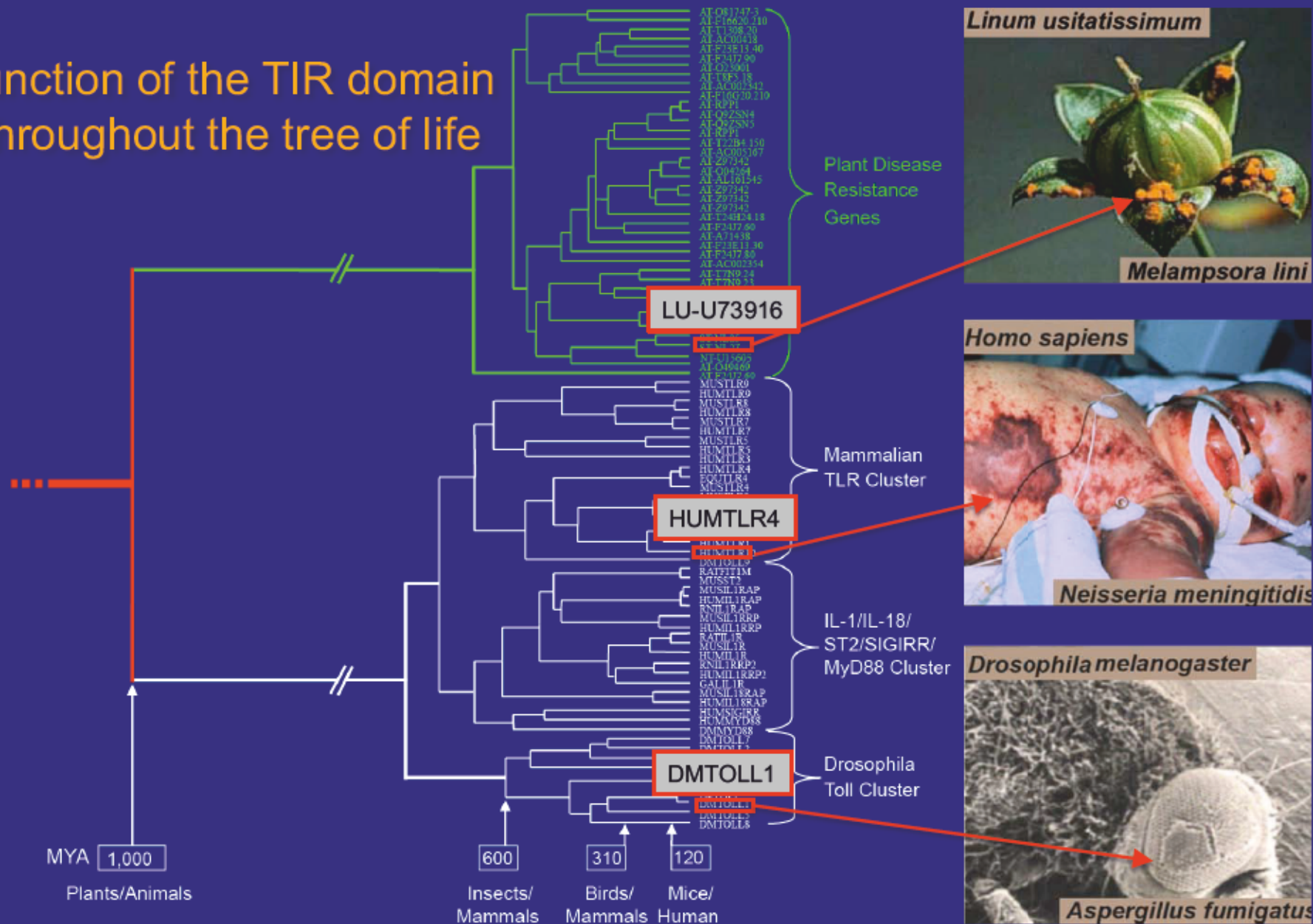




# Now, the mode of binding of several ligands to TLRs is understood

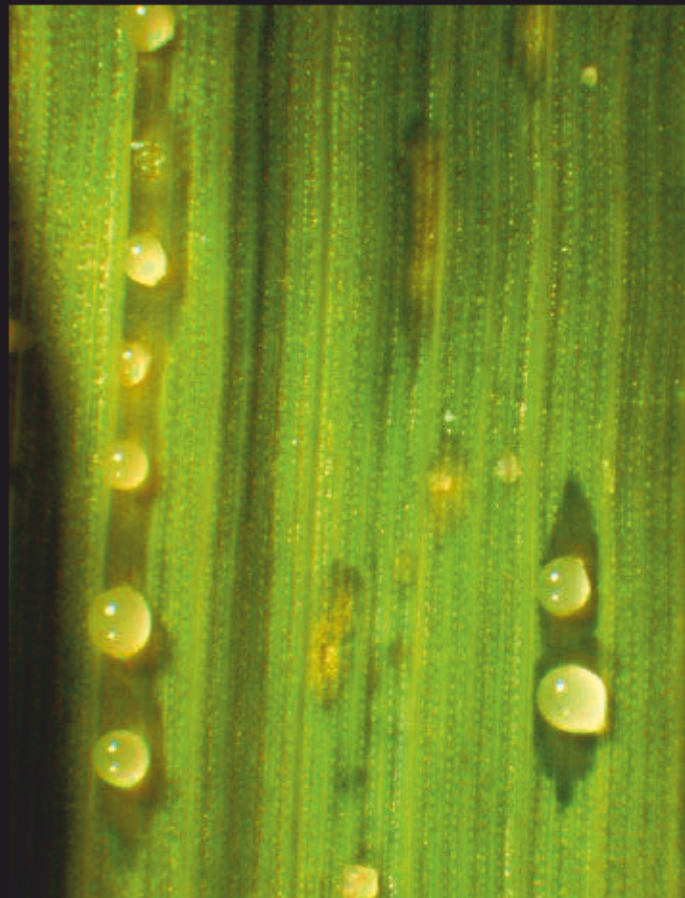


# Function of the TIR domain throughout the tree of life



# Bacterial blight of rice

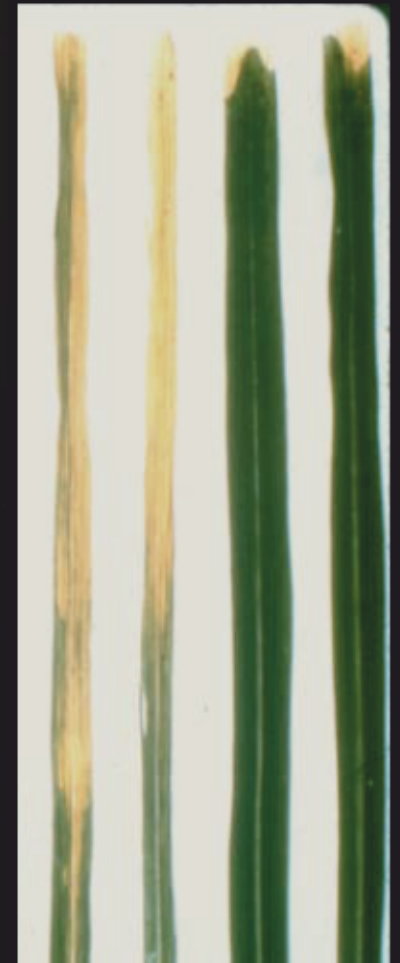
Pamela Ronald  
Discoverer of XA21

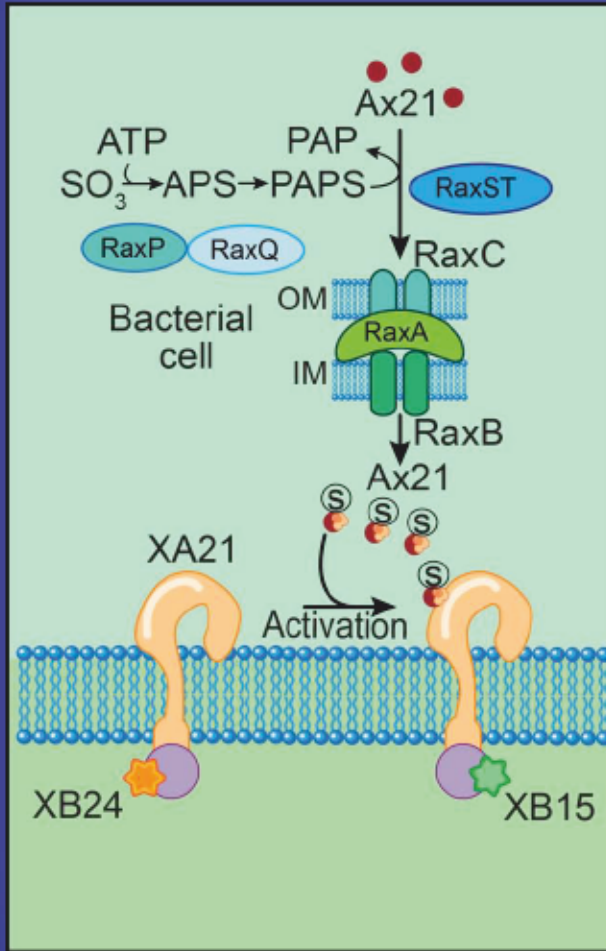


*Xanthomonas oryzae* (Xoo)

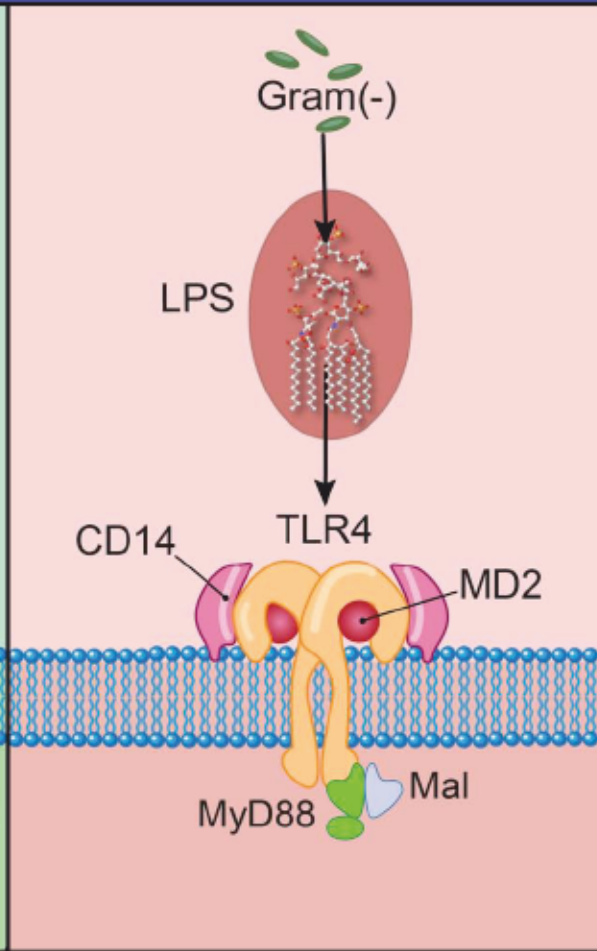
TP309

Xa21

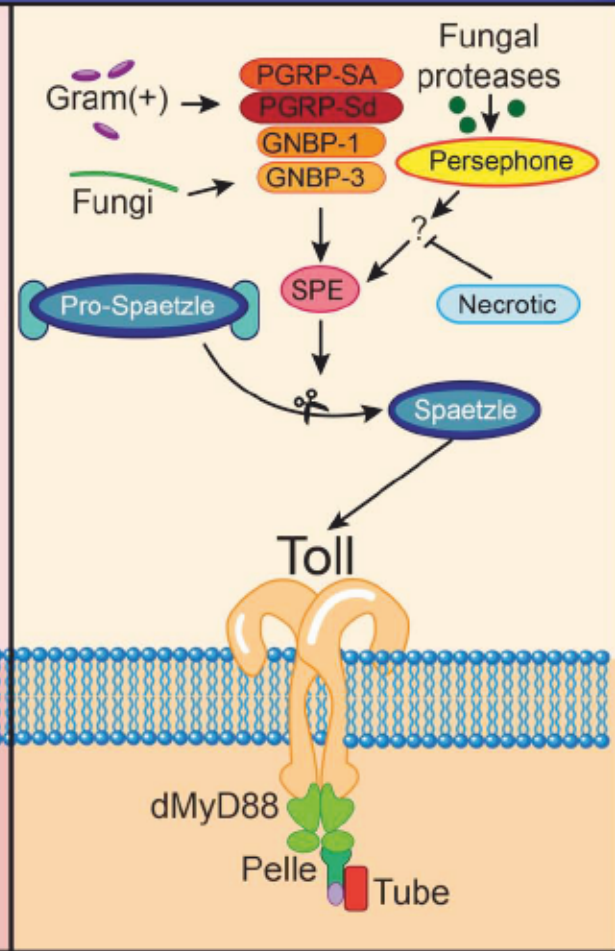




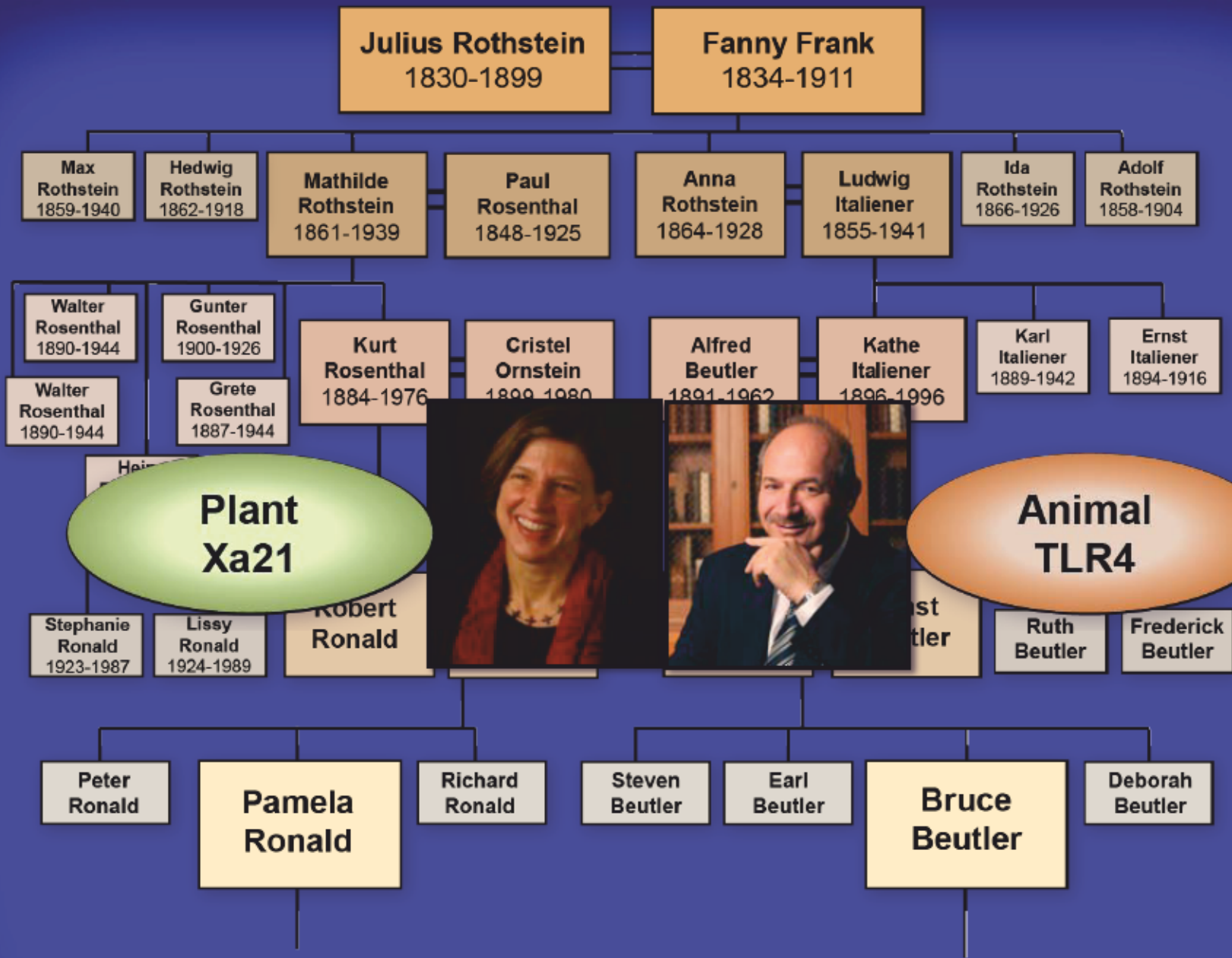
Monocots (Rice)



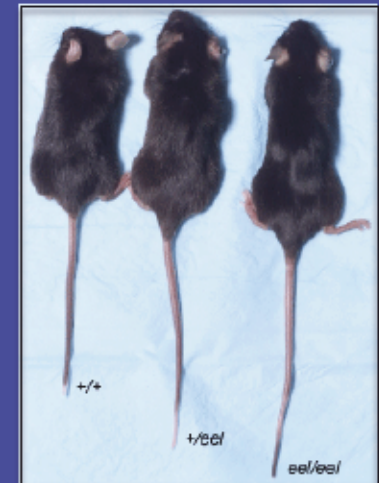
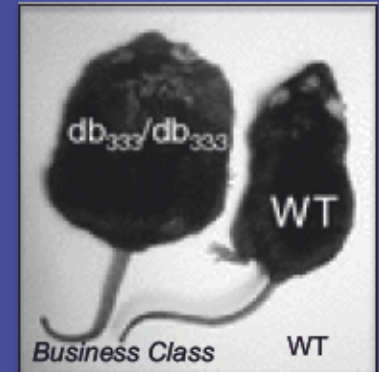
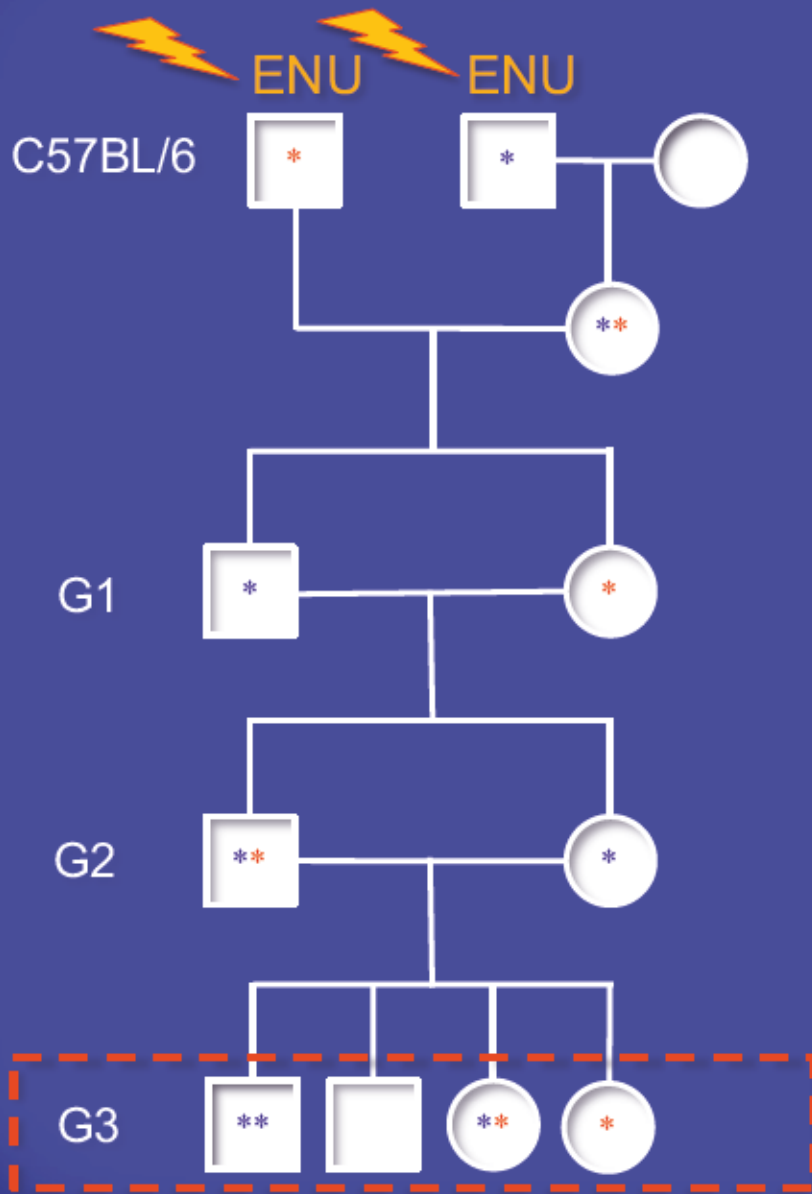
Vertebrates (Mouse)



Insects (Drosophila)



# Making new phenotypes in mice



As of 9/20/11, >151,000 G1 + G3 mice produced. ~60 coding/splicing changes per sperm.

The process of finding mutations was greatly accelerated by the sequencing and annotation of the mouse genome...

Never again was it necessary to make a BAC contig or search for genes.

# MGI genome browser (typical view)

**Search**  
Landmark or Region:  
   
Data Source  
  
**Overview**

Scroll/Zoom:        Flip

**Details**

**MGI Representative Transcripts**

**NCBI transcripts**

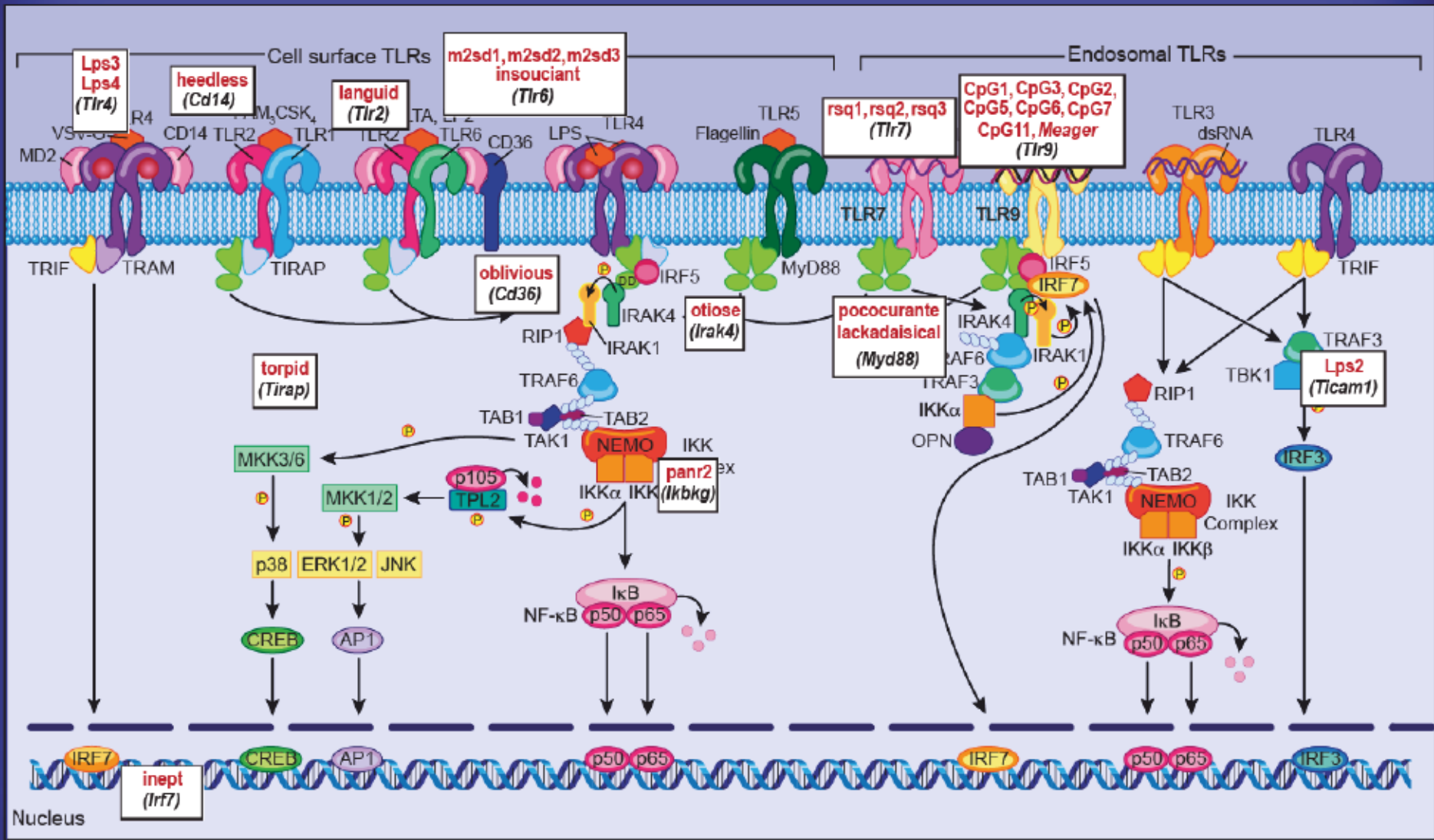
**ENSEMBL transcripts v53**

**VEGA transcripts v35**

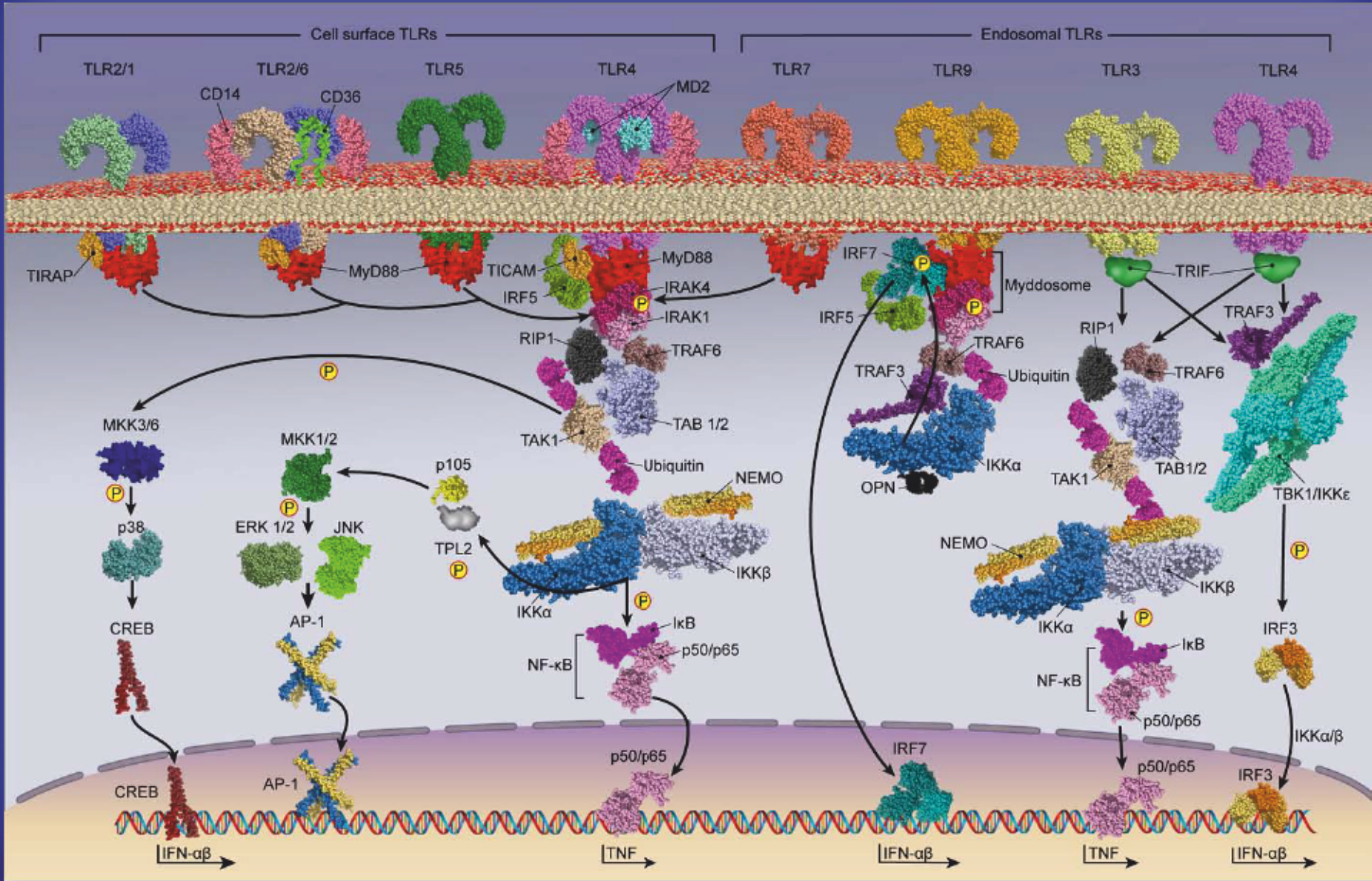


## Massively parallel short read sequencing has made targeted exon sequencing unnecessary in most cases.

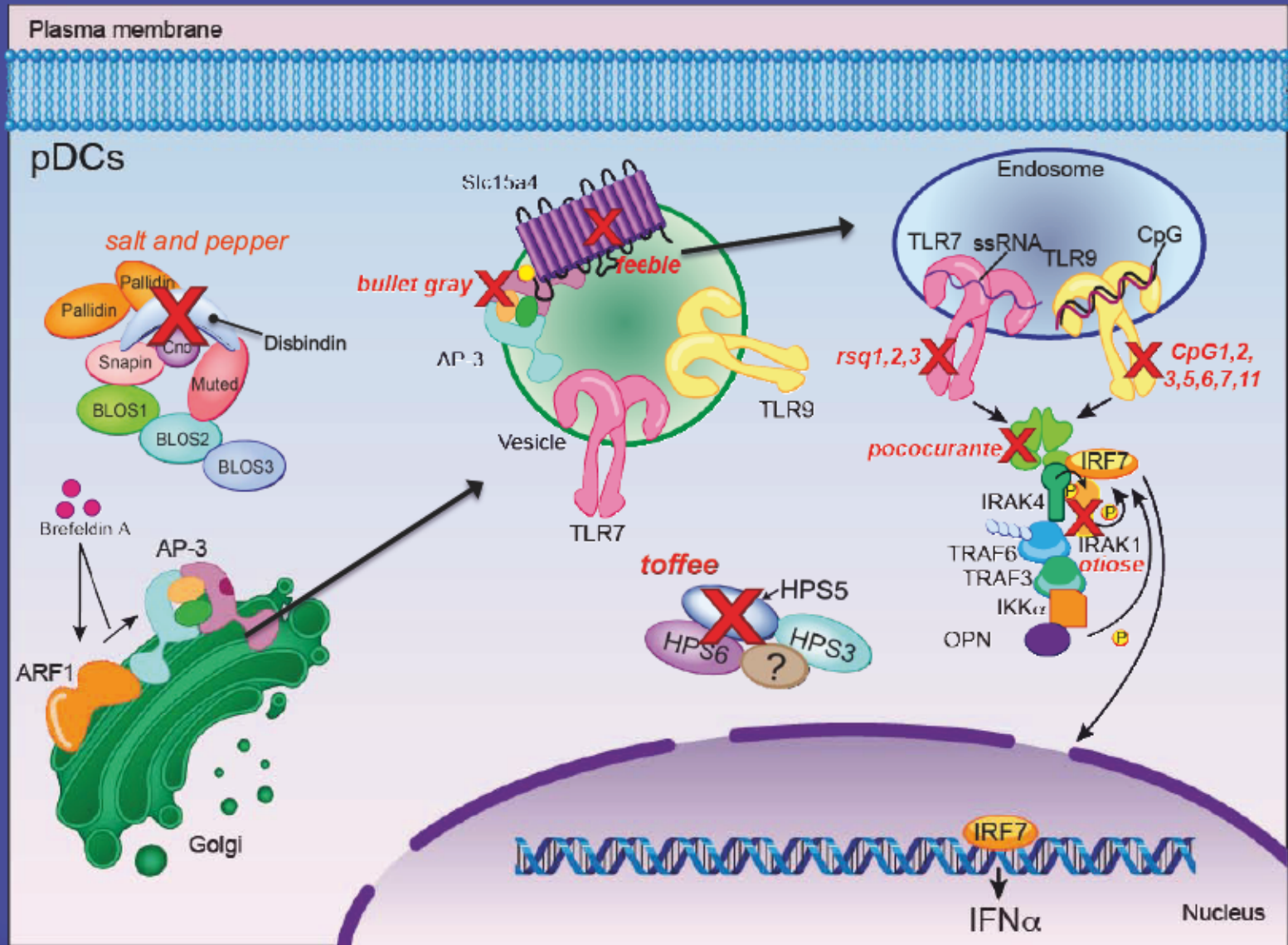
- The cost of sequencing a mouse genome to >90% coverage is now about \$3,000. About 4 mouse genomes can be sequenced per week in our lab.
- The price per base pair continues to drop by about 80% each year.
- With minimal mapping, it is possible to find mutations extremely quickly.

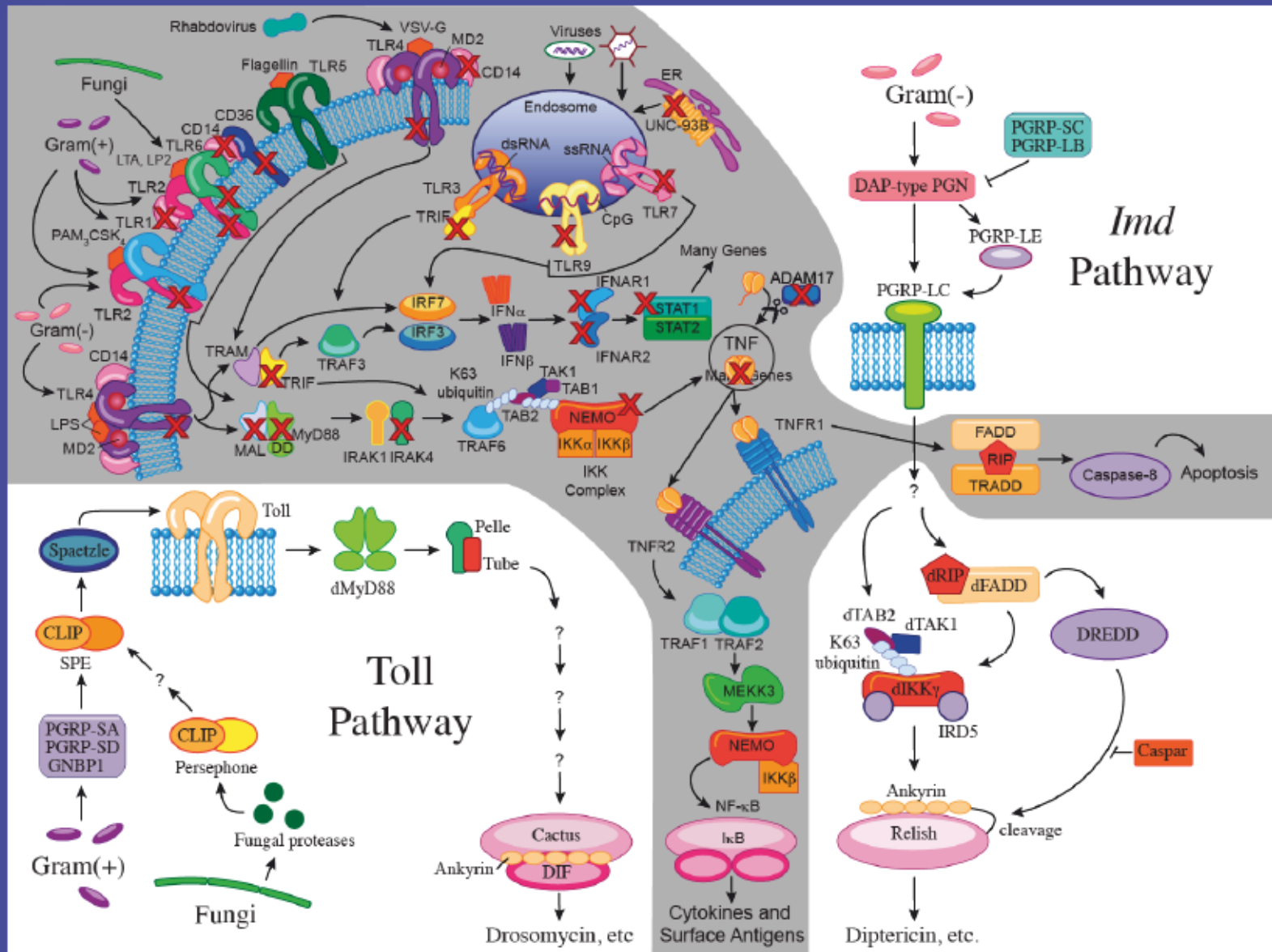


# From genetics to structures to mechanism...

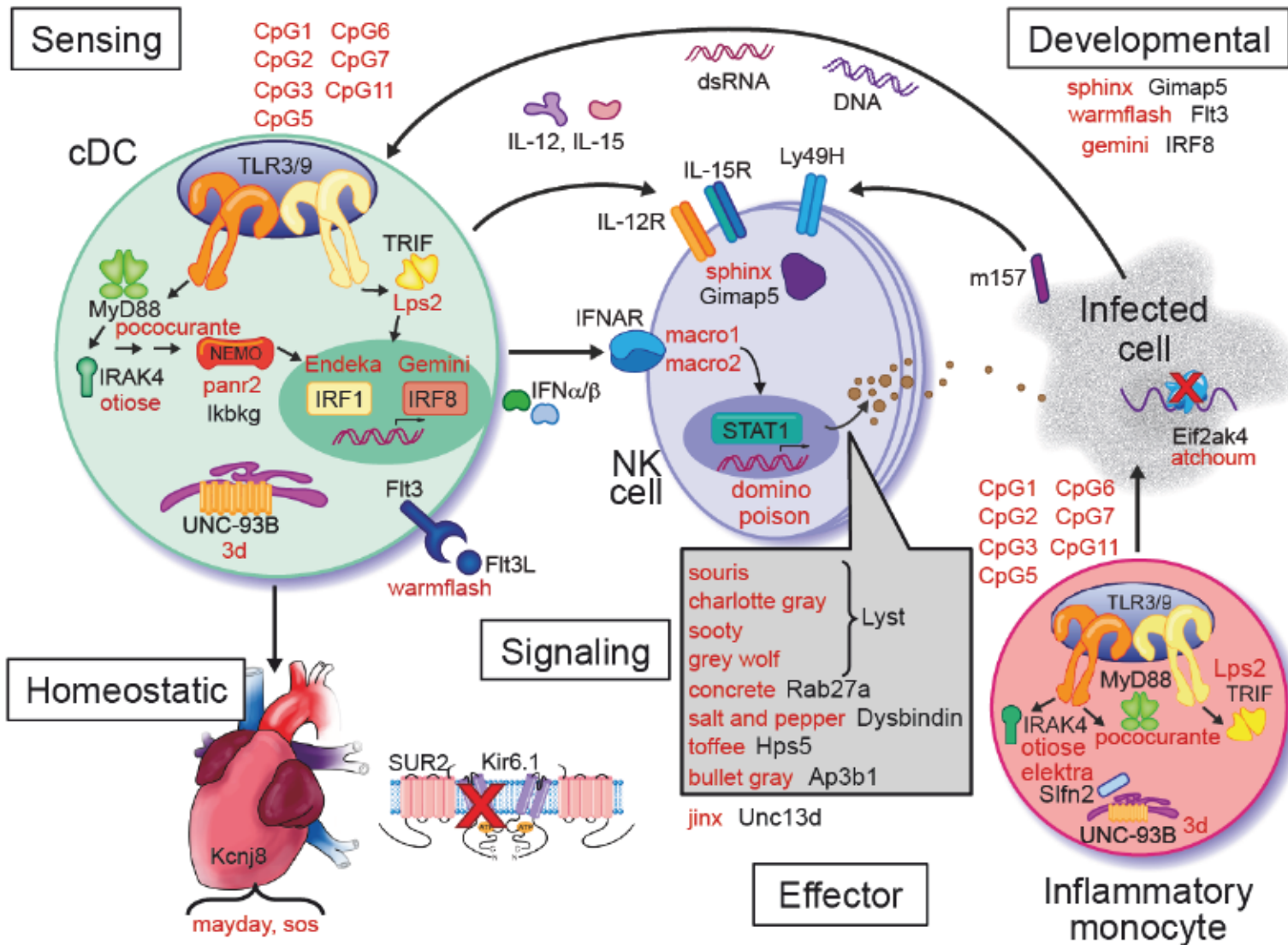


# In plasmacytoid dendritic cells, specialized machinery is needed for TLR signaling



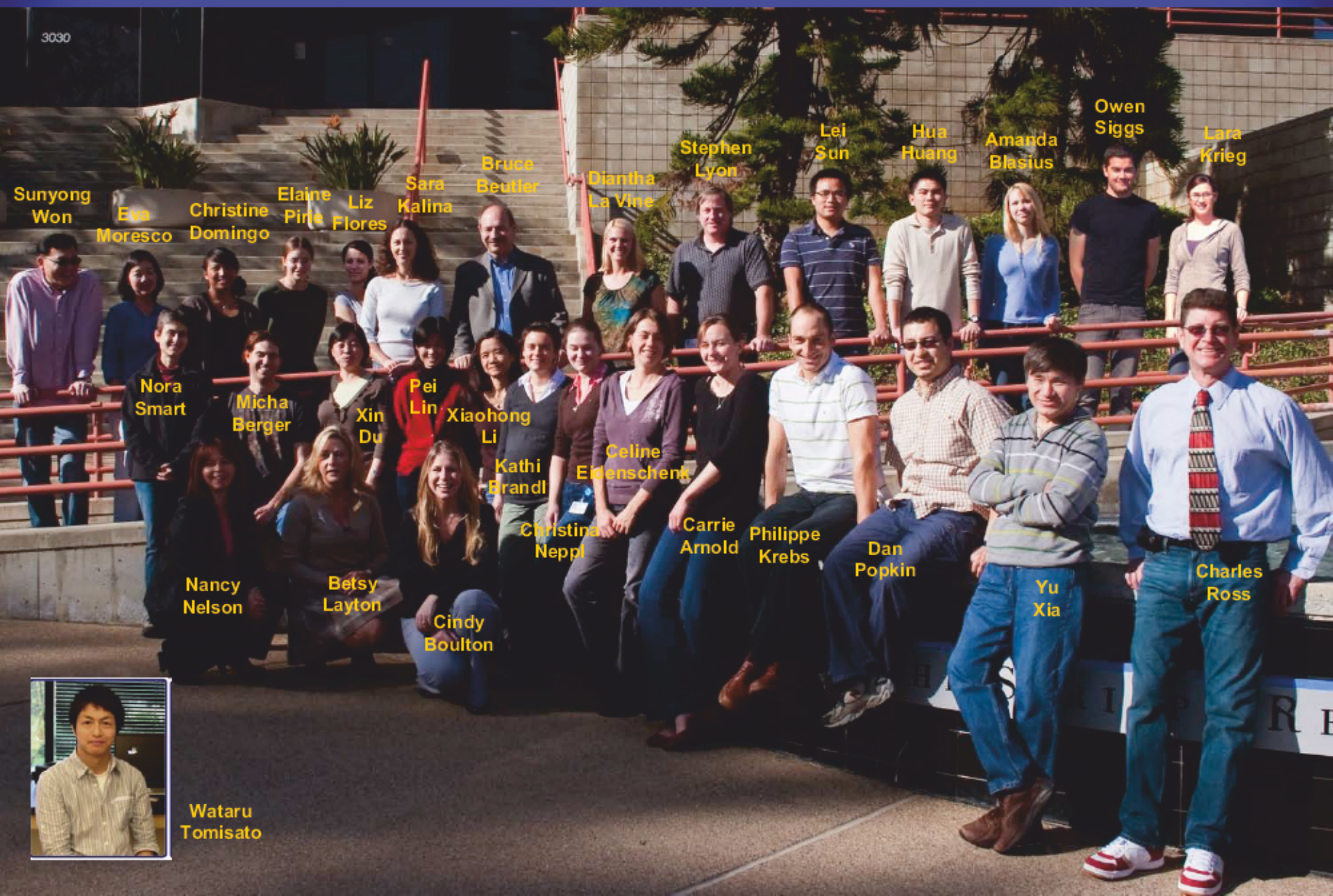


# Mutations we have found to cause MCMV susceptibility



# Beutler Lab, December 15, 2009

3030



Owen  
Siggs

Lara  
Krieg

Amanda  
Blasius

Hua  
Huang

Lei  
Sun

Stephen  
Lyon

Diantha  
La Vine

Bruce  
Beutler

Sara  
Kalina

Liz  
Flores

Elaine  
Pina

Christine  
Domingo

Eva  
Moresco

Sunyong  
Won

Nora  
Smart

Micha  
Berger

Xin  
Du

Pei  
Lin

Xiaohong  
Li

Celine  
Eidenschenk

Kathi  
Brandl

Carrie  
Arnold

Philippe  
Krebs

Dan  
Popkin

Yu  
Xia

Charles  
Ross

Cindy  
Boulton

Betsy  
Layton

Nancy  
Nelson



Wataru  
Tomisato

# Others not shown in the last slide

- Jiangfan Jiang
- Kasper Hoebe
- Ben Croker
- Koichi Tabet
- Karine Crozat
- Sophie Rutschmann
- Philippe Georgel



# For the cloning of the *Lps* locus

- Alexander Poltorak
- Irina Smirnova
- Christophe Van Huffel
- Betsy Layton
- Xiaolong He
- Mu-Ya Liu
- Xin Du
- Dale Birdwell
- Erica Alejos
  
- Chris Galanos and Marina Freudenberg (Max Planck Institute fur Immunbiologie, Freiburg)
  
- Paola Ricciardi-Castagnoli (University of Milan)