

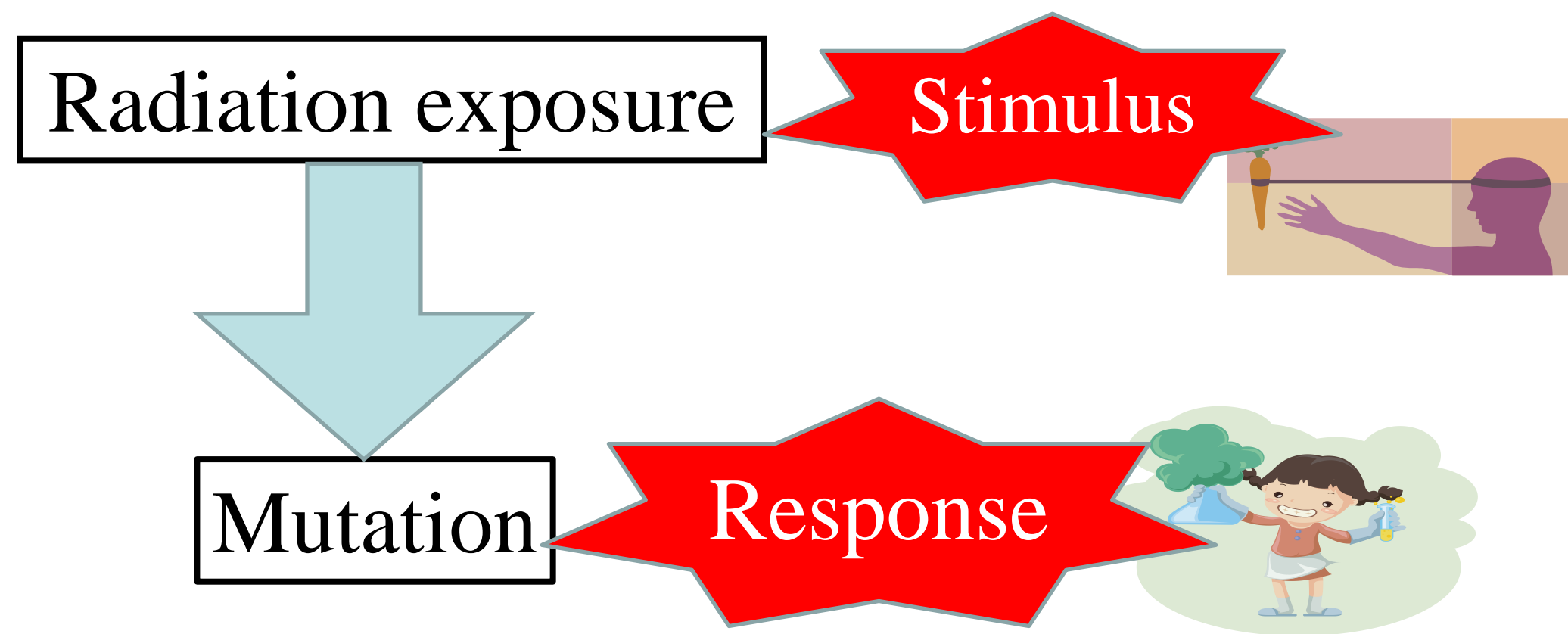
Whack-A-Mole Model: Towards a Unified Description of Biological Effects Caused by Radiation Exposure

Yuichiro Manabe¹, Takahiro Wada², Yuichi Tsunoyama³, Hiroo Nakajima⁴, Masako Bando^{5,6}

Quantum and Energy Engineering, Division of Sustainable Energy and Environmental Engineering, Graduate School of Engineering, Osaka University (Japan)¹,
 Faculty of Engineering Science, Kansai University(Japan)²,
 Division of Biology, Radioisotope Research Center, Kyoto University (Japan)³,
 Department of Radiation Biology and Medical Genetics, Graduate School of Medicine, Osaka University (Japan)⁴,
 Yukawa Institute for Theoretical Physics, Kyoto University (Japan)⁵, Research Center for Nuclear Physics, Osaka University (Japan)⁶

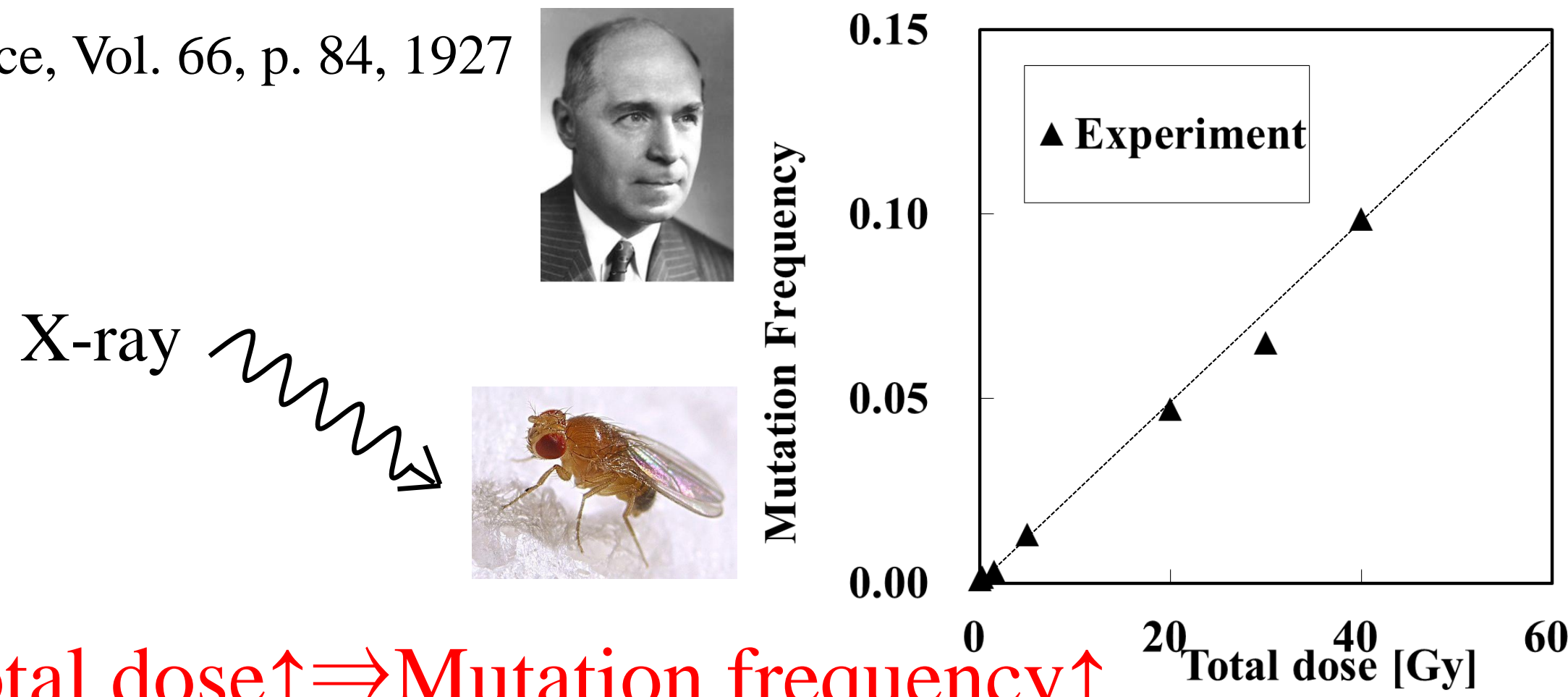
1. Introduction

Process of mutation



Drosophila experiment of Muller

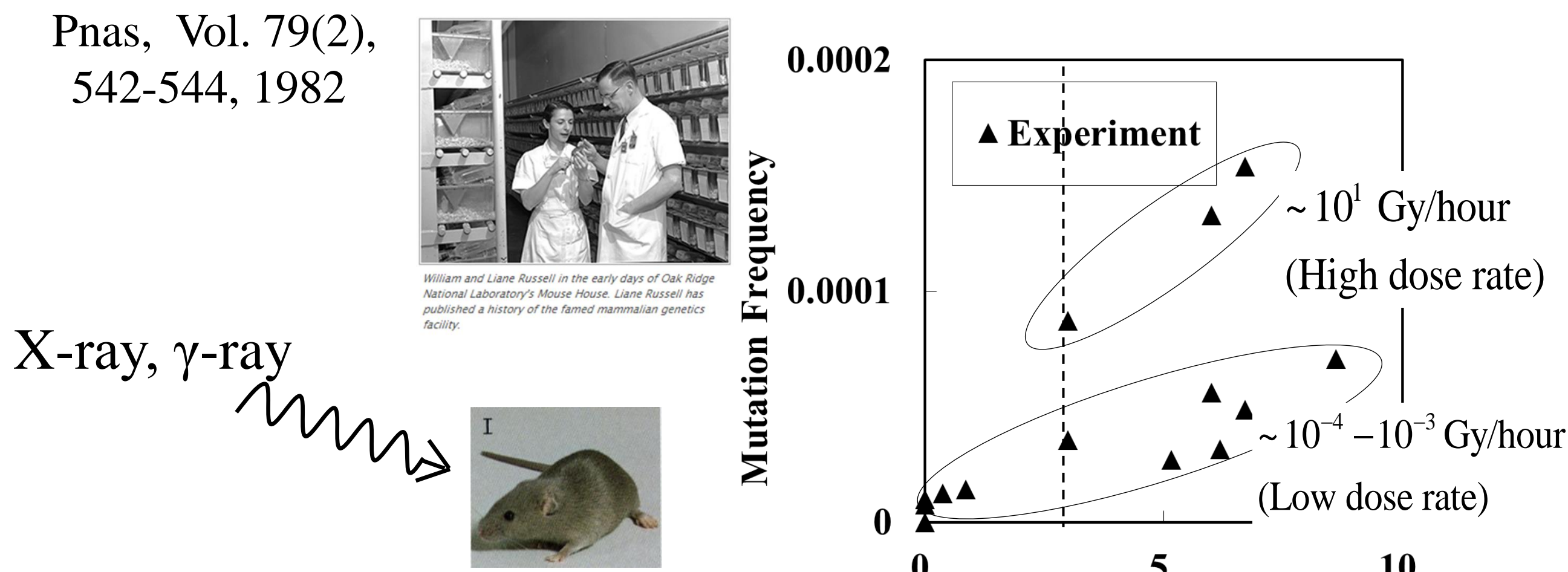
Science, Vol. 66, p. 84, 1927



Total dose $\uparrow \Rightarrow$ Mutation frequency \uparrow
 = LNT (Linear non-threshold)

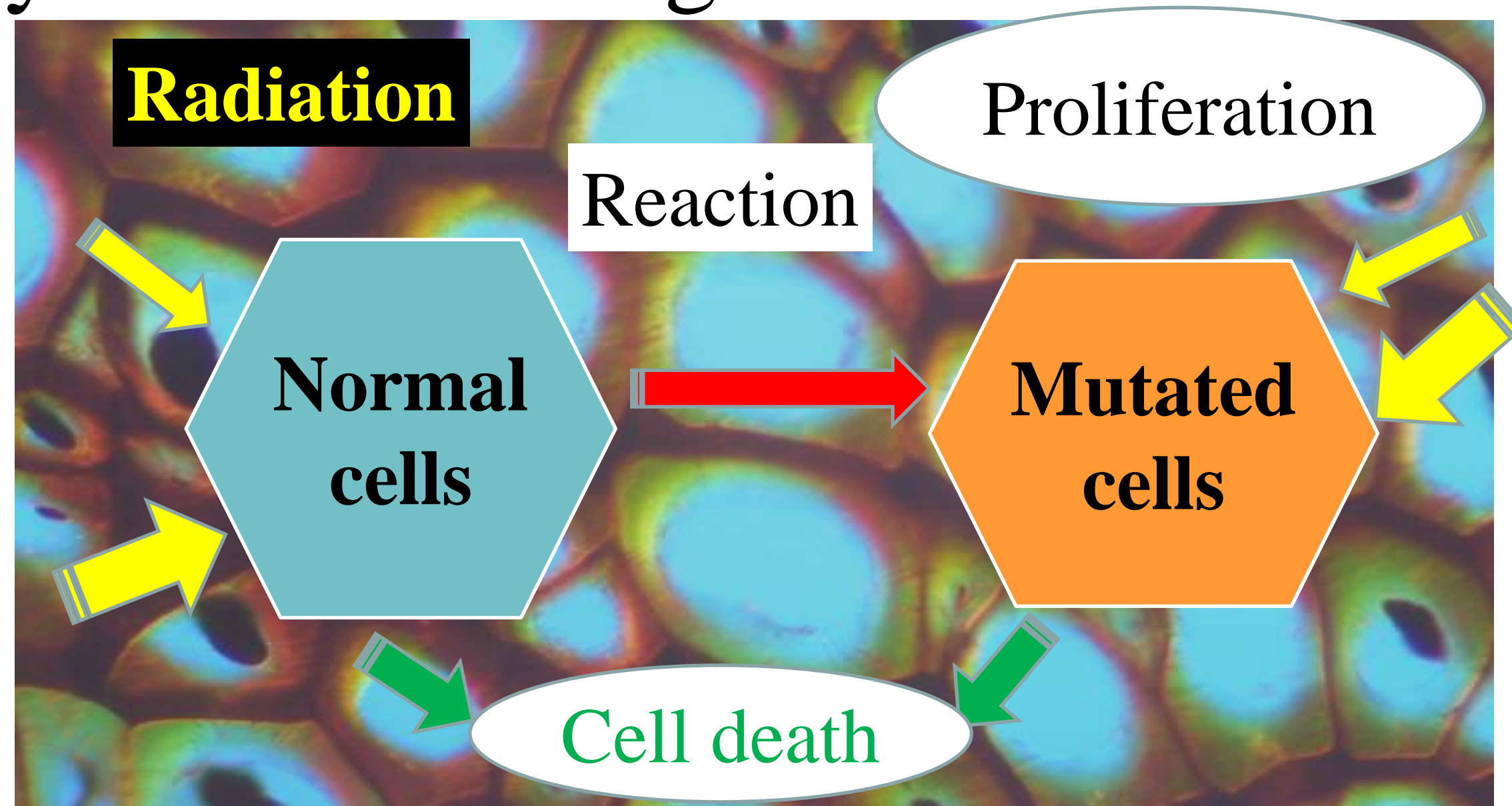
Mega-mouse experiment of Russell

Pnas, Vol. 79(2), 542-544, 1982



Dose-rate $\uparrow \Rightarrow$ Mutation frequency \uparrow

System consisting of cells



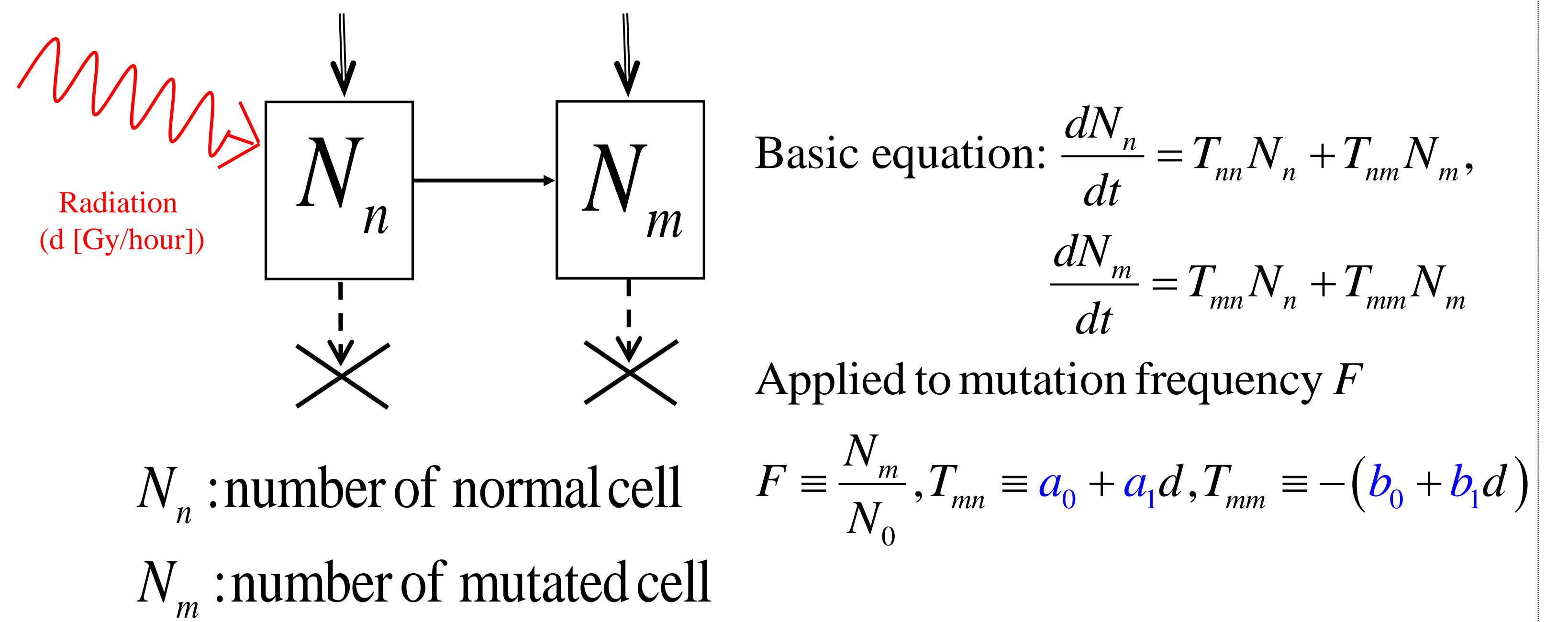
A variety of mechanisms works

We need a mathematical model considering dose rate dependence and a variety of mechanisms at cell level

2. Mathematical formula

Whack-A-Mole (WAM) Model

1. Important variable is dose-rate
2. Transition between Normal cell and mutated cell
3. Proliferation, cell death, recovery effect



Equation for mutation frequency

Preventative effects :

Mutation frequency is expressed as a differential equation with respect to time.

$$\frac{d}{dt} F(t) = (a_0 + a_1d) - (b_0 + b_1d) F(t)$$

~~$$\frac{dF}{dD} = c \frac{F_0}{D_0}$$~~

F : mutation frequency

d : Dose rate [Gy/hour]

D : Total dose [Gy]

Solution for mutation frequency

$$F(t) = F(\infty) \left(1 - e^{-(b_0 + b_1d)t}\right) + F(0) e^{-(b_0 + b_1d)t}$$

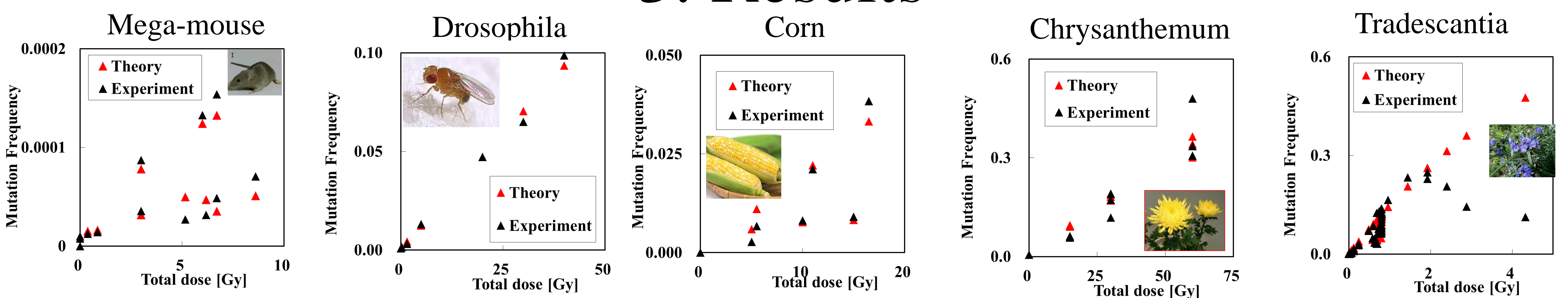
$$F(\infty) = \frac{(a_0 + a_1d)}{(b_0 + b_1d)}$$

a_0, a_1, b_0, b_1

4 parameters

We determine four parameters by χ fit

3. Results



4. Conclusion

1. We have analyzed the experimental data of five living organisms; mouse, drosophila, chrysanthemum, maize, and tradescantia.
2. Despite the difference between animal and plant, all these data reasonably fall on our predicted value.
3. We can apply this model to the scheduling of cancer therapy and radiation protection of nuclear plant workers.