

Computer modeling of radiation effects

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Table of Contents

1. Introduction
2. Simulation of DNA strand breaks by ionizing radiation
3. Molecular dynamical study of the DNA lesion repair
4. Modeling and simulation of the cellular level tumorigenesis
5. Conclusion

1. Introduction

❖ Radiation Effects ?

- ❖ Deterministic effect
 - organ/tissue damage (or death)
- ❖ Late time (stochastic) effect
 - radiation induced cancer

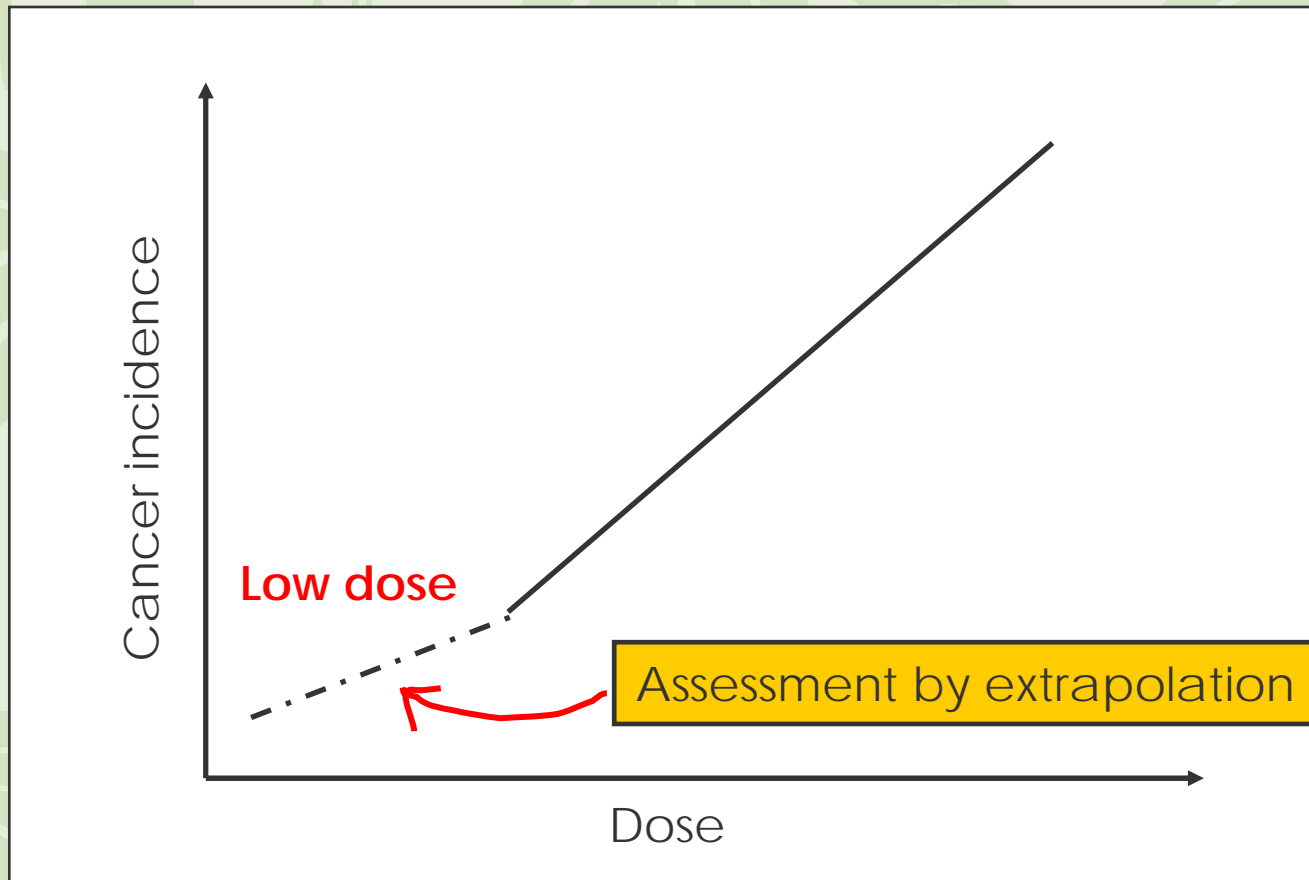
High Dose effect

At low dose region, quantitative risk estimation are not so easily obtained.

❖ Low dose radiation risk

- ❖ risk = probability of cancer incidence

Dose-Response



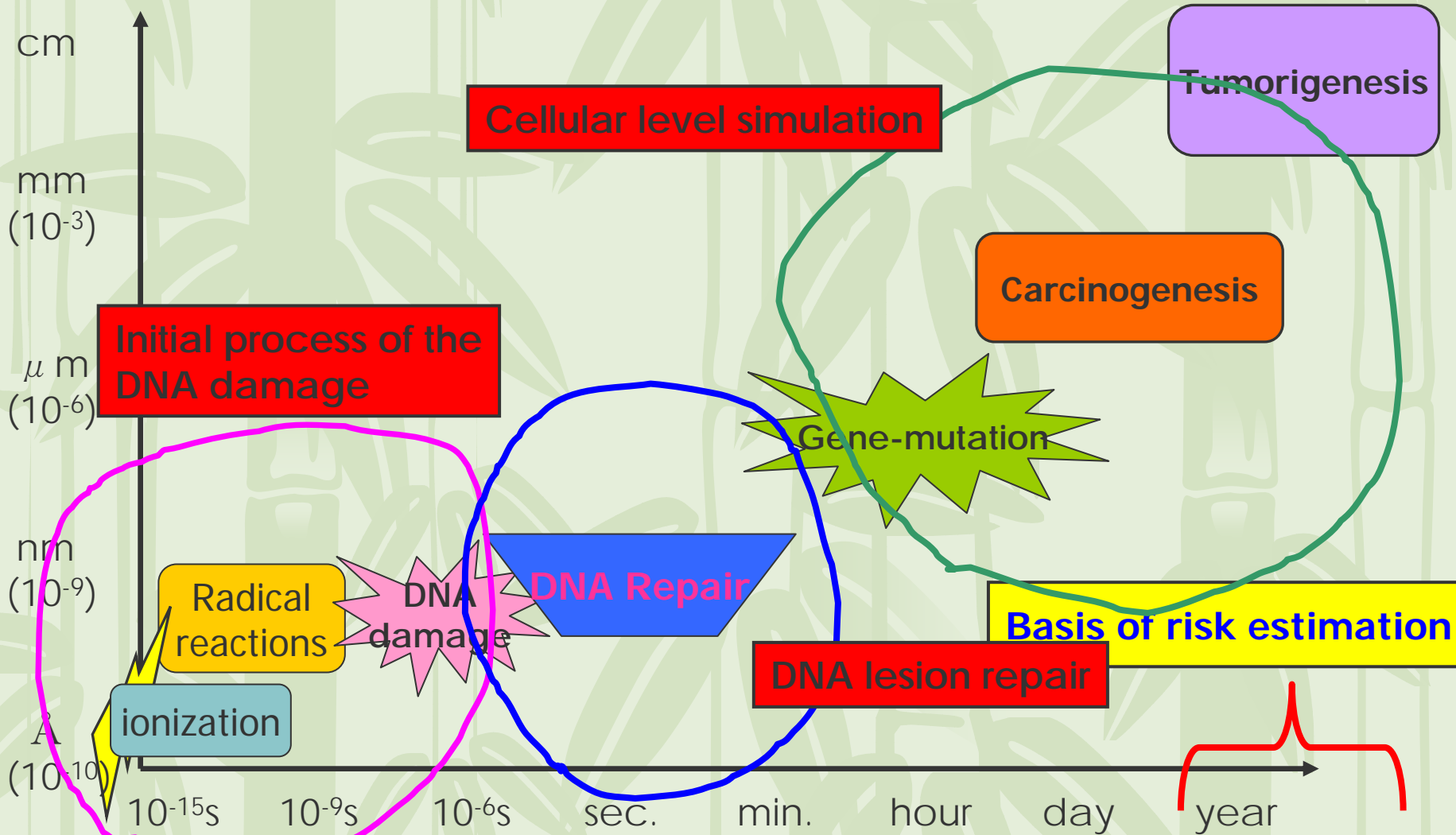
Risk estimation at low dose radiation needs further study based on the Biological mechanisms.

What is “low dose”?

- ❖ Experimental viewpoint
 - ❖ $< 100\text{mSv}$
 - ❖ Limit of the observation of radiation effects.
- ❖ Average annual effective dose of radiation workers = $\sim 15\text{mSv}$
- ❖ Various suggestions: $10\text{mSv} - 100\text{mSv}$

The definition of low dose is physically and operationally ambiguous, only some effect-based guidelines have been suggested.

Scale of the++



Check point #1

1

2. Initial process of radiation induced DNA Damage

Radiation to the cell nucleus causes damage to DNA

Biologically important damage



Single Strand Break (SSB)



Double Strand Break (DSB)

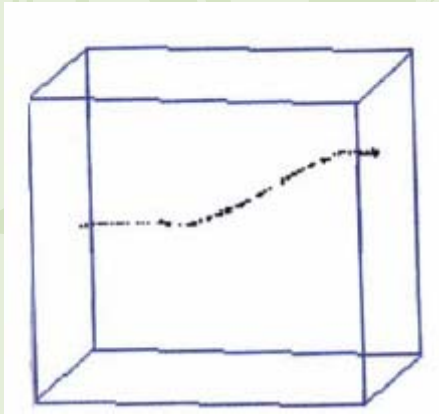
Question:

What kind of radiation with what type of track generate how much damages ?

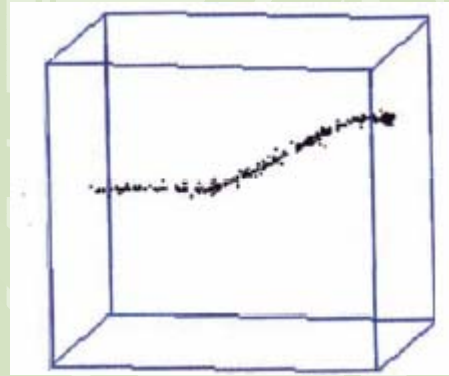


To clarify the relations between track structure and DNA strand breaks.

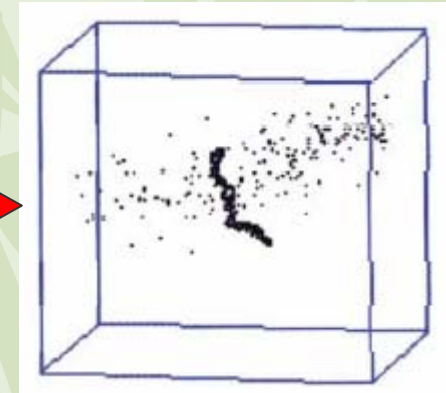
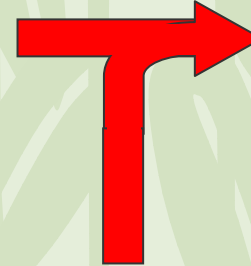
Simulation method



Track structure

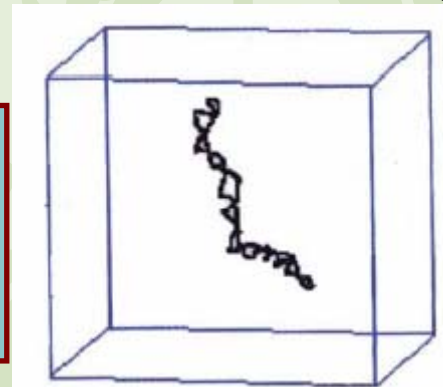


Radical production



Radical diffusion

1. Track structure calculation
2. Radical production
3. DNA modeling
4. Calculating DNA and radical reactions

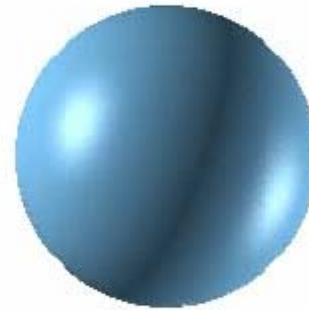


Target DNA modeling

Track structure: spatial distribution of energy deposition of ionizing radiation

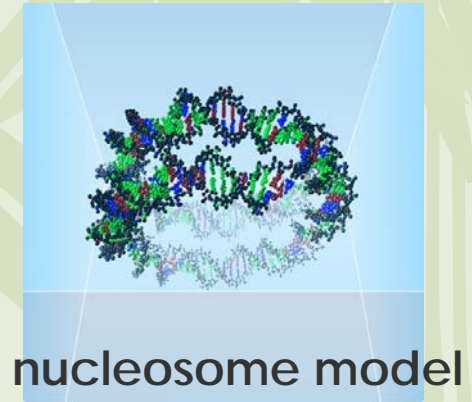
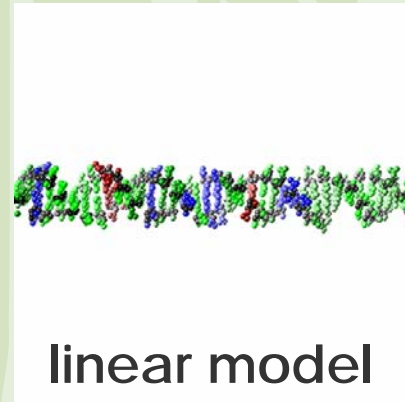
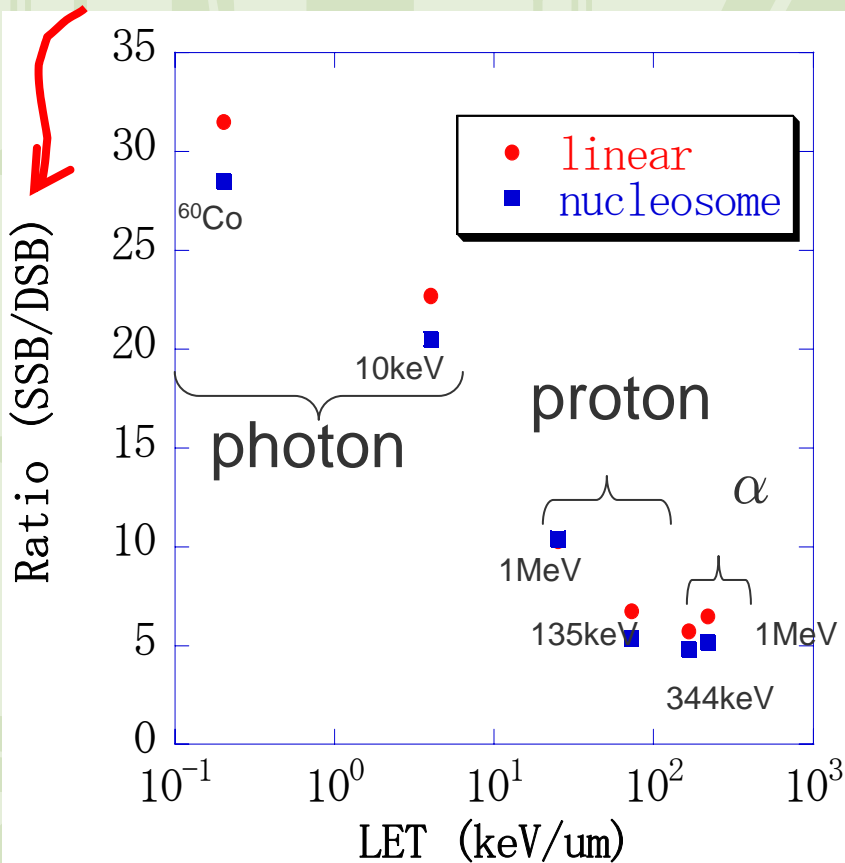
Simulation example

DNA damage induction
simulation
(proton + solenoid DNA)



Result [SSB/DSB ratio]

Indicator of complexity of DNA damage



LET [Linear Energy Transfer] -
energy deposition by the charged
particle per unit path length

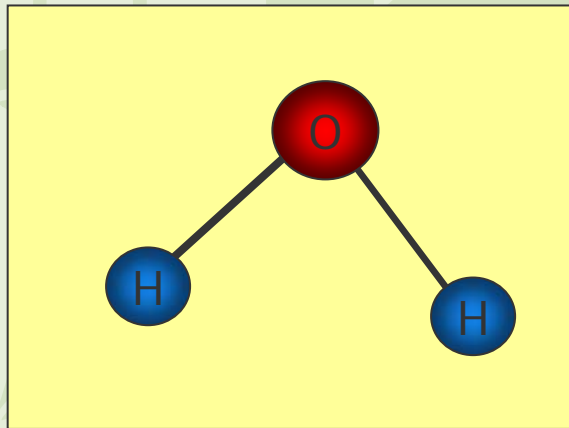
DSB yield increasing with LET up to 100 keV/ μ m

Check point #2

2

3. Molecular dynamical study of the DNA lesion repair

Molecular Dynamics simulation



$$F_i = -\frac{\partial V(\mathbf{r}_i)}{\partial \mathbf{r}_i} = m_i \frac{\partial^2 \mathbf{r}_i(t)}{\partial t^2}$$

\mathbf{r}_i Position of each atoms (i)

m_i mass

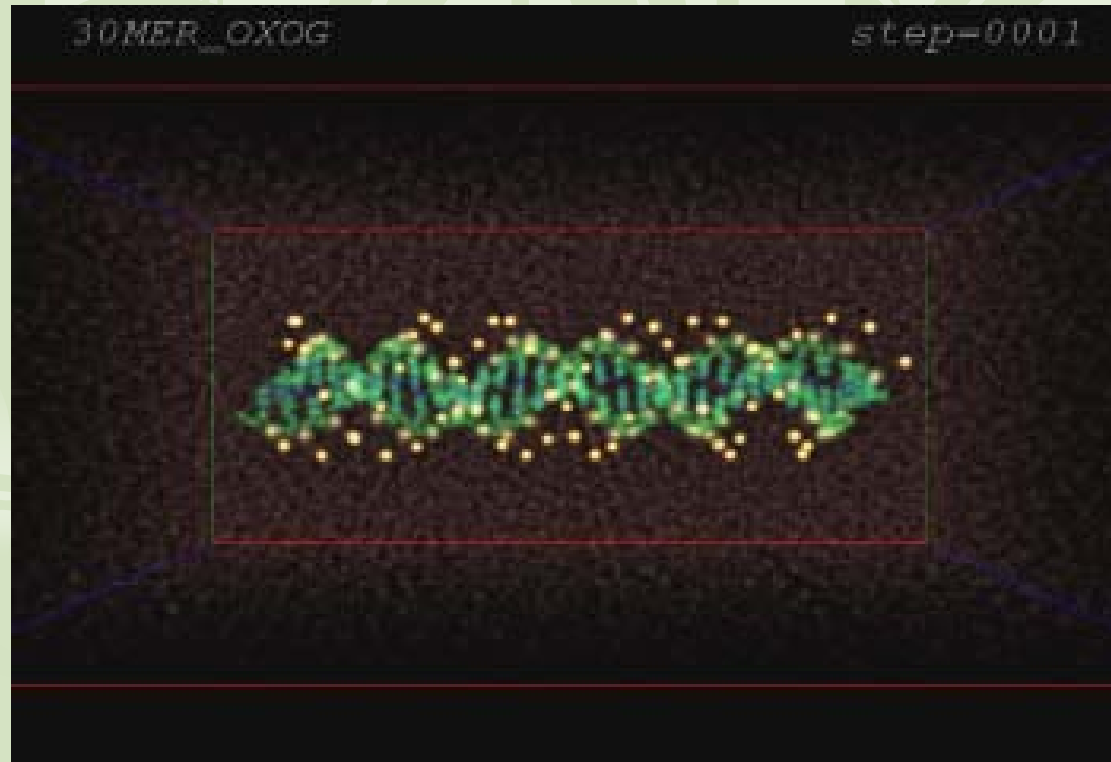
F_i Force acting on atom i

$V(\mathbf{r}_i)$ Potential energy of the system

$(\mathbf{r}_i(0), \dot{\mathbf{r}}_i(0))$ Initial condition (configuration)

To clarify a dependency between damaged DNA structural change and capability of the DNA repair.

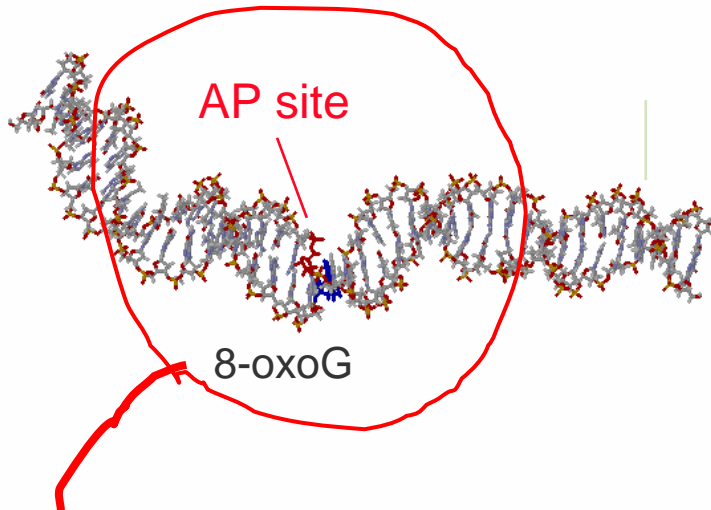
Simulation example



Shape change of damaged DNA

Damaged DNA: 8oxo-G + AP site

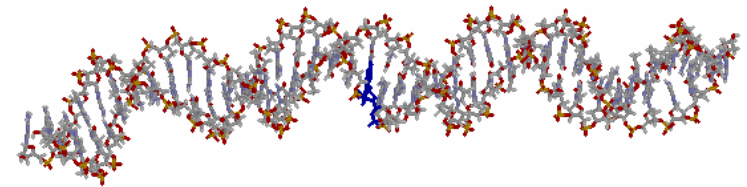
1.3 ns



Clustered damage

Native DNA (no damage)

2.0 ns



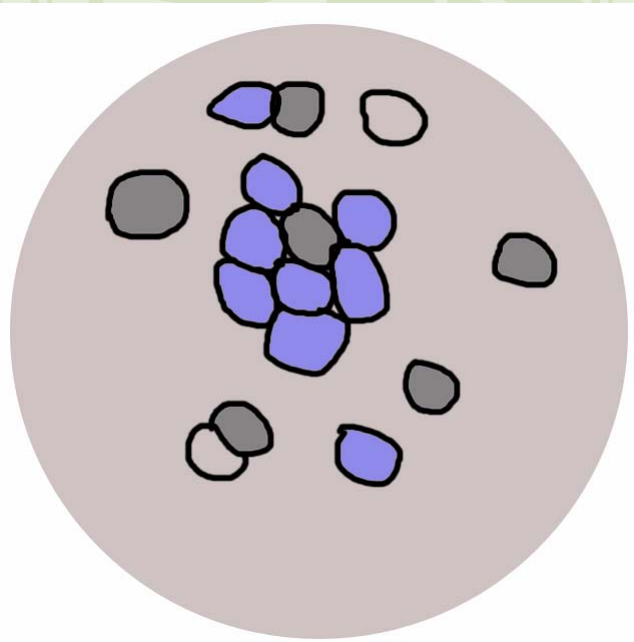
- Damaged DNA shows bending movement at lesioned site
- Dynamic analysis of DNA structure is ongoing.

Check point #3

3

3. Modeling and simulation of the cellular level tumorigenesis

The dynamics of the carcinogenesis is studied by the simulation of the cell group in the cell level.



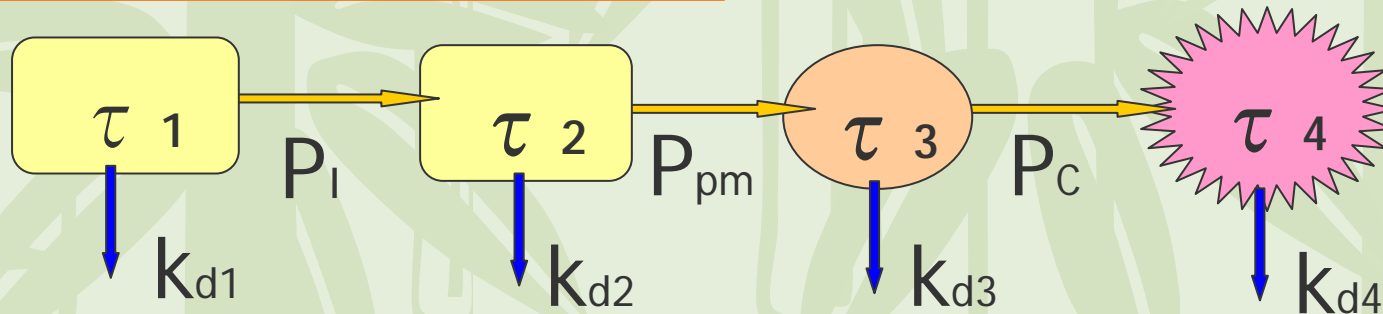
- ❖ Same configuration with Cell culture system
- ❖ Can study colony formation or tumorigenesis.
- ❖ Can introduce dynamical based group effect



- Easily comparable with the experiments.
- Molecular biologically based model.

Intracellular dynamics

Cell transformation



k_d : Prob. of cell death

P_I, P_{pm}, P_C : prob. of cell state change (genetic)

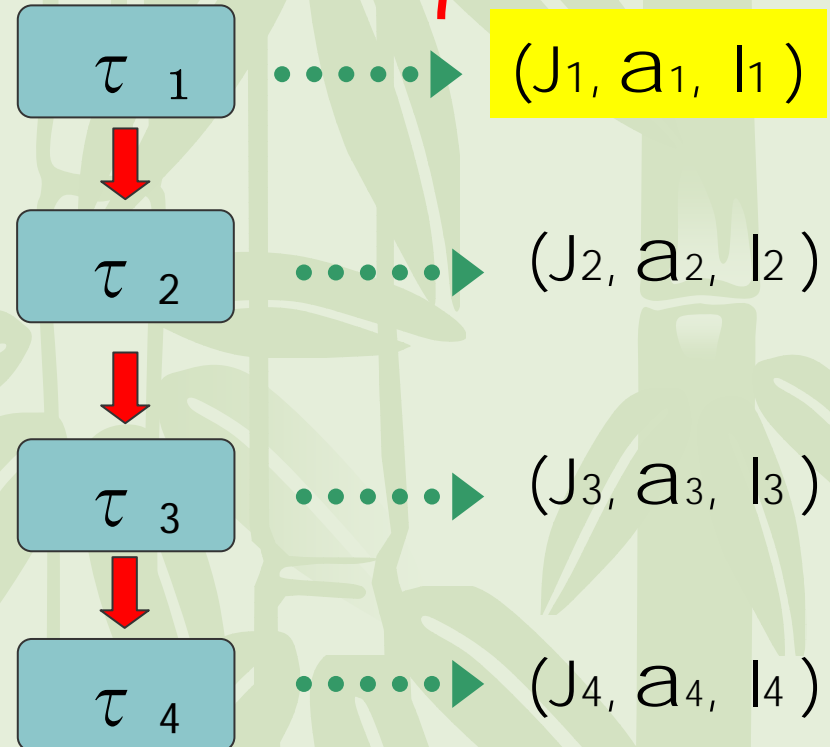
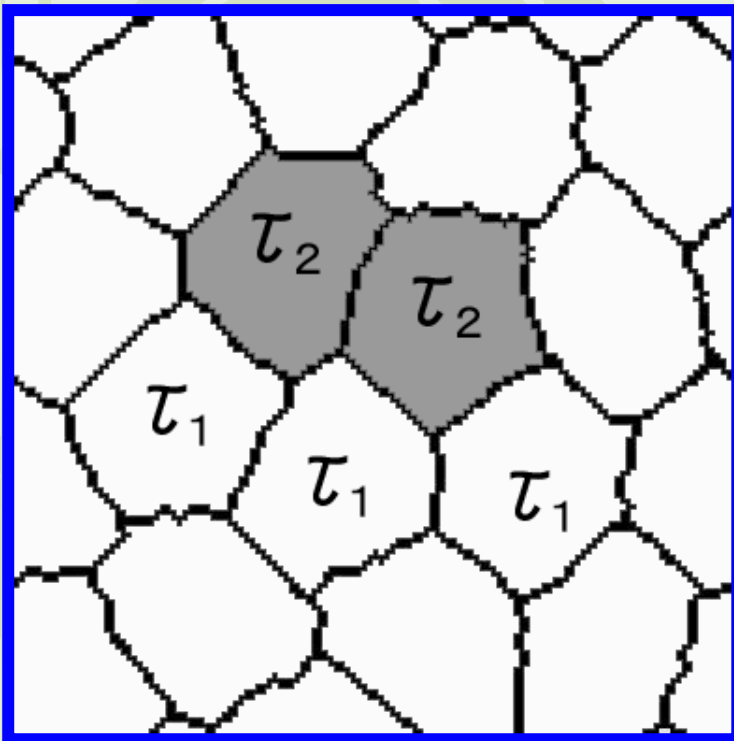
cell state τ : normal, initiation, promotion, cancer

Cell division

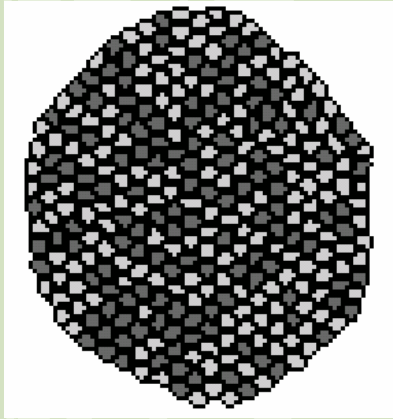
If $a(s) > a_c$ then cell division occur

Details of the model

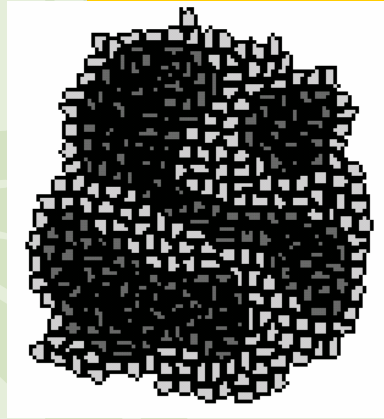
Intracellular state change affects the physical parameters (cell adhesion molecule, cell membrane)



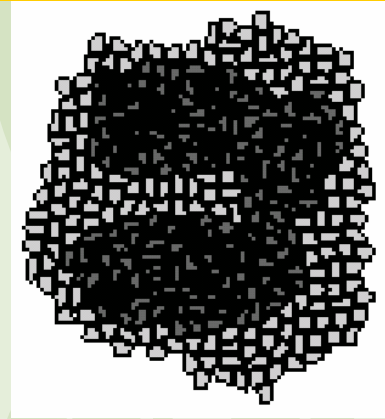
Spatial patterns (cell sorting)



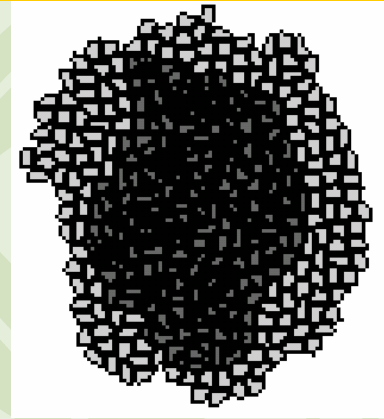
Initial



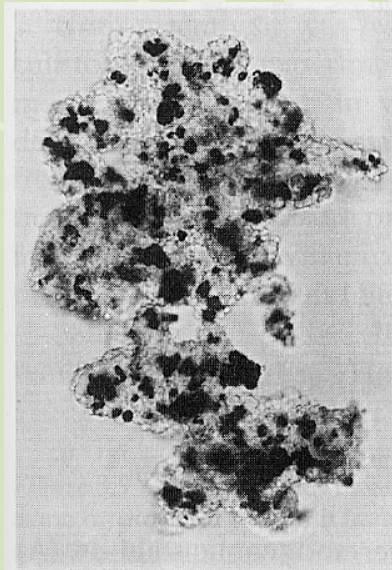
500steps



3000steps



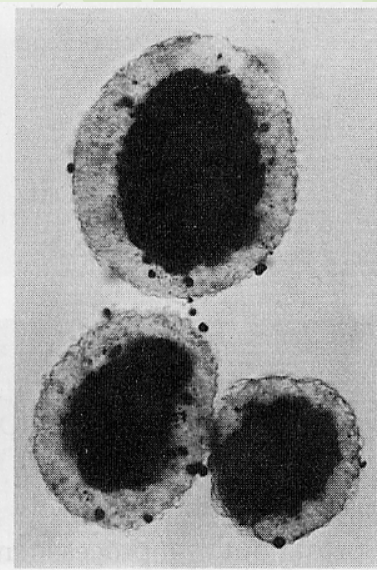
8000steps



(A)

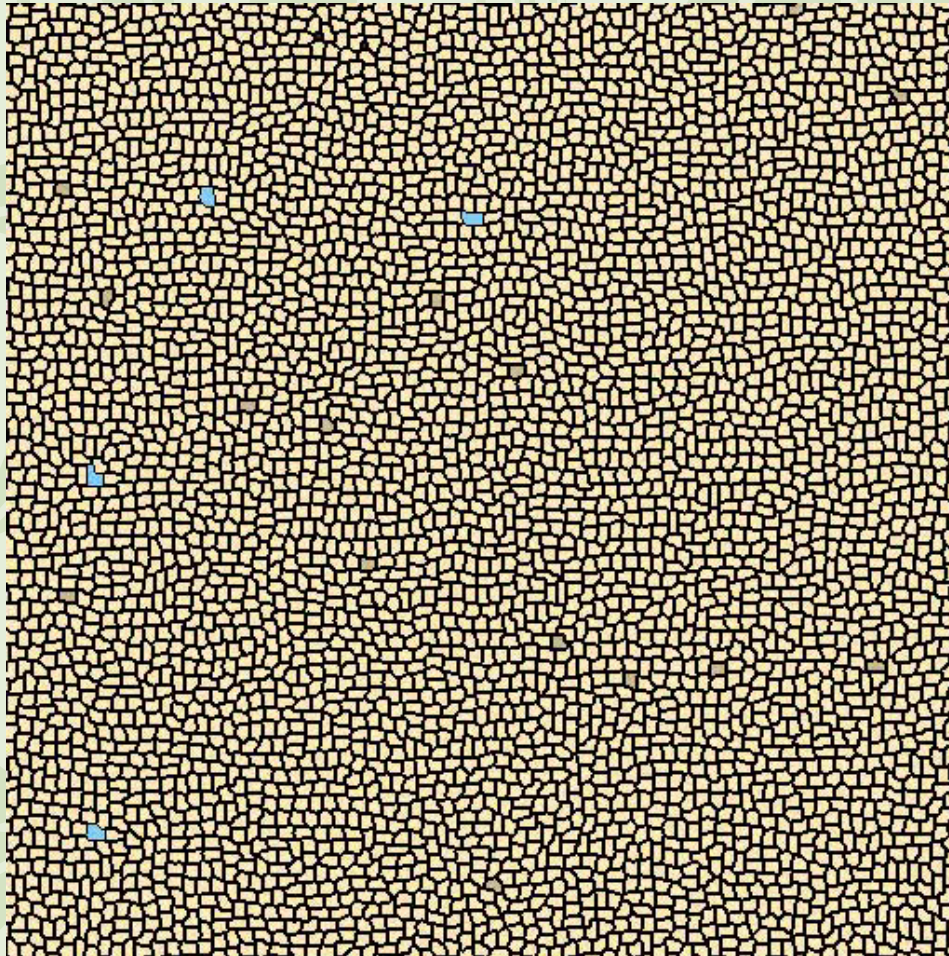


(B)



(C)

Simulation example



Medium



Normal cell

τ_1



Initiated cell τ_2



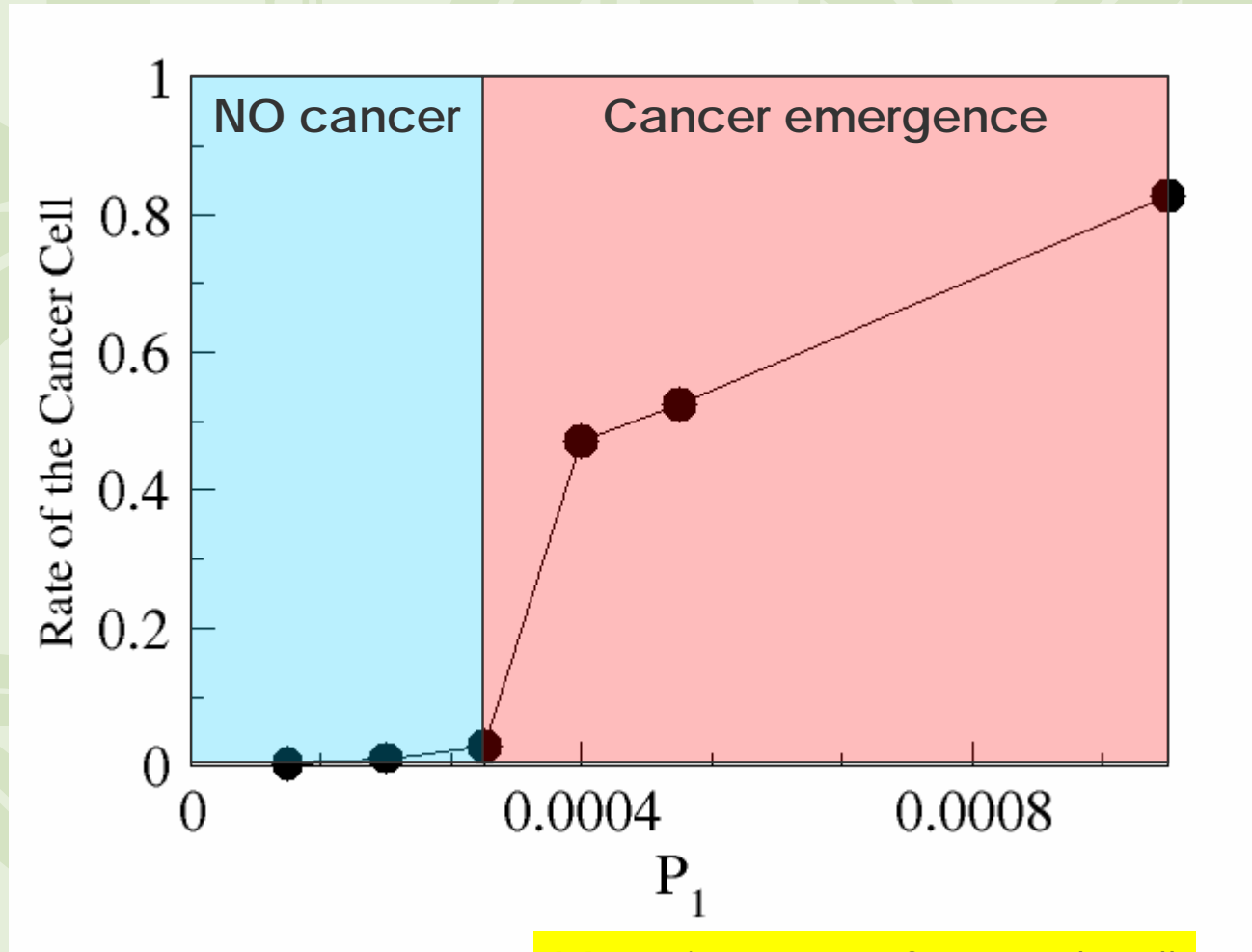
Progressed cell τ_3



Cancer cell τ_4

Large mutation rates are used for the time limitation.

Mutation rate vs. Cancer cell production



Mutation rate of normal cell

Conclusion

- ❖ Our ongoing study about initial to cellular level biological radiation effects using computer modeling and simulations is showed.
- ❖ LET dependency of the DNA damage complexity is studied.
- ❖ Relationship between structural change of damaged DNA and its repair is studied.
- ❖ Cellular level dynamics of the carcinogenesis is modeled and parameter (mutation rates) dependency is examined.

Thanks!

JAEA

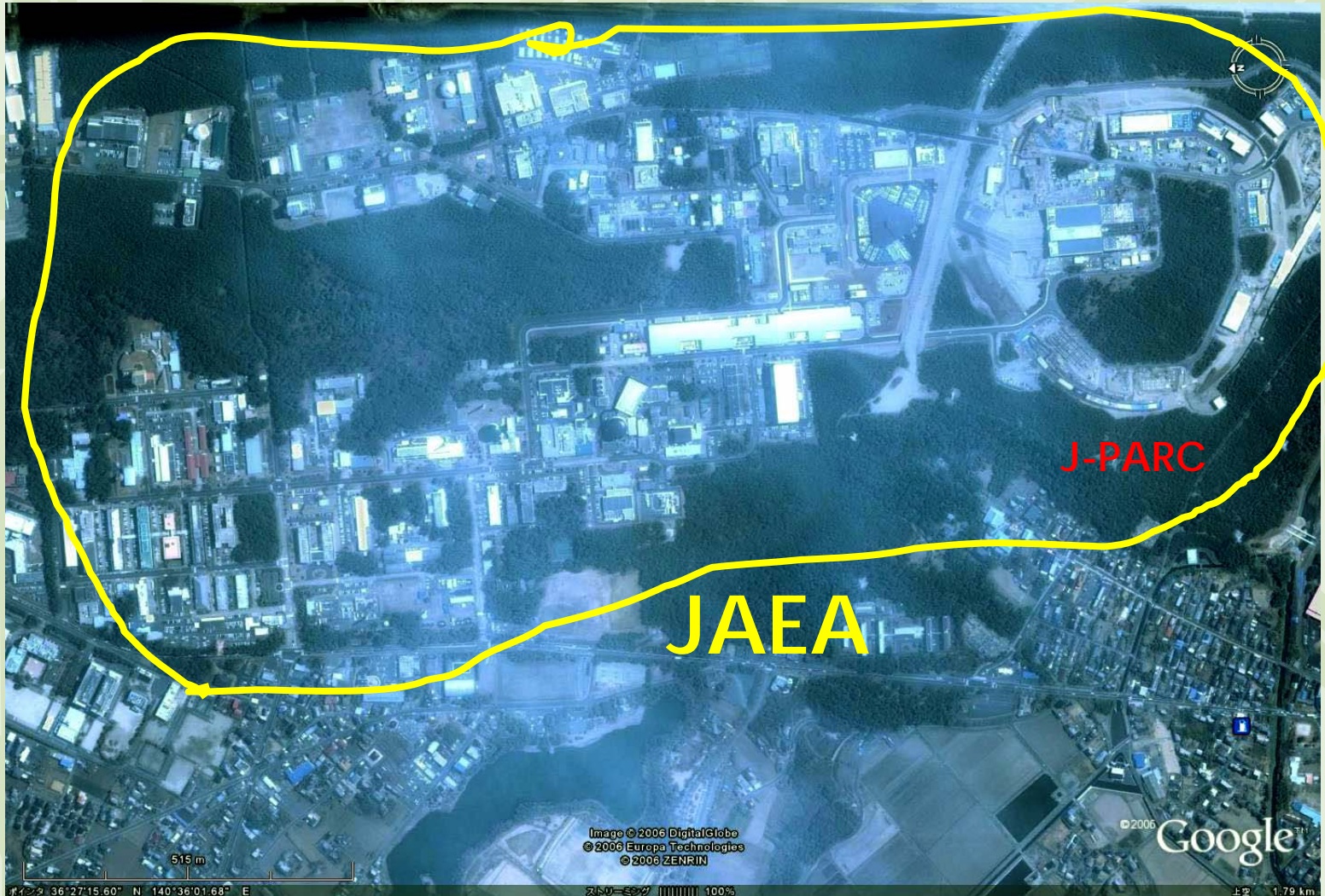
Dr. Ritsuko Watanabe : Simulation of DNA damage induction
Dr. Miroslav Pinak : Simulation of DNA repair
Dr. Julaj Kotulic Bunta : Simulation of Ku70/80 binding
Dr. Mariko Higuchi : Simulation of multiple lesioned DNA

NIID

Dr. Hideaki Maekawa : DNA damage induction experiment
Dr. Hirofumi Fujimoto : DNA repair simulation

NIRS

Dr. Manabu Koike : DNA repair experiment



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