

Pôle Physique



IN2P3  
Les deux infinis



Geant 4



# The IRIS Project: The long-term impact of low dose radiation on living systems

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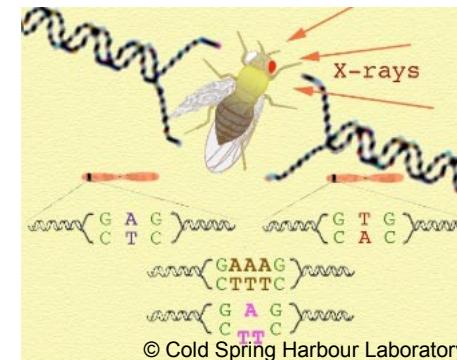
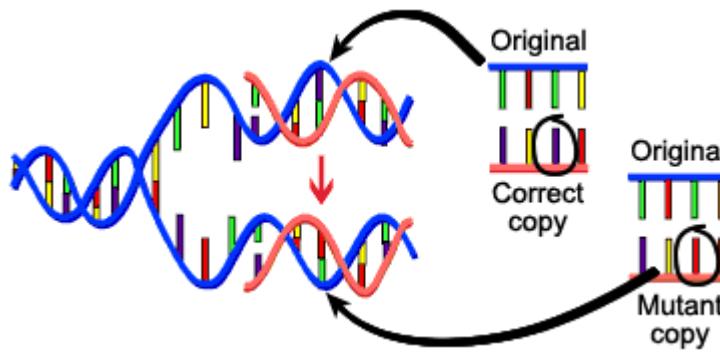
D.G. Biron, M. Coulon, M. Davidkova, T. Hindré,  
S. Incerti, L. Maigne, P. Micheau, **F. Piquemal,**  
**G. Warot**

DULIA-Bio, October 12, Canfrac, Spain

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# Motivation

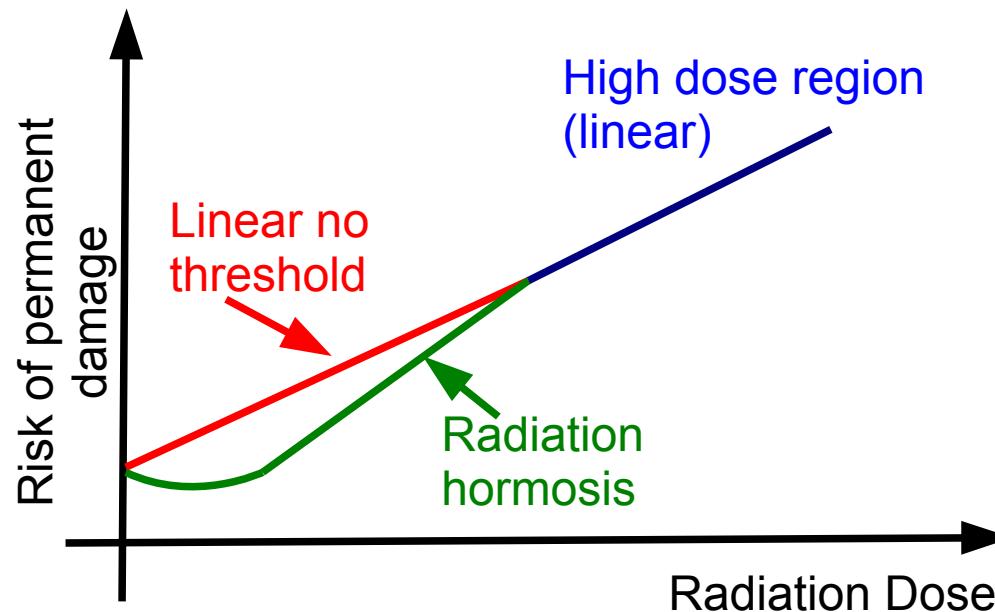
Mutations are the fundamental ingredient that drives evolution.



What role does radiation have to play in evolution compared to transcription and bio-chemistry.

# Motivation

How does natural radiation impact life?



Role in the oxidation response?

Triggering unlikely mutations?

Are radiation environments correlated with evolution events?

# Summary

- Biology in the LSM.
- What is a Long Term Evolution Experiment?
- The radiation background for Biology
- LTEE results from Clermont-Ferrand
- Simulating DNA damage with low background

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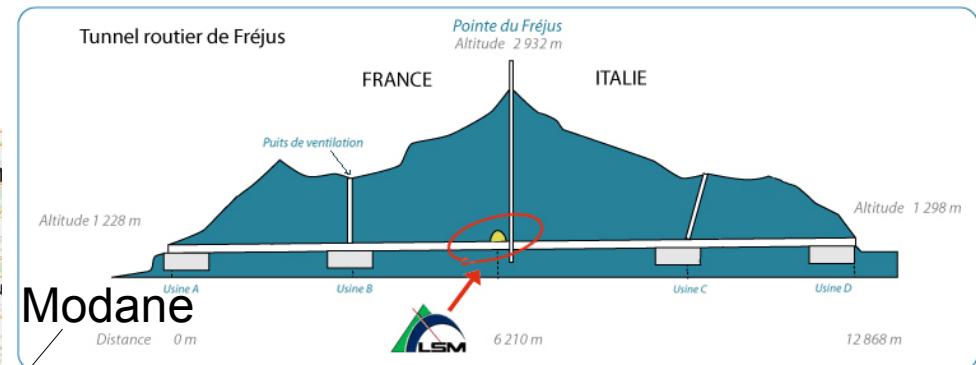
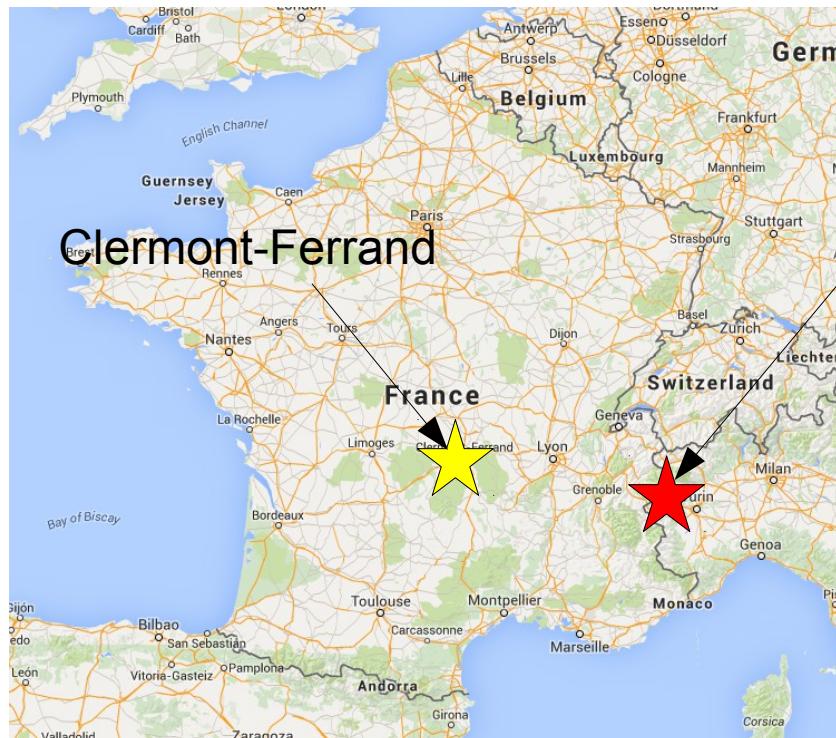


Environnement

Santé

**Biology at the LSM**

# The Laboratoire Souterrain de Modane



## The lab at a glance:

- Inside the Fréjus Tunnel, bordering Italy and France
- Under 1700m of rock (4800m water equivalent)
- Volume of  $3500 \text{ m}^3$
- Low radon level ( $15 \text{ Bq m}^{-3}$ )
- Home to:
  - 17 germanium detectors
  - EDELWEISS (dark matter)
  - SEDINE (dark matter)
  - SUPER-NEMO (neutrino-less double beta)

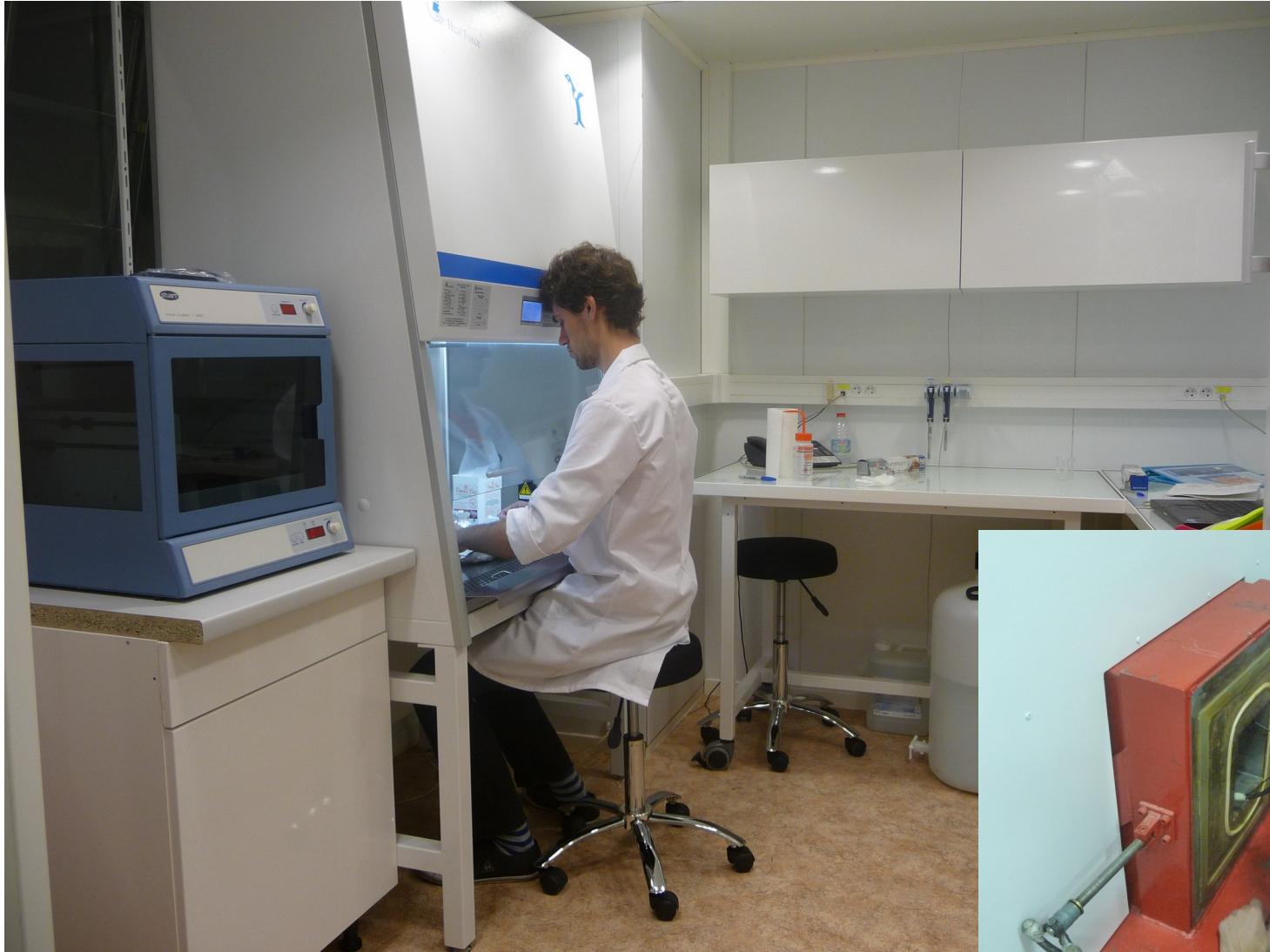
## Biology at the LSM



# Biology at the LSM



## Biology at the LSM



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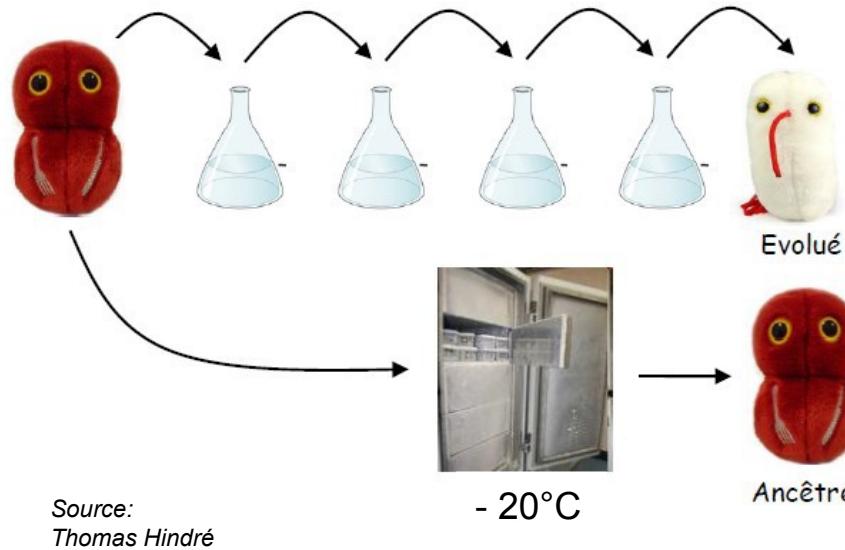


Santé

## Experimental Background

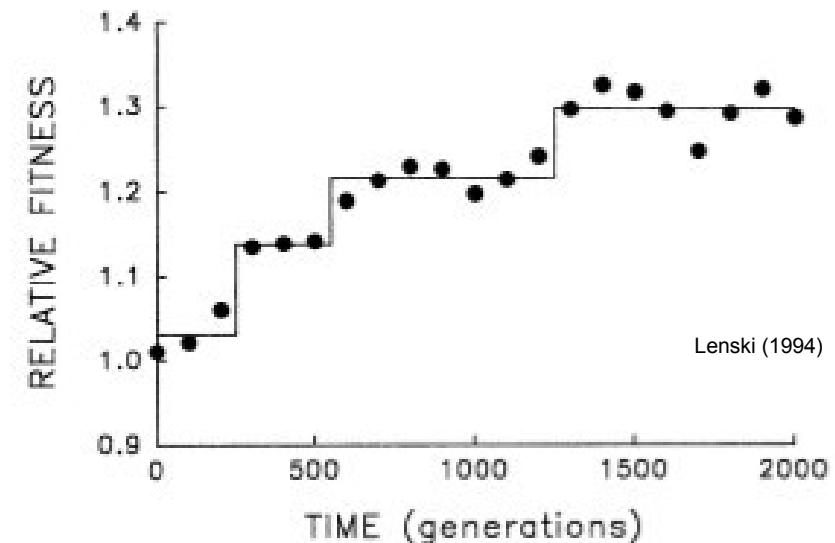
# Experimental Background

- The Lenski et al.(1991) Long Term Evolution Experiment is able to identify when beneficial mutations become dominant in a population.



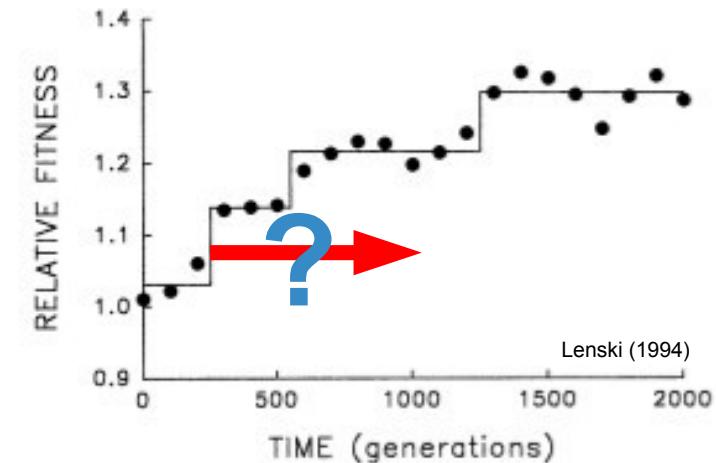
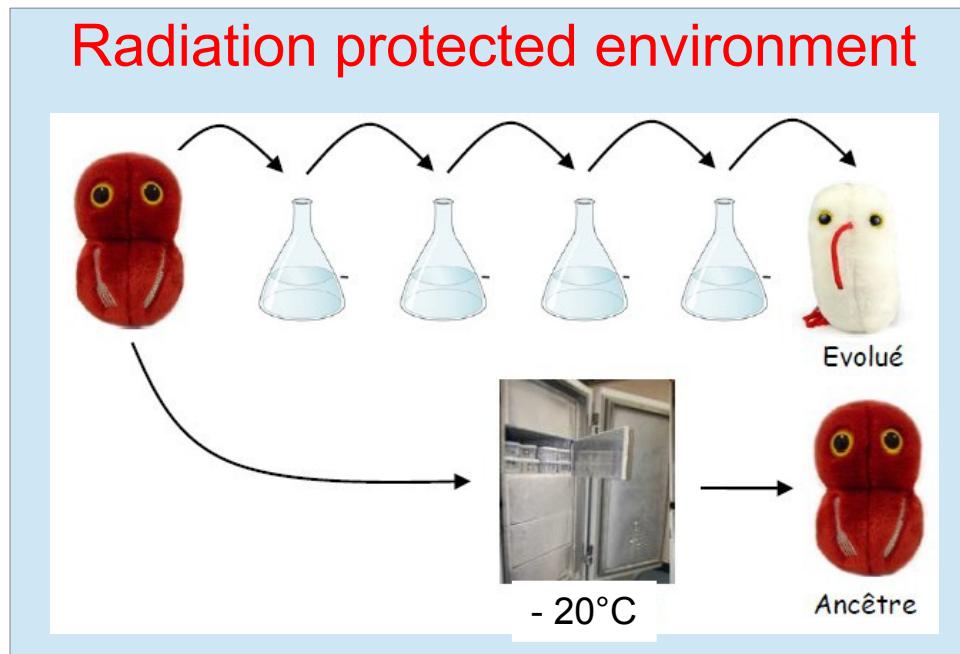
# Experimental Background

- Step-like changes in the bacterial behaviour can be noticed following this method.
- These come from a single mutation coming to dominate the entire population.
- They provide a measure of the beneficial mutation rate.



# Experimental Background

- Underground, radiation's role in DNA mutation and cell oxidation can be greatly reduced.
- How does this change the evolutionary behaviour?



# Our Experiment

- Run Long Term Evolution Experiment in multiple radiation environments
- 500 generations to begin
- Our radiation environments:

Active



Reduced Background



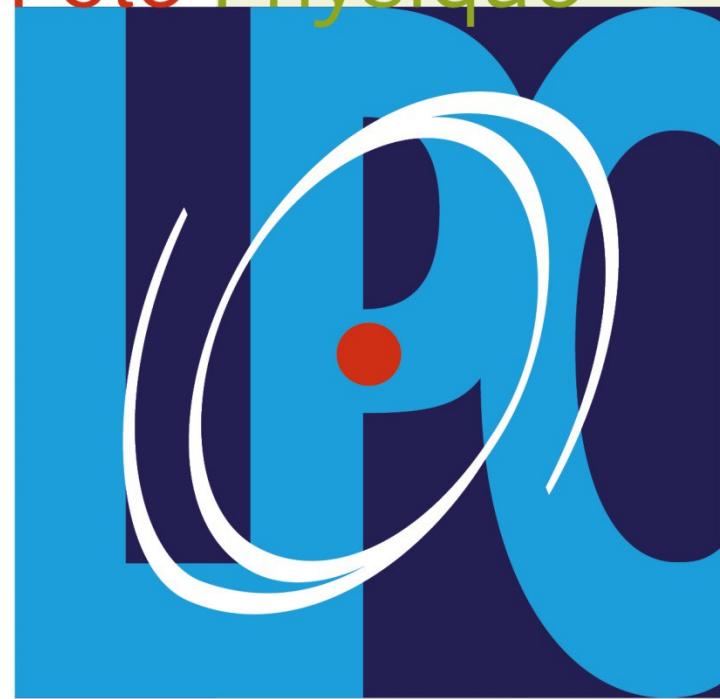
Ambient Background

Planned



Controlled Source

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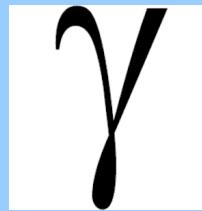


**Radiation background  
studies**

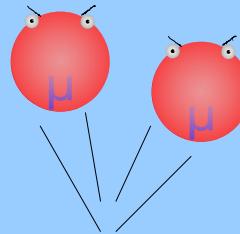
# Relevant Sources



(222)  
**Rn**  
86 RADON



K-40  
C-14  
Nutritional medium

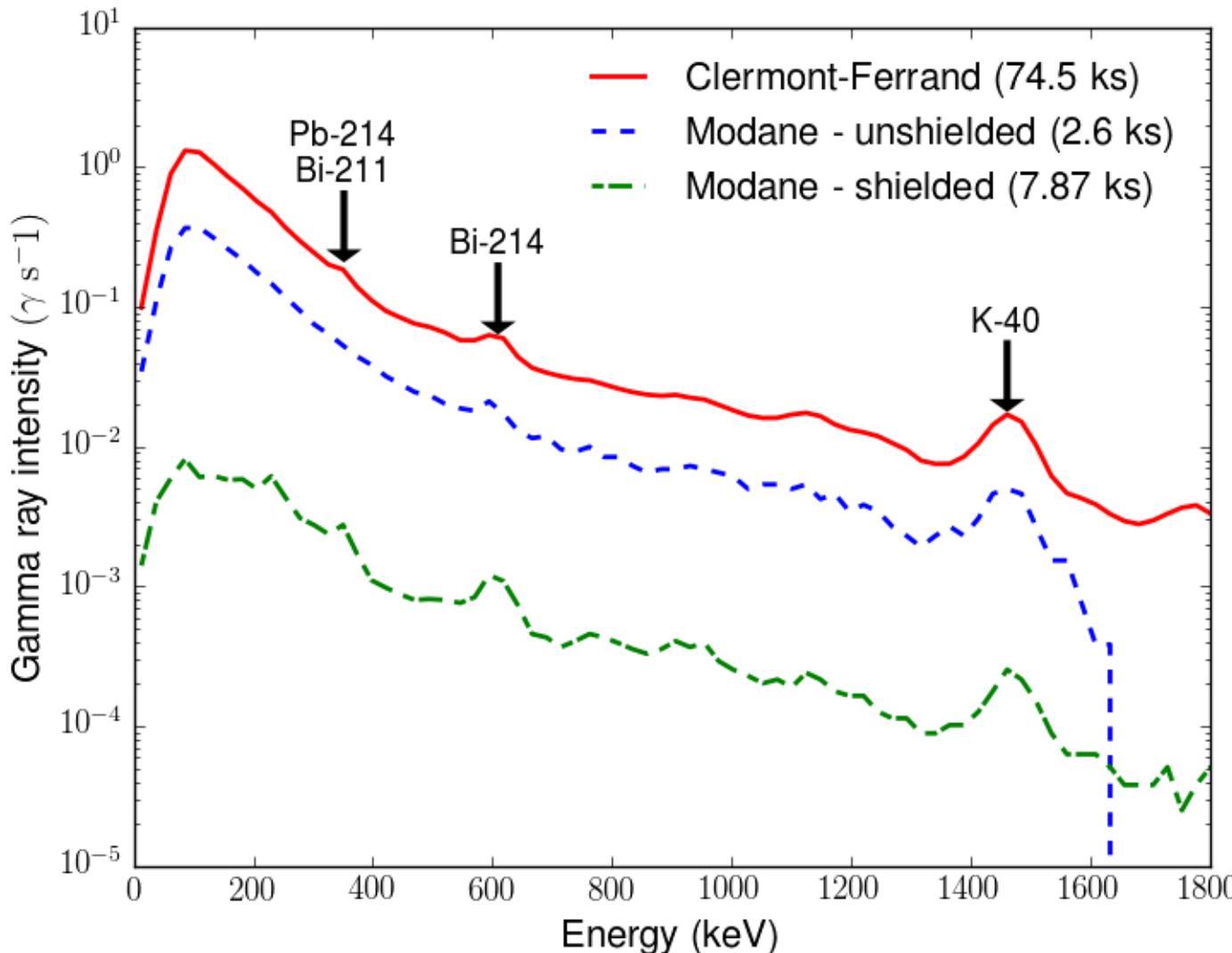


(222)  
~~Rn~~  
86 RADON



K-40  
C-14  
Nutritional medium

# Gamma Background



Nal measurements

# The macro: dosages

Source	Measurement Method	LPC Clermont (nGy/day)	Modane (nGy/day)	Modane (shielded) (nGy/day)
$\gamma$ background	Dosimeter measurement (rate varied by 10%)	2400	480	15
Muon flux	From theory	460	0	0
Potassium-40 ( $\gamma$ )	Simulations based on concentration	0.4	0.4	0.4
Potassium-40 ( $\beta$ )	Simulations based on concentration	74.4	74.4	74.4
Carbon-14 ( $\beta$ )	Simulations based on concentration	0.02	0.02	0.02
Total		2935	555	90

Using measured dosages, the background is reduced by a factor of 32

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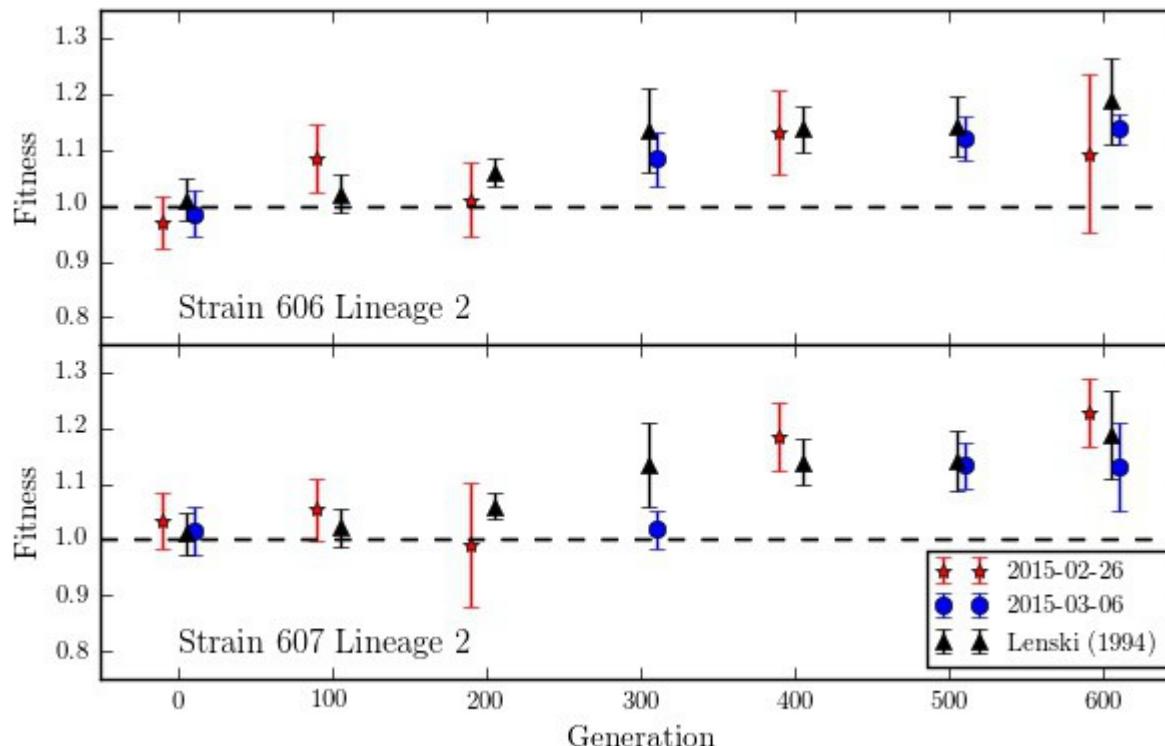
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Environnement

**Biological Results**

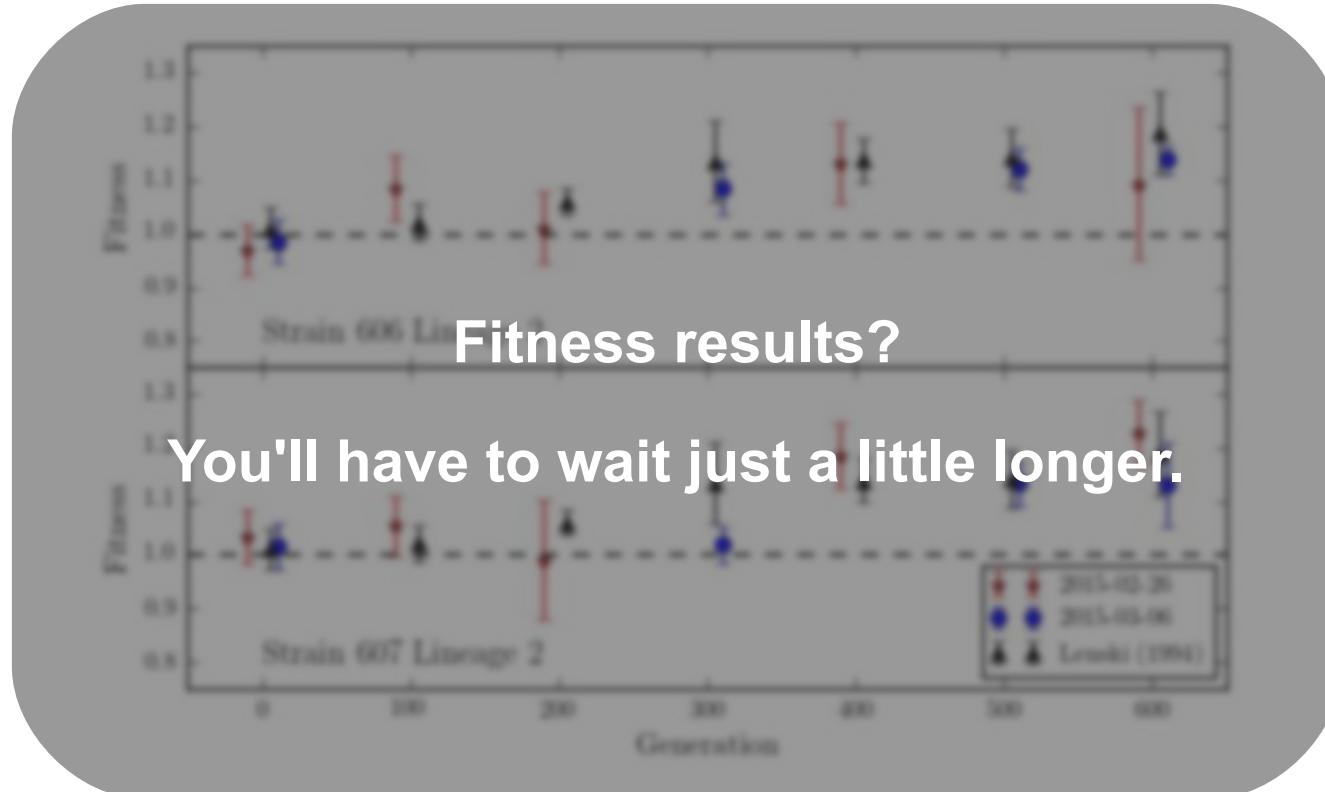
# Clermont-Ferrand

- Contamination-free growth to 500 generations
- Experimental protocols fully developed
- Observation of adaptation, as expected occurring near 300 generations

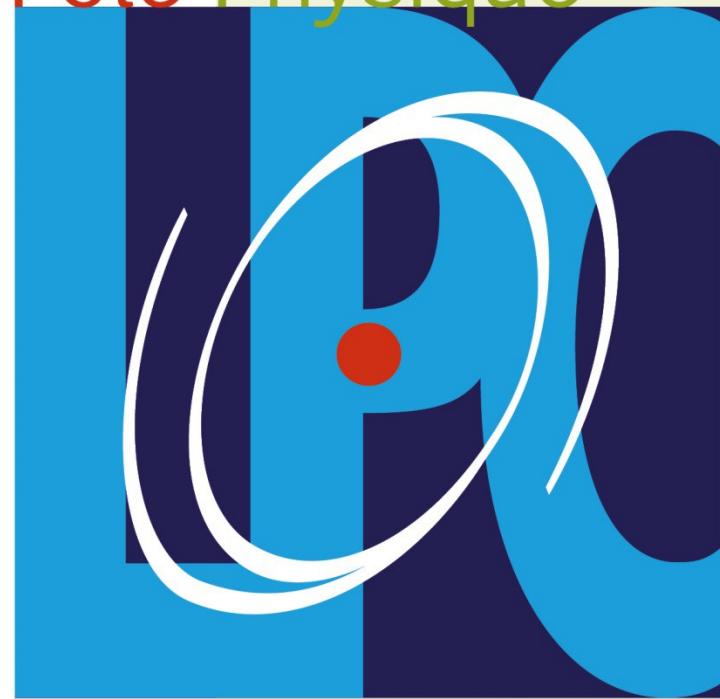


# Modane

- Contamination-free growth to 405 generations
- Development of protocols to measure fitness



Pôle Physique



**Simulating DNA damage  
in low background  
environments**

# Why Simulation?

- Allows us to test our theoretical understanding of the experiment, and of low background DNA damage
- Allows predictive estimates of how different radiation environments impact DNA
- Geant4 is a software platform that simulates the interactions of particles with matter
- Low energy electromagnetic simulations and biological effects can be simulated using models developed by the Geant4-DNA collaboration
- Geant4-DNA also permits the modelisation of chemistry

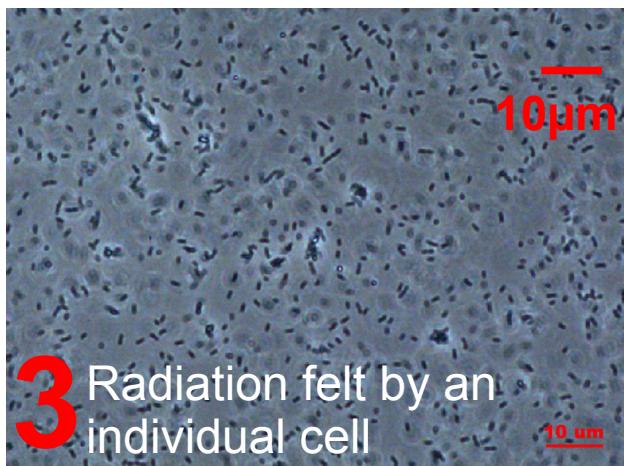
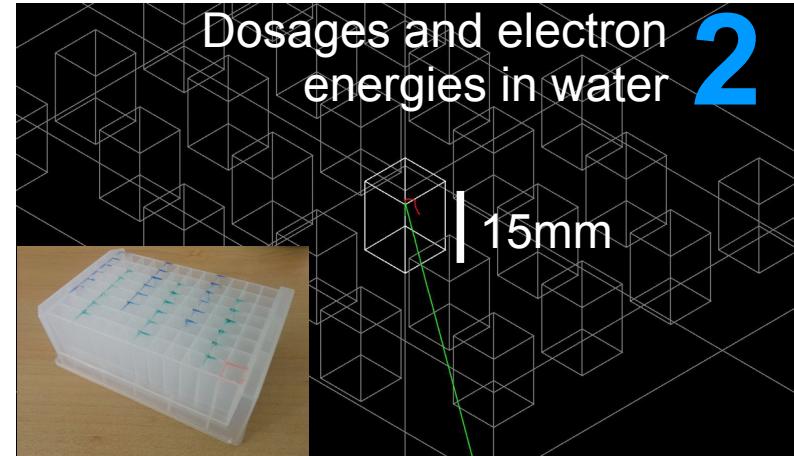
# From the metre to the nanometre

**1** Measurements of the environment

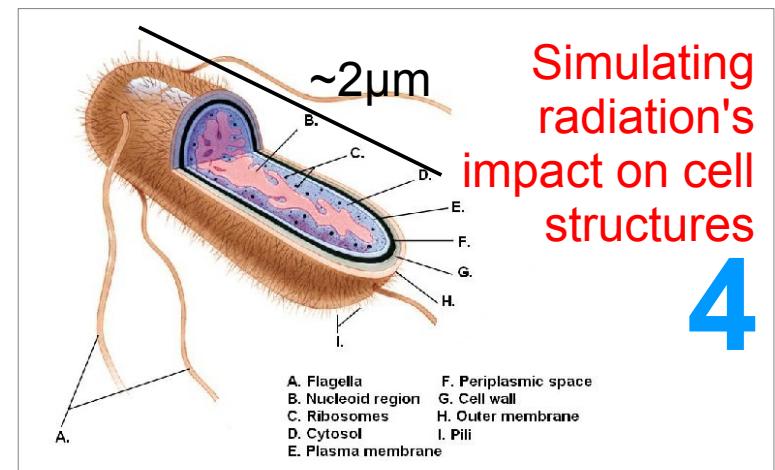


Dosages and electron energies in water

**2**

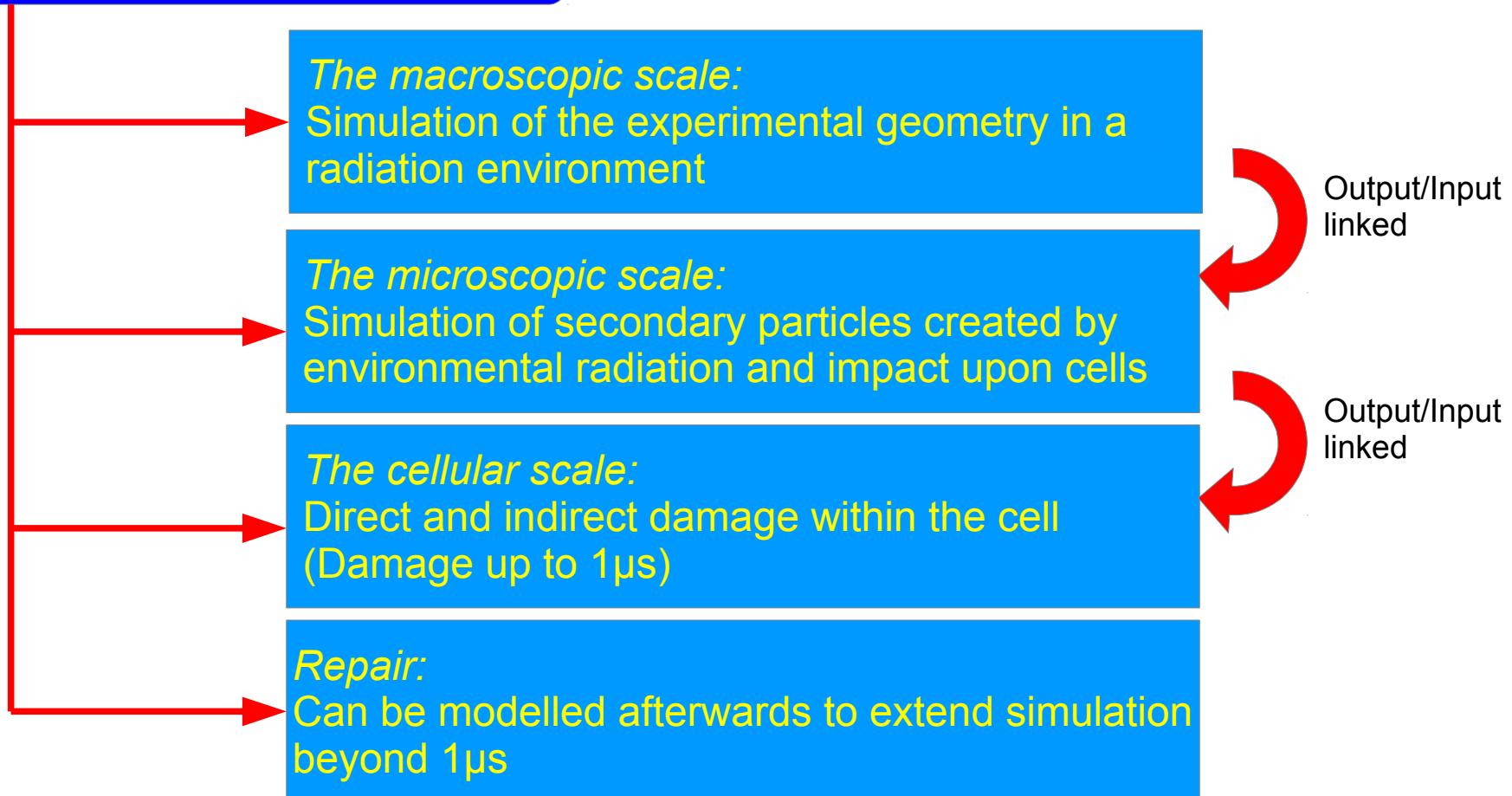


**3** Radiation felt by an individual cell

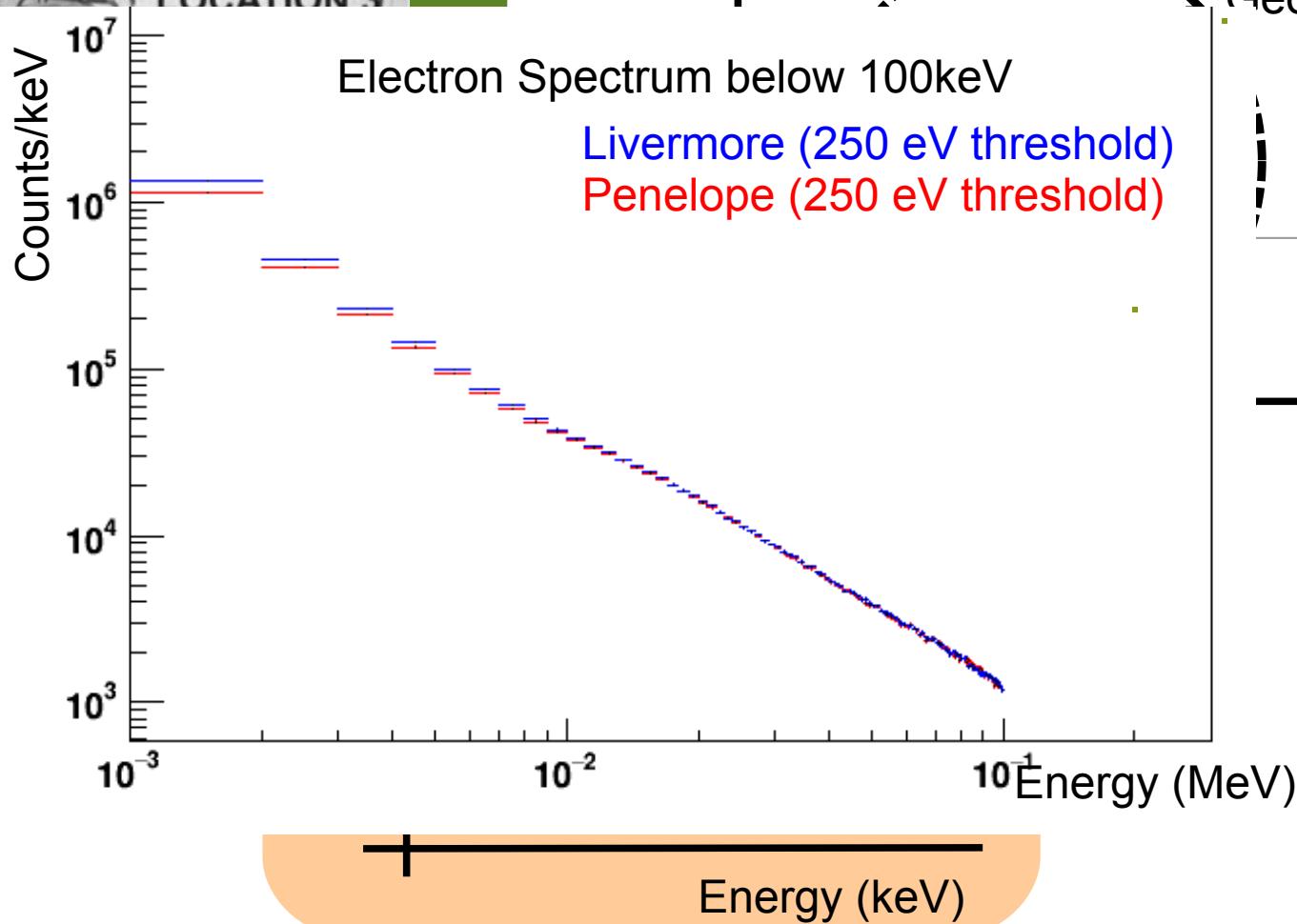


# A simulation workflow

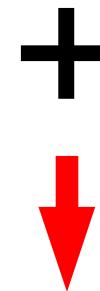
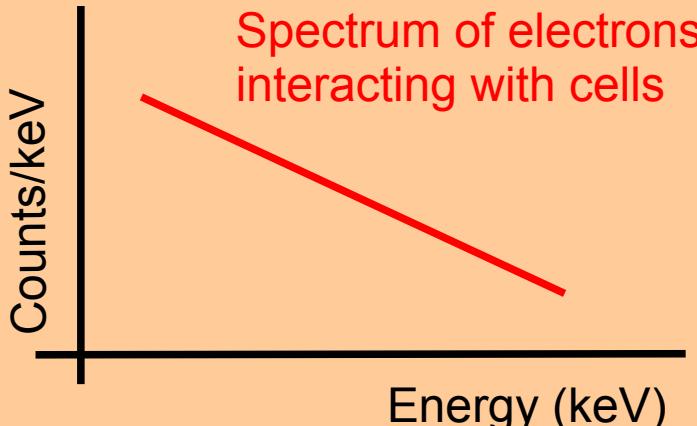
## *Simulating cellular damage from an environment*



# The macroscopic system

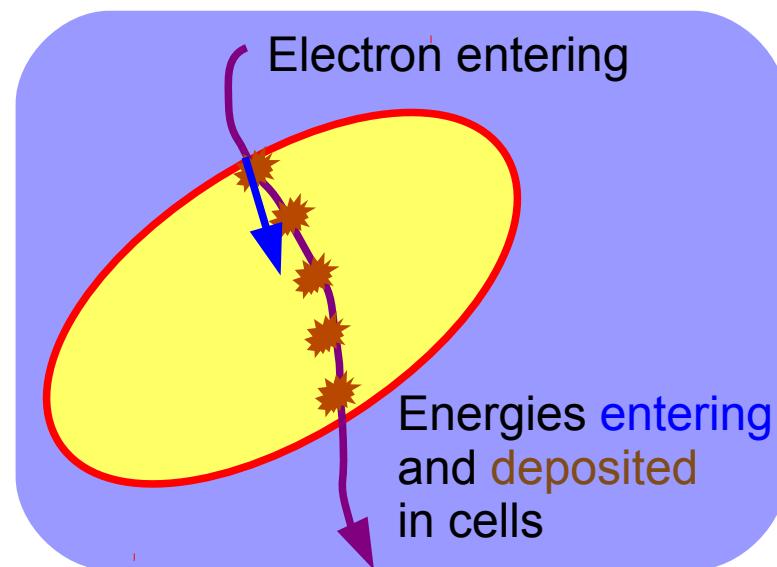
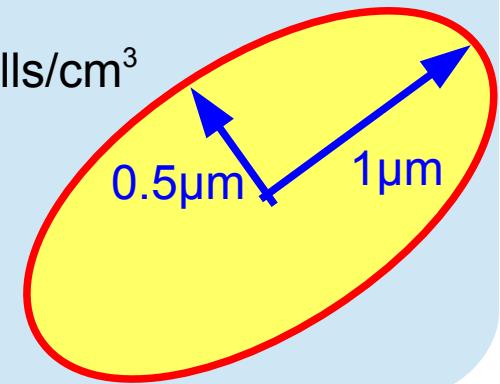


# The microscopic system

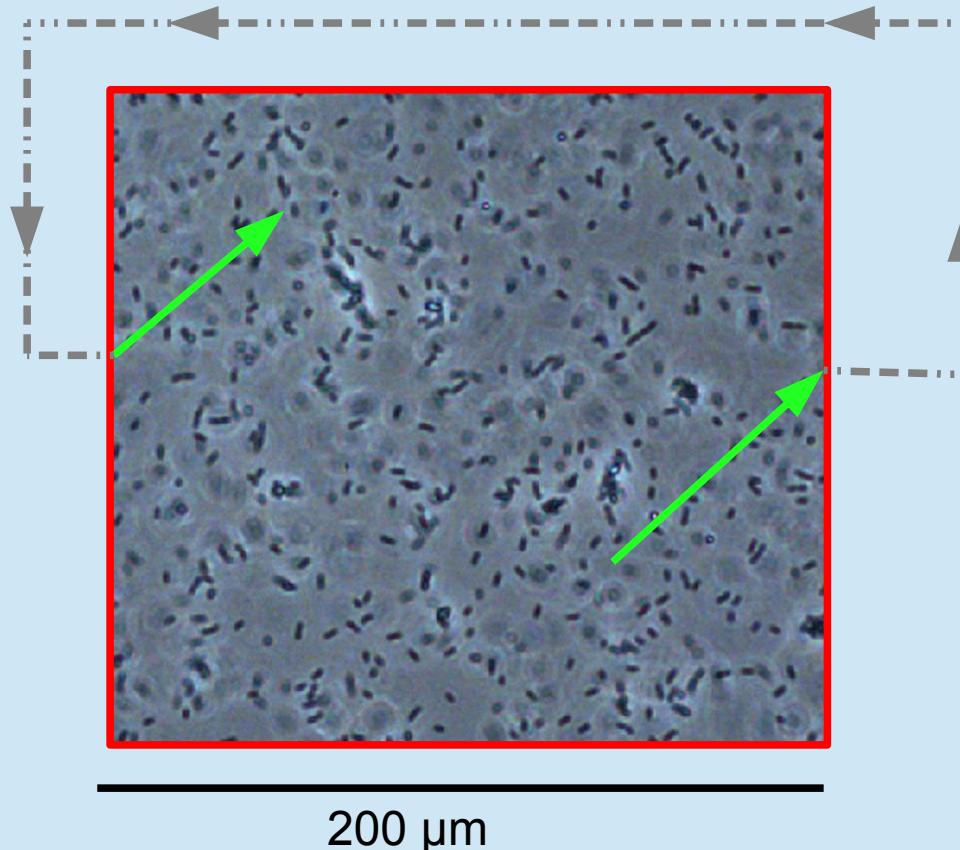


Broad cell details

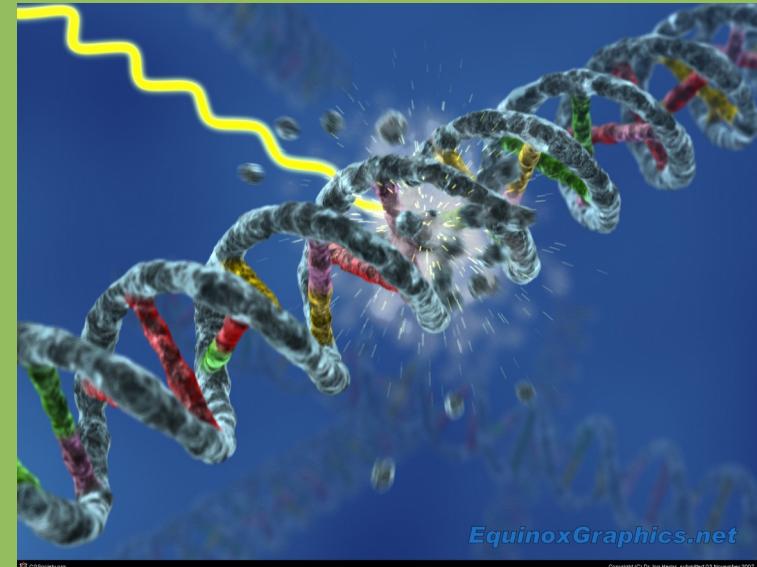
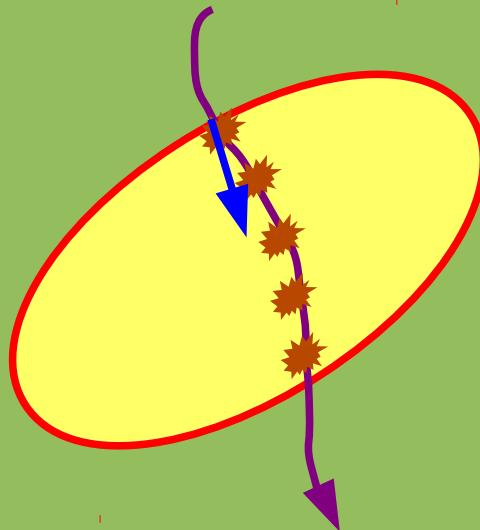
$10^7$  cells/cm<sup>3</sup>



# A periodic boundary to enable this



# Measuring DNA damage



EquinoxGraphics.net

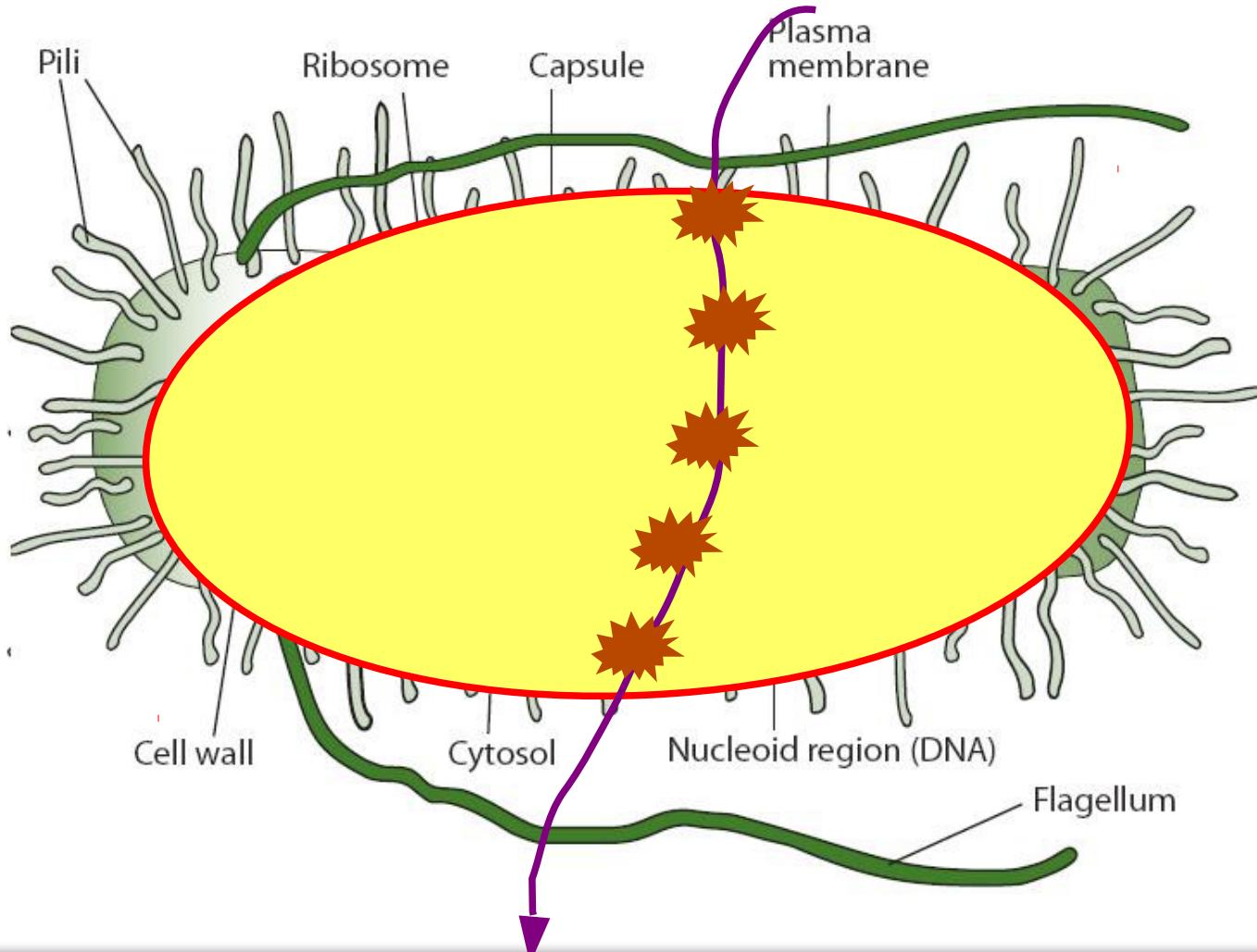
## Damage to consider

**Direct:**  
**Indirect:**

Radiation breaks one or both DNA strands  
Radical oxygen species damage the DNA Strand

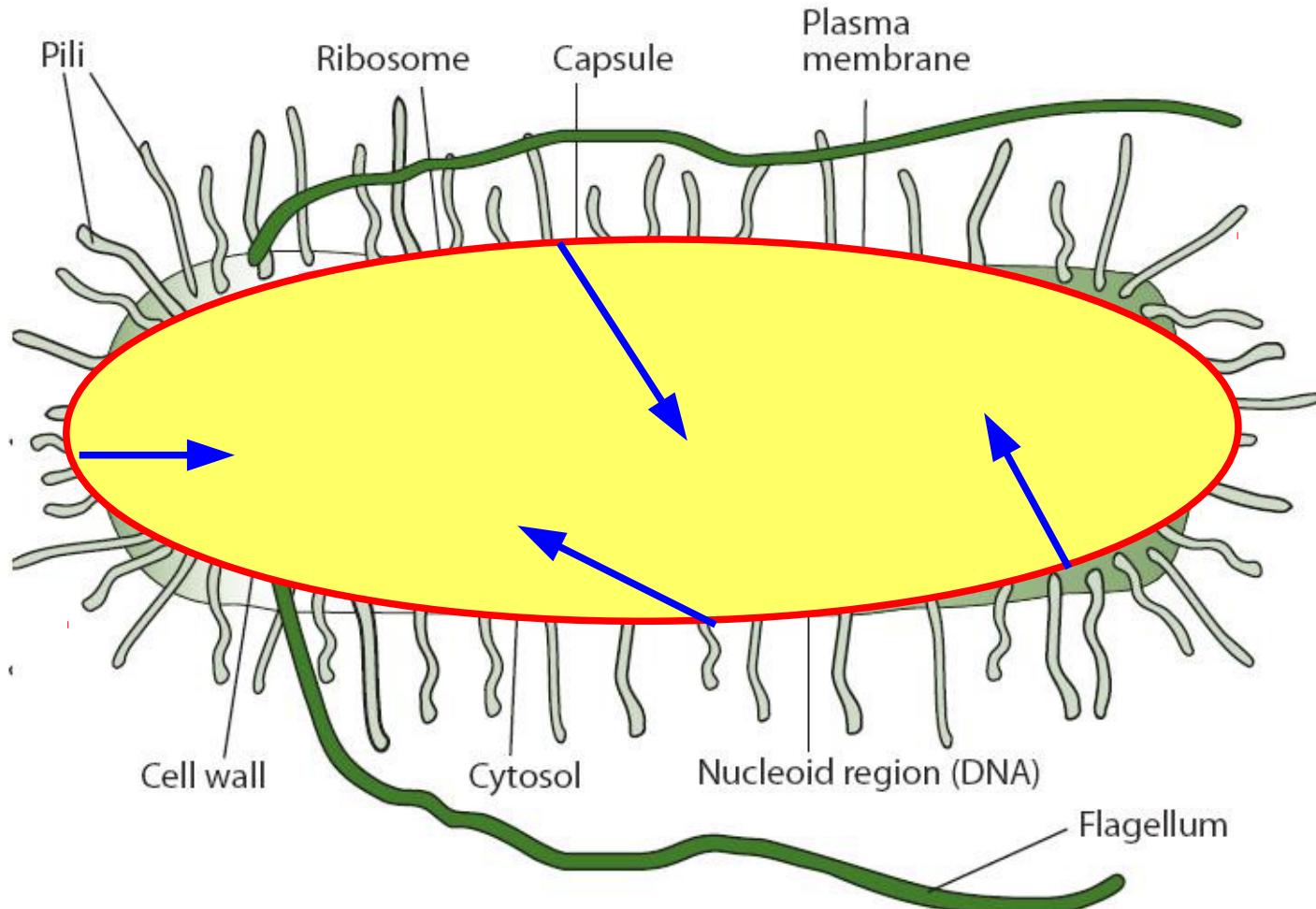
# Measuring DNA damage

**Method 1:** Correlate energy deposits with position



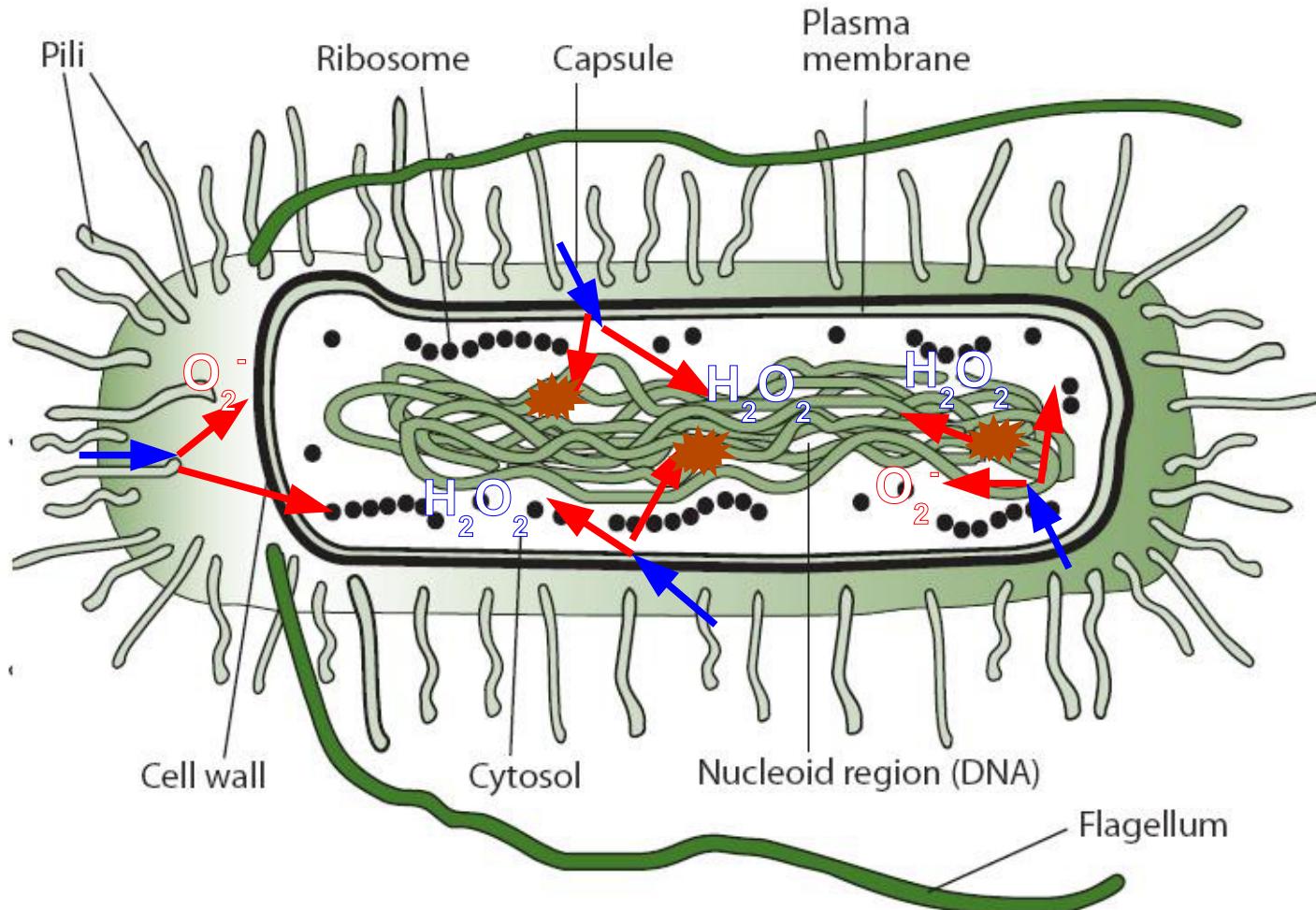
# Measuring DNA damage

**Method 2:** Build a bacteria model, and simulate chemistry



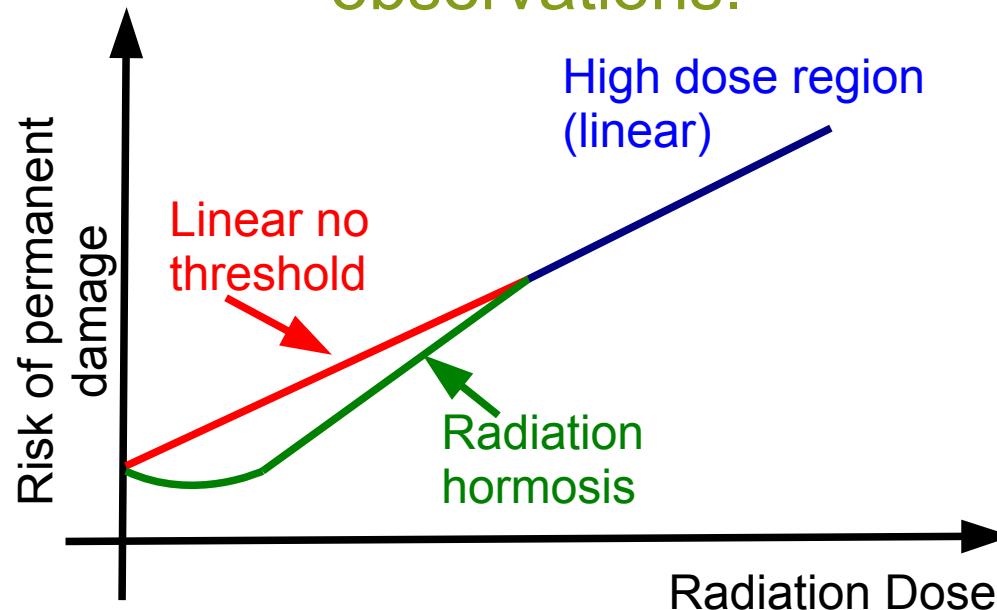
# Measuring DNA damage

**Method 2:** Build a bacteria model, and simulate chemistry



## Prospects: Simulation beyond the first $\mu\text{s}$

Simulating effects beyond the first  $\mu\text{s}$ , such as DNA repair allow us to probe the does-response curve at the low end, and build the theoretical support for LBE observations.

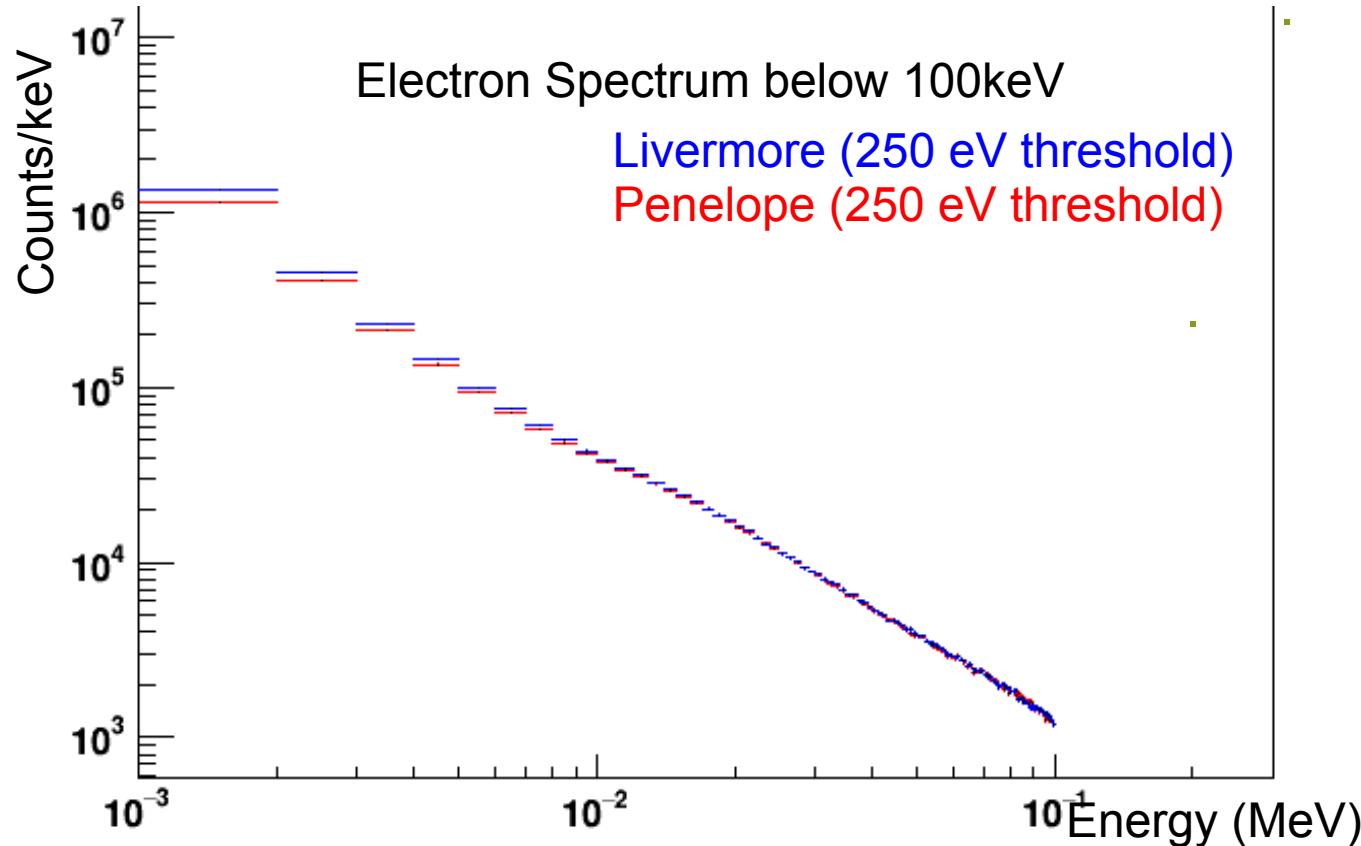


# Conclusions

- We have set up a biological laboratory at Modane
- There, we are conducting Long Term Evolution Experiments to see how radiation affects evolutionary behaviour
- We are building a robust simulation workflow that will permit the simulation of DNA damage in low background environments

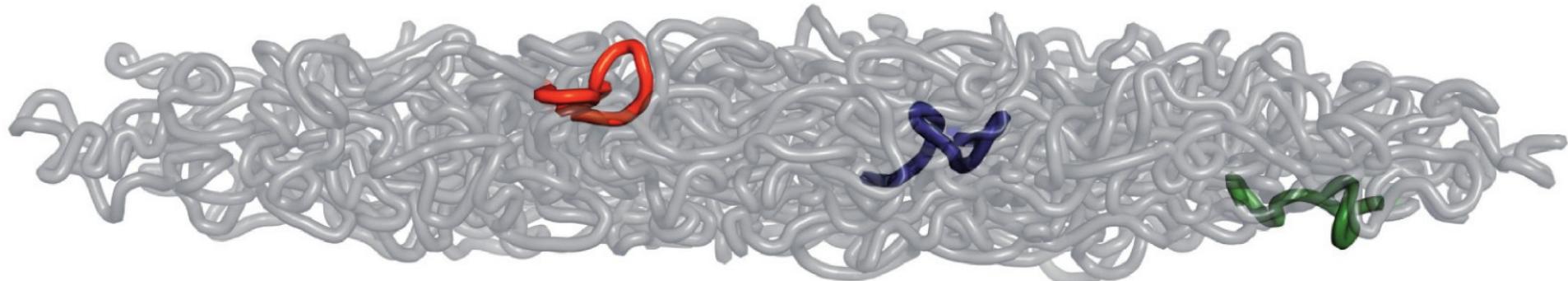
*That's all Folks!*

# The macroscopic system - output



# Bacterial DNA Organisation

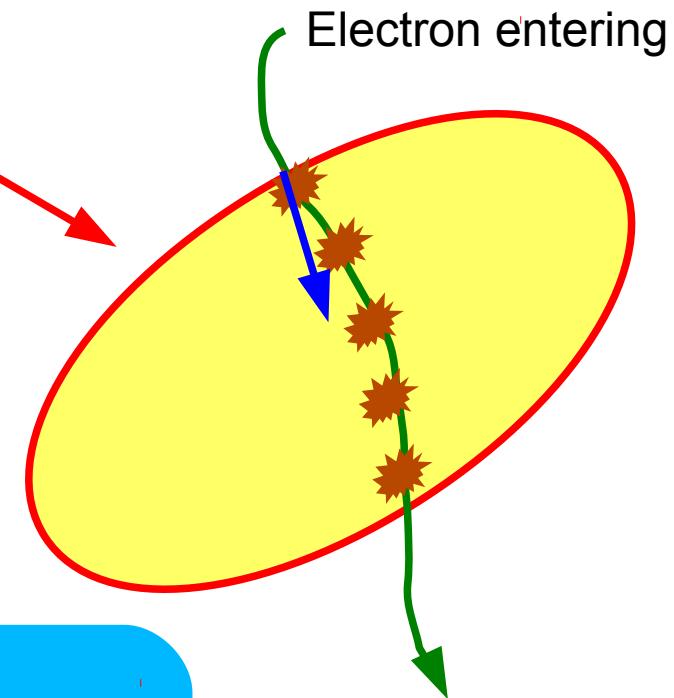
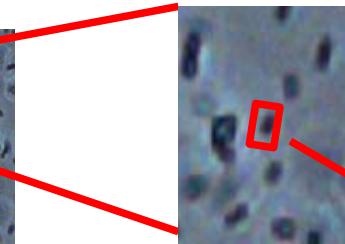
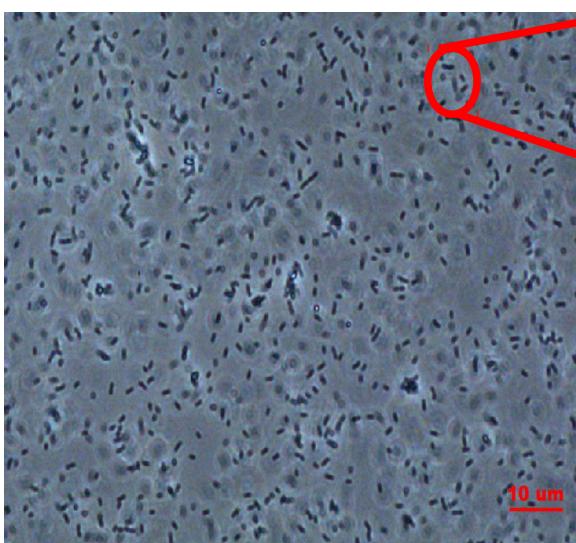
- Modelling the structure of DNA inside a bacterial nucleoid or an animal nucleus is a complex task
- Accuracy of the representation needs to be balanced with ease of modelling and desired outputs



The organisation of the bacterial nucleoid from polymer models (Fritzsche 2012), Genes that share the same transcription factor are clustered together.

# What does this tell us?

200  $\mu\text{m}$



For each cell, we learn:

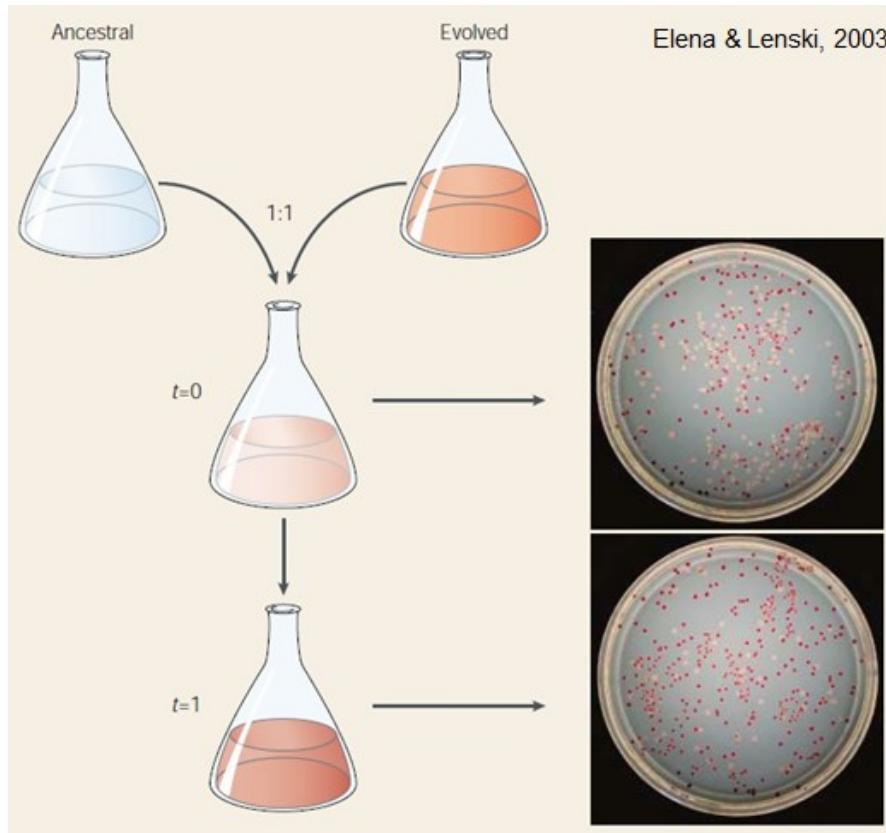


Position of energy depositions, and energy deposited



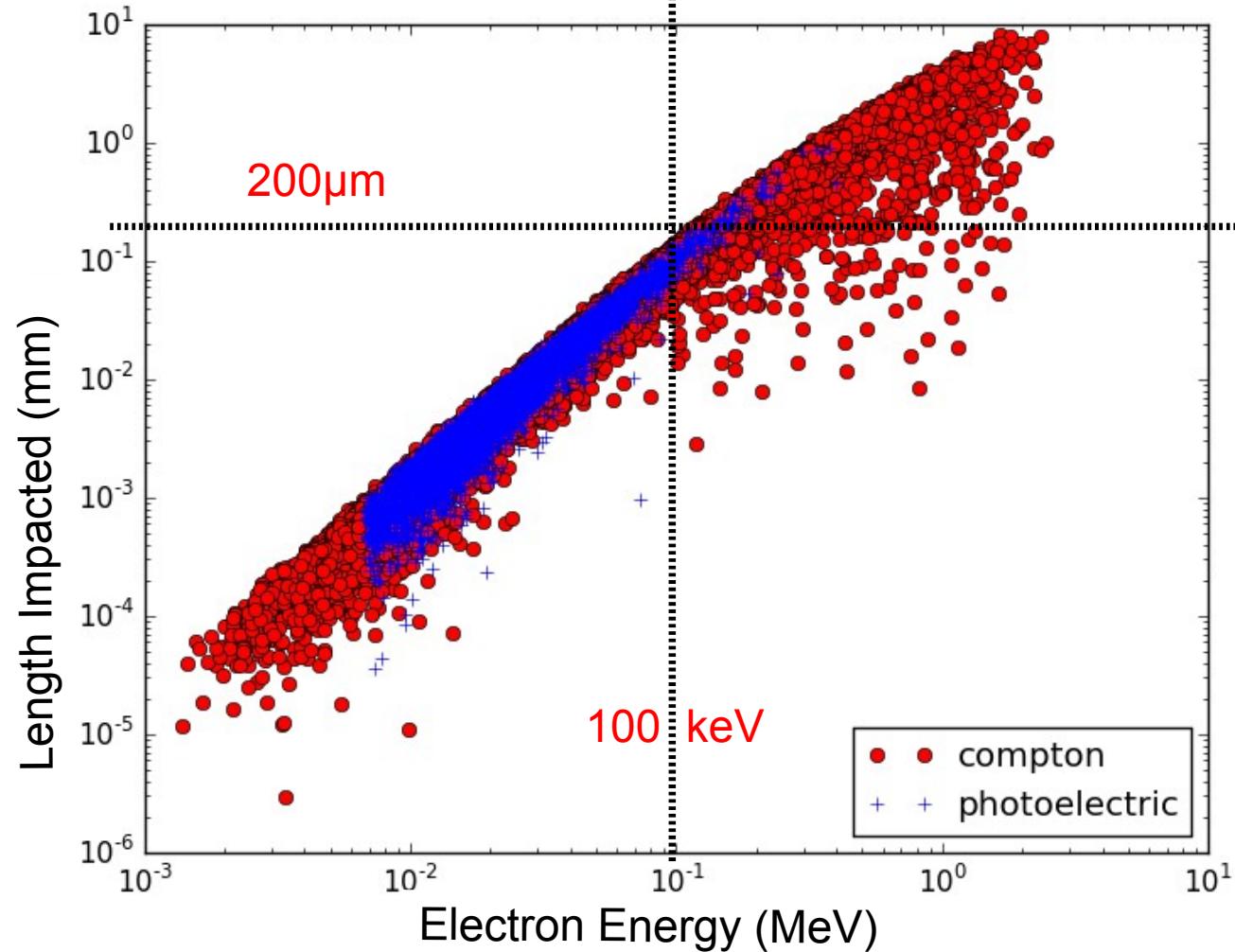
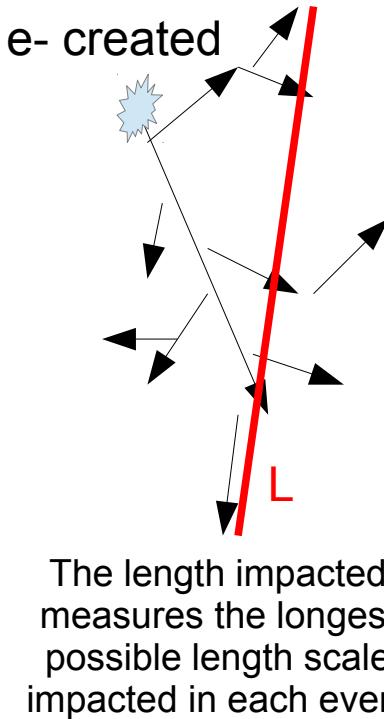
Energy, direction and position of entering electrons

# Experimental Background

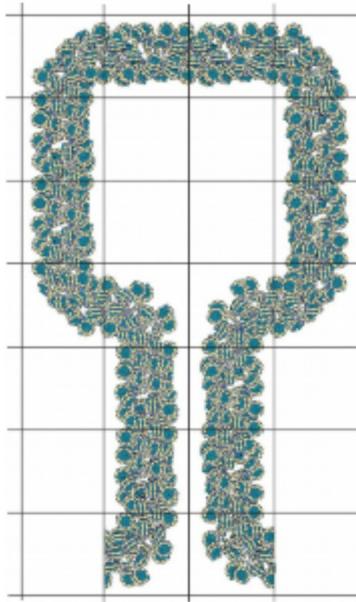


- Lenski, and us, measure bacterial fitness to measure the adaptation of the strain to its environment.
- An evolved strain is placed in competition with its ancestor, and the difference in their respective growths indicates the relative fitness

## Results: Electron energies in the well

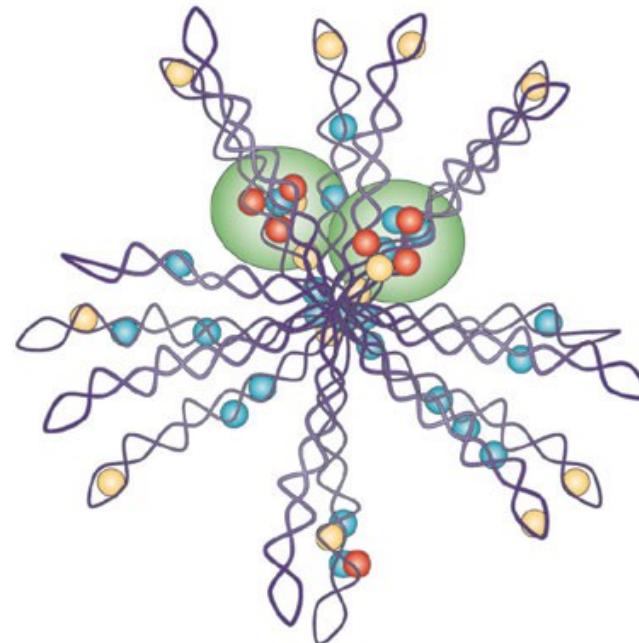


# The individual cell – Bacterial DNA



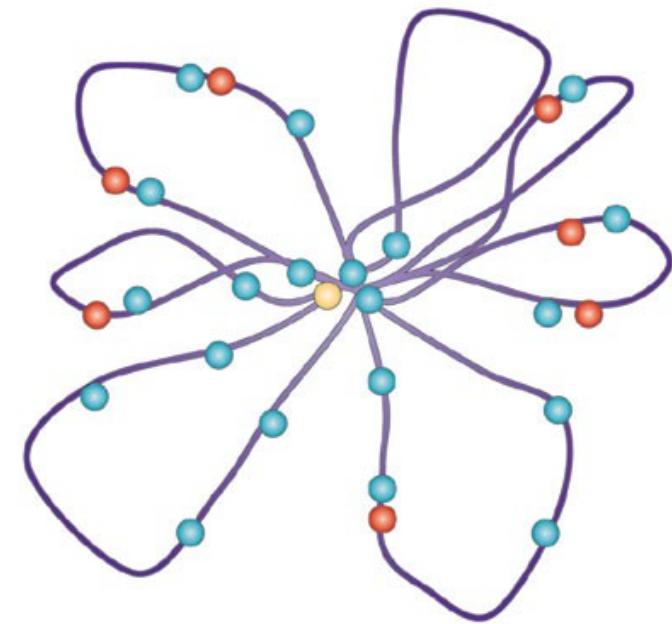
Eukaryotic DNA  
 model (Friedland et  
 al., 2011)

Bacterial DNA being duplicated  
 a Exponential phase of growth



● RNA polymerase at RNA promoters	● H-NS	● Transcription factories	● Fis
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DNA when bacteria is not replicating  
 b Stationary phase of growth

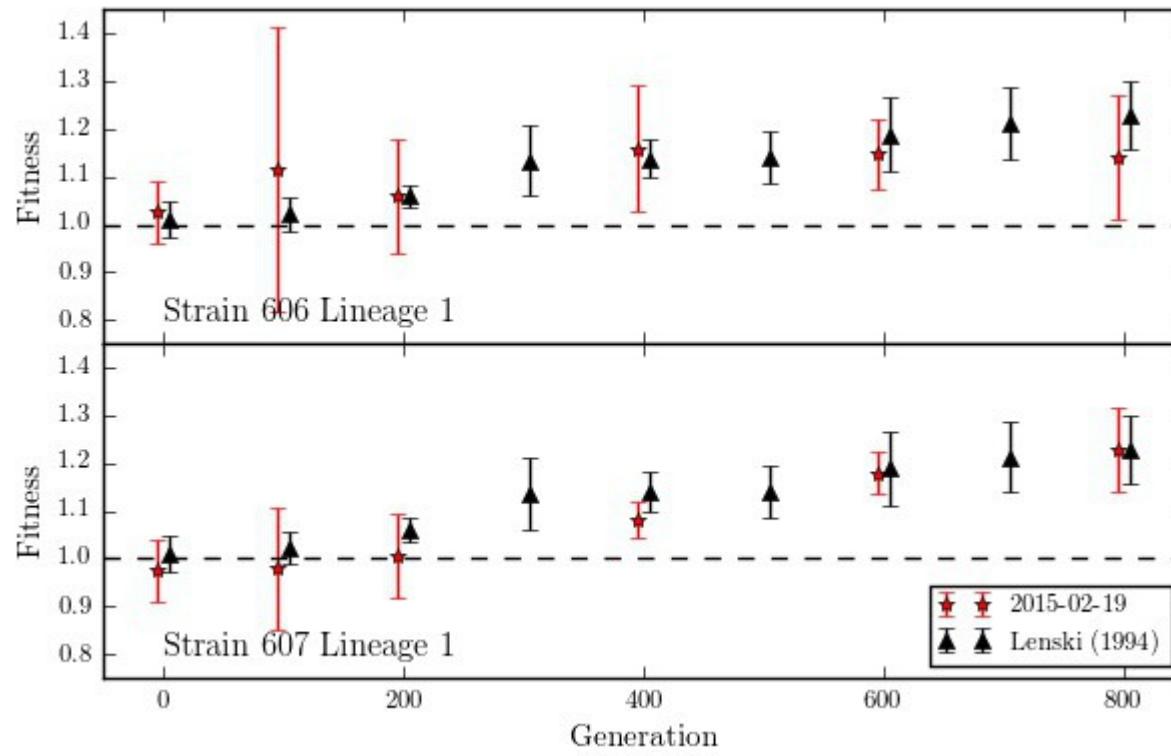


Dillon & Dorman (2010)

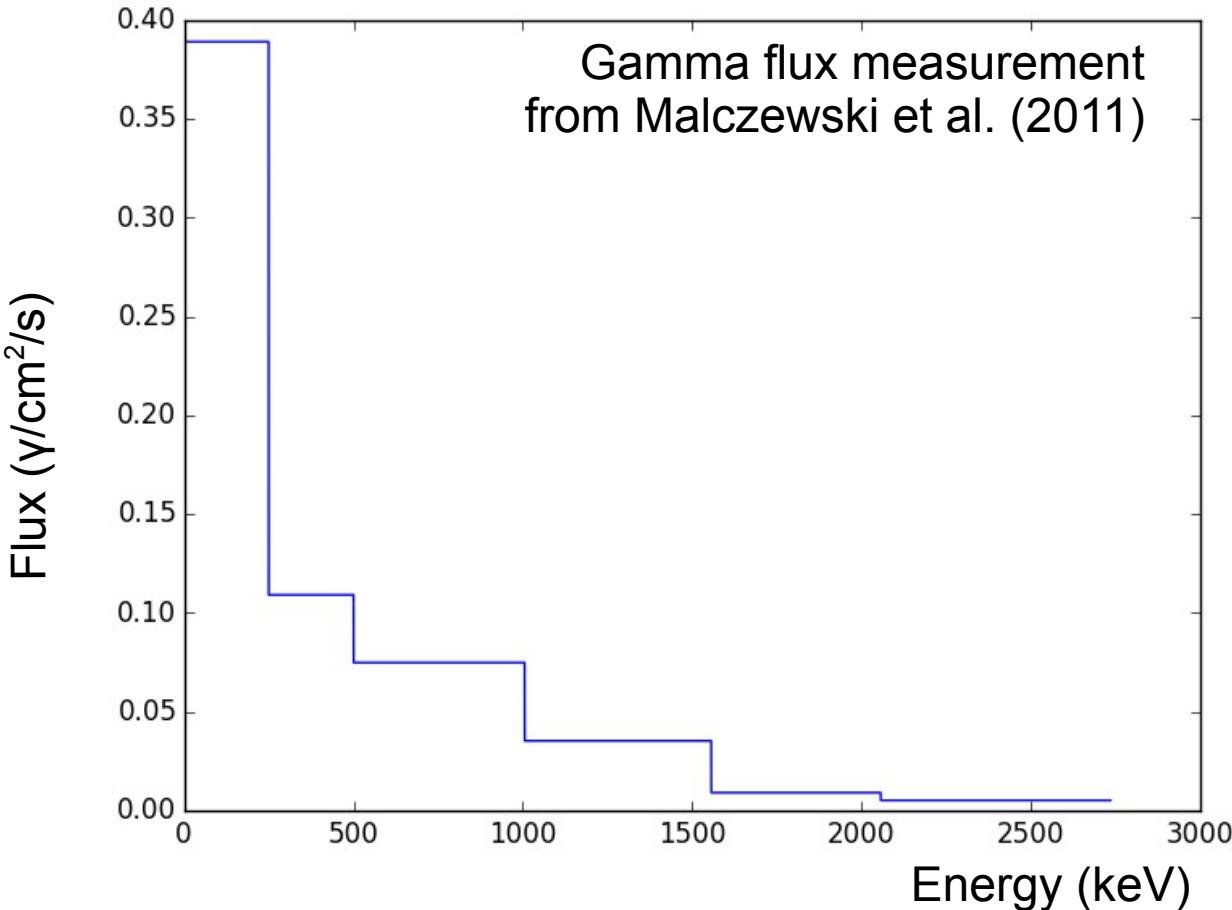
Bacterial DNA has a significantly different secondary and tertiary structure to eukaryotic DNA

Nature Reviews | Microbiology

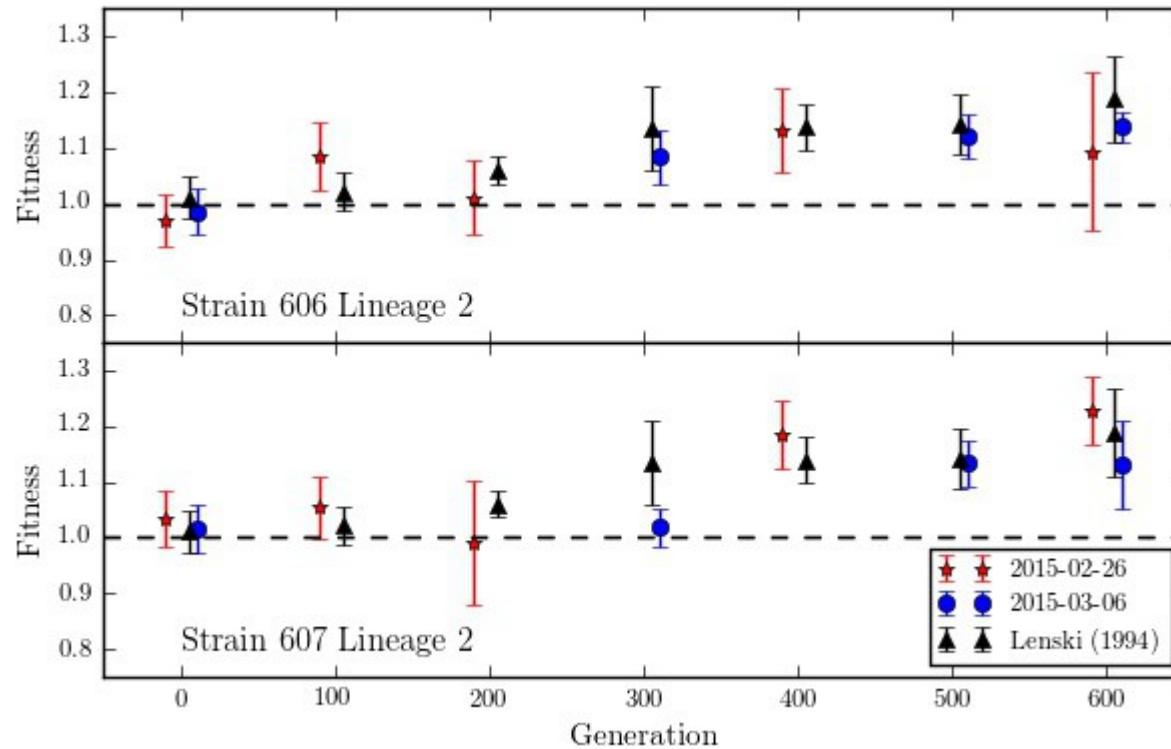
# Fitness measurements at CF



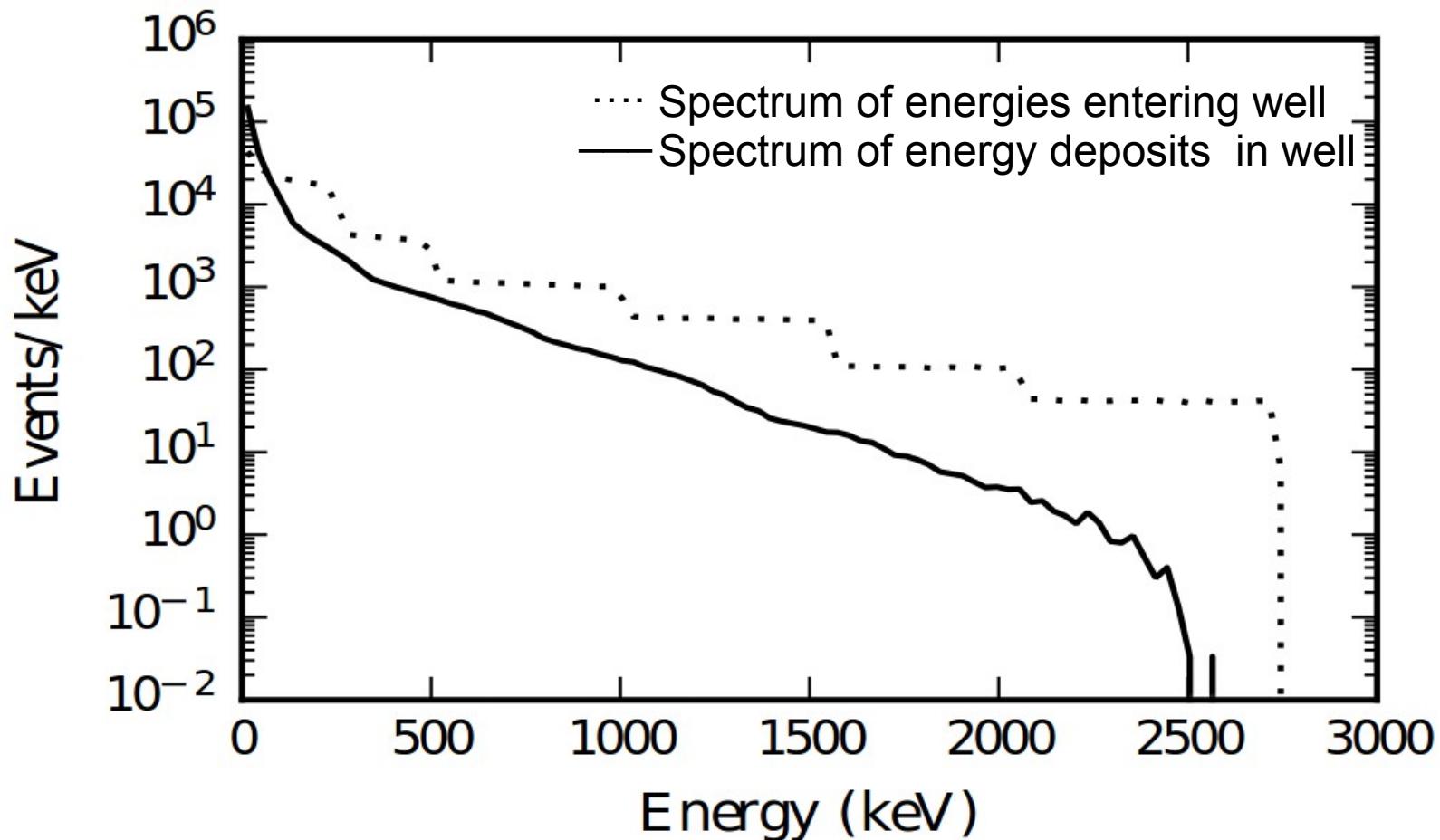
# Measurements of radiation in the environment



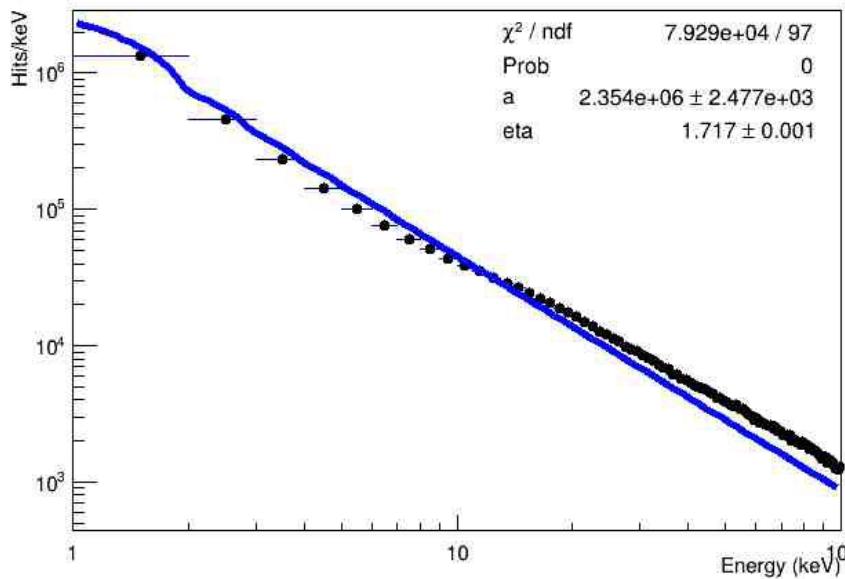
# Fitness measurements at CF



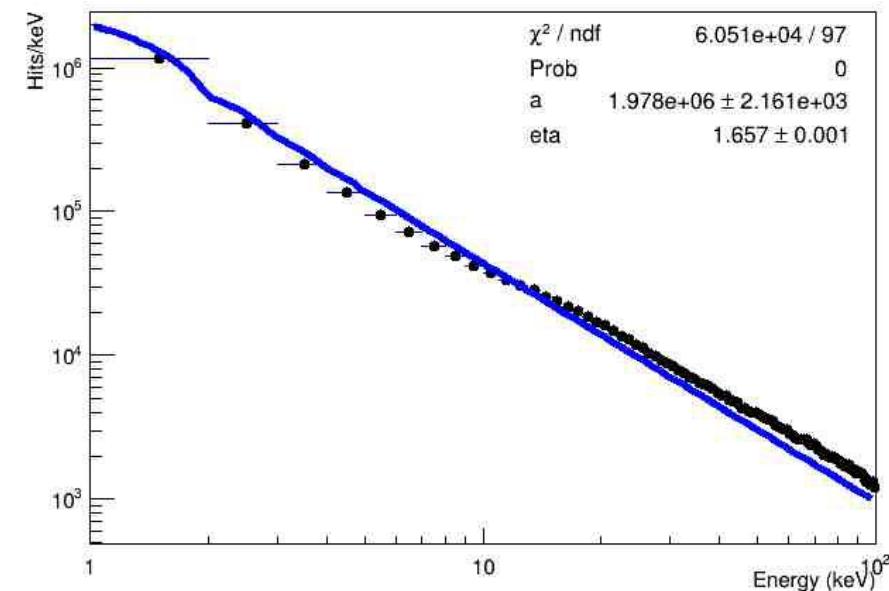
# Energy deposit spectrum in well



# A power law below 100keV



Livermore



Penelope

# From the environment to the cell

- We have now knowledge of the secondary electrons in the cell.
- Typical bacteria:

  - $\sim 10^8$  bacteria  $\text{cm}^{-3}$

- For us, 4000 bacteria in a cube with side length  $200\mu\text{m}^3$
- Putting  $10^8$  bacteria into memory is a bad idea, better to use a repeating boundary condition

