

Chemistry examples



geant4-dna.org

Chem3 example

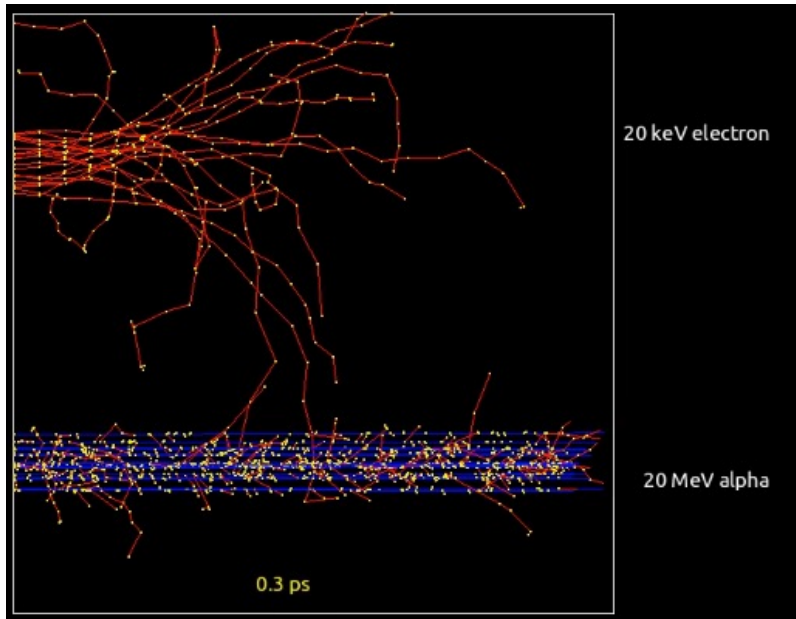
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Pavia, Italy
18/01/2024

Geant4.11.2.0
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Processes in the Physical Stage

Geant4-DNA simulation



		Branching ratio (%)			
Ionization	H ₂ O ⁺	H ₃ O ⁺ + ·OH	100	100	Ionization
	A ¹ B ₁	H ⁺ + ·OH	65	65	A1B1_DissociationDecay
H ₂ O + ΔE		35	35	No displacement	
Excitation	B ¹ A ₁	H ₃ O ⁺ + ·OH + e _{aq} ⁻	50	55	Auto-Ionization
		H ⁺ + ·OH	25.35	-	A1B1_DissociationDecay
		H ₂ + 2·OH	3.25	15	B1A1_DissociationDecay
		2H ⁺ + O(³ P)*	3.9	-	B1A1_DissociationDecay2
		H ₂ O + ΔE	17.5	30	No displacement
	Rydberg A+B, C+D, Diffuse bands	H ₃ O ⁺ + ·OH + e _{aq} ⁻	50	50	Auto-Ionization
		H ₂ O + ΔE	50	50	No displacement
Electron capture	DEA	OH ⁻ + ·OH + H ₂	100	100	Dissociative attachment
	Recombinat-ion	H ⁺ + ·OH	35.75	55	A1B1_DissociationDecay
		H ₂ + 2·OH	13.65	15	B1A1_DissociationDecay
		2H ⁺ + O(³ P)	15.6	-	B1A1_DissociationDecay2
		H ₂ O + ΔE	35	30	No displacement

W.-G. Shin, et al., Phys. Med. (2021)

Processes in the Chemical Stage

Water Molecules:

- Electron Hole Recombination
- Water Dissociation

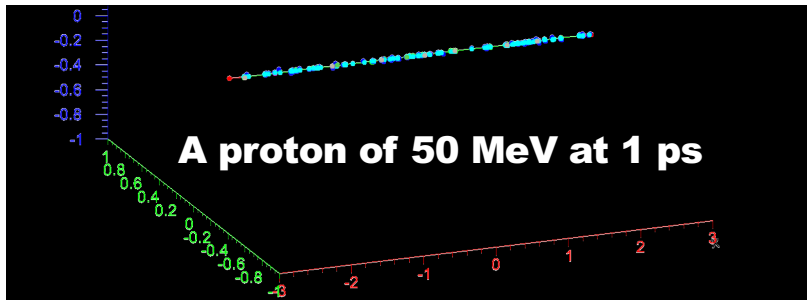
Chemical Species:

DIFFUSION

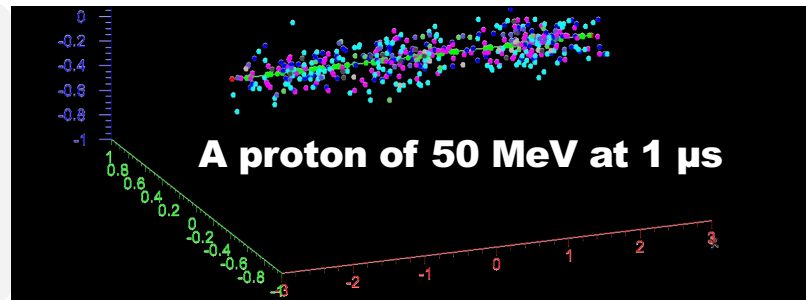
Species	Diffusion coefficient D ($10^{-9} \text{ m}^2 \text{ s}^{-1}$)
H_3O^+	9.0
$\text{H}\cdot$	7.0
OH^-	5.0
e^-_{aq}	4.9
H_2	5.0
$\cdot\text{OH}$	2.8
H_2O_2	1.4

REACTIONS

Reactants	Products	Rate [$1\text{e-3} * \text{m}^3 / (\text{mole} * \text{s})$]
$\text{e}^-_{\text{aq}} + \text{e}^-_{\text{aq}} + 2\text{H}_2\text{O}$	$\text{H}_2 + 2\text{OH}^-$	0.5e10
$\text{e}^-_{\text{aq}} + \cdot\text{OH}$	OH^-	2.95e10
$\text{e}^-_{\text{aq}} + \text{H}\cdot + \text{H}_2\text{O}$	$\text{H}_2 + \text{OH}^-$	2.65e10
$\text{e}^-_{\text{aq}} + \text{H}_3\text{O}^+$	$\text{H}\cdot + \text{H}_2\text{O}$	2.11e10
$\text{e}^-_{\text{aq}} + \text{H}_2\text{O}_2$	$\text{OH}^- + \cdot\text{OH}$	1.41e10
$\cdot\text{OH} + \cdot\text{OH}$	H_2O_2	0.44e10
$\cdot\text{OH} + \text{H}$	H_2O	1.44e10
$\text{H}\cdot + \text{H}\cdot$	H_2	1.20e10
$\text{H}_3\text{O}^+ + \text{OH}^-$	$2\text{H}_2\text{O}$	1.43e11



- proton
- e-
- e^-_{aq}
- OH^0
- H_3O^1
- H^0
- OH^1
- H_2^0
- H_2O_2^0



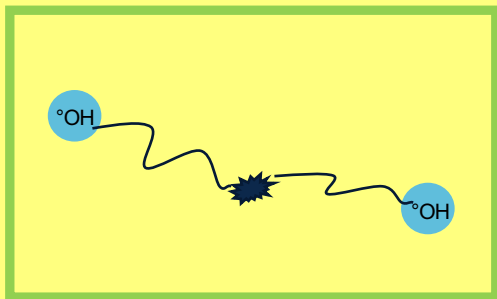
Chemistry Models

SBS model

At each step (bigger than a Minimum Time Step) the separation distance "d" of all pairs of reactants is checked.

Two species react with each other when d is below a given threshold "R", called reaction radius.

The Brownian bridge technique compensate for possible missed reactions.



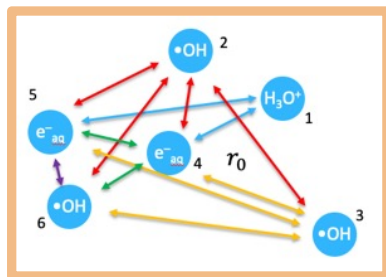
Karamitros, M. et al. (2014) Diffusion-controlled reactions modeling in Geant4-DNA. Journal of Computational Physics, 274, 841-882. <https://doi.org/10.1016/j.jcp.2014.06.011>

IRT model

An event table is constructed with the initial chemical species positions and reaction times (calculated with probability functions) for each reactant pair of interest.

Table's entries are sorted in ascending reaction time order and then processed.

Reaction product positions are randomly sampled within a sphere centred at the reaction site.



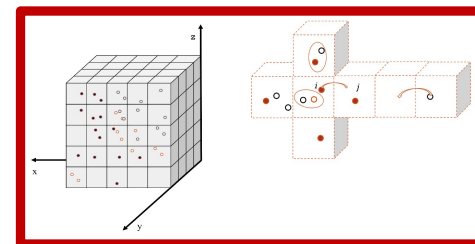
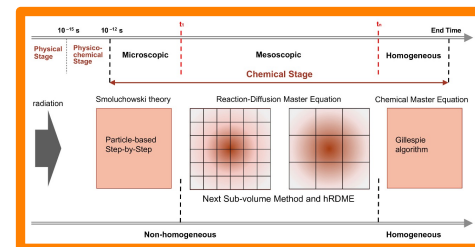
Karamitros, M. et al. (2020) Implementing the Independent Reaction Time method in Geant4 for radiation chemistry simulations. <https://doi.org/10.48550/arXiv.2006.14225>

Ramos-Méndez, J. et al. (2020) Independent reaction times method in Geant4-DNA: Implementation and performance. <https://doi.org/10.1002/mp.14490>

SBS-RDME model

Combination of:

- SBS model
- Compartment-based model using RDME (Reaction-Diffusion Master Equation)



Tran, H. N. et al. (2021) Geant4-DNA Modeling of Water Radiolysis beyond the Microsecond: An On-Lattice Stochastic Approach. International Journal of Molecular Sciences, 22(11), 6023. <https://doi.org/10.3390/ijms22116023>

The Chemistry Examples

example/extended/medical/dna

- The « chem1 » example illustrates how to activate the simulation of water radiolysis (step-by-step method).
- The « chem2 » example illustrates how to set minimum time step limits on water radiolysis (step-by-step method).
- The « chem3 » example illustrates how to implement user actions in the chemistry module (step-by-step method).
- The « chem4 » example illustrates how to compute radiochemical yields ("G") versus time, including a dedicated ROOT graphical interface (step-by-step method).
- The « chem5 » example illustrates how to compute radiochemical yields ("G") versus time, using alternative physics and chemistry lists (step-by-step method).
- The « chem6 » example illustrates how to compute radiochemical yields ("G") versus time and LET using IRT method.
- The « scavenger » example illustrates how to simulate scavenging using an easy-to-use interface and the IRT method.
- The « UHDR » extended/medical/dna example illustrates how to activate the chemistry mesoscopic model in combination with the step-by-step model, and allows to simulate chemical reactions beyond 1 us post-irradiation.

Chem3 example

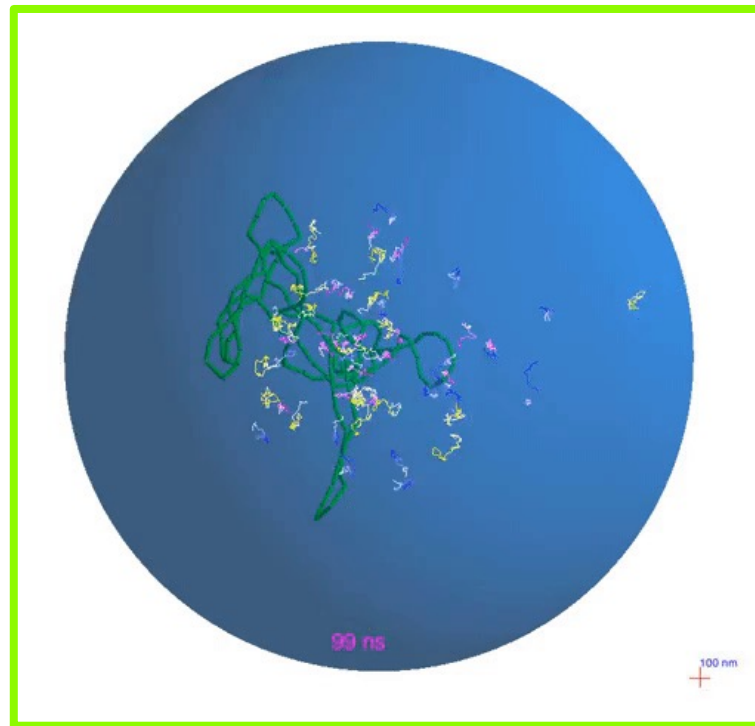
AUTHOR: Mathieu Karamitros

MAINTAINER: Hoang N. Tran

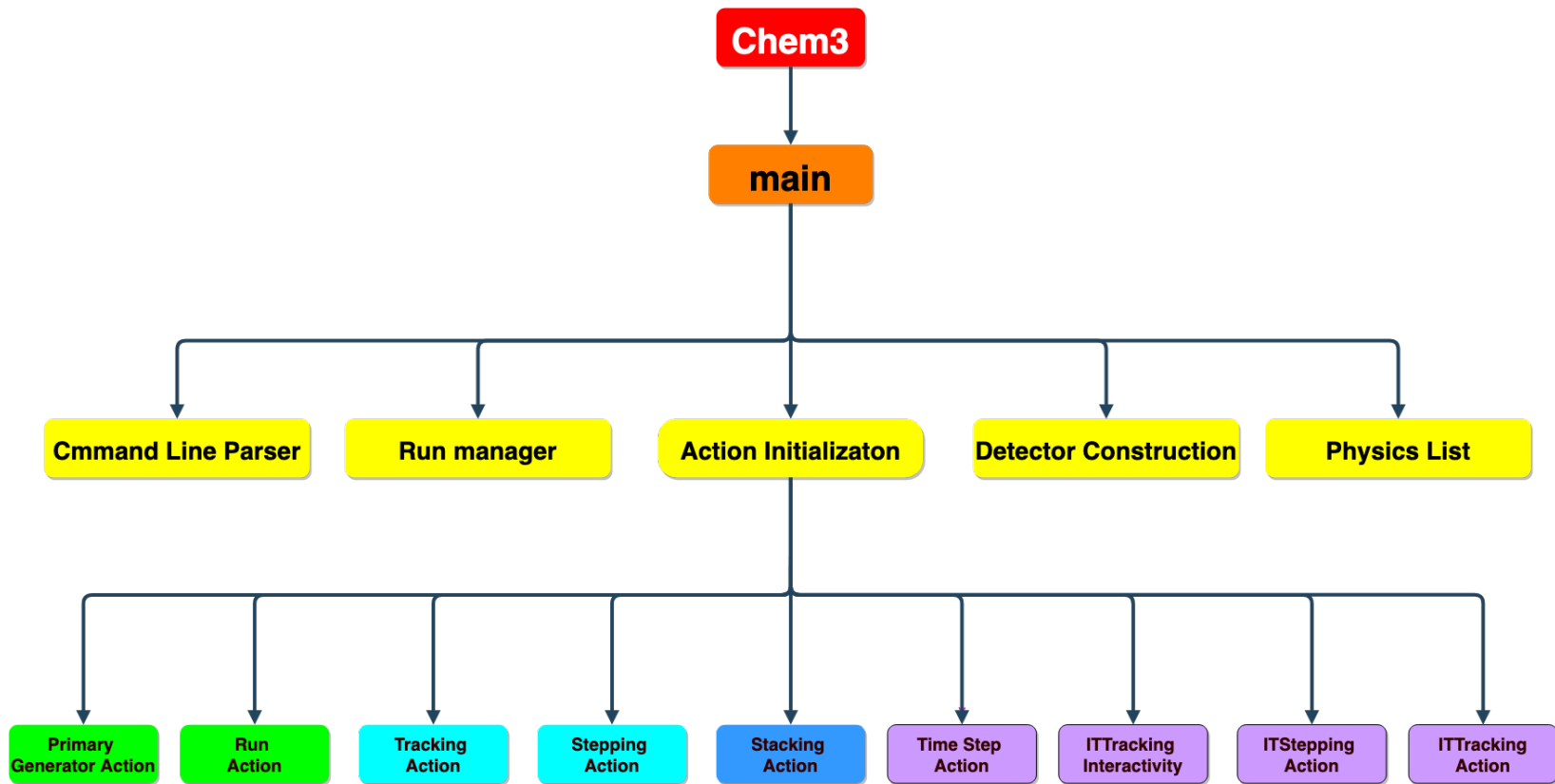
Chem3 Example

Illustrates how to implement **user actions** in the chemistry module using the **step-by-step** model.

Users can also **visualize** the **trajectories** of the chemical species in time and space using the **graphical user interface**.



Chem3 Application Code Structure



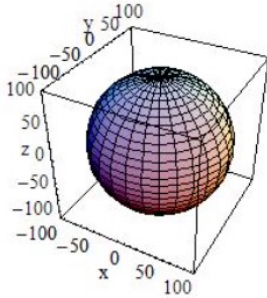
Chem3 Application Code Structure

chem3.cc	Main file ⇒ creation of RunManager , DetectorConstruction , PhysicsList , PrimaryGeneratorAction , ActionInitialization
→ RunManager	Management of all the calls to Geant4 kernel and the threads in MT mode
→ DetectorConstruction	Definition of the geometry: World as semi-transparent sphere
→ PhysicsList	Choice of the physics & chemistry lists Default Physics → G4EmDNAPhysics Default Chemistry → G4EmDNAChemistry Chemistry Model → SBS
→ ActionInitialization	Build() ⇒ creation of PrimaryGeneratorAction & optional user action classes: RunAction , TrackingAction , SteppingAction , StackingAction , TimeStepAction , ITrackingInteractivity , ITrackingAction , ITSteppingAction
→ PrimaryGeneratorAction	Choice of the primary particle. Default: 30 MeV Proton

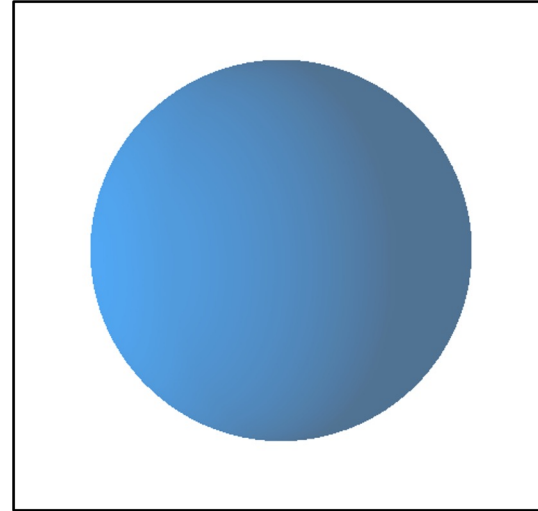
Chem3 Application Code Structure

→ RunAction	To retrieve information in the Beginning and in the End of Run. → In the EndOfRun gives in output the Seed used.
→ TrackingAction	To retrieve information of the Track. At the end of the Primary Track it gives in output the final kinetic energy of the primary.
→ SteppingAction	To retrieve information from a given "step".
→ StackingAction	When no more "physical tracks" remain, the method StackingAction::NewStage is called → The Chemistry Stage starts.
→ TimeStepAction	Allows the user to set minimal time step values and to retrieve information from a given time step (molecule names, reaction products, etc...).
→ ITTrackingInteractivity	Mandatory to visualize chemical tracks and use stepping/tracking actions.
→ ITTrackingAction	To retrieve information from a given track. Similar to TrackingAction but for chemistry.
→ ITSteppingAction	To retrieve information from a given track. Similar to SteppingAction but for chemistry.

Geometry Definition



In the picture:
pRmax = 100



- The World Volume is a G4ORB
- Two parameters define the geometry :
 - the material of the Sphere → for Geant4-DNA it has to be water.
 - the radius of the Sphere: double worldRadius = 0.3*micrometer;

Physics / Chemistry List

Physics Models Included:

- G4EmDNAPhysics ← default
- G4EmDNAPhysics_option1
- G4EmDNAPhysics_option2
- G4EmDNAPhysics_option3
- G4EmDNAPhysics_option4
- G4EmDNAPhysics_option5
- G4EmDNAPhysics_option6
- G4EmDNAPhysics_option7
- G4EmDNAPhysics_option8

Chemistry Models Included:

- G4EmDNAChemistry ← default
- G4EmDNAChemistry_option1

N.B.

If the CommandLineParser has the flag "chemOFF" the chemistry is switched off.

Primary Generator

- The primary kinematic consists of a single particle starting at the center of the sphere.
- Particle type
Particle energy
Particle initial position
Particle initial direction
are set in the PrimaryGeneratorAction class, and can be changed via the G4 build-in commands of G4ParticleGun class.
- The chemistry module is triggered in the StackingAction class when all physical tracks have been processed.

default: 30 MeV Proton

/gun/particle e-

/gun/energy 1 keV

/gun/position 0 0 0 micrometer

/gun/direction 0 0 1

TimeStepAction

In ActionInitialisation: `G4Scheduler::Instance()->SetUserAction(new TimeStepAction());`

Methods

- **TimeStepAction()**: Constructor of the TimeStepAction class. Inside it you can set the minimal time steps of your simulation.

<code>AddTimeStep(1 * picosecond, 0.1 * picosecond);</code>	<i>During the first simulated picosecond the minimal time step will be of 0.1 picosecond. If molecules are too close and can react before that time limit: Brownian bridge.</i>
<code>AddTimeStep(10 * picosecond, 1 * picosecond);</code>	<i>From 1 ps to 10 ps in simulation time, the minimal time step will be of 1 ps.</i>

TimeStepAction

In ActionInitialisation: `G4Scheduler::Instance()->SetUserAction(new TimeStepAction());`

Methods

- **StartProcessing()**: Beginning of the chemistry simulation.
- **EndProcessing()**: End of the chemistry simulation.
- **UserPreTimeStepAction()**: If the user wants to do something before the start of the current time step.
- **UserPostTimeStepAction()**: If the user wants to do something after the end of the current time step. Called once after stepping all the tracks.
- **UserReactionAction(Reactif1, Reactif2, Products)**: will be called just after a reaction happened.

How to start?

- Create your own simulation folder, i.e. `"my-simulations"`
- Copy the chem3 example in `"my-simulations"`
- Create a build folder for chem3, i.e. `"chem3-build"`
- Enter inside `"chem3-build"` and execute the cmake command:
`cmake path-of-geant4-install path of chem3`
- Source the geant4.sh:
`cd path-of-geant4-install/bin`
`source geant4.sh`
`cd -`
- Compile the example:
`make -jn` (with n = number of cores available in your machine)
- In the terminal window, inside your chem3-build folder:
`./chem3 -gui tcsh` or `./chem3 -gui qt`
- In the gui window:
`/control/execute beam.in`

beam.in Macro Commands

```
#/tracking/verbose 1
```

=> Print information

```
/gun/position 0 0 0 micrometer
```

```
/gun/direction 0 0 1
```

=> Define the source

```
/gun/particle e-
```

```
/gun/energy 1 keV
```

```
#
```

```
#=====
```

```
# NB: comment lines when the flag chemOFF is used
```

=> Print additional chemistry information

```
#/scheduler/verbose 2
```

```
#/scheduler/whyDoYouStop
```

```
#=====
```

```
#
```

=> Define the number of primary events

```
/run/beamOn 1
```

Output

Defined reaction table

Physics stage ends		
Number of chemical species involved in reactions = 6		
Reaction	Reaction Rate [dm3/(mol*s)]	Interaction Range for chosen reaction model [nm]
H3O ⁺ + e _{aq} ⁻ → H [•]	2.11e+10	0.200589
H3O ⁺ + OH ⁻ → No product	1.43e+11	1.34973
OH [•] + e _{aq} ⁻ → OH ⁻	2.95e+10	0.506256
OH [•] + OH [•] → H2O2 [•]	4.4e+09	0.207651
OH [•] + H [•] → No product	1.44e+10	0.194167
e _{aq} ⁻ + e _{aq} ⁻ → OH ⁻ + OH ⁻ + H ₂ [•]	5e+09	0.134838
e _{aq} ⁻ + H [•] → OH ⁻ + H ₂ [•]	2.65e+10	0.294265
e _{aq} ⁻ + H2O2 [•] → OH ⁻ + OH [•]	1.41e+10	0.295745
H [•] + H [•] → H ₂ [•]	1.2e+10	0.226528

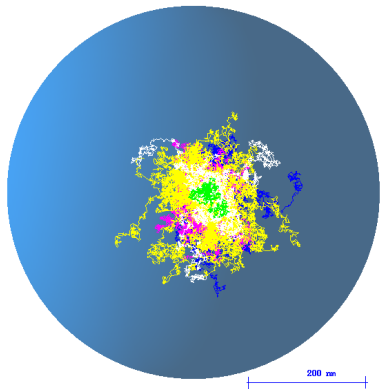
DNAMolecularStepByStepModel will be used

Chemical reactions

```

Physics stage ends
*** G4Scheduler starts processing
At time : 1 ps Reaction : OH^-1 (-112) + H3O^1 (-166) => No product
At time : 24.53 ps Reaction : OH^-1 (-145) + OH^-1 (-171) => H2O2^0 (-220)
At time : 37.218 ps Reaction : OH^-1 (-157) + e_aq^-1 (-72) => OH^-1 (-221)
At time : 58.705 ps Reaction : H^0 (-211) + H^0 (-215) => H_2^0 (-222)
At time : 84.673 ps Reaction : OH^-1 (-167) + OH^-1 (-175) => H2O2^0 (-223)
At time : 86.673 ps Reaction : OH^-1 (-153) + OH^-1 (-127) => H2O2^0 (-224)
At time : 90.673 ps Reaction : OH^-1 (-151) + H^0 (-217) => No product
At time : 132.29 ps Reaction : OH^-1 (-143) + OH^-1 (-183) => H2O2^0 (-225)
At time : 171.29 ps Reaction : H3O^1 (-148) + OH^-1 (-221) => No product
At time : 401.3 ps Reaction : OH^-1 (-185) + OH^-1 (-161) => H2O2^0 (-226)
At time : 486.46 ps Reaction : OH^-1 (-155) + e_aq^-1 (-20) => OH^-1 (-227)
At time : 489.46 ps Reaction : H3O^1 (-184) + OH^-1 (-227) => No product
At time : 549.27 ps Reaction : OH^-1 (-141) + e_aq^-1 (-94) => OH^-1 (-228)
At time : 582.27 ps Reaction : OH^-1 (-205) + e_aq^-1 (-99) => OH^-1 (-229)
At time : 637.1 ps Reaction : OH^-1 (-137) + OH^-1 (-218) => H2O2^0 (-230)
At time : 726.44 ps Reaction : e_aq^-1 (-190) + OH^-1 (-181) => OH^-1 (-231)
At time : 774.65 ps Reaction : OH^-1 (-216) + OH^-1 (-206) => H2O2^0 (-232)
At time : 1.003 ns Reaction : H3O^1 (-130) + OH^-1 (-229) => No product
At time : 1.4536 ns Reaction : OH^-1 (-197) + OH^-1 (-209) => H2O2^0 (-233)
At time : 2.0366 ns Reaction : H3O^1 (-182) + e_aq^-1 (-44) => H^0 (-234)
At time : 2.801 ns Reaction : OH^-1 (-179) + OH^-1 (-121) => H2O2^0 (-235)
At time : 3.804 ns Reaction : H3O^1 (-198) + e_aq^-1 (-73) => H^0 (-236)
At time : 3.4335 ns Reaction : e_aq^-1 (-199) + OH^-1 (-169) => OH^-1 (-237)
At time : 3.5283 ns Reaction : H3O^1 (-160) + OH^-1 (-237) => No product
At time : 3.9539 ns Reaction : H3O^1 (-186) + OH^-1 (-231) => No product
At time : 4.2798 ns Reaction : e_aq^-1 (-88) + H^0 (-203) => OH^-1 (-238) + H_2^0 (-239)
At time : 4.6319 ns Reaction : OH^-1 (-228) + H3O^1 (-134) => No product
At time : 4.776 ns Reaction : H3O^1 (-146) + e_aq^-1 (-39) => H^0 (-240)
At time : 5.1574 ns Reaction : OH^-1 (-147) + e_aq^-1 (-110) => OH^-1 (-241)
At time : 6.4734 ns Reaction : H3O^1 (-172) + e_aq^-1 (-53) => H^0 (-242)
At time : 7.4096 ns Reaction : e_aq^-1 (-188) + H^0 (-219) => OH^-1 (-243) + H_2^0 (-244)
At time : 8.1618 ns Reaction : OH^-1 (-113) + OH^-1 (-115) => H2O2^0 (-245)
At time : 11.903 ns Reaction : H3O^1 (-174) + OH^-1 (-241) => No product
At time : 13.103 ns Reaction : e_aq^-1 (-24) + H2O2^0 (-233) => OH^-1 (-246) + OH^-1 (-247)
At time : 16.403 ns Reaction : e_aq^-1 (-22) + OH^-1 (-165) => OH^-1 (-248)
At time : 17.403 ns Reaction : OH^-1 (-133) + OH^-1 (-214) => H2O2^0 (-249)
At time : 21.626 ns Reaction : e_aq^-1 (-87) + OH^-1 (-125) => OH^-1 (-250)
At time : 25.091 ns Reaction : OH^-1 (-187) + e_aq^-1 (-37) => OH^-1 (-251)
At time : 25.291 ns Reaction : H3O^1 (-132) + OH^-1 (-246) => No product
At time : 33.488 ns Reaction : H3O^1 (-195) + OH^-1 (-251) => No product
At time : 33.588 ns Reaction : H2O2^0 (-226) + e_aq^-1 (-50) => OH^-1 (-252) + OH^-1 (-253)
At time : 43.671 ns Reaction : H3O^1 (-142) + OH^-1 (-250) => No product
At time : 56.551 ns Reaction : e_aq^-1 (-85) + OH^-1 (-208) => OH^-1 (-254)
*** G4Scheduler ends at time : 100 ns
  
```

- G4Physics initialization and the defined reaction table are printed.
- ITStepManager processes the chemical stage time step after time step.
- Chemical reactions are printed.
- In the GUI window a cumulative trajectory of the chemical species is drawn.



Movie Macros Scheme

gui

In the terminal window, inside chem3-build folder:

```
./chem3 -gui tcsh
```

movie.mac

In the gui window:

```
/control/execute movie.mac
```

1

movie_prep.mac

beam.in

2

movie_physics.mac

movie_physics.loop

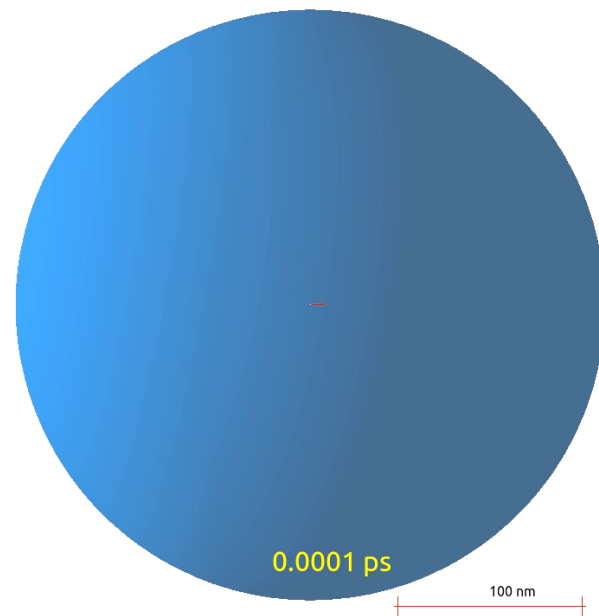
1

movie_chemistry.mac

movie_chemistry.loop

Time Evolution Visualization

- User can start a visualization of the chemical track evolution in time and space using `/control/execute movie.mac`
- Note, that in default setup the simulation requires machine with 6 GB of RAM.
- To lower memory requirements, either:
 - decrease energy of the incident electron in `beam.in`or/and
 - shorten the simulation using `SetEndTime` setting in `src/ActionInitialization.cc` → default 100 ns and in `movie_chemistry.mac`



More Information



GEANT4-DNA : EXTENDING THE GEANT4 MONTE CARLO SIMULATION TOOLKIT FOR RADIOBIOLOGY

Welcome to the web page of the Geant4-DNA project !

The [Geant4](#) general purpose particle-matter Monte Carlo simulation toolkit is being extended with processes for the **modeling of biological damage induced by ionising radiation at the DNA scale**. Such developments are on-going in the framework of the Geant4-DNA project. This project was originally initiated by the [European Space Agency \(ESA\)](#). Developments are undertaken by an [international collaboration](#), coordinated since 2008 by the [National Institute of Nuclear and Particle Physics \(IN2P3\)](#) of the [National Centre for Scientific Research \(CNRS\)](#) in France, in collaboration with the [Geant4@IN2P3](#) activities.

Once published, all developments are freely accessible in **full open access** through the [Geant4 toolkit](#) or through our freely accessible [Geant4 Virtual Machine](#).

Recent posts

June 27th, 2023 : [Geant4 11.1.2](#) LP2i Virtual Machine has been released, see [link](#).

Exercises

- **Exercise 1:** Run as it is the beam.in macro
- **Exercise 2:** Change the type of particle and re-run the beam.in
- **Exercise 3:** Change the particle energy and re-run beam.in
- **Exercise 4:** Change the dimension of the sphere and re-run the beam.in
- **Exercise 5:** Run as it is the movie.mac macro
- **Exercise 6:** Change the type of particle and re-run the movie.mac
- **Exercise 7:** Change the particle energy and re-run movie.mac
- **Exercise 8:** Change the dimension of the sphere and re-run the movie.mac
- **Exercise 9:** Change the “endTime” of the chemistry movie (suggestion: via macro)
- **Exercise 10:** Change the EndTime of the chemistry execution (suggestion: in the proper class...)

Exercise 1: Run as it is the beam.in macro

```
./chem3 -gui tcsh
```

chem3-build-11.2.0.beta — chem3 -gui tcsh — 126x50

```
Physics stage ends
Number of chemical species involved in reactions = 6
Reaction          Reaction Rate [dm3/(mol*s)]  Interaction Range for chosen reaction model [nm]
-----
H30^1 + e_aq^-1 -> H^0          2.11e+10          0.200589
H30^1 + OH^-1 -> No product      1.43e+11          1.34973
OH^0 + e_aq^-1 -> OH^-1         2.95e+10          0.506256
OH^0 + OH^0 -> H2O2^0          4.4e+09           0.207651
OH^0 + H^0 -> No product        1.44e+10          0.194167
e_aq^-1 + e_aq^-1 -> OH^-1 + OH^-1 + H_2^0 5e+09            0.134838
e_aq^-1 + H^0 -> OH^-1 + H_2^0  2.65e+10          0.294265
e_aq^-1 + H2O2^0 -> OH^-1 + OH^0  1.41e+10          0.295745
H^0 + H^0 -> H_2^0             1.2e+10           0.226528

DNAMolecularStepByStepModel will be used
*** G4Scheduler starts processing
At time : 1 ps Reaction : OH^0 (-204) + OH^0 (-205) -> H2O2^0 (-206)
At time : 61.718 ps Reaction : OH^0 (-192) + OH^0 (-112) -> H2O2^0 (-207)
At time : 140.32 ps Reaction : OH^0 (-122) + OH^0 (-164) -> H2O2^0 (-208)
At time : 146.32 ps Reaction : OH^0 (-124) + OH^0 (-126) -> H2O2^0 (-209)
At time : 229.64 ps Reaction : OH^0 (-144) + e_aq^-1 (-99) -> OH^-1 (-210)
At time : 335.08 ps Reaction : H2O2^0 (-206) + e_aq^-1 (-41) -> OH^-1 (-211) + OH^0 (-212)
At time : 341.08 ps Reaction : H30^1 (-196) + OH^-1 (-211) -> No product
At time : 367.13 ps Reaction : e_aq^-1 (-91) + OH^0 (-148) -> OH^-1 (-213)
At time : 388.11 ps Reaction : H30^1 (-187) + e_aq^-1 (-105) -> H^0 (-214)
At time : 473.16 ps Reaction : OH^0 (-183) + OH^0 (-172) -> H2O2^0 (-215)
At time : 559.35 ps Reaction : H30^1 (-149) + OH^-1 (-213) -> No product
At time : 661.87 ps Reaction : OH^0 (-188) + OH^0 (-170) -> H2O2^0 (-216)
At time : 661.87 ps Reaction : H30^1 (-159) + e_aq^-1 (-182) -> H^0 (-217)
At time : 732.01 ps Reaction : e_aq^-1 (-34) + H30^1 (-115) -> H^0 (-218)
At time : 793.47 ps Reaction : OH^-1 (-210) + H30^1 (-171) -> No product
At time : 1.4895 ns Reaction : OH^0 (-116) + H^0 (-218) -> No product
At time : 1.5895 ns Reaction : e_aq^-1 (-33) + OH^0 (-152) -> OH^-1 (-219)
At time : 1.8348 ns Reaction : H30^1 (-127) + e_aq^-1 (-60) -> H^0 (-220)
At time : 1.946 ns Reaction : H30^1 (-157) + OH^-1 (-219) -> No product
At time : 1.956 ns Reaction : H30^1 (-178) + e_aq^-1 (-80) -> H^0 (-221)
At time : 2.2587 ns Reaction : e_aq^-1 (-35) + OH^0 (-128) -> OH^-1 (-222)
At time : 2.2887 ns Reaction : OH^0 (-201) + OH^0 (-202) -> H2O2^0 (-223)
At time : 2.6779 ns Reaction : OH^0 (-150) + OH^0 (-166) -> H2O2^0 (-224)
At time : 3.5594 ns Reaction : e_aq^-1 (-108) + e_aq^-1 (-107) -> OH^-1 (-225) + OH^-1 (-226) + H_2^0 (-227)
At time : 4.2027 ns Reaction : e_aq^-1 (-58) + OH^0 (-180) -> OH^-1 (-228)
At time : 4.3075 ns Reaction : OH^0 (-134) + OH^0 (-158) -> H2O2^0 (-229)
```

beam.in

```
# /tracking/verbose 1
2 /gun/position 0 0 0 micrometer
3 /gun/direction 0 0 1
4 /gun/particle e-
5 /gun/energy 1 keV
6 #
7 #
8 # NB: comment lines related to scheduler when the flag chemOFF is used
9 /scheduler/verbose 2
10 /scheduler/whyDoYouStop
11 #
12 #
13 /run/beamOn 1
14
```

viewer-0 (OpenGLStoredX)

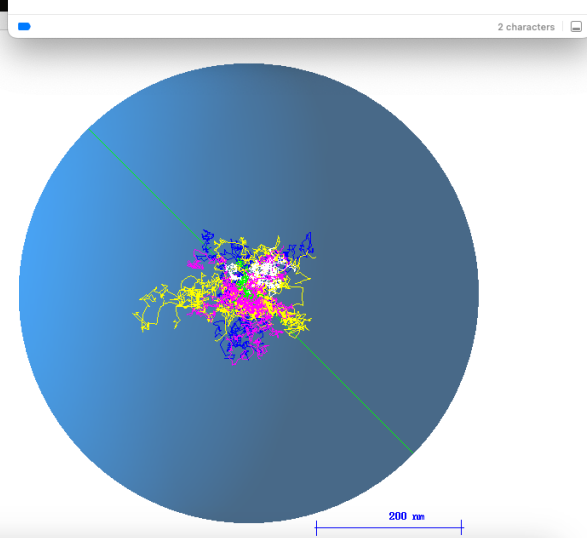
Exercise 2: Change the type of particle and re-run the beam.in

```
./chem3 -gui tcsh
```

```
chem3-build-11.2.0.beta -- chem3 -gui tcsh -- 126x50
hIoni: for p1- X$Type:3 SubType=2
dE/dx and range tables from 10 eV to 300 MeV in 140 bins
Lambda tables from threshold to 300 MeV, 20 bins/decade, spline: 1
StepFunction=(0.1, 0.05 nm); integ: 3; fluct: 1; linLossLim= 0.01
==== EM models for the G4Region DefaultRegionForTheWorld =====
ICRU7300 : Emin= 0 eV Emax=297.505 keV deltaVI
BetheBloch : Emin=297.505 keV Emax= 300 MeV deltaVI
End of tracking primary particle, its final energy is .0 eV
Physics stage ends
Number of chemical species involved in reactions = 6
Reaction Reaction Rate [dm3/(mol*s)] Interaction Range for chosen reaction model [nm]
-----
H30^1 + e_aq^-1 -> H^0 2.11e+10 0.200589
H30^1 + OH^.-1 -> No product 1.43e+11 1.34973
OH^0 + e_aq^-1 -> OH^.-1 2.95e+10 0.506256
OH^0 + OH^0 -> H2O2^0 4.4e+09 0.207651
OH^0 + H^0 -> No product 1.44e+10 0.194167
e_aq^-1 + e_aq^-1 -> OH^.-1 + OH^.-1 + H_2^0 5e+09 0.134838
e_aq^-1 + H^0 -> OH^.-1 + H_2^0 2.65e+10 0.294265
e_aq^-1 + H2O2^0 -> OH^.-1 + OH^0 1.41e+10 0.295745
H^0 + H^0 -> H_2^0 1.2e+10 0.226528
-----
DNAMolecularStepByStepModel will be used
*** G4Scheduler starts processing
At time : 2.132 ps Reaction : OH^0 (-54) + OH^0 (-49) -> H2O2^0 (-67)
At time : 2.4574 ps Reaction : e_aq^-1 (-25) + OH^0 (-62) -> OH^.-1 (-68)
At time : 949.97 ps Reaction : OH^.-1 (-68) + H30^1 (-37) -> No product
At time : 1.8943 ns Reaction : H^0 (-59) + OH^0 (-44) -> No product
At time : 5.7652 ns Reaction : e_aq^-1 (-18) + OH^0 (-40) -> OH^.-1 (-69)
At time : 9.4413 ns Reaction : OH^0 (-61) + OH^0 (-55) -> H2O2^0 (-70)
At time : 9.4513 ns Reaction : e_aq^-1 (-7) + OH^0 (-64) -> OH^.-1 (-71)
At time : 12.343 ns Reaction : H^0 (-50) + OH^0 (-36) -> No product
At time : 14.985 ns Reaction : OH^.-1 (-71) + H30^1 (-43) -> No product
At time : 17.072 ns Reaction : e_aq^-1 (-12) + OH^0 (-66) -> OH^.-1 (-72)
*** G4Scheduler ends at time : 200 ns

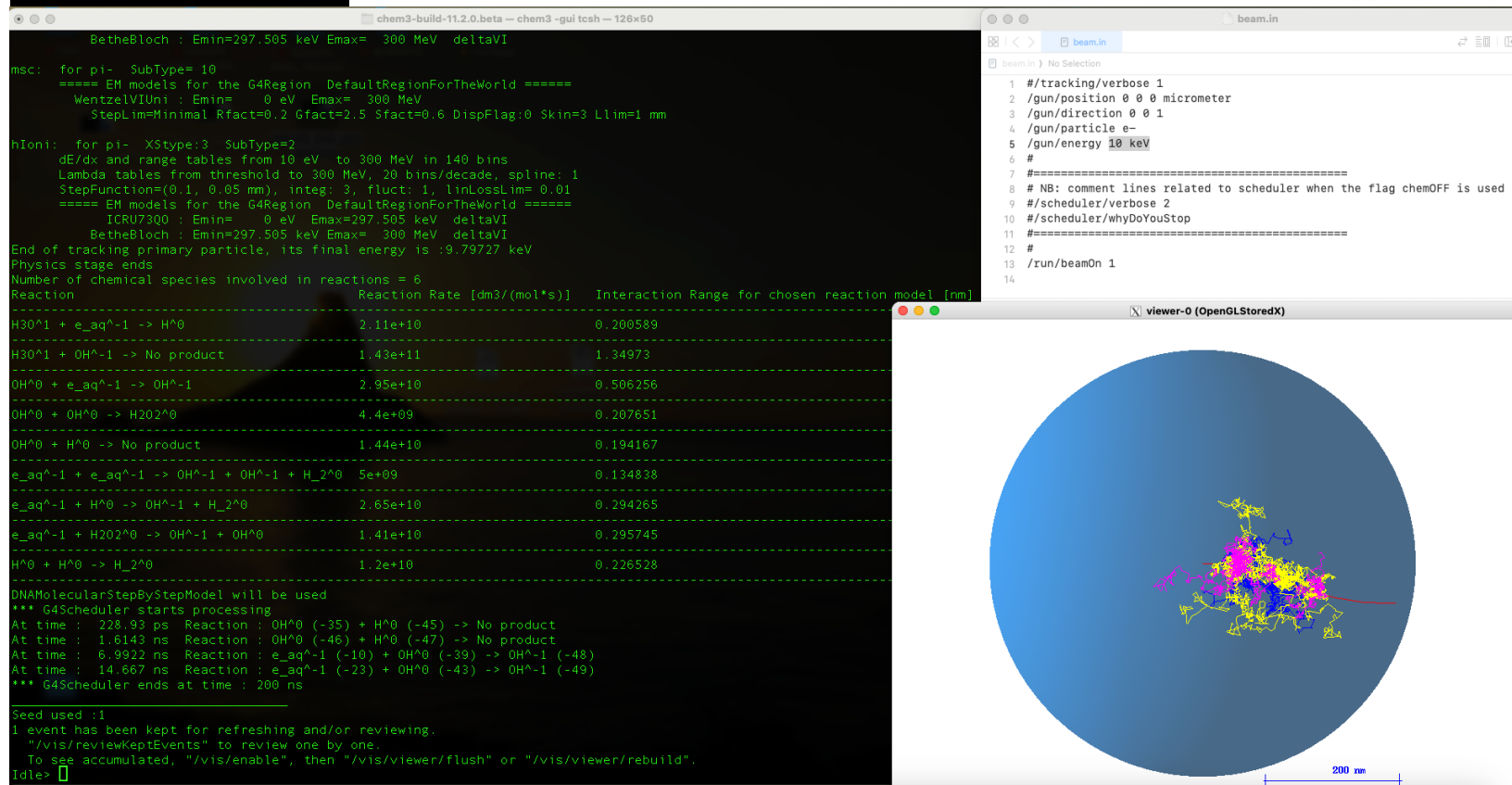
Seed used : 1
1 event has been kept for refreshing and/or reviewing.
"/vis/reviewKeptEvents" to review one by one.
To see accumulated, "/vis/enable", then "/vis/viewer/flush" or "/vis/viewer/rebuild".
Idle>
```

```
beam.in
1 #/tracking/verbose 1
2 /gun/position 0 0 0 micrometer
3 /gun/direction 0 0 1
4 /gun/particle e+
5 /gun/energy 1 keV
6 #
7 #=====
8 # NB: comment lines related to scheduler when the flag chemOFF is used
9 #/scheduler/verbose 2
10 #/scheduler/whyDoYouStop
11 #=====
12 #
13 /run/beamOn 1
14
```



Exercise 3: Change the particle energy and re-run beam.in

```
./chem3 -gui tcsh
```



The image shows a terminal window running the `chem3` program with the `-gui tcsh` option. The terminal output displays the configuration for the G4Region, the EM models, and the tracking of a primary particle. The particle's final energy is 9.79727 keV. A table of chemical reactions is shown, including the reaction of H₃O⁺ with e⁻ and OH⁻, and the reaction of e⁻ with H₂O⁺. The terminal also shows the DNAMolecularStepByStepModel being used and the G4Scheduler starting processing. The output ends with the seed used and the number of events kept for reviewing.

```
BetheBloch : Emin=297.505 keV Emax= 300 MeV deltaVI
msc: for pi- SubType=10
===== EM models for the G4Region_DefaultRegionForTheWorld =====
WentzelVIUni : Emin= 0 eV Emax= 300 MeV
StepLim=Minimal Rfact=0.2 Gfact=2.5 Sfact=0.6 DispFlag:0 Skin=3 Llim=1 mm

hIoni: for pi- XSType:3 SubType=2
dE/dx and range tables from 10 eV to 300 MeV in 140 bins
Lambda tables from threshold to 300 MeV, 20 bins/decade, spline: 1
StepFunction=(0.1, 0.05 mm), integ: 3, fluct: 1, linLossLim: 0.01
===== EM models for the G4Region_DefaultRegionForTheWorld =====
ICRU7300 : Emin= 0 eV Emax=297.505 keV deltaVI
BetheBloch : Emin=297.505 keV Emax= 300 MeV deltaVI
End of tracking primary particle, its final energy is :9.79727 keV
Physics stage ends
Number of chemical species involved in reactions = 6
Reaction Reaction Rate [dm3/(mol*s)] Interaction Range for chosen reaction model [nm]
-----
H30^+1 + e_aq^-1 -> H^0 2.11e+10 0.200589
H30^+1 + OH^-1 -> No product 1.43e+11 1.34973
OH^0 + e_aq^-1 -> OH^-1 2.95e+10 0.506256
OH^0 + OH^0 -> H2O2^0 4.4e+09 0.207651
OH^0 + H^0 -> No product 1.44e+10 0.194167
e_aq^-1 + e_aq^-1 -> OH^-1 + OH^-1 + H_2^0 5e+09 0.134838
e_aq^-1 + H^0 -> OH^-1 + H_2^0 2.65e+10 0.294265
e_aq^-1 + H2O2^0 -> OH^-1 + OH^0 1.41e+10 0.295745
H^0 + H^0 -> H_2^0 1.2e+10 0.226528
-----
DNAMolecularStepByStepModel will be used
*** G4Scheduler starts processing
At time : 228.93 ps Reaction : OH^0 (-35) + H^0 (-45) -> No product
At time : 1.6143 ns Reaction : OH^0 (-46) + H^0 (-47) -> No product
At time : 6.9922 ns Reaction : e_aq^-1 (-10) + OH^0 (-39) -> OH^-1 (-48)
At time : 14.667 ns Reaction : e_aq^-1 (-23) + OH^0 (-43) -> OH^-1 (-49)
*** G4Scheduler ends at time : 200 ns

Seed used :1
1 event has been kept for refreshing and/or reviewing.
"/vis/reviewKeptEvents" to review one by one.
To see accumulated, "/vis/enable", then "/vis/Viewer/flush" or "/vis/Viewer/rebuild".
Idle>
```

The `beam.in` file contains the following configuration:

```
1 #/tracking/verbose 1
2 /gun/position 0 0 0 micrometer
3 /gun/direction 0 0 1
4 /gun/particle e-
5 /gun/energy 10 keV
6 #
7 #=====
8 # NB: comment lines related to scheduler when the flag chemOFF is used
9 #/scheduler/verbose 2
10 #/scheduler/whyDoYouStop
11 #=====
12 #
13 /run/beamOn 1
14
```

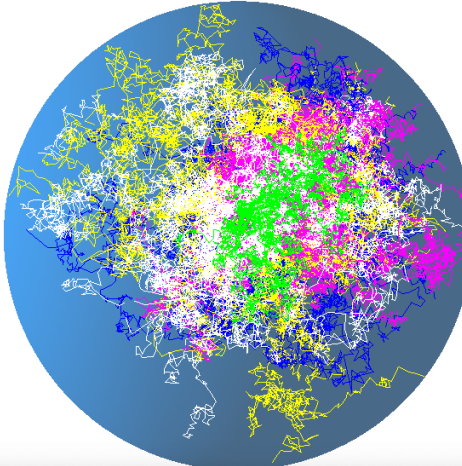
The 3D visualization window shows a molecular structure with a scale bar of 200 nm.

Exercise 4: Change the dimension of the sphere and re-run the beam.in

make

```
./chem3 -gui tcsh
```

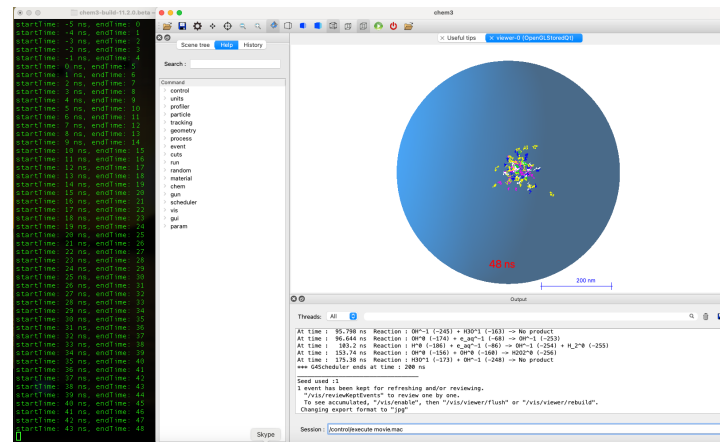
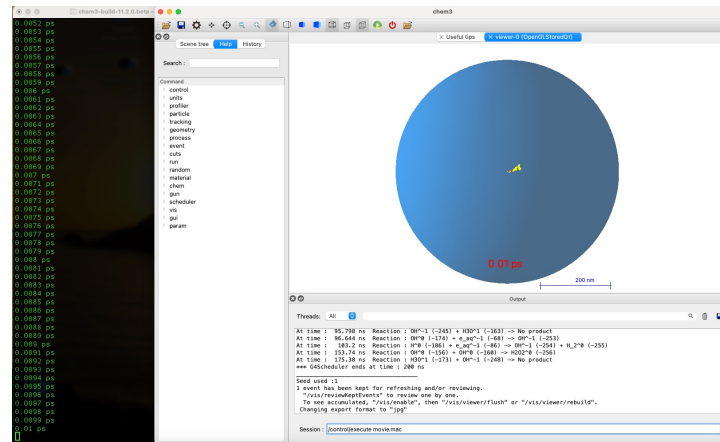
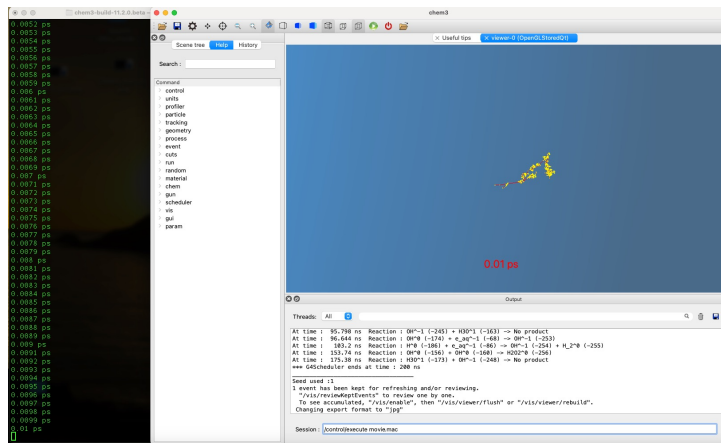
```
*** G4Scheduler starts processing
At time : 1 ps Reaction : OH^0 (-204) + OH^0 (-205) -> H2O2^0 (-206)
At time : 61.718 ps Reaction : OH^0 (-192) + OH^0 (-112) -> H2O2^0 (-207)
At time : 140.32 ps Reaction : OH^0 (-122) + OH^0 (-164) -> H2O2^0 (-208)
At time : 146.32 ps Reaction : OH^0 (-124) + OH^0 (-126) -> H2O2^0 (-209)
At time : 229.64 ps Reaction : e_aq^-1 (-99) + OH^0 (-144) -> OH^-1 (-210)
At time : 335.08 ps Reaction : H2O2^0 (-206) + e_aq^-1 (-41) -> OH^-1 (-211) + OH^0 (-212)
At time : 341.08 ps Reaction : H3O^1 (-196) + OH^-1 (-211) -> No product
At time : 367.13 ps Reaction : e_aq^-1 (-91) + OH^0 (-148) -> OH^-1 (-213)
At time : 388.11 ps Reaction : e_aq^-1 (-105) + H3O^1 (-187) -> H^0 (-214)
At time : 473.16 ps Reaction : OH^0 (-183) + OH^0 (-172) -> H2O2^0 (-215)
At time : 559.35 ps Reaction : OH^-1 (-213) + H3O^1 (-149) -> No product
At time : 661.87 ps Reaction : OH^0 (-188) + OH^0 (-170) -> H2O2^0 (-216)
At time : 661.87 ps Reaction : H3O^1 (-159) + e_aq^-1 (-182) -> H^0 (-217)
At time : 732.01 ps Reaction : e_aq^-1 (-34) + H3O^1 (-115) -> H^0 (-218)
At time : 793.47 ps Reaction : OH^-1 (-210) + H3O^1 (-171) -> No product
At time : 1.4895 ns Reaction : OH^0 (-116) + H^0 (-218) -> No product
At time : 1.5895 ns Reaction : e_aq^-1 (-33) + OH^0 (-152) -> OH^-1 (-219)
At time : 1.8348 ns Reaction : e_aq^-1 (-60) + H3O^1 (-127) -> H^0 (-220)
At time : 1.946 ns Reaction : H3O^1 (-157) + OH^-1 (-219) -> No product
At time : 1.956 ns Reaction : e_aq^-1 (-80) + H3O^1 (-178) -> H^0 (-221)
At time : 2.2587 ns Reaction : e_aq^-1 (-35) + OH^0 (-128) -> OH^-1 (-222)
At time : 2.2887 ns Reaction : OH^0 (-201) + OH^0 (-202) -> H2O2^0 (-223)
At time : 2.6779 ns Reaction : OH^0 (-150) + OH^0 (-166) -> H2O2^0 (-224)
At time : 3.5594 ns Reaction : e_aq^-1 (-107) + e_aq^-1 (-108) -> OH^-1 (-225) + OH^-1 (-226) + H_2^0 (-227)
At time : 4.2027 ns Reaction : e_aq^-1 (-58) + OH^0 (-180) -> OH^-1 (-228)
At time : 4.3075 ns Reaction : OH^0 (-134) + OH^0 (-158) -> H2O2^0 (-229)
At time : 4.3881 ns Reaction : OH^-1 (-228) + H3O^1 (-121) -> No product
At time : 4.7245 ns Reaction : e_aq^-1 (-37) + e_aq^-1 (-64) -> OH^-1 (-230) + OH^-1 (-231) + H_2^0 (-232)
At time : 4.9113 ns Reaction : OH^-1 (-231) + H3O^1 (-133) -> No product
At time : 7.2387 ns Reaction : H3O^1 (-139) + e_aq^-1 (-179) -> H^0 (-233)
At time : 8.3472 ns Reaction : OH^0 (-191) + OH^0 (-142) -> H2O2^0 (-234)
At time : 9.5549 ns Reaction : OH^0 (-132) + OH^0 (-136) -> H2O2^0 (-235)
At time : 11.324 ns Reaction : e_aq^-1 (-65) + e_aq^-1 (-73) -> OH^-1 (-236) + OH^-1 (-237) + H_2^0 (-238)
At time : 12.724 ns Reaction : H2O2^0 (-215) + e_aq^-1 (-82) -> OH^-1 (-239) + OH^0 (-240)
At time : 19.528 ns Reaction : OH^0 (-212) + e_aq^-1 (-54) -> OH^-1 (-241)
At time : 25.605 ns Reaction : OH^0 (-240) + e_aq^-1 (-84) -> OH^-1 (-242)
At time : 26.805 ns Reaction : OH^-1 (-236) + H3O^1 (-147) -> No product
At time : 29.187 ns Reaction : OH^0 (-114) + OH^0 (-118) -> H2O2^0 (-243)
At time : 30.287 ns Reaction : e_aq^-1 (-43) + e_aq^-1 (-63) -> OH^-1 (-244) + OH^-1 (-245) + H_2^0 (-246)
At time : 30.487 ns Reaction : OH^0 (-197) + OH^0 (-194) -> H2O2^0 (-247)
At time : 31.69 ns Reaction : e_aq^-1 (-77) + H2O2^0 (-209) -> OH^-1 (-248) + OH^0 (-249)
At time : 31.89 ns Reaction : OH^-1 (-239) + H3O^1 (-117) -> No product
At time : 34.908 ns Reaction : H3O^1 (-198) + OH^-1 (-237) -> No product
At time : 55.073 ns Reaction : e_aq^-1 (-45) + H2O2^0 (-247) -> OH^-1 (-250) + OH^0 (-251)
At time : 75.197 ns Reaction : OH^-1 (-245) + H3O^1 (-161) -> No product
At time : 91.857 ns Reaction : H3O^1 (-111) + OH^-1 (-242) -> No product
At time : 169.23 ns Reaction : OH^0 (-185) + OH^0 (-140) -> H2O2^0 (-252)
*** G4Scheduler ends at time : 200 ns
```



```
DetectorConstruction.cc
C:\DetectorConstruction
C:\DetectorConstruction:ConstructDetector()
97 G4VPhysicalVolume* DetectorConstruction::ConstructDetector()
98 {
99     G4Material *water = OtherMaterial("G4_WATER");
100
101     // WORLD VOLUME = an G4ORb FULL OF LIQUID WATER
102
103     double worldRadius = 0.1*micrometer;
104
105     G4Orb* solidWorld = new G4Orb("World", worldRadius);
106
107     G4LogicalVolume* logicWorld =
108         new G4LogicalVolume(solidWorld, //its solid
109                             water, //its material
```

Exercise 5: Run as it is the movie.mac macro

```
./chem3 -gui qt  
Idle>/control/execute movie.mac
```

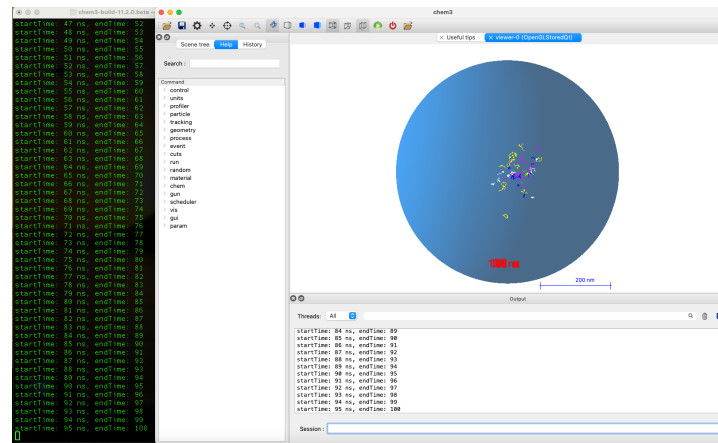
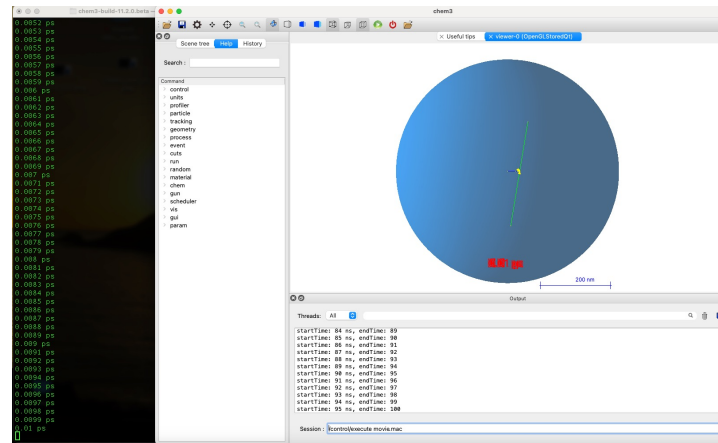
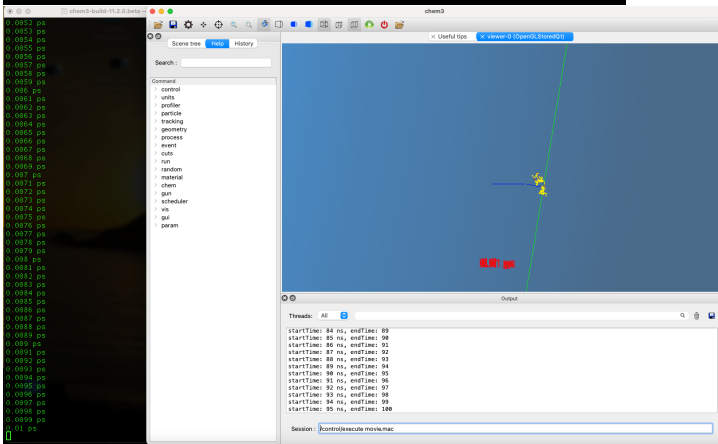


Exercise 6: Change the type of particle and re-run the movie.mac

```
beam.in
```

```
1 #/tracking/verbose 1
2 /gun/position 0 0 0 micrometer
3 /gun/direction 0 0 1
4 /gun/particle e+
5 /gun/energy 1 keV
6 #
7 #=====
8 # NB: comment lines related to scheduler when the flag chemOFF is used
9 #/scheduler/verbose 2
10 #/scheduler/whyDoYouStop
11 #=====
12 #
13 /run/beamOn 1
14
```

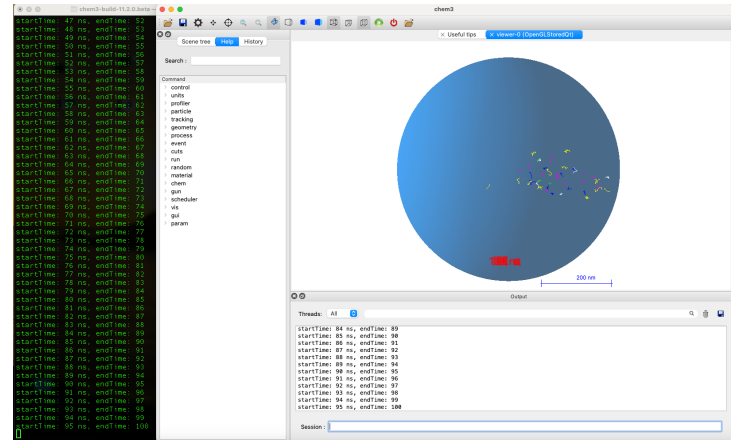
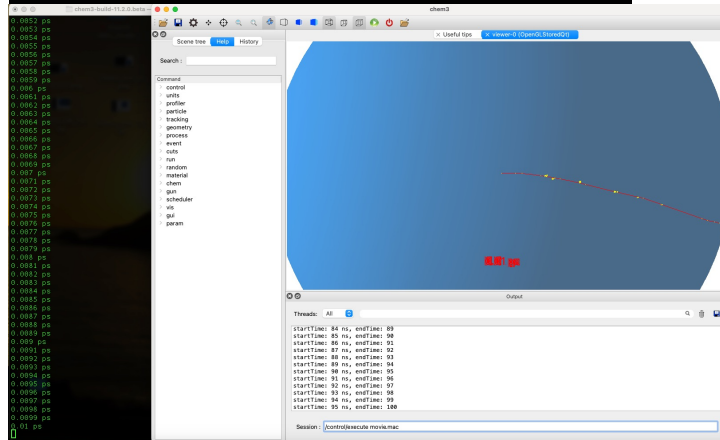
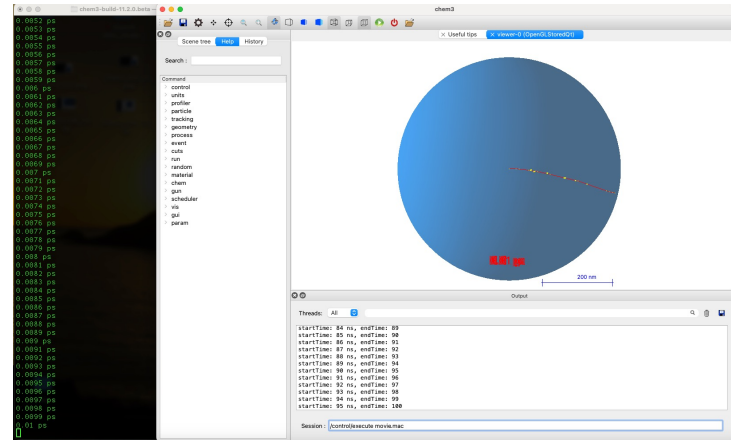
```
Idle>/control/execute movie.mac
```



Exercise 7: Change the particle energy and re-run movie.mac

```
beam.in
1 #/tracking/verbose 1
2 /gun/position 0 0 0 micrometer
3 /gun/direction 0 0 1
4 /gun/particle e-
5 /gun/energy 10[ keV
6 #
7 #=====
8 # NB: comment lines related to scheduler when the flag chemOFF is used
9 #/scheduler/verbose 2
10 #/scheduler/whyDoYouStop
11 #=====
12 #
13 /run/beamOn 1
```

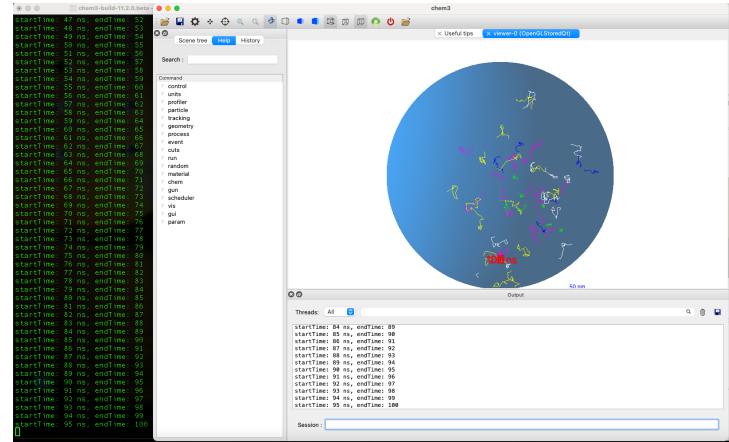
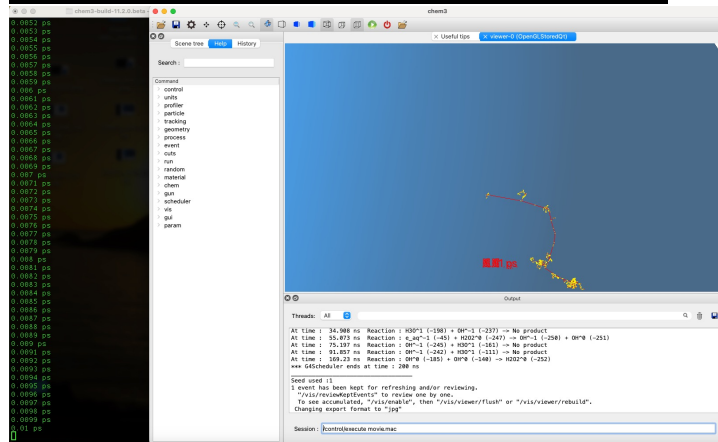
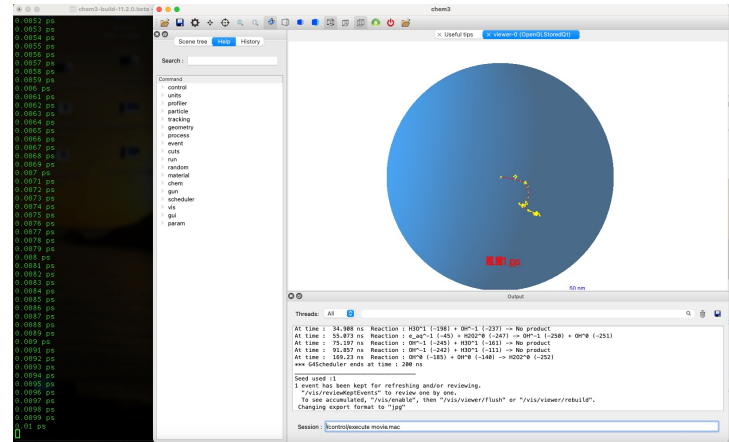
Idle>/control/execute movie.mac



Exercise 8: Change the dimension of the sphere and re-run the movie.mac

```
C# DetectorConstruction ) M DetectorConstruction::ConstructDetector()
94
95 //.....ooo0000oooo.....ooo0000oooo.....ooo0000oooo.....ooo0000oooo.....
96
97 G4PhysicalVolume* DetectorConstruction::ConstructDetector()
98 {
99     G4Material *water = OtherMaterial("G4_WATER");
100
101     // WORLD VOLUME = an G4ORB FULL OF LIQUID WATER
102
103     double worldRadius = 0.1*micrometer;
104
105     G4Orb* solidWorld = new G4Orb("World", worldRadius);
```

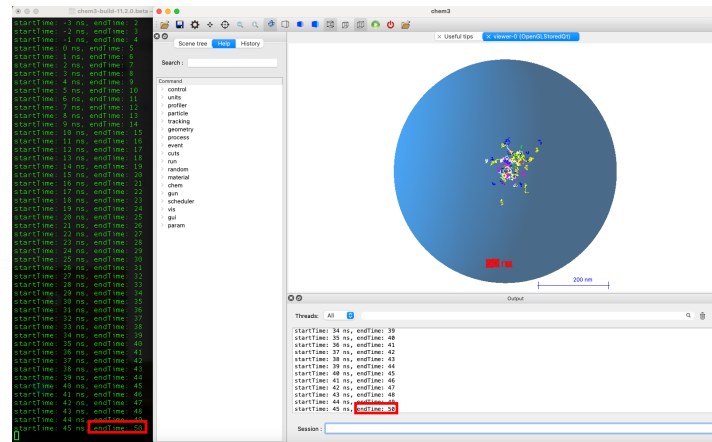
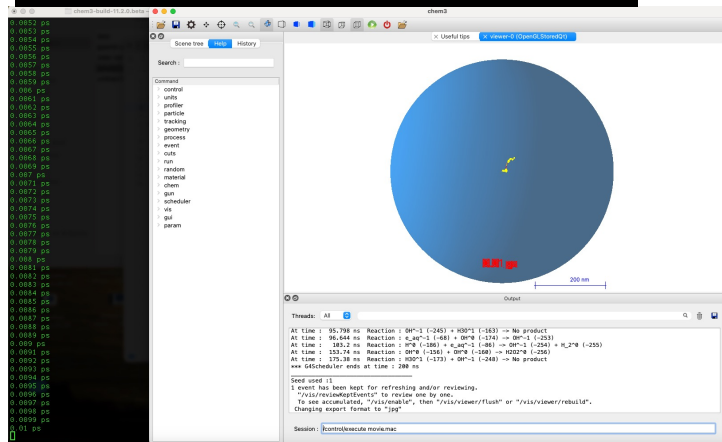
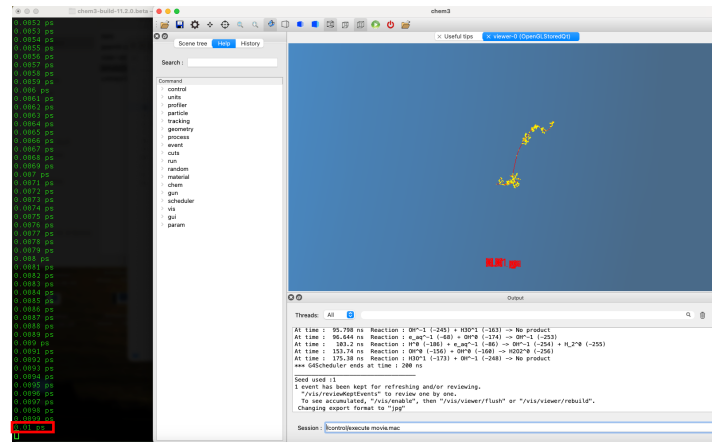
```
make
./chem3 -gui tcsh
Idle>/control/execute movie.mac
```



Exercise 9: Change the “endTime” of the chemistry movie

```
movie_chemistry.mac  
movie_chemistry.mac ) No Selection  
1 /vis/modeling/trajectories/drawByParticleID-0/default/setDrawStepPts false  
2  
3 # end time of chemistry in ns (change units in chemistry.loop  
4 /control/alias endTime 45  
5 /vis/viewer/set/timeWindow/startTime 0 ps  
6 #/vis/viewer/set/timeWindow/endTime 1 ps  
7  
8 /vis/viewer/set/timeWindow/fadeFactor 0  
9 /vis/viewer/set/timeWindow/displayHeadTime true -0.1 -0.8 24 1.0 0.0 0.0  
10 /control/alias timeWindow 5  
11 /control/loop movie_chemistry.loop startTime -{timeWindow} {endTime} 1  
12
```

Idle>/control/execute movie.mac



Exercise 10: change the EndTime of the chemistry execution

```
C:\ActionInitialization> ActionInitialization.cpp
72 //.....0000000000.....0000000000.....0000000000.....0000000000.....
74 ActionInitialization::ActionInitialization()
75 {}
77 //.....0000000000.....0000000000.....0000000000.....0000000000.....
79
80 void ActionInitialization::BuildForMaster() const
81 {
82     // In MT mode, to be clearer, the RunAction class for the master thread might
83     // be different than the one used for the workers.
84     // This RunAction will be called before and after starting the
85     // workers.
86     // For more details, please refer to :
87     // https://twiki.cern.ch/twiki/bin/view/Geant4/Geant4MTForApplicationDevelopers
88
89     // RunAction_runAction new RunAction();
90     // SetUserAction(runAction);
91 }
92
93 void ActionInitialization::Build() const
94 {
95     PrimaryGeneratorAction primGenAction = new PrimaryGeneratorAction;
96     SetUserAction(primGenAction);
97
98     // Set optional user action classes
99     SetUserAction(new RunAction());
100     SetUserAction(new TrackingAction());
101     SetUserAction(new SteppingAction());
102     SetUserAction(new StackingAction());
103
104     // chemistry part
105     if(G4DNACheckManager::IsActivated()){
106         G4Scheduler::Instance()->SetUserAction(new TimeStepAction());
107
108         // Uncomment and set to stop chemistry stage after:
109         // ..given number of time steps
110         //G4Scheduler::Instance()->SetMaxNbSteps(1000);
111
112         // ...ON reaching this time
113         G4Scheduler::Instance()->SetEndTime(50+nanosecond);
114     }
115 }
```

```
make
./chem3 -gui tcsh
Idle>/control/execute movie.mac
```

```
chem3-build-11.2.0.beta -- chem3 -gui qt -- 92x50
Physics stage ends
** G4Scheduler starts processing
At time: 1.6357 ns Reaction H30*1 (-232) + OH*-1 (-114) -> No product
At time: 1.6357 ps Reaction OH*0 (-153) + OH*0 (-209) -> H2O2*0 (-236)
At time: 1.6357 ps Reaction OH*0 (-233) + OH*0 (-115) -> H2O2*0 (-237)
At time: 23.697 ps Reaction H30*1 (-130) + OH*-1 (-120) -> No product
At time: 34.359 ps Reaction OH*0 (-214) + OH*0 (-229) -> H2O2*0 (-238)
At time: 36.359 ps Reaction OH*0 (-234) + OH*0 (-207) -> H2O2*0 (-239)
At time: 77.8 ps Reaction OH*0 (-173) + OH*0 (-123) -> H2O2*0 (-240)
At time: 89.796 ps Reaction OH*0 (-157) + OH*0 (-127) -> H2O2*0 (-243)
At time: 116.8 ps Reaction OH*0 (-213) + OH*0 (-212) -> H2O2*0 (-243)
At time: 128.8 ps Reaction OH*0 (-120) + H*0 (-231) -> No product
At time: 712.73 ps Reaction e_aq*-1 (-73) + OH*0 (-145) -> OH*-1 (-243)
At time: 931.26 ps Reaction OH*0 (-167) + OH*0 (-187) -> H2O2*0 (-244)
At time: 953.16 ps Reaction OH*0 (-139) + OH*0 (-159) -> H2O2*0 (-245)
At time: 965.16 ps Reaction OH*-1 (-243) + H30*1 (-186) -> No product
At time: 987.41 ps Reaction OH*0 (-155) + e_aq*-1 (-92) -> OH*-1 (-246)
At time: 1.0927 ns Reaction OH*0 (-151) + H*0 (-224) -> No product
At time: 1.2927 ns Reaction H30*1 (-126) + OH*-1 (-246) -> No product
At time: 1.5331 ns Reaction H*0 (-235) + e_aq*-1 (-52) -> OH*-1 (-247) + H_2*0 (-248)
At time: 1.7927 ns Reaction H30*1 (-183) + OH*-1 (-247) -> No product
At time: 1.8327 ns Reaction OH*0 (-177) + H*0 (-206) -> No product
At time: 1.8727 ns Reaction e_aq*-1 (-51) + OH*0 (-203) -> OH*-1 (-249)
At time: 1.8827 ns Reaction e_aq*-1 (-55) + OH*0 (-189) -> OH*-1 (-250)
At time: 1.9827 ns Reaction OH*0 (-175) + e_aq*-1 (-59) -> OH*-1 (-251)
At time: 2.0218 ns Reaction H30*1 (-192) + OH*-1 (-117) -> No product
At time: 2.7879 ns Reaction OH*0 (-225) + OH*0 (-205) -> H2O2*0 (-252)
At time: 2.976 ns Reaction H30*1 (-172) + OH*-1 (-251) -> No product
At time: 3.4618 ns Reaction OH*0 (-217) + OH*0 (-209) -> H2O2*0 (-253)
At time: 3.9918 ns Reaction OH*0 (-181) + e_aq*-1 (-75) -> OH*-1 (-254)
At time: 4.1371 ns Reaction e_aq*-1 (-65) + OH*0 (-228) -> OH*-1 (-255)
At time: 5.0284 ns Reaction e_aq*-1 (-83) + OH*0 (-118) -> OH*-1 (-256)
At time: 7.0573 ns Reaction H30*1 (-178) + OH*-1 (-248) -> No product
At time: 8.3057 ns Reaction H30*1 (-138) + OH*-1 (-256) -> No product
At time: 8.7761 ns Reaction OH*0 (-125) + e_aq*-1 (-79) -> OH*-1 (-257)
At time: 8.7761 ns Reaction H30*1 (-124) + OH*-1 (-257) -> No product
At time: 9.5049 ns Reaction e_aq*-1 (-111) + OH*0 (-197) -> OH*-1 (-258)
At time: 11.715 ns Reaction e_aq*-1 (-38) + e_aq*-1 (-202) -> OH*-1 (-259) + OH*-1 (-260) + H_2*0 (-261)
At time: 12.833 ns Reaction H30*1 (-164) + OH*-1 (-250) -> No product
At time: 14.433 ns Reaction e_aq*-1 (-66) + OH*0 (-223) -> OH*-1 (-262)
At time: 17.233 ns Reaction OH*0 (-149) + OH*0 (-131) -> H2O2*0 (-263)
At time: 20.923 ns Reaction H30*1 (-173) + OH*-1 (-254) -> No product
At time: 23.333 ns Reaction H30*1 (-122) + e_aq*-1 (-112) -> H*0 (-264)
At time: 34.29 ns Reaction e_aq*-1 (-85) + H30*1 (-144) -> H*0 (-265)
At time: 40.772 ns Reaction e_aq*-1 (-105) + OH*0 (-199) -> OH*-1 (-266)
** G4Scheduler ends at time: 40.772 ns
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