Electromagnetic Physics II Low Energy Models

Geant4 Tutorial at Chalk River

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using slides of Sebastien Incerti (CNRS, Bordeaux)

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Outline

- Context
- Physics Processes and Models
 - Livermore, including polarized photon models
 - Penelope models
 - Ion ICRU'73 model
 - Geant4-DNA processes and models (beyond physics)
 - MicroElec processes and models
 - Monash University models
 - Atomic de-excitation processes and models
- Implementing a physics list with low energy EM settings
- Documentation

Purpose

- Extend the coverage of Geant4 electromagnetic interactions with matter
 - for photons, electrons, positrons and ions
 - down to very low energies (sub-keV scale)
- Possible domains of applications
 - Space science
 - Medical physics
 - Underground physics
 - Microdosimetry and nanodosimetry for radiobiology and microelectronics
 - ...
- Main choices of physics models include
 - Livermore : electrons and photons [250 eV* GeV]
 - Penelope: electrons, positrons and photons [100 eV* 1 GeV]
 - Microdosimetry & nanodosimetry models
 - Geant4-DNA project: [eV ~ few 100 MeV]
 - MicroElec for Silicon : [eV 10 GeV]

Software design

- Identical to the software design proposed by the Standard EM working group
 - Applicable to all low energy electromagnetic software classes
 - Allows a coherent approach to the modelling of all electromagnetic interactions
- A physical interaction or process is described by a <u>PROCESS CLASS</u>
 - Naming scheme : « G4ProcessName »
 - Eg. : « G4ComptonScattering » for photon Compton scattering
- A physical process can be simulated according to several models, each model being described by a <u>MODEL CLASS</u>
 - Naming scheme : « G4ModelNameProcessNameModel »
 - Eg. : « G4LivermoreComptonModel » for the Livermore Compton model
 - Models can be alternative and/or complementary in certain energy ranges
- According to the selected model, model classes provide the computation of
 - the process total cross section & the stopping power
 - the process final state (kinematics, production of secondaries...)
- All required data files are located in the \$G4LEDATA directory

Livermore models

- Full set of models for electrons and gammas
- Based on publicly available evaluated data tables from the Livermore data library
 - EADL: Evaluated Atomic Data Library - Alternative set by Bearden for fluoresence lines
 - EEDL : Evaluated Electrons Data Library
 - EPDL97 : Evaluated Photons Data Library
 - EPICS2014 for photoelectric effect
 - Mixture of experiments and theories
 - Binding energies: Scofield
- Data tables are interpolated by Livermore model classes to compute
 - □ Total cross sections: photoelectric, Compton, Rayleigh, pair production, Bremsstrahlung
 - Shell integrated cross sections: photo-electric, ionization
 - Energy spectra: secondary e- and gamma
- Validity range (recommended): 250 eV (recommended)
 - Processes can be used down to 100 eV, with a reduced accuracy
 - Technically, down to ~10 eV
- Included elements from Z=1 to Z=100
 - Include atomic effects (fluorescence, Auger)
 - \blacksquare Atomic relaxation : Z > 5 (EADL transition data)
- Naming scheme: G4LivermoreXXXModel (eg. G4LivermoreComptonModel)

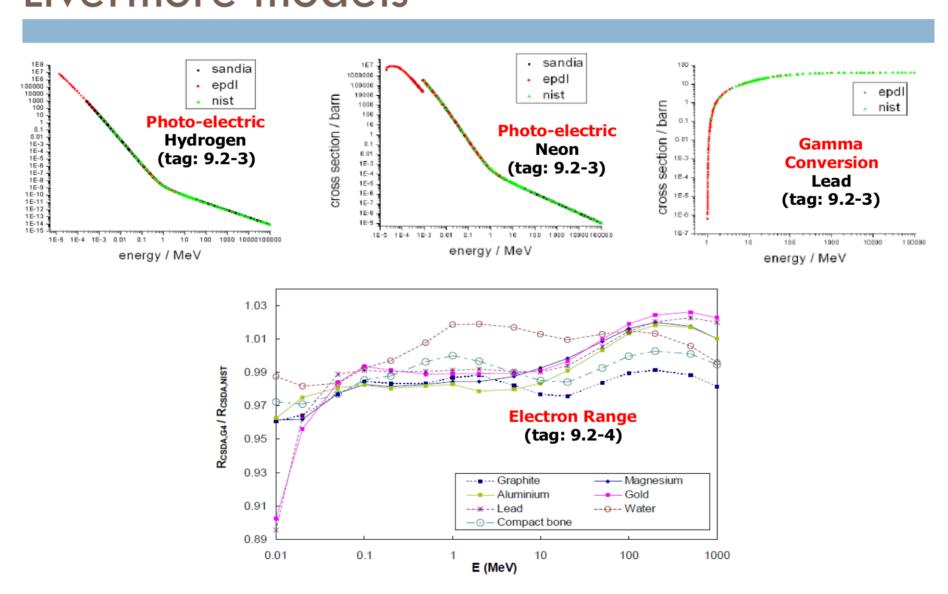
See http://www-nds.iaea.org/epdl97

Available Livermore models

Physics Process	Process Class	Model Class	Low Energy Limit	
Gammas				
Compton	G4ComptonScattering	G4LivermoreComptonModel	eV	
Polarized Compton	G4ComptonScattering	G4LivermorePolarizedComptonModel	eV	
Rayleigh	G4RayleighScattering	G4LivermoreRayleighModel	eV	
Polarized Rayleigh	G4RayleighScattering	G4LivermorePolarizedRayleighModel	250 eV (kill)	
Conversion	G4GammaConversion	G4Liver more Gamma Conversion Model	1.022 MeV	
Polarized Conversion	G4GammaConversion	G4LivermorePolarizedGammaConversion Model	1.022 MeV	
Photo-electric	G4PhotoElectricEffect	G4LivermorePhotoElectricModel	eV	
Polarized Photo-electric	G4PhotoElectricEffect	G4LivermorePolarizedPhotoElectricModel	eV	
Electrons				
Ionization	G4elonisation	G4LivermorelonisationModel	eV	
Bremsstrahlung	G4eBremsstrahlung	G4 Liver more Bremsstrahlung Model	10 eV	

Eg. of verification of Livermore models

Nucl. Instrum. and Meth. A 618 (2010) 315-322



Polarized Livermore models

- Describe in detail the kinematics of polarized <u>photon</u> interactions
- Based on the Livermore database
- Possible applications of such developments
 - design of space missions for the detection of polarized photons
- Naming scheme: G4LivermorePolarizedXXXModel
 - eg. G4LivermorePolarizedComptonModel
- More in the following publications

```
Nucl. Instrum. Meth. A 566 (2006) 590-597 (Photoelectric)
Nucl. Instrum. Meth. A 512 (2003) 619-630 (Compton and Rayleigh)
Nucl.Instrum. Meth. A 452 (2000) 298-305 (Pair production)
```

Penelope physics

 Geant4 includes the low-energy models for electrons, positrons and photons from the Monte Carlo code PENELOPE (PENetration and Energy LOss of Positrons and Electrons) version 2008

```
Nucl. Instrum. Meth. B 350 (2015) 41-48
Nucl. Instrum. Meth. B 207 (2003) 107-123
```

- Physics models
 - Specifically developped by the group of F. Salvat et al.
 - Great care dedicated to the low-energy description
 - Atomic effects, fluorescence, Doppler broadening...
- Mixed approach: analytical, parameterized & database-driven
 - Recommended applicability energy range: 100 eV 1 GeV
- Also include positrons
 - Not described by Livemore models
- G4PenelopeXXXModel (e.g. G4PenelopeComptonModel)

Available Penelope models

Physics Process	Process Class	Model Class	Low Energy Limit	High Energy Limit	
Gammas					
Compton	G4ComptonScattering	G4PenelopeComptonModel	eV	1 GeV	
Rayleigh	G4RayleighScattering	G4PenelopeRayleighModel	eV	1 GeV	
Conversion	G4GammaConversion	G4PenelopeGammaConversionModel	1.022 MeV	1 GeV	
Photo-electric	G4PhotoElectricEffect	G4PenelopePhotoElectricModel	eV	1 GeV	
Electrons/Positrons					
lonization	G4elonisation	G4PenelopelonisationModel	eV	1 GeV	
Bremsstrahlung	G4eBremsstrahlung	G4PenelopeBremsstrahlungModel	eV	1 GeV	
Positrons					
Annihilation	G4eplusAnnihilation	G4PenelopeAnnihilationModel	eV	1 GeV	

Ion energy loss model

- Describes the energy loss of ions heavier than Helium due to interactions with atomic electrons of target atoms
- This model computes
 - Cross sections for the <u>discrete</u> production of <u>delta</u> rays
 - Delta rays are only produced above the production threshold, which inherently also governs the discrete energy loss of ions
 - Restricted electronic stopping powers, that is the continuous energy loss of ions as they slow down in an absorber
 - Below the production threshold
- Mainly for medical and space applications
- See

Nucl. Instrum. Meth. B 268 (2010) 2343-2354

Ion energy loss model

Restricted stopping powers are calculated using 3 approaches

- $T < T_{Low}$: Free electron gas model
- $T_{Low} \le T \le T_{High}$: parameterization (ICRU'73) recommendation
- $T > T_{High}$: Bethe-Bloch formula (using an effective charge and higher order corrections)

ICRU'73 parameterization

- Large range of ion-materials combination
 - Incident ions: Li to Ar, and Fe
 - Targets: 25 elemental materials, 31 compounds
- Stopping powers based on the binary theory, effective charge approach for Fe
- Special case: water
 - Revised ICRU'73 tables by P. Sigmund
 - Mean ionization potential is 78 eV
- Energy limits
 - T_{High} = 1 GeV/nucleon
 - $T_{low} = 0.025 \text{ MeV/nucleon (lower boundary of ICRU'73 tables)}$

How to use the ion model?

- Model name: G4IonParametrisedLossModel
- Only applicable to ions with Z≥3
- Already included in Geant4 EM physics constructors
 - Low Energy EM: G4EmLivermorePhysics, G4EmLivermorePolarizedPhysics, G4EmPenelopePhysics, G4EmLowEPPhysics
 - Standard EM: G4EmStandard_option3, G4EmStandard_option4
- Designed to be used with the G4ionlonisation() process (from the Standard EM category)
 - Not activated by default when using G4ionlonisation
 - Users can employ this model by using the SetEmModel method of the G4ionlonisation process
- Restricted to one Geant4 particle type: G4Genericlon
 - The process G4ionlonisation is also applicable to alpha particles (G4Alpha) and He3 ions (G4He3), however the G4IonParametrisedLossModel model must not be activated for these light ions
 - Below Z<3, we use G4BraggModel (p) or G4BraggIonModel (alpha), and G4BetheBlochModel with the G4hlonisation and G4ionIonisation processes</p>

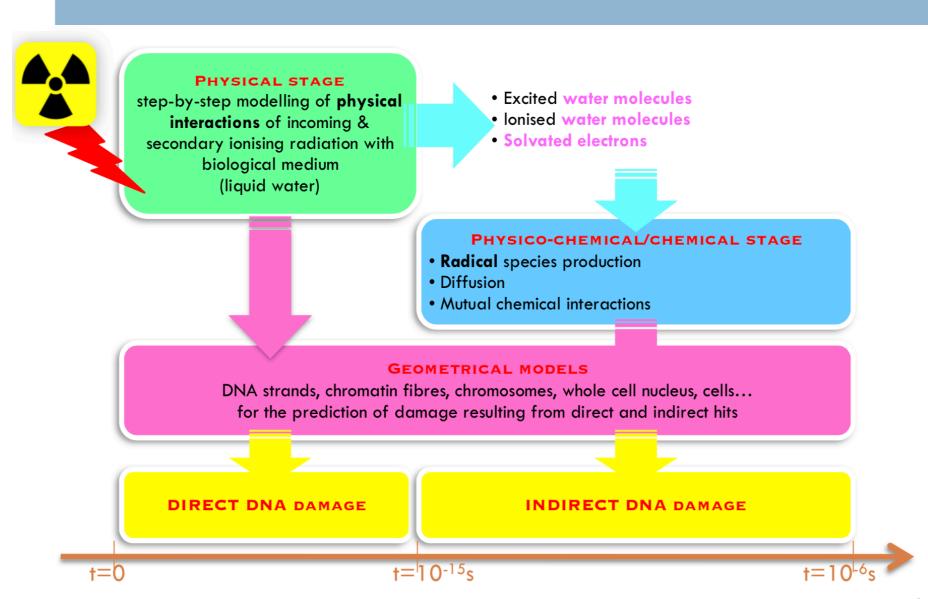
ICRU 73 data tables

- The ion model
 - uses ICRU'73 stopping powers, if the corresponding ion-material combinations are covered by the ICRU'73 report
 - otherwise applies a Bethe-Bloch based formalism
- Elemental materials are matched to the corresponding ICRU 73 stopping powers by means of the atomic number of the material. The material name may be arbitrary in this case.
- For compounds, ICRU 73 stopping powers are used if the material name coincides with the name of Geant4 NIST materials
 - e.g. "G4_WATER"
- For a list of applicable materials, refer to the ICRU 73 report
- All needed data files are in the \$G4LEDATA set of data

Geant4 for microdosimetry in radiobiology

- History
 - initiated in 2001 by Petteri Nieminen (European Space Agency / ESTEC) in the framework of the « Geant4-DNA » project
- Objective: adapt the general purpose Geant4 Monte Carlo toolkit for the simulation of interactions of radiation with biological systems at the cellular and DNA level (« microdosimetry for radiobiology »)
 - Early direct and non-direct radiation effects to DNA in cells
- A fully multidisciplinary activity of the Geant4 Low Energy Electromagnetic Physics working group, involving physicists, chemists, biophysicists...
- Applications
 - Radiobiology, radiotherapy and hadrontherapy
 - eg. early prediction of direct & non-direct DNA strand breaks from ionising radiation
 - Radioprotection for human exploration of Solar system

How can Geant4-DNA simulate early DNA damage?



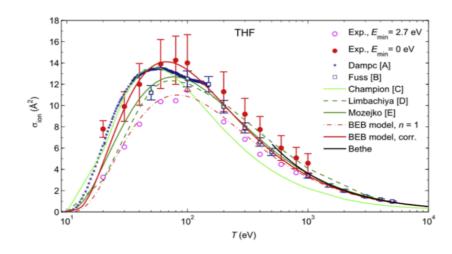
Geant4 for radiobiology

- Several models are available for the description of physical processes involving e⁻, p, H, He, He⁺, He²⁺, Li, Be, B, C, N, O, Si, Fe
- Include elastic scattering, excitation (electronic + vibrations), ionisation, charge exchange and molecular attachment
- These models are valid for liquid water medium and a few biological materials
- Models available in Geant4-DNA
 - are published in the literature
 - may be purely analytical or use interpolated cross section data
- They are all discrete processes!
- Can be combined with other EM categories
 - Standard, LowE thanks to common software design
- Many extended examples in the extended/medical/dna category

Other materials

- Part of the effort to extend Geant4-DNA models to other materials than liquid water
- Cross sections for biological materials are proposed since Geant4 10.4 Beta, applicable to DNA constituents
 - tetrahydrofuran (THF), trimethylphosphate (TMP), pyrimidine (PY) and purine (PU)
 - serving as models for the deoxyribose and phosphate groups in the DNA backbone as well as for the pyrimidine nucleobases, respectively
- For the following incident particles
 - electrons (12 eV-1keV, el. + exci. + ioni.): from measurements @ PTB, Germany
 - protons (70 keV-10 MeV, ioni.) from the HKS approach

Eg. total electron ionisation cross sections in THF



Rad. Phys. Chem. 130 (2017) 459-479

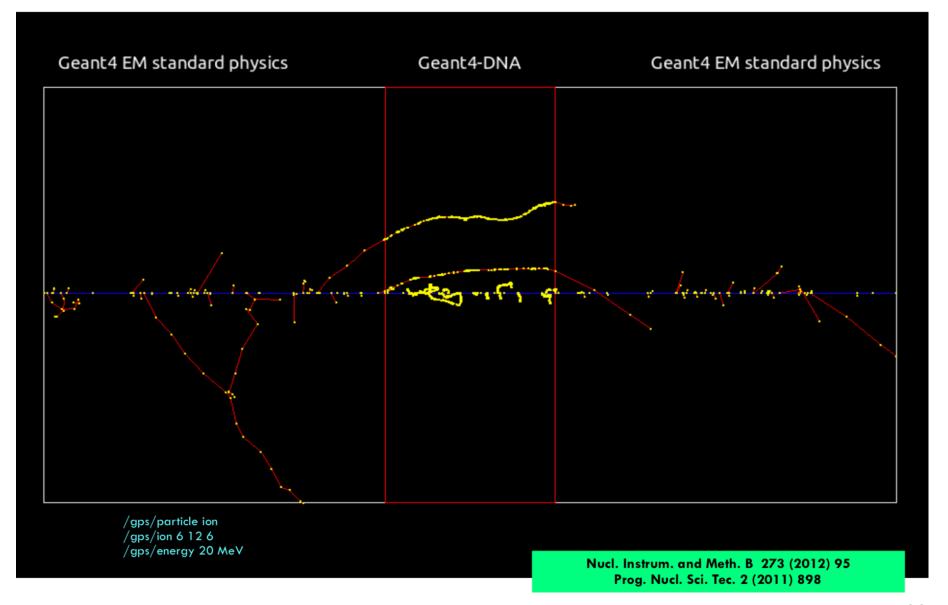
Multiscale combination of EM processes

Thanks to a unified software design, users can easily combine Geant4-DNA processes and models with existing Geant4 physics such as:

- Geant4 photon processes and models
 - Photoelectric effect, Compton sc., Rayleigh sc., pair production
 - Livermore (EPDL97) included by default
- Geant4 alternative electromagnetic processes and models for charged particles
 - lonisation, bremsstrahlung, etc...
 - Electrons, positrons, ions, etc...
- Geant4 atomic deexcitation (fluorescence + Auger emission, including cascades)
 - EADL97, Bearden
- ...and also Geant4 hadronic physics

G4 10.2

Mixed physics lists in geometrical regions: the « microdosimetry » extended example



Geant4-DNA Physics constructors

3 recommended constructors

Constructor name	Content	
G4EmDNAPhysics	Default models	
G4EmDNAPhysics_option1 (beta)	Same as G4EmDNAPhysics but uses New multiple scattering model G4LowEWentzelVIModel	
G4EmDNAPhysics_option2	Same as G4EmDNAPhysics but faster (usage of CDCS for ionisation processes)	
G4EmDNAPhysics_option3	Same as G4EmDNAPhysics (historical)	
G4EmDNAPhysics_option4	Electron ionisation and excitation models by loannina team	
G4EmDNAPhysics_option5 (beta)	Same but faster (usage of CDCS)	
G4EmDNAPhysics_option6	CPA100 models	

All are located in

\$G4INSTALL/source/physics_lists/constructors/electromagnetic

Physico-chemical stage

- During this stage, water molecules
 - Dissociate if ionized
 - Relax or dissociate if excited

Electronic state	Dissociation channels	Fraction (%)
All single ionization states	H ₃ O + + ° OH	100
Excitation state A1B1:	•OH + H•	65
(1b1) → (4a1/3s)	$H_2O + \Delta E$	35
Excitation state B1A1:	H ₃ O ⁺ + * OH + e ⁻ _{aq} (AI)	55
$(3a1) \rightarrow (4a1/3s)$	' OH + ' OH + H ₂	15
(301) 7 (401/38)	$H_2O + \Delta E$	30
Excitation state: Rydberg,	H ₃ O ⁺ + * OH + e ⁻ _{aq} (AI)	50
diffusion bands	$H_2O + \Delta E$	50
Dissociative attachment	*OH + OH- + H ₂	100

Products thermalize down to their energy of diffusion at equilibrium

Chemical stage

Species	Diffusion coefficient D (10 ⁻⁹ m ² s ⁻¹)
H ₃ O +	9.0
H•	7.0
OH-	5.0
e ⁻ aq	4.9
H ₂	5.0
•OH	2.8
H_2O_2	1.4

We followed the set of parameters published by the authors of the PARTRAC software (Kreipl et al., REB 2009). However, these parameters can be modified by the user.

Reaction	Reaction rate (10 ⁷ m ³ mol ⁻¹ s ⁻¹)
$H_3O^+ + OH^- \rightarrow 2 H_2O$	14.3
•OH + e ⁻ _{aq} → OH ⁻	2.95
$H^{\bullet} + e^{-}_{aq} + H_2O \rightarrow OH^{-} + H_2$	2.65
$H_3O^+ + e^{\alpha q} \rightarrow H^{\bullet} + H_2O$	2.11
H• + •OH → H ₂ O	1.44
$H_2O_2 + e^{aq} \rightarrow OH^- + \bullet OH$	1.41
$H^{\bullet} + H^{\bullet} \rightarrow H_2$	1.20
e- _{aq} + e- _{aq} + 2 H ₂ O→ 2 OH- + H ₂	0.50
•OH + •OH → H ₂ O ₂	0.44



A new interface to describe geometries in Geant4-DNA

- PDB : Protein Data Bank
 http://www.rcsb.org/pdb/
 - 3D structure of molecules
 - Proteins
 - Nucleic acids
- Description of DNA molecules
 - □ 1FZX.pdb
 - Dodecamer
 - 12 DNA base pairs
 - $(2.8 \times 2.3 \times 4.01 \text{ nm}^3)$
 - 1ZBB.pdb
 - Tetranucleosome
 - 2 nucleosomes: 347 pairs of bases
 - $(9.5 \times 15.0 \times 25.1 \text{ nm}^3)$

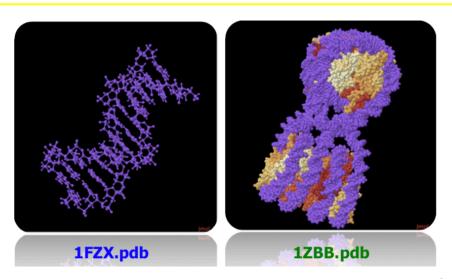
```
HEADER STRUCTURAL PROTEIN/DNA 08-APR-05 1ZBB TITLE STRUCTURE OF THE 4_601_167 TETRANUCLEOSOME ...

ATOM 1 05' DAI 1 70.094 16.969 123.433 0.50238.00 0 ATOM 2 C5' DAI 1 70.682 18.216 123.054 0.50238.00 C ATOM 3 C4' DAI 1 69.655 19.289 122.776 0.50238.00 C ...

TER 14223 DT J 347 ...

HELIX 1 1 GLY A 44 SER A 57 1 14 HELIX 2 2 ARG A 63 ASP A 77 1 15 ...

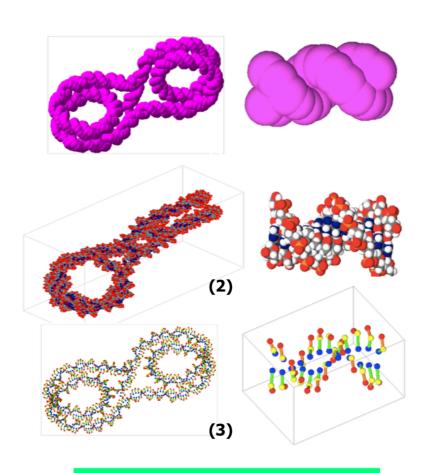
SHEET 1 A 2 ARG A 83 PHE A 84 0 SHEET 2 A 2 THR B 80 VAL B 81 1 O VAL B 81 N ARG A 83
```





« pdb4dna » extended example

- 1) A C++ library
 - Reading of PDB files
 - Build bounding boxes from atom coordinates
 - Search for closest atom from a given point
 - Geometry and visualization: 3 granularities
 - (1) Barycenter of nucleotides
 - (2) Atomistic
 - (3) Barycenter of nucleotide components
- 2) A Geant4-DNA example
 - Water box surrounding the molecule
 - The output results consists in a ROOT file, containing for each event:
 - energy deposit in bounding boxes
 - number of single strand breaks (SSB)
 - number of double strand breaks (DSB)



Geant4-DNA examples included in Geant4

Example code name	Purpose	Location	
dnaphysics	 Usage of Geant4-DNA Physics processes variable density 	\$G4INSTALL/examples/extended/medical/dna	
microdosimetry	Combination of Standard EM or Low Energy EM processes with Geant4-DNA Physics processes	\$G4INSTALL/examples/extended/medical/dna	
range	Range simulation with Geant4-DNA	\$G4INSTALL/examples/extended/medical/dna	
slowing	Calculation of electron slowing down spectra	\$G4INSTALL/examples/extended/medical/dna	
spower	Calculation of stopping power	\$G4INSTALL/examples/extended/medical/dna	
svalue	Usage of Geant4-DNA Physics processes in spheres	\$G4INSTALL/examples/extended/medical/dna	
wvalue	Calculation of W values	\$G4INSTALL/examples/extended/medical/dna	
clustering	Clustering code	\$G4INSTALL/examples/extended/medical/dna	
icsd	Usage of alternative materials	\$G4INSTALL/examples/extended/medical/dna	
chem1, chem2, chem3, chem4	Usage of Geant4-DNA chemistry	\$G4INSTALL/examples/extended/medical/dna	
wholeNuclearDNA	Cell nucleus	\$G4INSTALL/examples/extended/medical/dna	
pdb4dna	Interface to PDB database	\$G4INSTALL/examples/extended/medical/dna	
microbeam	3D cellular phantom	\$G4INSTALL/examples/advanced	
neuron	3D neural network	\$G4INSTALL/examples/extended/medical/dna	
TestEm12	DPK	\$G4INSTALL/examples/extended	
TestEm14	Extraction of cross sections	\$G4INSTALL/examples/extended	

Geant4-DNA website

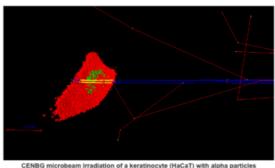
http://geant4-dna.org

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

Welcome to the Internet 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 early biological damage induced by ionising radiation at the DNA scale. Such developments are on-going in the framework of the Geant4-DNA project - originally initiated by the European Space Agency/ESTEC - and are undertaken by an international collaboration.

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



robeam irradiation of a keratinocyte (HaCaT) with alpha particles see the « microbeam » Geant4 advanced example -- movie courtesy of L. Garnier (CNRS) -





New processes and models for microelectronics

- Purpose
 - extend Geant4 with processes and models for the simulation of particlematter interactions in highly integrated microelectronic components
 - for electrons, protons, heavy ions in Silicon
- They use the same step-by-step approach as Geant4-DNA processes and models
 - Similarly based on the complex dielectric function theory
- Applicable to the « G4_Si » NIST material
- Named as « MicroElec » for microelectronics

New processes and models for microelectronics

Processes and models

Physics Process	Process Class	Model Class	Low Energy Limit	High Energy Limit
Electrons				
Elastic scattering	G4MicroElecElastic	G4MicroElecElasticModel	5 eV (kill < 16.7 eV)	100 MeV
Ionization	G4MicroElecInelastic	G4MicroElecInelasticModel	16.7 eV	100 MeV
Protons and heavy ions				
Ionization	G4MicroElecInelastic	G4MicroElecInelasticModel	50 keV	10 GeV

- A dedicated advanced user example is available (« microelectronics »)
- Validation range

■ Electrons: 50 eV – 50 keV

Protons: 50 keV – 23 MeV

Nucl. Instrum. Meth B 288 (2012) 66 – 73 Nucl. Instrum. Meth B 287 (2012) 124 – 129 IEEE Trans. Nucl. Sci. 59 (2012) 2697 – 2703

Improved Compton model

- Monash U. (J. M. C. Brown) recently proposed to improve the accuracy of Livermore gamma models
 - Unpolarized Compton scattering off atomic bound electrons in the relativistic impulse approximation, derived from Livermore Compton model
 - Polarized version is also available
 - As an alternative to Compton scattering models (Livermore and Penelope) developed from Ribberfor's Compton scattering framework
 - More accurate electron ejection direction algorithms below 5 MeV
 - Special relativistic formalism + energy & momentum conservation, in order to compute
 - Energy and angular distribution of Compton scattered photons off non-stationary atomic bound electrons
 - Energy and ejected angular distributions of Compton electrons

Improved Compton model

Nucl. Instrum. Meth A 835 (2016) 186 – 225

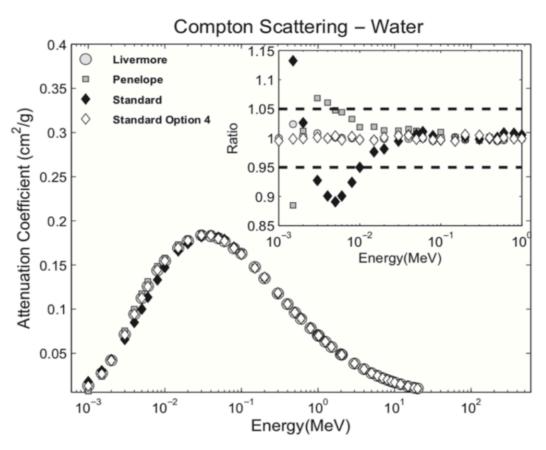


Fig. 11. Compton scattering attenuation coefficient, calculated for different Geant4 models. G4LowEPComptonMode1 is used in the Option4 EM physics configuration. The inset shows the ratio of the coefficient calculated using each alternative Geant4 electromagnetic physics list to the value from NIST XCOM [49]. The dashed lines correspond to a \pm 5% difference.

Improved Compton model

- Model class is G4LowEPComptonModel
 (or G4LowEPPolarizedComptonModel for the polarized version)
- You can register it easily to your Physics list

```
G4ComptonScattering* cs = new G4ComptonScattering;

cs->SetEmModel(new G4KleinNishinaModel(),1);

G4VEmModel* theLowEPComptonModel = new G4LowEPComptonModel();

theLowEPComptonModel->SetHighEnergyLimit(20*MeV);

cs->AddEmModel(0, theLowEPComptonModel);

ph->RegisterProcess(cs, particle);
```

- You can also use two Physics constructors
 - G4EmLowEPPhysics identical to G4EmLivermorePhysics except for Compton
 - G4EmStandard_option4 « best » of Geant4 EM

Atomic de-excitation effects

- Atomic de-excitation is initiated by other EM processes
 - E.g.: photo-electric effect, Compton, ionisation by e- and ions
 - Leave the atom in an excited state
- EADL data contain transition probabilities
 - radiative: fluorescence
 - non-radiative:
 - Auger e-: inital and final vacancies in different sub-shells
 - Coster-Kronig e-: identical sub-shells
- Alternative set for fluorescence lines by Bearden et al. (1967)
 - X-Ray Data Booklet
- Thanks to a common interface (G4UAtomicDeexcitation), atomic de-excitation is compatible with both Standard & Low Energy Electromagnetic physics categories
 - See more in

NIMB 372 (2016) 91-101 X-Ray Spec. 40 (2011) 135-140

Including atomic effects

Activation of atomic effects can be easily done directly via UI commands

```
/process/em/fluo true
/process/em/auger true
/process/em/augerCascade true
/process/em/pixe true
/run/initialize
```

- Boolean parameters in the "/process/em/" deexcitation commands correspond to activation of fluorescence, Auger, Auger cascade, and PIXE respectively
- Note that fluorescence is activated by default in the G4EmDNAPhysics, G4EmLivermorePhysics, G4EmLivermorePolarizedPhysics, G4EmLowEPPhysics, G4EmPenelopePhysics, G4EmStandard_option3 and G4EmStandard_option4 physics constructors while Auger production and PIXE are not
- To select Bearden et al. (1967) fluorescence lines instead of EADL, use (before /run/initialize): /process/em/fluoBearden true (or G4AtomicTransitionManager::Instance()->SetFluoDirectory("fluor_Bearden"); in your Physics list)
- As an example, look into \$G4INSTALL/examples/extended/electromagnetic/TestEm5 and macro pixe.mac

Note on production thresholds

- Remember that production cuts for secondaries are specified as range cuts.

 These are converted at initialisation time into energy thresholds for secondary gamma, electron, positron and proton production.
- A range cut value is set by default to 1.0 mm in Geant4 reference physics lists. This value can be specified in the optional SetCuts() method of the user physics list or via UI commands:
 - for eg. to set a range cut of 10 micrometers, one can use /run/setCut 0.01 mm
 - or, for a given particle type (for e.g. electron) /run/setCutForAGivenParticle e- 0.01 mm
- If a range cut equivalent to an energy lower than 990 eV is specified, then the energy cut is still set to 990 eV.

In order to decrease this value (for eg. down to 250 eV, to see low energy emission lines of the fluorescence spectrum), one can use the UI command:

```
/cuts/setLowEdge 250 eV
```

or alternatively directly in the user physics list, in the optional SetCuts() method, using: G4ProductionCutsTable::GetProductionCutsTable()->SetEnergyRange(250*eV, 1*GeV);

In addition, independently, one can also fully deactivate production cuts for the simulation of all atomic deexcitation products

/process/em/deexcitationIgnoreCut true

In your macro, these UI commands should be put before the UI command

/run/initialize

Recent EM UI commands

- Transport of electrons
 - /process/em/lowestElectronEnergy X ex

In all ionization processes which simulate energy loss along step, a lowest energy limit is introduced, which forces full energy deposition at a step independently on material. Its value may be changed via a new UI commands or through the interface class G4EmParameters.

The default value is 100 eV, for Opt3, Opt4, Livermore, Penelope and LowEEP physics constructors for e+- (see source code)

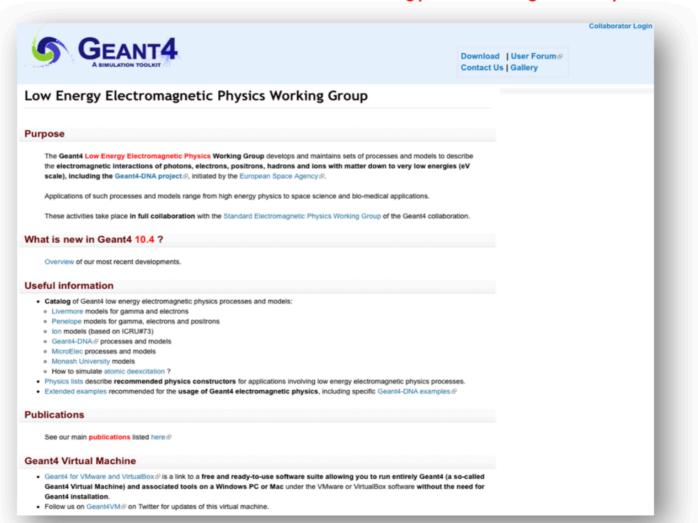
- Configuration per G4Region for PAI, MicroElec, or Geant4-DNA models
 - Done on top of any EM physics constructor
 - Number of G4Regions is not limited
 - See example for PAI models configuration in /example/extended/ electromagnetic/TestEm8
 - Ul commands
 - /process/em/AddPAIRegion proton MYREGION pai
 - /process/em/AddMicroElecRegion MYREGION
 - /process/em/AddDNARegion MYREGION opt0

Web sites

- A unique reference web page on Geant4 EM Physics
 - http://geant4.cern.ch/collaboration/EMindex.shtml
- From there, links to
 - Geant4 Standard Electromagnetic Physics working group pages
 - Geant4 Low Energy Electromagnetic Physics working group pages
- Also from Geant4 web site
 - http://geant4.org
 - Who we are
 - Standard Electromagnetic Physics
 - Low Energy Electromagnetic Physics

Low Energy EM WG TWiki

Geant4 → Collaboration → Low Energy Electromagnetic Physics



Summary: when/why to use the "Low Energy" EM models

- Use Low-Energy models (Livermore or Penelope), as an alternative to Standard models, when you:
 - need precise treatment of EM showers and interactions at low-energy (keV scale or below)
 - are interested in atomic effects, as fluorescence X-rays, Doppler broadening, etc.
 - can afford a more CPU-intensive simulation
 - want to cross-check another simulation (e.g. with a different Physics List)
 - are interested in specific low energy applications (Geant4-DNA, MicroElec)
- Do not use when you are interested in EM physics > MeV
 - same results as Standard EM models
 - strong performance penalty