

<http://geant4.org>

# GEANT4-DNA HANDS ON

KIT Tutorial, October 25-26, 2011, Karlsruhe

# 3 examples

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- We are going to try **3** Geant4 examples which make use of Geant4-DNA Physics processes
  - advanced examples located in [\\$G4INSTALL/examples/advanced](#)
    - dnaphysics
    - microdosimetry
  - extended examples located in [\\$G4INSTALL/examples/extended](#)
    - TestEm1 2

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dnaphysics

# dnaphysics advanced example

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Geant4 9.5 BETA

- Located in `$G4INSTALL/examples/advanced/dnaphysics`
- This example teaches to Geant4 users how to use the Geant4-DNA physics processes and models in a liquid water volume
  - Produces ROOT histograms to visualize track structures
  - The PhysicsList uses the default Physics builder called G4EmDNAPhysics which contains the recommended Geant4-DNA physics, so you do not need to code the physics yourself
- It also explains how to easily change the density of the target material (liquid water) : « variable density material » new feature of Geant4
  - Possibility to investigate density change effects
    - eg. 1.06 g/cm<sup>3</sup> average density of cell nucleus (cf. PARTRAC)

# Install dnaphysics

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- For usage with Geant4 9.4, download the example from <http://geant4.in2p3.fr>
  - Tutorials & Teachings
  - Download « Dnaphysics Hands On »
  - Save it on your desktop
  
- On your Geant4 virtual machine
  - Open a terminal
  - `mv /mnt/hgfs/Desktop/dnaphysics.tar`
  - `tar -xvf dnaphysics.tar`
  - `cd dnaphysics`
  - `snavigator &`
  - `gmake`

# Run dnaphysics

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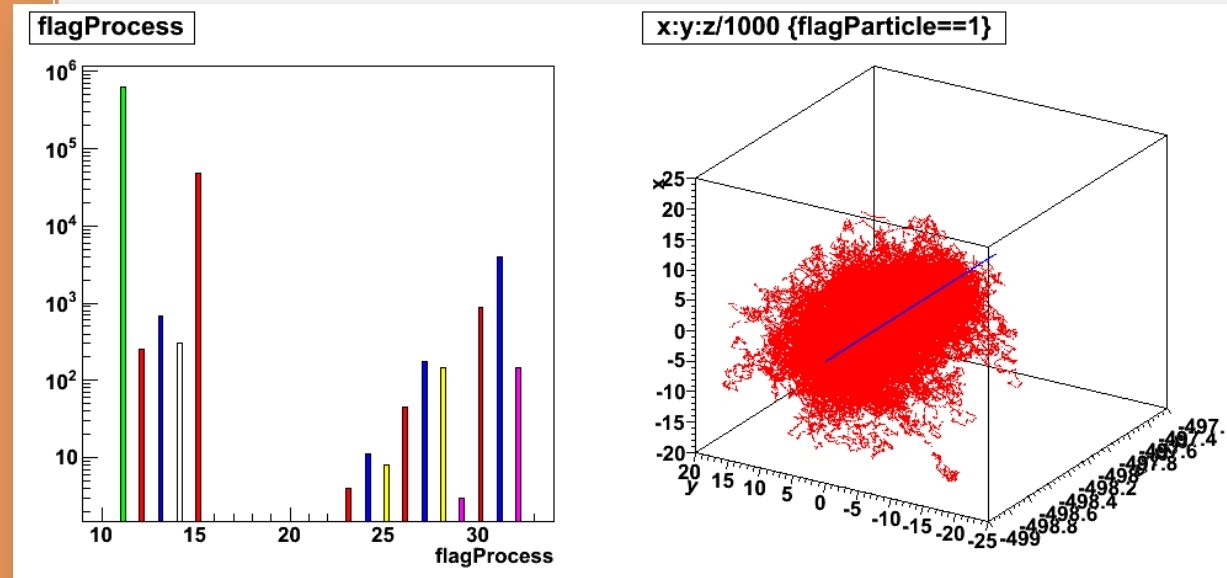
- On your terminal
  - ▣ `$G4WORKDIR/bin/$G4SYSTEM/DNAPhysics`
    - The macro file dna.mac is read
    - 100 electrons of 1 keV are shot
    - No visualization by default
    - Results are saved in dna.root file
  - ▣ `exit`
  - ▣ `root plot.C`
    - ROOT is already installed on your system
    - ROOT macro is read to plot histograms from dna.root file

# Output of dnaphysics

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|                          |    |
|--------------------------|----|
| e-_G4DNAElastic          | 11 |
| e-_G4DNAExcitation       | 12 |
| e-_G4DNAIonisation       | 13 |
| e-_G4DNAAttachment       | 14 |
| e-_G4DNAVibExcitation    | 15 |
|                          |    |
| proton_G4DNAExcitation   | 17 |
| proton_G4DNAIonisation   | 18 |
| proton_G4DNAChargeDec.   | 19 |
|                          |    |
| hydrogen_G4DNAExcitation | 20 |
| hydrogen_G4DNAIonisation | 21 |
| hydrogen_G4DNAChargeInc. | 22 |
|                          |    |
| alpha_G4DNAExcitation    | 23 |
| alpha_G4DNAIonisation    | 24 |
| alpha_G4DNAChargeDec.    | 25 |
|                          |    |
| alpha+_G4DNAExcitation   | 26 |
| alpha+_G4DNAIonisation   | 27 |
| alpha+_G4DNAChargeDec.   | 28 |
| alpha+_G4DNAChargeInc.   | 29 |
|                          |    |
| helium_G4DNAExcitation   | 30 |
| helium_G4DNAIonisation   | 31 |
| helium_G4DNAChargeInc.   | 32 |

Eg. one 100 keV He<sup>+</sup>



# Do more

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- Try yourself to
  - Set number of incident electron to 1
  - Activate visualization
  - Zoom display by 10000
  - run again
  - Try to shoot one proton, then one hydrogen, then one helium with 10 keV incident energy
- In [snavigator](#), look at the following classes
  - **PhysicsList**
    - Usage of G4EmDNAPhysics builder
  - **SteppingAction**
    - Recording of particles & processes
    - Positions of pre-step points
    - Energy deposit for each step
  - **DetectorConstruction**
    - Change of value of water density
  - **HistoManager**
    - Handling of histograms (creation, filling up, saving)



# PhysicsList header

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```
#include "G4VModularPhysicsList.hh"
#include "globals.hh"
class G4VPhysicsConstructor;
class PhysicsList: public G4VModularPhysicsList
{
public:
    PhysicsList();
    ~PhysicsList();
    void ConstructParticle();
    void ConstructProcess();
    void SetCuts();
    void SetCutForGamma(G4double);
    void SetCutForElectron(G4double);
    void SetCutForPositron(G4double);

private:
    G4double cutForGamma;
    G4double cutForElectron;
    G4double cutForPositron;
    G4double currentDefaultCut;
    G4VPhysicsConstructor* emPhysicsList;
};
```

# PhysicsList implementation

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- In constructor
  - `emPhysicsList = new G4EmDNAPhysics();`
- In `ConstructParticle()`
  - `emPhysicsList->ConstructParticle();`
- In `ConstructProcess()`
  - `AddTransportation();`
  - `emPhysicsList->ConstructProcess();`
- Production cuts are **not used** by Geant4-DNA processes
  - All interactions are explicitly simulated
- Look at the Geant4-DNA Physics builder itself
  - `$G4INSTALL/source/physics_list/builders/src/G4EmDNAPhysics.cc`

# What is included in the G4EmDNAPhysics builder ?

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```
void G4EmDNAPhysics::ConstructParticle()  
{  
  // bosons  
  G4Gamma::Gamma();  
  
  // leptons  
  G4Electron::Electron();  
  G4Positron::Positron();  
  
  // baryons  
  G4Proton::Proton();  
  G4GenericIon::GenericIonDefinition();  
  G4DNAGenericIonsManager * genericIonsManager;  
  genericIonsManager=G4DNAGenericIonsManager::Instance();  
  genericIonsManager->GetIon("alpha++");  
  genericIonsManager->GetIon("alpha+");  
  genericIonsManager->GetIon("helium");  
  genericIonsManager->GetIon("hydrogen");  
}
```

Proton and  $\text{He}^{2+}$  nuclei can  
gain electrons and become  
 $\text{H}$ ,  $\text{He}^+$ ,  $\text{He}^0$

# G4EmDNAPhysics: Physics processes & models for electrons

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```
void G4EmDNAPhysics::ConstructProcess()
{
    G4PhysicsListHelper* ph = G4PhysicsListHelper::GetPhysicsListHelper();
    theParticleIterator->reset();
    while( (*theParticleIterator)() )
    {
        G4ParticleDefinition* particle = theParticleIterator->value();
        G4String particleName = particle->GetParticleName();

        if (particleName == "e-") {
            // *** Elastic scattering (two alternative models available) ***
            G4DNAElastic* theDNAElasticProcess = new G4DNAElastic("e-_G4DNAElastic");
            theDNAElasticProcess->SetModel(new G4DNAChampionElasticModel());

            // or alternative model
            //theDNAElasticProcess->SetModel(new G4DNAScreenedRutherfordElasticModel());
            ph->RegisterProcess(theDNAElasticProcess, particle);

            // *** Excitation ***
            ph->RegisterProcess(new G4DNAExcitation("e-_G4DNAExcitation"), particle);
            // *** Ionisation ***
            ph->RegisterProcess(new G4DNAIonisation("e-_G4DNAIonisation"), particle);
            // *** Vibrational excitation ***
            ph->RegisterProcess(new G4DNAVibExcitation("e-_G4DNAVibExcitation"), particle);
            // *** Attachment ***
            ph->RegisterProcess(new G4DNAAttachment("e-_G4DNAAttachment"), particle);
        }
    }
}
```

We have two models for electron elastic scattering in liquid water

PROCESS



MODEL



# G4EmDNAPhysics: Physics processes & models for other Geant4-DNA particles

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```
} else if ( particleName == "proton" ) {  
    ph->RegisterProcess(new G4DNAExcitation("proton_G4DNAExcitation"), particle);  
    ph->RegisterProcess(new G4DNAIonisation("proton_G4DNAIonisation"), particle);  
    ph->RegisterProcess(new G4DNAChargeDecrease("proton_G4DNAChargeDecrease"), particle);  
  
} else if ( particleName == "hydrogen" ) {  
    ph->RegisterProcess(new G4DNAExcitation("hydrogen_G4DNAExcitation"), particle);  
    ph->RegisterProcess(new G4DNAIonisation("hydrogen_G4DNAIonisation"), particle);  
    ph->RegisterProcess(new G4DNAChargeIncrease("hydrogen_G4DNAChargeIncrease"), particle);  
  
} else if ( particleName == "alpha" ) {  
    ph->RegisterProcess(new G4DNAExcitation("alpha_G4DNAExcitation"), particle);  
    ph->RegisterProcess(new G4DNAIonisation("alpha_G4DNAIonisation"), particle);  
    ph->RegisterProcess(new G4DNAChargeDecrease("alpha_G4DNAChargeDecrease"), particle);  
  
} else if ( particleName == "alpha+" ) {  
    ph->RegisterProcess(new G4DNAExcitation("alpha+_G4DNAExcitation"), particle);  
    ph->RegisterProcess(new G4DNAIonisation("alpha+_G4DNAIonisation"), particle);  
    ph->RegisterProcess(new G4DNAChargeDecrease("alpha+_G4DNAChargeDecrease"), particle);  
    ph->RegisterProcess(new G4DNAChargeIncrease("alpha+_G4DNAChargeIncrease"), particle);  
  
} else if ( particleName == "helium" ) {  
    ph->RegisterProcess(new G4DNAExcitation("helium_G4DNAExcitation"), particle);  
    ph->RegisterProcess(new G4DNAIonisation("helium_G4DNAIonisation"), particle);  
    ph->RegisterProcess(new G4DNAChargeIncrease("helium_G4DNAChargeIncrease"), particle);  
}  
}
```



PROCESS

# More on Geant4-DNA Physics

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- Some DNA models « kill » the particle below a certain energy threshold because the models are not validated or defined below this threshold
  - In this case, tracking is stopped and kinetic energy is locally deposited
  - Electrons below 4 eV (by `G4DNAChampionElasticModel`) or below 9 eV (by `G4DNAScreenedRutherfordElasticModel`)
  - Protons & H below 100 eV by `G4DNAIonisation`
  - $\text{He}^{2+}$ ,  $\text{He}^+$  and  $\text{He}^0$  below 1 keV by `G4DNAIonisation`
  
- See other hints from the Geant4-DNA web page
  - <https://twiki.cern.ch/twiki/bin/view/Geant4/LoweMigratedDNAProcesses>
  - How to access total cross sections
  - How to « kill » particles below a certain energy threshold
  - ...

# SteppingAction

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- Flag of particle
  - `if (s->GetTrack()->GetDynamicParticle()->GetDefinition()->GetParticleName() == "e-")  
flagParticle = 1;`
  
- Flag of process
  - `if (s->GetPostStepPoint()->GetProcessDefinedStep()->GetProcessName() == "hydrogen_G4DNAExcitation")  
flagProcess = 20;`
  
- Step information
  - `x=s->GetPreStepPoint()->GetPosition().x()/nanometer;`
  - `y=s->GetPreStepPoint()->GetPosition().y()/nanometer;`
  - `z=s->GetPreStepPoint()->GetPosition().z()/nanometer;`
  - `s->GetTotalEnergyDeposit()/eV`

# DetectorConstruction

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## □ Definition of **water material**

- `// Water is defined from NIST material database`
- `G4NistManager * man = G4NistManager::Instance();`
- `G4Material * H2O = man->FindOrBuildMaterial("G4_WATER");`

## □ **Change of density value** (from Geant4 9.5 BETA)

- `// If one wishes to test other density value for water material, one should use instead:`
- `// G4Material * H2O = man->BuildMaterialWithNewDensity("G4_WATER_MODIFIED", "G4_WATER",  
1000*g/cm/cm/cm);`
- `// Note: any string for "G4_WATER_MODIFIED" parameter is accepted`
- `// and "G4_WATER" parameter should not be changed`

## □ **Display density**

- `G4cout << "-> Density of water material (g/cm3)=" << waterMaterial->GetDensity() / (g/cm/cm/  
cm) << G4endl;`



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# microdosimetry

# microdosimetry advanced example

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- Shows the multi-scale combination of
  - Geant4 Standard EM Physics (condensed/discrete)
    - Geant4 Standard EM
  - Purely discrete processes
    - Geant4-DNA
  - For protons/hydrogen
  - In two regions (the « World » and « Target »)
  
- Useful when the user is interested in simulating efficiently high energy incident particles with Geant4 EM Standard Physics, providing a space phase input for Geant4-DNA simulations applied to much smaller volumes
  
- Also shows how to create a process to kill particles below a certain energy threshold
  - G4ElectronCapture

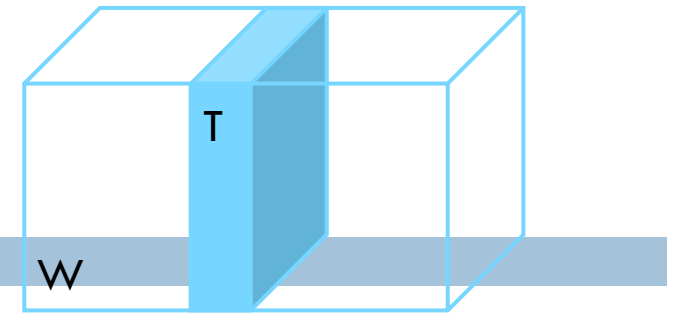
# Install microdosimetry

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- ❑ Open a terminal on your virtual machine
- ❑ Type the following commands:
  - ❑ `cp -R $G4INSTALL/examples/advanced/microdosimetry .`
  - ❑ `cd microdosimetry`
  - ❑ `gmake`
  - ❑ `$G4WORKDIR/bin/$G4SYSTEM/Microdosimetry`
    - ❑ A 5 MeV proton is shot (see [microdosimetry.mac](#))
  - ❑ `exit`
  - ❑ `root plot.C`

# Geometry

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- A **50  $\mu\text{m}$  thick** « Target » volume placed in a **1 mm thick** « World » mother volume
- Both contain liquid water only
- We would like to use
  - **EM Standard** models in the « World »
  - **Geant4-DNA** models in the « Target » below 1 MeV for e-

# 1) Create a **Region** for the **target** volume in DetectorConstruction

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```
fRegion = new G4Region("Target");
```

```
G4ProductionCuts* cuts = new G4ProductionCuts();
```

```
G4double defCut = 1*nanometer;
```

```
cuts->SetProductionCut(defCut,"gamma");
```

```
cuts->SetProductionCut(defCut,"e-");
```

```
cuts->SetProductionCut(defCut,"e+");
```

```
cuts->SetProductionCut(defCut,"proton");
```

```
fRegion->SetProductionCuts(cuts);
```

```
fRegion->AddRootLogicalVolume(logicTarget);
```

## 2) Define Physics for the World

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- In the loop over particles, activate EM standard processes
  - ▣ they will be ACTIVE IN THE WORLD

```
if (particleName == "e-") {
```

```
    // msc is active in the world
```

```
    G4eMultipleScattering* msc = new G4eMultipleScattering();
```

```
    pmanager->AddProcess(msc, -1, 1, 1);
```

```
    // standard ionisation is active in the world
```

```
    G4elonisation* eion = new G4elonisation();
```

```
    eion->SetEmModel(new G4MollerBhabhaModel(), 1);
```

```
    pmanager->AddProcess(eion, -1, 2, 2);
```

# Define Physics for the World

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- then **INACTIVATE Geant4-DNA processes** using a **G4DummyModel** and the **SetModel** method. They will be **INACTIVE IN THE WORLD**.

```
// DNA elastic is not active in the world
G4DNAElastic* theDNAElasticProcess = new G4DNAElastic("e-_G4DNAElastic");
theDNAElasticProcess->SetModel (new G4DummyModel(),1);
pmanager->AddDiscreteProcess(theDNAElasticProcess);

// DNA excitation is not active in the world
G4DNAExcitation* dnaex = new G4DNAExcitation("e-_G4DNAExcitation");
dnaex->SetModel (new G4DummyModel(),1);
pmanager->AddDiscreteProcess(dnaex);

// DNA ionisation is not active in the world
G4DNAIonisation* dnaioni = new G4DNAIonisation("e-_G4DNAIonisation");
dnaioni->SetModel (new G4DummyModel(),1);
pmanager->AddDiscreteProcess(dnaioni);

// DNA attachment is not active in the world
G4DNAAttachment* dnaatt = new G4DNAAttachment("e-_G4DNAAttachment");
dnaatt->SetModel (new G4DummyModel(),1);
pmanager->AddDiscreteProcess(dnaatt);

// DNA vib. excitation is not active in the world
G4DNAVibExcitation* dnavib = new G4DNAVibExcitation("e-_G4DNAVibExcitation");
dnavib->SetModel (new G4DummyModel(),1);
pmanager->AddDiscreteProcess(dnavib);
```

### 3) Define Physics for the Target

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- Inactivate EM Standard processes in the TARGET Region below 1 MeV by registering corresponding EM Standard models to the G4EmConfigurator

```
G4EmConfigurator* em_config = G4LossTableManager::Instance()->EmConfigurator();
```

```
G4VEmModel* mod;
```

```
mod = new G4UrbanMscModel93();
```

```
mod->SetActivationLowEnergyLimit(1*MeV);
```

```
em_config->SetExtraEmModel("e-","msc",mod,"Target");
```

```
mod = new G4MollerBhabhaModel();
```

```
mod->SetActivationLowEnergyLimit(1*MeV);
```

```
em_config->SetExtraEmModel("e-","eloni",mod,"Target",0.0,100*TeV,new G4UniversalFluctuation());
```

Specify  
low energy limit  
of activation

Specify particle name, process  
name, model and region name  
and energy interval  
(option if more than one models  
are used)



# Define Physics for the Target

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- **Activate Geant4-DNA processes in the TARGET Region** by registering the corresponding Geant4-DNA models to the **G4EmConfigurator**

```
mod = new G4DNAChampionElasticModel();  
em_config->SetExtraEmModel("e-","e-_G4DNAElastic",mod,"Target",0.0,1*MeV);
```

```
mod = new G4DNABornIonisationModel();  
em_config->SetExtraEmModel("e-","e-_G4DNAIonisation",mod,"Target",11*eV,1*MeV);
```

```
mod = new G4DNABornExcitationModel();  
em_config->SetExtraEmModel("e-","e-_G4DNAExcitation",mod,"Target",9*eV,1*MeV);
```

```
mod = new G4DNAMeltonAttachmentModel();  
em_config->SetExtraEmModel("e-","e-_G4DNAAttachment",mod,"Target",4*eV,13*eV);
```

```
mod = new G4DNASancheExcitationModel();  
em_config->SetExtraEmModel("e-","e-_G4DNAVibExcitation",mod,"Target",2*eV,100*eV);
```

**Specify energy range  
of applicability**



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# TestEm12

# TestEm 1 2 purpose

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- Shows how to get **energy deposit distribution** for electrons shot within the center of a liquid water sphere
  
- Uses **Physics builders**
  - ▣ Standard EM
  - ▣ Low Energy EM
  - ▣ **Geant4-DNA**
    - To be run with **macro dna.mac**
  
- Produces histograms
  
- Can be fully run & configured through **UI commands**

# Install and run TestEm12

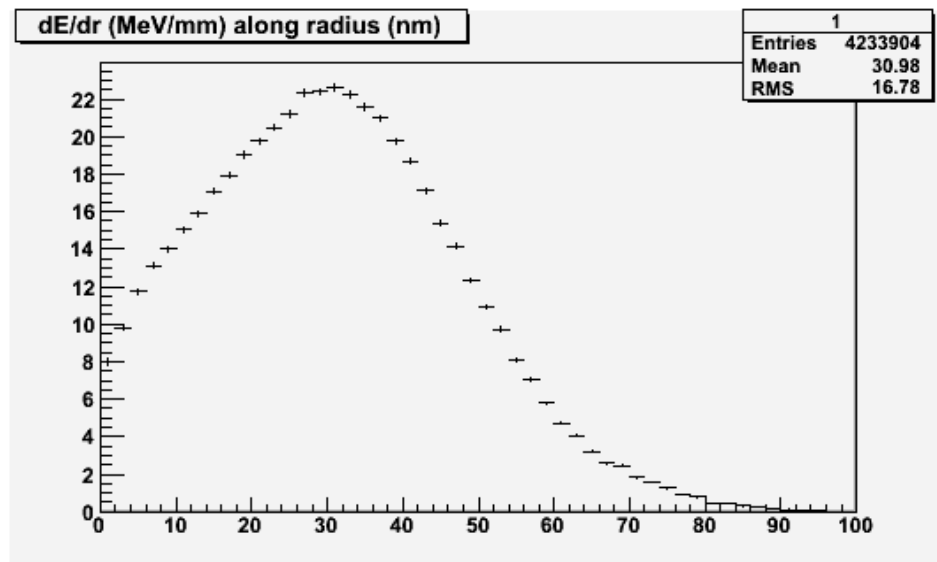
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- Open a terminal on your virtual machine
- Type the following commands:
  - `cp -R $G4INSTALL/examples/extended/electromagnetic/TestEm12 .`
  - `cd TestEm12`
  - `gmake`
  - `$G4WORKDIR/bin/$G4SYSTEM/TestEm12 dna.mac`
    - 5000 electrons of 1 keV are shot from the center of a liquid water sphere of radius 100 nm (see [dna.mac](#))
  - `exit`
  - `root`
  - `Tbrowser g`
- Open the `dna.root` file in the browser

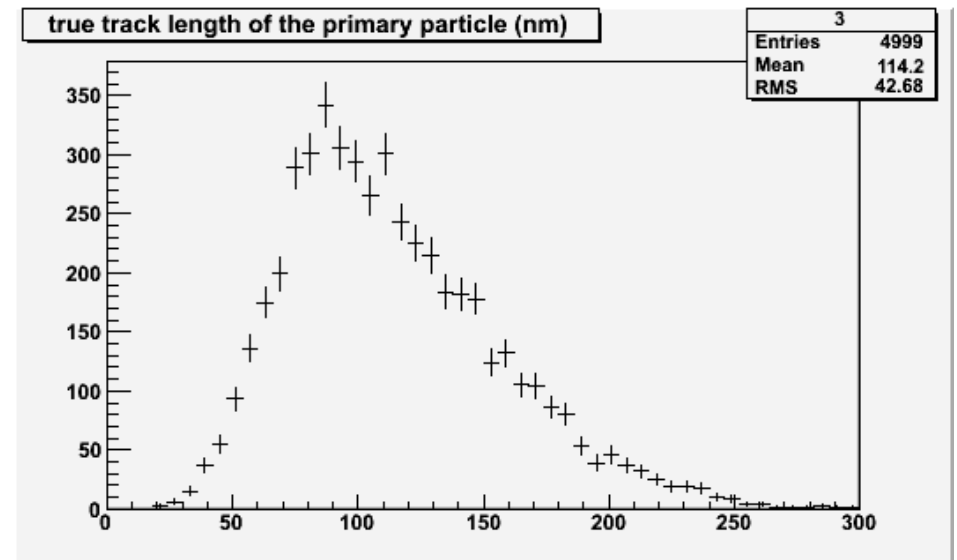
# Example of results

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$dE/dr$  along radius



Track length of primary



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Thank you for your attention