



Exercise

Biassing

# Definition of problem

## Goal

Study the effect of different biasing options using the example from the activation exercise

## Beam characteristics and Geometry (*identical to activation exercise*)

- 28.5 GeV electron beam with a beam power of 20W, Gaussian profile with  $(\sigma_x, \sigma_y) = (0.5, 1.0)$  mm
- cylindrical tunnel section with copper target (25cm long, 3cm in radius) and downstream stainless steel sample (2cm long, 1cm in radius)

## Biasing options to be studied

### 1) **Inelastic interaction length biasing for photons**

- switch off biasing and calculate star density and fluence maps for as many primary particles as needed to obtain the same CPU time as with interaction length biasing
- calculate star density and fluence maps and compare both cases

### 2) **Leading particle biasing in target**

- calculate star density and fluence maps with and without leading particle biasing
- choose the number of primaries such that both runs need the same CPU time

### 3) **Region importance biasing into the sample**

- calculate residual dose rate around the sample with and without region importance biasing
- choose the number of primaries such that both runs need the same CPU time

### 4) **Number of "replicas" in the simulation of the decay radiation**

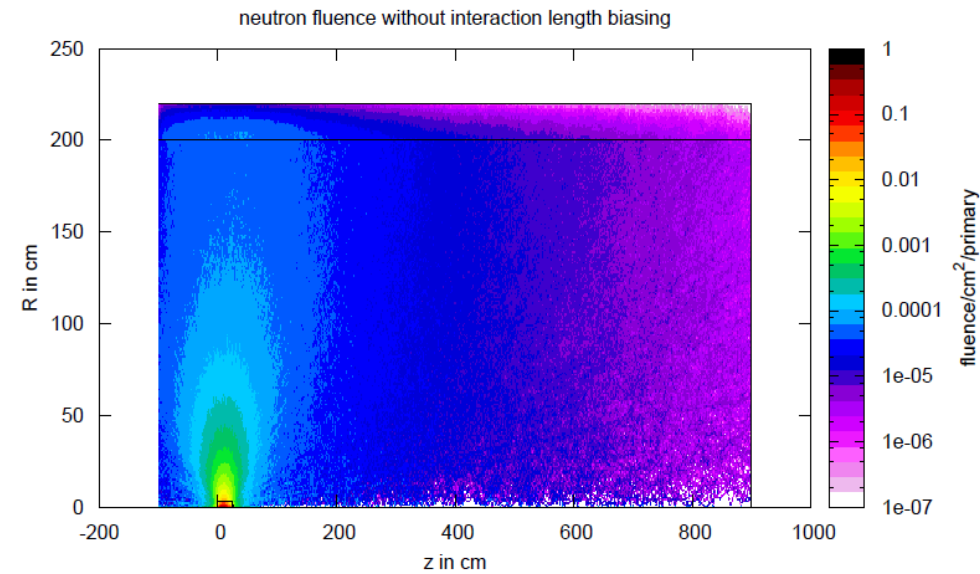
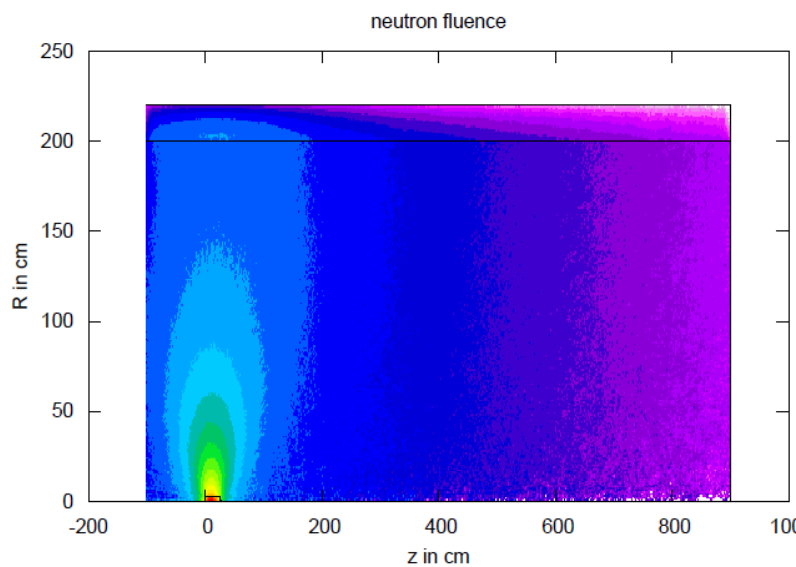
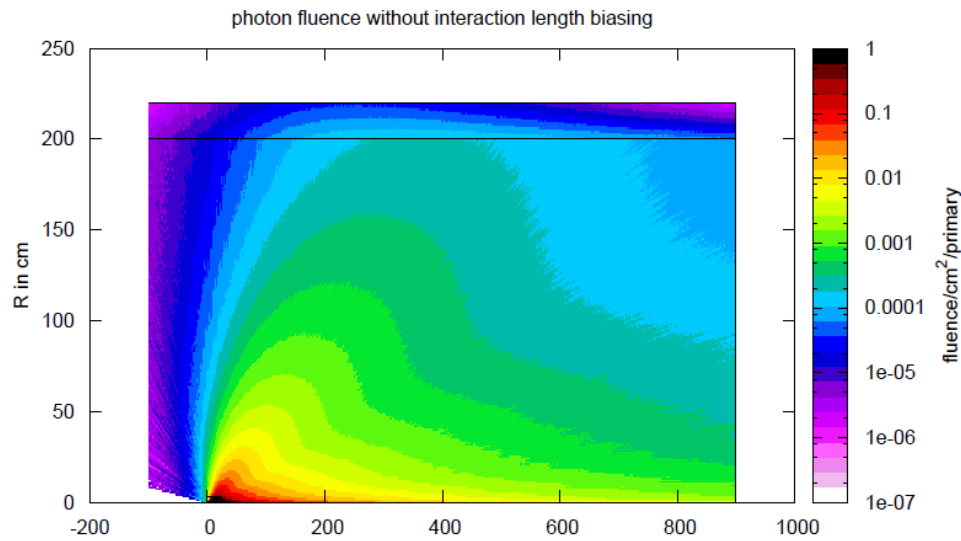
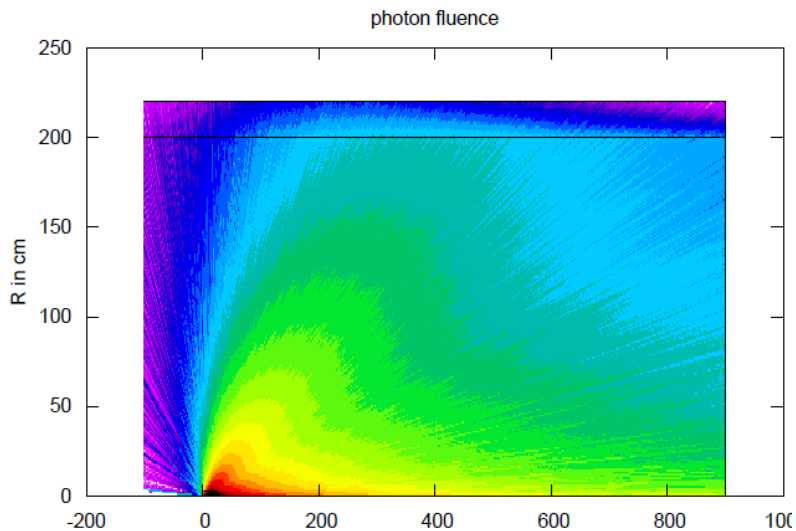
- calculate residual dose rate around the sample for different number of "replicas" (*e.g.*, 3 and 10)

# Interaction length biasing

500 primaries  
w ilb

(same total CPU time!)

6500 primaries  
w/o ilb



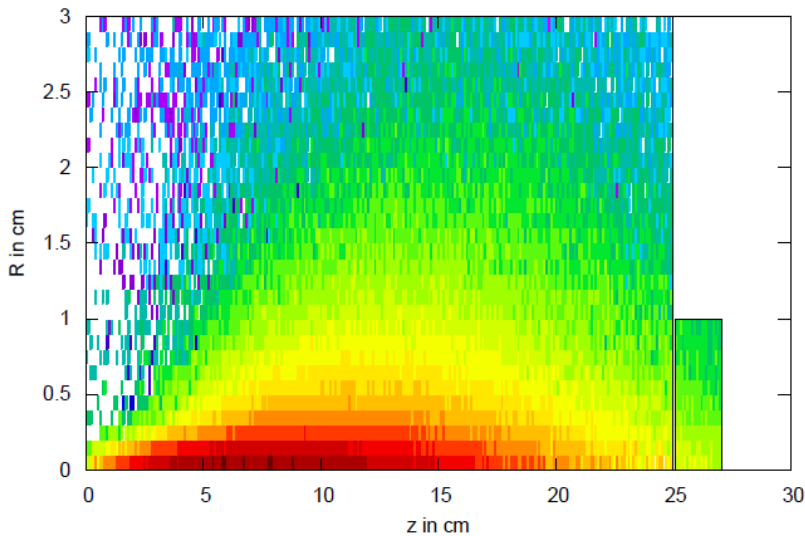
# Interaction length biasing

500 primaries  
w/ ilb

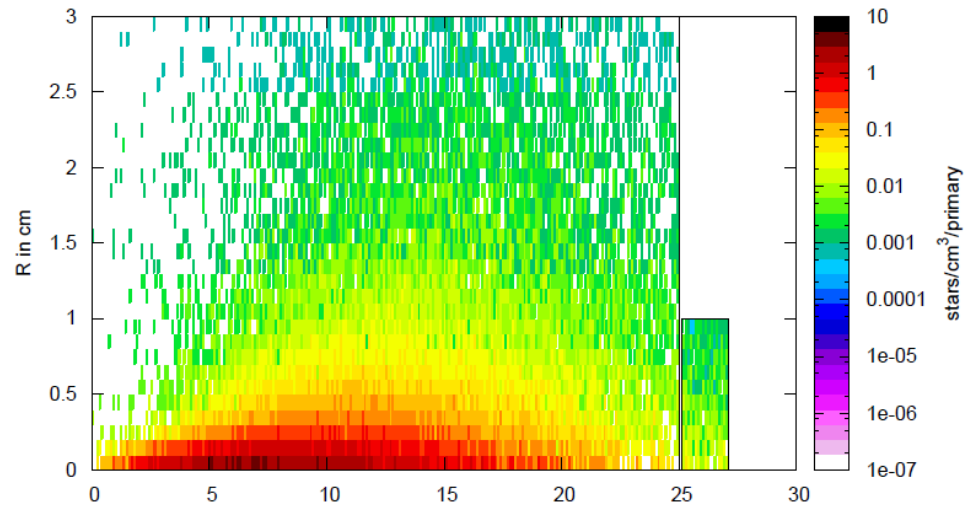
(same total CPU time!)

6500 primaries  
w/o ilb

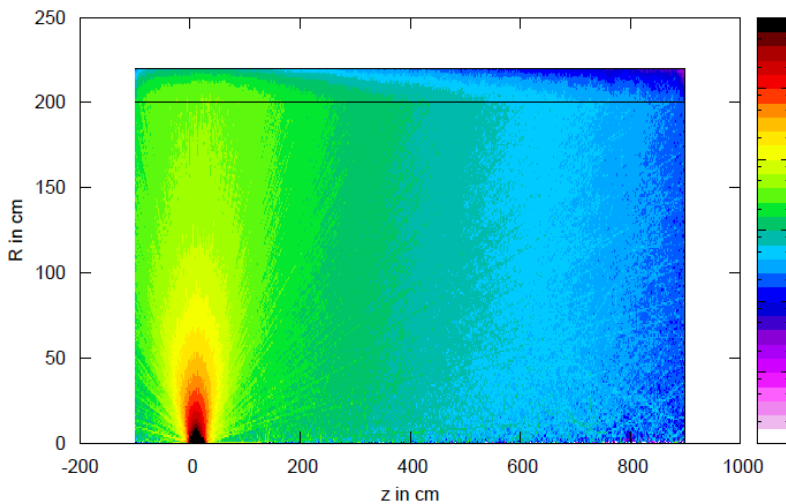
star density target and sample



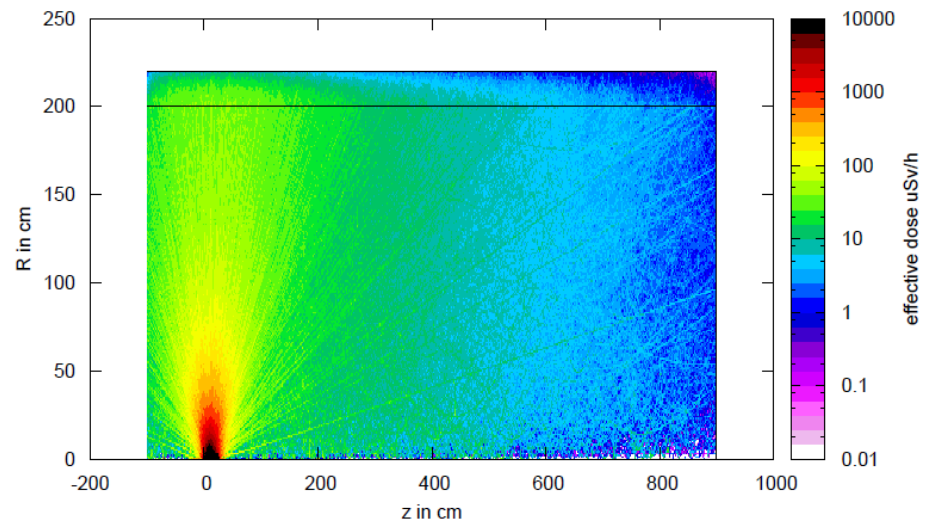
star density target and sample without interaction length biasing



effective dose 1w



effective dose 1w without interaction length biasing

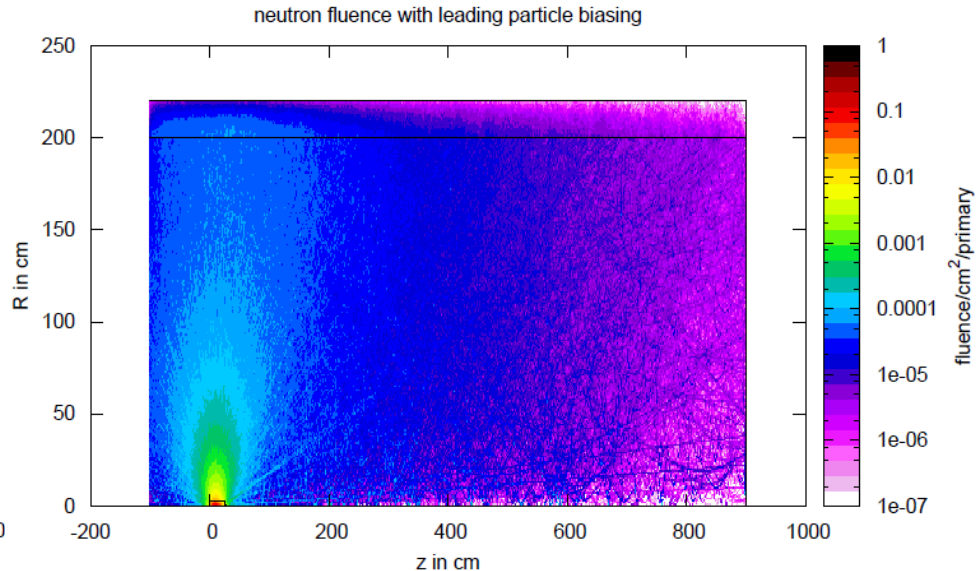
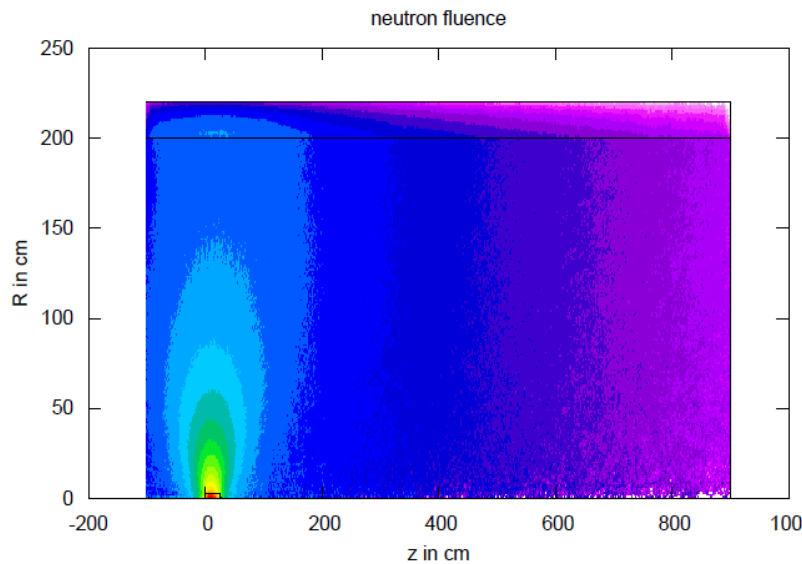
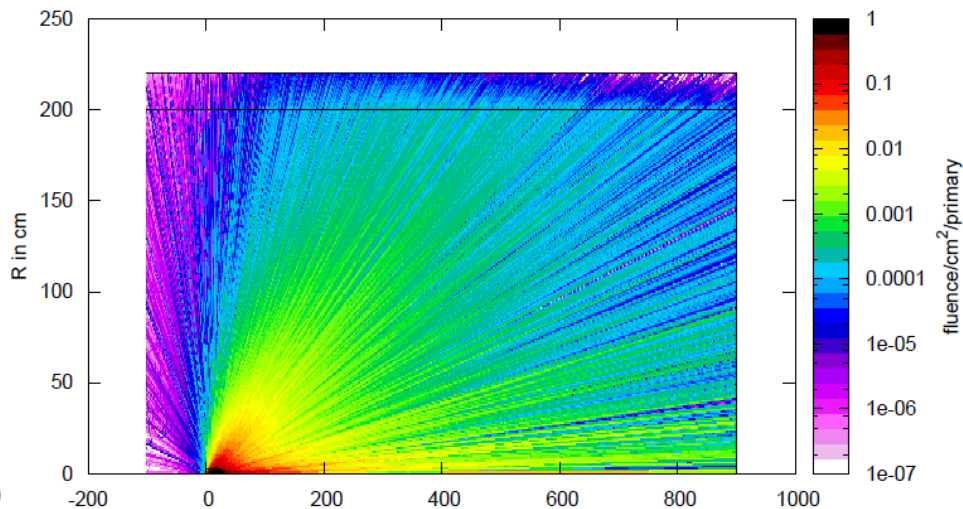
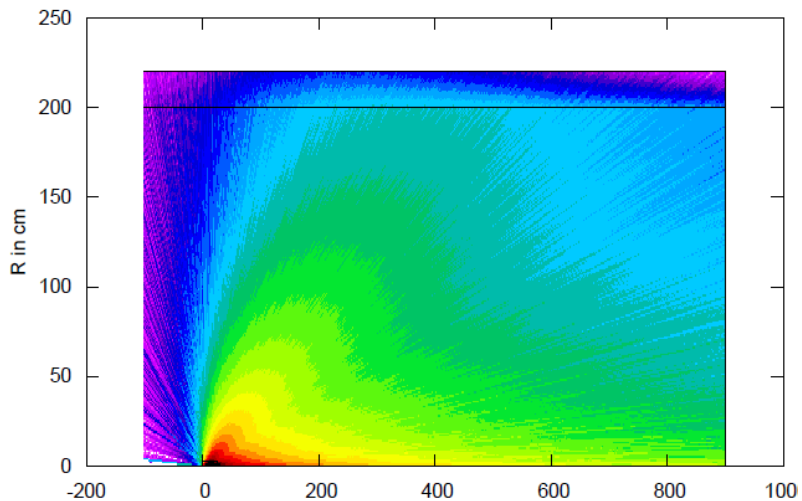


# Leading particle biasing

500 primaries  
w/o lpb  
photon fluence

(same total CPU time!)

10000 primaries  
w/ lpb  
photon fluence with leading particle biasing



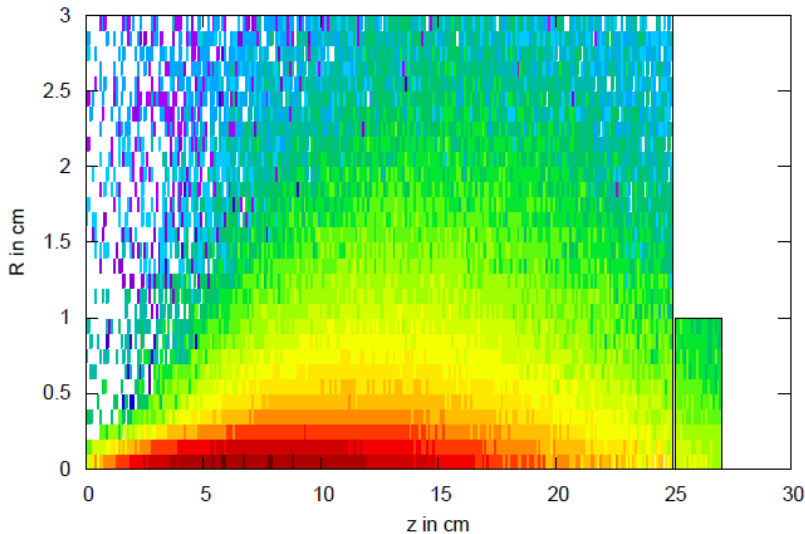
# Leading particle biasing

500 primaries  
w/o lpb

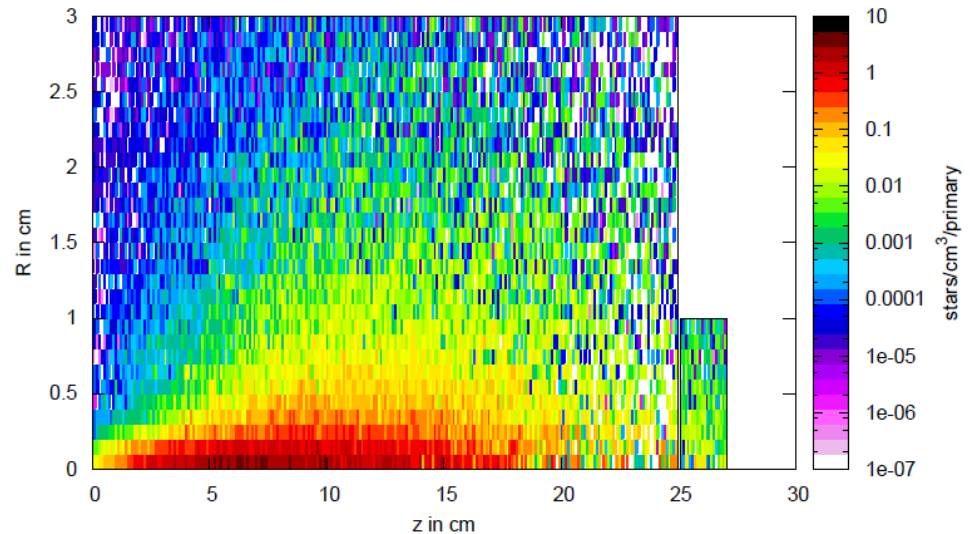
(same total CPU time!)

10000 primaries  
w/ lpb

star density target and sample



star density target and sample with leading particle biasing



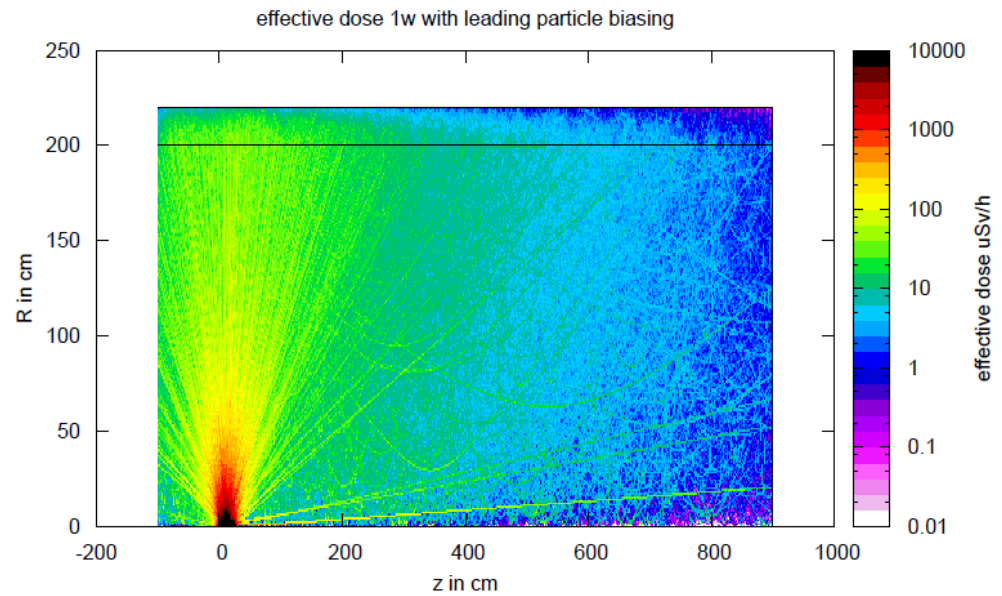
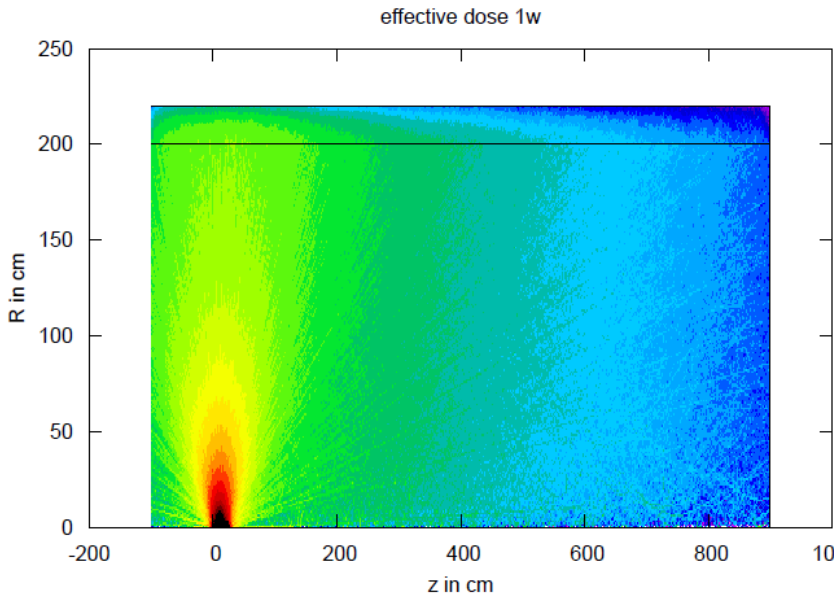
- Leading particle biasing introduces significant weight fluctuations
- Weight windows could be defined to reduce them
- In the present example, leading particle biasing introduces a bias towards high energy photonuclear interactions that have lower cross sections than GDR interactions. In addition the latter contribute significantly to the nuclide yield.
- Thus, better not use leading particle biasing in the present example...

# Leading particle biasing

500 primaries  
w/o lpb

(same total CPU time!)

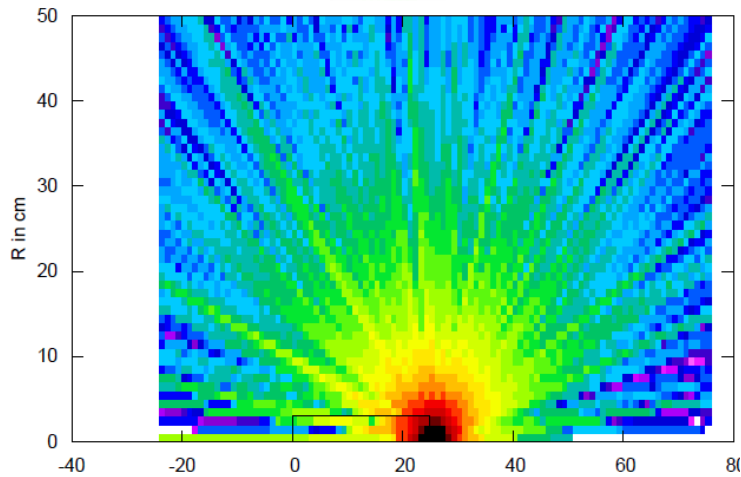
10000 primaries  
w/ lpb



# Region importance biasing

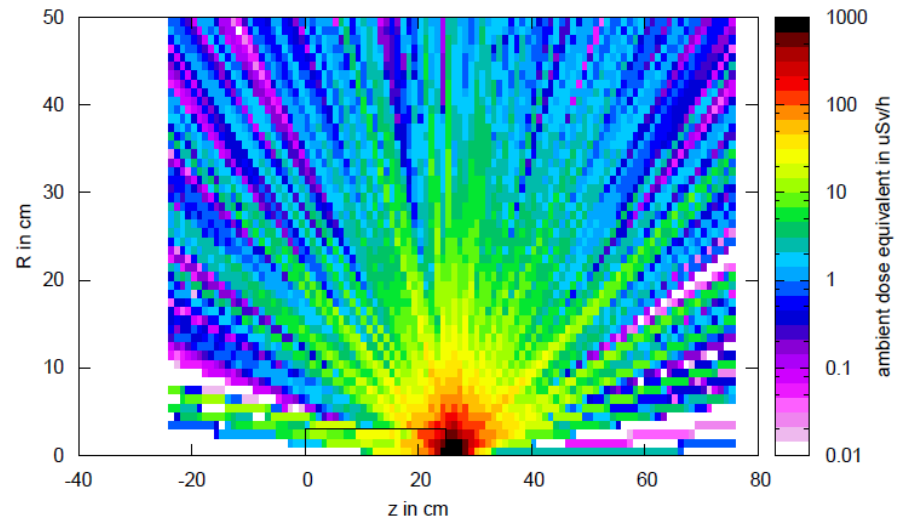
1000 primaries (same total CPU time!) 2000 primaries

With region importance biasing



Without region importance biasing

ambient dose equivalent 1w, no region importance biasing

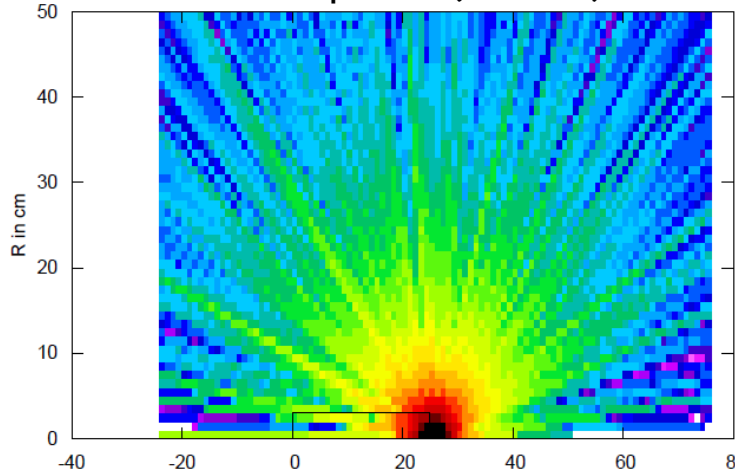


- Here: region importance in sample set equal to 5 (the maximum relative value possible) , all other regions =1
- It requires sufficient statistics in the particle fluence entering the sample.

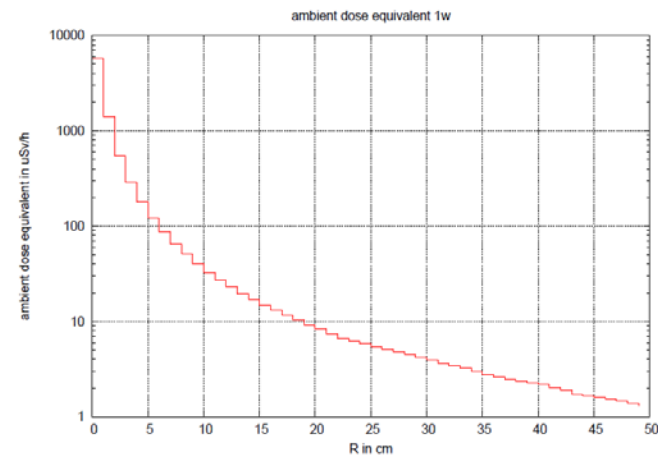
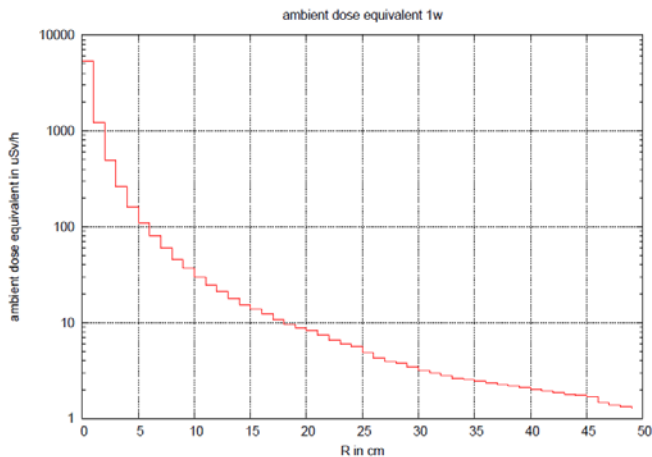
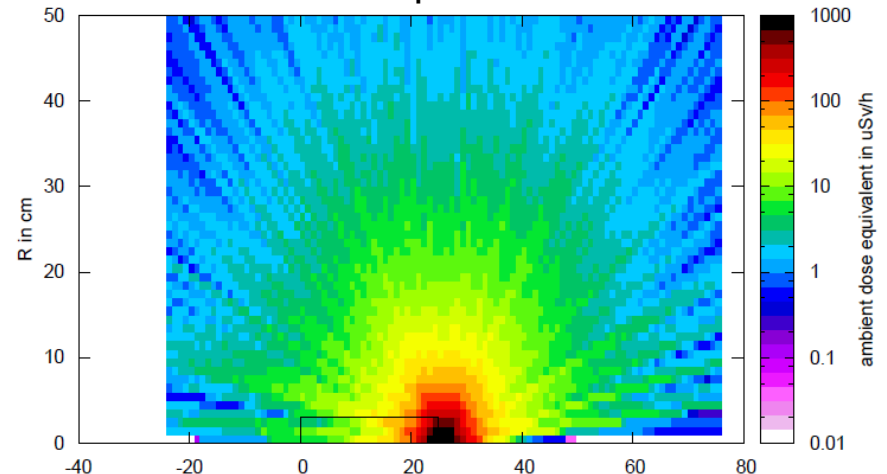


# Number of "replicas"

3 replicas (default)



10 replicas



- Increasing the number of replicas requires sufficient statistics in the production of nuclides (check, e.g., star density or activity maps!)
- In any case, don't overdo it...