

Computational methods for Medical Physics

Lecture 4: The convolution operation and its use in proton dose calculation

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LUDWIG-MAXIMILIANS-UNIVERSITÄT MÜNCHEN



- Lectures 1-3:
 - MC integration technique
 - Random sampling
 - Sampling from distributions
 - MC particle transport examples
- This Lecture:
 - Definition of Convolution
 - Convolution and its role in image processing and restoration
 - Basic properties and theorems
 - The pencil beam dose calculation algorithm in proton therapy

• We define the convolution operation f * g as:

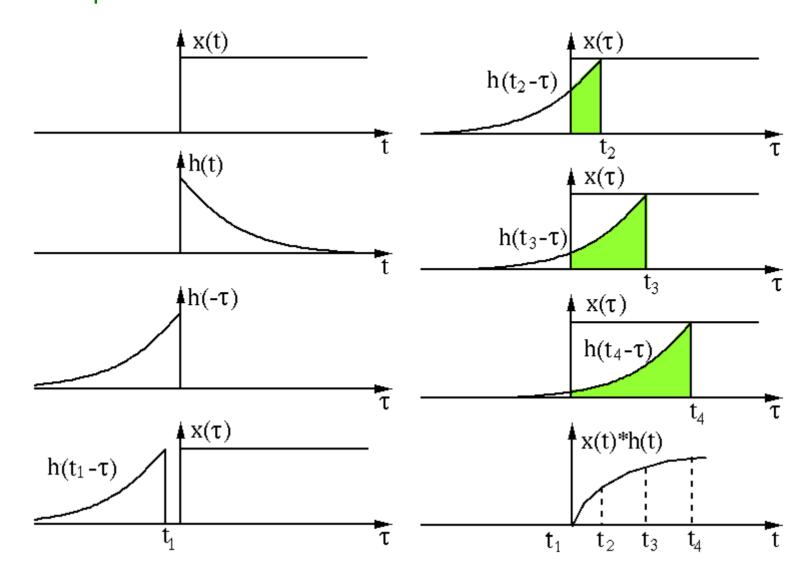
$$h(t) = (f * g)(t) = \int_{-\infty}^{\infty} f(\tau)g(t - \tau)d\tau$$

and we say that the function h is the convolution of the functions f and g.

• The operation is similarly defined in higher dimensional spaces

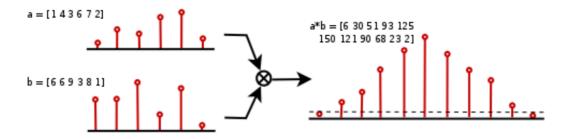
^{*}The condition of Lebesgue integrability is sufficient but not necessary

Schematic representation of convolution x*h



- Convolution can also be defined for discrete signals
- In the majority of the time this is the type of signals we deal with

$$(f * g)[n] = \sum_{m=-\infty}^{\infty} f[m] \cdot g[n-m]$$



• The length of the final vector is: length (f) + length (g) - 1

f = [1, 4, 7]

Example:

$$h[n] = (f * g)[n] = \sum_{m=-\infty}^{\infty} f[m] \cdot g[n-m]$$

$$(f * g)[0] = \sum_{m=-\infty}^{\infty} f[m] \cdot g[-m] = f[0]g[0] = 3$$

$$g = [3, 5, 2]$$

$$[ndex 0] \quad [ndex 2]$$

$$[ndex 1]$$

$$(f * g)[1] = \sum_{m=-\infty}^{\infty} f[m] \cdot g[1-m] = f[0]g[1] + f[1]g[0] = 1*5 + 4*3 = 17$$

$$(f * g)[2] = \sum_{m=-\infty}^{\infty} f[m] \cdot g[2-m] = f[0]g[2] + f[1]g[1] + f[2]g[0] = 1*2 + 4*5 + 7*3 = 43$$

$$(f * g)[3] = \sum_{m=0}^{\infty} f[m] \cdot g[3-m] = f[0]g[3] + f[1]g[2] + f[2]g[1] + f[3]g[0] = 0 + 4 \cdot 2 + 5 \cdot 7 + 0 = 43$$

$$(f * g)[4] = \sum_{m=0}^{\infty} f[m] \cdot g[4-m] = f[2]g[2] = 2*7 = 14$$

$$(f * g)[5] = \sum_{m=0}^{\infty} f[m] \cdot g[5-m] = 0$$

$$h = (f * g) = [3,17,43,43,14]$$



- The potential of a field caused by an extended body (Physics)
- The distribution of the sum of two random variables (Probability)
- Filtering (Image processing)
- Image deblurring (Image processing)
- Spectra unfolding (Radiation detection)
- Dose calculation in radiation therapy (Medical Physics)



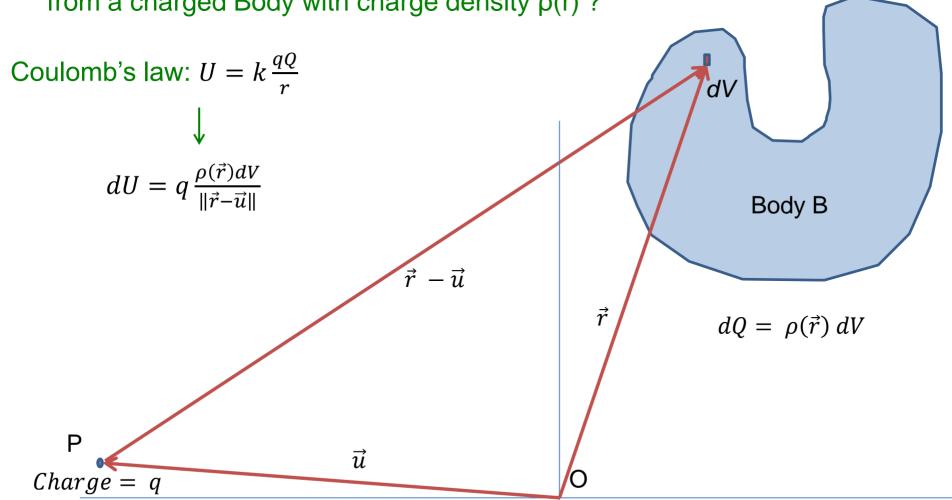
- The convolution operation appears in many fields and applications:
 - The potential of a field caused by an extended body (Physics)
 - The distribution of the sum of two random variables (Probability)
 - Filtering (Image processing)
 - Image deblurring (Image processing)
 - Spectra unfolding (Radiation detection)
 - Dose calculation in radiation therapy (Medical Physics)
- Why the convolution operation appears in so many, seemingly completely different problems?

The electrostatic problem:

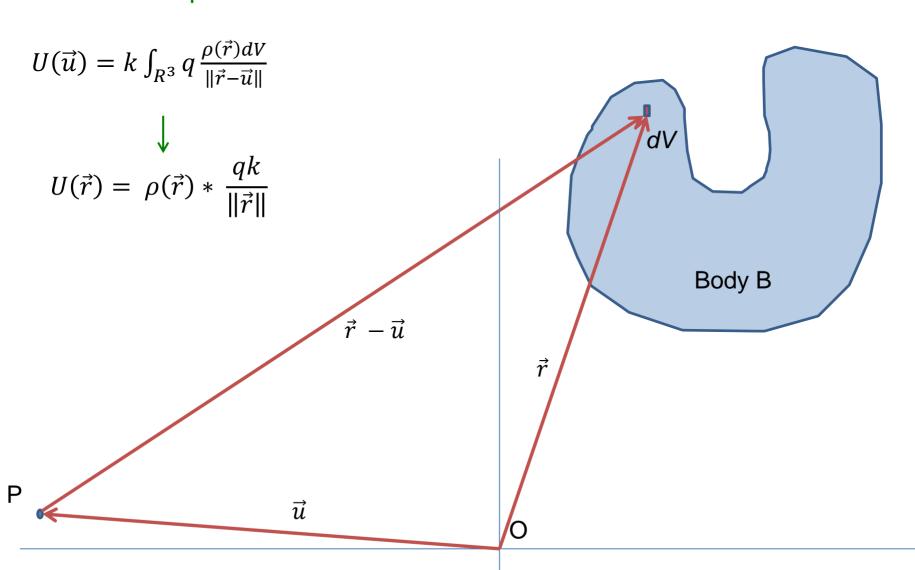
What is the electrostatic potential of experienced by a point charge P

from a charged Body with charge density $\rho(r)$?

MEDICAL PHYSICS

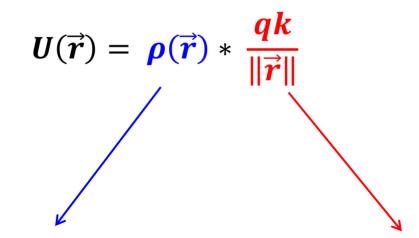


• The electrostatic problem:



The convolution operation General example

The electrostatic problem:
 What is the electrostatic potential of experienced by a point charge P from a charged Body with charge density ρ(r)?



This is an <u>arbitrary</u> charge distribution (it may be different in every problem)

This is Coulomb's law (a Universal Law)

Properties of convolution

Commutative:

$$(f * g) = (g * f)$$

Associative:

$$(f * g)*h = f * (g * h)$$

Distributive:

$$f * (g + h) = (f * g) + (f * h)$$

With delta function:

$$f * \delta = f$$

With derivation:

$$\partial_{x}(f*g) = (\partial_{x}f*g)$$

Complexity of the convolution operation:

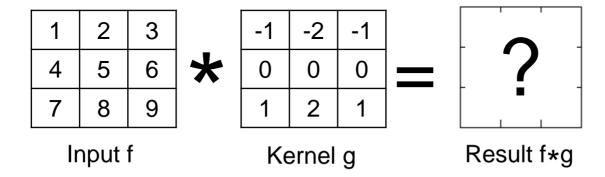
$$(f * g)[n] = \sum_{m=-\infty}^{\infty} f[m] \cdot g[n-m]$$

- The complexity of the above operation is O(n²)
- It roughly means that the CPU/memory/time/storage scales with n²

 Just as in the continuous case, convolution can also be defined for discrete signals of higher dimensions:

$$(f * g)[x, y] = \sum_{m=-\infty}^{\infty} \sum_{n=-\infty}^{\infty} f[m, n] \cdot g[x-m, y-n]$$

• Example:



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The convolution operation 2D

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X	\y 0	1	2	x\y	0	1	2
0	1	2	3	0	-1	-2	-1
xample:	4	5	6	*	0	0	0
2	7	8	9	2	1	2	1

$$\int \left(f * g \right) [x, y] = \sum_{m=-\infty}^{\infty} \sum_{n=-\infty}^{\infty} f[m, n] \cdot g[x-m, y-n]$$

Input f

Kernel g

$$(f * g)[0,0] = \sum_{m=-\infty}^{\infty} \sum_{n=-\infty}^{\infty} f[m,n] \cdot g[-m,-n] = \sum_{m=-\infty}^{\infty} (f[m,0] \cdot g[-m,0]) = f[0,0] \cdot g[0,0] = -1$$

$$(f * g)[0,1] = \sum_{m=-\infty}^{\infty} \sum_{n=-\infty}^{\infty} f[m,n] \cdot g[-m,1-n] = \sum_{m=-\infty}^{\infty} (f[m,0] \cdot g[-m,1] + f[m,1] \cdot g[-m,0])$$

$$= f[0,0] \cdot g[0,1] + f[0,1] \cdot g[0,0] = -4$$

Continue like that until we fill a matrix with dimensions $x' = x_f + x_g - 1$ and $y' = y_f + y_g - 1$

• Example:

$$(f * g)[x, y] = \sum_{m=-\infty}^{\infty} \sum_{n=-\infty}^{\infty} f[m, n] \cdot g[x-m, y-n]$$

			1				,	-1	-4	-8	-8	-3
1	2	3		-1	-2	-1		-4	-13	-20	-17	-6
4	5	6	*	0	0	0		-6	-18	-24	-18	-6
7	8	9		1	2	1		4	13	20	17	6
Ir	nput	f	Kernel g			7	22	32	26	9		

• Example (with a trick):
$$(f * g)[x, y] = \sum_{m=-\infty}^{\infty} \sum_{n=-\infty}^{\infty} f[m, n] \cdot g[x-m, y-n]$$

Take kernel g

-1	-2	-1
0	0	0
1	2	1

Kernel g

• Example (with a trick):
$$(f * g)[x, y] = \sum_{m=-\infty}^{\infty} \sum_{n=-\infty}^{\infty} f[m, n] \cdot g[x-m, y-n]$$

Take kernel g, flip it in both dimensions

1	2	1
0	0	0
-1	-2	-1

Kernel g

- Example (with a trick): $(f * g)[x, y] = \sum_{m=-\infty}^{\infty} \sum_{n=-\infty}^{\infty} f[m, n] \cdot g[x-m, y-n]$
- Take kernel g, flip it in both dimensions

1	2	1
0	0	0
-1	-2	-1

Kernel g

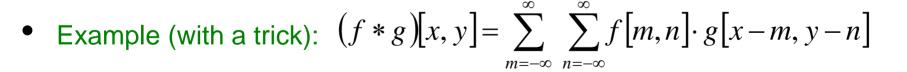
1	2	1	
0	0 1	0 2	3
-1	-2 4	-1 5	6
	7	8	9

- Example (with a trick): $(f * g)[x, y] = \sum_{n=0}^{\infty} \sum_{n=0}^{\infty} f[m, n] \cdot g[x m, y n]$ $m=-\infty$ $n=-\infty$
- Take kernel g, flip it in both dimensions

1	2	1
0	0	0
-1	-2	-1

Kernel g

	1	2	1
	0 1	0 2	0 3
	-1 4	-2 5	-1 6
ľ	7	8	9



Take kernel g, flip it in both dimensions

1	2	1
0	0	0
-1	-2	-1

Kernel g

	1	2	1		
1	0 2	0 3	0		
4	-1 5	-2 6	-1	=	-17
7	8	9		'	

- Example (with a trick): $(f * g)[x, y] = \sum_{m=-\infty}^{\infty} \sum_{n=-\infty}^{\infty} f[m, n] \cdot g[x-m, y-n]$
- Take kernel g, flip it in both dimensions

1	2	1
0	0	0
-1	-2	-1

Kernel g

1	2 1	1 2	3
0	0 4	<mark>0</mark> 5	6
-1	- <mark>2</mark> 7	-1 8	9

• Example (with a trick): $(f * g)[x, y] = \sum_{m=-\infty}^{\infty} \sum_{n=-\infty}^{\infty} f[m, n] \cdot g[x-m, y-n]$

Take kernel g, flip it in both dimensions

1	2	1
0	0	0
-1	-2	-1

Kernel g

Apply it on the input (element by element multiplication and addition)

$\begin{bmatrix} 4 & 5 & 6 \\ 7 & 8 & 9 \end{bmatrix} * \begin{bmatrix} 0 & 0 & 0 \\ 1 & 2 & 1 \end{bmatrix} = ?$	1	2	3		-1	-2	-1		
	4	5	6	*	0	0	0	_	7
	7		ı		1	2	1		Ī

Input f

Kernel g

- Example (with a trick): $(f * g)[x, y] = \sum_{m=-\infty}^{\infty} \sum_{n=-\infty}^{\infty} f[m, n] \cdot g[x-m, y-n]$
- Take kernel g, flip it in both dimensions

1	2	1
0	0	0
-1	-2	-1

Kernel g

		ī	1		ī .	·	•	-1	-4	-8	-8	-3
1	2	3		-1	-2	-1		-4	-13	-20	-17	-6
4	5	6	*	0	0	0	_	-6	-18	-24	-18	-6
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1	2	1
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Kernel g

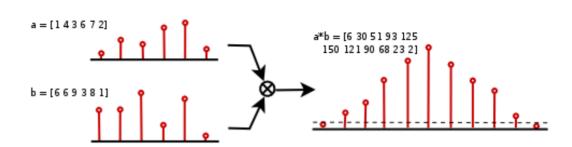
Apply it on the input (element by element multiplication and addition)

							-1	-4	-8	-8	-3	
1	2	3		-1	-2	-1		-4	-13	-20	-17	-6
4	5	6	*	0	0	0	_	-6	-18	-24	-18	-6
7	8	9		1	2	1		4	13	20	17	6
Input f Kernel g						7	22	32	26	9		

(Image filtering)

Filtering: a signal (or filter, or kernel) g is used to modify another signal f

$$(f * g)[n] = \sum_{m=-\infty}^{\infty} f[m] \cdot g[n-m]$$



Some examples from image filtering

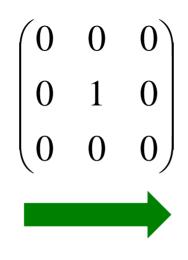




• Filtering:



Original image



What is the result of this filter?

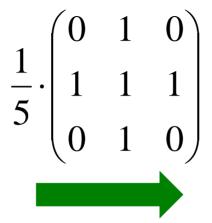


Original image





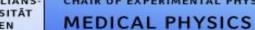
Original image



What is the result of this filter?



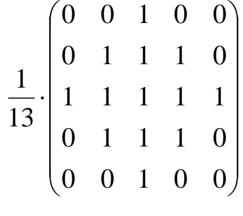
Blurring



Filtering:



Original image





What is the result of this filter?

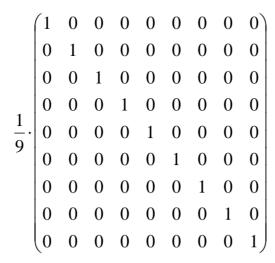


(More) Blurring

Filtering:



Original image





What is the result of this filter?



Motion Blurring

Sobel operators or Sobel filters

$$Gx = \begin{pmatrix} -1 & 0 & 1 \\ -2 & 0 & 2 \\ -1 & 0 & 1 \end{pmatrix} \qquad Gy = \begin{pmatrix} 1 & 2 & 1 \\ 0 & 0 & 0 \\ -1 & -2 & -1 \end{pmatrix} \qquad G = \sqrt{Gx^2 + Gy^2}$$

$$Gy = \begin{pmatrix} 1 & 2 & 1 \\ 0 & 0 & 0 \\ -1 & -2 & -1 \end{pmatrix}$$

$$G = \sqrt{Gx^2 + Gy^2}$$







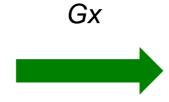
Sobel operators or Sobel filters

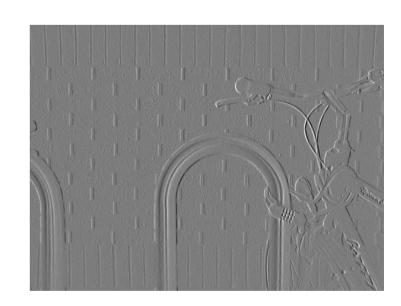
$$Gx = \begin{pmatrix} -1 & 0 & 1 \\ -2 & 0 & 2 \\ -1 & 0 & 1 \end{pmatrix} \qquad Gy = \begin{pmatrix} 1 & 2 & 1 \\ 0 & 0 & 0 \\ -1 & -2 & -1 \end{pmatrix} \qquad G = \sqrt{Gx^2 + Gy^2}$$

$$Gy = \begin{pmatrix} 1 & 2 & 1 \\ 0 & 0 & 0 \\ -1 & -2 & -1 \end{pmatrix}$$

$$G = \sqrt{Gx^2 + Gy^2}$$







Sobel operators or Sobel filters

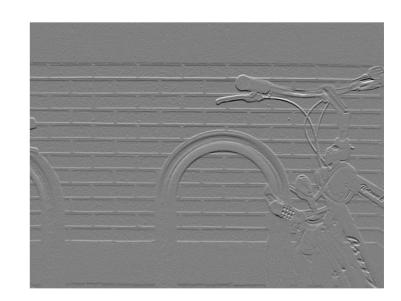
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$$Gy = \begin{pmatrix} 1 & 2 & 1 \\ 0 & 0 & 0 \\ -1 & -2 & -1 \end{pmatrix}$$

$$G = \sqrt{Gx^2 + Gy^2}$$







Convolution based photon dose calculation

- Treatment planning of external beam radiation therapy changed drastically in the last two decades of the 20th century
 - Computed tomography (CT) scanners began being actively used in clinics world wide (3D anatomy)
 - Computer manufacturers begun providing solutions that were attractive to the clinical environment (increased computational power)
- Physicists realized that the combination of CT and computer technology there could improve the accuracy of the dose calculations in clinical practice
- In the last 30 years several dose calculation algorithms were proposed in order to achieve the goal of "accurate" treatment planning

Convolution based photon dose calculation

- The 90's and early 00's was the Pencil Beam (PB) algorithm was developed
- Of course there is also Monte Carlo. Highly accurate but too slow to be used
- The idea of the convolution methods:
 - Instead of doing a full Monte Carlo dose calculation
 - Precalculate via Monte Carlo the relevant physical information common to all particular cases
- The problem of dose calculation uses two types of info:
 - "Universal" physical information (dose deposition patterns)
 - Specific physical information (fluences, materials, geometries)

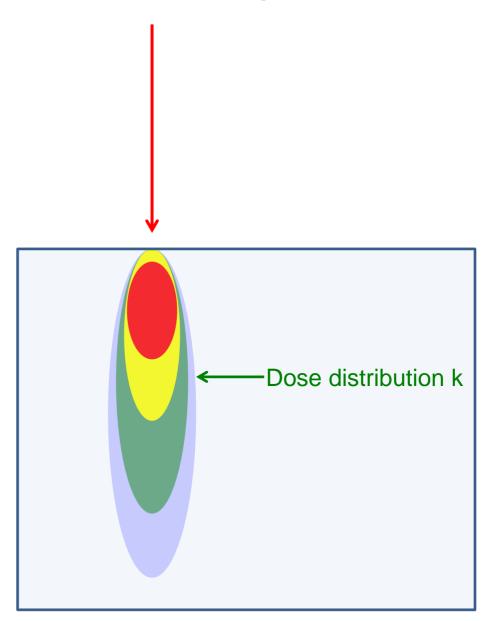
Convolution based photon dose calculation

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Convolution based photon dose calculation

- In the 80's the center of development was what is now called the Convolution/superposition method
- The 90's and early 00's was the time of the Pencil Beam Algorithm
- Of course there is also Monte Carlo. Highly accurate but too slow to be used
- The idea of the convolution methods:
 - Instead of doing a full Monte Carlo dose calculation
 - Precalculate via Monte Carlo the relevant physical information common to all particular cases
- The problem of dose calculation uses two types of info:
 - "Universal" physical information (dose deposition patterns)
 - Specific physical information (fluences, materials, geometries)

Consider the case of an infinitesimal pencil beam on a water target



- A bit more realistic:
 - An extended beam
 - The dose contribution in point P
 - If each PB has a different intensity/fluence

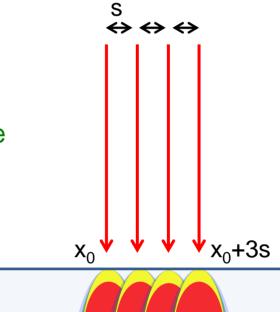
$$d(x,z) = \sum_{i=0}^{3} \phi(i)k(x-i\cdot s, z)$$

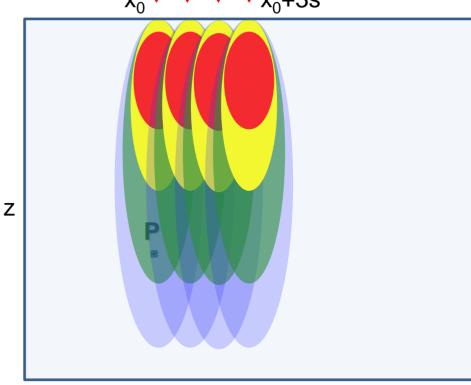
Then the dose in position P is:

$$d(x,z) = \phi(x) * k(x,z)$$

• Similarly to:

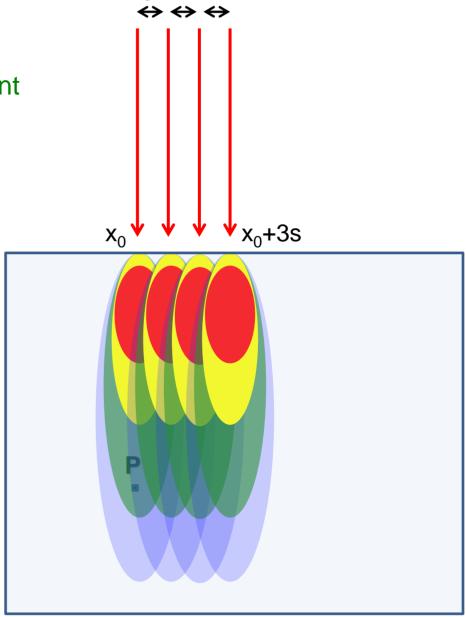
$$h[n] = (f * g)[n] = \sum_{m=-\infty}^{\infty} f[m] \cdot g[n-m]$$



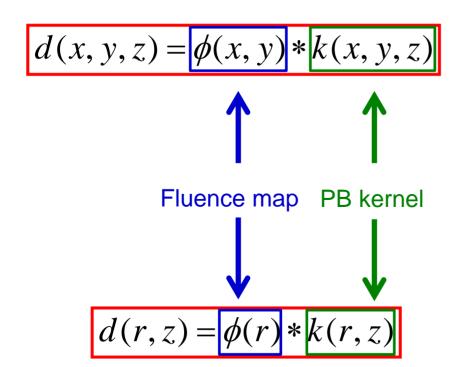


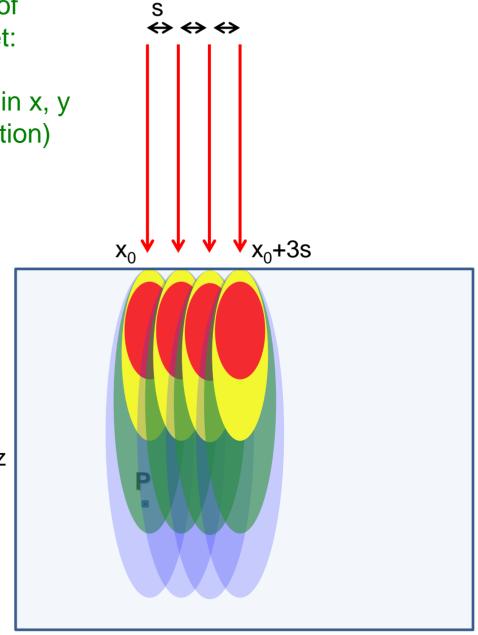
- It shouldn't be surprising that convolution has appeared:
 - 1. The pencil beam kernel is shift invariant
 - 2. The absorbed dose obeys linearity

- Convolution is a superposition operation with a constant shift invariant kernel
- The kernel is a constant (for the same energy the dose kernel remains the same)
- The intensity/fluence of each PB is the specific information (could be different in every case)



- For an extended photon beam composed of infinitesimal PBs on a homogeneous target:
 - Assuming that the PBs are symmetric in x, y (plane perpendicular to the beam direction)



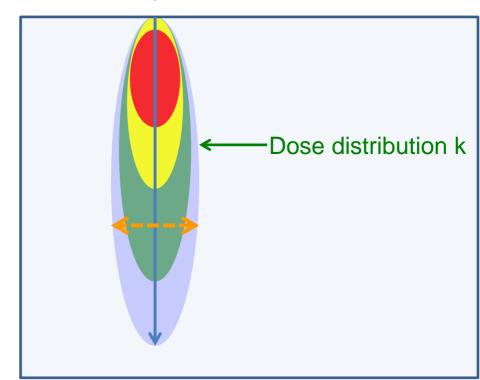


- Description of the dose kernel k:
 - The dose kernel is precalculated with MC and consists of dose in water from PBs of different E (basic input data to the TPS)

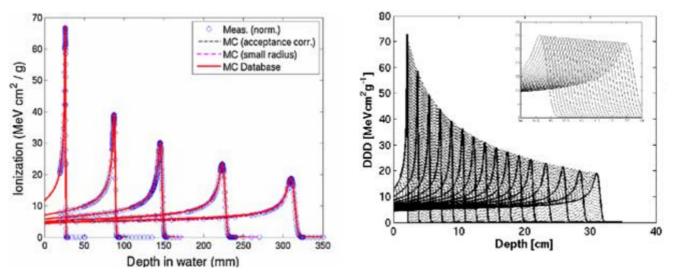
 Based on those data, the TPS splits the dose kernel into two terms: a central-axis term C and

an off-axis term O, so that:

$$k(x, y, z) = C(z) \cdot O(x, y, z)$$



The central-axis term is taken from the laterally integrated PB input data



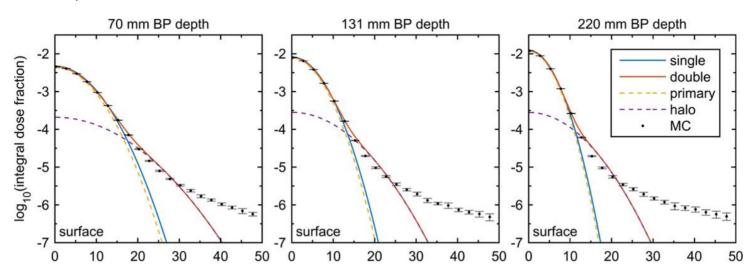
Parodi et al, Phys Med Biol, Volume 57, Number 12

 Scaling of the Bragg peaks as calculated into water for the cases of different materials is based on water equivalent path length

 The off-axis term was initially parameterized as a Gaussian shape of the lateral profile as a function of depth, for each PB

$$O(x, y, z) = \frac{1}{2\pi [\sigma_{tot}(z)]^2} \exp\left(-\frac{x^2 + y^2}{2[\sigma_{tot}(z)]^2}\right)$$

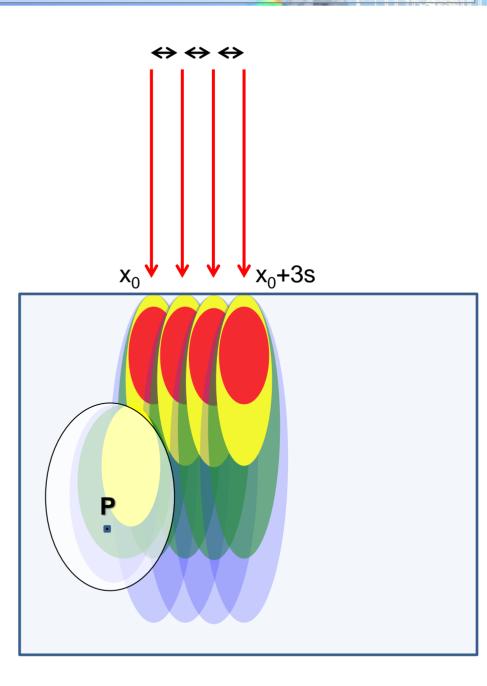
- Where σ_{tot} is the quadratic sum of all contributions broadening the beam
- The single Gaussian parameterization is not adequate.
 Therefore, double Gaussian is used:



- A patient is clearly not a water tank
- What happens with heterogeneites?

$$d(x, y, z) = \phi(x, y) * k(x, y, z)$$
Fluence map PB kernel
$$d(r, z) = \phi(r) * k(r, z)$$

 The kernel is not shift invariant anymore (takes a position dependent shape)



- What happens in presence of heterogeneities?:
- Tabulate the dose kernel k(r,z) in water (as for the homogeneous case)
- using CT data, apply a correction factor A(z_{rad}) (1D)

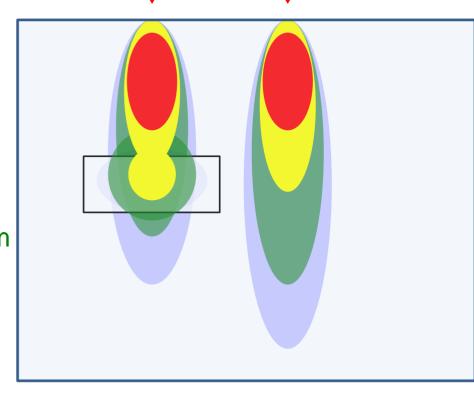
$$k'(x, y, z) = A(z_{rad})k(x, y, z)$$

• or $A(x,y,z_{rad})$ (3D)

$$k'(x, y, z) = A(x, y, z_{rad})k(x, y, z)$$

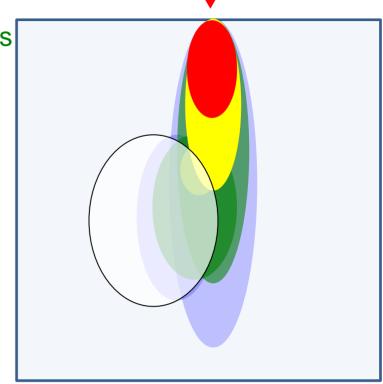
- This function is not shift invariant:
- Convolution does not describe the problem
- It becomes a more general superposition

$$d(x, y, z) = \sum_{i=0}^{N} \phi_i k_i'(x, y, z)$$



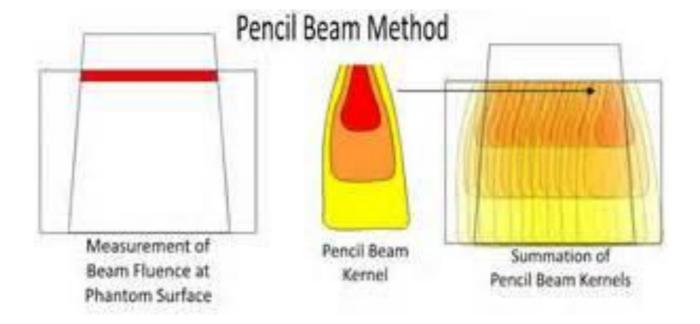


- The golden years of the pencil beam algorithm: 1995 2005
- The PB algorithm for photons appeared in commercial implementations of several manufacturers
- The main reason of the popularity of PB algorithms was the need for a fast dose calculation for IMRT
- Nevertheless, the algorithm fails to give a good description in the presence of strong heterogeneities
- Especially 1D scaling extremely fast but not accurate enough



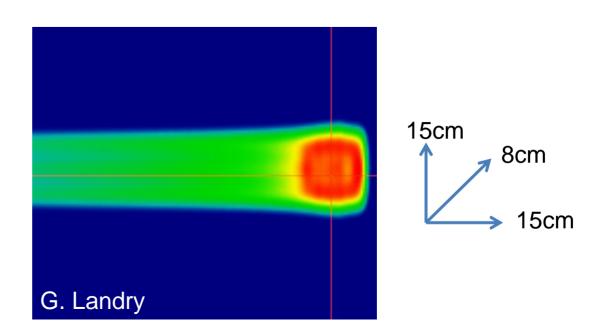
A summarizing sketch of the PB algorithm:

•





- Programming Exercise 5:
 - Calculate the dose distribution using the PB algorithm according to the basic data (dose kernel) available on the lecture's website
 - Assume homogeneous water geometry (15cmx15cmx8cm – voxel size 1mmx1mmx1mm) and a single Gaussian lateral spread parameterization
 - For PB energies in between the nominal ones, use linear interpolation for both the laterally integrated dose and the lateral spread σ
 - The dose should look like:



- Programming Exercise 5:
 - Report on the calculation time and plot a longitudinal and a lateral dose profile

