

WE-B-AUD C CE-Therapy
July 30, 2008

Dose calculation algorithms in 3DCRT and IMRT

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Sweden

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Honoring the past
Celebrating the present
Preparing for the future
Houston, Texas • July 27 - 31, 2008



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Objectives

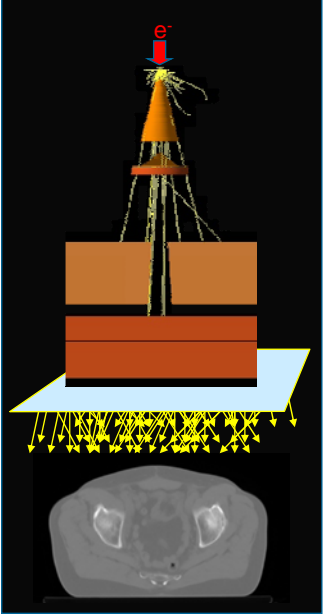
1. To provide an educational review of the physics and techniques behind convolution algorithms
2. To review the methods used to improve the simulation efficiency i.e. pencil beam and collapsed cone convolutions
3. To briefly review the performance of codes currently used for clinical treatment planning.
4. To discuss the issues associated with experimental verification of dose calculation algorithms.
5. To briefly review the potential clinical implications of accurate calculated dose distributions.

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The Problem

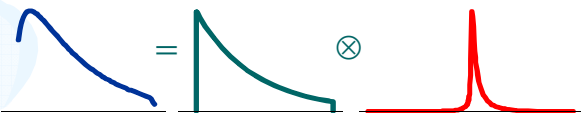
- Modelling the linac
 - Energy fluence
 - Source models
 - Monte Carlo
- Modelling of dose in patients
 - Interpolation and correction of measured data
 - Fluence to dose modelling
 - Monte Carlo



3

1

Fluence to dose Convolution



[1D convolution]

$$D(x) = \int T(x') \cdot K(x-x') dx'$$

$$D(\mathbf{r}) = \tilde{T}(\mathbf{r}) \otimes K(\mathbf{r})$$

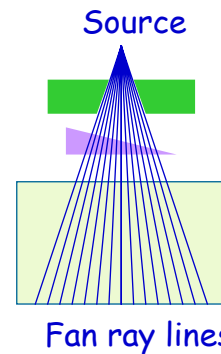
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This idea was explored by several papers at the ICCR 1984

Modelling primary photon energy fluence and loss

1

- Ray-tracing Total Energy Released in Mass (TERMA)
- Similar to determining effective or radiological depth



$$T(E, z) = \frac{\mu_E}{\rho} \cdot \Phi(E, z) \cdot E = \frac{\mu_E}{\rho} \Phi_0(E, 0) \cdot e^{-\mu_E \cdot z_{eq}} \cdot E$$

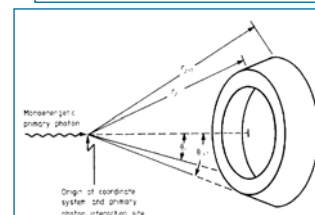
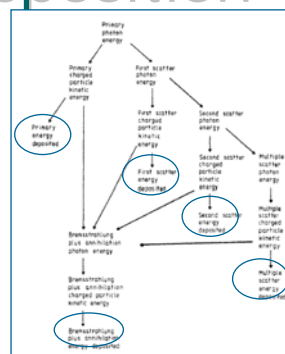
$$z_{eq} = \frac{1}{\rho_{water}} \int_0^z \rho(z') \cdot dz'$$

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Modelling dose deposition

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- Dose distribution around a single interaction point – **Point dose kernel**
- Separate primary, 1st scatter, 2nd scatter, multiple residual scatter dose kernels

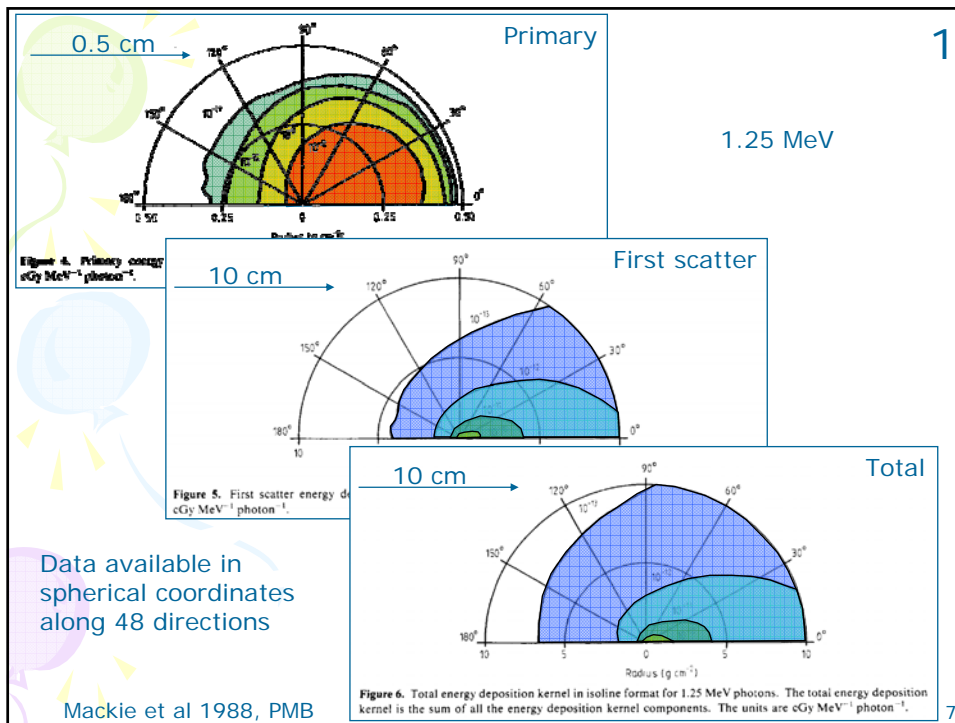


Phys. Med. Biol., 1988, Vol. 33, No 1, 1-20. Printed in the UK

Generation of photon energy deposition kernels using the EGS Monte Carlo code

T R Mackie¹], A F Bielajew², D W O Rogers³ and J J Battista⁴

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Convolve!

- Apply the dose kernel to each TERMA point
- Integrate over the whole volume i.e. a convolution

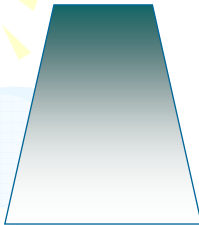
[1D convolution]

$$D(x) = \int T(x') \cdot K(x-x') dx'$$

$$D(\mathbf{r}) = T(\mathbf{r}) \otimes K(\mathbf{r})$$

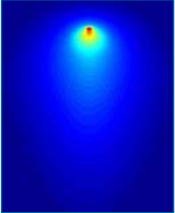
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Convolution in 2D



TERMA

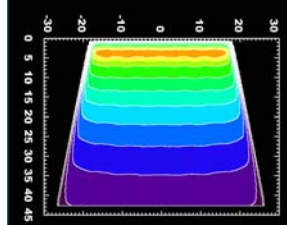
\otimes



Dose
Deposition
Kernel

[CMS]

$=$



Absorbed Dose

Convolution is efficiently solved by Fast Fourier Transform techniques

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Example: Point kernel convolution - CMS

- Re-sampling of Mackie's kernels to Cartesian coordinates
- FFT solution
- Two separate calculations:
 - A **primary kernel** for which the calculation is performed at high-resolution but over a small region – **high gradient – short range**
 - A **scatter kernel**, where the calculation is performed at a lower resolution but over a larger area – **low gradient – long range**
 - Time saving of about 65 % by this technique

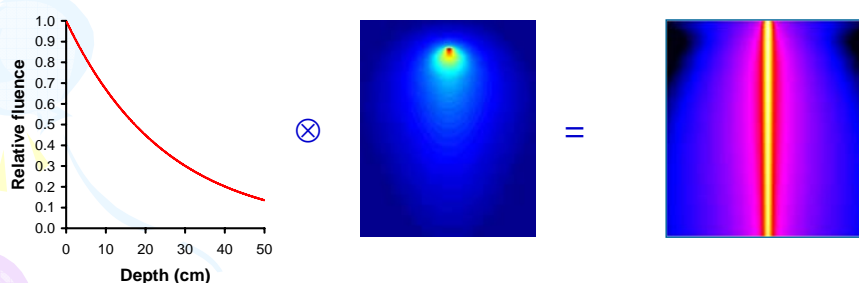
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Limitations of convolution

- Kernels are not invariant in space
 - Energy distribution varies with position in beam
 - Beam softening laterally
 - Beam hardening longitudinally
- Kernels vary with density
- Divergence leading to tilted kernels
- Pre-calculated kernels won't make it!!!
- FFT not suitable – analytical methods must used – time consuming
- Approximate methods required

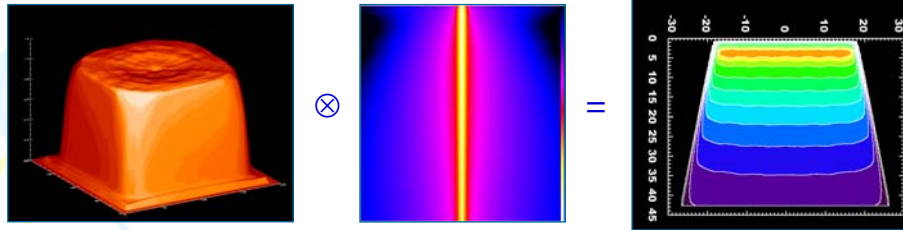
1st approximation Pencil Beam

- Reduce the dimensionality of the problem by pre-convolving in the depth dimension



- => Pencil beams (PB)
- Superposition of pencil beams in 2D => Faster

Illustration of Pencil Beam superpositioning (convolution) 2



Energy fluence

⊗

Dose
Deposition
Kernel

=

Absorbed Dose

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Construction of Pencil Kernels

Extraction of pencil beam kernels by the deconvolution method

Chen-Shou Chui and Radhe Mohan
 Department of Medical Physics, Memorial Sloan-Kettering Cancer Center, New York, New York 10021
 (Received 15 June 1987; accepted for publication 14 December 1987)

De-convolved from
measurements



Experimental determination of the dose kernel in high-energy x-ray beams

From measurements
by differentiating

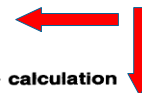


Crister P. Coberg and Bengt E. Bjärngård
 Department of Radiation Oncology, Roger Williams Medical Center, Brown University, Providence,
 Rhode Island 02908-4735
 Timothy C. Zhu
 Department of Radiation Oncology, Shands Cancer Center, University of Florida, Gainesville,
 Florida 032610-0385
 (Received 26 June 1995; accepted for publication 12 December 1995)

Use of fast Fourier transforms in calculating dose distributions for irregularly shaped fields for three-dimensional treatment planning^{a)}

Radhe Mohan and Chen-Shou Chui
 Department of Medical Physics, Memorial Sloan-Kettering Cancer Center, New York, New York 10021
 (Received 30 June 1986; accepted for publication 22 October 1986)

Calculated by
Monte Carlo
methods



A pencil beam model for photon dose calculation

Anders Ahnesjö
 Department of Radiation Physics, Karolinska Institute, Stockholm, Sweden and Helax AB, Box 1704,
 S-751 47 Uppsala, Sweden
 Mikael Saxner and Avo Trepp
 Helax AB, Box 1704, S-751 47 Uppsala, Sweden
 (Received 29 October 1990; accepted for publication 1 July 1991)

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Example: Pencil beam model – Nucletron

2

- Pencil beams based on MC calculated point kernels, integrated and fitted to a limited number of depth doses
- Separates “primary” and “scatter” dose
- Heterogeneities handled via effective path length – **only longitudinal** scaling
- Extensive beam modelling

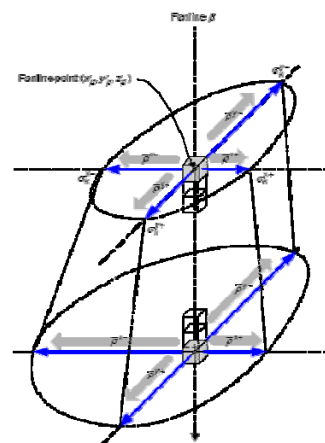
Nucletron (former MDS Nordion, Helax-TMS)

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Example: Pencil beams model - Eclipse

2

- Uses pencil beams extracted from measurements (SPB) or from Monte Carlo calculation (AAA)
- Heterogeneities handled via effective path length – longitudinal
- AAA adds a scaling of the spread of the pencil based on density – **lateral**
- AAA also have an extensive beam modelling

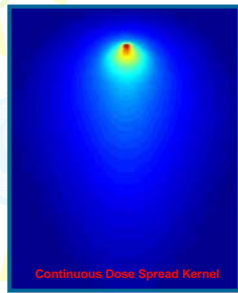


Analytical Anisotropic Algorithm

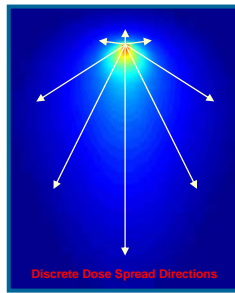
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2nd approximation Collapsed cone convolution

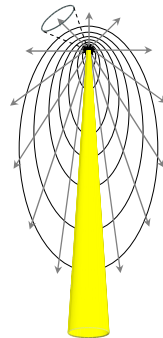
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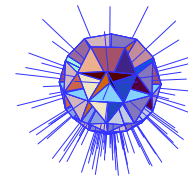
Continuous Dose Spread Kernel



Discrete Dose Spread Directions



Quantization in cones



Collapsed Cones

Kernels are discretised

Collapsed cone convolution of radiant energy for photon dose calculation
in heterogeneous media

Anders Ahnesjö¹
Department of Radiation Physics, Karolinska Institute and University of Stockholm, Box 40211, S-141 86
Stockholm, Sweden
(Received 15 August 1988; accepted for publication 3 May 1989)

Collapsing removes the inverse
square law – only exponential
attenuation is left

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Number of collapsed cones or directions

2

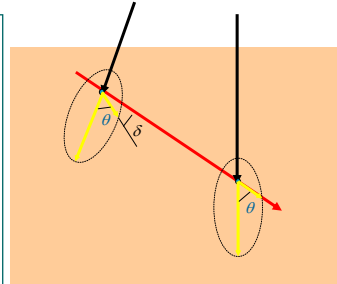
- Sufficient density of cones to distribute energy to all voxels
 - Not possible but at least while the energy is significant
 - ~100 (Mackie et al, 1996 Summer school)
 - Voxels will be missed at large distances
 - very low energy contribution
- 128 CC are used in CMS (48 for the fast version)
- 106 CC are standard in OMP

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Implementation issues

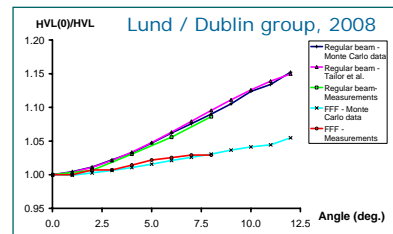
Accounts for

- **Heterogeneities**
Kernels scaled for different tissues
- **Lateral energy transport**
- **Beam Hardening and Off-axis spectrum softening**
Included in Ray Trace process
- **Tilt of kernels**
Included in Transport



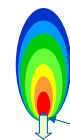
Polyenergetic Spectrum accounted for by weighted sum of monoenergetic kernels calculated by Monte Carlo

Weights determined by comparison with measured data



Examples: Collapsed cone

- Pinnacle
 - Polyenergetic weighted kernels, total energy
 - Off-axis/tilting considered during TERMA
 - Collecting dose or dose point of view
- CMS
 - Two kernels, Primary electron dose and scattered photon dose
 - No Off-axis/tilting
 - Collecting dose or dose point of view
- Nucletron
 - Two Kernels are used:
 - One for Collision Kerma into Primary Dose
 - One for 'Scerma' into Phantom Scatter Dose
 - Kernels parameterised and fitting parameters stored for run time
 - Off-axis/tilting



These are 'iso-scatter' lines.

They link points producing equal scatter to here.



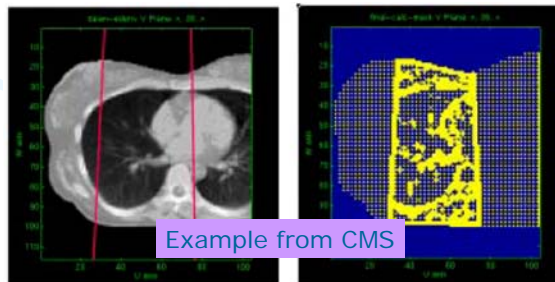
Primary interaction point

These are 'isodose' lines

From Deshpande, Philips

Further approximation

- Multigrid solution (CMS)
 - Only calculate dose using superposition at points where it is necessary, and at all other points use interpolation to get a reasonable estimate of dose
- Adaptive CCC (Pinnacle)
 - Only performs convolution at every 4th point in the TERMA array
 - Gradient search performed on TERMA array
 - Dose in between is interpolated if gradient low (i.e. TERMA doesn't change much)
 - Convolution performed at every point if TERMA gradient high



Example from CMS

Conclusions

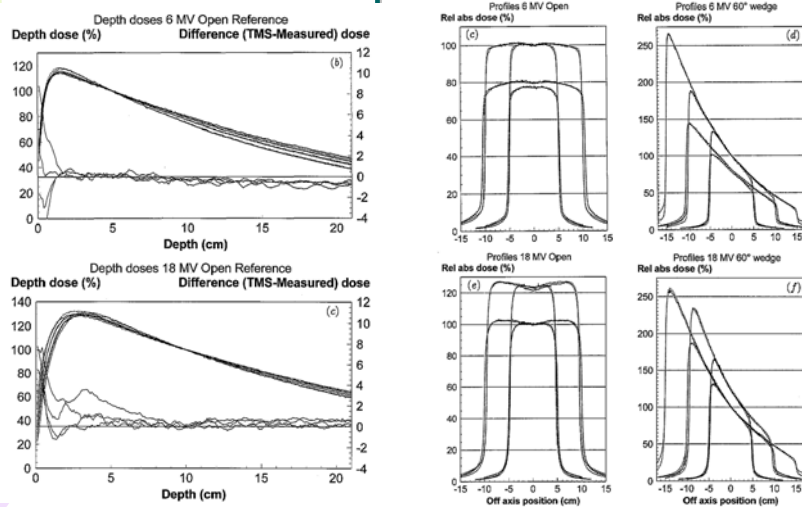
- Inhomogeneities are handled by scaling the kernels rectilinearly with electron density according to the theorem by O'Connor 1957
- **Type a** – Models primarily based on EPL scaling for inhomogeneity corrections.
 - Eclipse/SPB, OMP/PB, PPLAN, XiO/Convolution
 - **LONGITUDINAL** scaling
- **Type b** – Models that in an approximate way consider changes in lateral electron transport
 - Pinnacle/CC, Eclipse/AAA, OMP/CC, XiO/Superpositioning.
 - **LONGITUDINAL** and **LATERAL** scaling

Performance of convolution models

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Comparison in homogeneous water phantoms

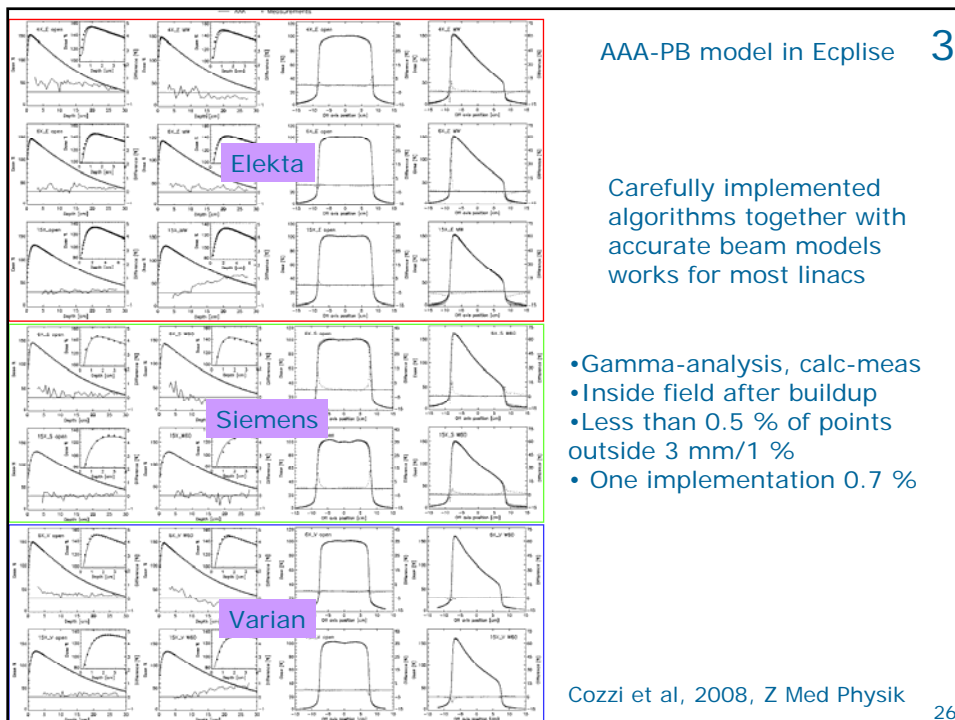
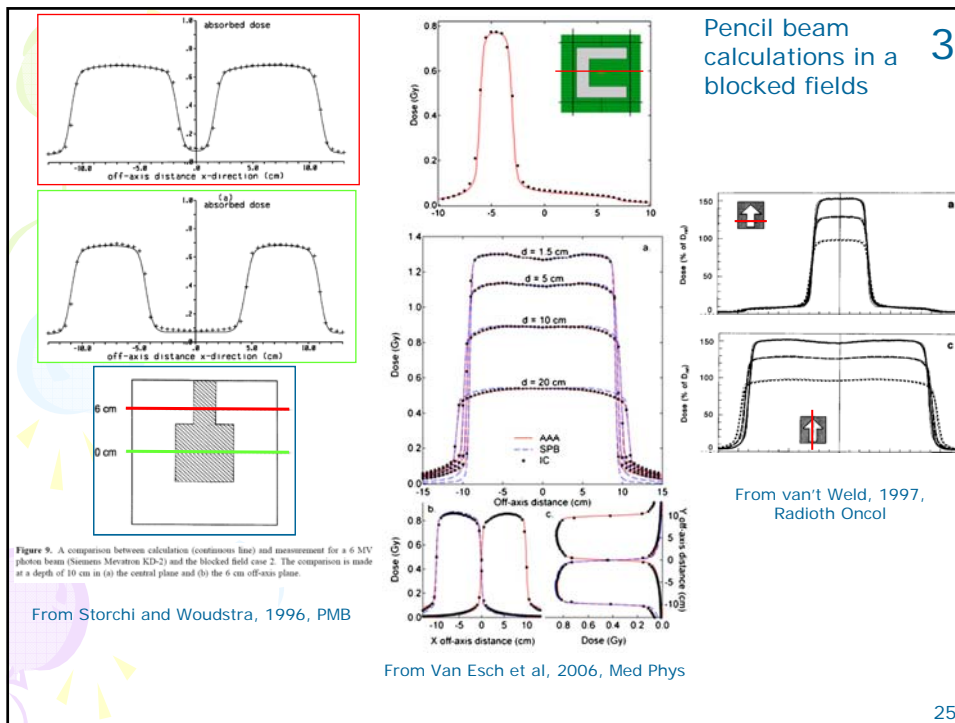
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All systems are expected to work excellent in homogenous water

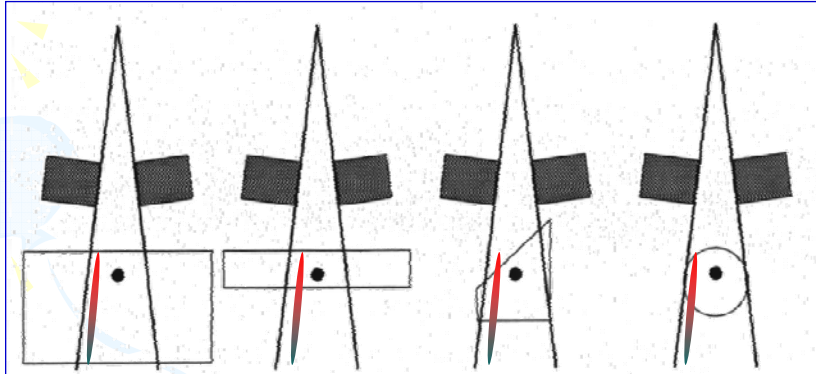
Knöös et al, 1994, PMB

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A problem using pencil beams Irregular geometries

2



The same dose to ● in all geometries since the PB is pre-integrated to a certain depth/length

See also Hurkmans et al, 1986, RO

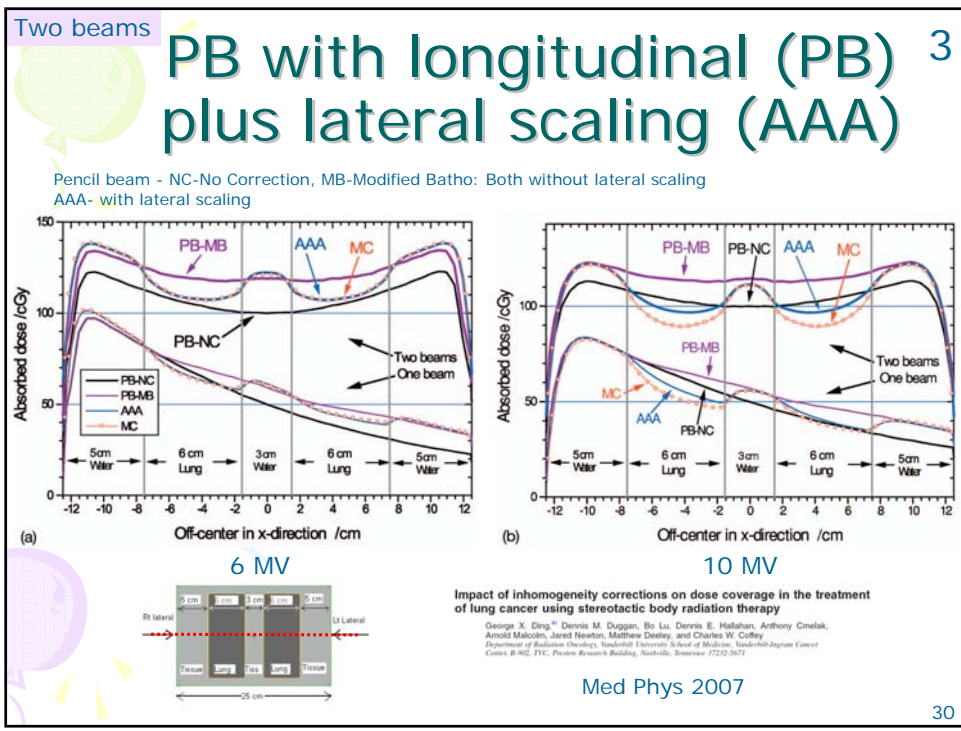
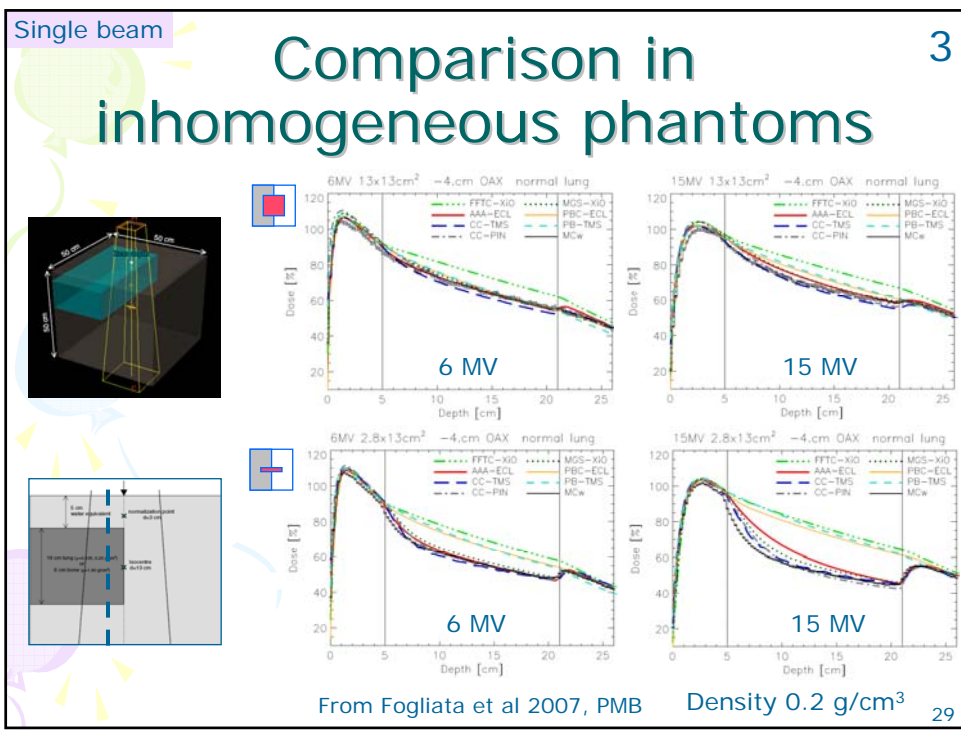
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Convolution methods in homogeneous water

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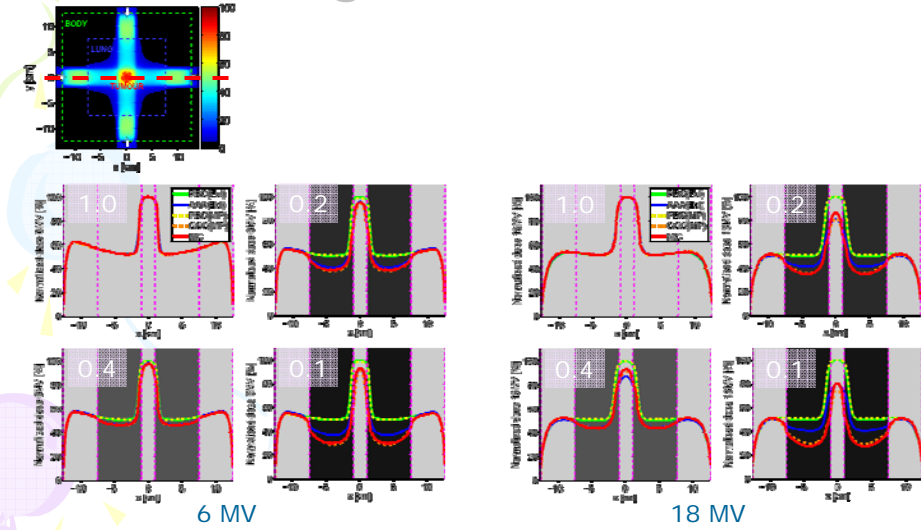
- Differences in beam modelling
 - Head scatter
 - Electron contamination
 - Wedges/Blocks
 - MLC
- May lead to slightly different accuracy
- Basically all models perform well in water
 - Point, pencil or collapsed cone implementations

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Multiple beams

PB w/wo lateral scaling and CC vs MC

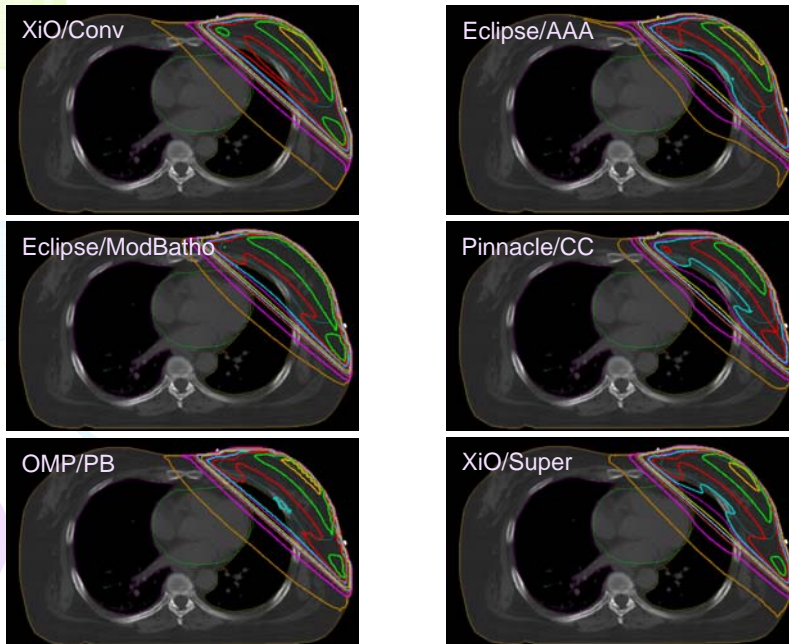


From Lasse Rye Aarup, Copenhagen

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Tangential treatment of breast 3

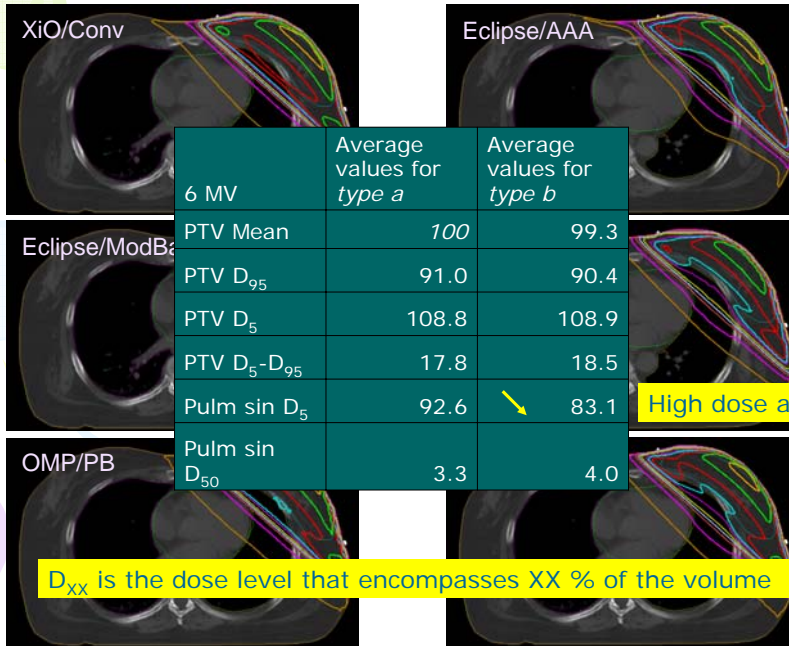
Knöös et al., 2006, PMB



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Tangential treatment of breast 3

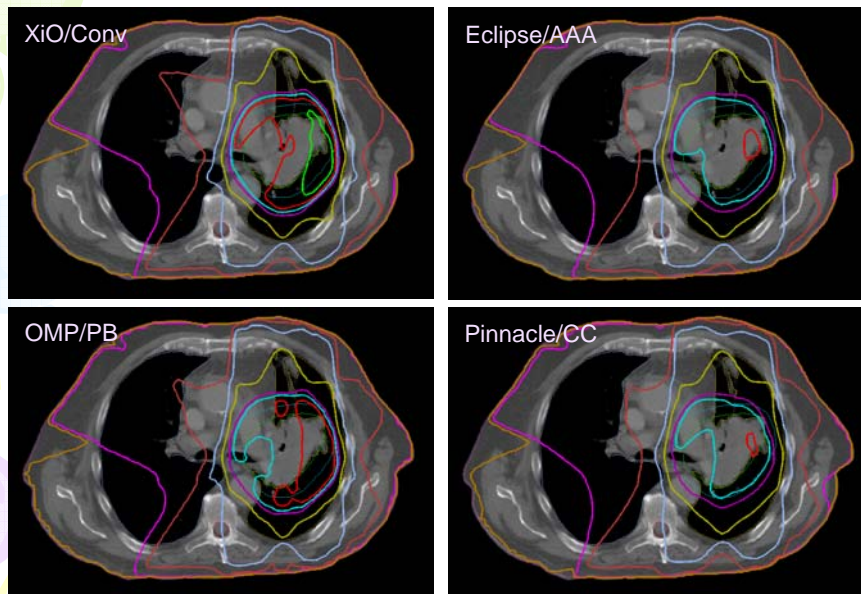
Knöös et al, 2006, PMB



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5 field 18 MV – lung 3

Knöös et al, 2006, PMB



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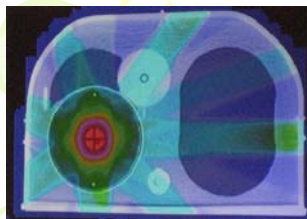
5 field 18 MV – lung

3

	XiO/Conv		Eclipse/AAA	
	Average values for type a	Average values for type b	Average values for type a	Average values for type b
PTV Mean	100	97.5	100	96.3
PTV D ₉₅ ~max	92.7	91.3	95.2	91.5
PTV D ₅ ~min	106.2	102.8	104.4	99.8
PTV D ₅ -D ₉₅	13.5	11.6	9.2	8.3
Pulm Sin D ₅₀	14.2	19.7	15.7	20.6
Pulm Sin D ₅	103.7	96.3	101.3	91.9
Pulm Sin D ₁	107.9	100.0	104.3	95.9

Knös et al, 2006, PMB

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Results from RPC thorax phantom

- 15 cases planned with **type a**
 - 84% ± 16% of the pixels met the criteria (5%/5mm)
- 30 cases planned with **type b**
 - 99% ± 4% of the pixels met the criteria (5%/5mm)

AAPM 2008 TU-C-AUD B-3 P Alvarez et al

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Conclusions – Dose changes

- Prostate
 - non-significant
- H&N
 - none (depending on accuracy of scatter integration) and air cavities (air or low dense water)
- Breast
 - slightly lower dose to breast and especially in lung in proximity to the target however larger irradiated lung volume
- Lung - PTV
 - 2-4 % lower average dose
 - Wider penumbra
- Lung (treated side)
 - 10 % lower dose to the highest irradiated parts of the lung
 - 5 % higher dose (15 => 20 %) to the lung (D₅₀)
- Lung (healthy side)
 - Average dose identical (9.8-10.7 %)

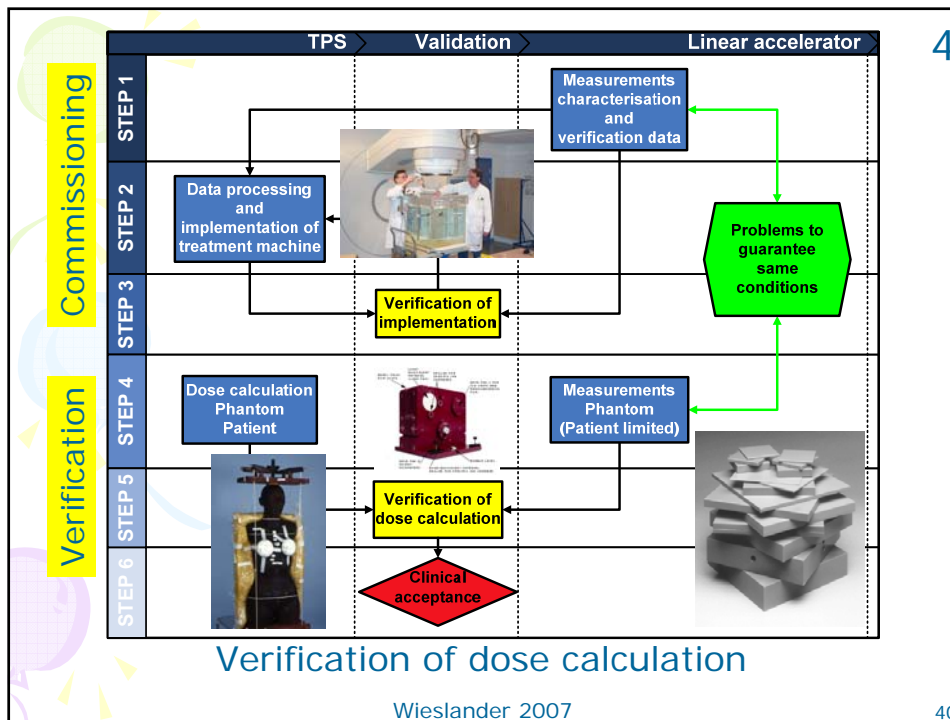
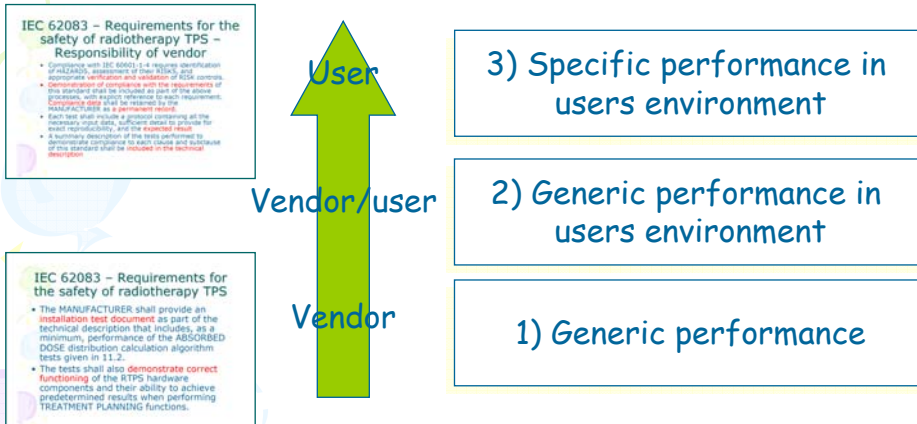
Knöös et al, 2006, PMB

How to verify dose calculation algorithms?

The collage includes the following documents:

- AAPM REPORT NO. 48**: Comprehensive QA for Radiation Oncology
- CEI IEC 62083**: Norme Internationale Standard
- ESTRO**: Guidelines for the Verification of EMU
- ESTRO**: Dose Unit Calculation High Energy Photon Beams
- IAEA**: Commissioning a Quality Assurance Computerized Planning Systems for Radiation Treatment of Cancer (Reports Series No. 430)
- IAEA**: Quality Assurance of Treatment Planning Systems - Practical Examples for Non-IMRT Photon Beams
- IAEA**: Specification and Acceptance Testing of Radiotherapy Treatment Planning Systems
- American Association of Physicists in Medicine Radiation Therapy Committee Task Group 53**: Quality assurance for clinical radiotherapy treatment planning (Med Phys 1998)

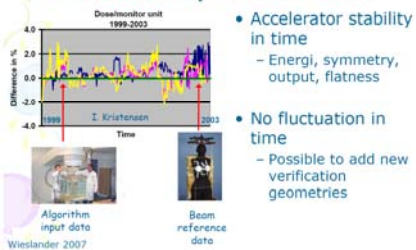
Process of acceptance



Verification of dose calculation

MC methods facilitate verification

Consistency of data sets

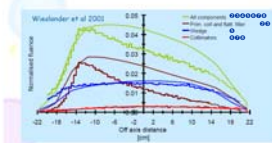


- Accelerator stability in time
 - Energi, symmetry, output, flatness
- No fluctuation in time
 - Possible to add new verification geometries

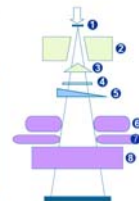
Wieslander 2007

Study of beam models

Labelling in MC codes to keep track of interaction history



Wieslander 2007

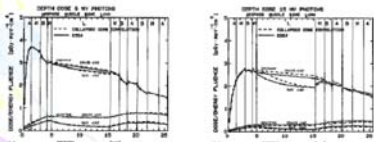


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Studies of dose components

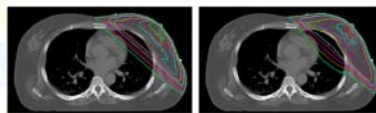
Separation of dose components in phantoms primary and scatter dose



Inhomogeneous slab geometry 6 and 15 MV photons
A. Ahnesjö, Med. Phys., 1989 EGS4
Wieslander 2007

Virtual linear accelerator

Comparisons in patient geometry



TPS calculated dose distribution based on the virtual accelerator

MC calculated dose distribution based on the virtual accelerator

Wieslander 2007

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Implications of introducing new and more accurate algorithms ⁵

- Significant changes in dose to target volumes and surrounding tissues especially when lung is involved
 - Consequences for assessment of dose-effect relationships
- Careful analysis of changes is required before adopting new algorithms
 - Retrospectively re-calculate plans when clinical outcome is known?
 - Construct new plans with older algorithms and re-calculate?
 - New plans with old prescriptions and new algorithms?
 - Optimize plans to the same biological effect on PTV and/or OAR?

Morgan et al 2008

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Implications of introducing new and more accurate algorithms ⁵

- Significant changes in dose to target volumes and surrounding tissues especially when lung is involved
 - Consequences for assessment of dose effect relationships
- Careful analysis of changes is required before adoption of new algorithms
 - When is it necessary to construct new plans with older algorithms and re-calculate?
 - New plans with old prescriptions and new algorithms?
 - Optimize plans to the same biological effect on PTV and/or OAR?

Discussion is needed between physicists and oncologists to fully understand the effects and potential consequences

Morgan et al 2008

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Conclusion

- Convolution methods are accurate
 - For low density regions – use models with lateral scaling
- Verification
 - Also Vendor's responsibility!
- Be careful when transferring to more accurate models but...
- **Important to do this!**

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