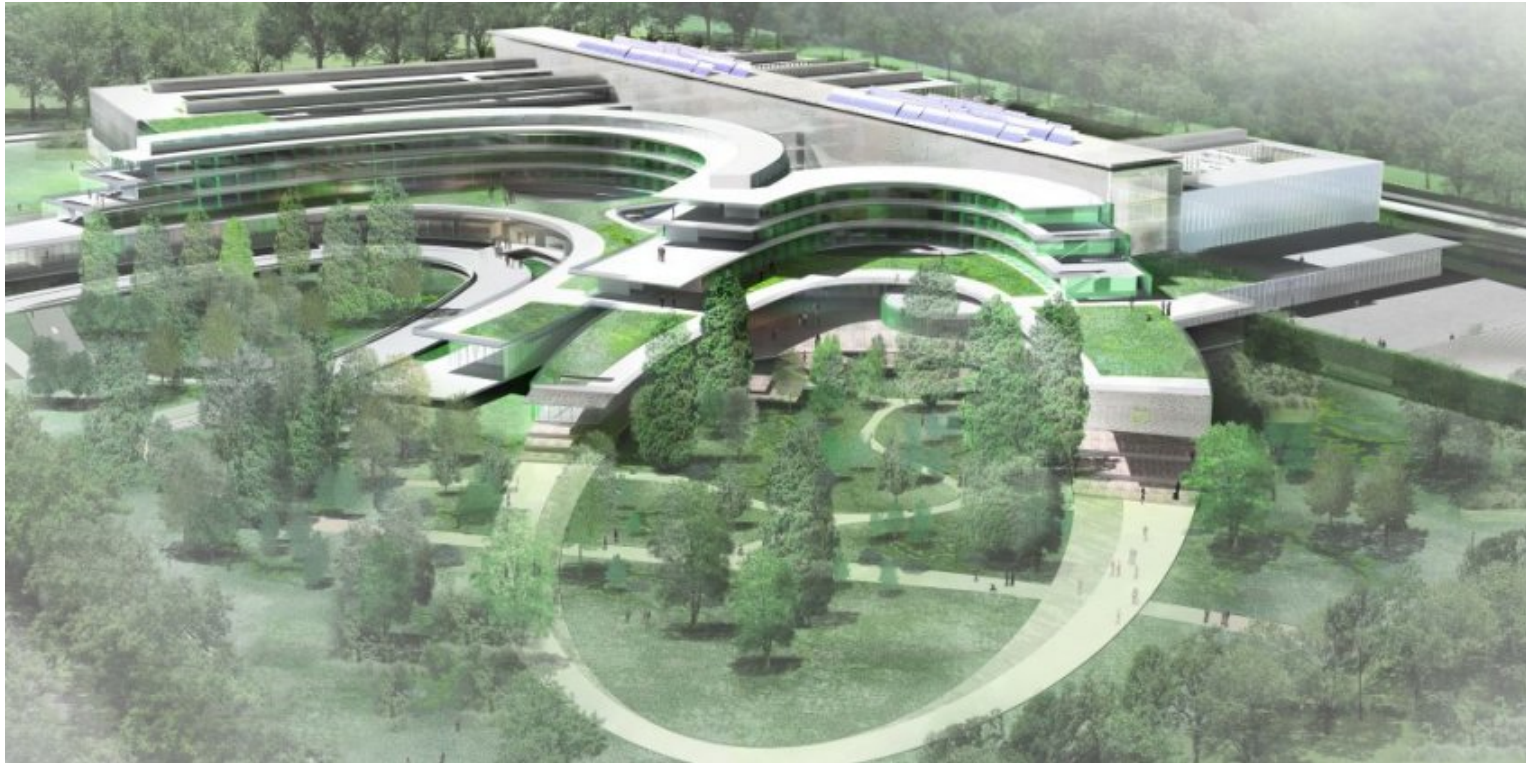


# From Model-based to Patient-specific dosimetry in Nuclear Medicine



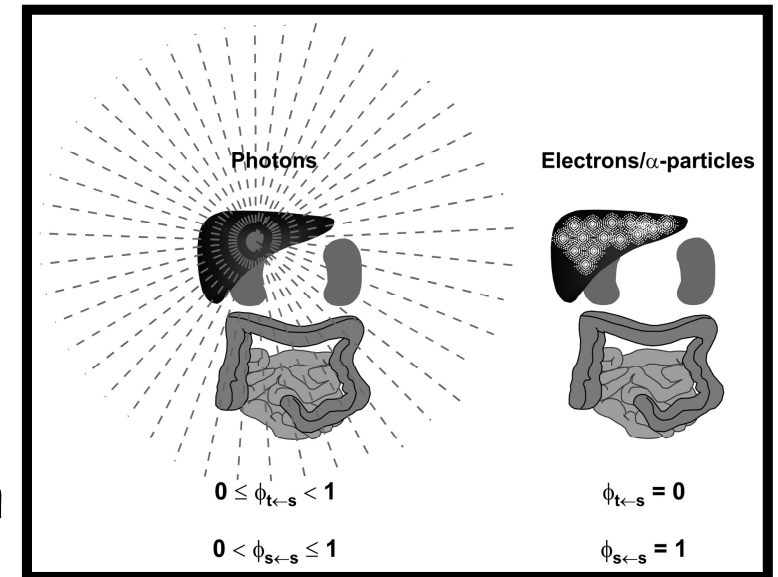
Manuel Bardiès ([manuel.bardies@inserm.fr](mailto:manuel.bardies@inserm.fr))

Centre de Recherches en Cancérologie de Toulouse, France

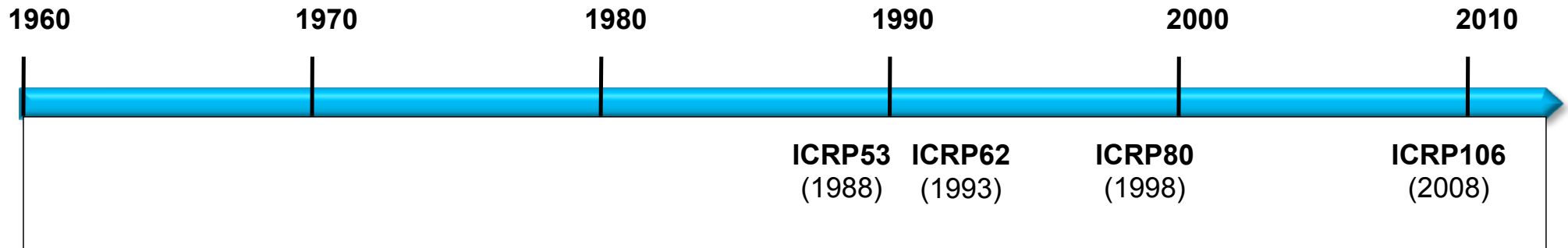
# Nuclear Medicine Dosimetry

$$\bar{D}_k = \sum_h \tilde{A}_h \times S_{(k \leftarrow h)}$$

- MIRD formalism
  - $\tilde{A}$ : Cumulated activity
    - Quantitative Imaging
    - Time-Activity Curve integration
  - $S$ : Absorbed Dose Calculation
- And... global accuracy relies on both terms:
  - Improving  $\tilde{A}$  requires improving  $S$  (and vice-versa)

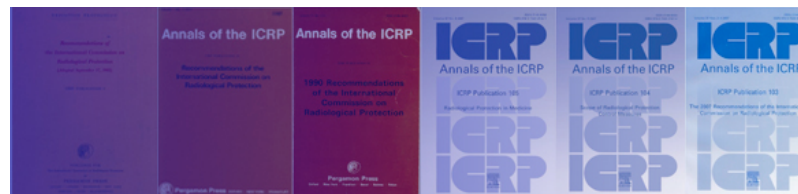


# Diagnositics dosimetry

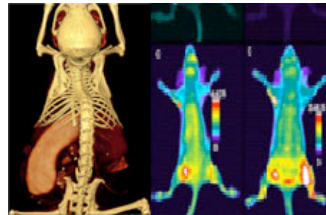
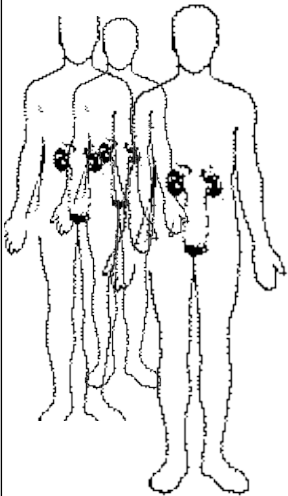


«Radiation dose to patients from radiopharmaceuticals»

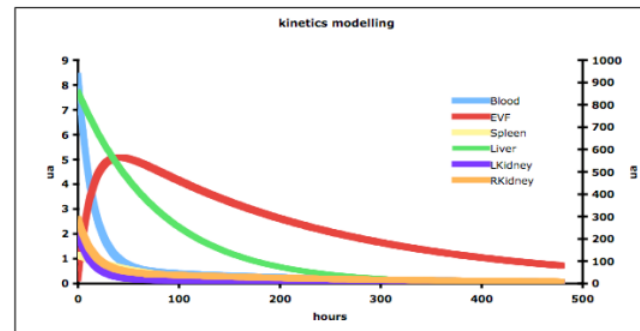
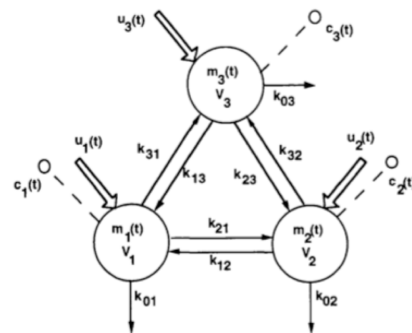
- 1988 ICRP Publication 53. Ann. ICRP 18 (1-4)
- 1993 Addendum 1 to ICRP Publication 53. Ann. ICRP 22(3)
- 1998 Addendum 2 to ICRP Publication 53. Ann. ICRP 28 (3)
- 2008 Addendum 3 to ICRP Publication 53. Ann. ICRP 38 (1-2)



# ICRP Approach (Diagnostics)

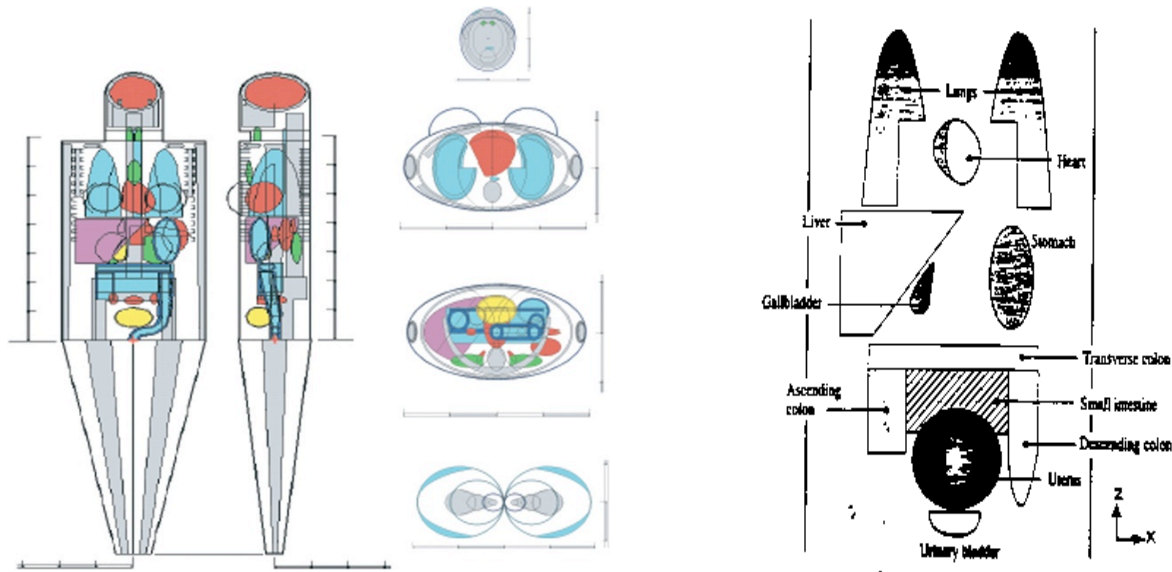


Ä

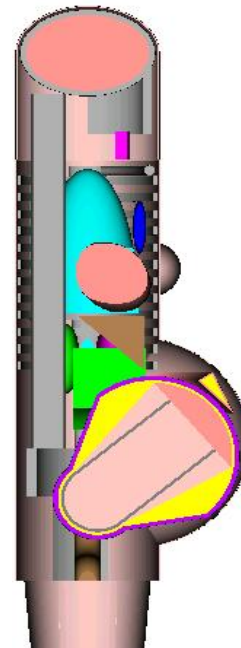
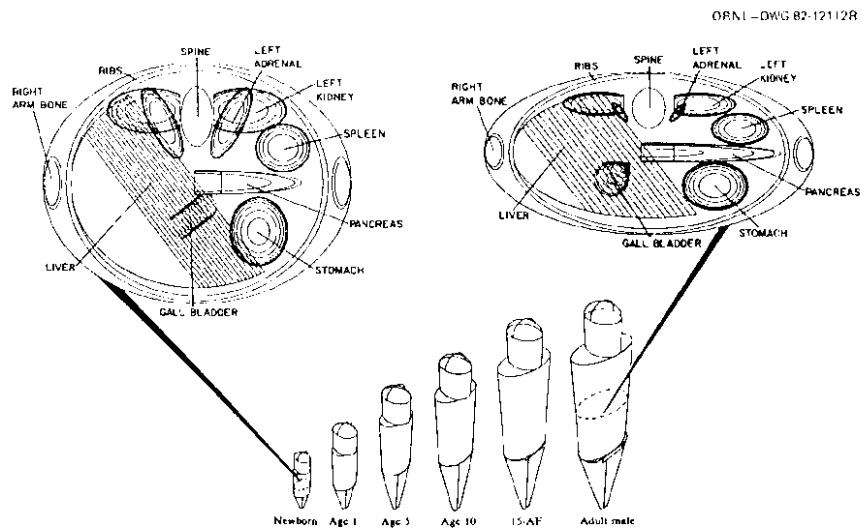




# ICRP Approach (Diagnostics)



S



# Computing models

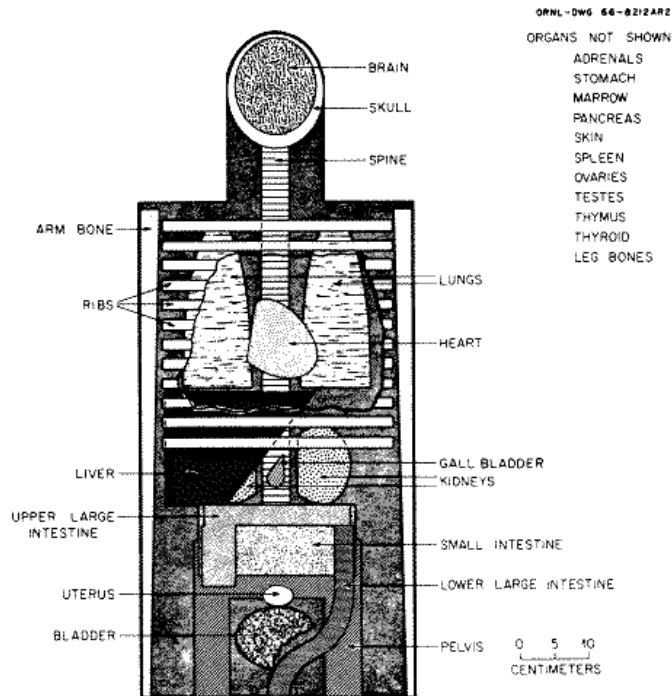
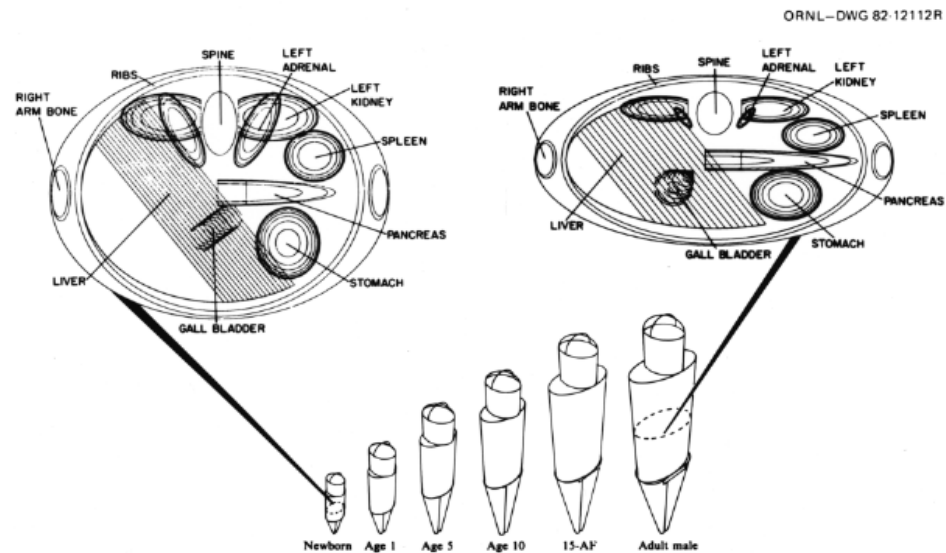
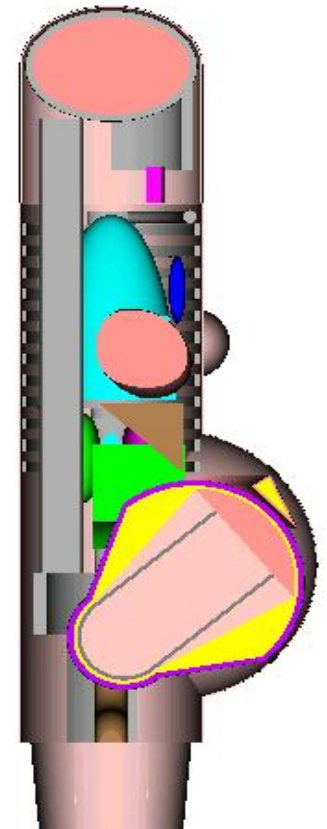


Fig. A-3. Anterior view of the principal organs in the head and trunk of the adult phantom developed by Snyder et al. (1974). Although the heart and head have been modified in this report, this schematic illustrates the simplicity of the geometries of the organs.

Snyder 1975

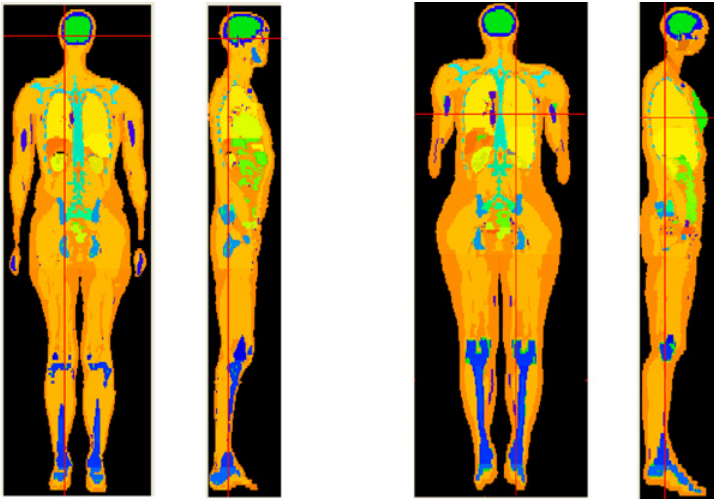


Cristy & Eckerman 1987

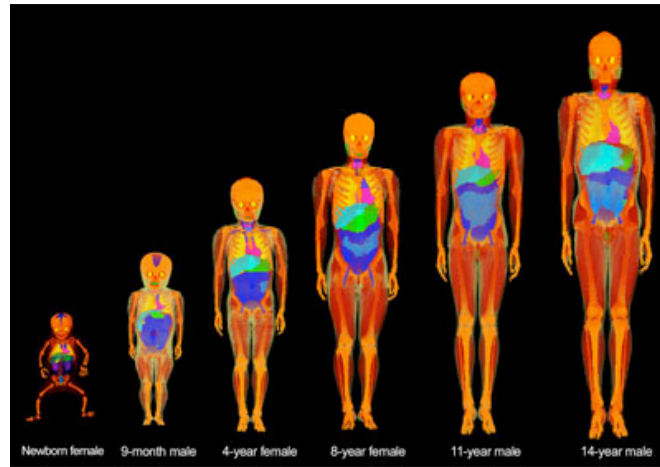


Stabin 1995

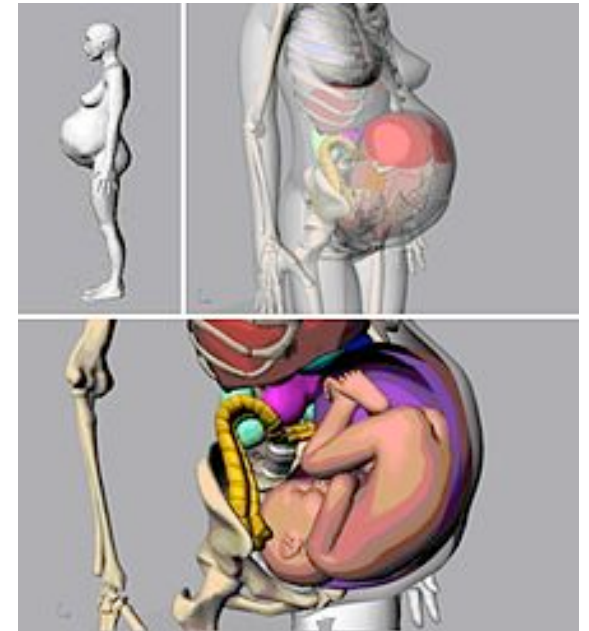
# Computing models



Reference Adult  
male/female  
ICRP 110



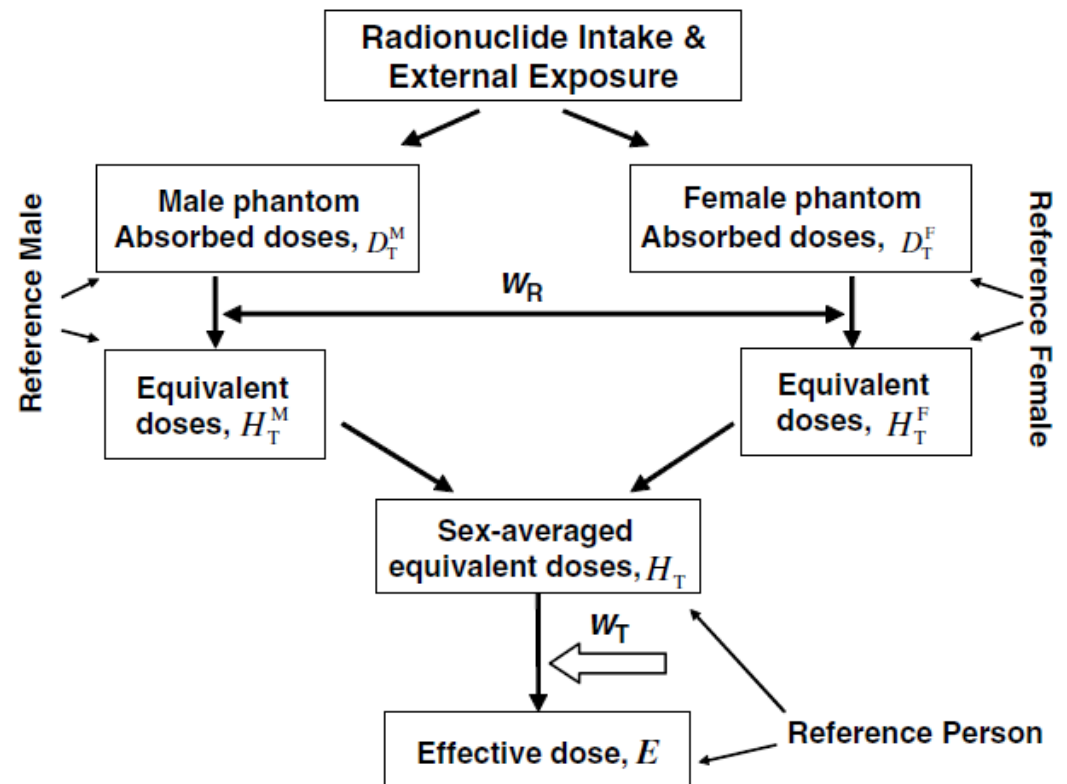
Paediatric series  
Lee et al. (2010)  
PMB 55(2):339-363



Pregnant female  
Guo et al. (2010)  
RPD 138(1):20-28

# ICRP Evolution

- Recent reference report (ICRP 103)
- New computing models (ICRP 110 + ... ?)
- New calculation scheme
- New weighting factors
- Transition phase! (ex: ICRP 106)





# New ICRP 110 models

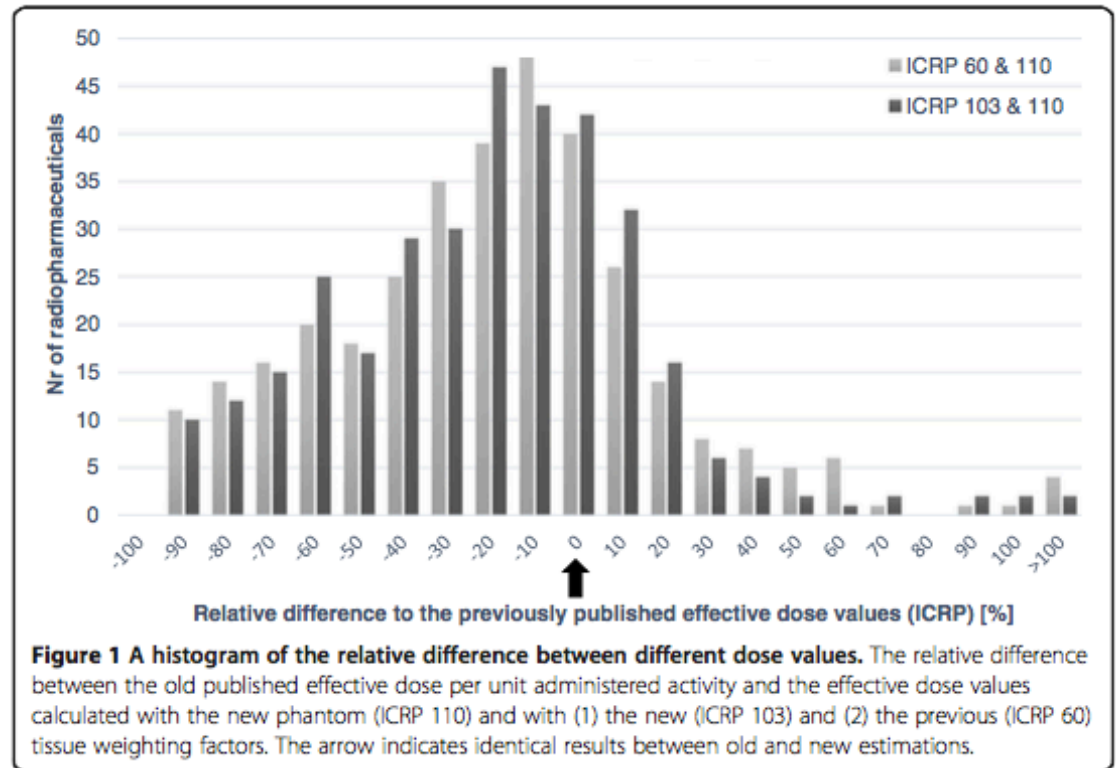
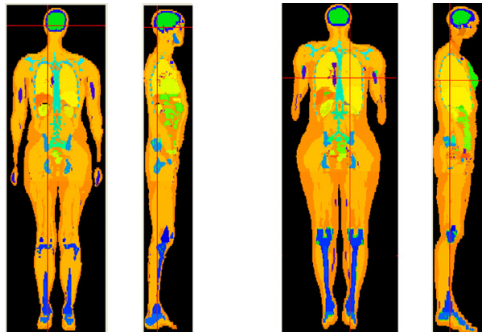
ORIGINAL RESEARCH

Open Access

Effective dose to adult patients from 338 radiopharmaceuticals estimated using ICRP biokinetic data, ICRP/ICRU computational reference phantoms and ICRP 2007 tissue weighting factors

Martin Andersson<sup>1\*</sup>, Lennart Johansson<sup>2</sup>, David Minarik<sup>1</sup>, Sigrid Leide-Svegborn<sup>1</sup> and Sören Mattsson<sup>1</sup>

Andersson et al. EJNMMI Physics 2014 1:9

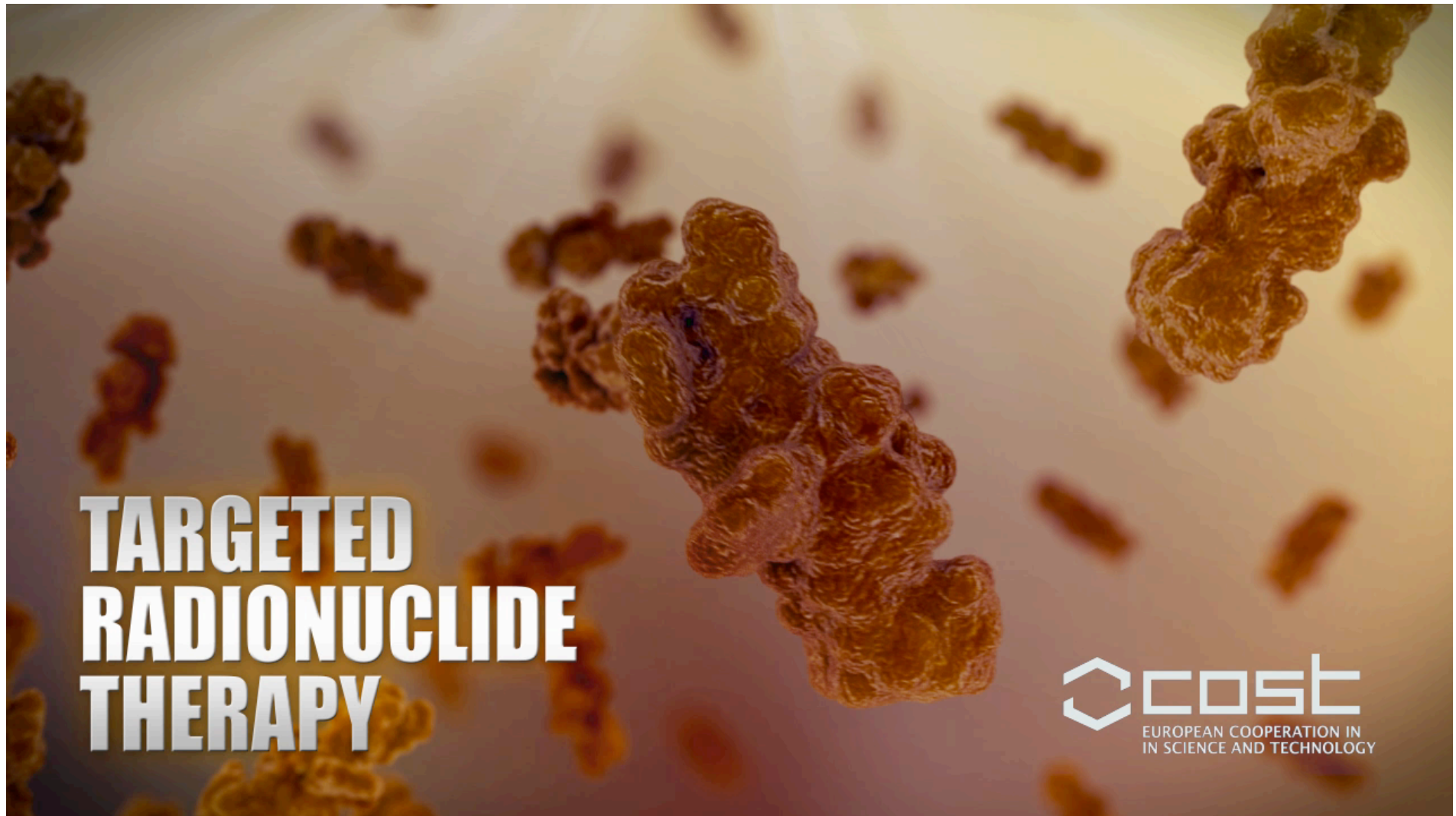


- Zankl et al. «Electron specific absorbed fractions for the adult male and female ICRP/ICRU reference computational phantoms» Phys Med Biol 2012, 57(14):4501–4526
- Andersson et al. «An internal radiation dosimetry computer program, IDAC2.0, for estimation of patient dose for radiopharmaceuticals» Radiat Prot Dosimetry 2013; doi: 10.1093/rpd/nct337

# Diagnosics dosimetry: Conclusion

Group	Model	Model ICRP - MIRD DER

# Molecular Radiotherapy



<http://www.youtube.com/watch?v=GRRmX5eTa8s>

# Dosimetry for MRT:

Group	Model	Model ICRP - MIRD DER
Specific		

- *Patient-specific dosimetry requires AT LEAST a specific determination of  $\tilde{A}_h$*



# Quantitative imaging: $\tilde{A}_h$

Is quantitative imaging for dosimetric purposes different from 'conventional' quantitative imaging in NM?

Quick answer: No...

...but some aspects are specific...

What kind of quantitative imaging is required for dosimetry?

# Quantitative imaging: $\tilde{A}_h$

- What quantitative imaging implies:
  - On principle: Absolute quantification
  - Activity concentration in all voxels (Bq/cc)
  - Corrections OK for the whole FOV
- For the whole patient (space)
- Follow radiopharmaceutical kinetics (time)

MIRD Phamphlet 16 (Siegel *et al.* JNM 40, 37s-61s, 1999)

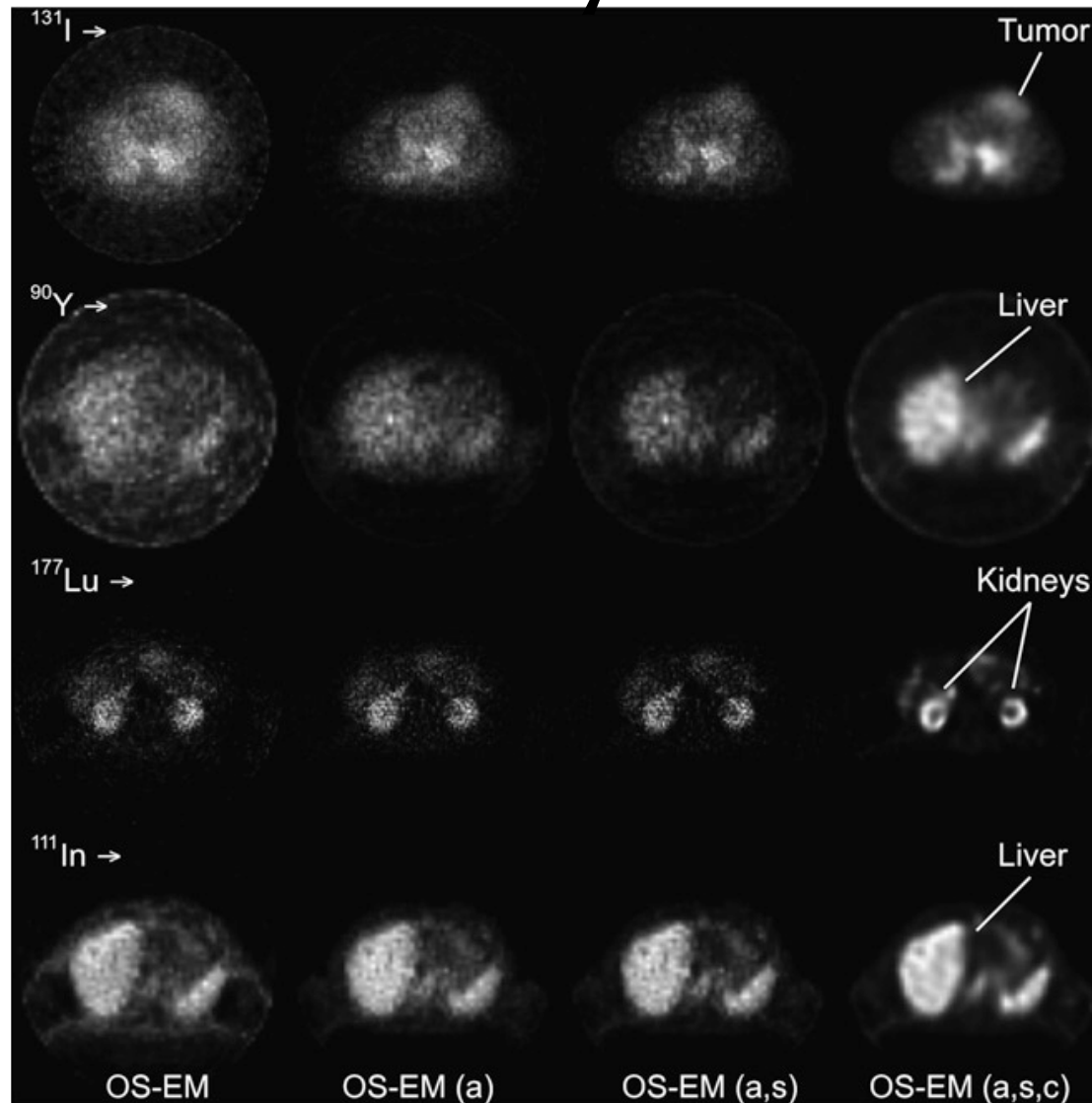
# Assessing errors: the main issue?

- Methodologies have been proposed to correct for several effects that degrade the quantitative content of NM images
- Many references are available in the literature!
  - Some approaches were implemented in clinical practice
  - Most remain as 'one centre' approach
- So who's right?

Dewaraja YK et al. **2012, MIRD pamphlet No. 23:**

Quantitative SPECT for patient-specific 3-dimensional dosimetry in internal radionuclide therapy. *J Nucl Med* 53(8), pp. 1310-25

# SPECT: currently used isotopes



Dewaraja YK et al. **2012, MIRD pamphlet No. 23:**

Quantitative SPECT for patient-specific 3-dimensional dosimetry in internal radionuclide therapy. *J Nucl Med* 53(8), pp. 1310-25



# SPECT: currently used isotopes

Study	Radionuclide	System	Reconstruction	accuracy
Zeintl et al., 2010 (18)	$^{99m}\text{Tc}$	SPECT/CT	OS-EM, CDR, CT-derived AC, energy window-based SC, PVC	<6.8% error for 0.5- to 16-mL spheres
Dewaraja et al., 2010 (37)	$^{131}\text{I}$	SPECT/CT	OS-EM, CDR, CT-derived AC, energy window-based SC	<17% error for 8- to 95-mL spheres; 31% for 4-mL sphere
Assie et al., 2010 (23)	$^{111}\text{In}$	SPECT and CT separate	OS-EM, CT-derived AC, energy window-based SC, PVC	<20% error for organs and 2- to 32-mL spheres; 48% error for 0.5-mL sphere
Shcherbinin et al., 2008 (49)	$^{99m}\text{Tc}$ , $^{111}\text{In}$ , $^{123}\text{I}$ , $^{131}\text{I}$	SPECT/CT	OS-EM, CDR, CT-derived AC, analytic scatter modeling	3%–5% error for 32-mL bottles
Minarik et al., 2008 (95)	$^{90}\text{Y}$	SPECT/CT	OS-EM, CDR, CT-derived AC, ESSE	<11% error for liver and 100-mL sphere
Willowson et al., 2008 (19)	$^{99m}\text{Tc}$	SPECT/CT	OS-EM, CT-derived AC, transmission-dependent SC, PVC	<4% error for liver and cardiac chambers
de Wit et al., 2006 (59)	$^{166}\text{Ho}$	SPECT	OS-EM, CDR, $^{153}\text{Gd}$ transmission source-derived AC, Monte Carlo scatter modeling	16% average error for 220-mL bottles
Du et al., 2006 (62)	$^{123}\text{I}$	SPECT/CT	OS-EM, CDR, CT-derived AC, ESSE, PVC	<2% error for putamen and caudate regions of brain phantom
He at al, 2005 (52)	$^{111}\text{In}$	SPECT/CT	OS-EM, CDR, CT-derived AC, ESSE, PVC	<12% error for organs and 8- to 23-mL spheres
Koral et al., 2005 (50)	$^{131}\text{I}$	SPECT and CT separate	OS-EM, CDR, CT-derived AC, energy window-based SC, PVC	<7% average error for 100-mL sphere

Dewaraja YK et al. **2012, MIRD pamphlet No. 23:**

Quantitative SPECT for patient-specific 3-dimensional dosimetry in internal radionuclide therapy. *J Nucl Med* 53(8), pp. 1310-25

# Dosimetry for MRT:

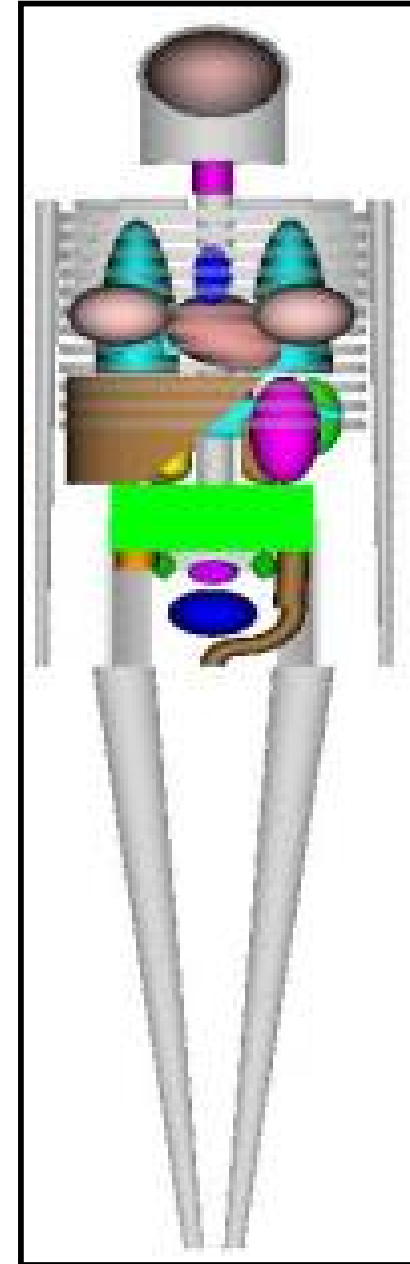
Group	Model	Model ICRP - MIRD DER
Specific		

- *Possibly the most important source of uncertainty?*

# S factor calculation: $S_{(k \leftarrow h)}$

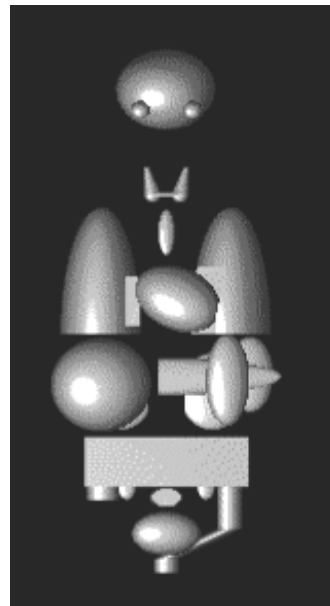
- From 'old' MIRD phantoms

MIRDOSE3  
Olinda

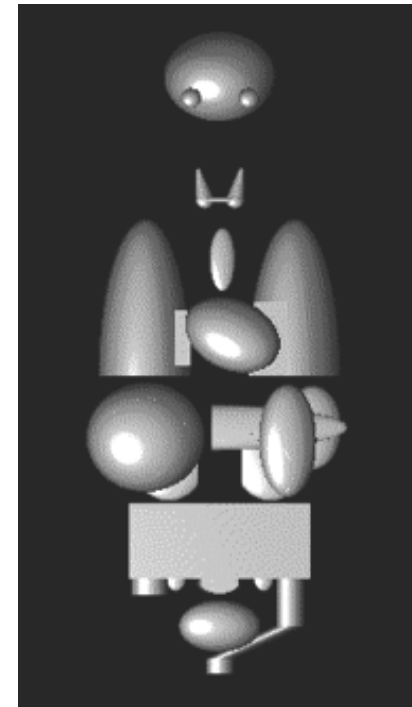


# S factor calculation: $S_{(k \leftarrow h)}$

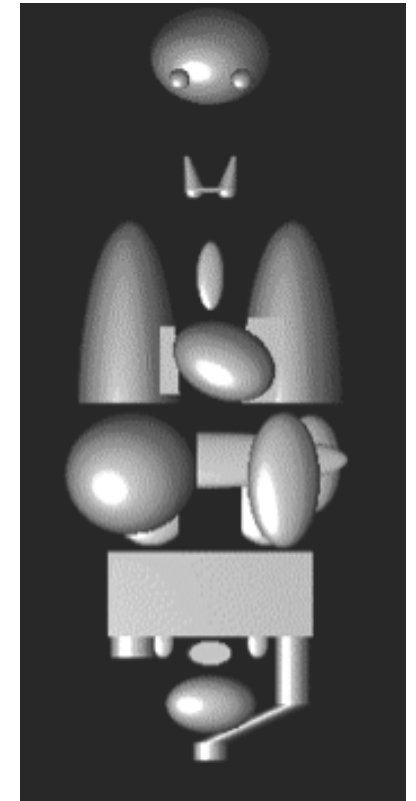
- From 'old' MIRD phantoms
- To more refined phantoms



150 cm



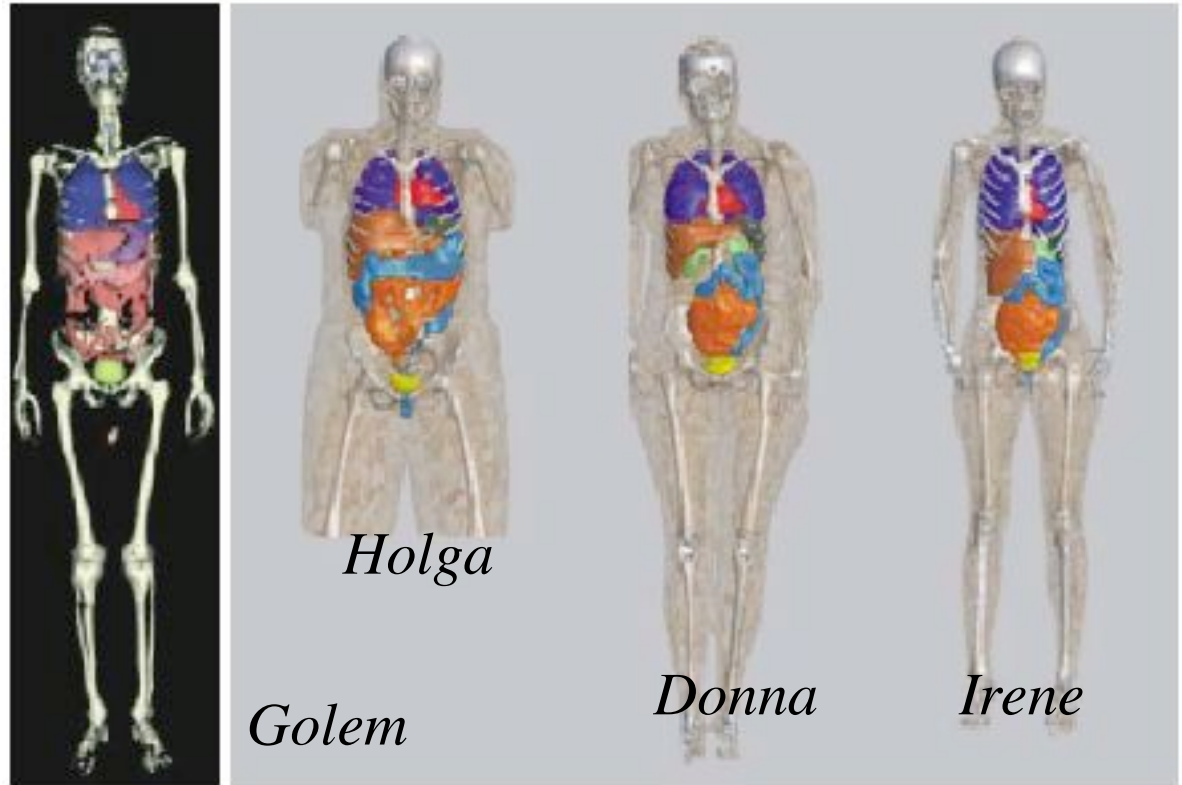
160 cm



170 cm

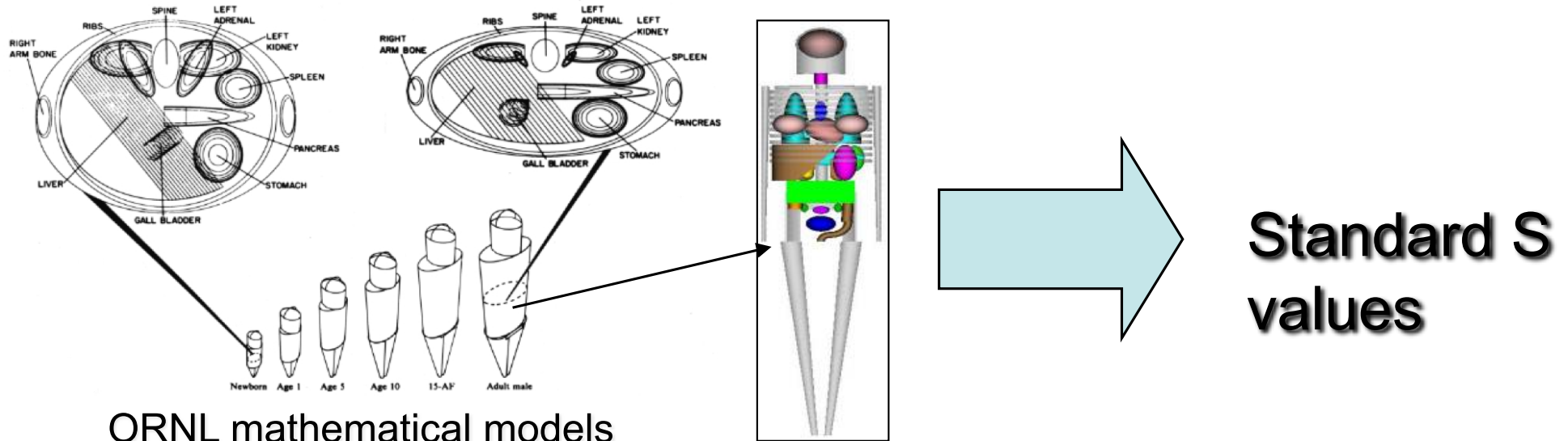
# S factor calculation: $S_{(k \leftarrow h)}$

- From 'old' MIRD phantoms
- To more refined phantoms
- To voxel-based phantoms



*Radiat. Env. Biophys* (2001) 40:153-162  
*PMB* (2002) 47:89-106

# Mass Adjustment



ORNL mathematical models  
(ORNL/8381)

For SELF Irradiation Only

$$S_{r \leftarrow r}(\textit{patient}) = S_{r \leftarrow r}(\textit{standard}) \cdot \frac{\textit{Mass}_r(\textit{standard})}{\textit{Mass}_r(\textit{specific})}$$

# OLINDA mass adjustment

Input Data:

pha

## Model to adjusted-model!

1420.0	Brain	1120.0	Red Marrow
351.0	Breasts	120.0	Osteogenic Cells
10.5	Gallbladder Wall	3010.0	Skin
167.0	LLI Wall	183.0	Spleen
677.0	Small Intestine	39.1	Testes
158.0	Stomach Wall	20.9	Thymus
220.0	ULI Wall	20.7	Thyroid
316.0	Heart Wall	47.6	Urinary Bladder Wall
299.0	Kidneys	79.0	Uterus
1910.0	Liver	0.0	Fetus
1000.0	Lungs	0.0	Placenta
28000.0	Muscle	73700.0	Total Body
8.71	Ovaries		

Alpha Weight Factor

5.0

Beta Weight Factor

1.0

Photon Weight Factor

1.0

Reset organ values

Multiply all masses by:

1.0

DONE



# Dosimetry for MRT:

Group	Model	Model ICRP - MIRD DER
Specific	Model $\pm$ adjusted	Model $\pm$ realistic

- *Still «model-based» dosimetry - but easily implemented in a clinical environment!*

# Dosimetry for MRT:

Group	Model	Model ICRP - MIRD DER
Specific	Model $\pm$ adjusted	Model $\pm$ realistic
Specific		

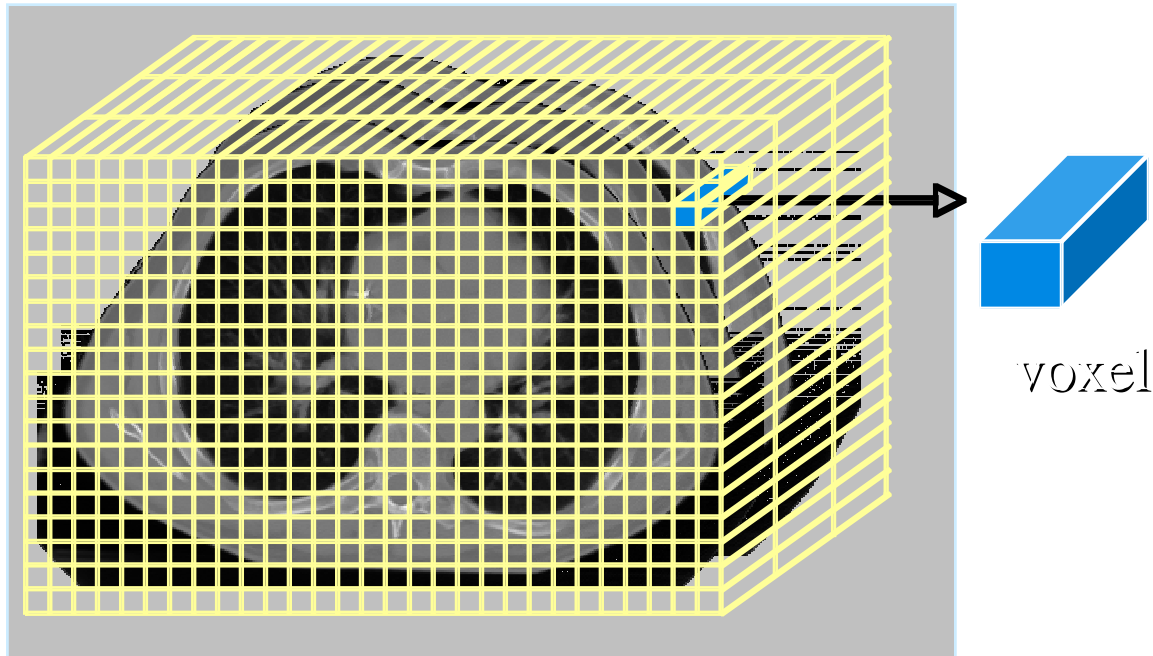
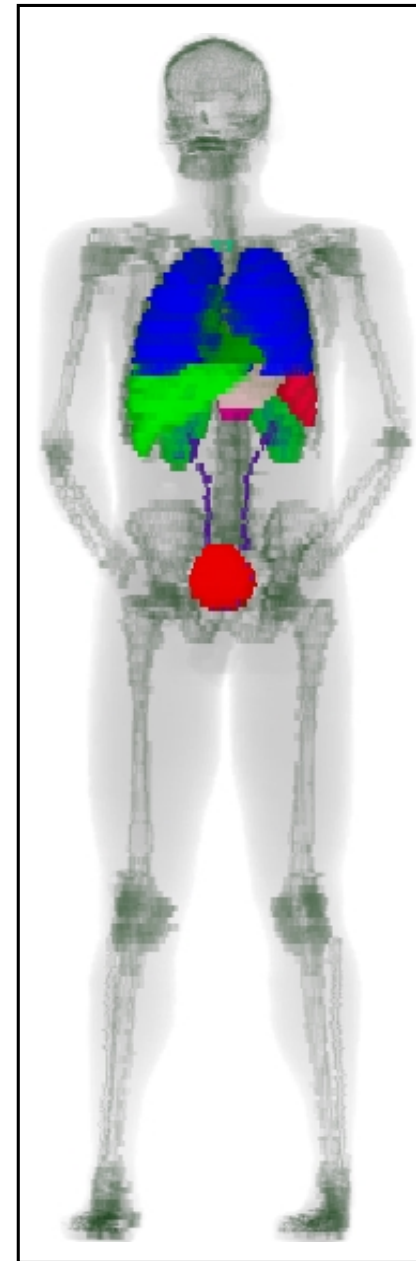
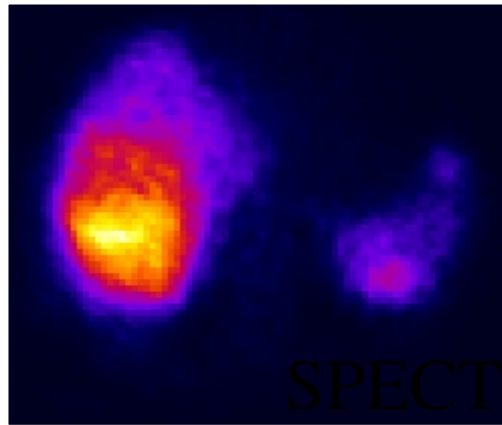
- *Patient-specific dosimetry requires AT LEAST a specific determination of  $\tilde{A}_h$*

# Dosimetry for MRT:

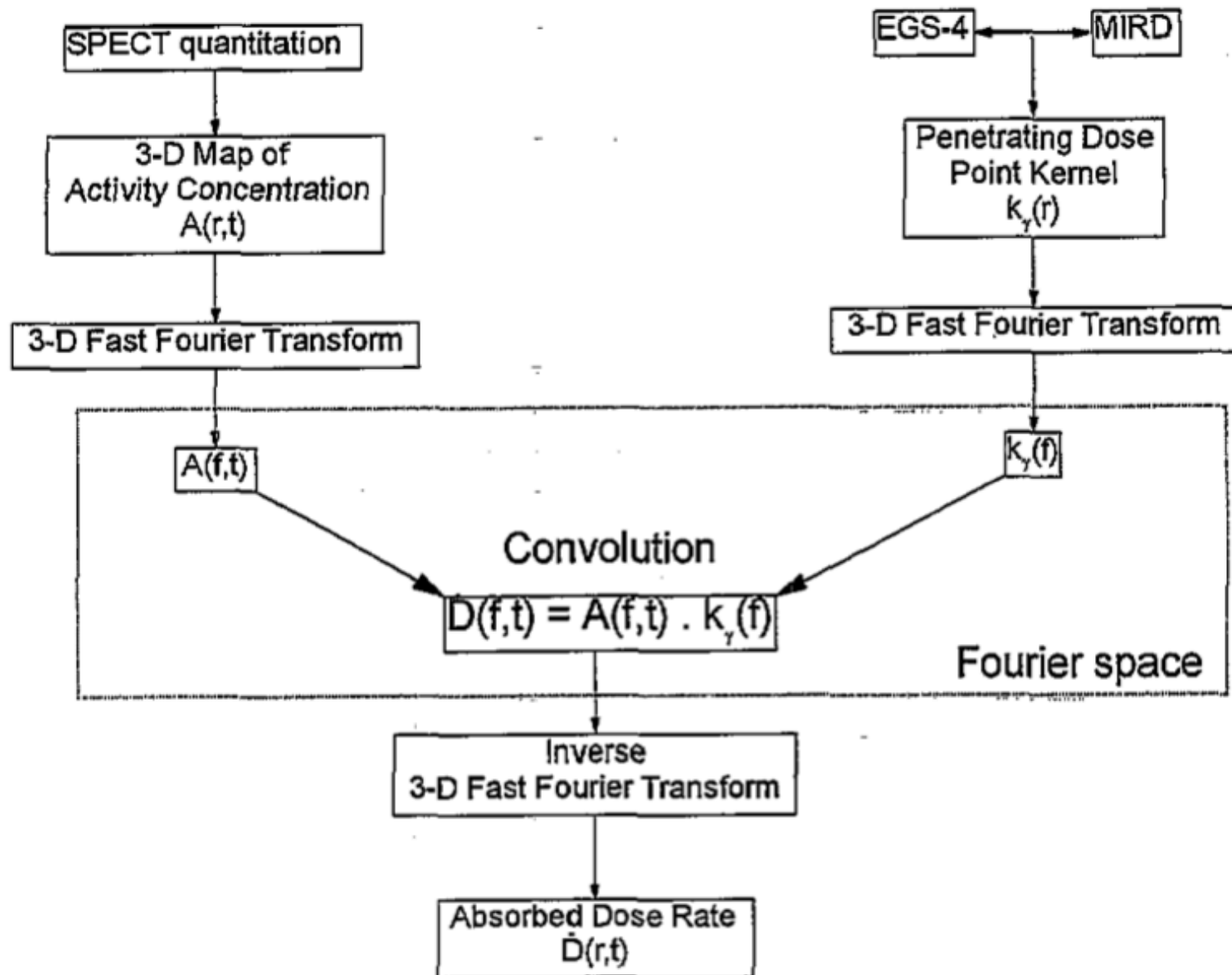
Group	Model	Model ICRP - MIRD DER
Specific	Model $\pm$ adjusted	Model $\pm$ realistic
Specific	Specific	

- *Specific S factor determination requires patient-specific geometry assessment*

# Patient-Specific dosimetry:



# Patient specific dosimetry



# Patient specific dosimetry

The screenshot displays the 3D-ID software interface for patient-specific dosimetry. The main window is titled "3D-ID DEFINE REGIONS OF INTEREST (version 0.1.1)". It features a menu bar with "File", "Contours", "Volume", "View", and "Help". Below the menu, there are input fields for "Patient name: CC 49 female" and "Patient Id: 999999".

The central area is divided into two panels. The left panel shows a CT scan slice of a liver with a yellow contour drawn around a region. The right panel shows a corresponding PET scan slice with a red and orange heatmap overlay, indicating the dose distribution. Above the right panel, there is a "Dual Modality Image Slices" toolbar with icons for "slice 5" through "slice 11".

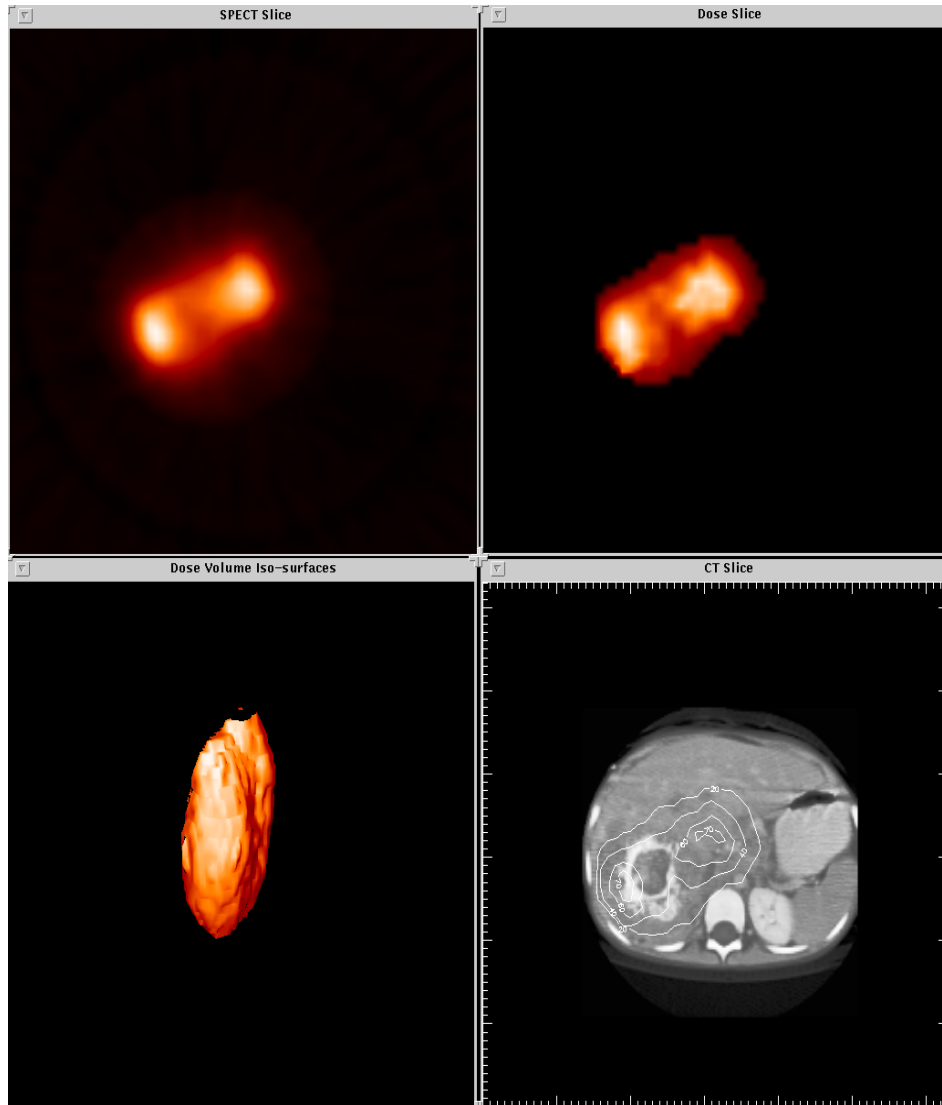
On the right side, there is an "ROI VOLUME" panel. It contains a list of regions of interest: "liver", "tumor 1", and "tumor 2". The "tumor 2" entry is selected. Below the list, it displays "Volume: 160,270 cc" and a "Dismiss" button.

At the bottom of the interface, there are two horizontal sliders for "Image 1: level" (ranging from 1 to 88) and "Image 2: level" (ranging from 1 to 95). Below these are "Contour functions" (Draw, Remove, Accept, Help) and "Point value" (048) and "Point coords" (084, 282) fields. A "Dismiss" button is also present in the bottom right corner.

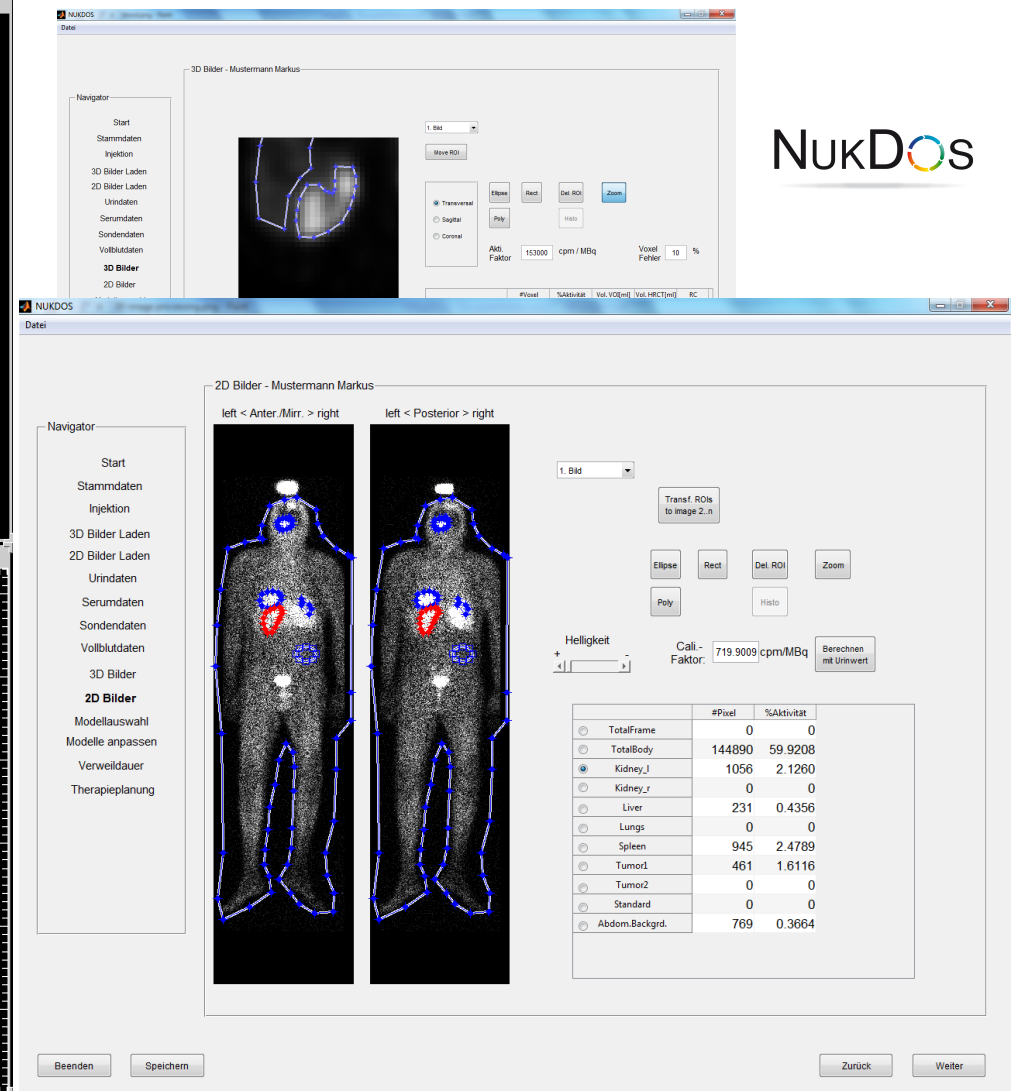
The text "3D-ID MSKCC" is visible in the bottom right corner of the interface.



# Therapy dosimetry

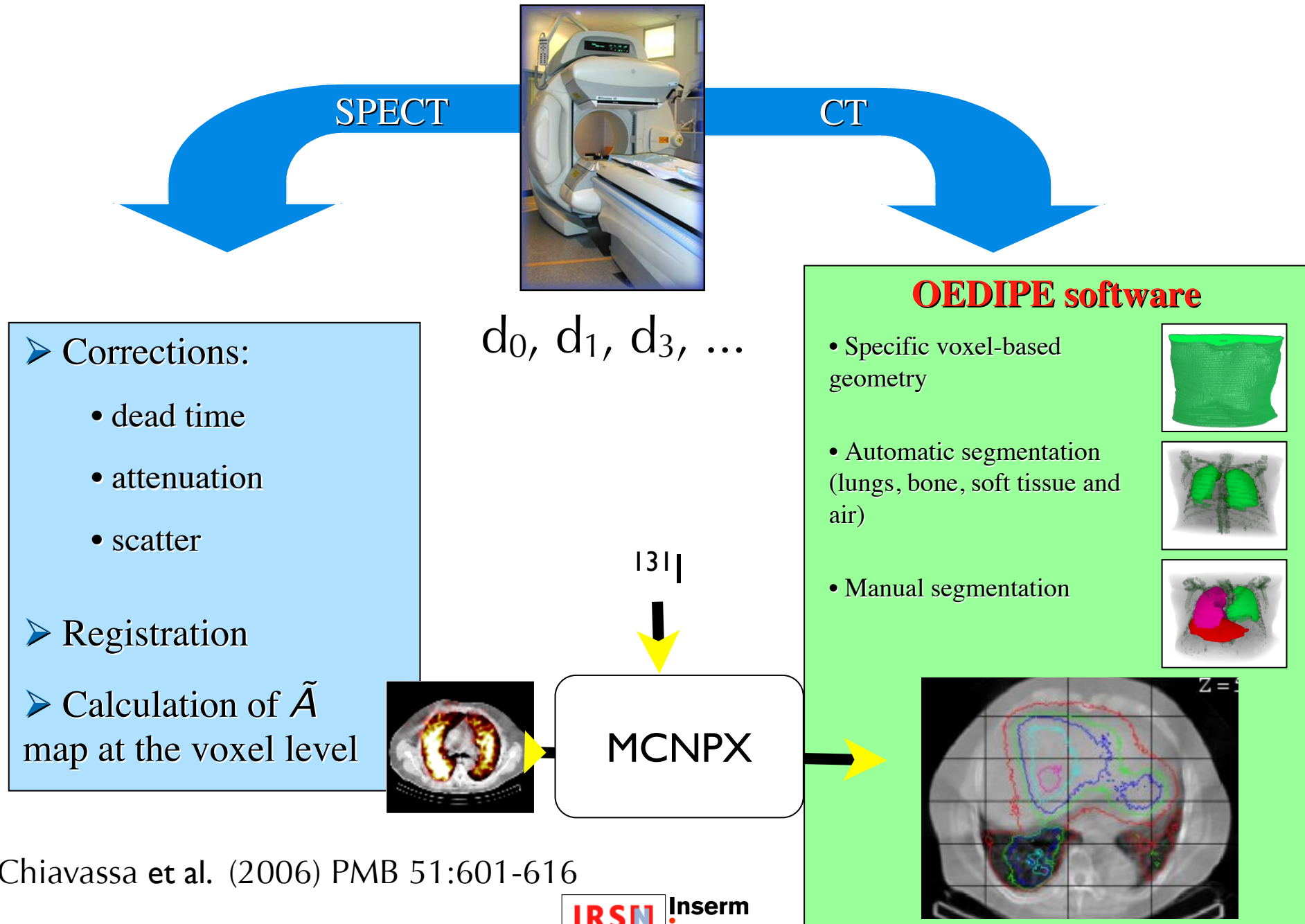


RMDP (M Guy, RMH)

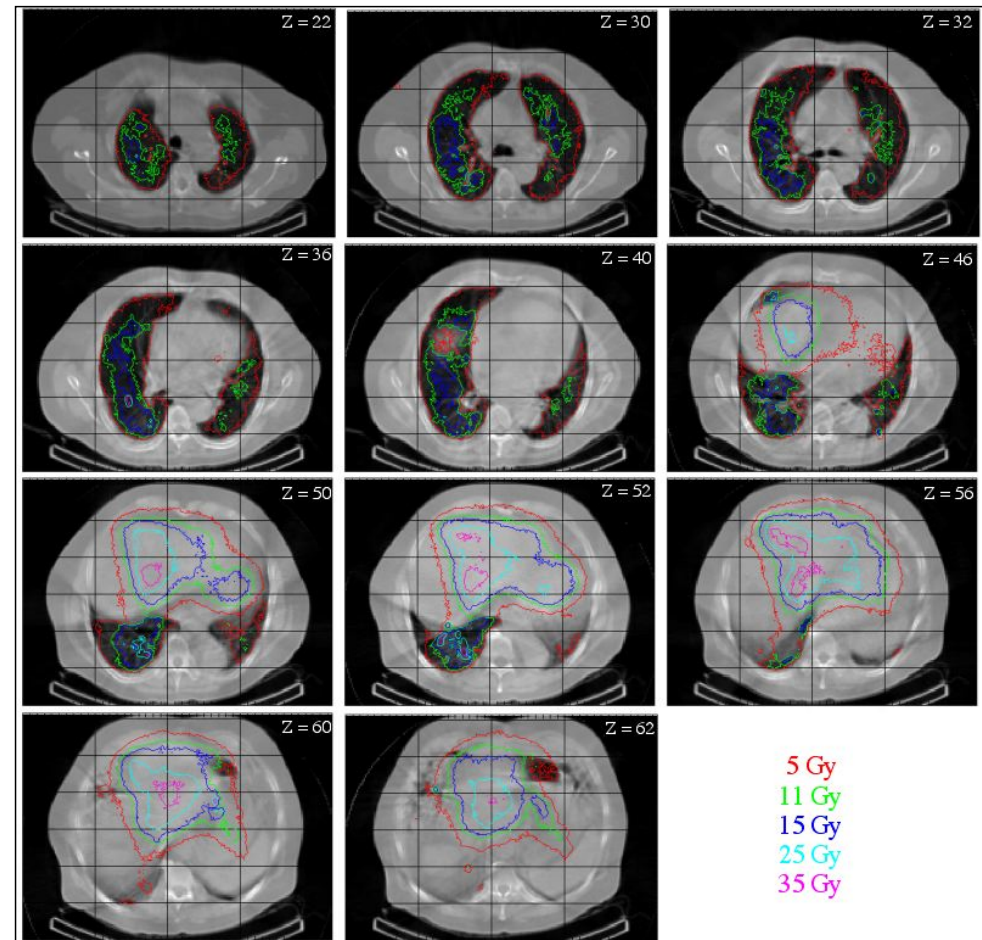
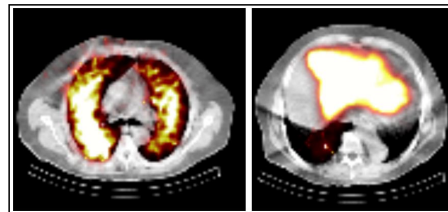
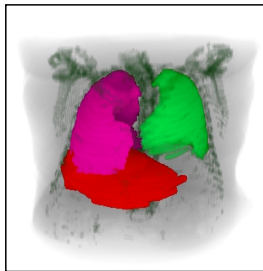


NukDos (M Laßmann, UKW)

# Patient-specific clinical dosimetry



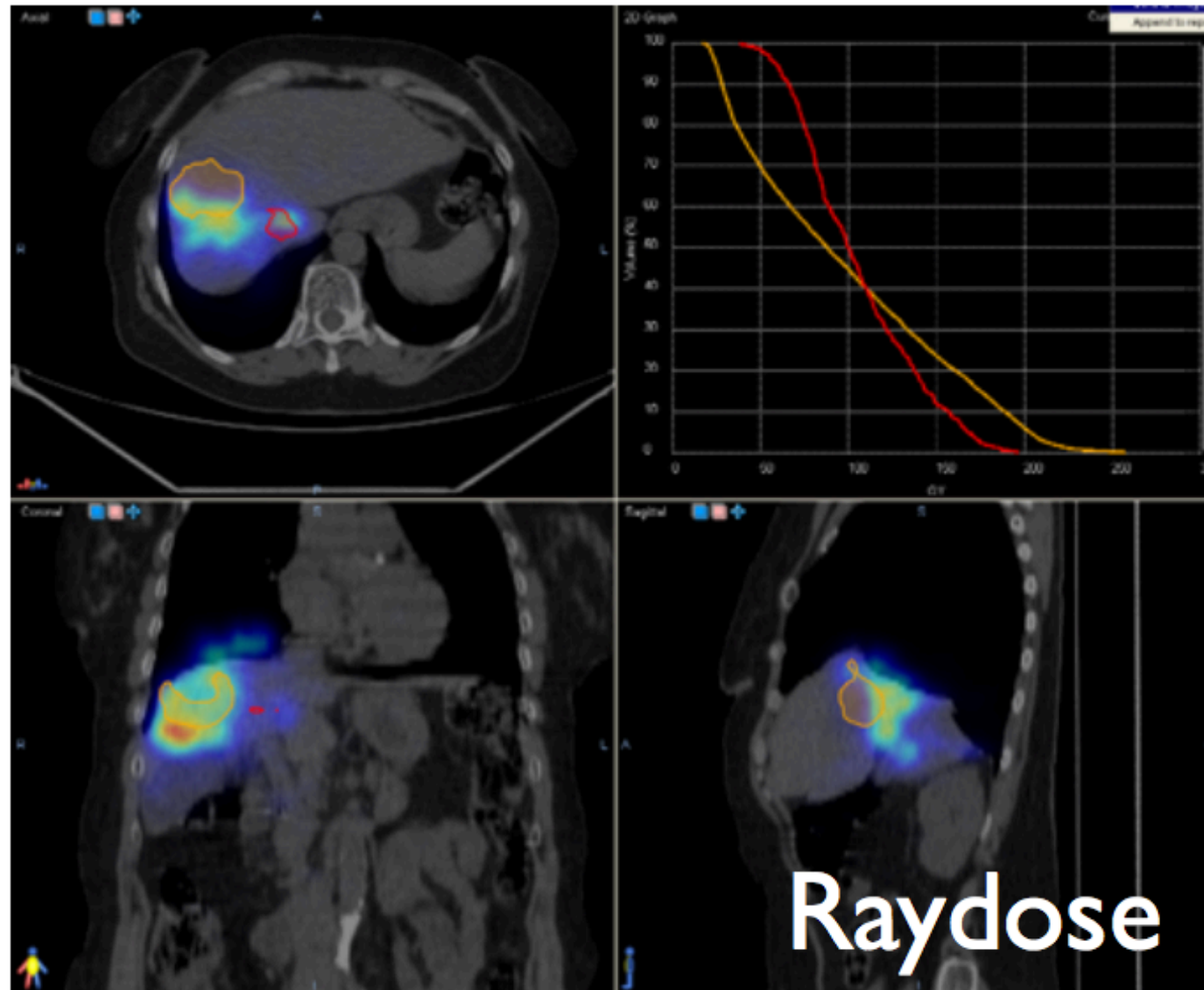
# APPLICATION: LIPIOCIS™



- 194 x 140 x 90 voxels
- (2.21 x 2.21 x 4.42 mm<sup>3</sup>)
- Organ: 45 min ( $\sigma < 2\%$ )
- Voxel: 3.8 d ( $\sigma < 10\%$ )

S Chiavassa et al. (2006) PMB 51:601-616

# Monte Carlo based dosimetry

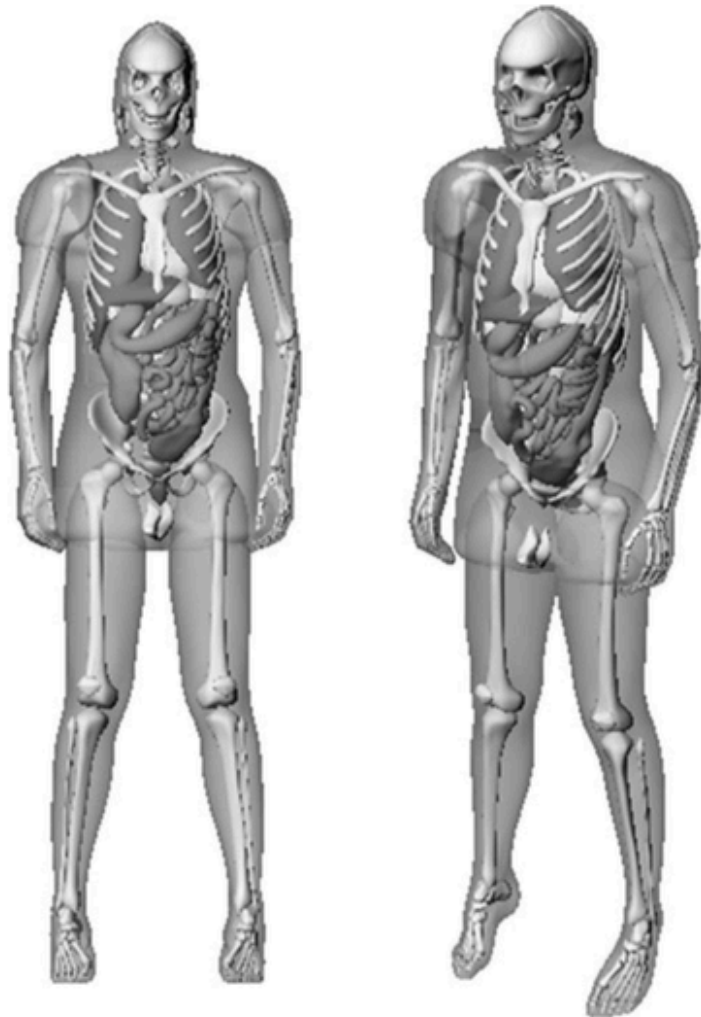


Courtesy: E Spezi (Velindre, Cardiff)

Marcatili et al. Phys Med Biol 2013 58 2491-2508



# Monte Carlo based dosimetry



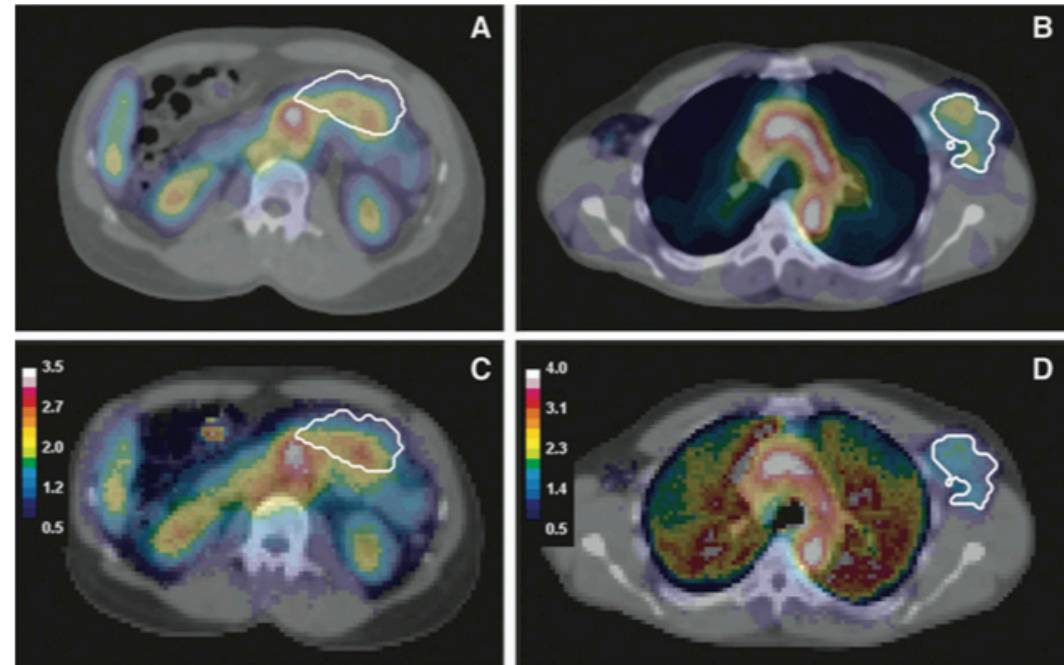
**FIG. 3.** Anterior views of the RADAR adult male NURBS phantom. NURBS, Non-Uniform Rational B-Spline; RADAR, Radiation Dose Assessment Resource.

CANCER BIOTHERAPY AND RADIOPHARMACEUTICALS  
Volume 30, Number 1, 2015  
© Mary Ann Liebert, Inc.  
DOI: 10.1089/cbr.2014.1713

**Original Article**

## VIDA: A Voxel-Based Dosimetry Method for Targeted Radionuclide Therapy Using Geant4

Susan D. Kost,<sup>1</sup> Yuni K. Dewaraja,<sup>2</sup> Richard G. Abramson,<sup>3</sup> and Michael G. Stabin<sup>3</sup>



**FIG. 5.** Fused SPECT/CT images for patient 1 (A) and patient 2 (B) with matching 3D dose maps overlaid on CT for patient 1 (C) and patient 2 (D). The dose maps are displayed in units of Gy. Color images available online at [www.liebertpub.com/cbr](http://www.liebertpub.com/cbr)

# Dosimetry for MRT:

Group	Model	Model ICRP - MIRD DER
Specific	Model $\pm$ adjusted	Model $\pm$ realistic
Specific	Specific	Specific

- *Patient-specific dosimetry: ALL steps must be patient-specific*



# Conclusion

- Patient-specific dosimetry is feasible
- Huge literature in quantitative imaging/absorbed dose calculation (the methodology is there!)
- Patient-specific dosimetry requires ALL steps to be patient-specific
- BUT the biological/clinical end-point conditions the kind of approach that needs to be implemented!

# Acknowledgements

- L Ferrer (CLCC & CHU, Nantes)
- Glenn Flux (ICR/RMH, Sutton)
- EANM Dosimetry & Therapy Committees

Special issue:

«Dosimetry in nuclear medicine therapy»

*The Quarterly Journal of Nuclear Medicine  
and Molecular Imaging* 55(1-2), 2011



manuel.bardies@inserm.fr

**Thank you :-)**

# Radiopharmaceutical dosimetry: Introduction & MIRD scheme



**Manuel Bardières, UMR 1037/UPS, Toulouse**  
**[manuel.bardies@inserm.fr](mailto:manuel.bardies@inserm.fr)**

# Nuclear Medicine Dosimetry

- For many years: diagnostic only

  - For new radiopharmaceuticals

- $^{131}\text{I}$  Thyroid therapy

- Targeted Radionuclide Therapy (or MRT)

  - 🔧 Radioimmunotherapy (RIT)

  - 🔧 mIBG, PRRT,

  - 🔧 Bone pain palliation agents (Xofigo™), etc...

  - 🔧 Microspheres (SirSpheres/TheraSpheres),

# Nuclear Medicine Dosimetry

## ○ Diagnostic procedures

- 🔹 Low amount of radiation
- 🔹 Stochastic effects of radiations
- 🔹 Radiation safety (ALARA)

## ○ Therapeutic procedures

- 🔹 Deterministic effects
- 🔹 Normal (critical) organ absorbed dose
- 🔹 Tumour absorbed dose



# Therapy vs. Diagnostic

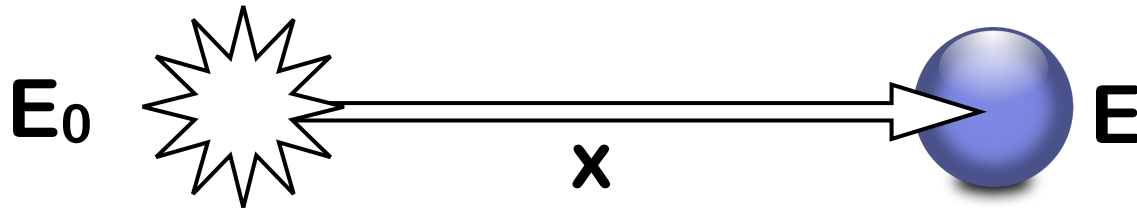
- The goals are NOT the same
  - The required accuracy is NOT the same
- For diagnostics: an estimate is OK
- For therapy:
  - Comparative studies
  - Increase treatment efficacy/toxicity ratio
    - Pre-therapeutic study or during the treatment
  - Patient follow-up (absorbed dose accumulation)
    - Absorbed dose - effect relationship?

# The MIRD Scheme

**MIRD = Medical Internal Radiation Dose Committee**

- **Committee of the Society of Nuclear Medicine (USA)**
    - **Mix group (physicians + physicists)**
  - **Publication via the SNM (JNM):**
    - **25 Pamphlets**
    - **20 Dose estimate reports**
    - **3 Books**
- from 1968 to now...**
- **No web server (see [www.snm.org](http://www.snm.org))**
  - **Main achievement: a global formalism for absorbed dose calculations in Nuclear Medicine**

# Formalism



$$\phi(x, E_0) = \frac{E}{E_0}$$

**AF: Absorbed Fraction, dimensionless**

$$\Phi(x, E_0) = \frac{\phi(x, E_0)}{dm}$$

**SAF: Specific Absorbed Fraction, in  $g^{-1}$**

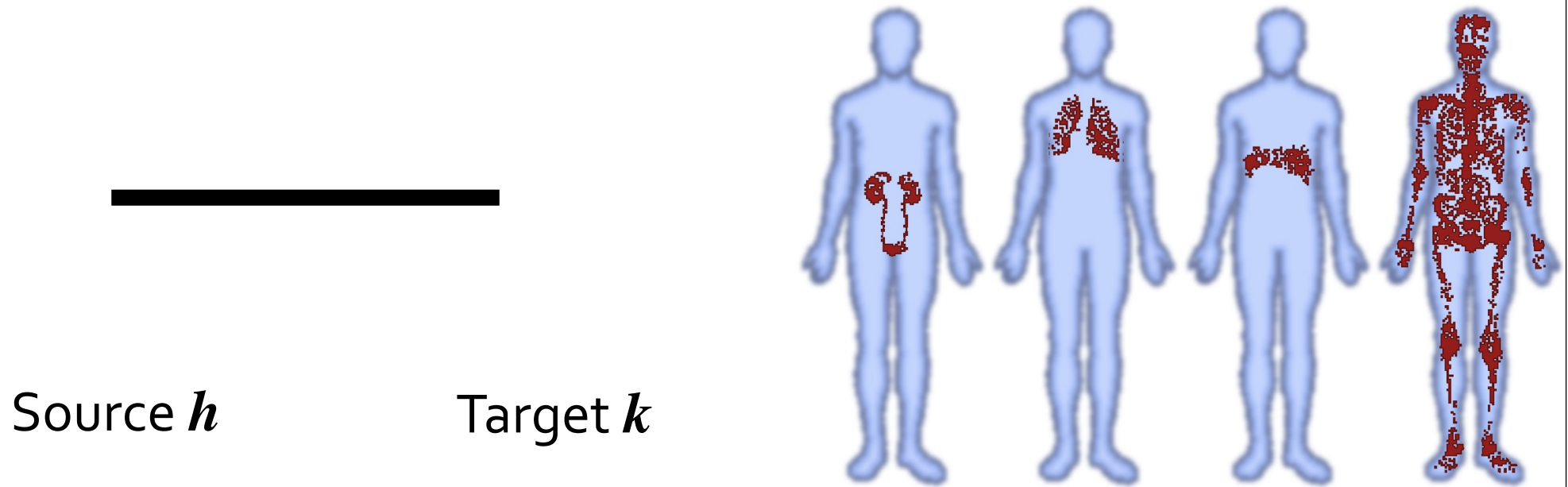
# Absorbed dose definition

$$\bar{D} = \frac{E}{dm} = \frac{\phi(x, E_0) \cdot E_0}{dm} \quad \bar{D} = \Phi(x, E_0) \cdot E_0$$

**Mean absorbed dose in Gy (J/kg)**

**This is obtained without simplifying hypothesis,  
Always true!**

# Volume Generalisation



$$\bar{D}(k \leftarrow h) = \frac{E}{m_k} = \frac{\phi(k \leftarrow h) \cdot E_0}{m_k} = \Phi(k \leftarrow h) \cdot E_0$$

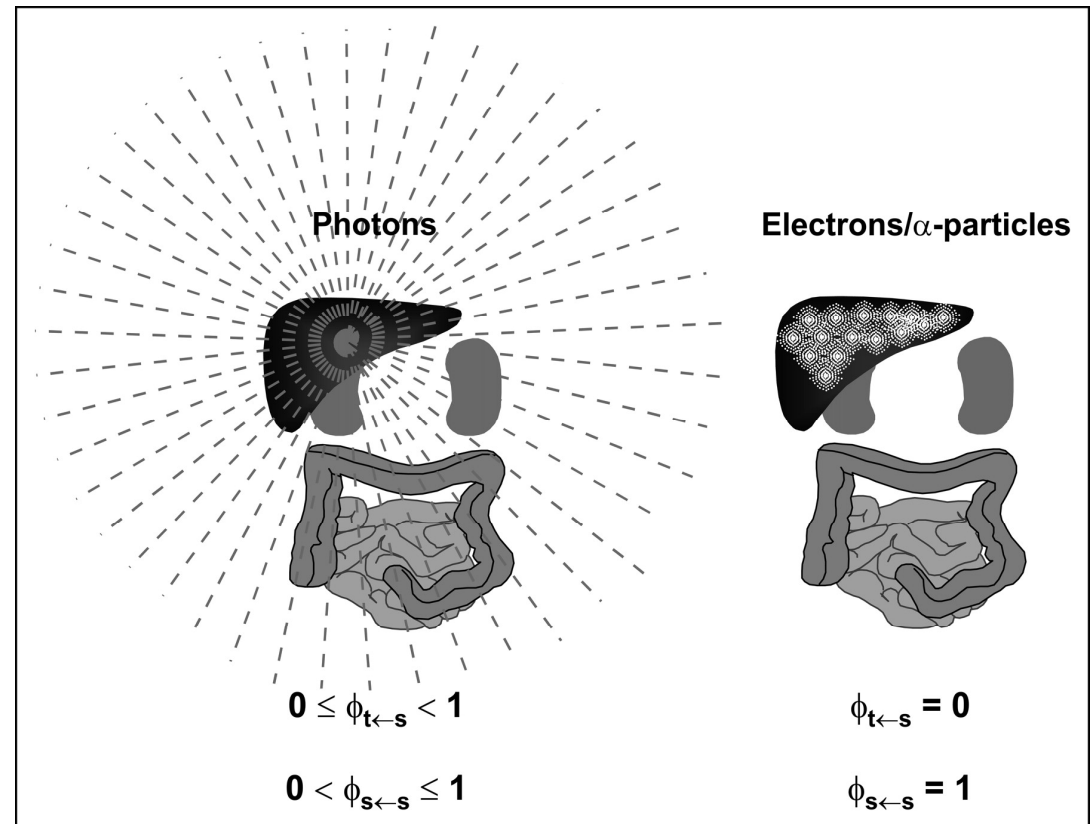
$\bar{D}$  Mean Absorbed Dose over target volume

# Non penetrating radiation

Depends on:  
 Organ size  
 Particle range

$$\phi_i(k \leftarrow h) = 0 \text{ if } k \neq h$$

$$\phi_i(k \leftarrow h) = 1 \text{ if } k = h$$



$$\bar{D}(k \leftarrow k) = \frac{\phi(k \leftarrow k) \cdot E_0}{m_k} = \frac{E_0}{m_k}$$

$$\bar{D}(k \leftarrow h) = 0$$

# Radionuclide generalisation

The absorbed dose rate is the sum of all contributions:

$$\overline{\dot{D}}(t)_{(k \leftarrow h)} = K \cdot A_h(t) \cdot \sum_i n_i E_i \cdot \Phi_i(k \leftarrow h)$$

Sometimes seen as:

$$\overline{\dot{D}}(t)_{(k \leftarrow h)} = A_h(t) \cdot \Delta \cdot \Phi(k \leftarrow h)$$



# Integration over time

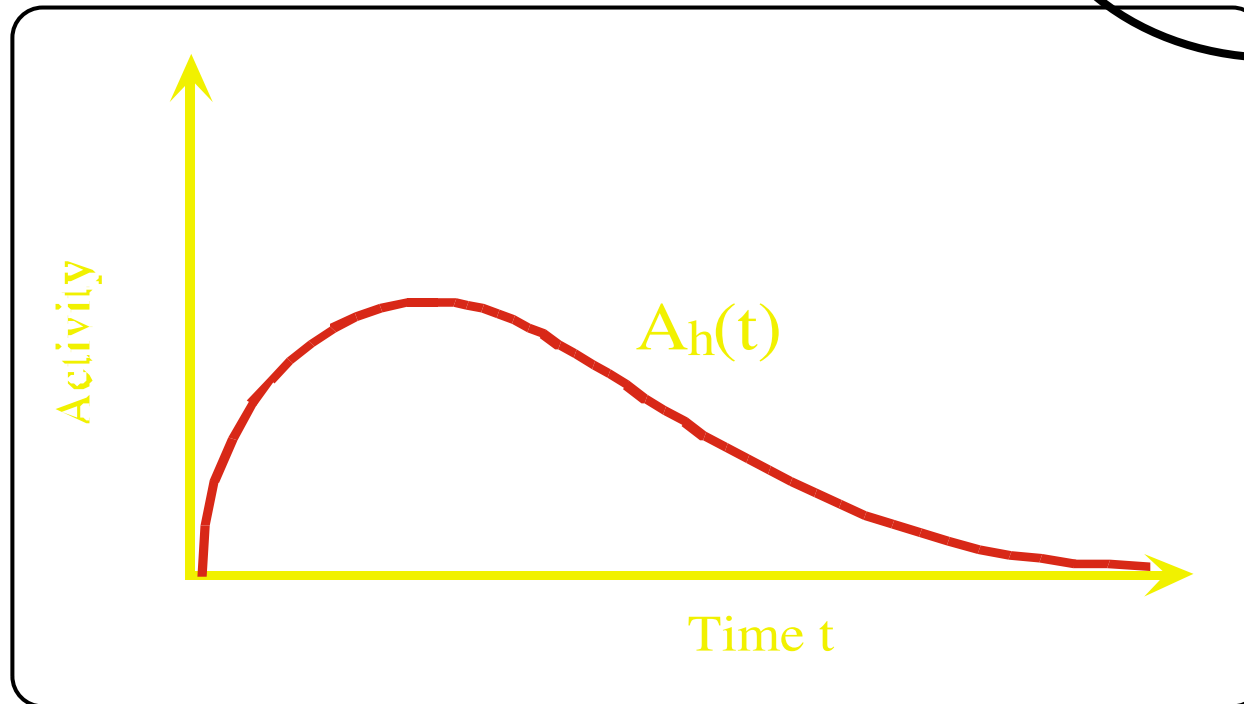
$$\bar{D}_{(k \leftarrow h)} = \int_{t_1}^{t_2} \bar{\dot{D}}(t)_{(k \leftarrow h)} dt$$

$$\bar{D}_{(k \leftarrow h)} = \int_{t_1}^{t_2} K \cdot A_h(t) \cdot \sum_i n_i E_i \cdot \Phi_i(k \leftarrow h) dt$$

$\bar{D}_{(k \leftarrow h)}$  Mean absorbed dose (Gy)  
in target k from source h

# Integration over time (2)

$$\bar{D}_{(k \leftarrow h)} = K \cdot \sum_i n_i E_i \cdot \Phi_i(k \leftarrow h) \cdot \int_{t_1}^{t_2} A_h(t) dt$$



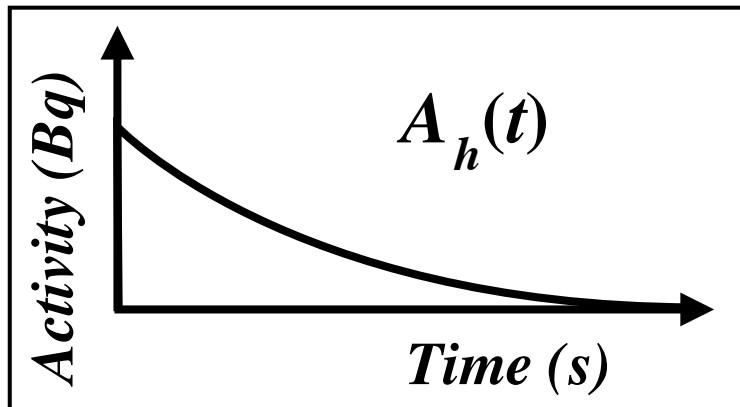
$$\tilde{A}_h = \int A_h(t) dt$$

**Cumulated activity (Bq.s or  $\mu\text{Ci.h}$ )  
'time integral of the activity'**

# Cumulated activity

*Activity detected decreases because:*

- *Vector washout (biological half-life)*
- *Radioactive decay (physical half-life)*



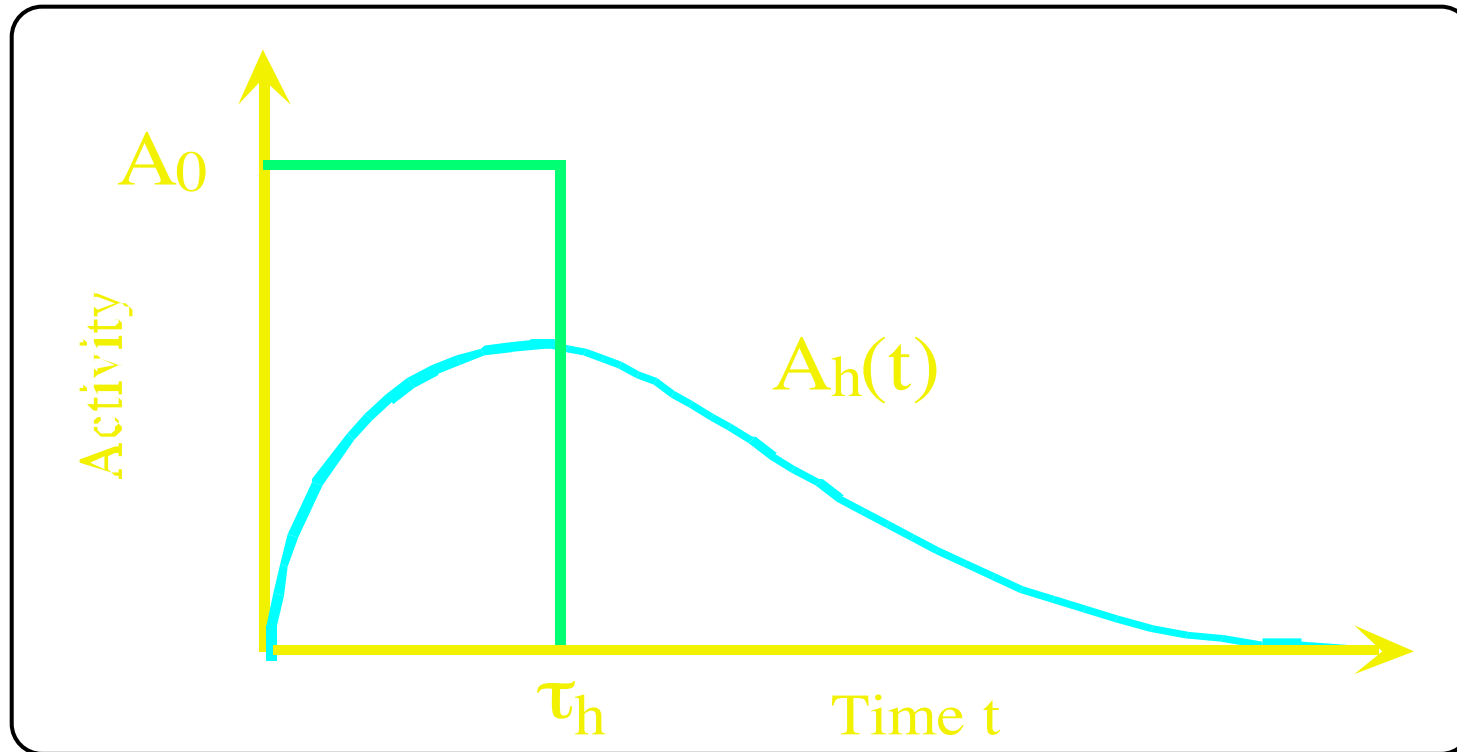
$$A_h(t) = A_0 \times e^{-(\lambda_{phy} + \lambda_{bio})t}$$

$$\tilde{A}_h = 1,443 \times A_0 \times T_{eff}$$

$$\frac{1}{T_{eff}} = \frac{1}{T_{bio}} + \frac{1}{T_{phy}}$$

$T_{eff}$  is the effective half-life

# Residence time: $\tau_h$



$A_0$  is the injected activity

$$\tau_h = \frac{\tilde{A}_h}{A_0}$$

$\tilde{A}_h$  in Bq.s  
 $A_0$  in Bq

}  $\tau_h$  in s

# Fundamental MIRD equation



Source  $h$

Target  $k$

$$\overline{D}_{(k \leftarrow h)} = K \cdot \tilde{A}_h \cdot \sum_i n_i E_i \cdot \Phi_i(k \leftarrow h)$$

**Summary: mean absorbed dose (Gy)**

**Source  $h$**

**Target  $k$**

**$\tilde{A}_h$  nuclear transitions in source  $h$  (Bq.s)**

# Simplified MIRD equation

$$\bar{D}_{(k \leftarrow h)} = K \cdot \tilde{A}_h \cdot \sum_i n_i E_i \cdot \Phi_i(k \leftarrow h)$$

Group all terms independent of time:

$$S_{(k \leftarrow h)} = K \cdot \sum_i n_i E_i \cdot \Phi_i(k \leftarrow h)$$

**MIRD Simplified Equation:**

$$\bar{D}_{(k \leftarrow h)} = \tilde{A}_h \cdot S_{(k \leftarrow h)} \quad \text{or:} \quad \frac{\bar{D}_{(k \leftarrow h)}}{A_0} = \tau_h \cdot S_{(k \leftarrow h)}$$

## Absorbed dose calculation:

$$\overline{D}_{(k \leftarrow h)} = \tilde{A}_h \cdot S_{(k \leftarrow h)}$$

- Determination of  $\tilde{A}_h$ 
  - 📌 Quantitative imaging
  - 📌 TAC fitting
- Use the relevant S factor
  - 📌 Absorbed dose calculations





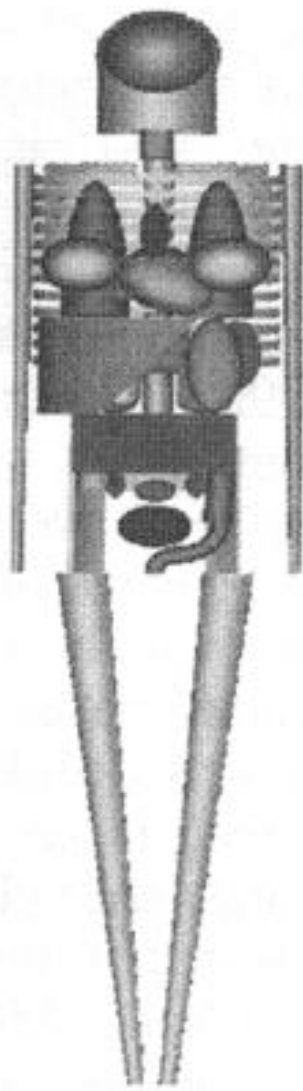
# Work of the MIRD committee

- Calculation scheme for radiopharmaceutical dosimetry
- S value calculations
  - For several radionuclides
  - For several geometries
  - Using anthropomorphic phantoms
- MIRD pamphlet 11

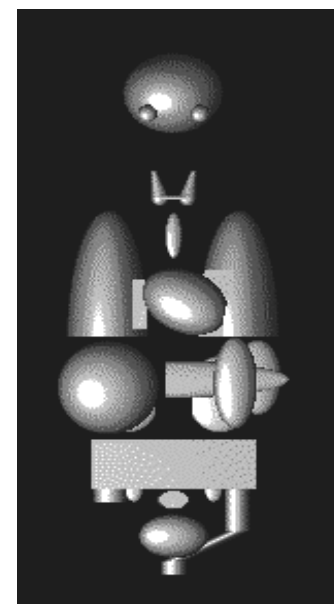
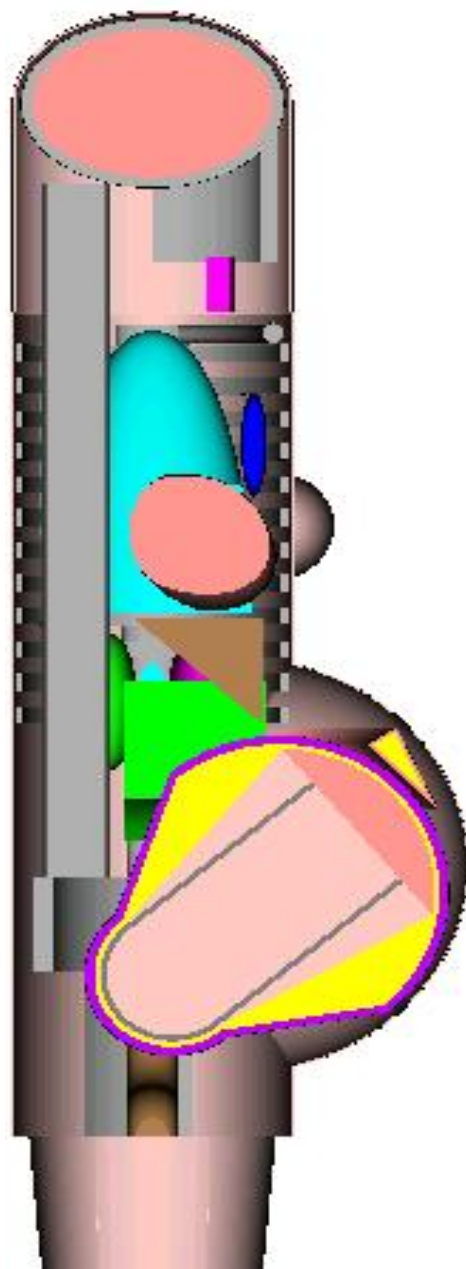
# Anthropomorphic phantoms



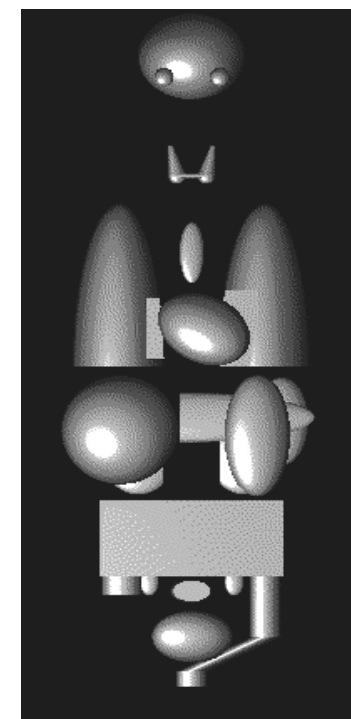
(a)



(b)

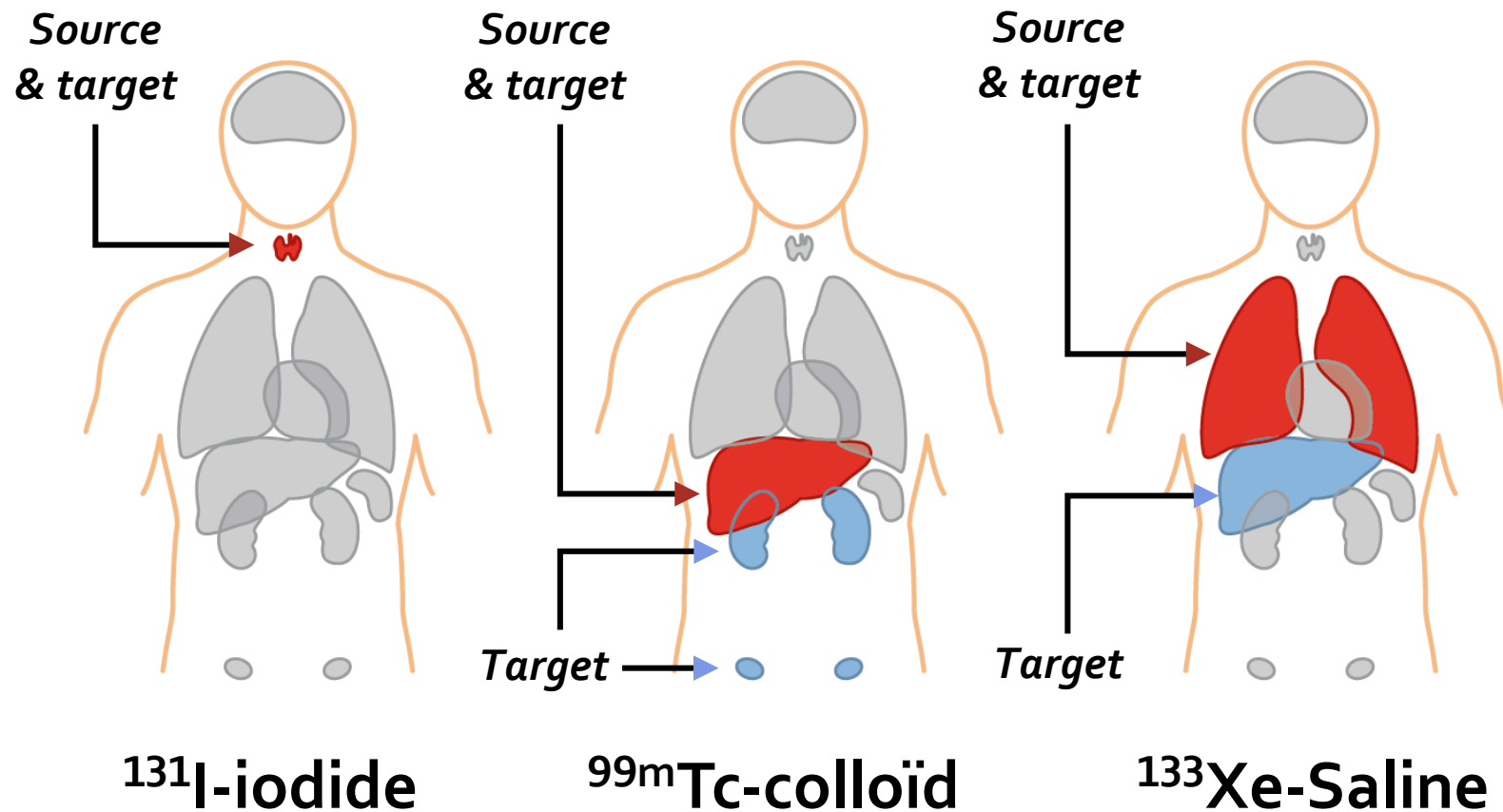


150 cm



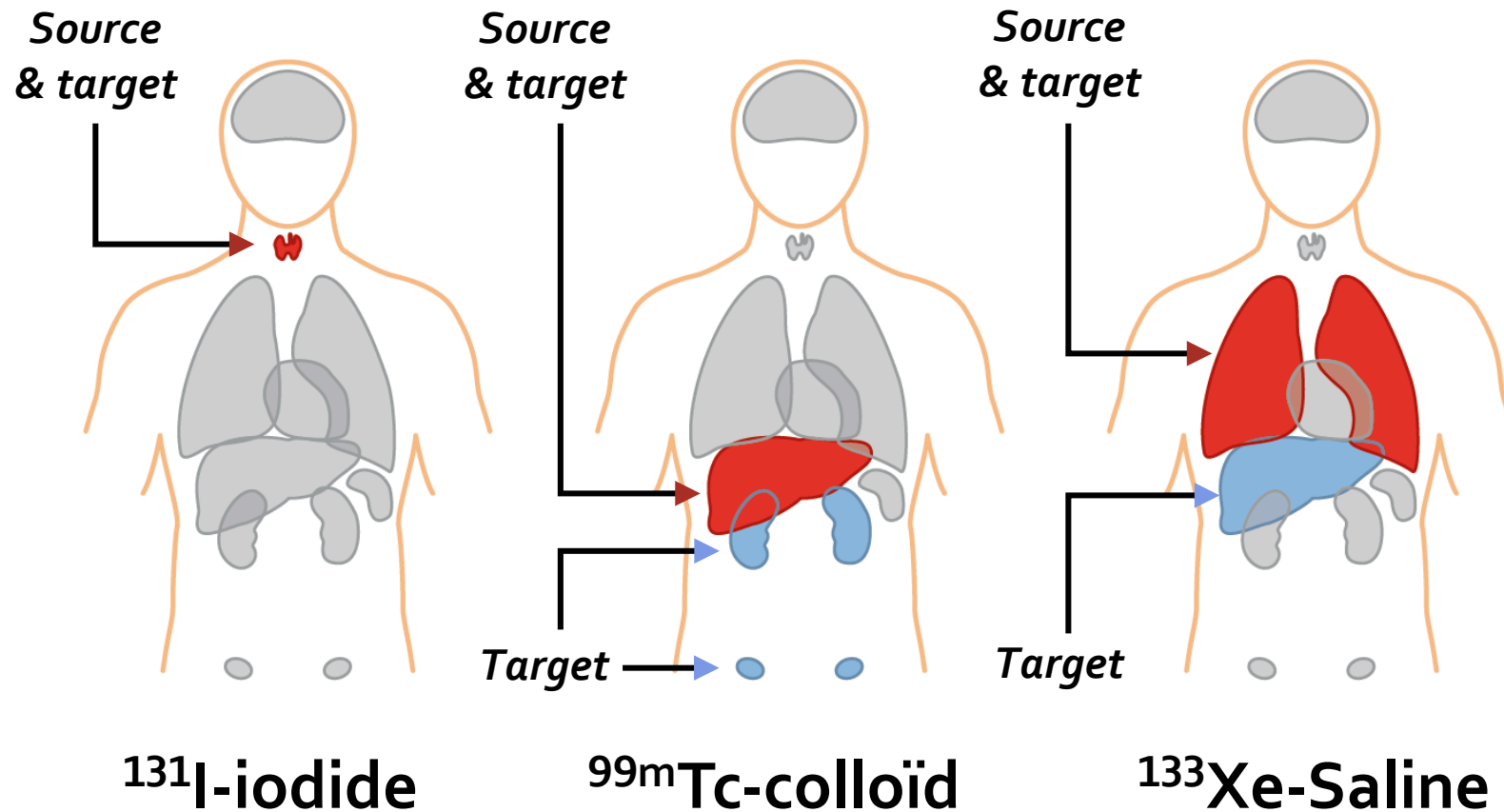
170 cm

# Using the MIRD scheme



In «real» life: there will be different radiation sources,  
And absorbed dose needs to be calculated for  $\neq$  targets...  
Depends on the application...

# Using the MIRD scheme



$$\bar{D}_k = \sum_h \tilde{A}_h \times S_{(k \leftarrow h)}$$

# MIRD Pamphlet 21: New nomenclature

**TABLE 1.** Quantities, Parameters, Symbols, and Units Used in the MIRD and ICRP Dosimetry Schema (Listed in Order of Appearance in Equations 1–17)

Quantity or parameter	MIRD Pamphlet 21	MIRD Primer (1991) (4)	ICRP publications (7,8,18)	Units or special name
Source region (or tissue)	$r_S$	$r_h$	$S$	
Target region (or tissue)	$r_T$	$r_k$	$T$	
Absorbed dose rate to target region	$\dot{D}(r_T, t)$	$\dot{D}(r_k)$ or $\dot{D}_k$	$\dot{D}_{T,R}$	Gy s <sup>-1</sup>
Activity in source region	$A(r_S, t)$	$A_h(t)$	$q_S(t)$	Bq
Absorbed dose rate per unit activity	$S(r_T \leftarrow r_S, t)$	$S(r_k \leftarrow r_h)$	Not defined	Gy (Bq s) <sup>-1</sup>
Dose-integration period	$T_D$	Assumed to be $\infty$	$\tau$	s
Absorbed dose to target	$D(r_T, T_D)$	$\bar{D}(r_k)$ or $\bar{D}_k$	$D_{T,R}$	Gy
Administered activity	$A_0$	$A_0$	$q_0$	Bq
Fraction of administered activity in the source region	$a(r_S, t) = A(r_S, t)/A_0$	$f_h(t)$	Not defined	Unitless
Absorbed dose coefficient	$d(r_T, T_D)$	Not defined	$d_T(\tau)$	Gy Bq <sup>-1</sup>
Mean energy of the i <sup>th</sup> transition	$E_i$	$E_i$	$E_i$	J or MeV
Number of i <sup>th</sup> transitions per nuclear transformation	$Y_i$	$n_i$	$Y_i$	(Bq s) <sup>-1</sup>
Mean energy of the i <sup>th</sup> transition per nuclear transformation	$\Delta_i$	$\Delta_i$	$\Delta_i$	J (Bq s) <sup>-1</sup> or MeV (Bq s) <sup>-1</sup>
Absorbed fraction	$\phi(r_T \leftarrow r_S, E_i, t)$	$\phi(r_k \leftarrow r_h)$	$AF(T \leftarrow S, E_i)$	Unitless
Mass of target region	$M(r_T, t)$	$m_k$	$m_T$	kg
Specific absorbed fraction	$\Phi(r_T \leftarrow r_S, E_i, t)$	$\Phi(r_k \leftarrow r_h)$	$SAF(T \leftarrow S, E_i)$	kg <sup>-1</sup>

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Quantity or parameter	MIRD Pamphlet 21	MIRD Primer (1991) (4)	ICRP publications (7,8,18)	Units or special name
Time-integrated activity in source region*	$\tilde{A}(r_S, T_D)$	$\tilde{A}_h$	$U_S$	Bq s
Time-integrated activity coefficient†	$\tilde{a}(r_S, T_D)$	$\tau$	Not defined	s
Equivalent dose to target	$H(r_T, T_D)$	Not defined	$H_T$	Sv
Radiation weighting factor	$w_R$	Not defined	$w_R$	Unitless
Absorbed dose to target by radiation type R	$D_R(r_T, T_D)$	Not defined	$D_{T,R}$	Gy
Radiation-weighted S	$S_w(r_T \leftarrow r_S, t)$	Not defined	$SEE(T \leftarrow S)$	Sv (Bq s) <sup>-1</sup>
Equivalent dose coefficient	$h(r_T, T_D)$	Not defined	$h_T(\tau)$	Sv Bq <sup>-1</sup>
Effective dose	$E$	Not defined	$E$	Sv

\*This quantity was termed *cumulated activity* in 1991 MIRD Primer.

†This quantity was termed *residence time* in 1991 MIRD Primer.

# Quick discussion on the new nomenclature

$$\bar{D}_{(r_k)} = \tilde{A}_h \cdot S_{(r_k \leftarrow r_h)}$$

$$\bar{D}(r_T, T_D) = \tilde{A}(r_S, T_D) \cdot S(r_T \leftarrow r_S, t)$$

- Explicit mention of irradiation time ( $T_D$ )
- More “ICRP compliant” (radiation weighting factor, Effective Dose,...)

$\tilde{A}(r_S, T_D)$  « Time-integrated activity » vs. « cumulated activity »

$\tilde{a}(r_S, T_D)$  « Time-integrated activity coefficient » vs. « residence time »

**Be careful with the new nomenclature...**



# Conclusion

- **The MIRD FORMALISM is valid for both diagnostics and therapy...**
- **MIRD S values:**
  - 📌 **Impressive database**
  - 📌 **Can be used (for diagnostic) easily (tables)**
  - 📌 **For radiation safety**
  - 📌 **For a model rather than YOUR patient**
- **Therapy requires patient-specific dosimetry:**
  - 📌 **Quantitative imaging ( $\tilde{A}$ )**
  - 📌 **Patient-specific S values**



# Conclusion

## **MIRD formalism $\neq$ MIRD S Factors**

*One can use the MIRD formalism  
AND compute one's OWN S Factors*

*As a consequence: Writing «dosimetry was performed using the MIRD formalism is NOT sufficient!*

**The dosimetric approach should be described:**

**How  $\tilde{A}$  was obtained**

**How S was obtained**

*Cf: EANM Dosimetry Committee Guidance document (2010): «Good practice of clinical dosimetry reporting»*

# Reference books

**MIRD primer for absorbed dose calculations** Loevinger R, Budinger TF and Watson EE, The society of nuclear medicine, N.Y 1988, rev 1991

- Describes the MIRD scheme

**MIRD radionuclide data and decay schemes** Eckerman KF, Endo A, The society of nuclear medicine, N.Y. 2008

- Radionuclide data, particle type, energy, physical half-life, etc...

**MIRD Cellular S Values** Goddu SM, Howell RW, Bouchet LG, Bolch WE et Rao DV. The society of nuclear medicine, N.Y. 1998

- S values for many radionuclides at the cell level