

*IN SEARCH OF
EXOTIC EVENTS FOR PET:
A GAMMASPHERE EXPERIMENT*

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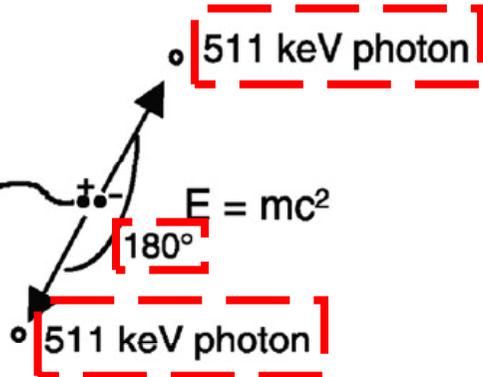
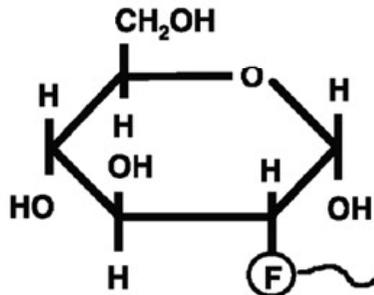
Argonne National Laboratory, 9700 S. Cass Avenue, Argonne IL 60439, USA



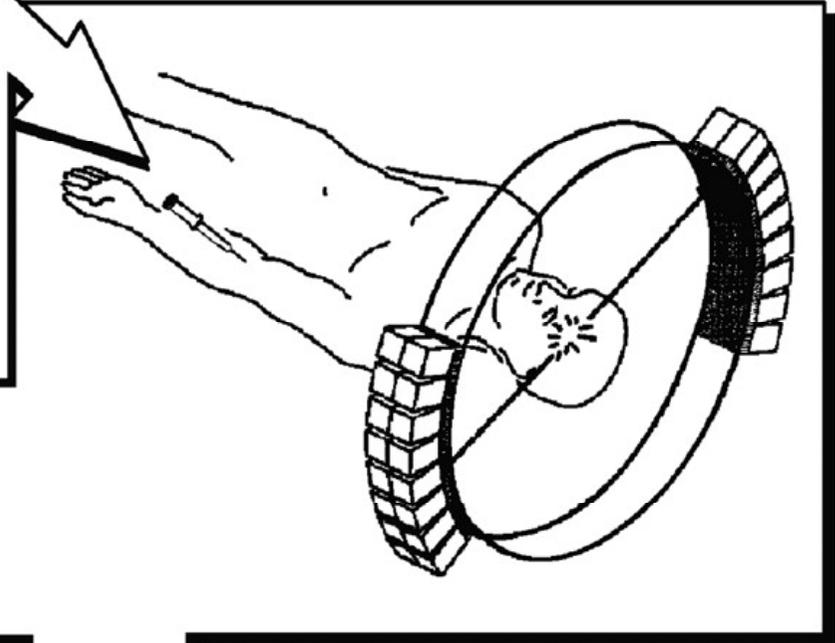
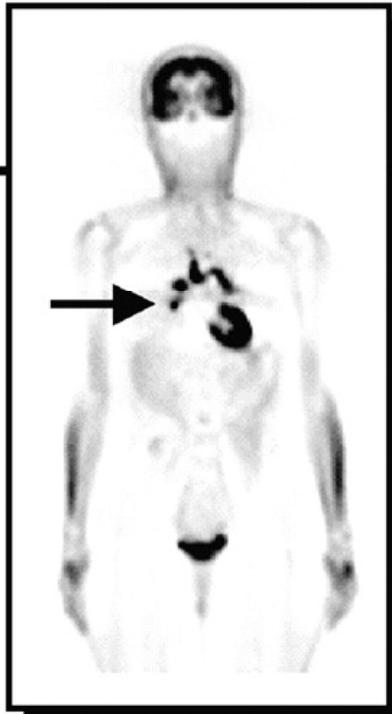
UNIVERSITY OF
SURREY

e⁺ EMITTER: CLINICAL SOURCE

2-[¹⁸F]Fluoro-2-Deoxy-D-Glucose (FDG)



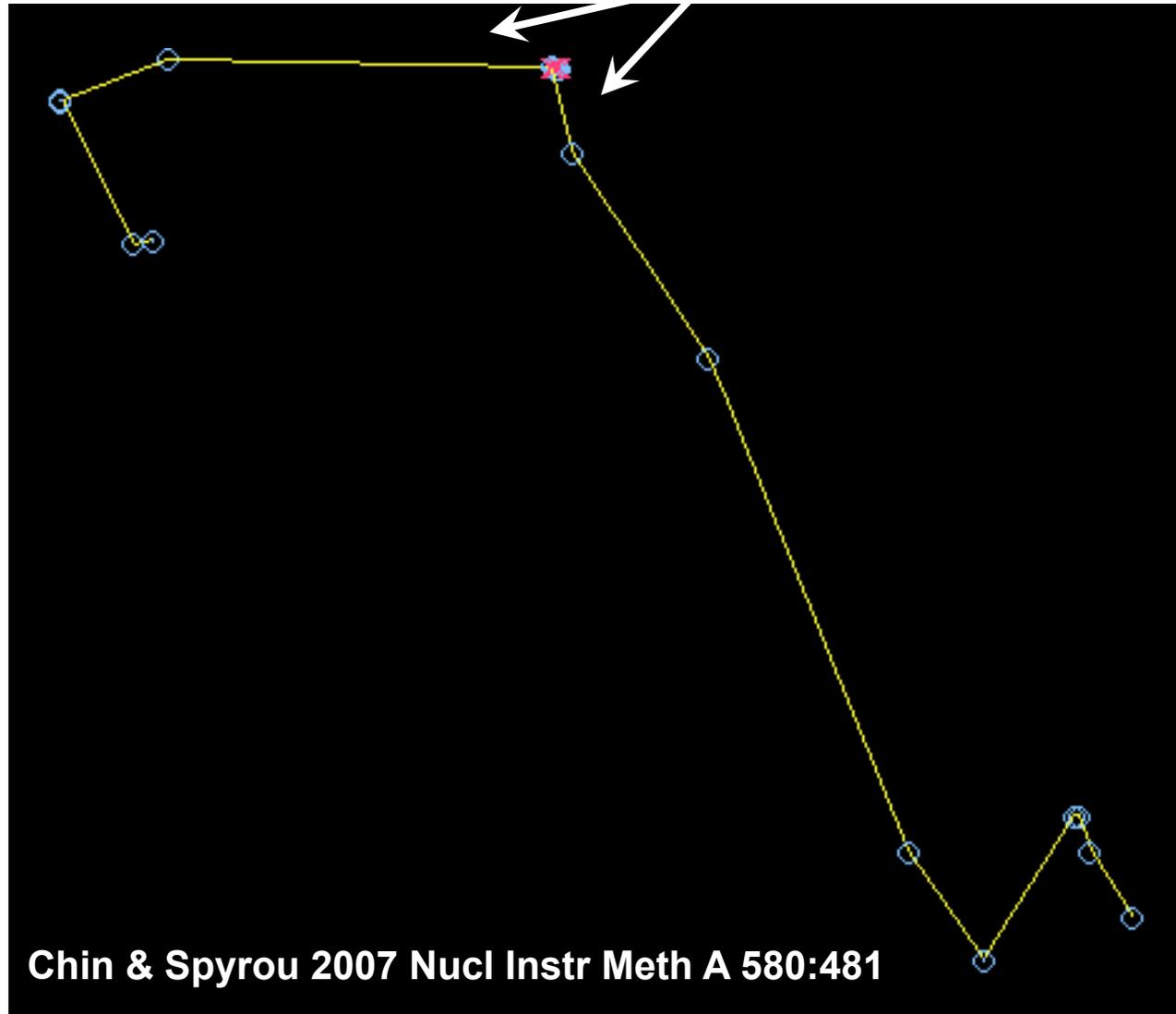
A POSITRON'S LIFE IS FAR MORE EVENTFUL



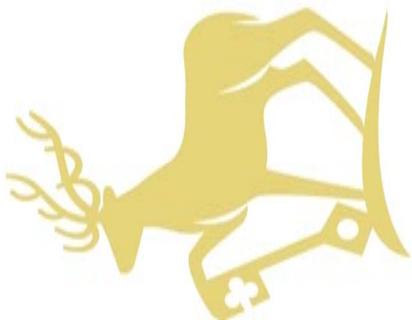
“ PRINCIPLES OF PET ”

Positron about to decay in flight	1	0.606	1	2	0.155,0.023,500.314	-0.122,-0.592, 0.797
Resulting photons	1	0.979	0	2	0.155,0.023 500.314	-0.311, 0.020, 0.950
	2	0.649	0	2	0.155,0.023 500.314	0.282,-0.935,-0.213

NOT 0.511 NOT BACK-TO-BACK!



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IN SEARCH OF
EXOTIC EVENTS FOR PET:
A GAMMASPHERE EXPERIMENT

PREVIOUS WORK (previous slide)

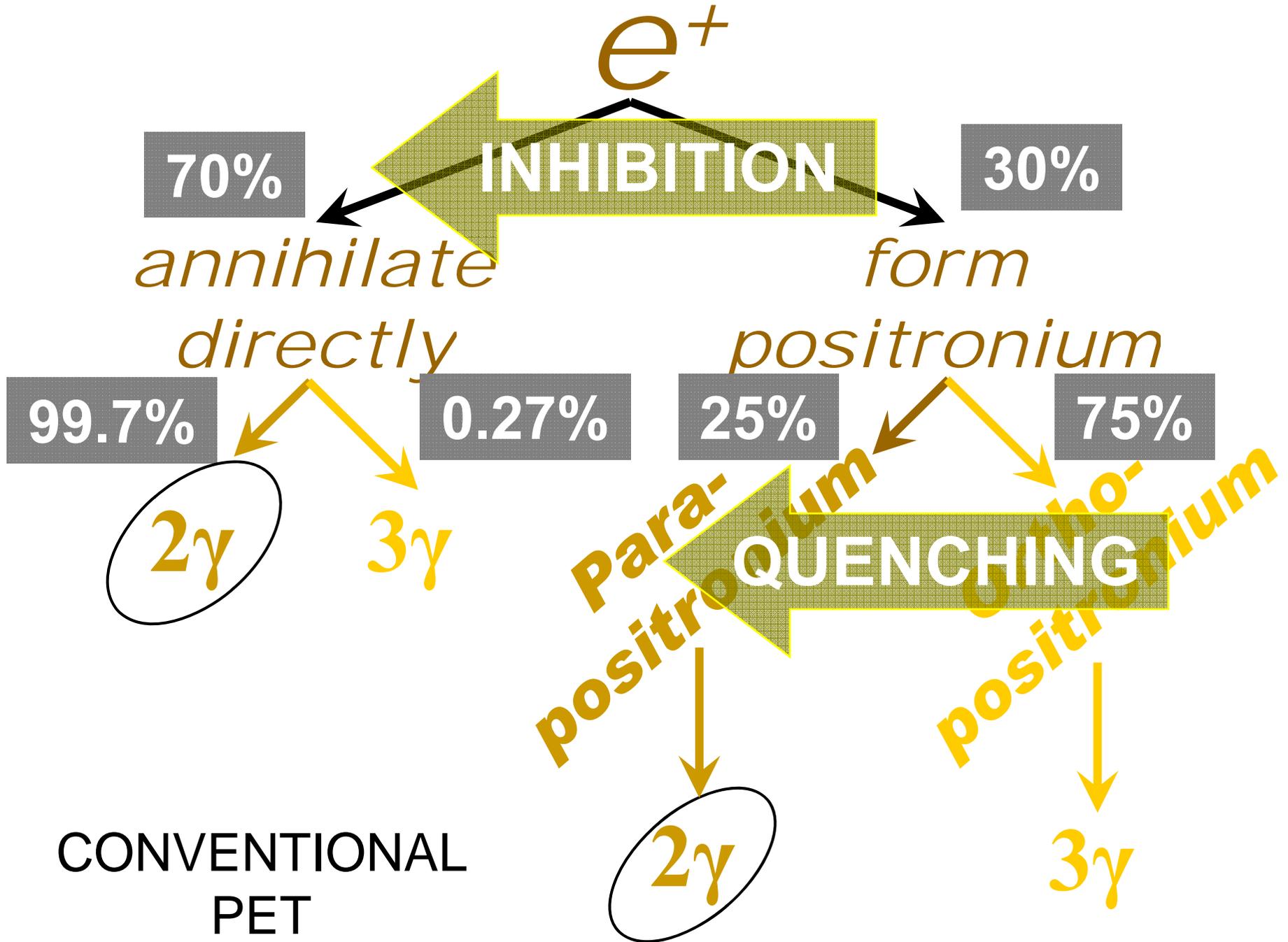
NOT 511 keV & NOT BACK-TO-BACK

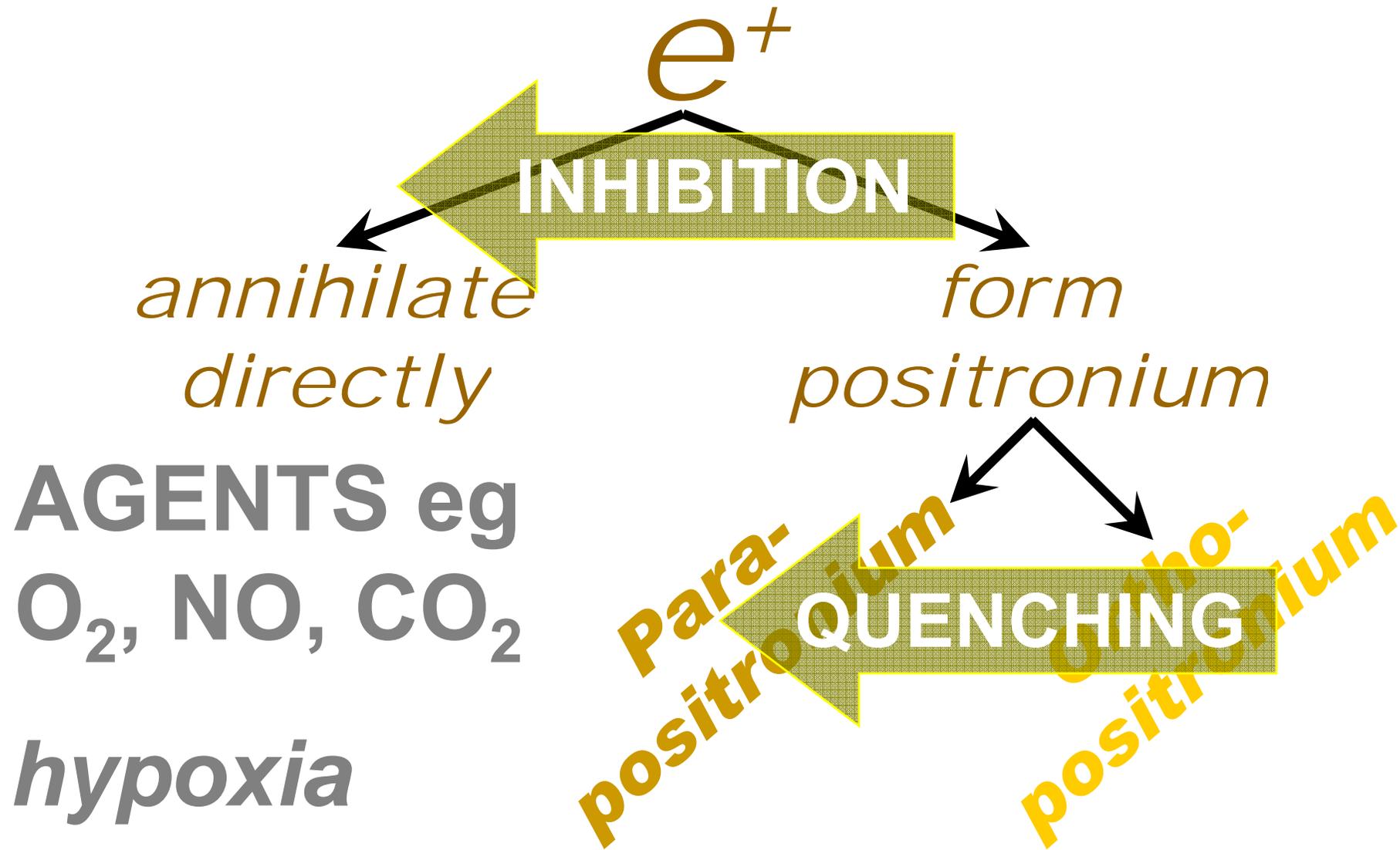
Monte Carlo

THIS WORK

NOT TWO BUT THREE PHOTONS

Real !





PROSPECTS FOR MOLECULAR IMAGING

MEASURE $3\gamma/2\gamma$ YIELD



FIND OUT O_2 IN BODY TISSUES?

**PROSPECTS FOR
MOLECULAR IMAGING**

PROPOSAL

**Three-Gamma Annihilation Imaging in Positron
Emission Tomography**

Krzysztof Kacperski*, Nicholas M. Spyrou, and F. Alan Smith

Abstract—It is argued that positron annihilation into three photons, although quite rare, could still be used as a new imaging modality of positron emission tomography. The information gained when the three decay photons are detected is significantly higher than in the case of 511 keV two-gamma annihilation. The performance of three-gamma imaging in terms of the required detector properties, spatial resolution and counting rates is discussed. A simple proof-of-principle experiment confirms the feasibility of the new imaging method.

Index Terms—Positron emission tomography, three-gamma annihilation.

5. Conclusions

REFUTAL

Three-gamma imaging is potentially more powerful than standard PET because each event bears the complete position information enabling the localization of the activity distribution without use of back-projection tomographic techniques. However, only a subfraction of the three-photon events are usable as each photon energy must be above the detection threshold and only total energy deposition events can be used, as a tight total energy window and time window (<5 ns) must be applied (for typical source strengths used in imaging) to reduce the strong background from two-photon decay pile-up events. However, these conditions can be met with high-resolution semiconductor detectors as pointed out in Kacperski and Spyrou (2005), and construction of a detection system with the required attributes for such studies is not beyond the reach of current technology.

The present work answers one question raised by Kacperski and Spyrou (2005) concerning the potential biological sensitivity of the three-photon imaging. Unfortunately, one should not expect any sensitivity to the level of dissolved O_2 . Our results indicate that the overall three-photon yield is about 0.5% in all samples. These conclusions assume that the direct three-photon yield is identical to that for free e^- s. Only direct measurement of the three-photon yield can determine if this assumption is correct (Seweryniak 2006).

Table 1. The delayed (F_3^{de}) and total (F_3) three-photon yields as well as the fit parameters (K_p and K_{cap}/λ_d) for the various liquid samples (HSA is for human serum albumin).

Material	Oxygen	F_3^{de} (%)	F_3 (%)	K_p (ns ⁻¹)	$R = K_{\text{cap}}/\lambda_d$
Iso-octane	Low	0.58	0.85	1.34	2.15
	High	0.39	0.65	2.27	2.06
Water	Low	0.26	0.52	2.77	3.05
	High	0.25	0.51	2.84	3.07
Saline	Low	0.24	0.51	2.86	3.14
	High	0.24	0.51	2.98	3.07
HSA	Low	0.25	0.51	2.64	3.30
	High	0.25	0.51	2.95	2.96
Blood	Venous	0.25	0.52	2.86	3.02

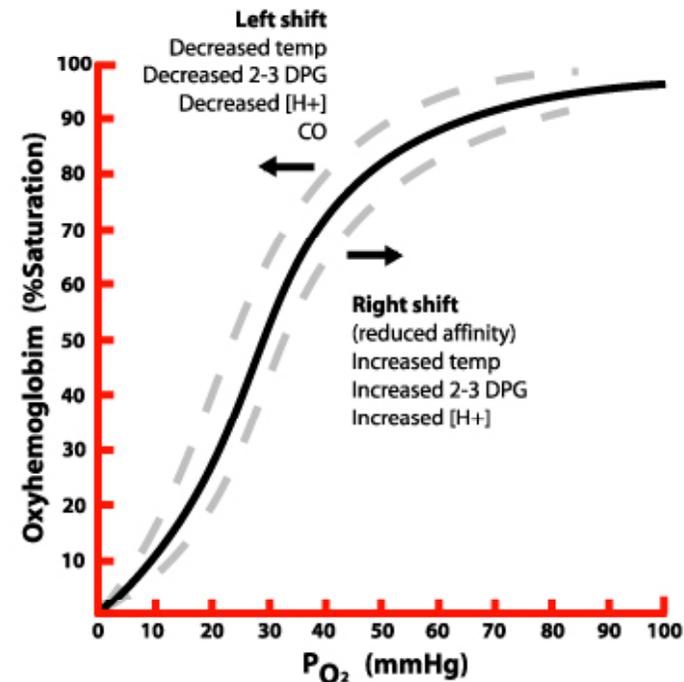
The present work answers one question raised by Kacperski and Spyrou (2005) concerning the potential biological sensitivity of the three-photon imaging. Unfortunately, one should not expect any sensitivity to the level of dissolved O₂. Our results indicate that the overall three-photon yield is about 0.5% in all samples. These conclusions assume that the direct three-photon yield is identical to that for free e⁻s. Only direct measurement of the three-photon yield can determine if this assumption is correct (Seweryniak 2006).

MEASURE $3\gamma/2\gamma$ YIELD



FIND OUT O_2 IN BODY TISSUES?

WHAT DO WE MEAN?
LOOSE OR BOUND?

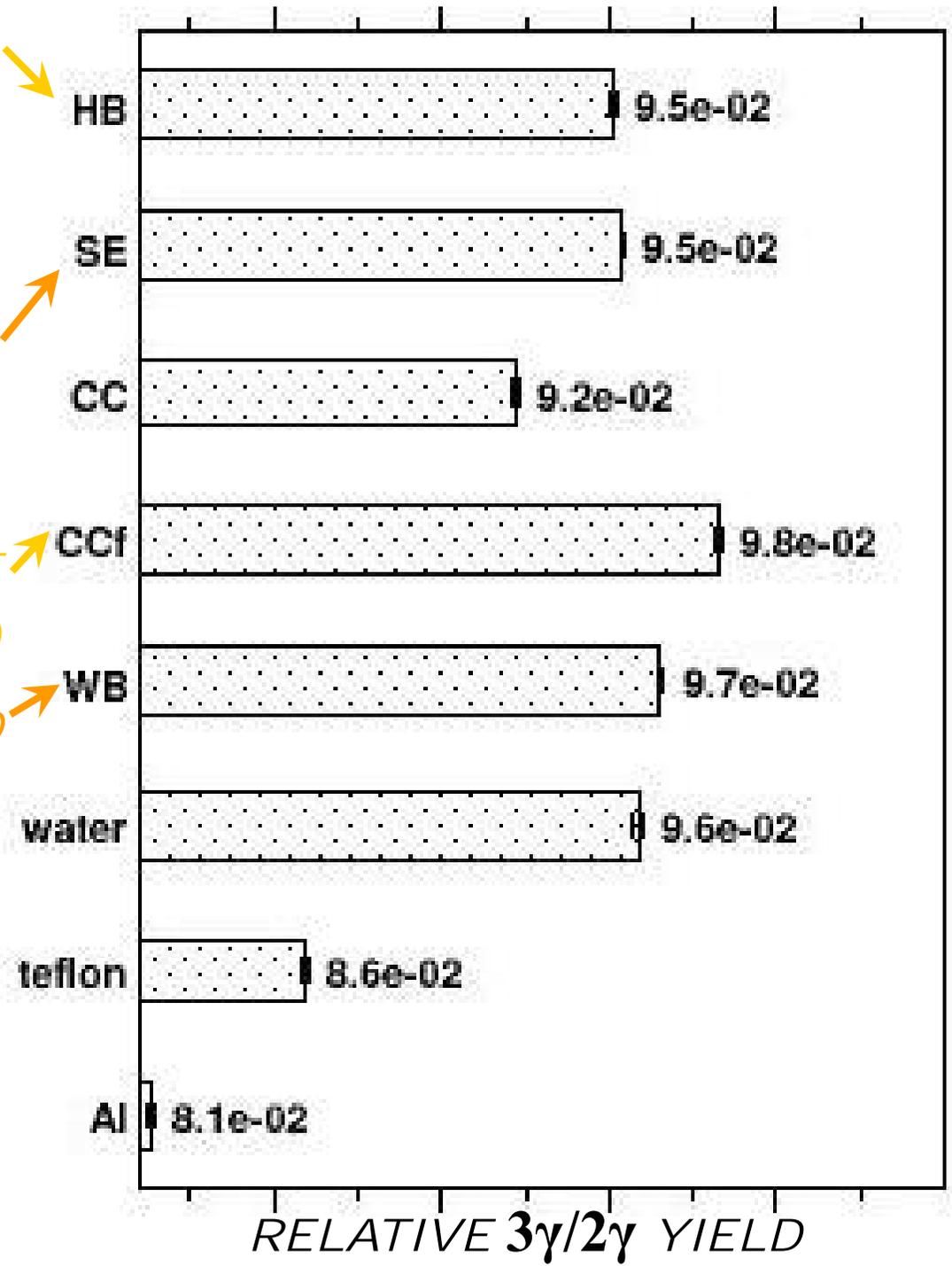


HAEMOLYSED BLOOD
(HAEMOGLOBINS HAVE
BEEN LIBERATED FROM
RED BLOOD CELLS)

SERUM
(LIKE PLASMA BUT NO
CLOTTING FACTORS)

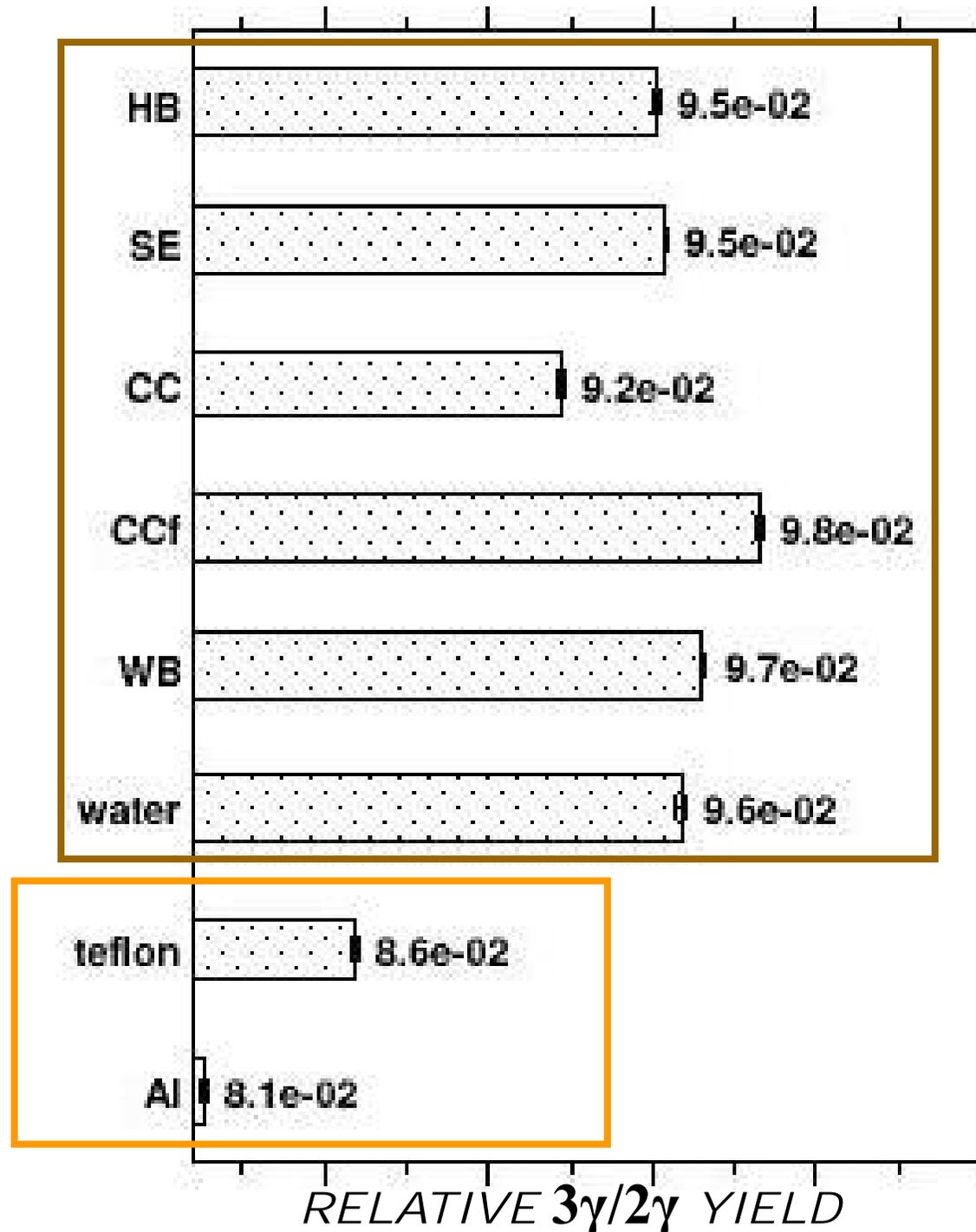
CELL CONCENTRATE
(NO PLASMA)

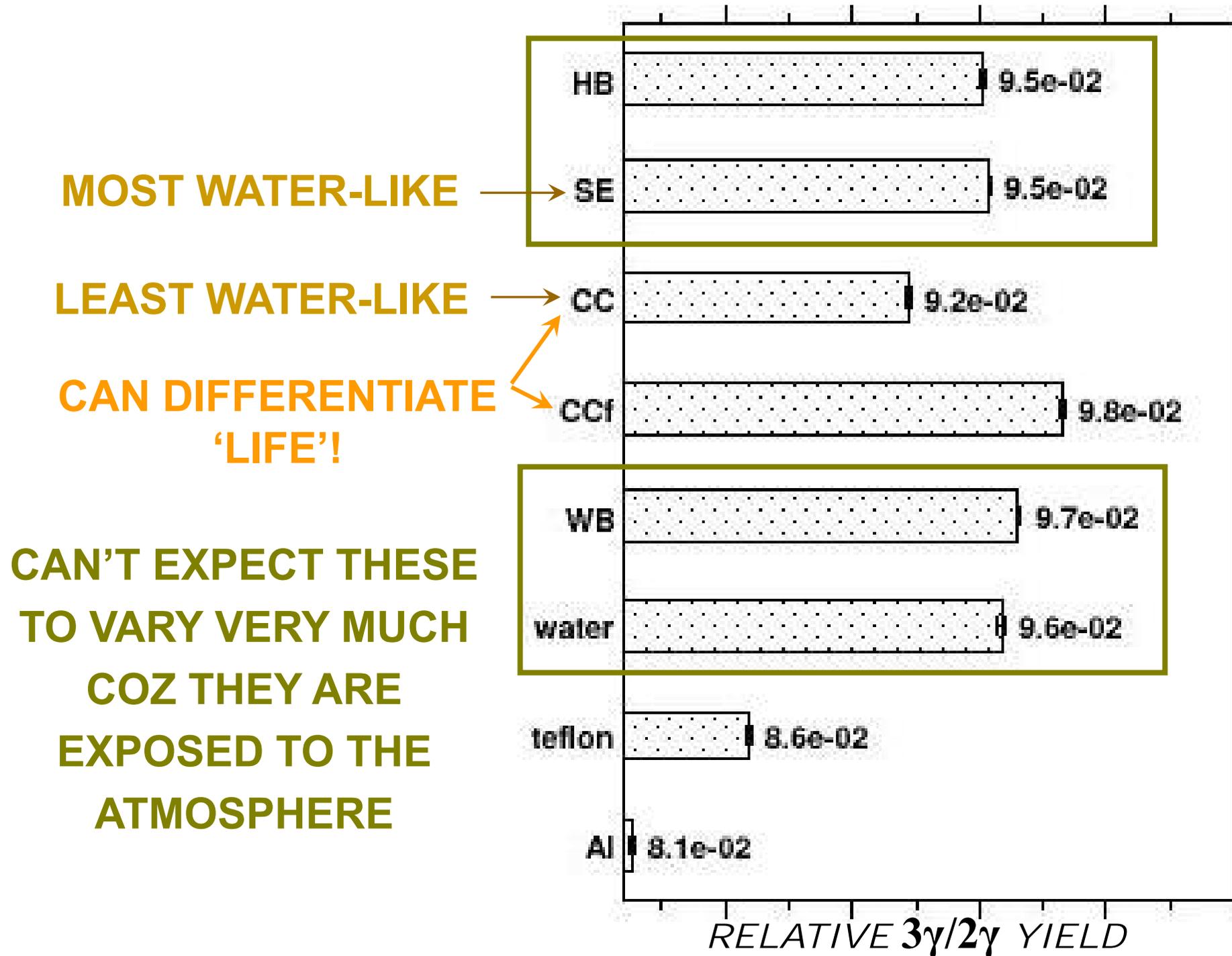
WHOLE BLOOD



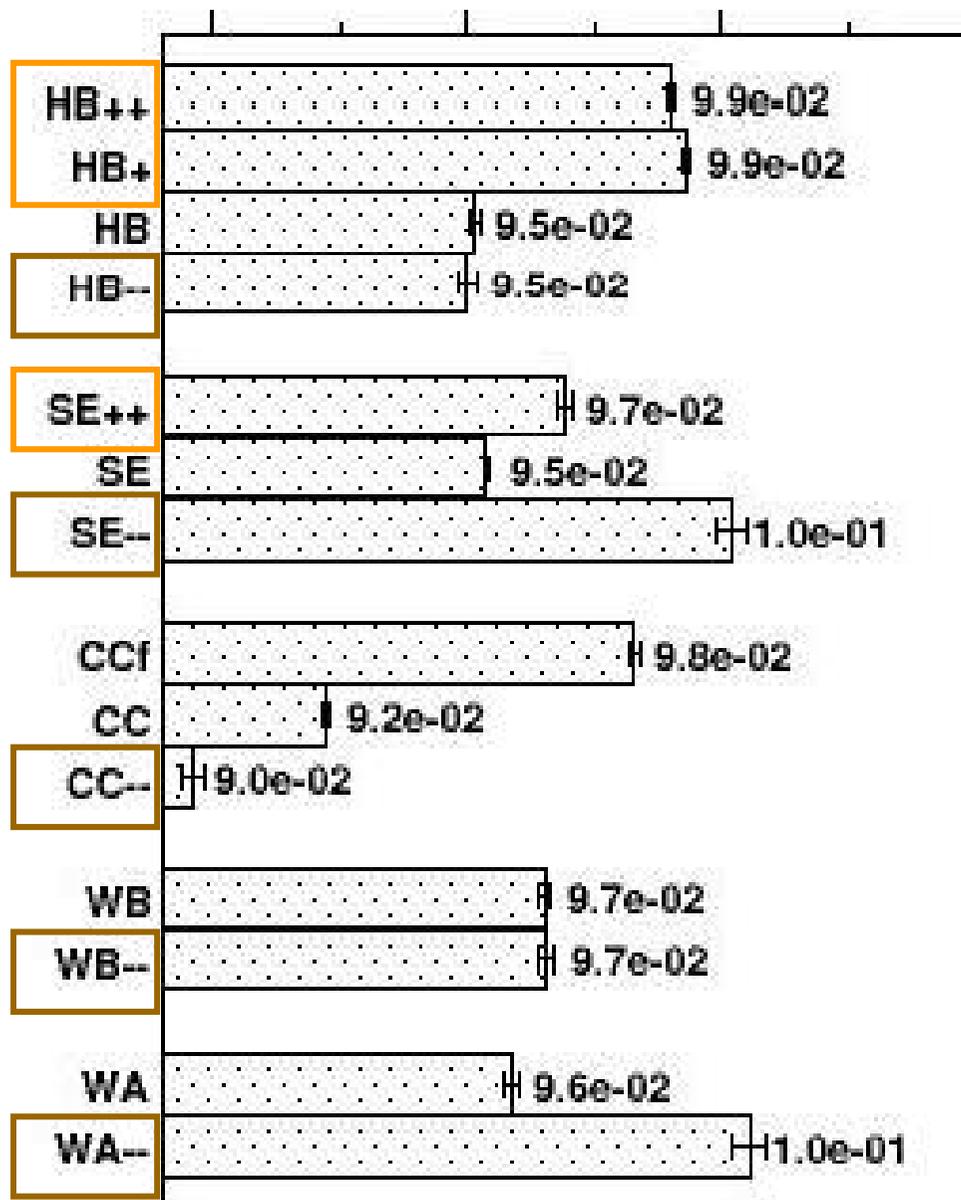
THE FACT THAT
THEY VARY IS
ENCOURAGING – AT
LEAST THE METHOD IS
SENSITIVE TO
SOMETHING

CONFIRMS VALUES IN
THE LITERATURE





OXYGENATED



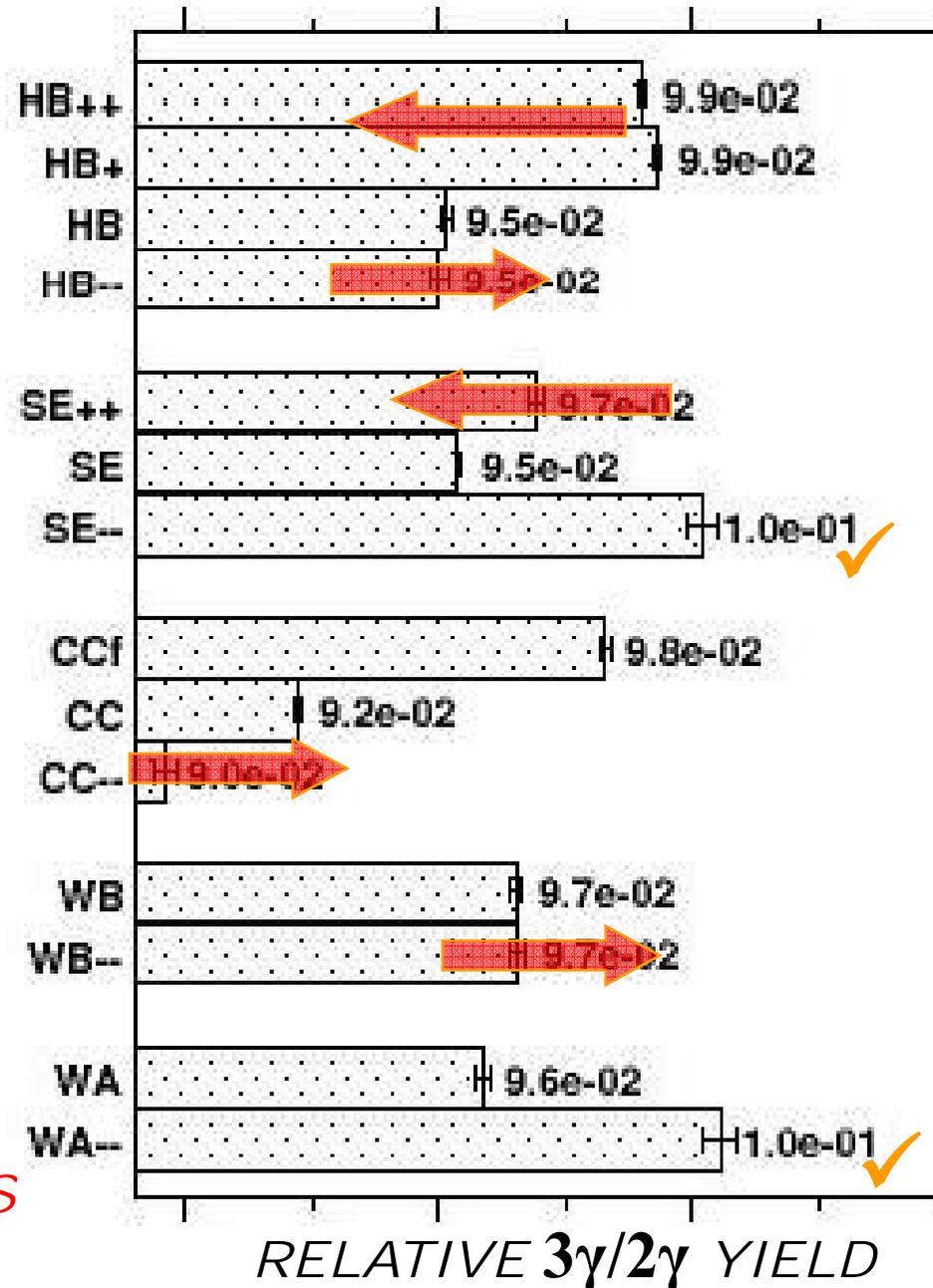
DEOXYGENATED

RELATIVE $3\gamma/2\gamma$ YIELD

IF O₂ WERE THE ONLY FACTOR

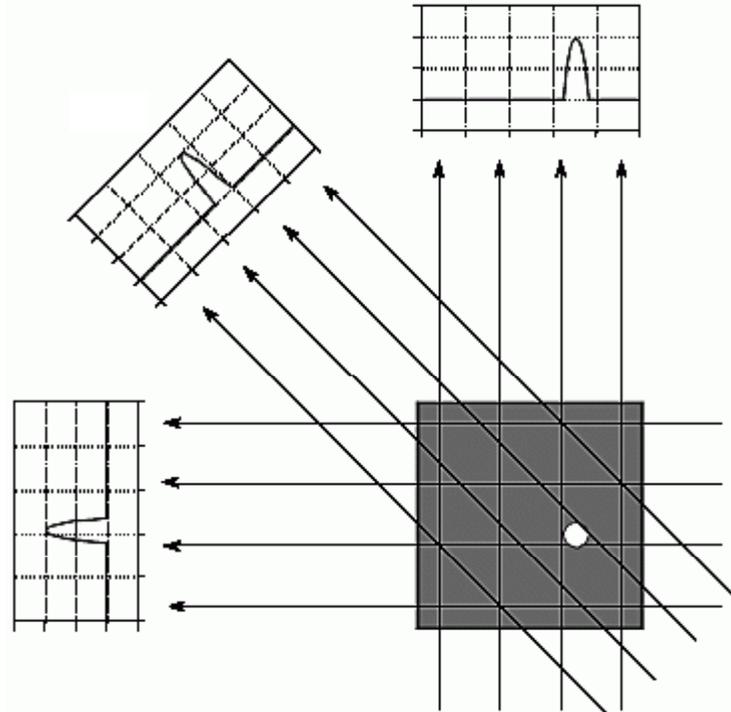
EXPECTED
O₂ ↓ YIELD ↑

UNEXPECTED
CONFOUNDING FACTORS



2 γ PET

IMAGE RECONSTRUCTION



WE HAVE TO BACK-PROJECT
COZ WE DON'T KNOW AT WHICH
POINT THE ANNIHILATION
HAPPENED

LINE OF RESPONSE

3 γ PET

Local chemistry ✓

By conservations of
energy & momentum

KNOWN

(E_1, α_1)

(E_2, α_2)

(E_3, α_3)

UNKNOWNNS

(x_1, y_1, z_1)

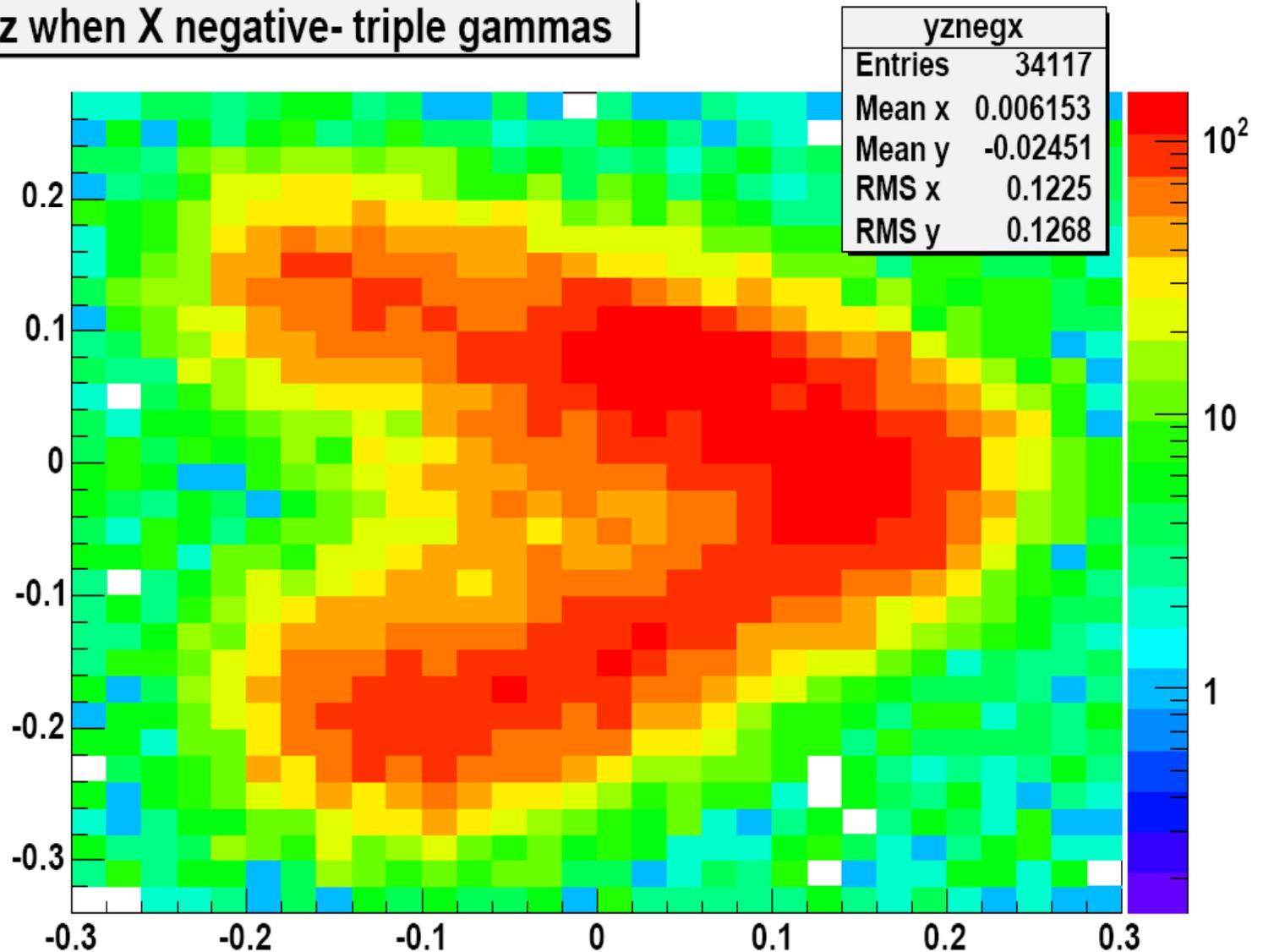
solve simultaneous eqs

POINT OF RESPONSE



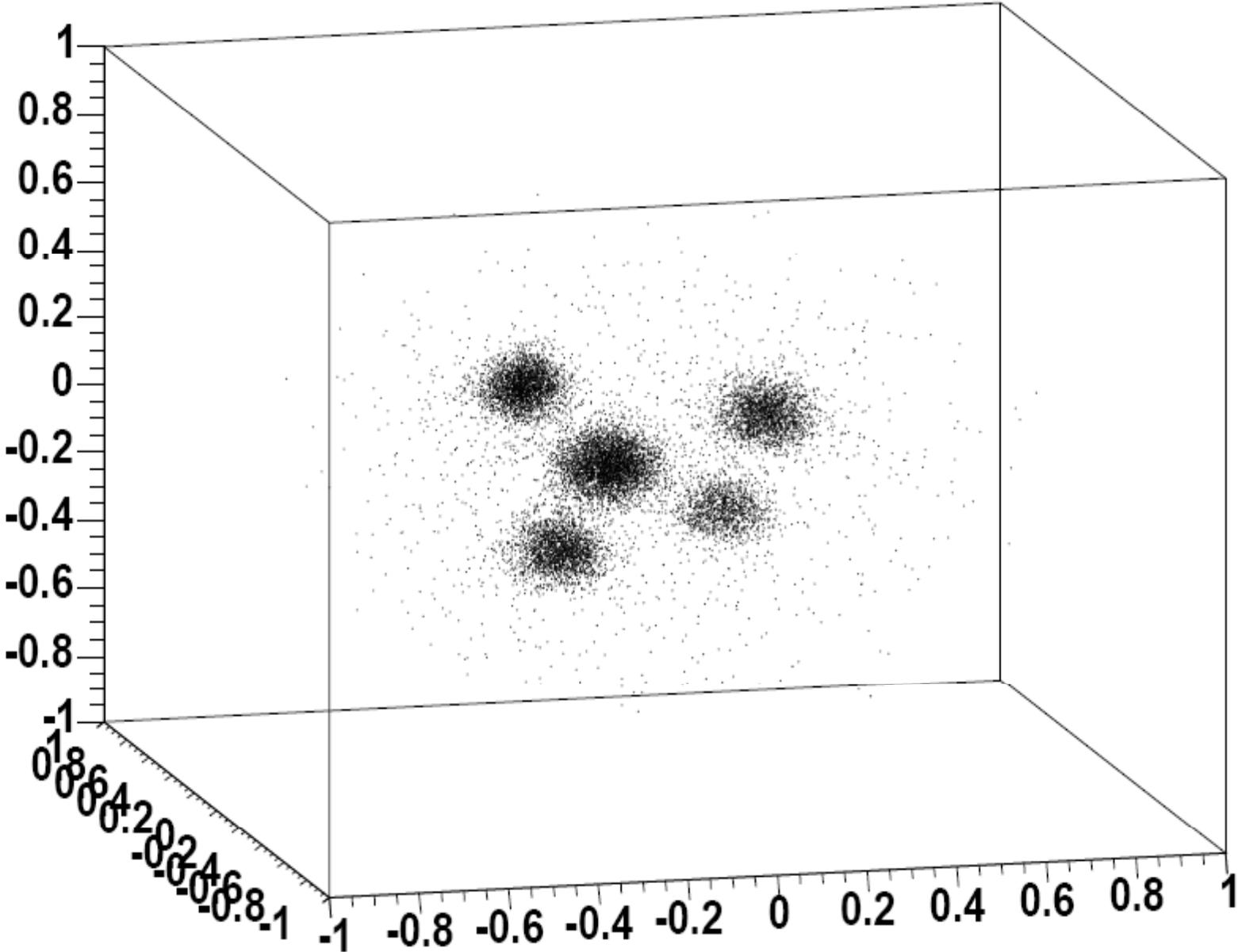
LOGO PAINTED WITH BLOOD ON STYROFOAM CUBE

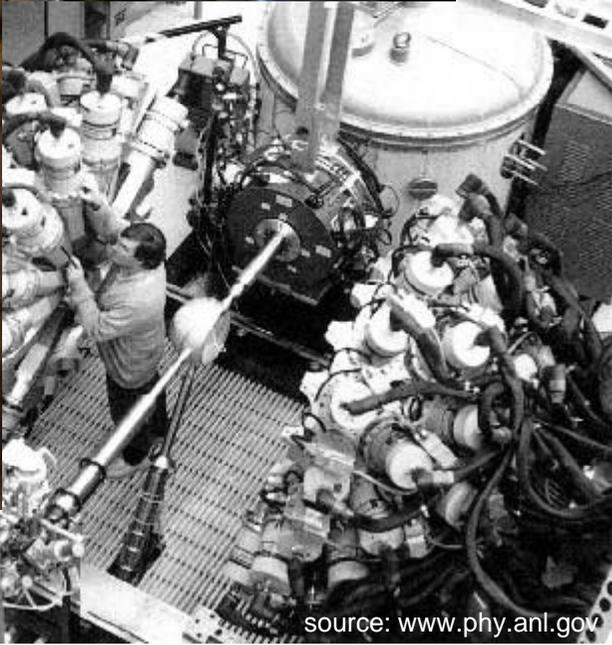
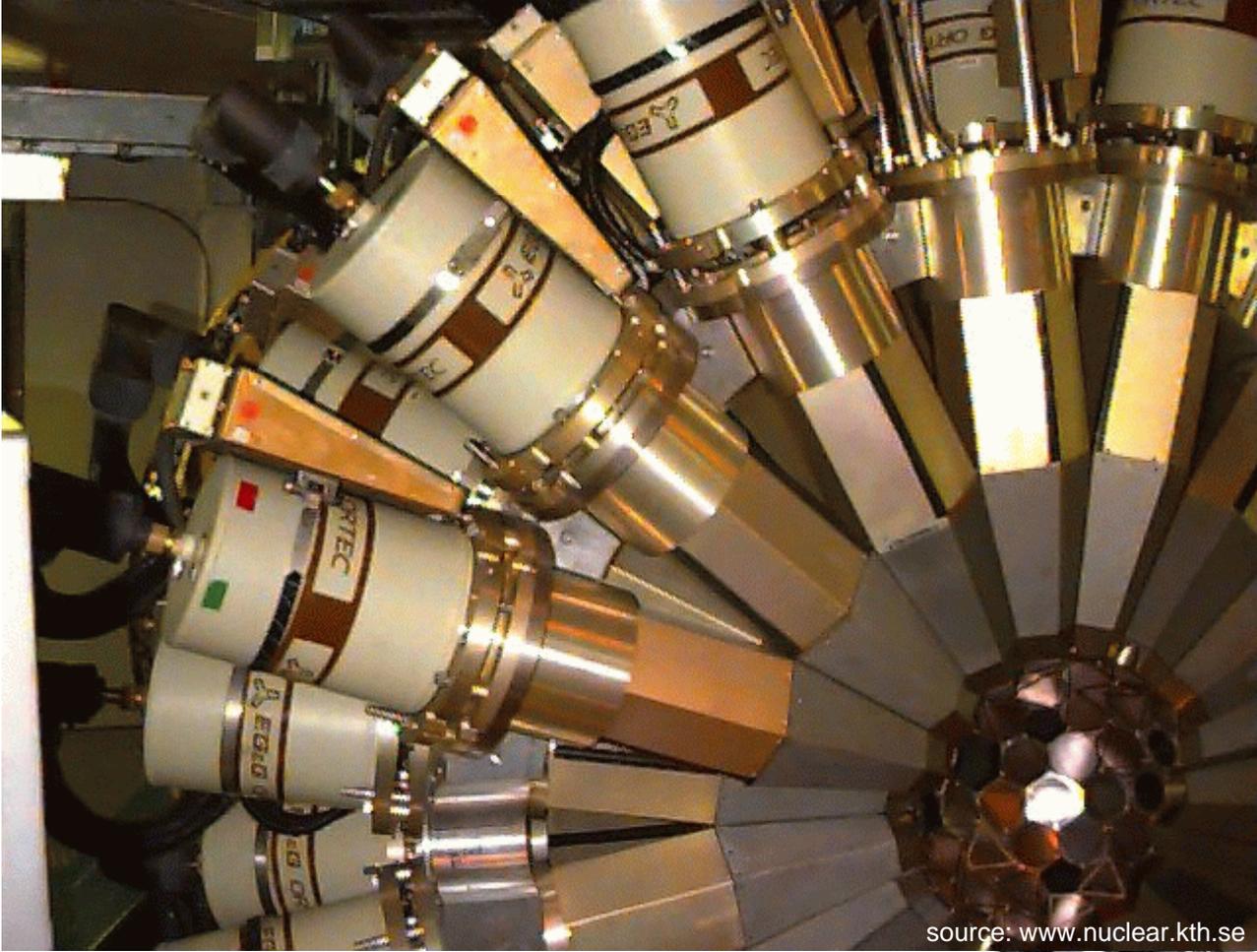
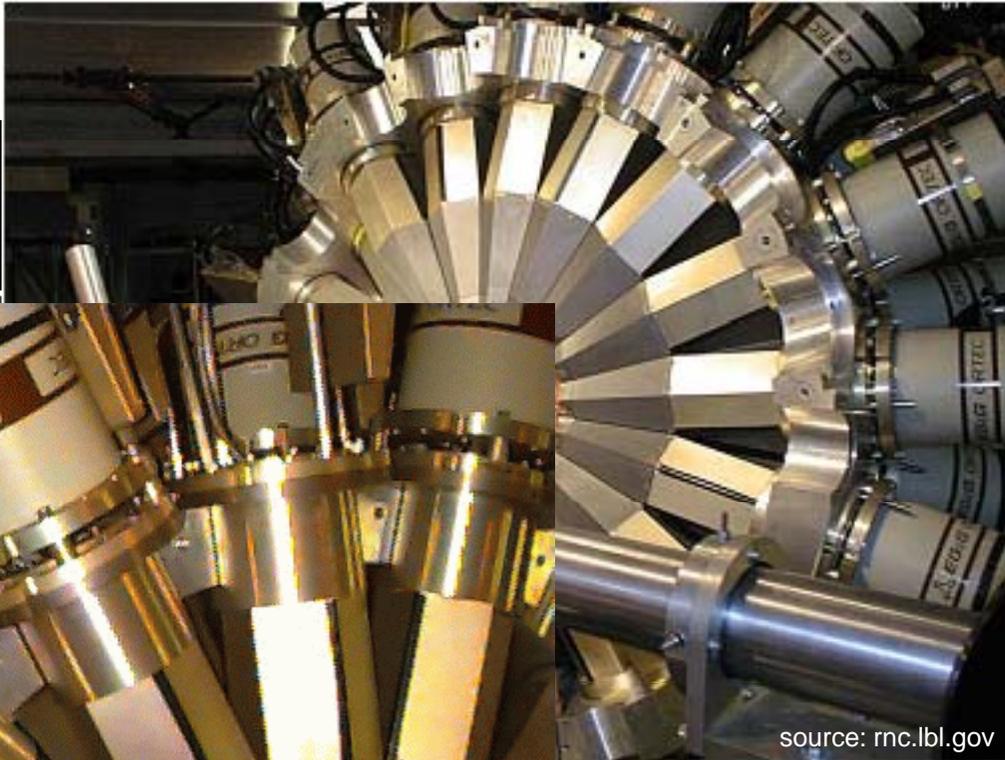
yz when X negative- triple gammas



x, y, z for triple events

5 SOURCES INSIDE
STYROFOAM CUBE





GAMMASPHERE: AN UNPRECEDENTED LUXURY FOR US

LUXURY #1 NEAR-4 π SOLID ANGLE

LUXURY #2 MULTI-DIMENSIONAL DATA WRITTEN OUT

```
GAMMASPHERE - EVENT
```

```
DATA BIT - EVENT PROCESSING CONTROL BITS
1  gcmode          gain correction on/off
2  tmmode          time veto on/off
3  hmode           adjacent detector veto on/off
```

```
The EVE BIT - EVENT DATA CONTROL BITS FOR EACH DETECTOR
```

```
typedef 4  writeget          output ge time on/off
        { 5  writegef          output full ge data on/off
          u_s 6  writebgo          output of BGO data on/off
          u_s
          u_s BIT - EVENT DATA CONTROL BITS FOR EACH EVENT
          u_s
          u_s 7  writeallge          output of dirty ge data on/off
          u_s 8  writeallbgo          output of clean bgo data on/off
          u_s
          u_s BIT - MISC
          u_s
          u_s 9  writeIsomerTag
          u_s 10 rf_timing          ge times calculated vs rf pulses
          u_s                          (subtract tac2 on the fly)
        }
EVENT_BUFFER;
```

GAMMASPHERE: AN UNPRECEDENTED LUXURY FOR US

LUXURY #1 NEAR-4 π SOLID ANGLE

1	hpid	bgo hit pattern	(0xfe00)	1111 1110 0000 0000	[1]
		and ge hit bit	(0x0100)	0000 0001 0000 0000	
		and id register	(0x00ff)	0000 0000 1111 1111	
2	ge_high	14 bit high res ge	(0x3fff)	0011 1111 1111 1111	
		and over-range bit	(0x4000)	0100 0000 0000 0000	
		and Pileup bit	(0x8000)	1000 0000 0000 0000	
3	ge_side	12 bit side ch ge	(0x0fff)	0000 1111 1111 1111	[2]

The appropriate mask to extract the information is shown both in hex and binary formats.

The rest of the events depend on what EFF write out options are on:

If the "writeget" [4] (germanium time) or "writegef" [5] (trap + lowres signals) are set

4	ge_time	12/13 bit[5] ge time	(0x1fff)	0001 1111 1111 1111	
---	---------	----------------------	----------	---------------------	--

If "writegef" [5] (trap + lowres signals) is set:

5	ge_trap	12 bit trap corr	(0x0fff)	0000 1111 1111 1111	
6	ge_low	12 bit low res ge	(0x0fff)	0000 1111 1111 1111	

If "writebgo" [6] (clean bgo) is set:

7	bgo_time	12 bit bgo time	(0x0fff)	0000 1111 1111 1111	[3]
8	bgo_low	12 bit bgo energy	(0x0fff)	0000 1111 1111 1111	[3]

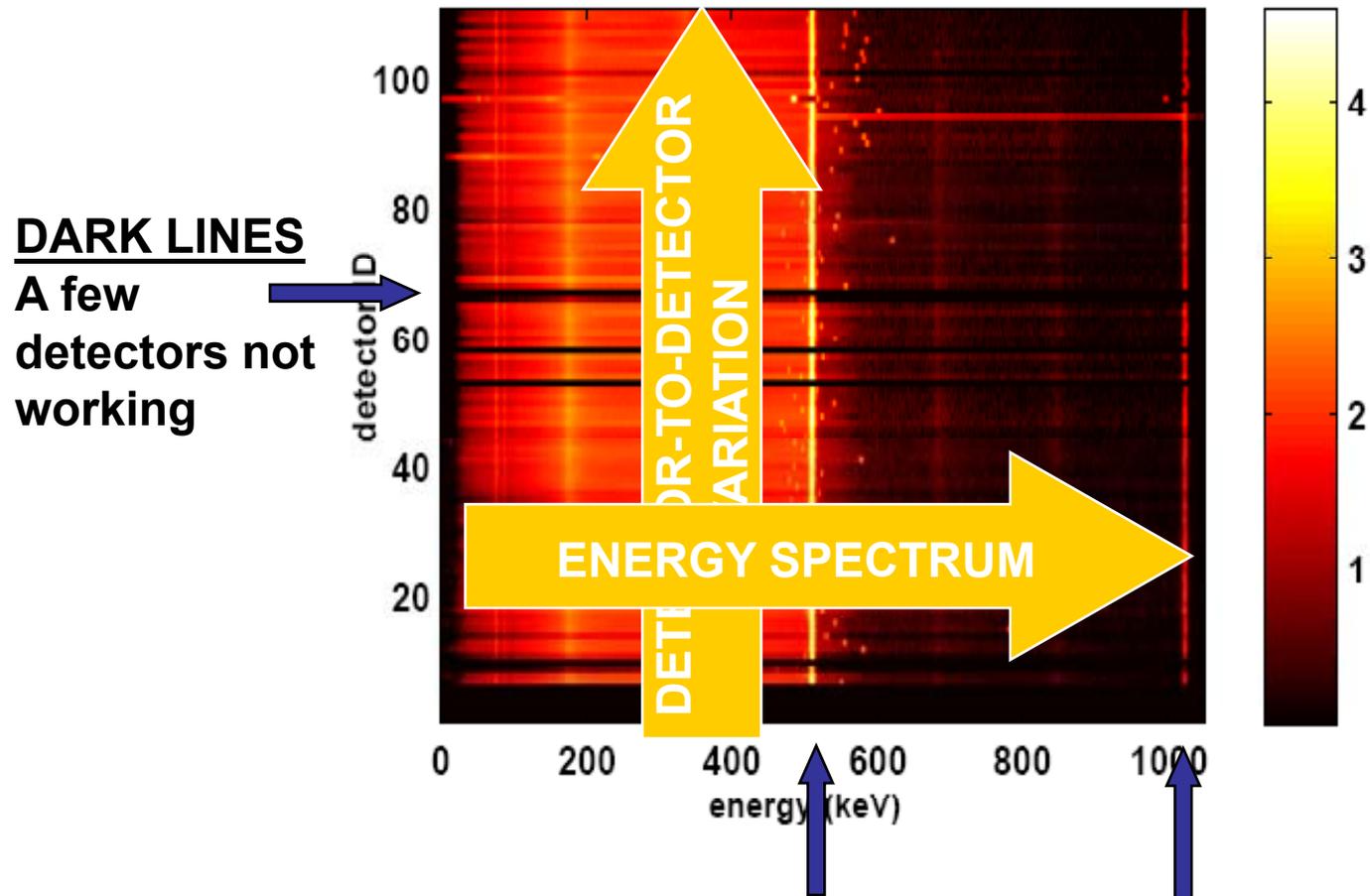


Fig. 3. The energy spectra reported by 110 individual detectors. The colour scale at each pixel denotes $\log_{10} N_{i,e}$, where $N_{i,e}$ is the number of hits in energy bin e recorded by the i th detector. A horizontal line profile drawn across the map would be the energy spectrum for the specific detector. A vertical profile across the map would be the detector-to-detector variation for the specific energy bin. Only time-gated clean hits are counted. The summed peak at 1022 keV is three orders of magnitude lower than the 511 keV peak.

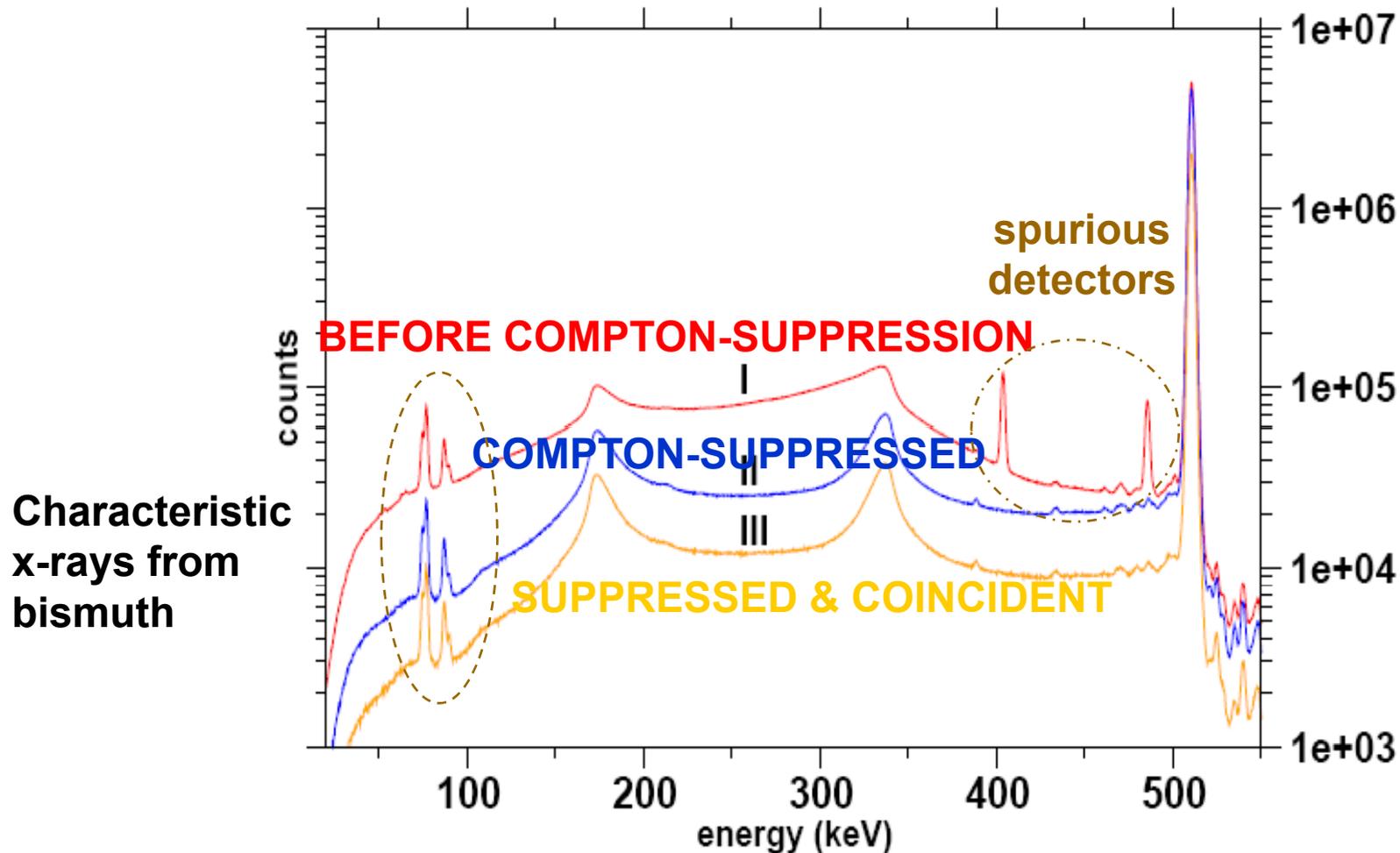
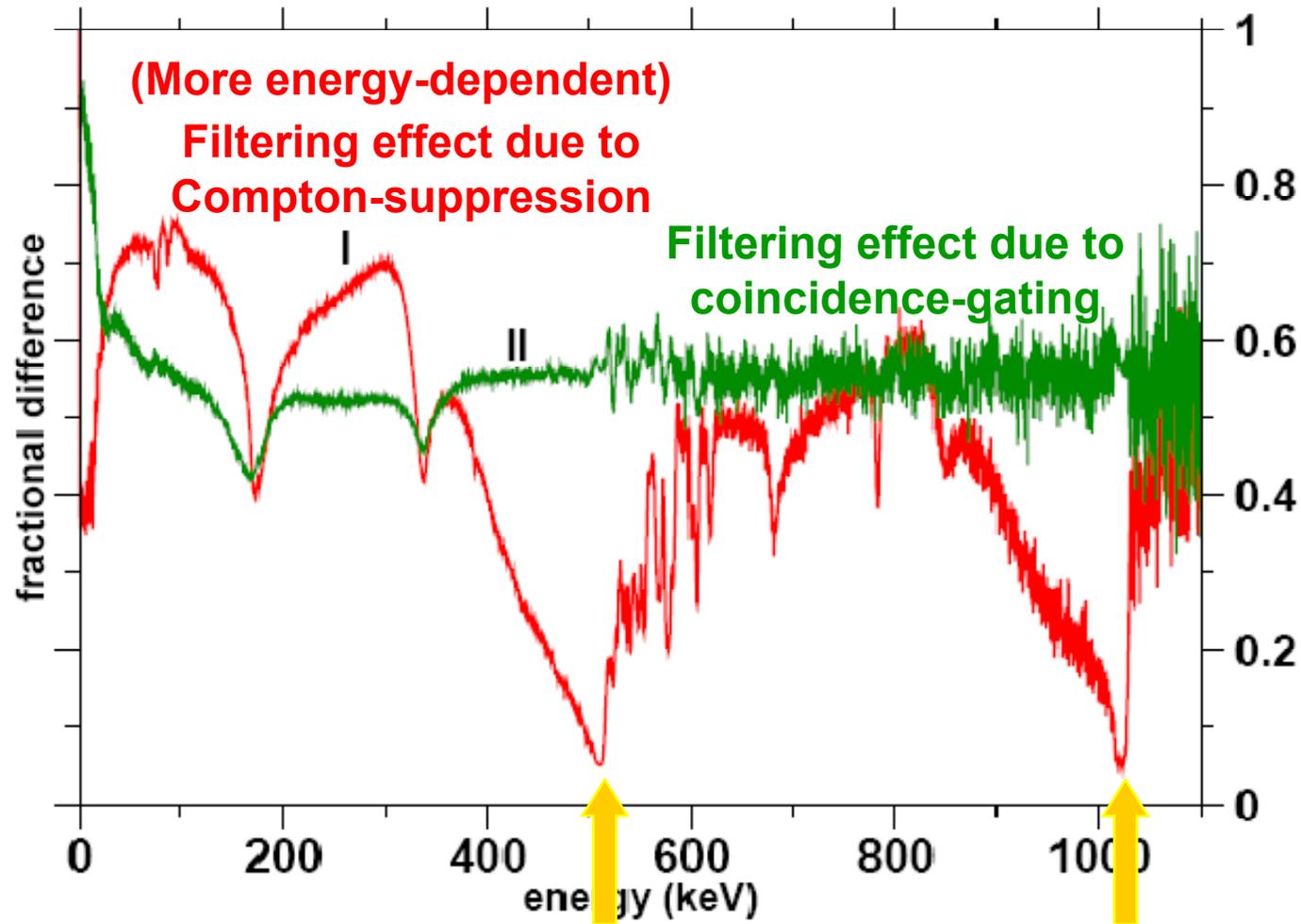


Fig. 1. Gamma spectra from the FDG source with the reference sample in place: (I) all hits from all detectors; (II) clean hits from non-outlying detectors only; (III) time-gated clean hits from non-outlying detectors only. The energy axis has been truncated.



FILTERED THE LEAST: FULL-ENERGY PHOTO-PEAK & SUMMED PEAK

Fig. 2. Variation in filtering effects with energy: fractional difference in counts per energy bin (I) with and without Compton-suppression; (II) before and after time-gating.

OURS versus OTHERS' 3γ WORK

$^{18}\text{F-FDG}$

WE USED THIS

CLINICAL SOURCE

MORE
CHALLENGING

^{22}Na

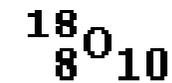
LABORATORY
SOURCE

MORE
CONVENIENT

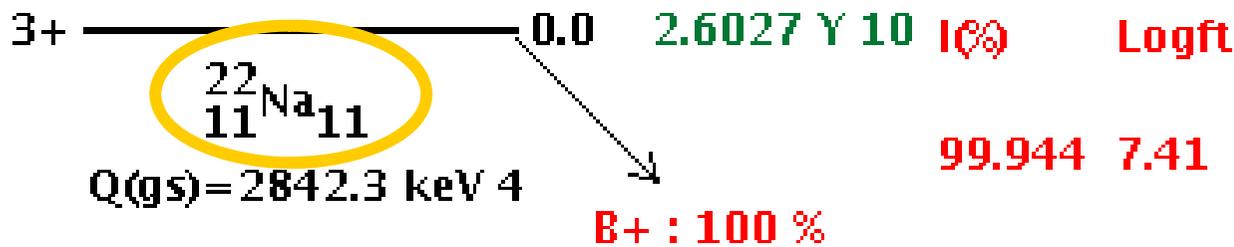


100 % Logft

100 3.5700 ————— 0 STABLE



VERSUS

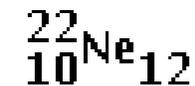


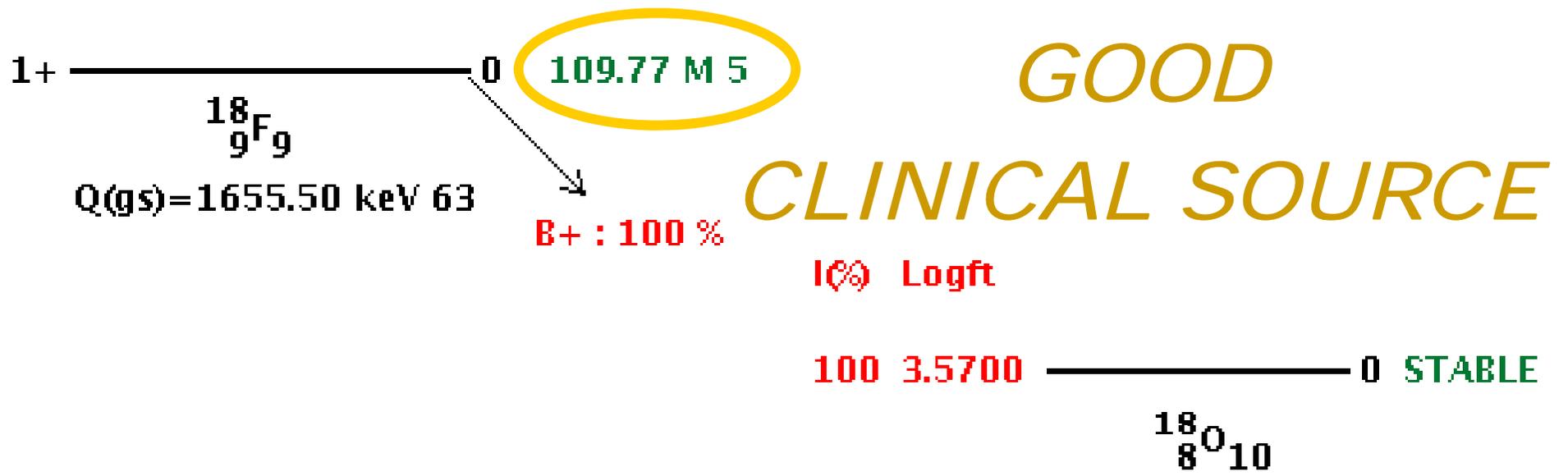
99.944 7.41

0.056 14.91

1274.5

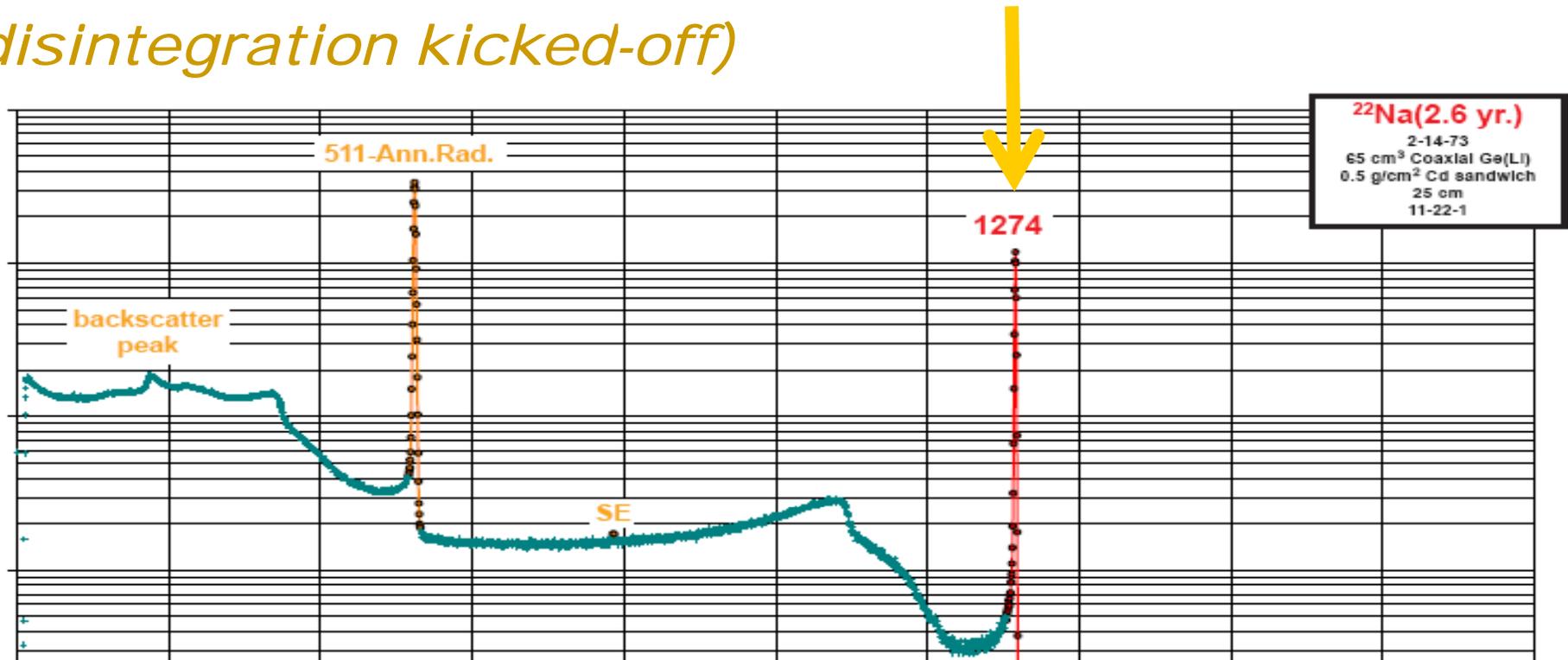
STABLE





THIS EXTRA PEAK (not found in ^{18}F)

- ALLOWS NORMALISATION BETWEEN 2 DATA ACQUISITIONS (we know how many disintegrations took place)
- PROVIDES REFERENCE TIME TRIGGER FOR LIFETIME MEASUREMENTS (we know when the disintegration kicked-off)



DETECTION OF $3\gamma/2\gamma$ YIELDS

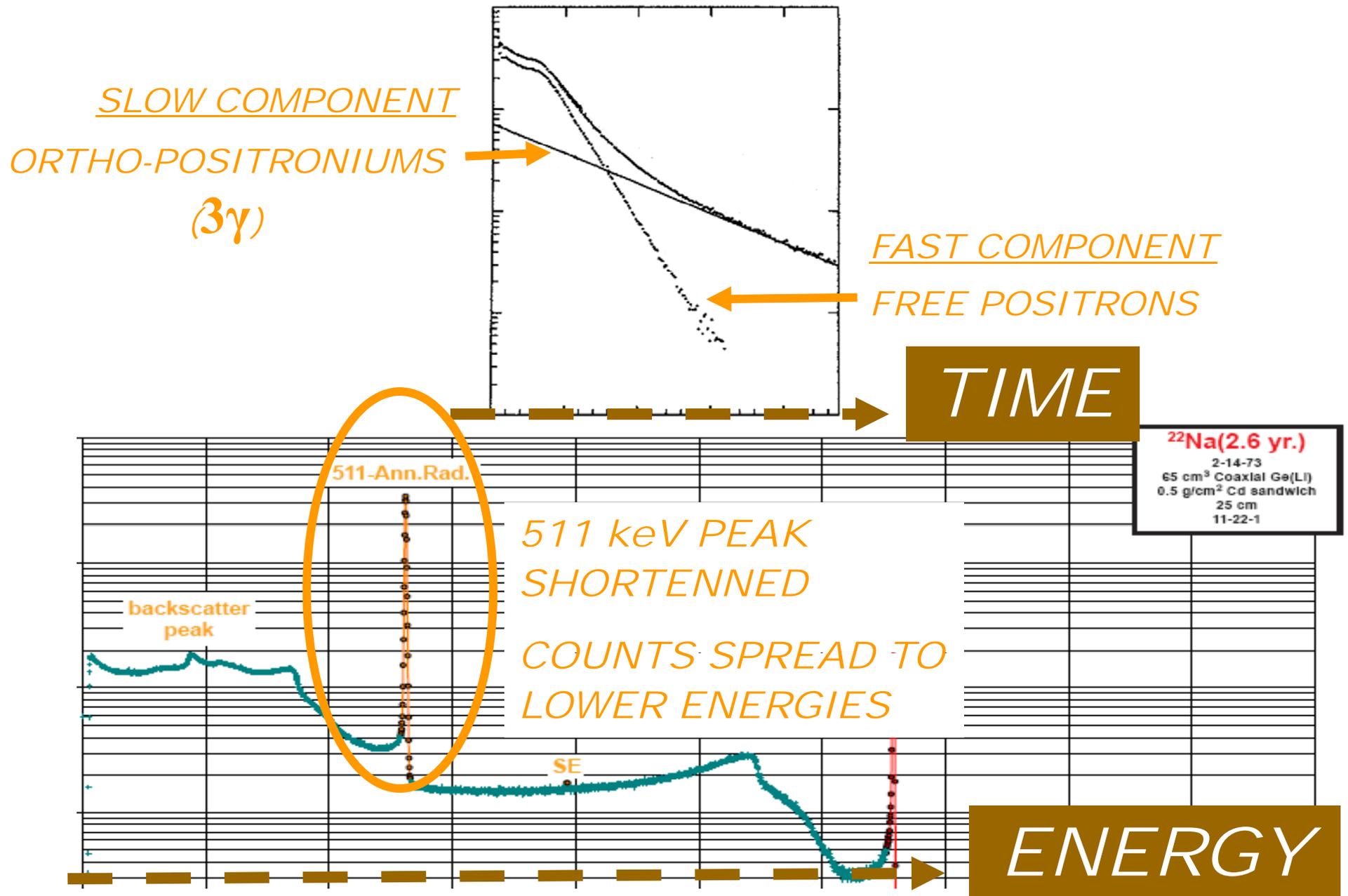
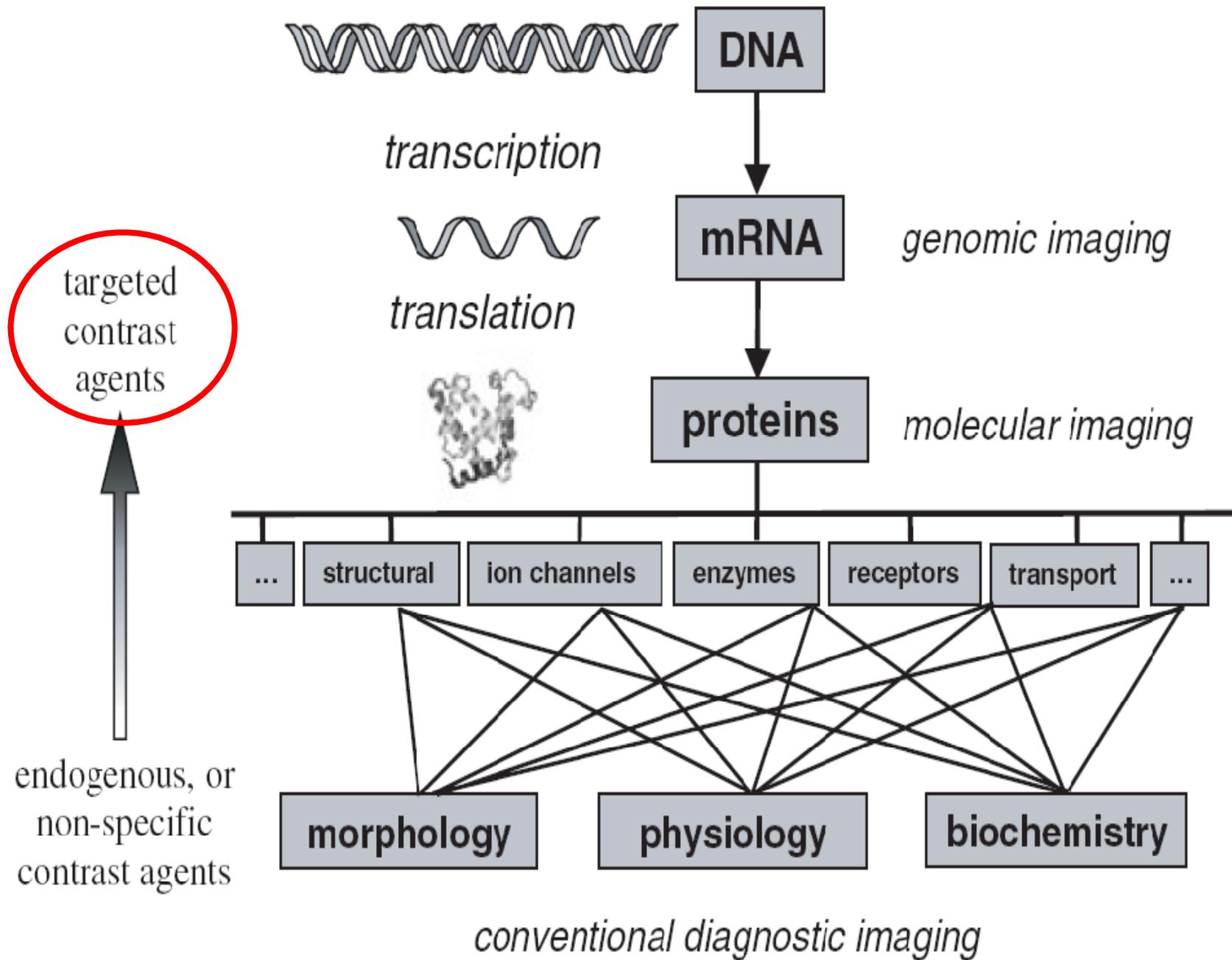


TABLE I.—*Main non-invasive in vivo imaging modalities used in molecular and cellular imaging studies.*

Imaging modality	Form of energy used and variable assessed	Main imaging agent/contrast	Primary use	Type of information
Positron emission tomography (PET)	Annihilation photons Radioactivity distribution	^{18}F , ^{11}C , ^{15}O , ^{64}Cu , ^{124}I	Whole-body clinical and research applications	Physiologic cellular molecular
Single photon emission computed tomography (SPECT)	γ photons Radioactivity distribution	$^{99\text{m}}\text{Tc}$, ^{111}In , ^{123}I , ^{125}I , ^{201}Tl	Whole-body clinical and research applications	Physiologic cellular molecular
Bio-luminescence imaging (BLI)	Visible to infrared photons Luminescence	Firefly and Renilla luciferase	Reporter gene expression, cell tracking	Cellular molecular
Fluorescence imaging (FLI)	Visible to infrared photons Fluorescence	GFP, RFP, NIR fluorochromes. Quantum dots	Reporter gene expression, cell tracking,	Cellular molecular
Magnetic resonance imaging/spectroscopy (MRI/MRS)	Radiofrequency waves Tissue molecular composition	Chelated gadolinium, SPIO, Nanoparticles	Whole-body high contrast clinical general imaging and spectroscopy	Morphologic physiologic cellular
Computer tomography (CT)	X-rays Tissue density	Iodinated, blood pool contrast agent	Whole-body general imaging. Small animal phenotyping	Morphologic



REPRODUCED FROM CHERRY SR PHYS. MED. BIOL. 2004;49:R13-48

CONCLUSION

$3\gamma/2\gamma$ YIELDS ARE ABLE TO DIFFERENTIATE
BIOLOGICAL SAMPLES

THE METHOD IS SENSITIVE TO CHEMICAL SPECIES
IN THE SAMPLES

$3\gamma/2\gamma$ YIELD DOES NOT DEPEND ON O_2 ALONE

PROSPECTS NOT JUST FOR HYPOXIA-IMAGING
ALONE BUT MOLECULAR IMAGING
ON A WIDER SCALE

FURTHER WORK

TO INVESTIGATE THE DIFFERENT CONFOUNDING
FACTORS PERTURBING $3\gamma/2\gamma$ YIELDS

DEAD TIME ISSUES

GAMMA-TRACKING INSTEAD OF
COMPTON-SUPPRESSION

ACKNOWLEDGEMENTS

Dr K Kacperski carried out the experiments while he was employed as a post-doc by the University of Surrey

This work was supported in part by the US Department of Energy, Office of Nuclear Physics, under contract No. DE-AC02-06CH11357

	Total plasma N	Urea N	NPN	Amino N	Sugar	Inorganic phosphorus	Hemo- globin	Cell
	<i>mg. %</i>	<i>mg. %</i>	<i>mg. %</i>	<i>mg. %</i>	<i>mg. %</i>	<i>mg. %</i>	<i>gm. %</i>	<i>vol. %</i>
Cattle	1177.6 (2)	12.2 (2)	30.0 (27)	7.19 (36)	53.0 (39)	5.7 (2)	10.90 (52)	34.3 (36)
Horse	1296.0 (23)	18.7 (21)	25.8 (31)		109.0 (31)			33.0 (51)
Sheep	1054.4 (22)	13.0 (26)	28.5 (33)	4.76 (37)	63.90 (40)	7.1 (44)		
Dog	1131.2 (22)	11.7 (28)	30.8 (28)	6.70 (28)	82.00 (28)	3.5 (45)	14.11 (47)	47.7 (47)
Rabbit	1072.0 (23)	13.0 (26)	31.0 (32)		124.0 (43)	4.5 (46)		
Swine	1216.0 (22)	17.3 (29)	31.4 (34)		128.0 (34)		11.95 (48)	47.8 (48)
Rat	1040 (23)	15.6 (29)	42.0 (35)		122 (42)			48.0 (40)
Hen		4.53 (25)	38.5 (25)		212 (25)	3.96 (2)		
Cat	1347.2 (22)	30.3 (26)	52.6 (32)					
Man	1120 (24)	17.1 (30)	35.6 (30)	6.40 (38)	112.0 (30)			45.6 (50)
Goat*	1128	22.31	48.52	9.60	59.1	7.70	9.26	28.8

REPRODUCED FROM HOUCHIN OB et al J. DAIRY SCI. 1939;22:241-50

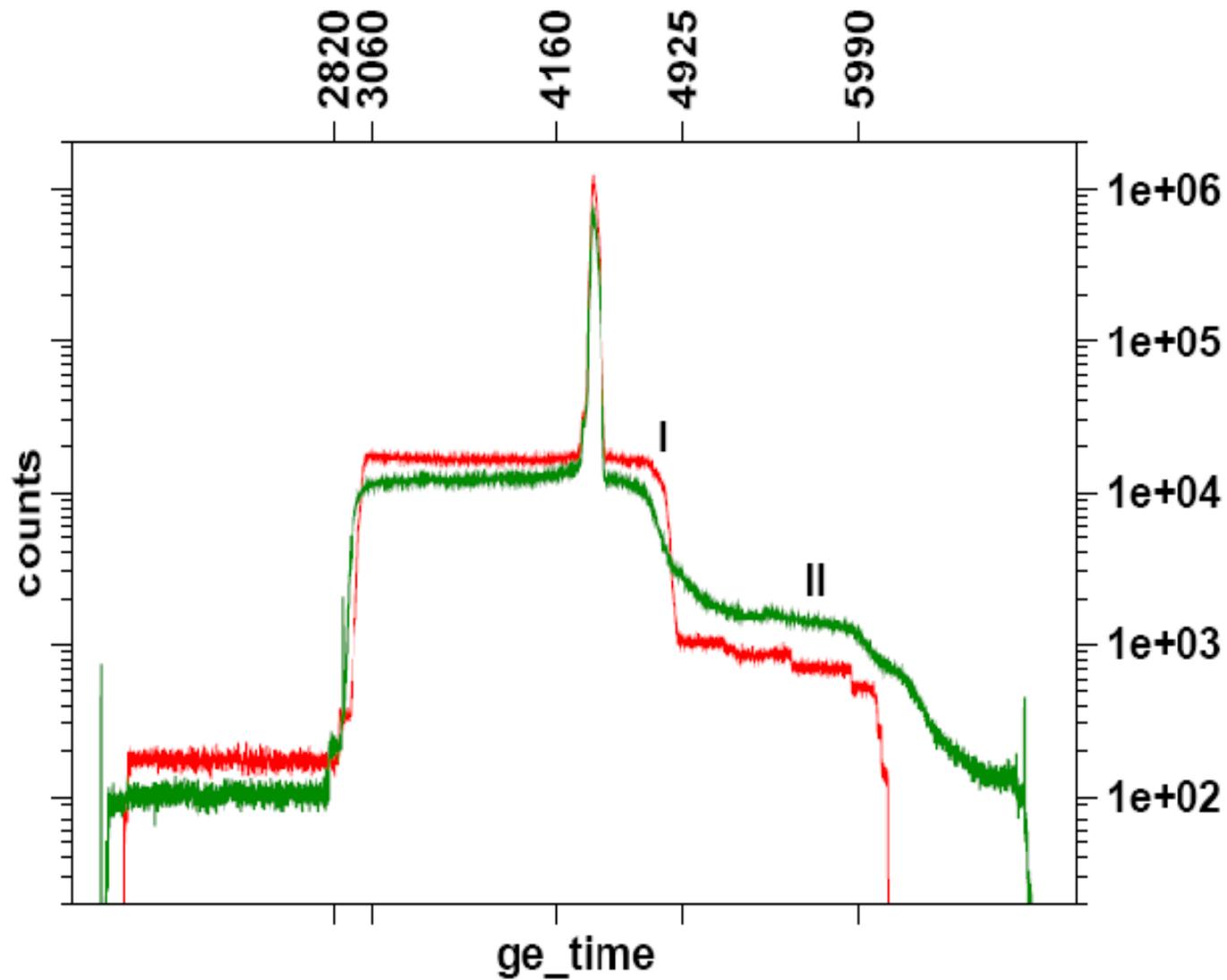


Fig. 4. Time spectra of (I) clean hits and (II) dirty hits. The FWHM of the peak is equivalent to ~ 5 ns. Hits from outlying detectors were not counted.