The dicentric assay in triage mode as a reliable biodosimetric scoring strategy for population triage in large scale radiation accidents

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The aim of the project

After an unclear radition exposure, the conventional chromosome analysis based on the dicentric assay is method of choice for dose reconstruction. But the method is time consuming and needs well trained scorers (One Scorer needs 1 week for a carefull analysis of 1000 cells).

Strategies, to receive fast results in a large scale radiation accident :

- > Network for mutual assitance, increasing capacity
- Faster scoring strategies, scoring in triage mode, quick scan, web based
- Improvement of the method, automation

Here: 8 laboratories are involved in an inter-comparison to set up a new network, first step is to validate the scoring in triage mode









Dicentric chromosome assay performed in triage mode





Mitosis in first cell division, Fluoreszence plus Giemsa staining

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IRPA 13, Glasgow, May 13th -18th 2012

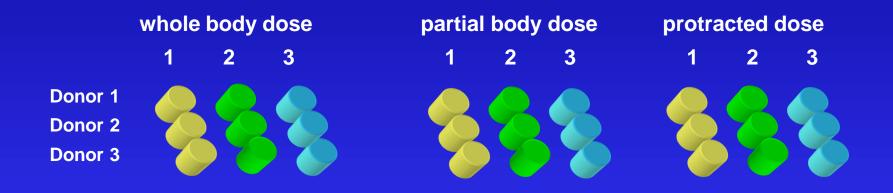






Each Lab perform conventional scoring as usual

Use of standard culture conditions Conventional scoring of 50 cells / slide, 2 slides / sample



Plus 6 shame controls

Status:

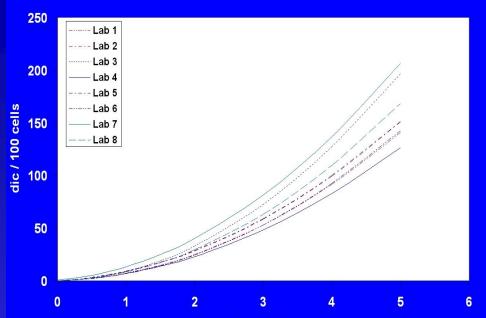
33 blood samples were distributed in total. Scoring in triage mode (20, 30 & 50 cells) performed and doses estimated







The γ -ray dose effect curves of participants used for dose estimations $Y = C + \alpha^*D + \beta^*D^2$ Curves are not significant different Mean $\alpha = 0.023$, $\beta = 0.059$ SE $\alpha = 28 \pm 4 \%$, $\beta = 7 \pm 2 \%$



dose (Gy)

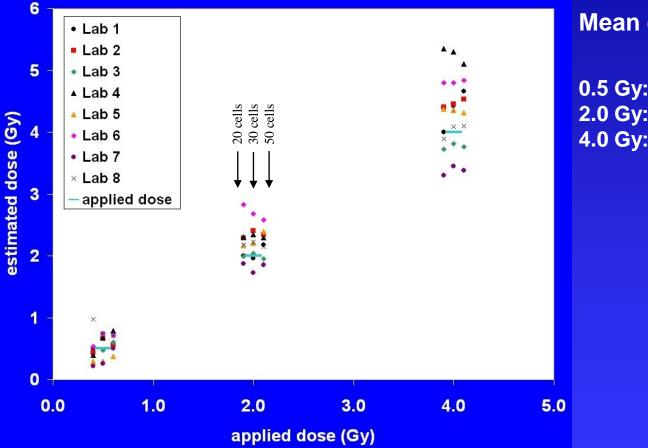
| С | SE | alpha | SE | beta | SE | radiation quality | dose rate (Gy / min) |
|--------|--|--|--|--|--|--|--|
| 0.0002 | 0.0001 | 0.0187 | 0.0047 | 0.0527 | 0.0039 | Caesium 137 | 0.42 |
| 0.0007 | 0.0004 | 0.0375 | 0.0085 | 0.0531 | 0.0054 | Cobalt 60 | 0.30 |
| 0.0005 | 0.0005 | 0.0142 | 0.0044 | 0.0759 | 0.0027 | Cobalt 60 | 0.50 |
| 0.0011 | 0.0008 | 0.0228 | 0.0048 | 0.0460 | 0.0020 | Cobalt 60 | 0.28 |
| 0.0010 | 0.0004 | 0.0338 | 0.0101 | 0.0536 | 0.0044 | Cobalt 60 | 0.50 |
| 0.0006 | 0.0003 | 0.0135 | 0.0043 | 0.0544 | 0.0034 | Cobalt 60 | 0.25 |
| 0.0072 | 0.0063 | 0.0538 | 0.0310 | 0.0716 | 0.0163 | Caesium 137 | 0.40 |
| 0.0013 | 0.0005 | 0.0210 | 0.0052 | 0.0631 | 0.0040 | Cobalt 60 | 1.07 |
| | 0.0002 0.0007 0.0005 0.0011 0.0010 0.0006 0.0072 | 0.00020.00010.00070.00040.00050.00050.00110.00080.00100.00040.00060.00030.00720.0063 | 0.0002 0.0001 0.0187 0.0007 0.0004 0.0375 0.0005 0.0005 0.0142 0.0011 0.0008 0.0228 0.0010 0.0004 0.0338 0.0006 0.0003 0.0135 0.0072 0.0063 0.0538 | 0.00020.00010.01870.00470.00070.00040.03750.00850.00050.00050.01420.00440.00110.00080.02280.00480.00100.00040.03380.01010.00060.00030.01350.00430.00720.00630.05380.0310 | 0.00020.00010.01870.00470.05270.00070.00040.03750.00850.05310.00050.00050.01420.00440.07590.00110.00080.02280.00480.04600.00100.00040.03380.01010.05360.00060.00030.01350.00430.05440.00720.00630.05380.03100.0716 | 0.00020.00010.01870.00470.05270.00390.00070.00040.03750.00850.05310.00540.00050.00050.01420.00440.07590.00270.00110.00080.02280.00480.04600.00200.00100.00040.03380.01010.05360.00440.00060.00030.01350.00430.05440.00340.00720.00630.05380.03100.07160.0163 | CSEalphaSEbetaSEquality0.00020.00010.01870.00470.05270.0039Caesium 1370.00070.00040.03750.00850.05310.0054Cobalt 600.00050.00050.01420.00440.07590.0027Cobalt 600.00110.00080.02280.00480.04600.0020Cobalt 600.00100.00040.03380.01010.05360.0044Cobalt 600.00060.00030.01350.00430.05440.0034Cobalt 600.00720.00630.05380.03100.07160.0163Caesium 137 |







Dose estimates based on 20, 30 and 50 cells Acute whole body exposure (HDR) with 0.5, 2.0 and 4.0 Gy



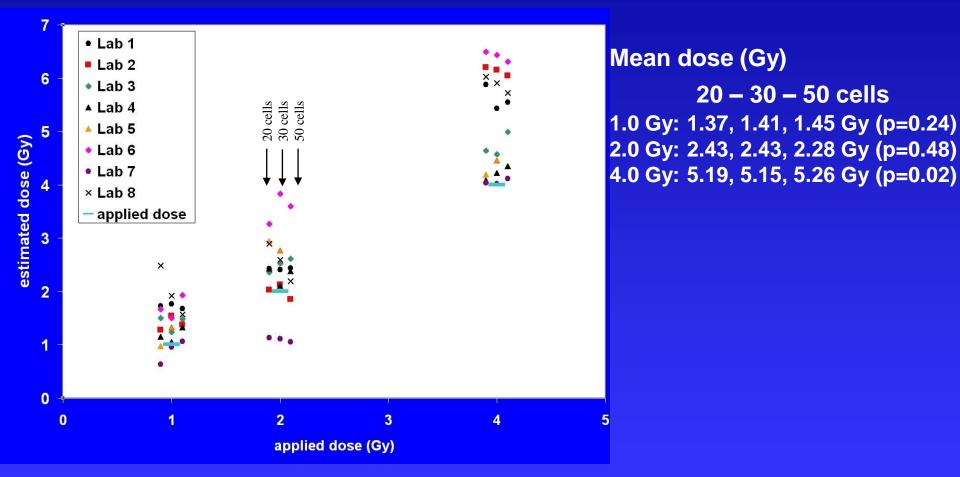
Mean dose (Gy) 20 - 30 - 50 cells 0.5 Gy: 0.51, 0.57, 0.60 Gy (p=0.74) 2.0 Gy: 2.21, 2.20, 2.22 Gy (p=0.64) 4.0 Gy: 4.26, 4.33, 4.34 Gy (p=0.53)







Dose estimates based on 20, 30 and 50 cells Protracted exposure (LDR) with 1.0, 2.0 and 4.0 Gy





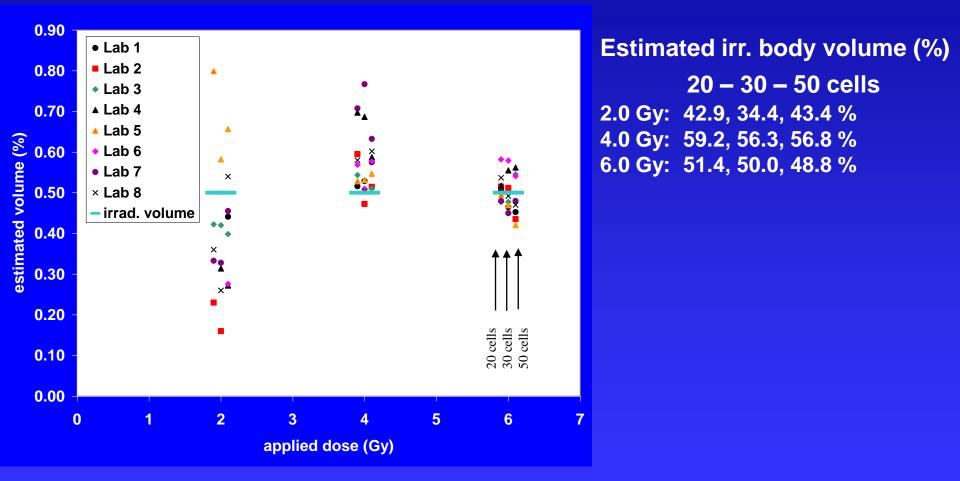


COOPERATION

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% irr. body volume estimates based on 20, 30 and 50 cells Partial body exposure (PAR) with 1.0, 2.0 and 6.0 Gy



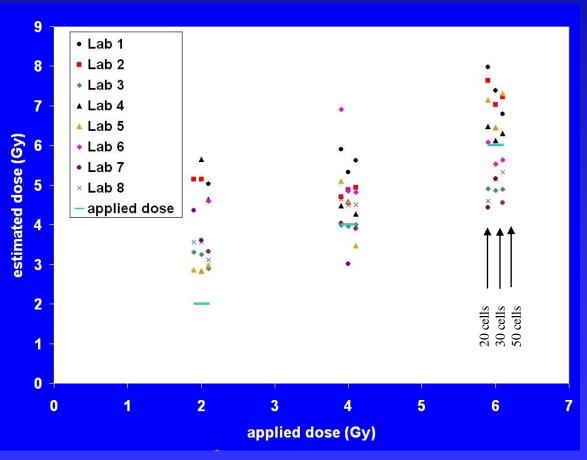
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COOPERATION



Dose estimates based on 20, 30 and 50 cells Partial body exposure (PAR) with 1.0, 2.0 and 6.0 Gy



Mean dose (Gy) 20 - 30 - 50 cells 2.0 Gy: 1.67, 1.36, 2.33 Gy (p=0.02) 4.0 Gy: 3.71, 3.63, 4.18 Gy (p=0.54) 6.0 Gy: 5.16, 5.78, 5.88 Gy (p=0.84)





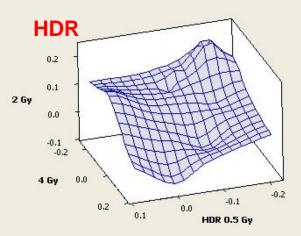
Youden plots showing the uncertainties caused by random

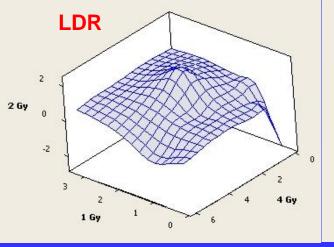
and systematic errors

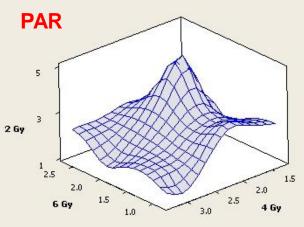
z - values on the straight surface but far from the origin indicate large systematic errors, z - values far from the origin and not on the surface indicate large random errors.

HDR: This figure nicely illustrates the spread of dose estimates across the labs, which increases with increasing dose, but decreases with increasing numbers of cells scored.

LDR is fairly flat, but the higher z-values are related to the systematic trend of dose over-estimation with Increasing protracted dose. PAR: The increased z-values indicate systematic errors and the bumps indicate random errors, highlighting the need to score more cells when assessing partial body exposures.







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Summary

- > All labs were well trained during this exercise, using the tools for the different scenarios
- > 20 cells can give a rough indication of dose, 50 cells are more accurate, but not different
- > The results after acute exposure were good
- There was an unexpected trend to dose overestimations after protracted exposure, may be explained by uncertainties in the alpha term
- Limitations of scoring in triage mode after partial body exposure were observed, therefore we recommend to increase the cell number in cases, where there are signs of potential inhomogeneous exposure in order to obtain more reliable data
- With respect to the homogeneity of the conventional scoring results, each lab was able to manage the challenges in biological dosimetry
- In the frame of the MULTIBIODOSE project, further investigations with these samples will be performed to improve the automated scoring procedure and to validate new scoring strategies







Thank you for your attention



This work was funded by EU project FP7-SEC-241536-MULTIBIODOSE





