



Homeland Defense & Security Information Analysis Center

HDIAC Medical Webinar Briefing: Current High Throughput Cytogenetic Techniques for Radiation Biodosimetry August 25, 2015

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To understand the cellular basis of chronic and acute ionizing radiation injuries and their impact on human health.





Natural sources

Man-Made

- Cosmic rays
- Radon radioactive gas resulting from the decay of Uranium
- Potassium-40 and Carbon-14 from food
- Ingesting contaminated air and water

- Diagnostic X-rays
- Radioisotopes
- Occupational exposure (Radium/Uranium)
- Nuclear accidents
- TVs, computers & mobile phones
- Luminous markers
- Radioactive watch dials



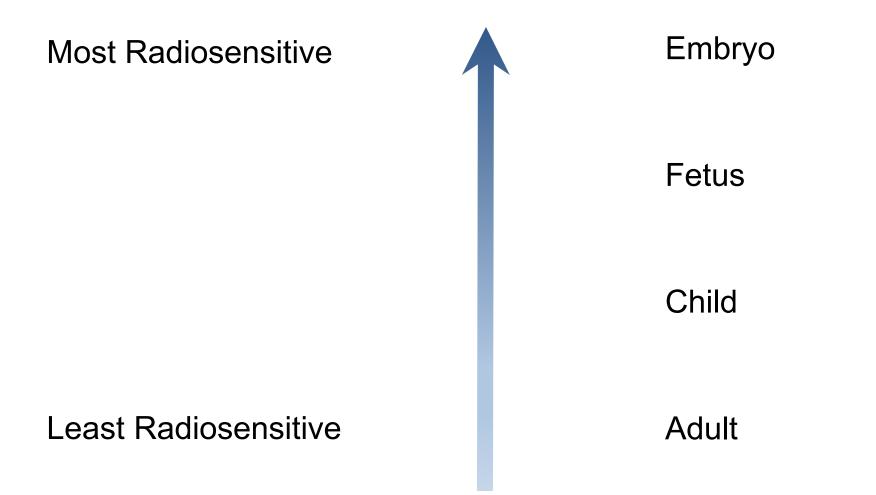


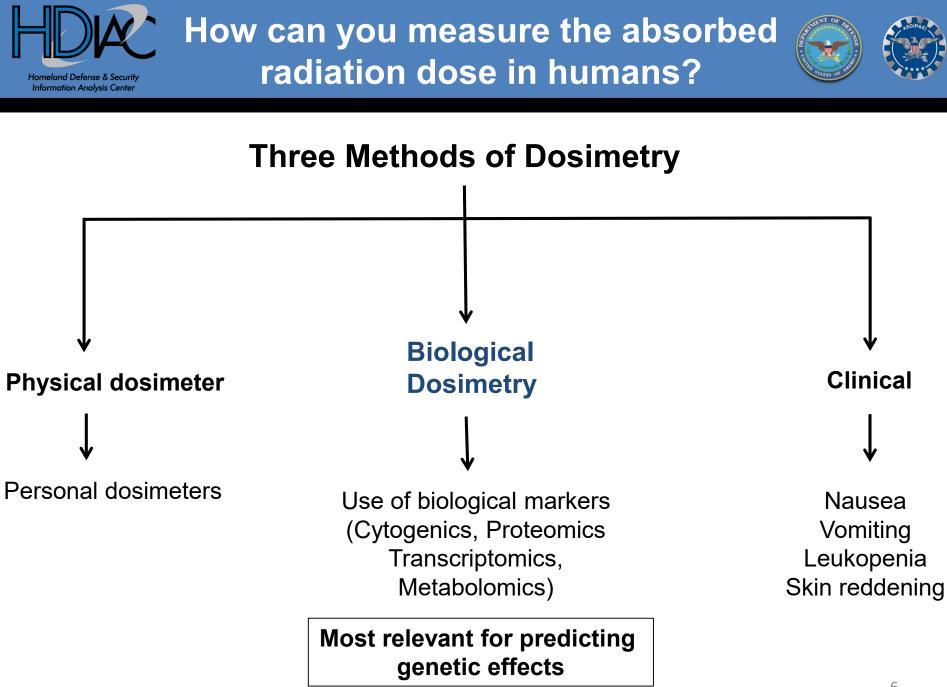
- Total dose
- 🔿 Dose rate
- Volume of tissue or anatomical body part irradiated
- → Type of radiation (Low and high LET)
- Pre-existing physical conditions, trauma, illness, burns
- Genetic predisposition inherent radio sensitivity due to genetic mutations



Human Sensitivity to Ionizing Radiation











- Nuclear Accidents (Chernobyl, Goiania and Fukushima)
- Nuclear Device Detonation
- Dirty Bomb
- Radiological Terrorism

In the above scenarios, several hundreds of people may be exposed to radiation.







- Individualized dose assessment will help in identifying the degree of exposure (low vs high)
- Predict the long-term health effects including tissue degeneration and cancer risks
- To avoid psycho-social effects in low dose exposed human population
 - Reassurance that the effects are not lethal







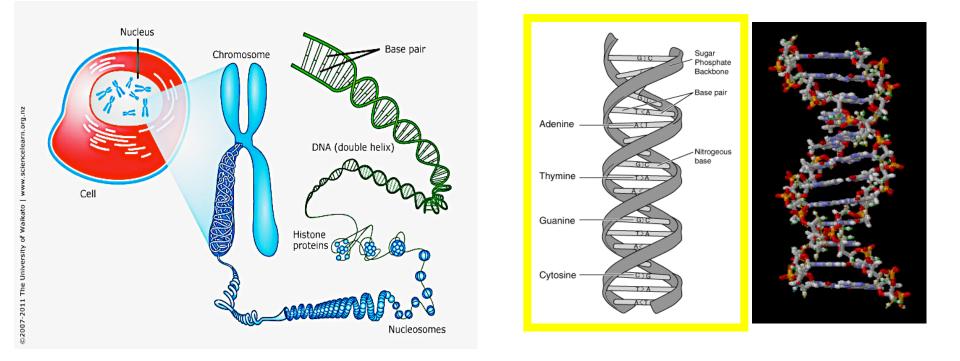
- Biological samples need to be minimally invasive
- Sensitive and specific for radiation dose assessment
- Rapid processing and analysis of large scale samples
- Stability of biomarkers over time
- Reliable and reproducible





What is the Biological Target for Ionizing Radiation?





Chromosomal DNA is susceptible to ionizing radiation induced DNA double strand breaks.



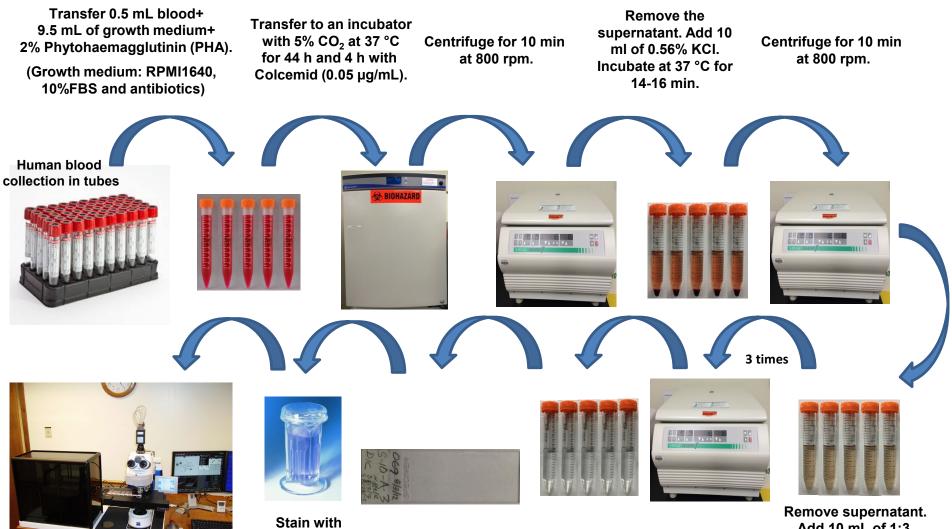


- Cytogenetics is the study of chromosome structure and function
- Each human cell nucleus has 46 chromosomes (22 pairs of autosomes and 1 pair of sex chromosomes)
- Chromosomes are made up of genes and genes are made up of DNA complexed with histone proteins
- Genes are crucial for our disease-free survival
- Ionizing radiation exposure produces breaks in the DNA thereby disrupting the chromosome structure and function
- Misrejoining of broken chromosomes result in dicentrics, rings and translocations



Preparatory Steps for Chromosome Analysis





Imaging and analysis in Metafer

5% Giemsa, rinse in water & air dry. Drop 50 μL of cell suspension on slide.

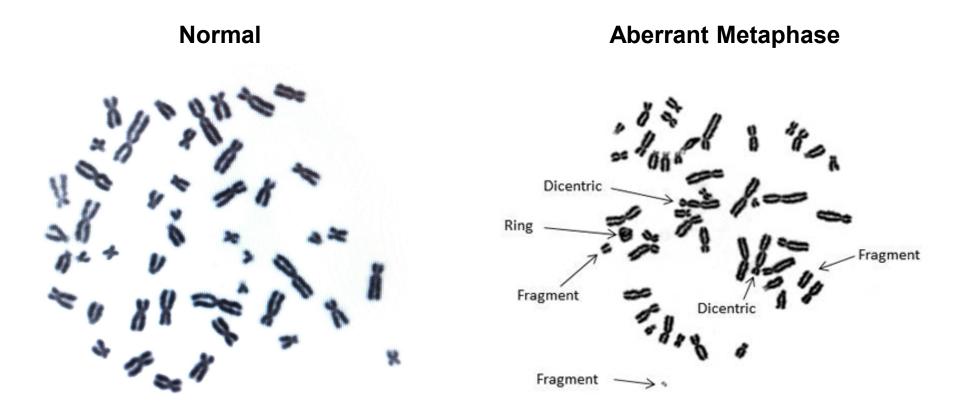
Remove supernatant. Suspend the pellet in 500 µL of fixative. Centrifuge for 10 min at 800 rpm.

Remove supernatant. Add 10 mL of 1:3 acetic acid:methanol Fixative.12



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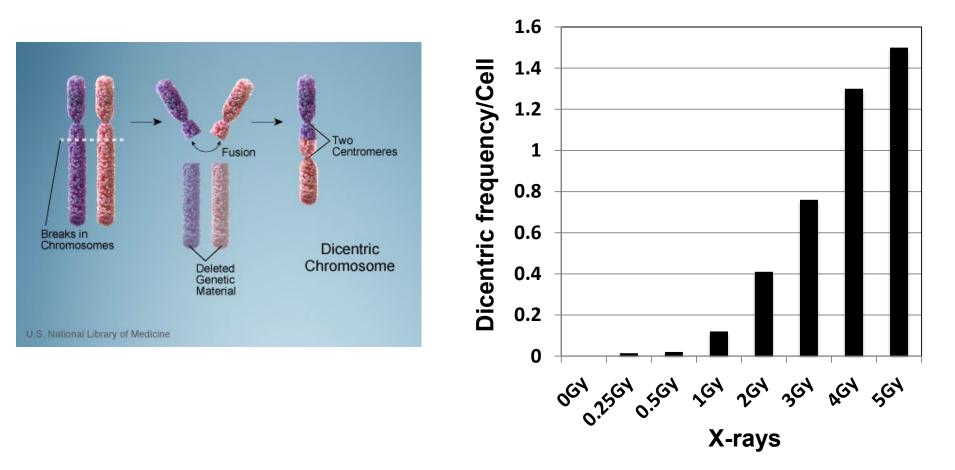
- Ionizing radiation induces chromosome aberrations.
- Radiation causes the formation of dicentric chromosomes, ring fragments and translocations due to misrejoining of strand breaks.



Types of Cytogenetic Biodosimeters



1. Dicentric Chromosome



Dicentric frequency is radiation dose dependent.





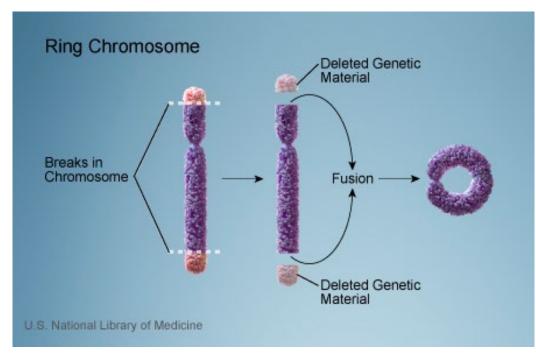
Dicentric Assay is the "gold standard" for dose assessment.

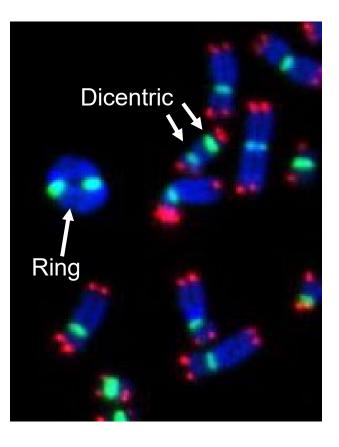
- Low background rate (1 per 1,000 cells)
- Independent of age and gender
- Ease of detection
- Sensitivity range is 0.20 to 5.0 Gy
- Reproducible dose response
- Proven in accidents over four decades







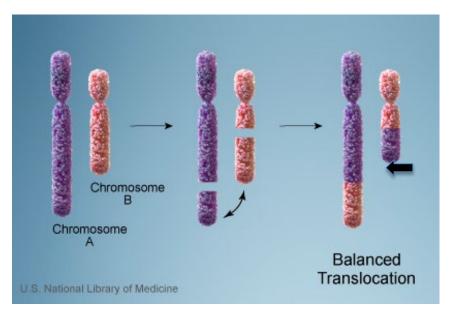


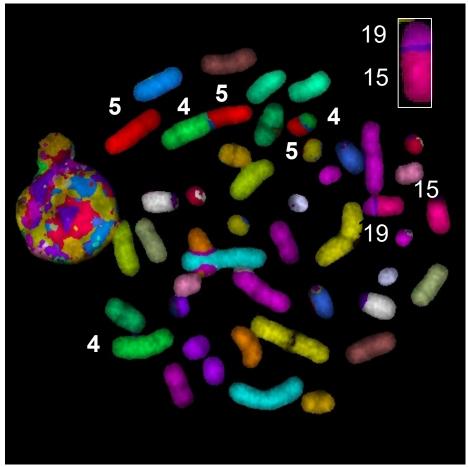




Translocation









Micronuclei



Micronuclei is a useful biomarker for radiation exposure.

Micronucleus formation CYTOCHALASIN-B BLOCK CYTOKINESIS-BLOCK Nucleoplasmic bridge formation

M. Fenech et al. Mutagenesis 2011; 26:125-132

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Difficult to increase high throughput sampling because cytogenetic assays are usually done in 15 ml conical tubes

Processing time is long (48-72hr) and labor intensive

Manual analysis is time consuming

Automation of high throughput techniques for sample processing, imaging and analysis is critical for biodosimetry in case of radiological/nuclear incident(s)

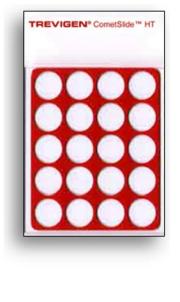


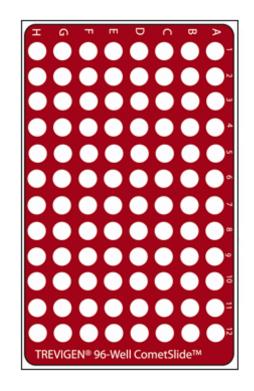
Maximizing the Sample Processing





Use of bar coded 1.2 mL tubes can be used for high throughput sample processing for cytogenetic analysis using as little as 50 µL of blood

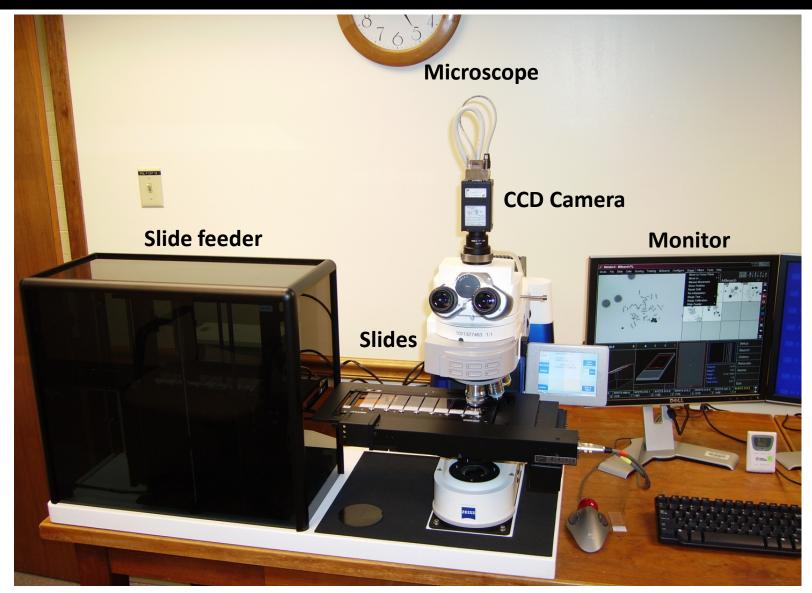




Samples can be spotted on multichambered glass slides for high throughput imaging and analysis



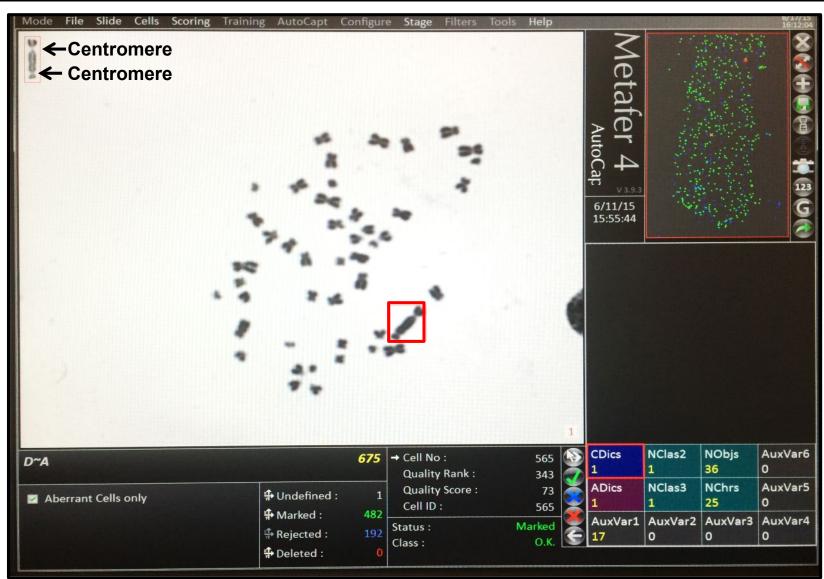




Automated Detection of Human Dicentric Chromosomes Using Metafer

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Timeline



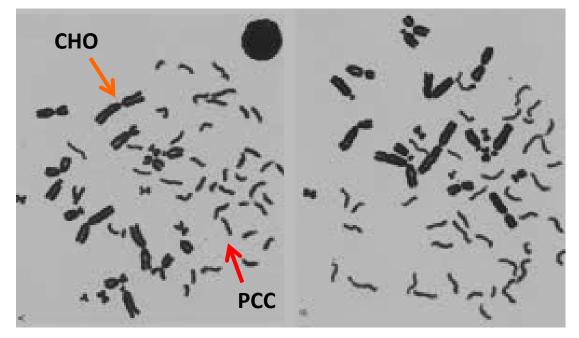
Time Line for Single Metaphase Image Scanning, Capture and Dicentric Analysis		
	Anonymous scorer	Metafer
	(Manual)	(Automated)
Spotting a metaphase spread		
at 10X Objective	10-15 sec	~0.4 sec
Analysis at 63X Objective		Capture/analysis time
Cell with 0 dicentrics	30 sec	$\frac{captarcyanarysis time}{8.5 \pm 0.1 \text{ sec}}$
Cell with 1-2 dicentrics	35 sec	8.5 ± 0.1 sec
Cell with 2-4 dicentrics	45 sec	8.5 ± 0.1 sec
Cell with 6-8 dicentrics	1 min	8.5 ± 0.1 sec
Cell with > 8 dicentrics	1 min 15 sec	8.5 ± 0.1 sec
Average time for a metaphase	49 sec	8.5 sec
Average time for 1000 metaphases	s 13.61 h	2.47 h



Premature Chromosome Condensation (PCC) Technique



Fusion of mitotic CHO cells with human G0 lymphocytes

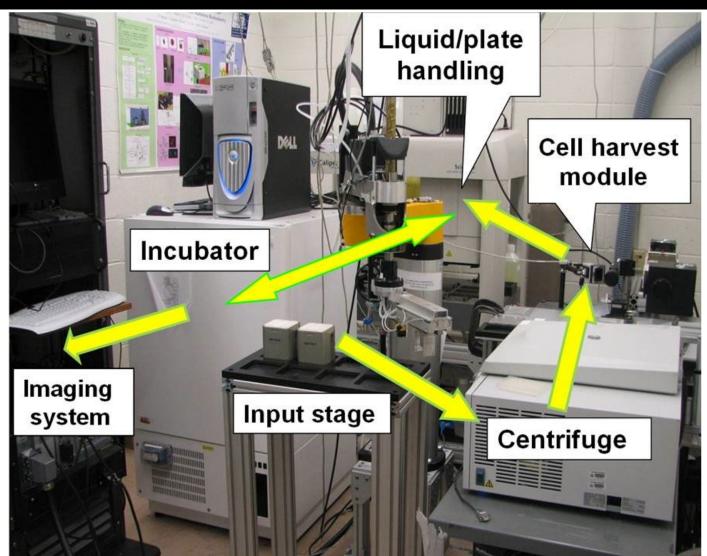


Dicentrics, rings and translocations can be detected on PCC by FISH Dicentric assay time can be cut short to 6h after blood <u>collection</u>

Rapid Automated Biodosimetry Tool (RABiT) Homeland Defense & Security

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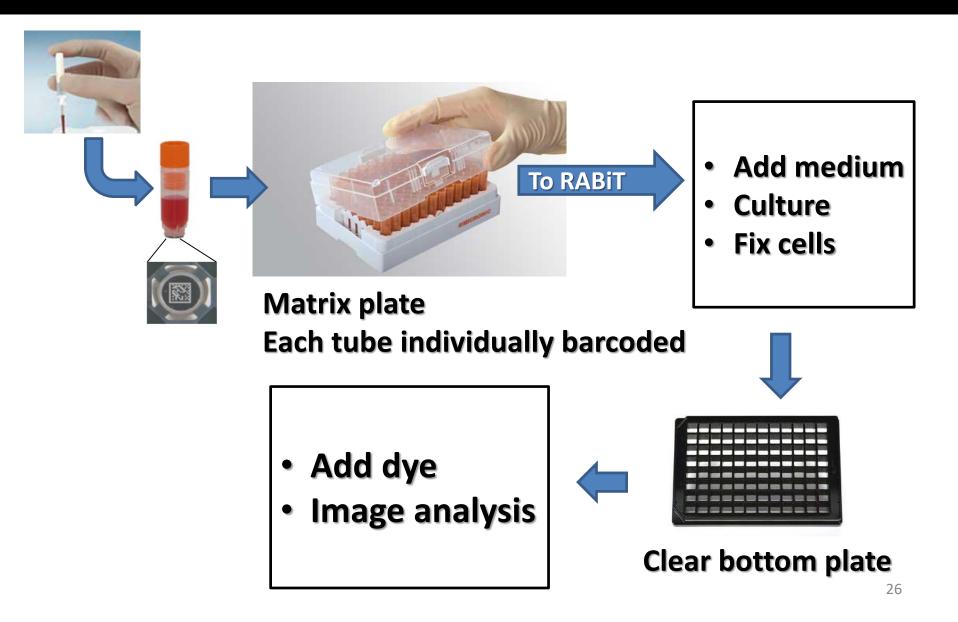


CUMC NY

https://www.youtube.com/watch?feature=player embedded&v=ESeT63F3YQY#t=0







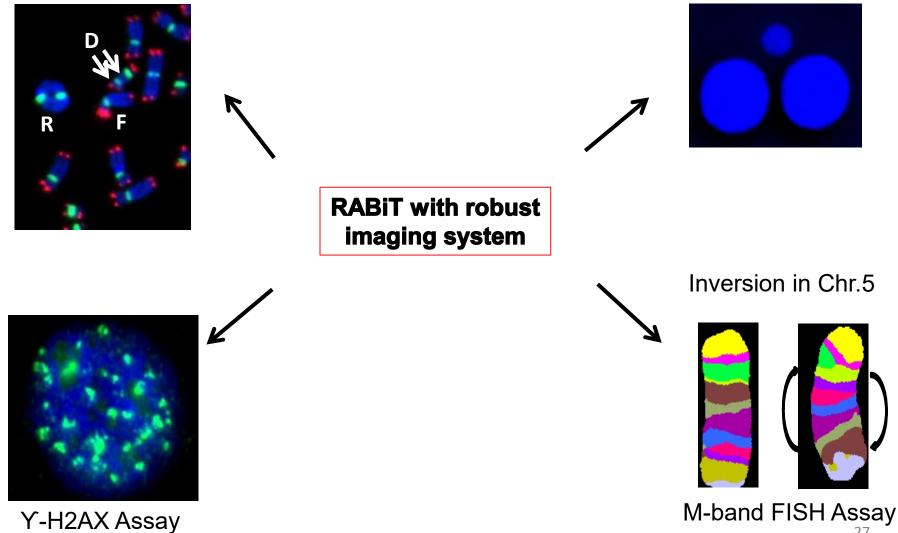


Dicentric Assay

Multiple High Throughput Radio Biodosimetry Assays



Micronucleus Assay







An automated imaging system for radiation biodosimetry. Microsc Res Tech. 2015 Jul;78(7):587-98. doi: 10.1002/jemt.22512. Epub 2015 May 4.





Biomarkers for multiple organ dysfunction syndrome (MODS) after radiation exposure *in vivo*.

- 1. Hematopoietic system/Bone marrow aplasi Flt3-L
- 2. Gastrointestinal tract
- 3. Liver and Cardiovascular
- 4. Salivary gland

- Citrulline
- Oxysterols
- Amylase

Flt3-L has been successively used to distinguish between total and partial body irradiation in mice.





- Recent years have seen a tremendous improvements in developing multiple genomic, proteomic, metabolomics and transcriptomic biomarkers for assessing radiation exposure and dose prediction.
- All these diverse biomarkers will be further defined, validated and available for future use in case of nuclear accidents!
- Thanks to all of those radiobiology scientists who tirelessly work towards achieving the goals for human health and welfare.



Acknowledgements





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