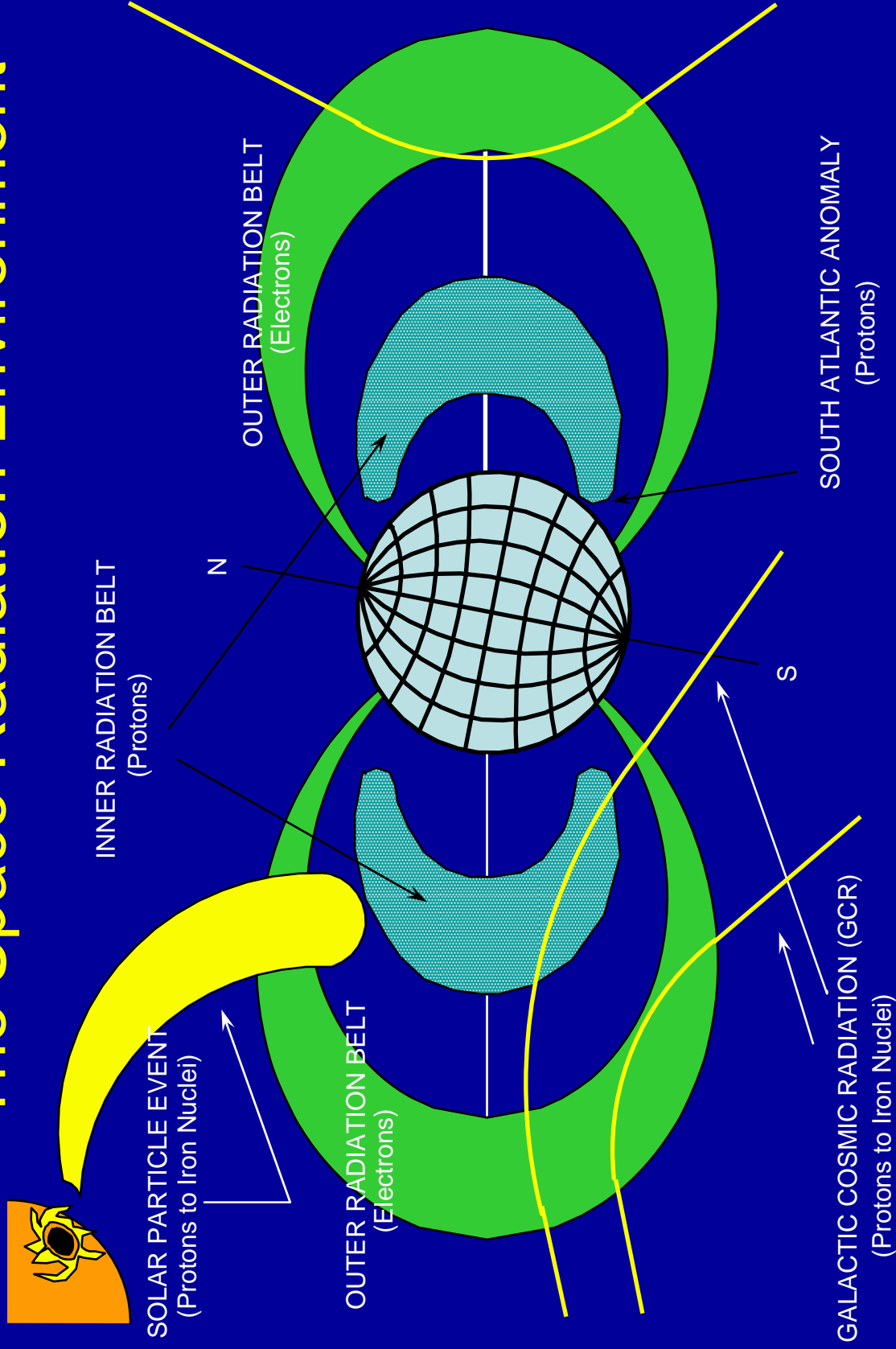




Painting analysis of chromosome aberrations induced by energetic heavy ions in human cells

Honglu Wu, Megumi Hada and Francis Cucinotta
NASA Johnson Space Center

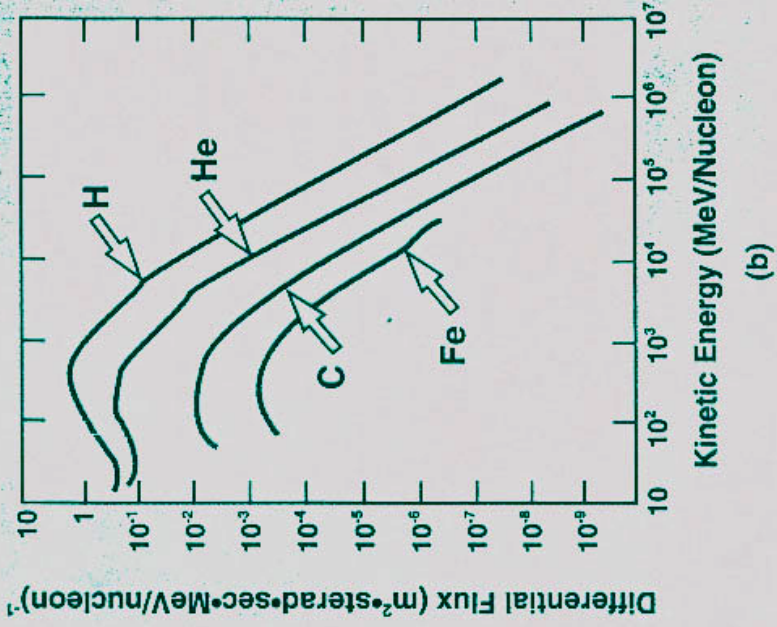
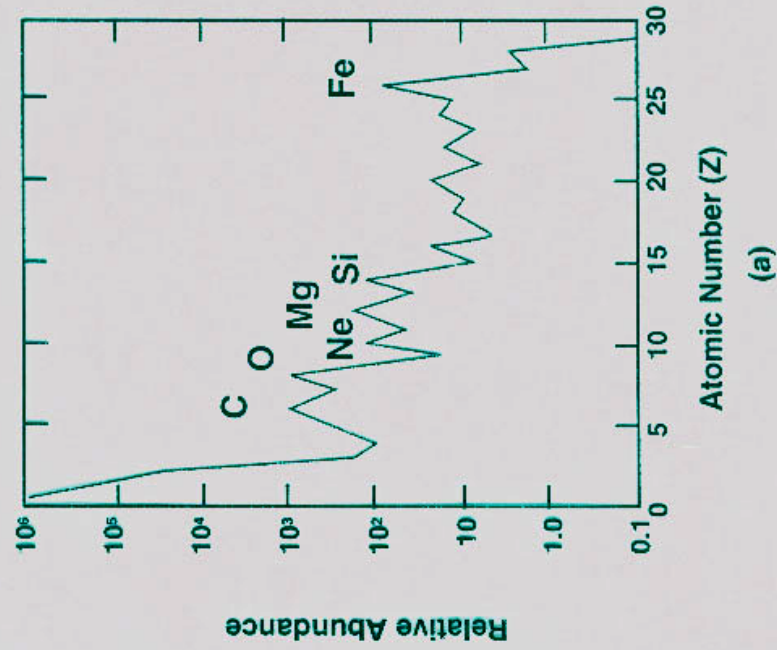
The Space Radiation Environment



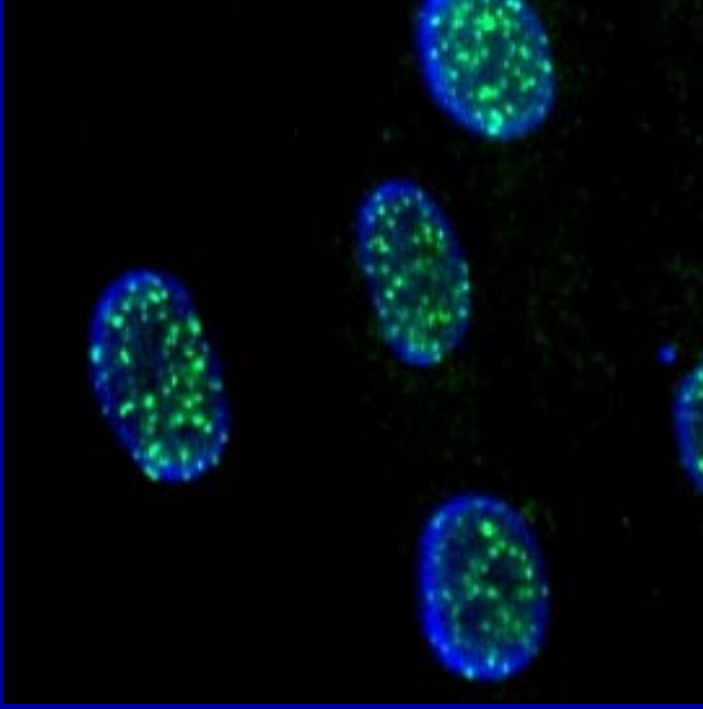
Representation of the major sources of ionizing radiation of importance to manned missions in low-Earth orbit. Note the spatial distribution of the trapped radiation belts.

Galactic cosmic radiation

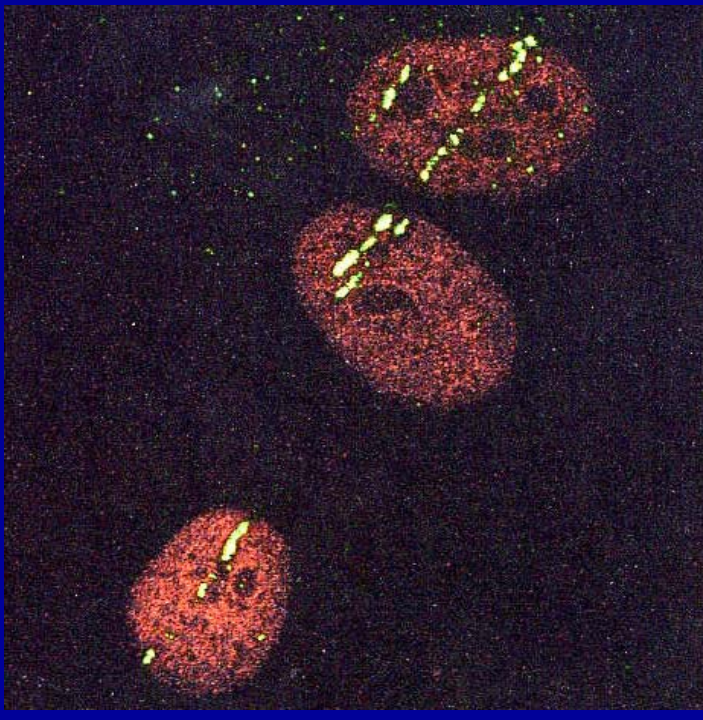
Figure D.1. Abundances (a) and Energy Spectra (b) of GCR



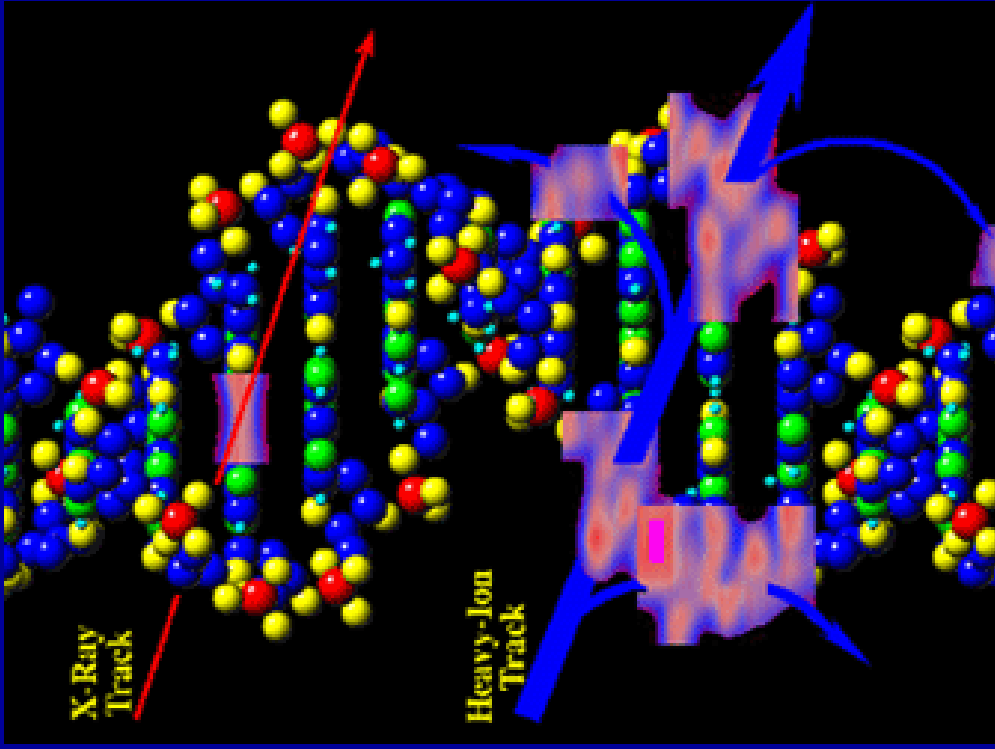
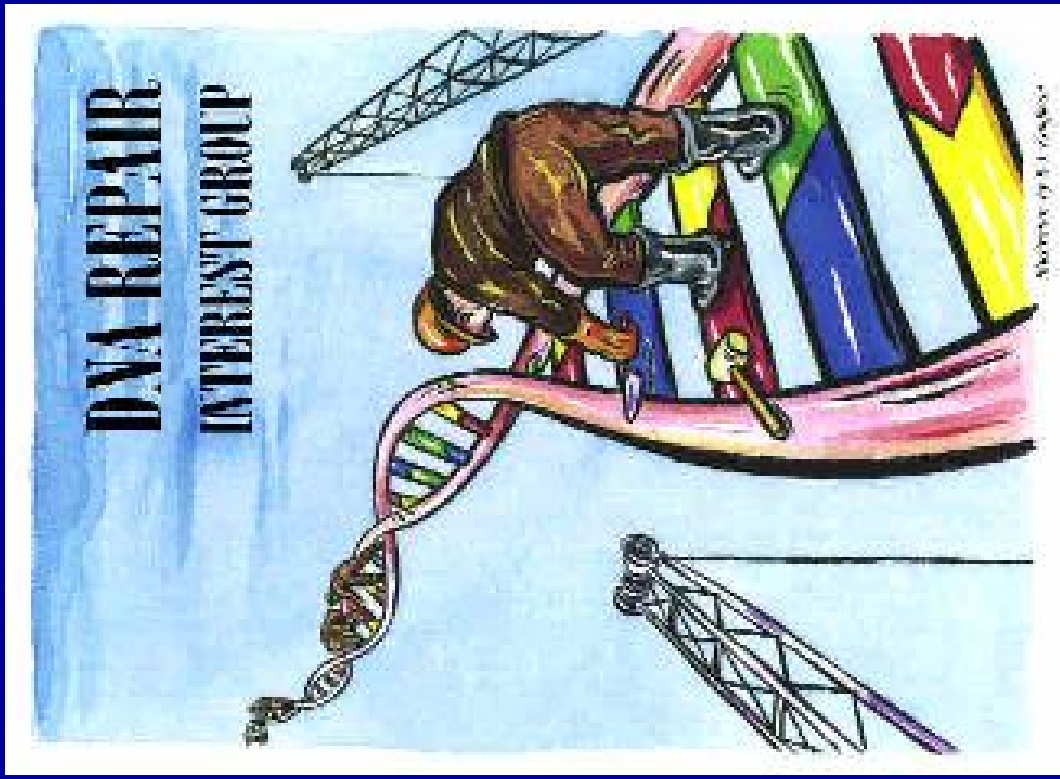
DSB induction



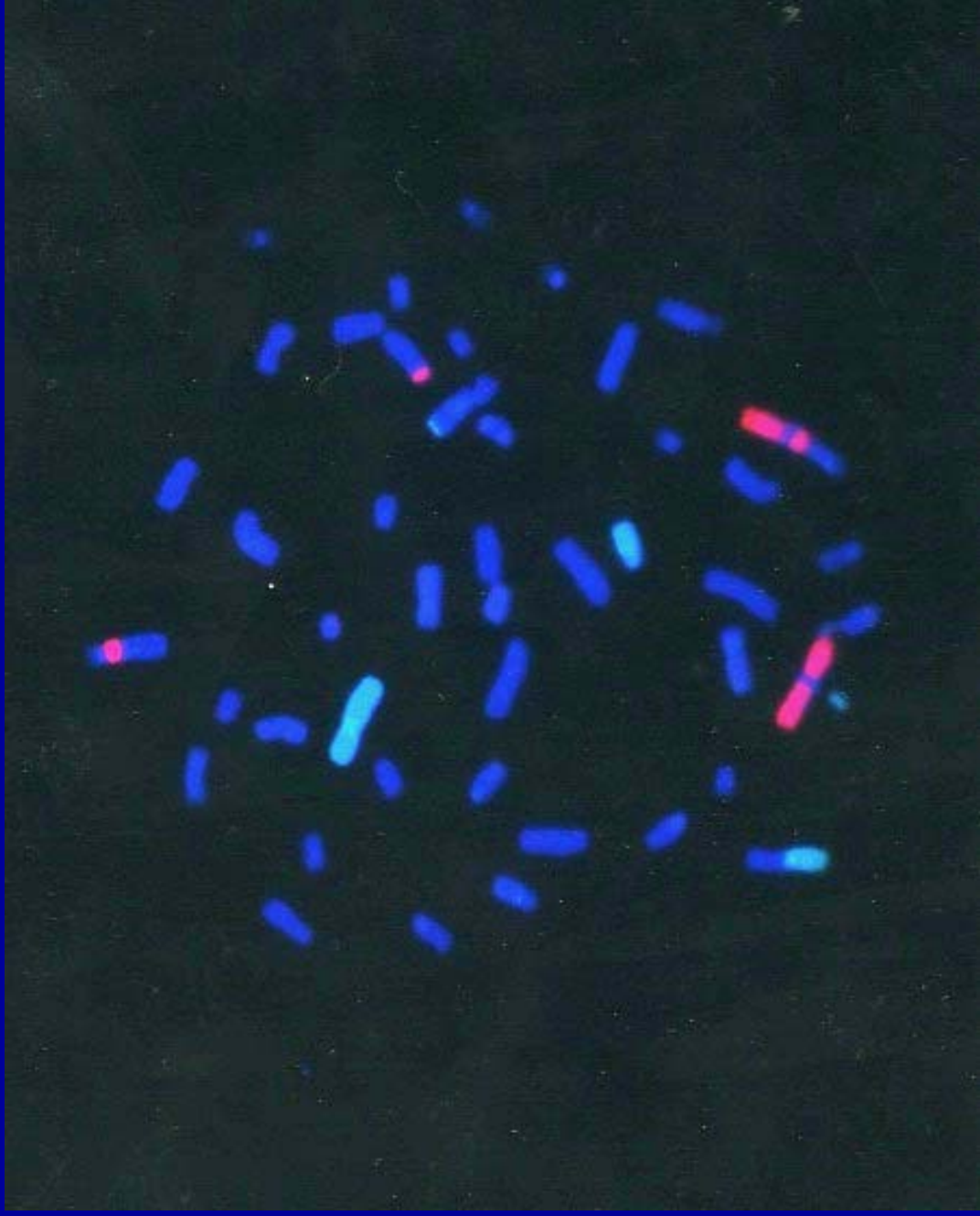
Low-LET



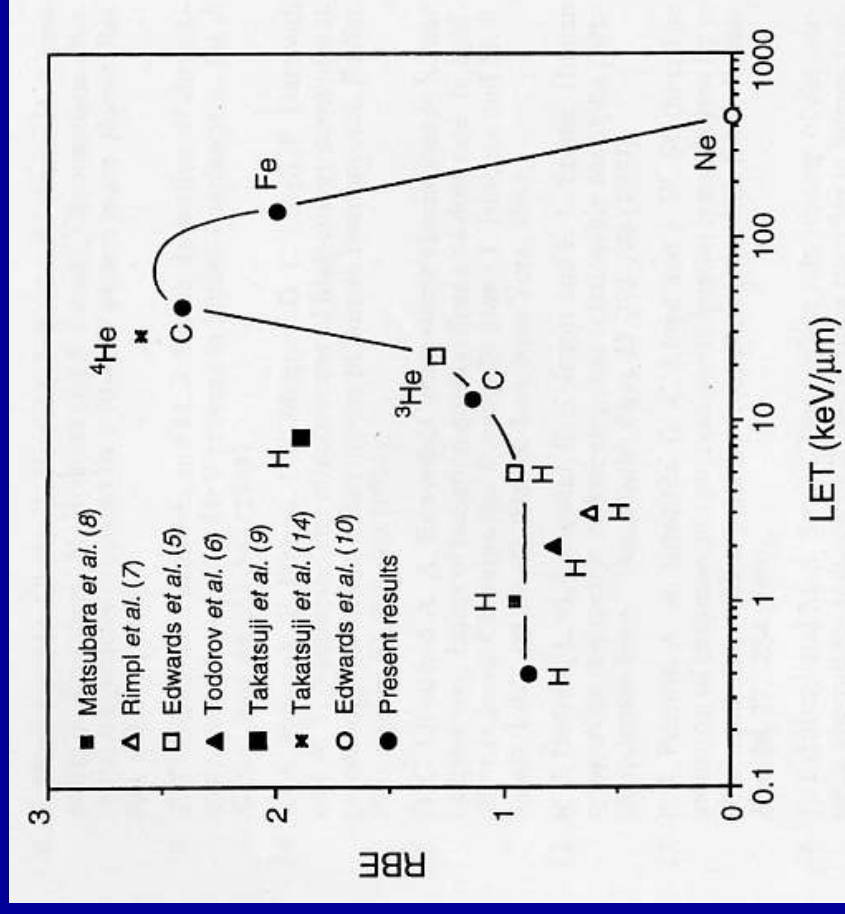
High-LET



Complex aberrations

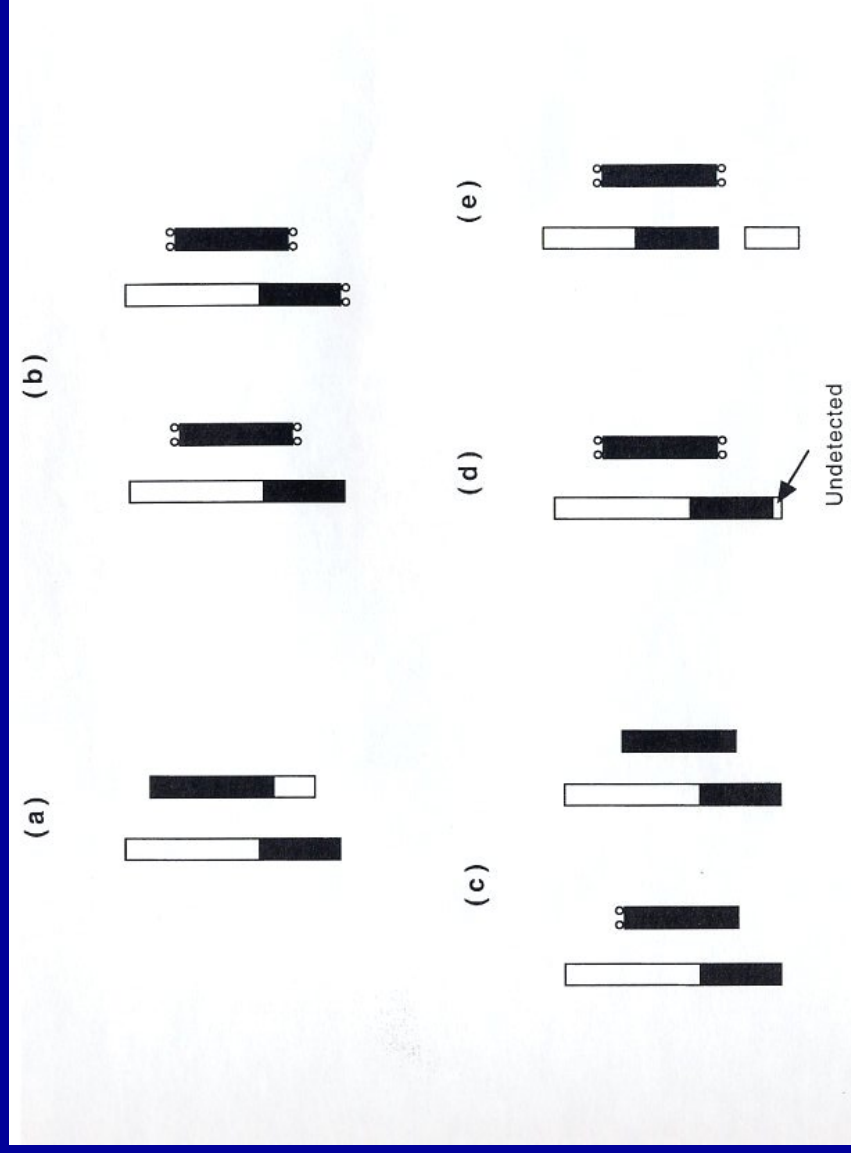


Radiation-induced chromosome aberrations in lymphocytes in vitro



Wu, Durante, George and Yang, *Radiat. Res.* (1997)

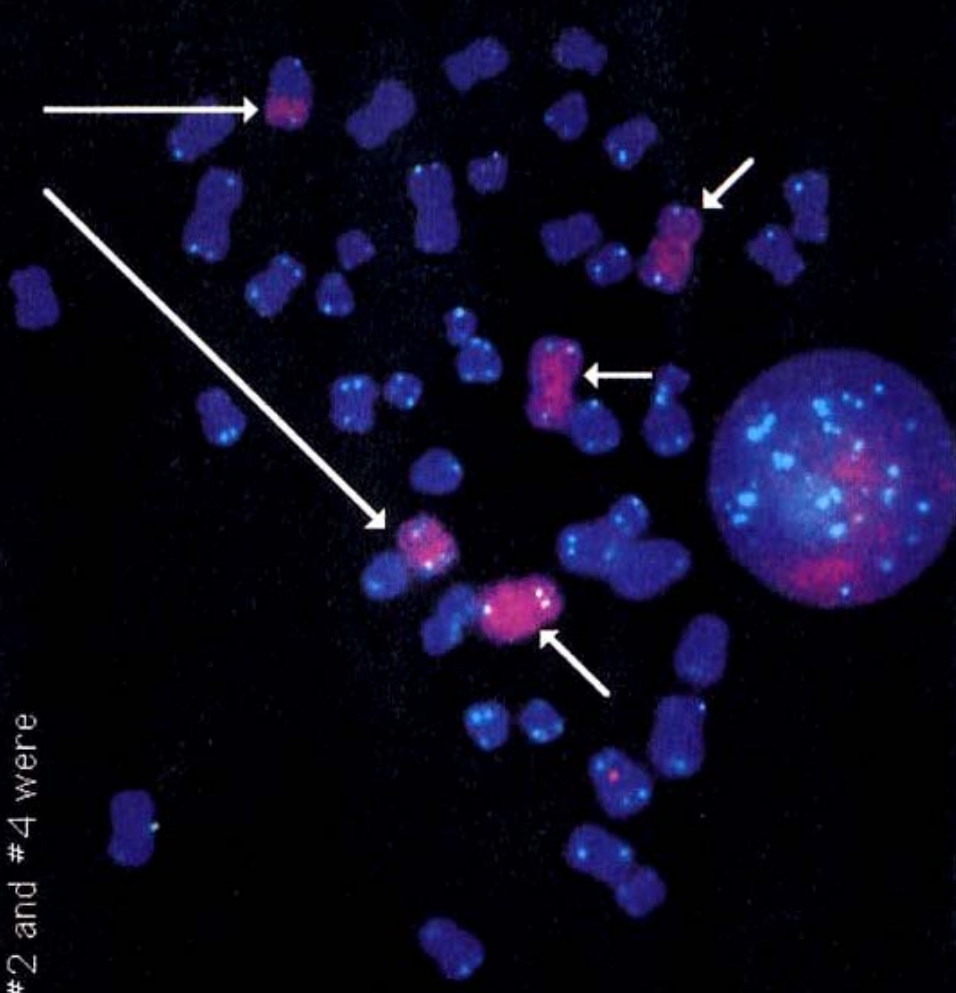
Analysis of truly incomplete exchanges using telomere probes



Telomere Analysis

Human lymphocytes exposed to 2 Gy gamma rays. Chromosomes #2 and #4 were painted.

False incomplete exchange



Most of the incomplete exchanges analyzed with FISH are actually complete

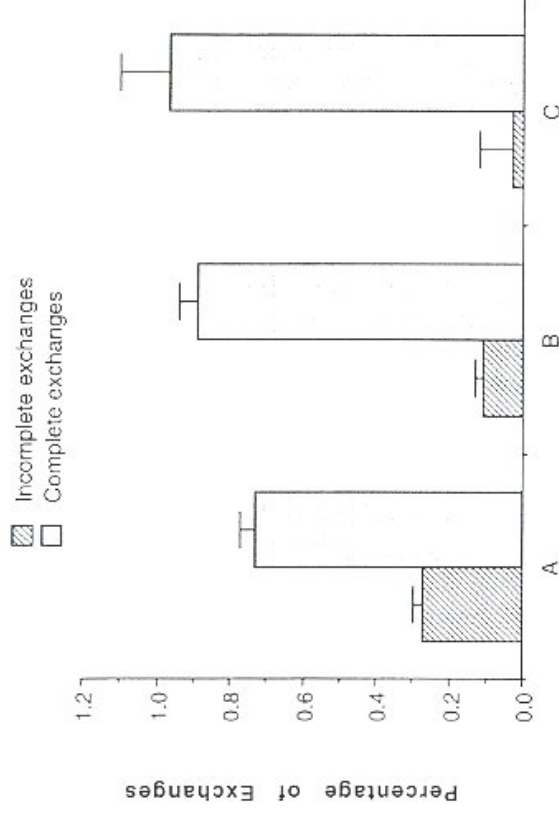
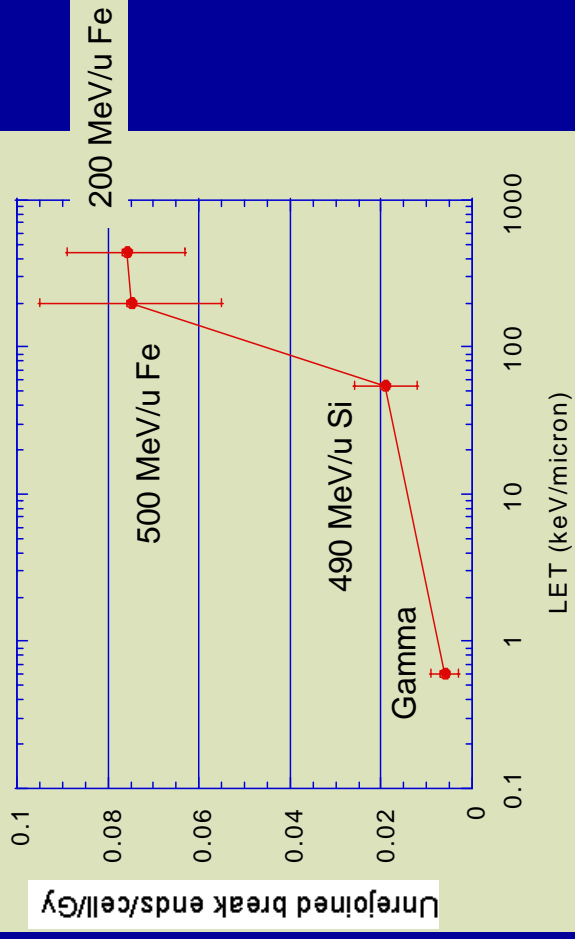


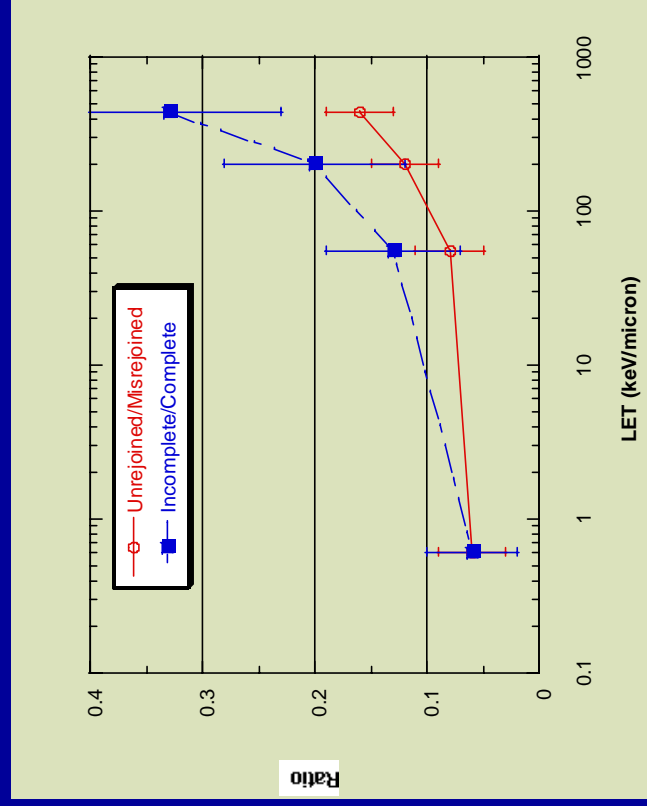
Figure 4. Percentage of complete and incomplete exchanges from the sum of the data. (A) The percentage of incomplete exchanges was 27% without the consideration of telomere probes. (B) With false incoherent telomere probes included as complete, the percentage of incomplete exchanges decreased to 11%. (C) The estimated percentage of true incomplete exchange was 3%. (bar = 1 SD)

Wu, George and Yang, IJRB (1998, 1999)



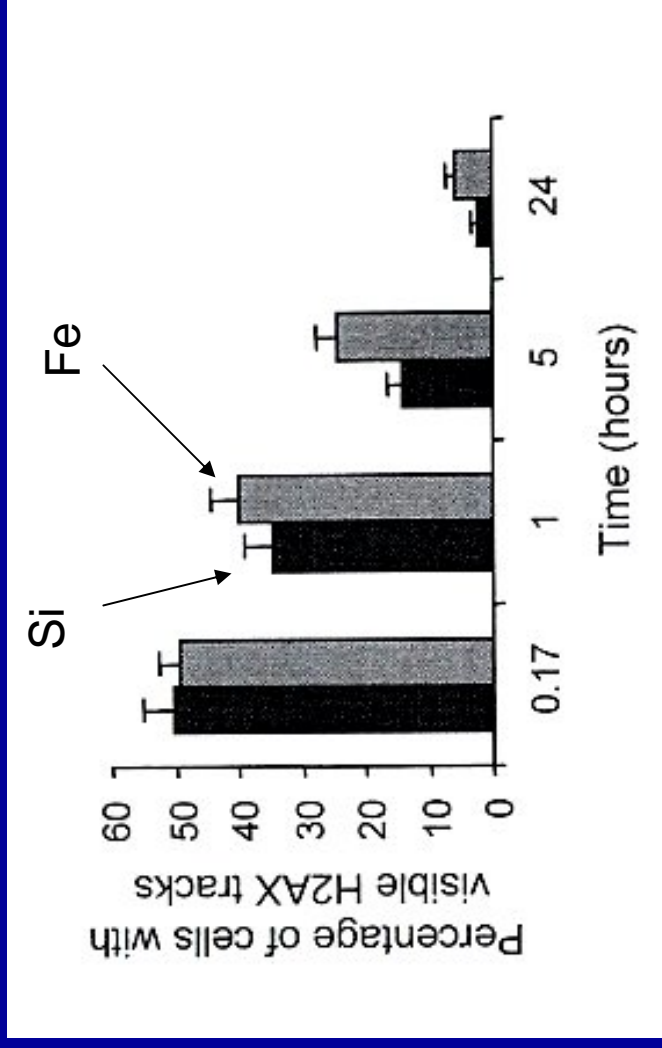
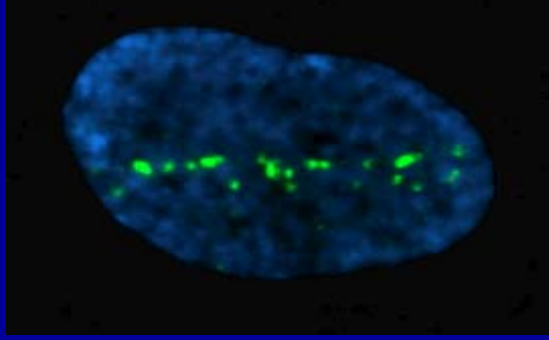
- The fraction of unrejoined chromosome breaks are higher for high LET

- Unrejoined breaks and incomplete chromosomal exchanges are possible biosignatures of high-LET radiation



Wu, Durante, Furusawa, George, Kawata and Cucinotta, Rad. Res. (2003)

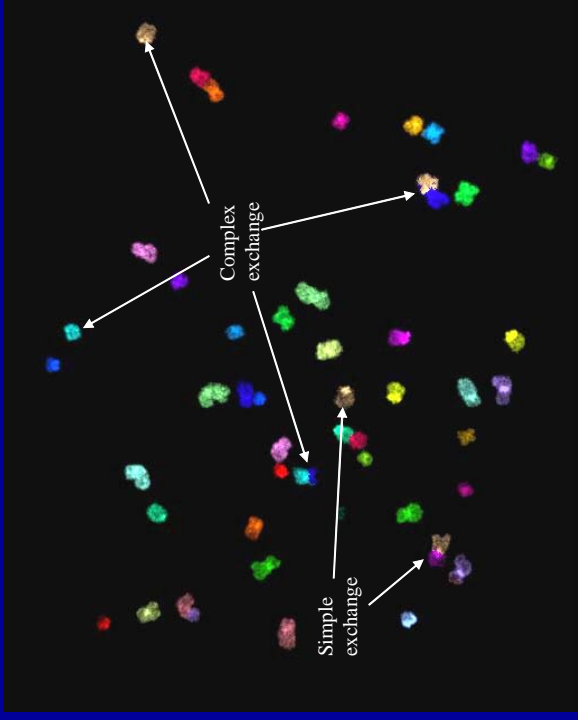
High-LET radiation induces more unrejoined DNA double strand breaks



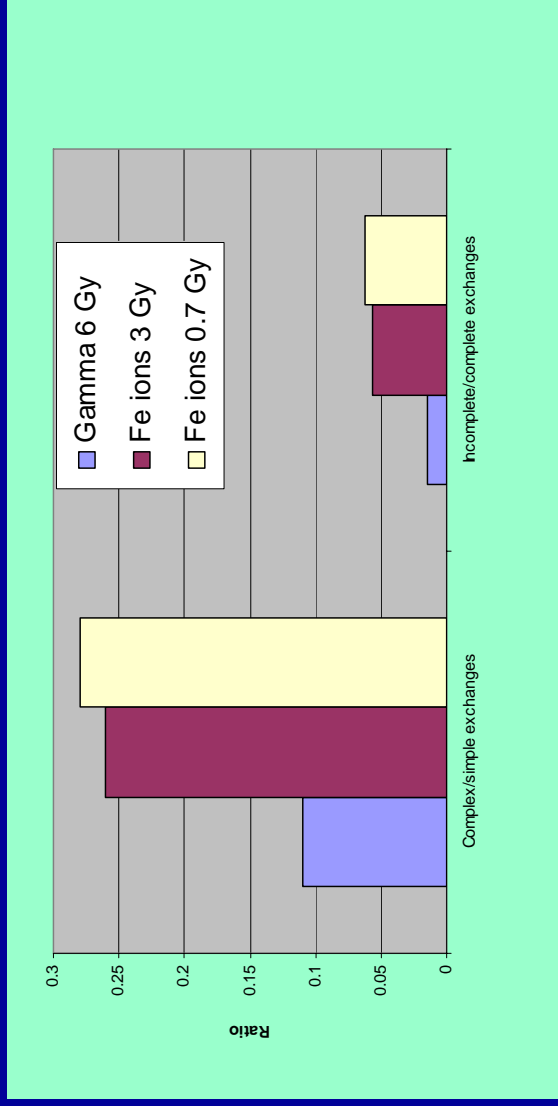
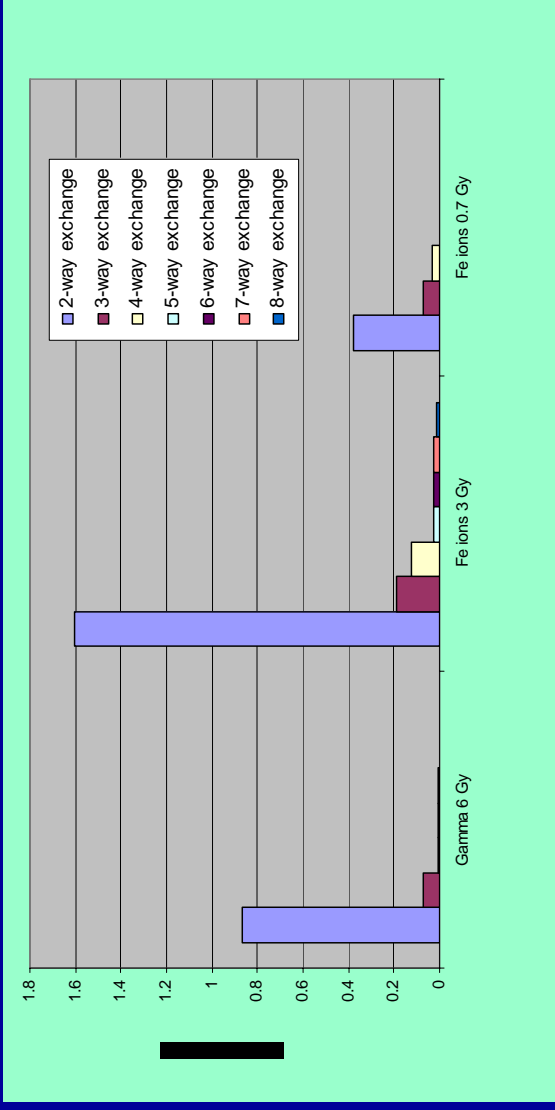
Desai, Davis, O'Neill, Durante, Cucinotta and Wu, Rad. Res. 2005

Complex aberrations -- mFISH analysis

BIOSIGNATURE OF HIGH-LET RADIATION

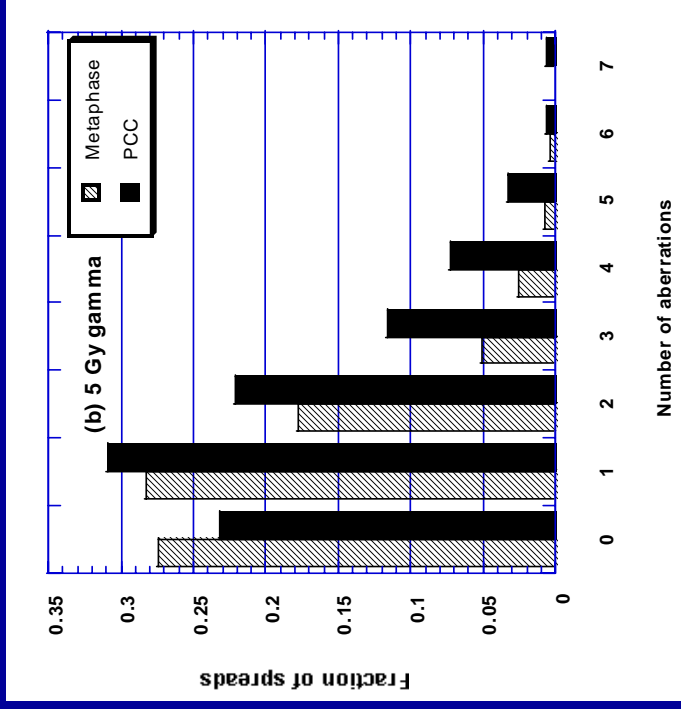


Complex type aberrations



mFISH showed a higher fraction of complex and incomplete exchanges for high-LET

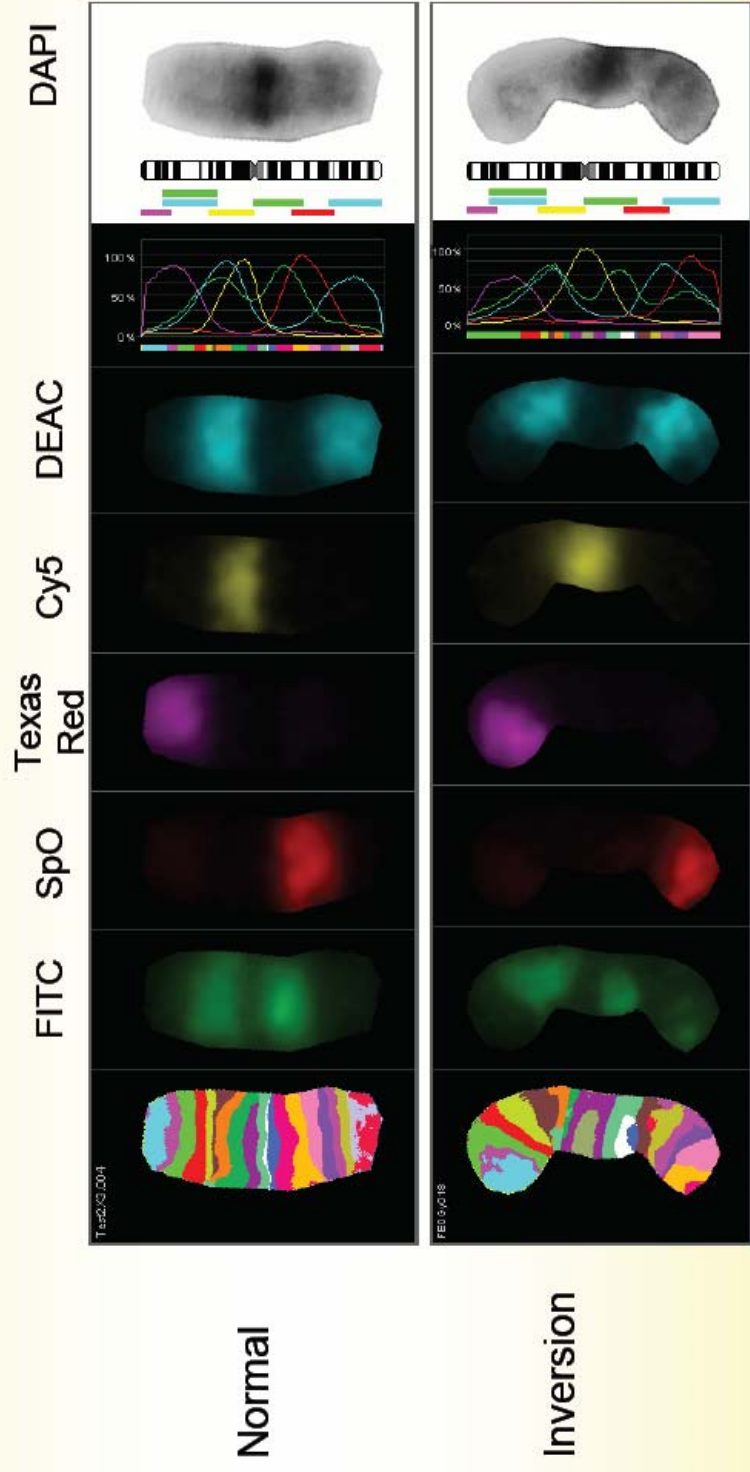
Interphase vs. metaphase: Issues of biosignature



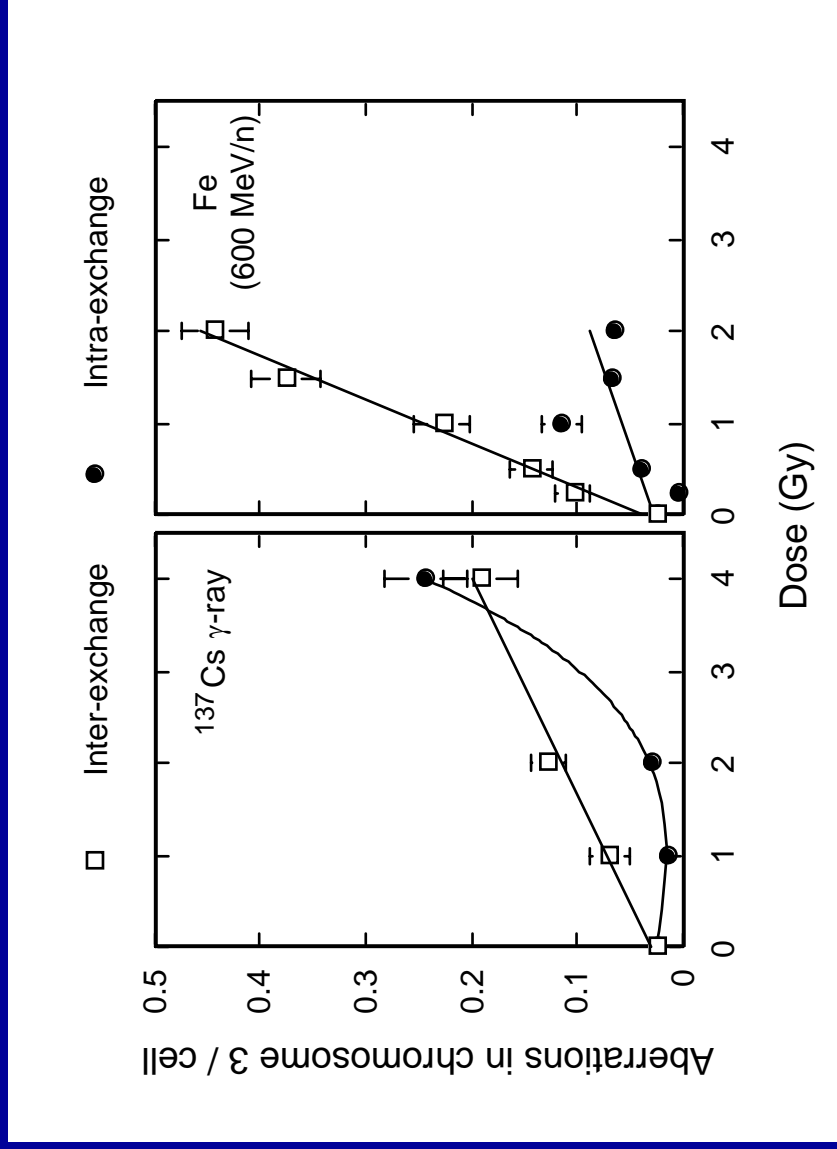
Centromere probes were used.

Radiation	Dose (Gy)	Harvest method	F ratio
γ ray	2	PCC	15.3±6.3
γ ray	2	Meta	12.5±5.9
γ ray	5	PCC	8.2±2.0
γ ray	5	Meta	9.1±2.5
1 GeV/u Fe	3	PCC	5.2±0.9
1 GeV/u Fe	3	Meta	9.1±2.2

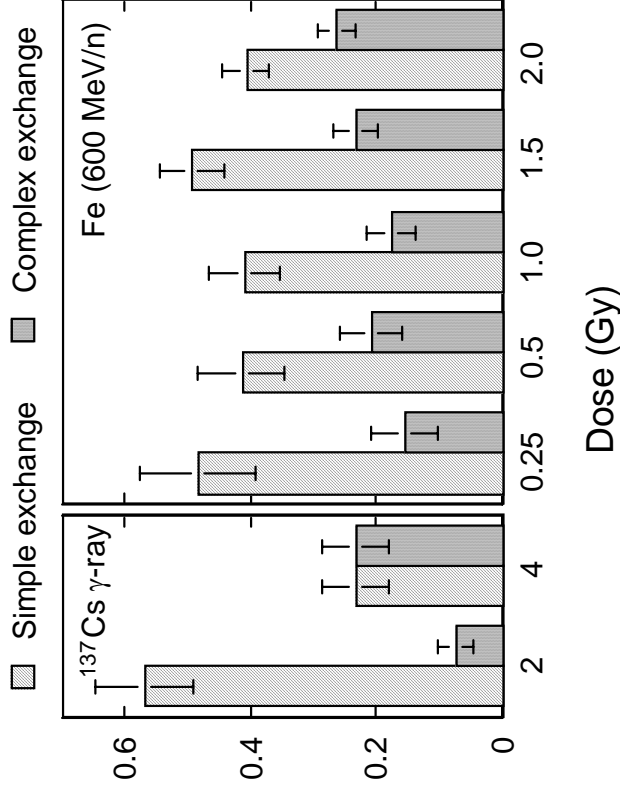
mBAND analysis



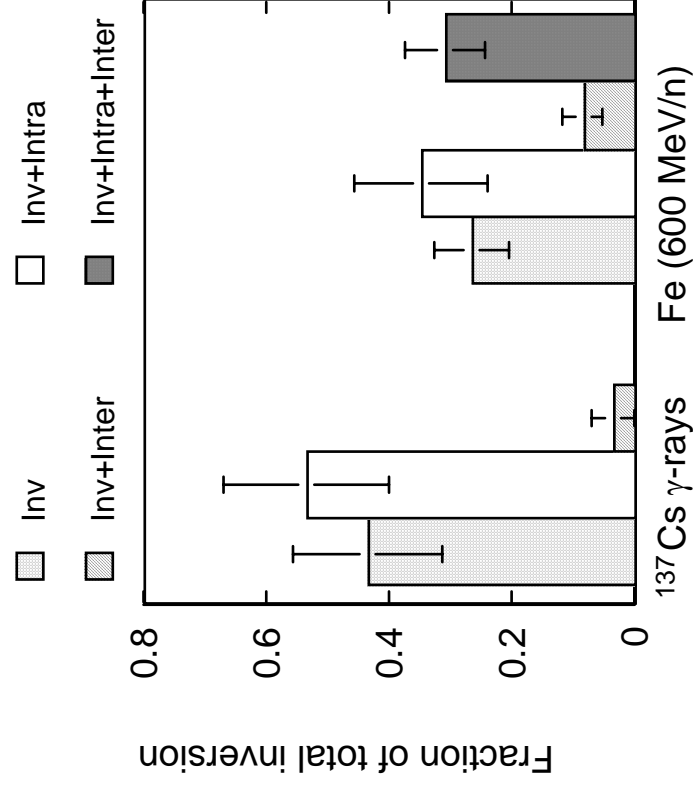
Inter- vs. intra chromosome exchanges (mBAND)



Fraction in damaged chromosome 3



mBAND analysis



Most inversions were involved with other inter- and/or intra-chromosome rearrangements

Summary

- FISH, mFISH, mBAND, telomere and centromere probes have been used to study chromosome aberrations induced in human cells exposed to low- and high-LET radiation in vitro
- High-LET induced damages are mostly a single track effect
- Unrejoined chromosome breaks (incomplete exchanges) and complex type aberrations were higher for high-LET
- Biosignatures may depend on the method the samples are collected
- Recent mBAND analysis has revealed more information about the nature of intra-chromosome exchanges

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Marco Durante
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Veronica Willingham

