

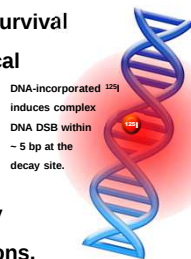
CHROMOSOME ABERRATIONS INDUCED BY THE AUGER ELECTRON EMITTER ^{125}I

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Introduction

- DNA-associated Auger electron emitters (AEE) induce cellular damage leading to high-LET type cell survival curves and possess enhanced relative biological effectiveness.
- DNA double strand breaks (DSB) induced by Iodine-125-deoxyuridine (^{125}I -UdR) decays are claimed to be very complex, thus efficiently leading to cell transformation and gene mutations.
- To elucidate the assumed genotoxic potential of DNA-associated AEE, chromatid aberrations (CA) were analyzed in ^{125}I -UdR-exposed human blood lymphocytes (PBL).



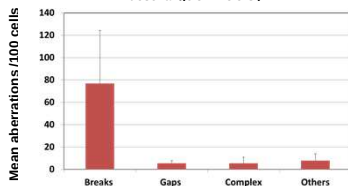
Summary

- ^{125}I -UdR has an explicit genotoxic capacity in human PBL, even at very low doses of about 0.2 Gy (29 decays/cell).
- Cells bearing a critical amount of unrepaired breaks, representing accumulated lethal damages due to high activity concentrations of ^{125}I -UdR, die due to apoptosis.
- Every fifth ^{125}I decay in the DNA gives rise to a single CA, which is in good agreement with data from Sedelnikova et al. (2002) and Yasui et al. (2004), stating that one decay gives rise to ~ 0.26 DSBs.
- Each ^{125}I -UdR-induced DSB is converted into one CA.**

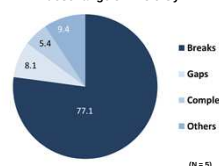
Results

Induction of chromatid aberrations in lymphocytes

A Frequency of chromatid-type aberrations dose range 3.1 – 3.5 Gy

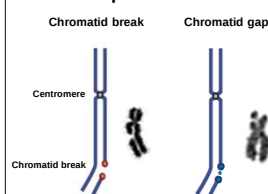


B Distribution of all detected CA dose range 3.1 – 3.5 Gy



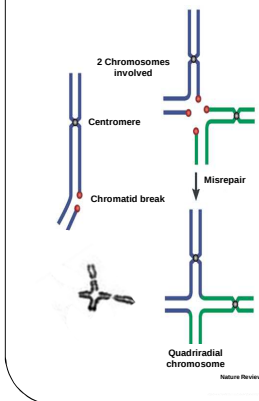
Aberration types

A Simple aberrations

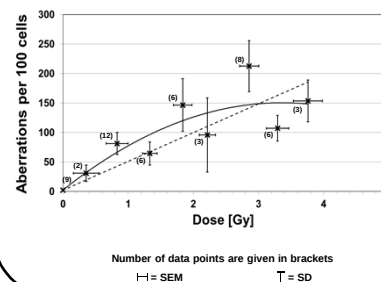


B Complex aberration

Asymmetrical chromatid interchange

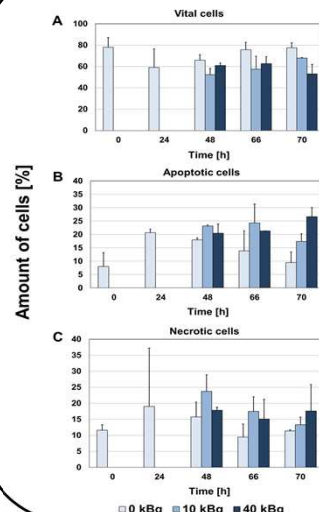


Dose-Response-Relationship



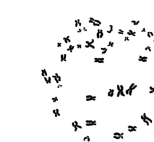
Dose-response relationship for chromatid-type aberrations induced by I-125-UdR (N=9). The solid line represents the polynomial fit ($Y = -13.11x^2 + 88.10x + 2.43$; $R^2 = 0.72$), whereas the dashed line shows the linear fit ($Y = 48.72x + 2.43$; $R^2 = 0.58$).

Induction of Apoptosis

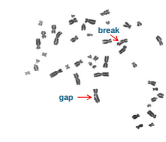


Percent of viable (A), apoptotic (B) and necrotic (C) cells at different time points: After isolation (0 h) of PBL, after stimulation with PHA and before incubation with I-125-UdR (24 h), after removing of I-125-UdR (48 h), immediately before addition of Colcemid (66 h), and before harvesting (70 h) of PBL. Error bars represent the SD.

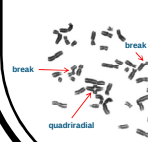
A Control metaphase



B Moderately damaged



C Severely damaged



D Highly damaged



Material and Methods

PBL were isolated from whole blood and stimulated with chromosome medium containing phytohaemagglutinin (PHA). After 24 h cultures were labelled with I-125-UdR for 18 h (1- 45 kBq) during the S-phase of the cell cycle. After removal of radioactive medium and washing steps, cells were cultured in stimulation medium for further 24 h. Colcemid was added 5.5 h before harvest of cells followed by fixation for aberrations at 71.5 h post-stimulation. All slides were stained with 10 % Giemsa, and at least 100 metaphases were analyzed microscopically for each activity concentration. Apoptosis was measured by Annexin V-FITC Assay using flow cytometry.