

X-射线离体照射人体淋巴细胞诱发的染色体畸变：剂量一效应关系的研究

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摘要

本文研究了180千伏X-线离体 (in vitro) 照射人体淋巴细胞诱发染色体畸变的剂量一效应关系。全血样本在离体条件下 (恒温37±0.5 0C) 接受不同剂量(0-465拉特) 照射。用微量血培养技术培养48-56小时, 观察和分析的中期细胞均为受射后的第一次有丝分裂。按WHO的标准识别各类染色体畸变, 用最小二乘法作泊森方差和加权回归分析, 对实验资料配以4个剂量一效应关系模式: $Y=a_1+b_1D$, $Y=a_2+c_2D^2$, $Y=a_3+b_3D+c_3D^2$ 和 $Y=b_4D^n$ 。结果表明, 双着丝点体最适于配二次多项式, $a_3=0$ (回归线通过原点), $Y=(0.52+0.18) \cdot 10^{-3}D + (4.71 \pm 0.67) \cdot 10^{-6}D^2$; 也适于配幂函数, $Y=7.10 \cdot 10^{-5}D^{1.59+0.08}$ 。双着丝点体+着丝点环同样适于配以上二个模式, 分别为 $Y=(0.51+0.21) \cdot 10^{-3}D + (5.02+0.77 \cdot 10^{-6}D^2)$ 和 $Y=6.50 \cdot 10^{-1.61+0.05}$ 。在二次多项式中, 样本回归系数b,和c, 与总体回归系数为“0”有显著差别; 在幂函数中, 剂量幂次 $n \sim 0$, $n \sim 2$, 而是界于1和2之间。对一次击中畸变 (末端缺失), 最适于配直线模式, $a_1=0$ (回归线通过原点), $Y=(5.86+0.55) \cdot 10^{-4}D$; 也可配以幂函数, $Y=1.47 \cdot 10^{-3.81+0.153}$ 相互易位十臂间倒位仅能配以幂函数, $Y=3.34 \cdot 10^{-4}D^{0.87+0.20}$ 。

文中较详细地分析了电离辐射诱发的各类染色体畸变和剂量之间的关系, 以及影响畸变量的诸因素。最后作者认为, 双着丝点体或双着丝点体+着丝点环最适于用作生物剂量测定; 而在较低剂量的情况下, 或可用末端缺失作为剂量测定的指标, 但有待进一步的研究。

关键词

分类号

X-RAY INDUCED CHROMOSOME ABERRATIONS IN HUMAN PERIPHERAL BLOOD LYMPHOCYTES IRRADIATION IN VITRO: STUDIES ON DOSE-RESPONSE RELATIONS

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Abstract

The relationship between X-ray induced chromosome aberrations in human peripheral blood lymphocytes in vitro and radiation dose has been studied. Whole blood samples were irradiated with various (0-465 rads) of 180 kV X-rays at 37±0.50C, and cultured for 46--54 hr by microtechnique. In all cases aberrations were scored in metaphase cells observed in their first post-irradiation mitotic division in culture All chromosome-type aberrations were classified into: polycentrics, centric rings, acentric rings, minutes, reciprocal translocations, pericentric inversions and terminal deletions. Depending on WHO's criteria, the resulting data were fitted to the four models, $Y=a_1 + b_1D$, $Y = a_2 + c_2 D^2$, $Y = a_3 + b_3D + c_3D^2$ and $Y = b_4D^n$ by least square regression analysis using Poisson variance and weights. The dicentric data gave the best to the second degree polynomial model with $a_3 = 0$ (the regression was constrained to pass through the origin), $Y = (0.52 + 0.18) \cdot 10^{-3} D + (4.71 + 0.67) \cdot 10^{-6} D^2$. An equally

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good fit was obtained with the potential function, $Y=7.10 \cdot 10^{-5} D^{1.59}+0.08$. The dicentric plus centric ring data follow similar models to those for dicentric, $Y=(0.51 \pm 0.21) \cdot 10^{-3} D^{5.02 \pm 0.77}-10^{-6} D^2$ and $Y=6.50 \cdot 10^{-5} D^{1.61}+0.05$ respectively. The data for one hit aberration (terminal deletion) gave best fit to linear model with $a=0$ (the regression was constrained to pass through the origin). An equally good fit was obtained with the potential function, $Y=1.47 \cdot 10^{-3} D^{0.81}+0.15$. But the reciprocal translocation plus pericentric inversion was only fitted to the potential function, $Y=3.34 \cdot 10^{-4} D^{0.87}+0.02$. The relationship between aberration yield (Y) and radiation dose (D) was concerned in details. The possible importance of other technical factors in influencing the aberration yield was also discussed. From this analysis, we believe that the dicentric or dicentric plus centric rings may be the most efficient indicator for dose estimation at higher doses, however, the terminal deletions would be more useful than the asymmetric exchanges at lower doses for same aim.

Key words

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