

U-shaped Dose-Responses at Low Doses: Explanation with a New Model for *in vitro* Neoplastic Transformation

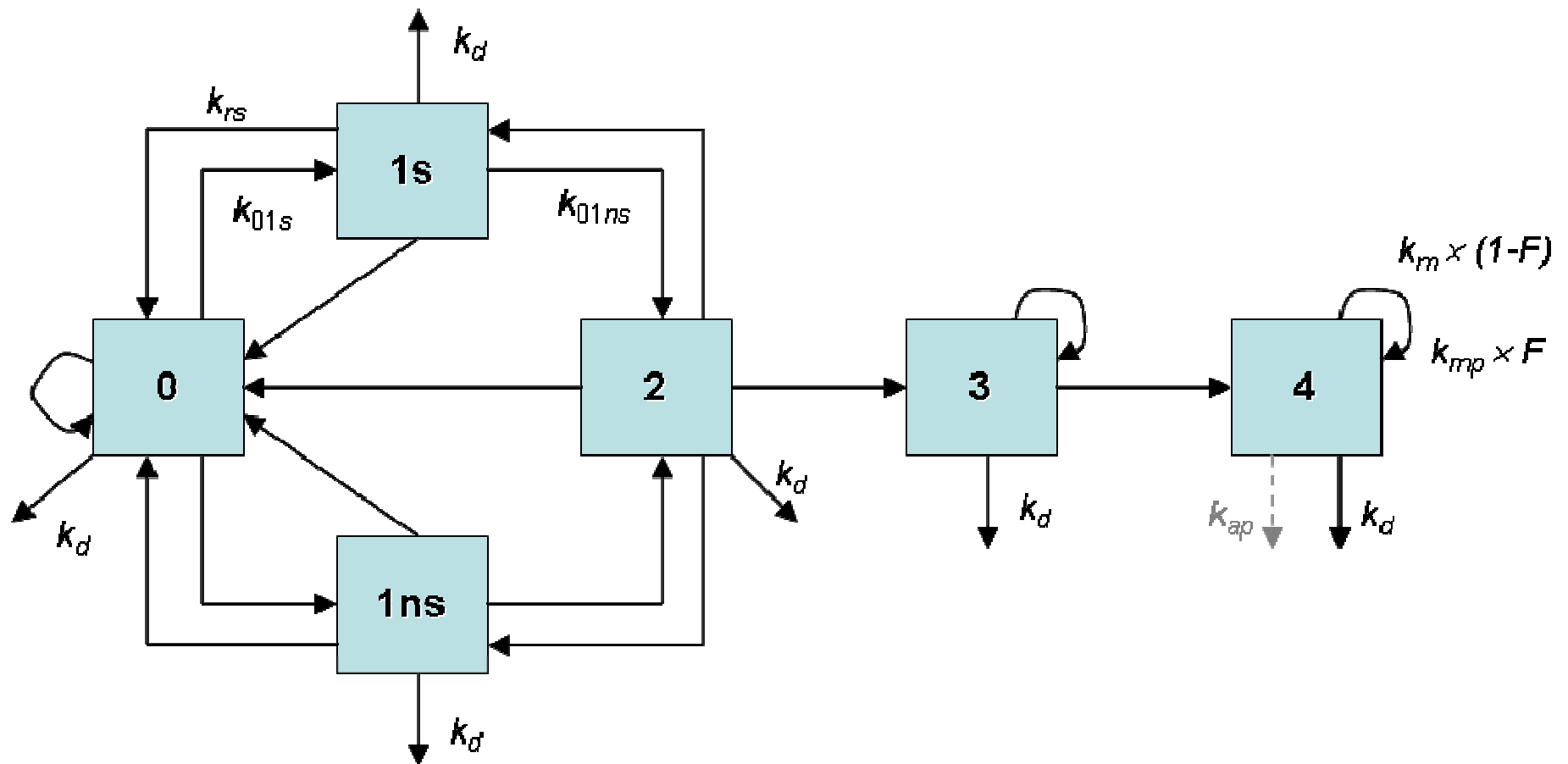
Helmut Schöllnberger



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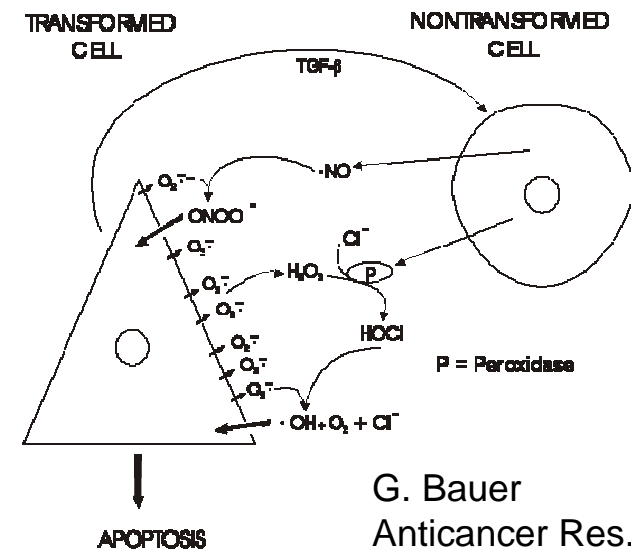
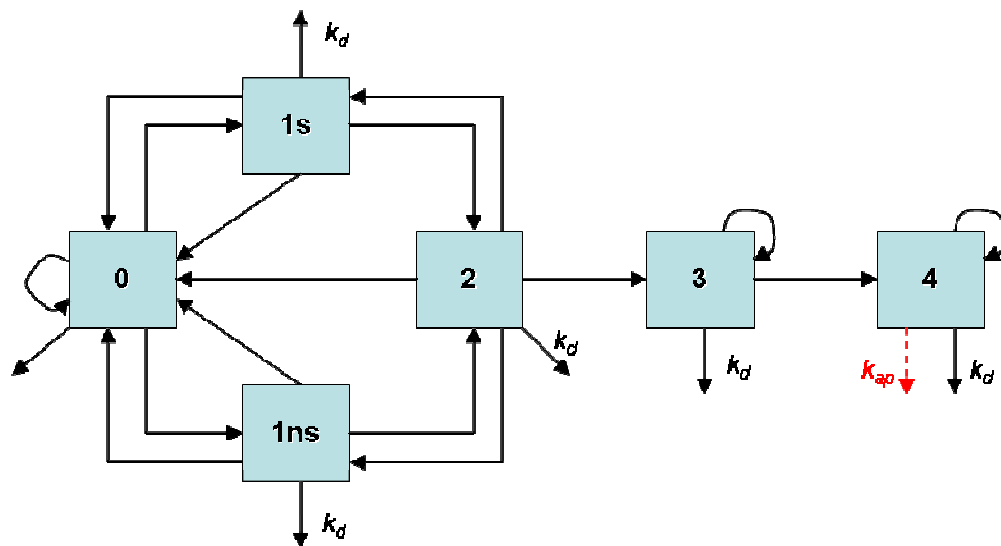
State Vector Model



Bystander-induced apoptosis

For low-LET radiation

- **P**rotective **A**poptosis-**M**ediated process (PAM)
- PAM can eliminate cells in State 4



Data by Redpath et al.

Radiat. Res. 2001

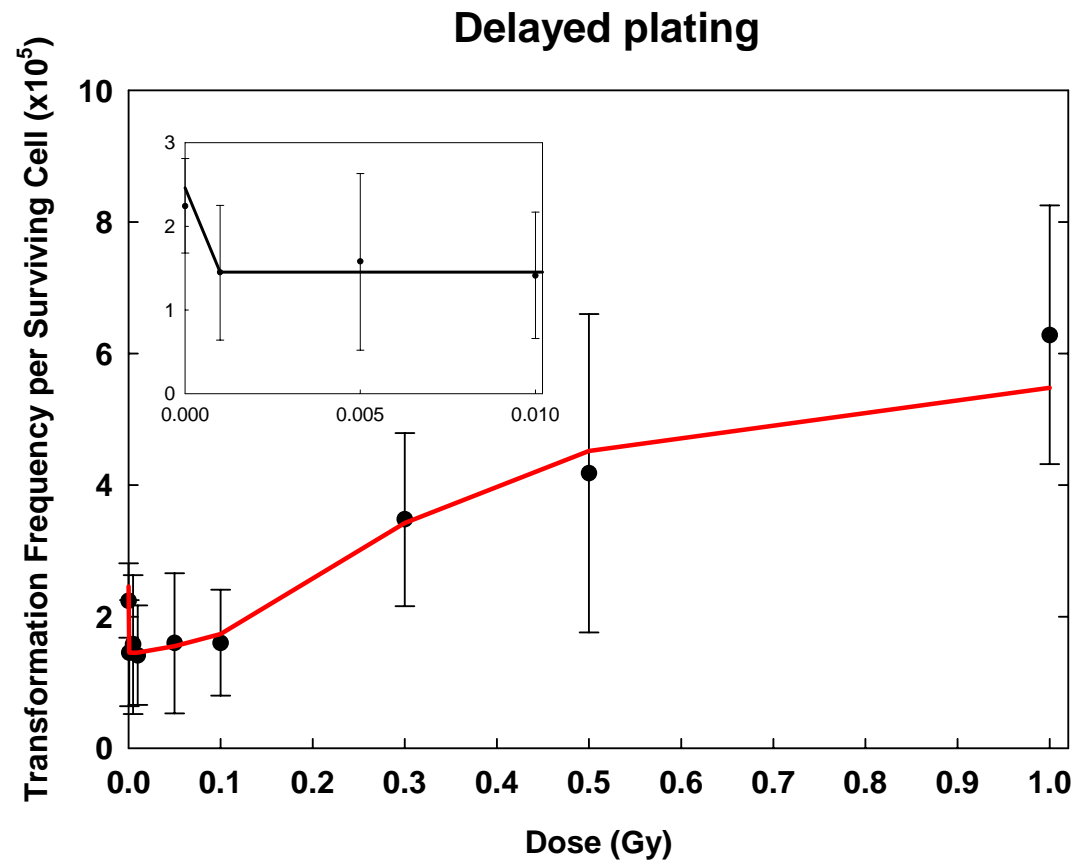
- CGL1 cells, γ -rays, neoplastic transformation
- **Irradiation period:** 3.3 mGy/min for $D \leq 100$ mGy
cell doubling time of 20 hrs
- **1 day holding period:** 20 hrs
- **10 days exponential growth:** 20 hrs
- **Confluent growth until day 26:** 38 days



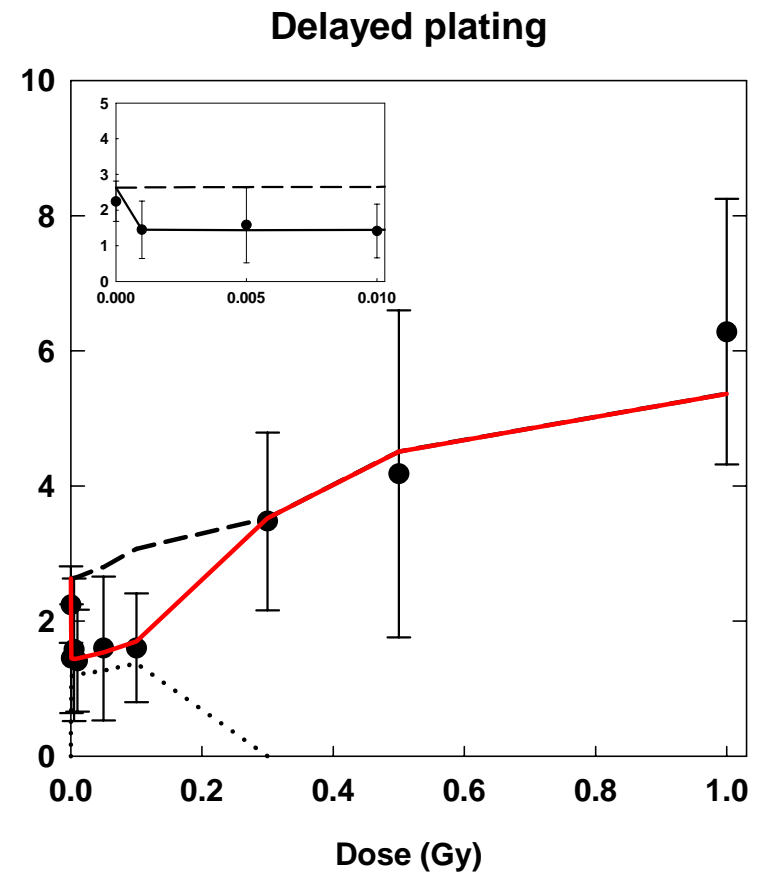
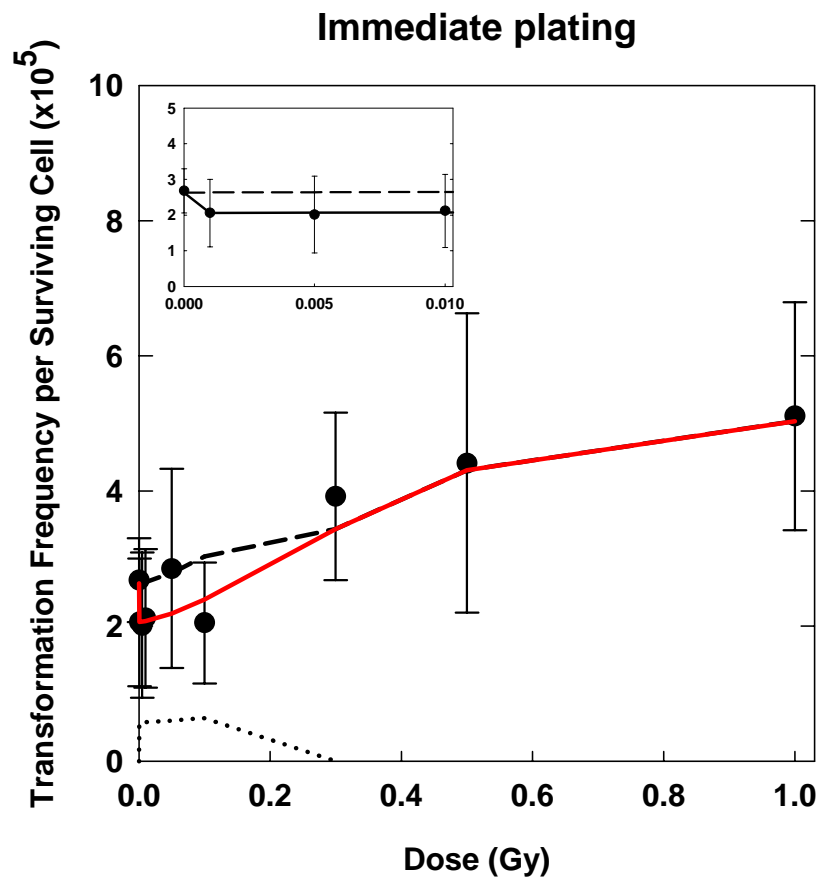
Fit approach

- TFSC =
$$\frac{N_4(t_{end})}{N_0(t_{end}) + N_{1s}(t_{end}) + N_{1ns}(t_{end}) + N_2(t_{end}) + N_3(t_{end}) + N_4(t_{end})}$$
- **Fit model without PAM** to control and high dose data („direct contribution“)
- **Fit model with PAM** to all data points for delayed plating: 1 to 3 free parameters: k_{ap} , t_{ap_on} , t_{ap_off} („total contribution“)

Data by Redpath et al.



Data by Redpath et al.



„Where is uncertainty?“

Perform local model fit → calculate the 95% CI for best estimated values with

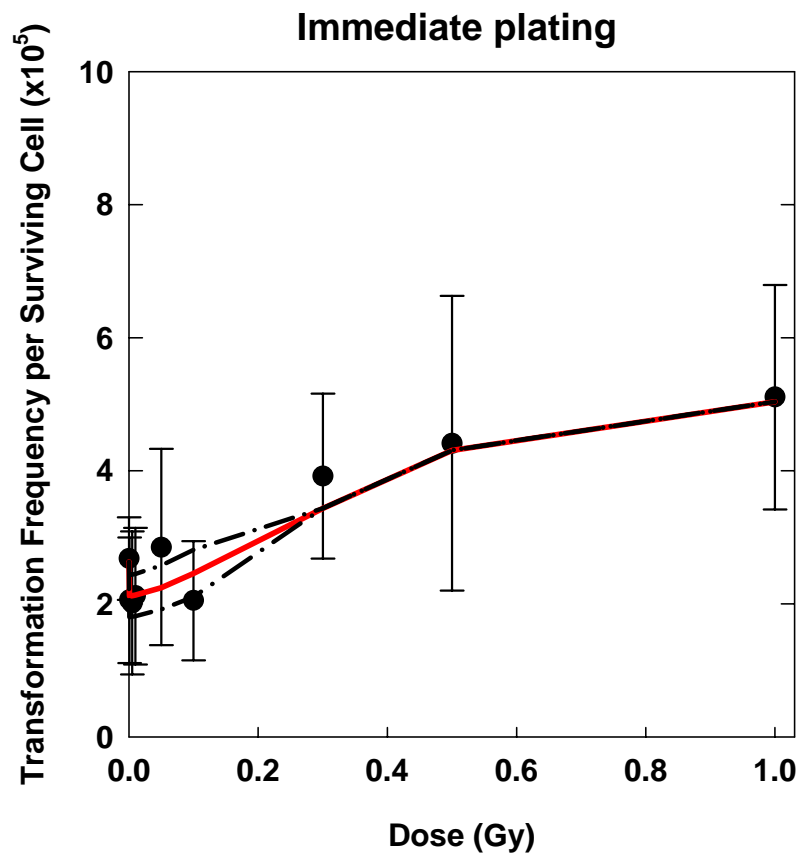
- the Jacobian matrix, $J_f = \left(\frac{\partial f_i}{\partial \hat{\beta}_j} \right)_{i=1, \dots, m; j=1, \dots, n}$
 - numerical model solutions, f_i , (i.e. TFSC_{*i*}),
 - best estimated values, $\hat{\beta}_j$, of free model parameters
- the residuals (measured TFSC – model predicted TFSC)

i... # of data points (i.e. residuals), *j*... # of free parameters

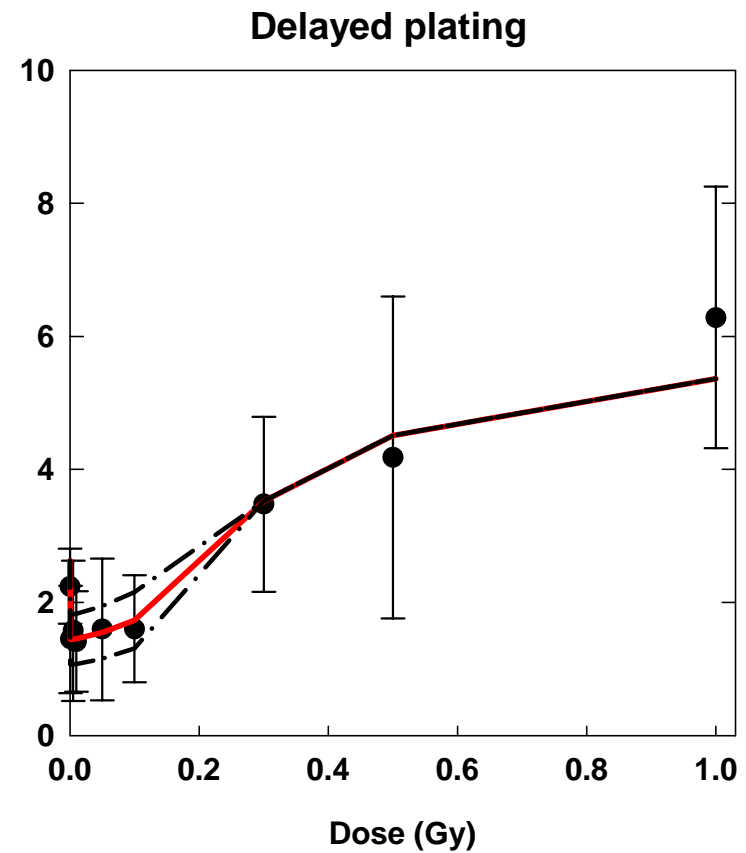
$\delta \propto$ 95%-tile of Student's t distribution with *v* degrees of freedom; *v* = # of residuals - # of free parameters.

$$95\% \text{ CI} = [\hat{\beta} - \delta; \hat{\beta} + \delta]$$

Data by Redpath et al.



$$k_{ap} = 0.054/\text{day} (0.031-0.078)$$

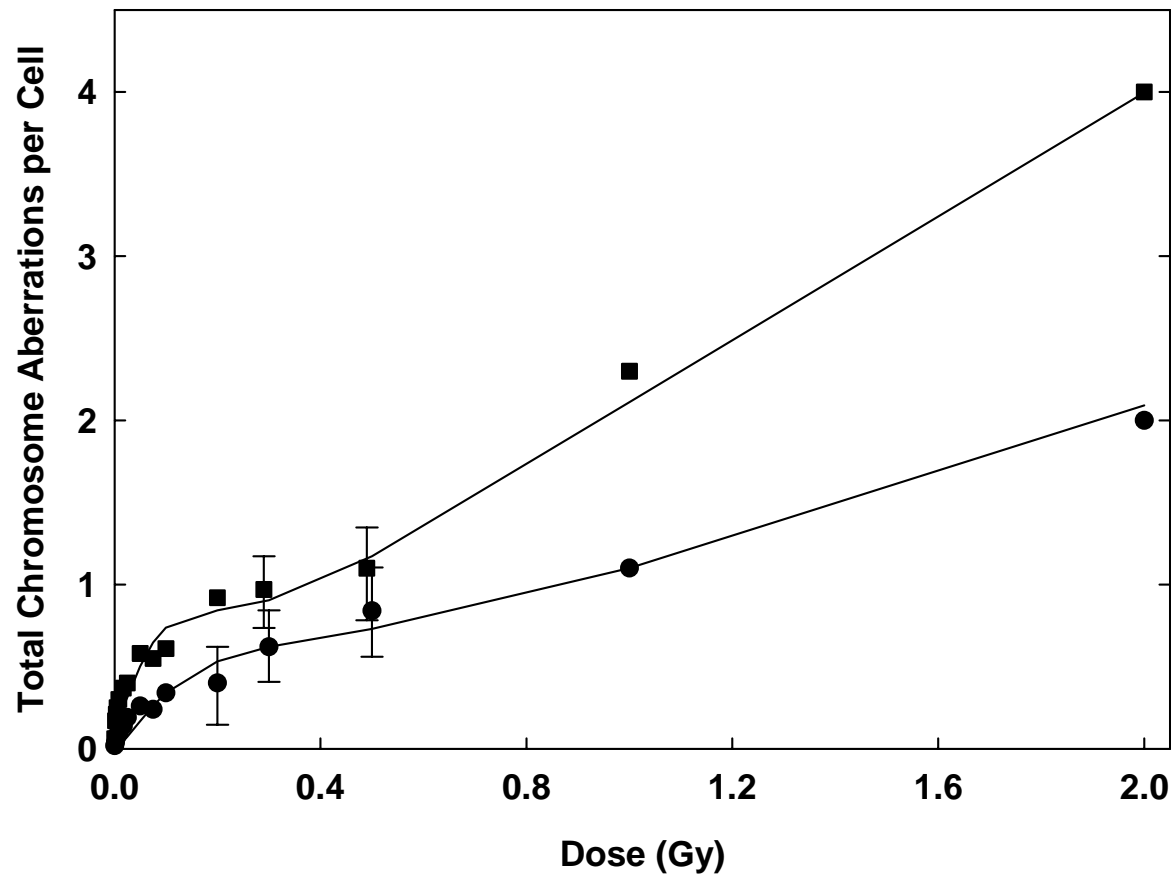


$$k_{ap} = 0.022/\text{day} (0.007-0.036)$$

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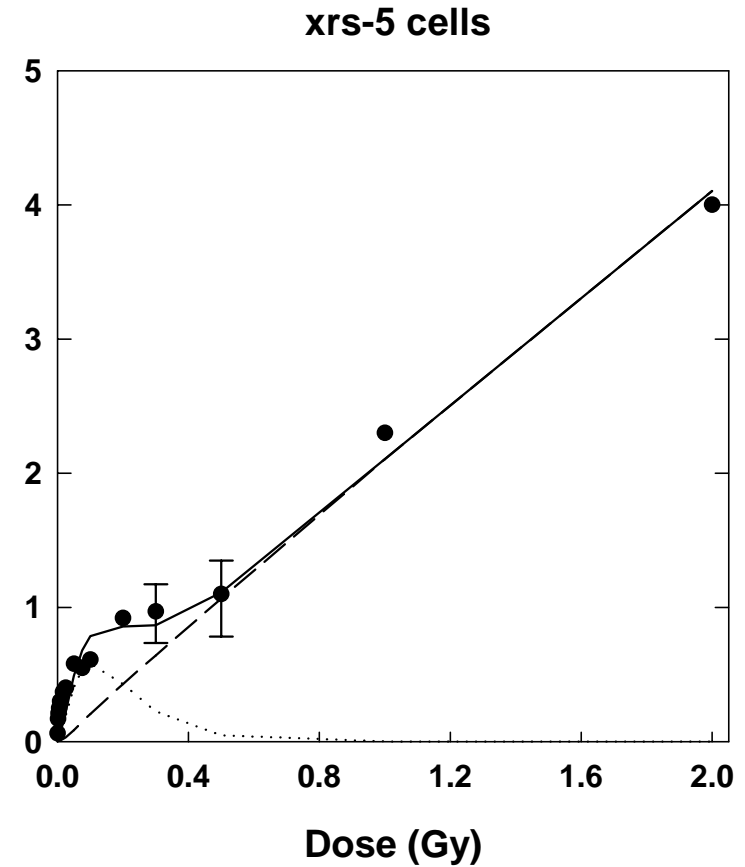
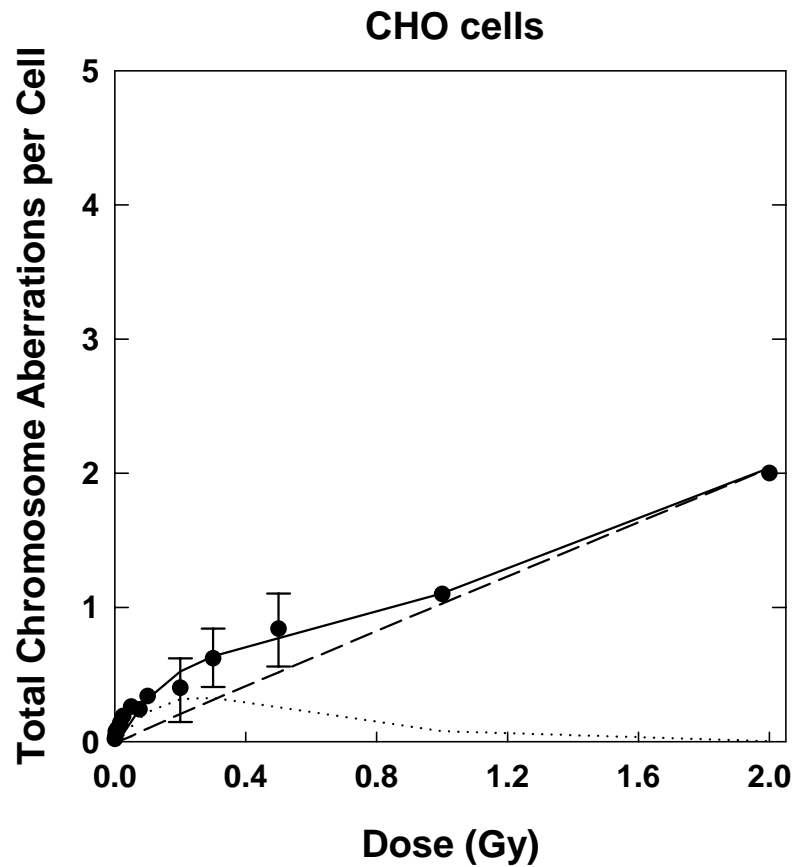
Data by Nagasawa and Little

Mutat Res. 2002, α -particles (112 keV/ μ m), CHO and xrs-5

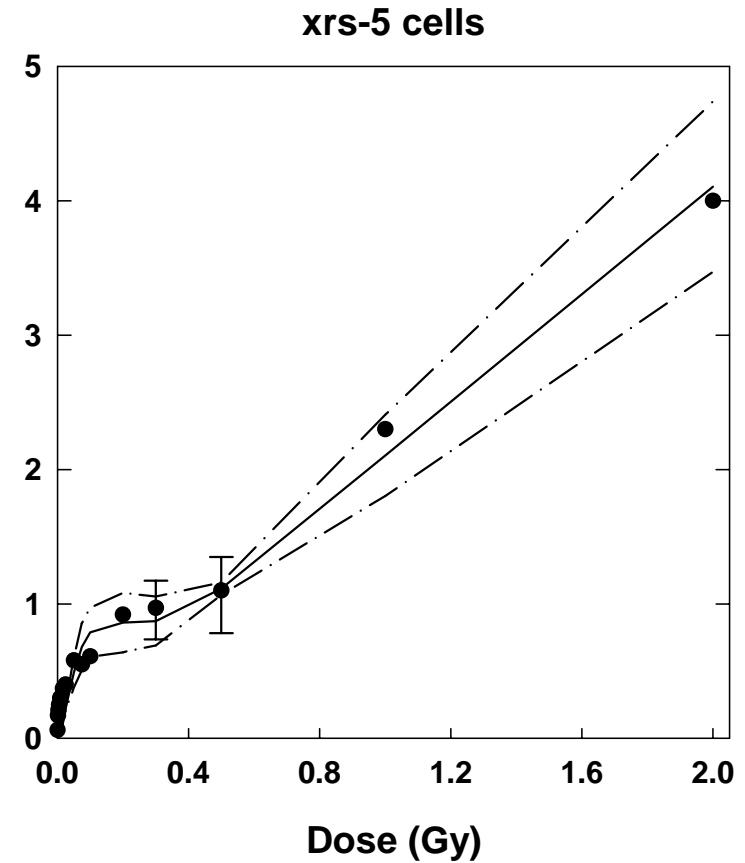
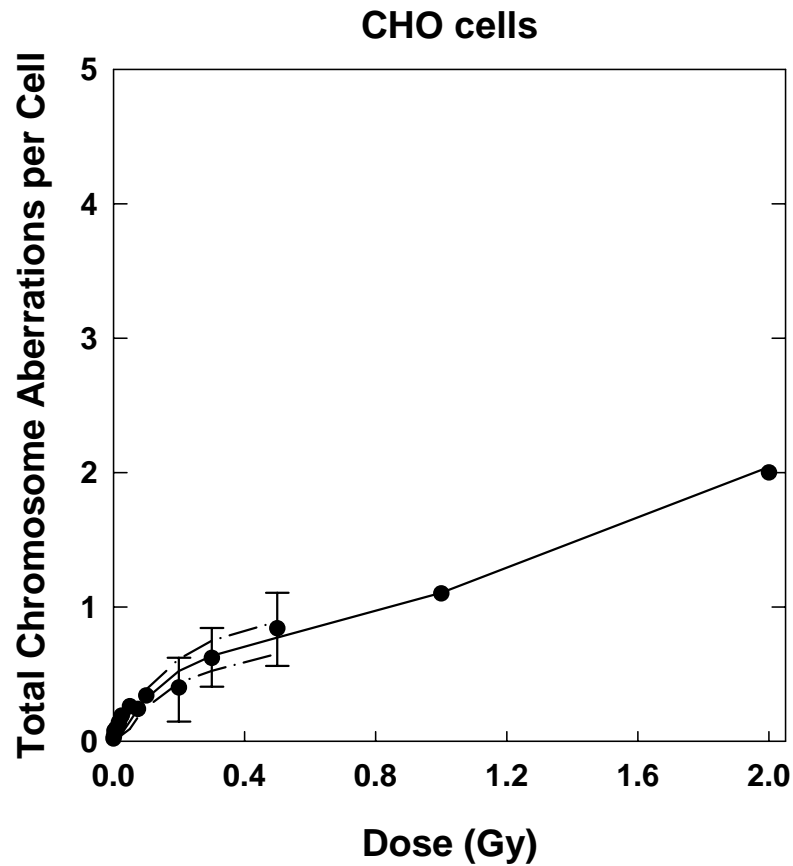


$$k_{01b_{by}} \times e^{-\lambda_{1by} \times D}$$

Data by Nagasawa and Little

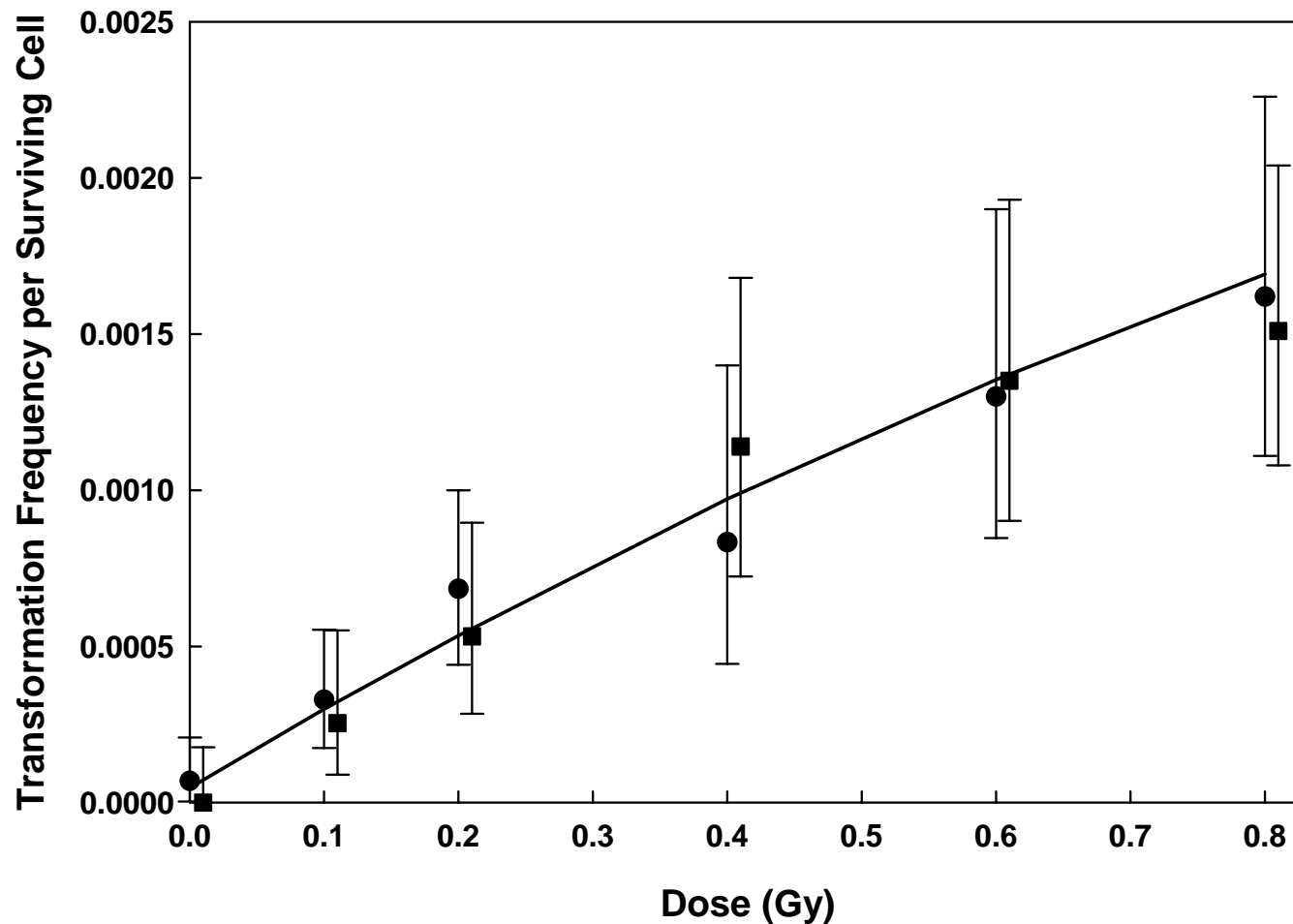


Data by Nagasawa and Little

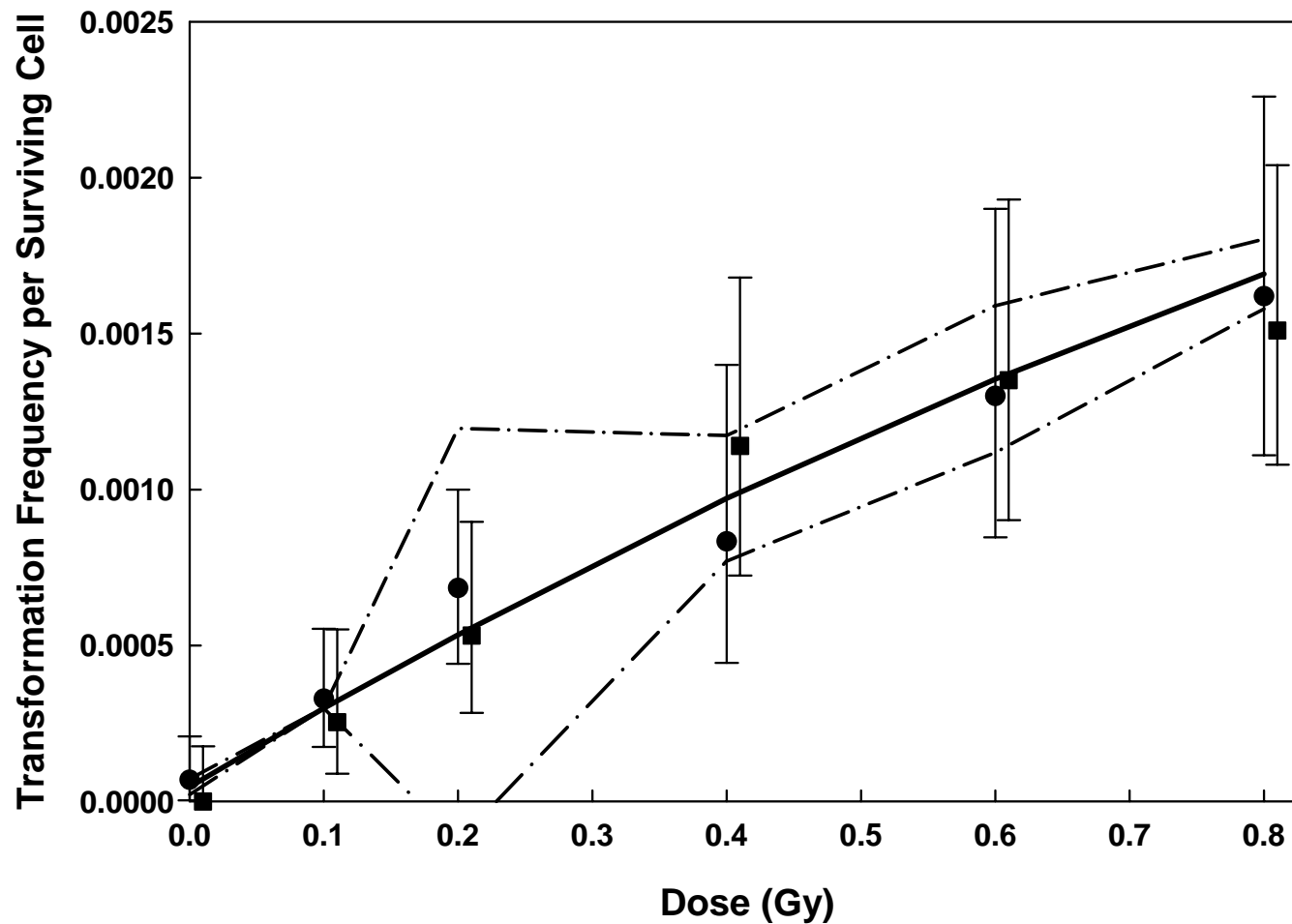


Data by R.C. Miller et al.

Radiat Res. 1995, α -particles (150 keV/ μ m), C3H 10T1/2

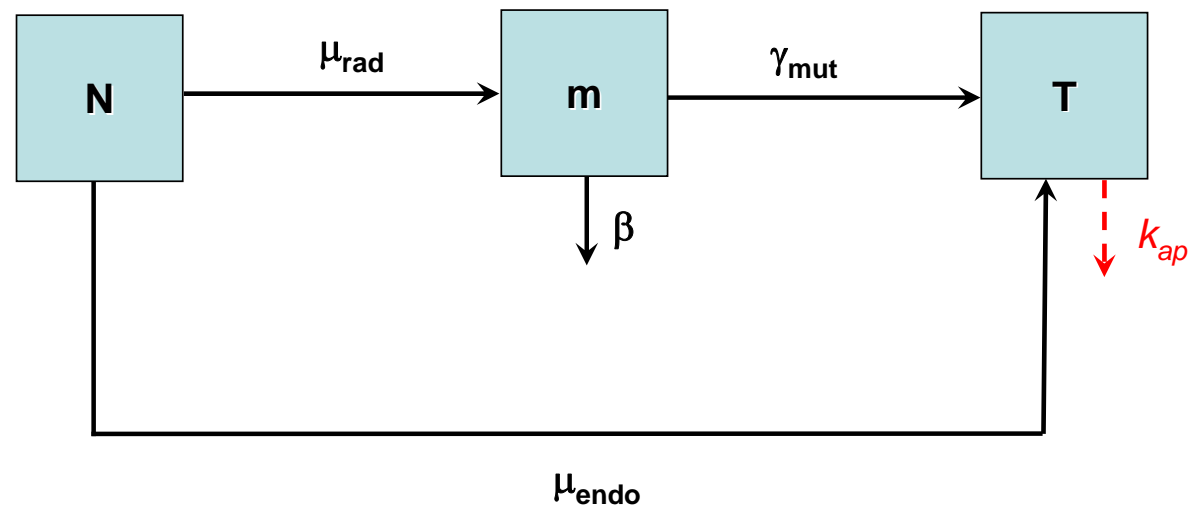


Data by R.C. Miller et al.



New model for neopl. transformation

Cellular level



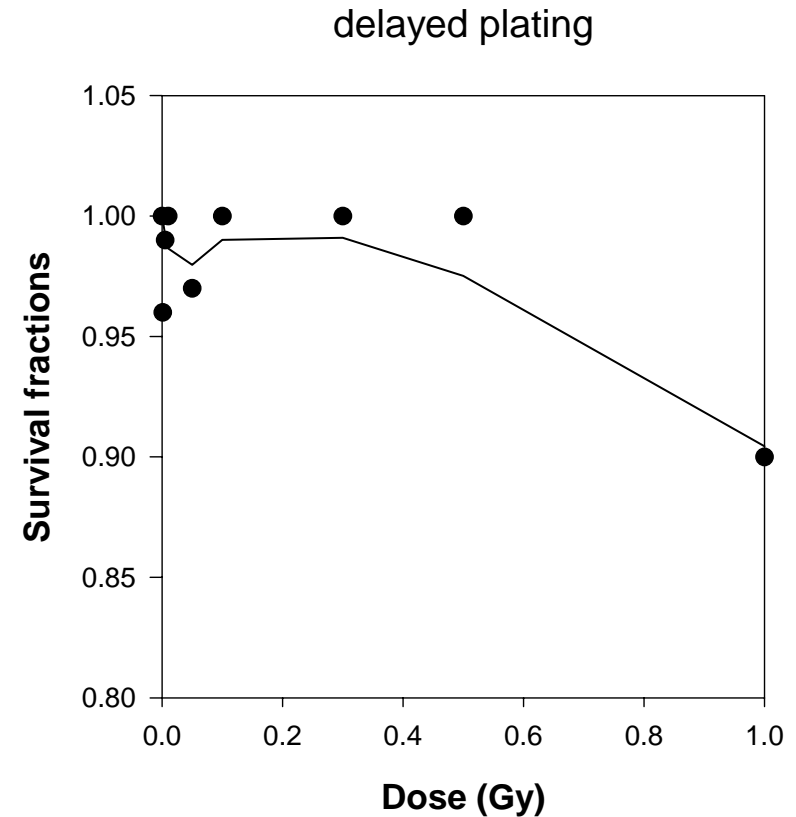
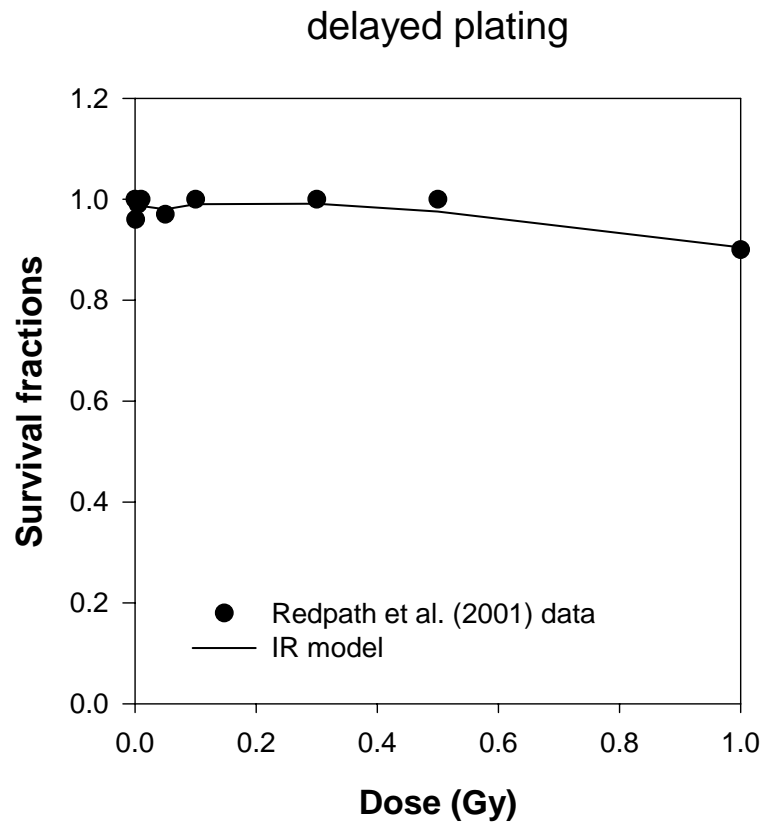
Molecular level

DNA damage production: $\Sigma_{sl}^{\text{endo}}, \Sigma_{cl}^{\text{endo}}, \Sigma_{sl}^{\text{rad}}, \Sigma_{cl}^{\text{rad}}$

DNA repair: $\lambda_{sl}, \lambda_{HR}, \lambda_{NHEJ}$

DNA misrepair: $\Phi_{sl}^{\text{endo}}, \Phi_{cl}^{\text{endo}}, \Phi_{HR}, \Phi_{NHEJ}$

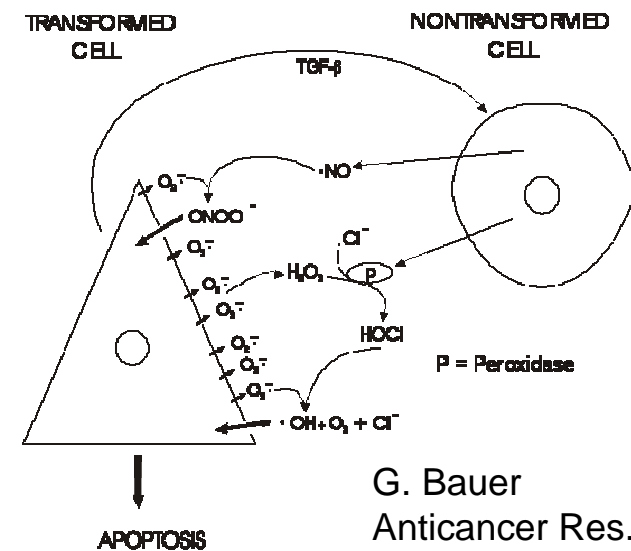
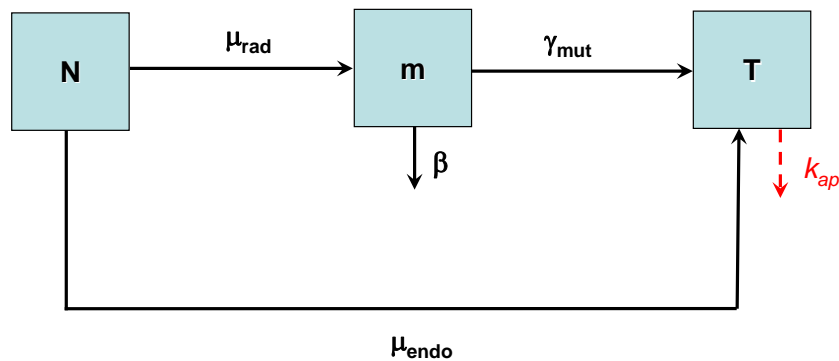
Low Dose HRS and IRR



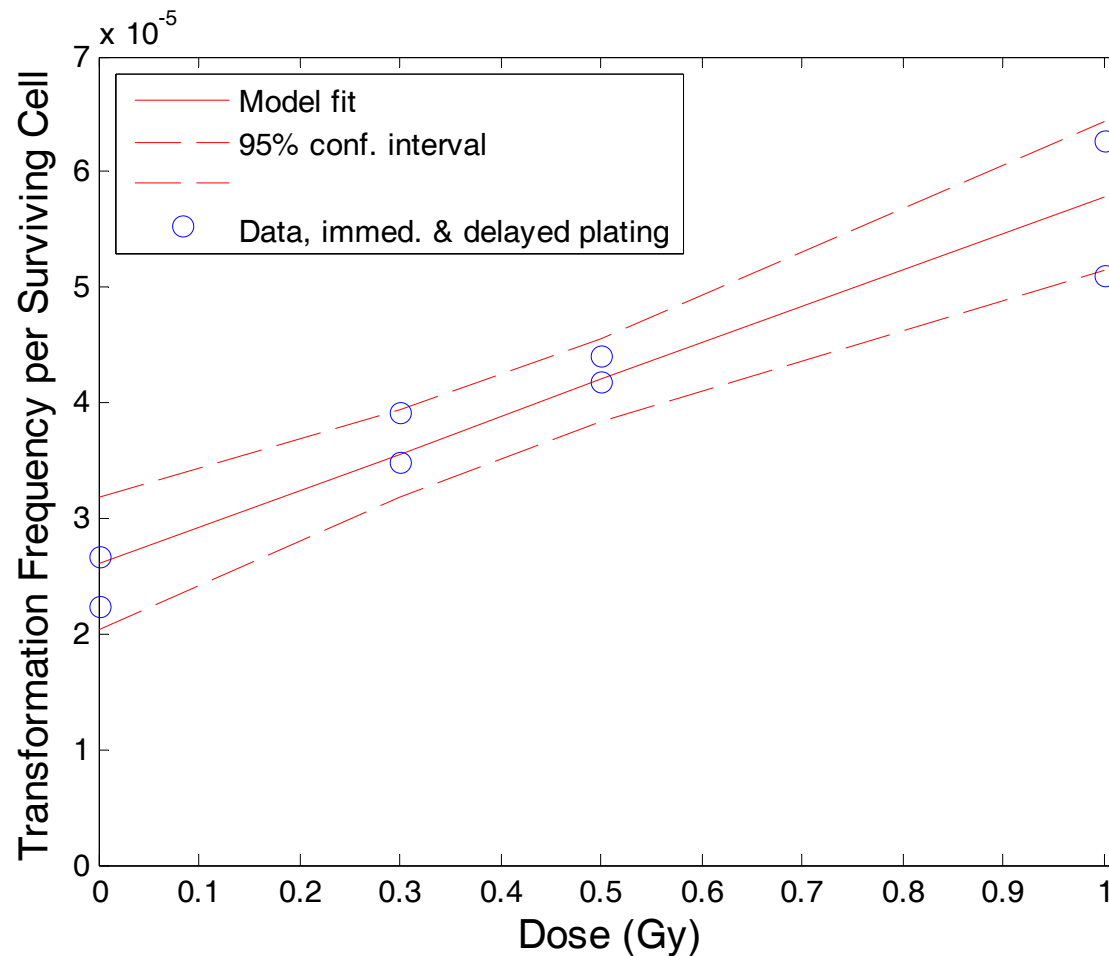
$$\text{IR model: } SF = \exp\{-\alpha_r[1 + ((\alpha_s/\alpha_r) - 1)\exp(-D/D_c)]D - \beta_1 D^2\}$$

Bystander-induced apoptosis

- **P**rotective **A**poptosis-**M**ediated process (PAM)
- PAM can eliminate transformed cells



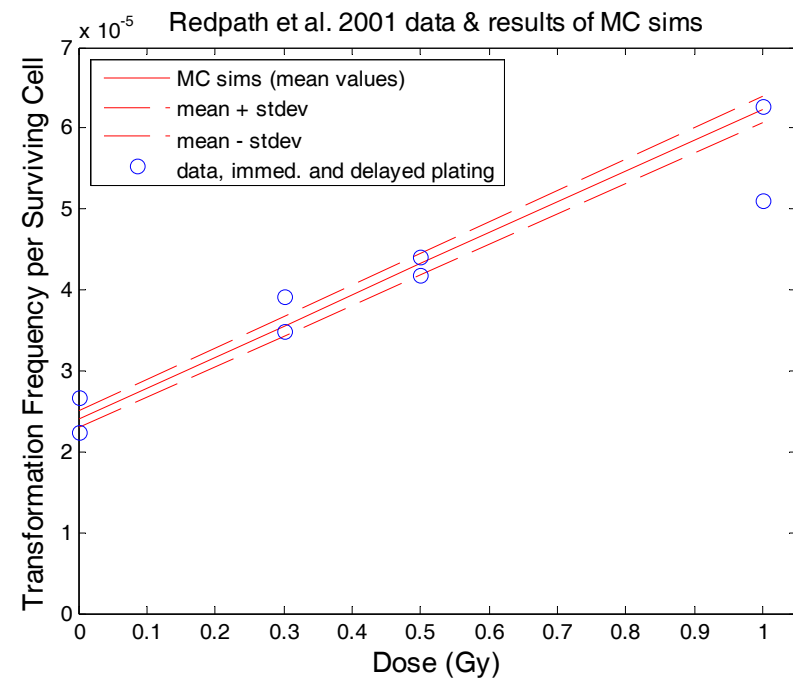
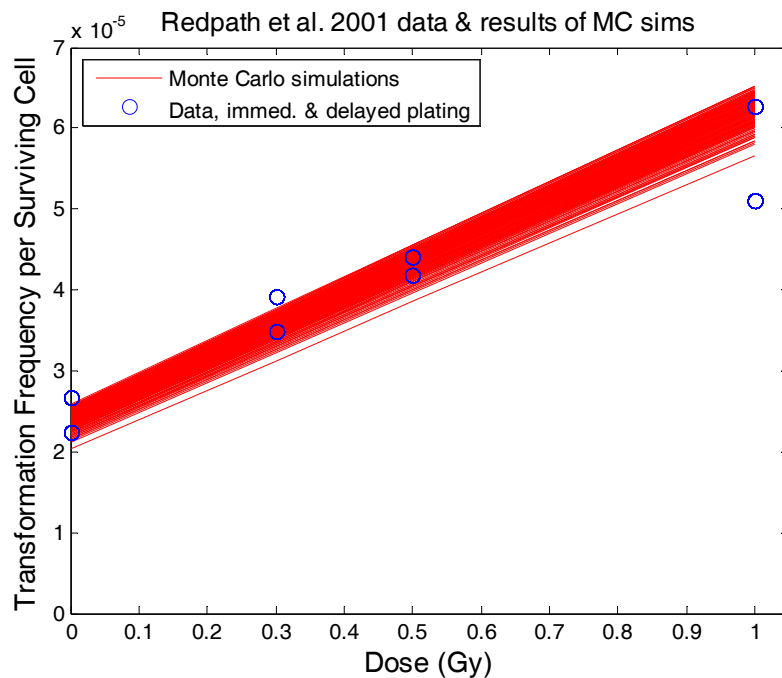
Model fits



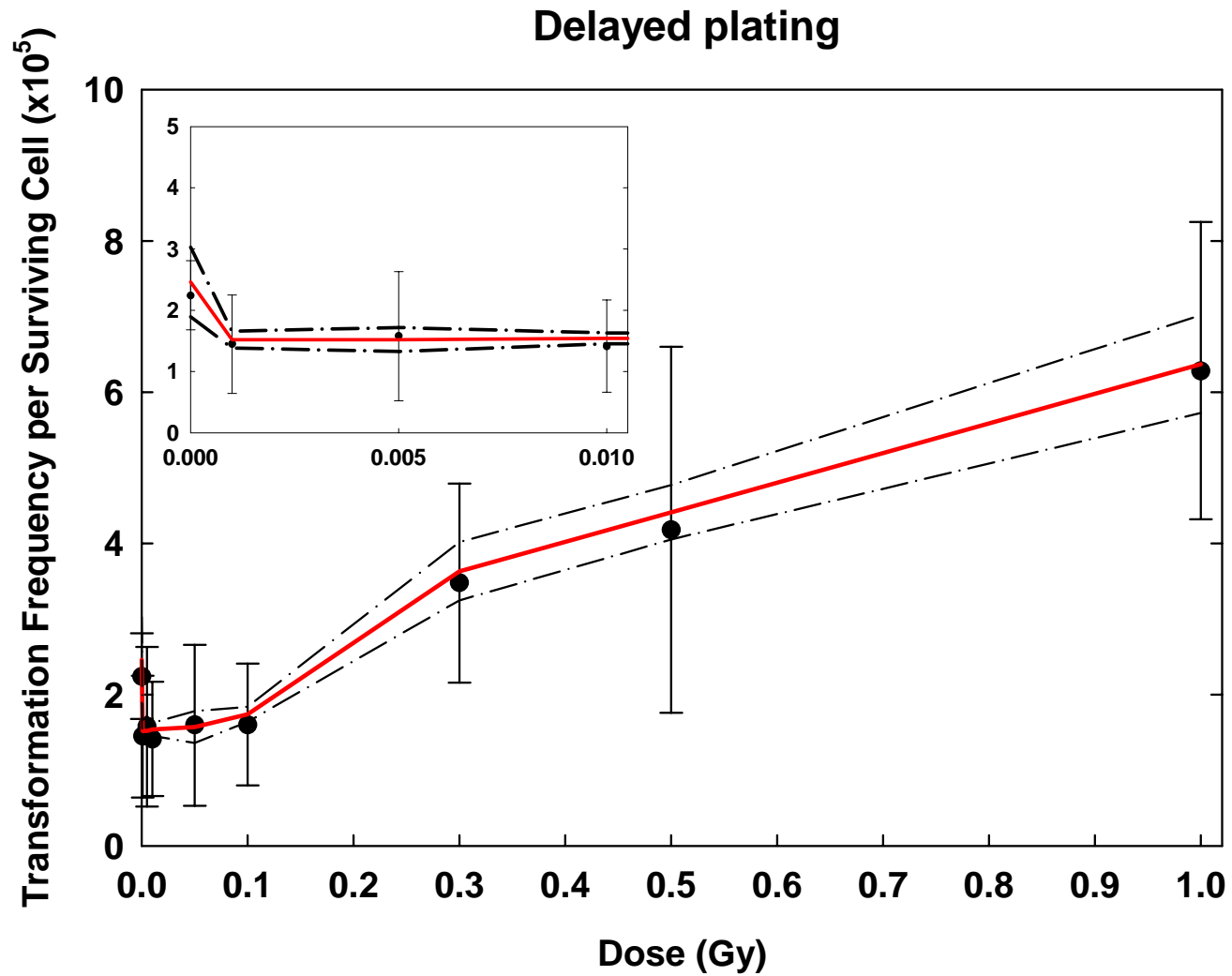
$$\Omega = 1.2 \times 10^{-5} (9.3 \times 10^{-6} - 1.4 \times 10^{-5})$$

$$\gamma_{\text{mut}} = 1.4 \times 10^{-3}/\text{day} (7.2 \times 10^{-4} - 2.2 \times 10^{-3})$$

Monte Carlo simulations



200 forward simulations with allowed uncertainties in λ_{HR} , λ_{NHEJ} , and λ_{SI} .



$$k_{ap} = 2.0/\text{day} \quad (-1.7 - 5.7)$$

Conclusions

- SVM successfully equipped with algorithms that allow to calculate 95% CI for model predictions and best estimated values
- New compartment model for *in vitro* neoplastic transformation
- Model has system-biological aspects and contains
 - endogenous and radiation-induced lesions
 - DSB repair via HR and NHEJ
 - misrepair

Conclusions

- low dose HRS/IRR
- bystander-induced apoptosis
- Model successfully equipped with algorithms that allow to calculate 95% CI for model predictions and best estimated values
- Model made stochastic via Monte Carlo methods

Outlook

- Work towards model that contains all mechanisms that are rate-limiting and essential at low doses:
 - Low dose HRS/IRR ✓
 - Genomic Instability
 - endogenous DNA lesions ✓
 - inducible repair
 - radical scavenging
 - protective bystander effects ✓
 - detrimental bystander effects
 - Stochasticity ✓
 - Uncertainty ranges ✓

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 FWF