



310/1828

310/9

Workshop on Biomedical Applications of High Energy Ion Beams

Co-sponsored by: ICGEB and University of Surrey

12-16 February 2007

Venue: Adriatico Guest House Giambiagi Lecture Hall ICTP, Trieste, Italy

Modelling Patient Outcomes to Radiotherapy

Norman KIRKBY University of Surrey, U.K.



11:00 – 12:30 Modelling Patient Outcomes to Radiotherapy

Dr. Norman Kirkby, University of Surrey, U.K.



The Abdus Salam International Centre for Theoretical Physics







Wednesday 14th Feb 2007



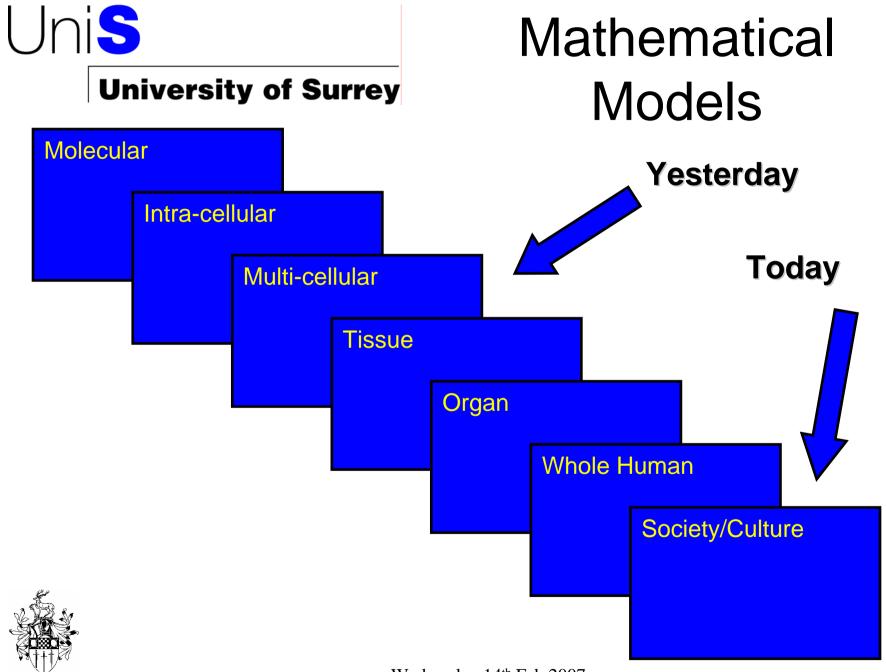
- Background
- BJJK model
 - Model of a patient & treatment
 - Model of a population of patients
 - Optimisation: fitting to clinical trials
- Results
- Conclusions and Future Work
- Questions





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Engineering and Physical Sciences Research Council

Background

- Discipline Hopping grant from the Life Sciences Interface of the Engineering and Physical Sciences, BBSRC and MRC
- One year working 'out of discipline'
- What can I take to the new discipline
- What can I return with from the experience?







"What committee in its right mind would take a Chemical Engineer and use taxpayers money to put him in a clinical neuro-oncology team for a year?"



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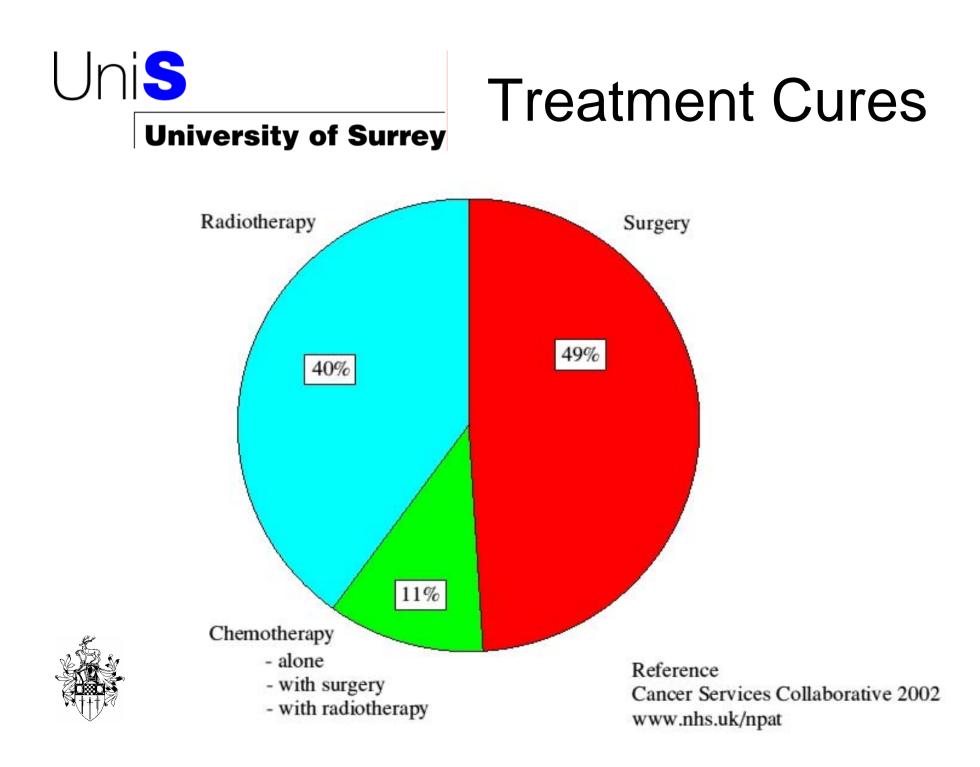


There are about 350,000 new cases of cancer in the UK per annum

- Lung, breast, colorectal and prostate account for the vast majority.
- There are about 4000 new cases per year of brain cancer of which 2000 are grade 4 glioblastoma.
- Average years of life lost per patient is higher for GBM than any other cancer









Typical GBM (Grade 4 glioma)

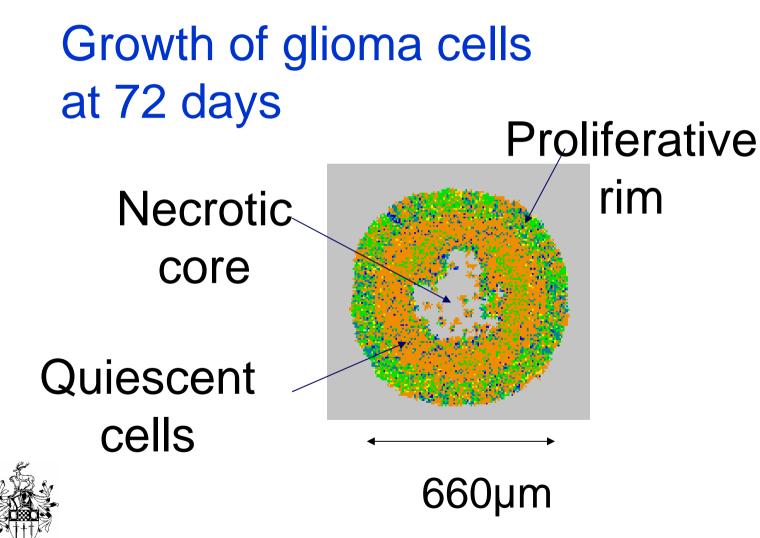




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Cellular Automata Models



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GBM Patients

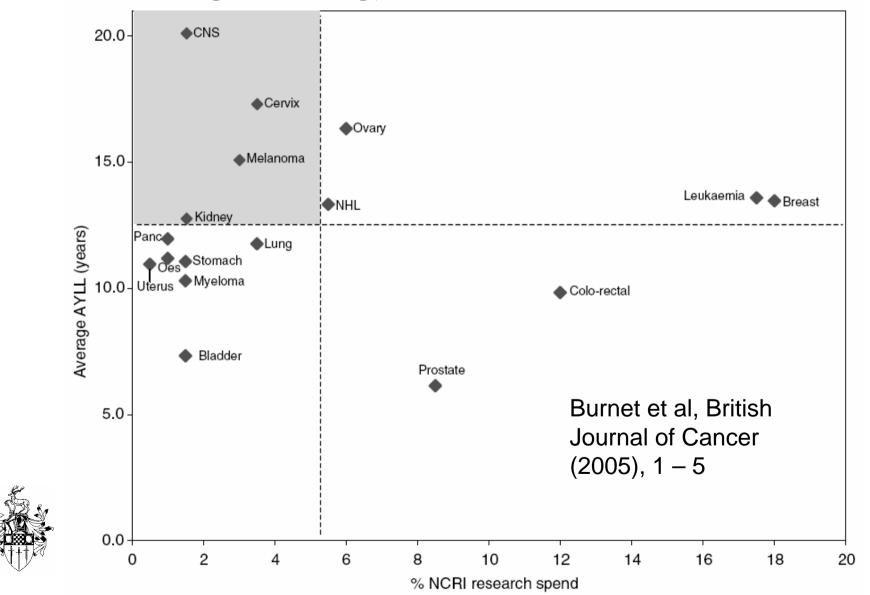
- The outcome has hardly changed over a long period of time
- Median survival < 300 days
- Long-term survival < 3%
- Average years of life lost per patient is highest for all main cancers
- Spending on research is low...





AYLL vs Spend

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GBM Patients

- Symptoms at presentation: headache, neurological impairment, fits.
- Metastatic spread is rare
- Recurrence at original site is common
- Proliferative rim includes invasion of normal brain





GBM Patients: Cause of Death

- Intra-cranial pressure
- Destruction of vital section of normal brain
- Toxic burden of necrosis?
- Other?





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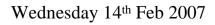
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Patient Outcome Models

- A model to address direct clinical issues
- What total dose to give?
- How to fractionate?
 - Given finite resources, waiting lists etc
- How to plan therapy?
- How to control side effects?
- Can we extract biological data from clinical studies?





Approach

- A model of 'a patient'
 - Tumour growth
 - Normal brain cell damage
 - Response to radiotherapy
 - Delay before treatment
- Monte Carlo simulation to generate a population of patients
- Simulated annealing to fit onto clinical data





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Model of a Patient

- Some reaction engineering...
- Normal brain cells, n_n
- Cancer cells, n_c

Tumour Growth

$$C \xrightarrow{k_c} 2C \quad r_c = k_c n_c$$

Destruction of Normal Brain

 $N + C \xrightarrow{k_n} C \quad r_n = k_n n_c n_n$



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First Approximation

- Exceedingly crude tumour growth kinetics
- Does damage to the normal brain continue if the tumour is not getting any bigger
- A better approximation might be to make damage related to growth rate

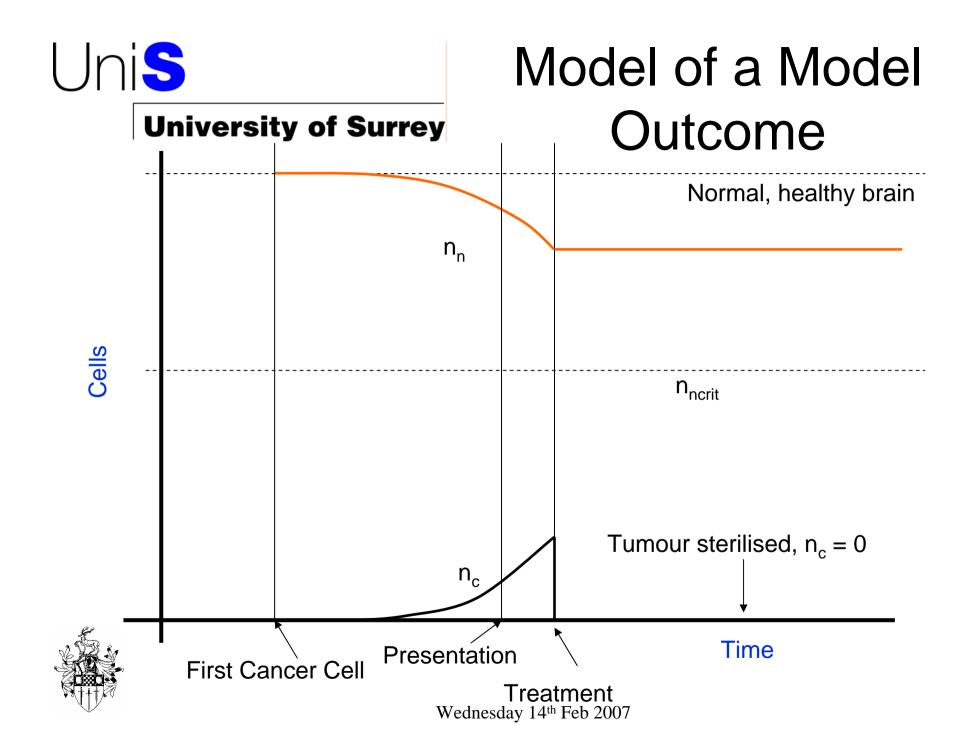


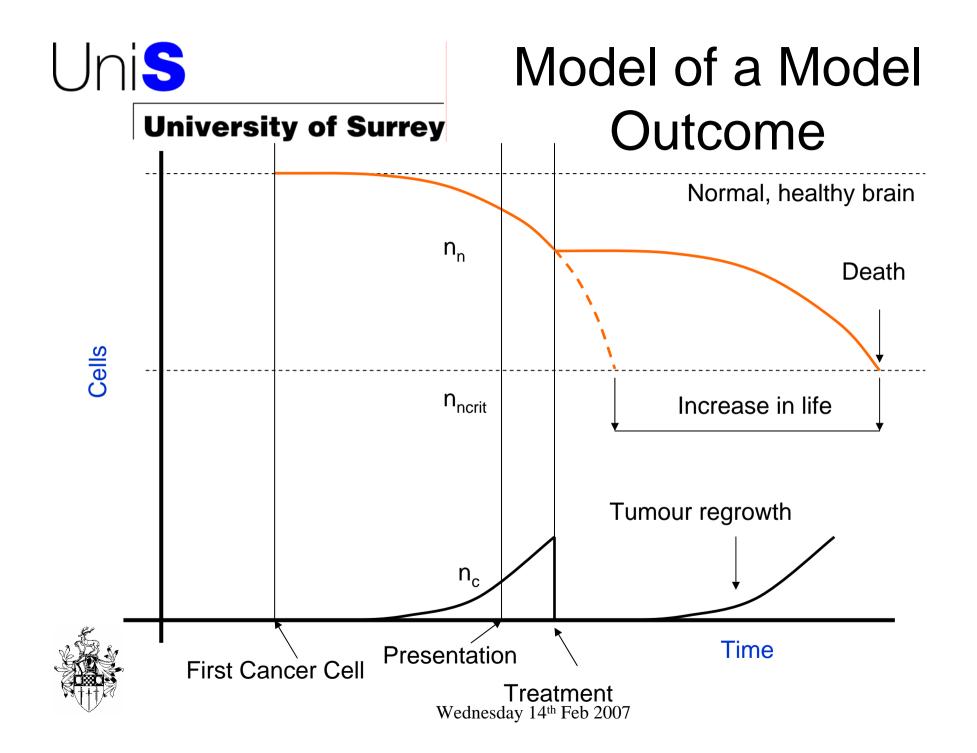


Model of a Patient

- Patient presents with some tumour, n_{c0}, and some brain left, n_{n0}
- When n_n drops below n_{ncrit} the patient dies
- We assume radiotherapy reduces n_c but has no effect on n_n.
- We assume no other loss of normal brain cells, e.g. with age.









- Assume the tumour is a closed system
- Perform a number balance on the system

$$\frac{dn_{c}}{dt} = r_{c} = k_{c}n_{c}$$
 Tumour

$$\frac{dn_n}{dt} = -r_n = -k_n n_n n_c \qquad \text{Normal brain}$$





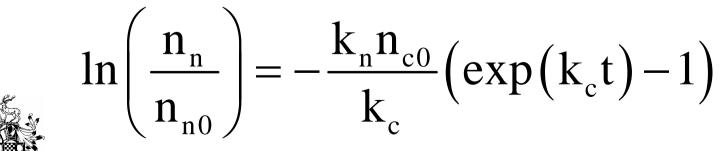
Model of a Patient

• Solve analytically

Tumour

$$n_{c}(t) = n_{c0} \exp(k_{c}t)$$

Normal brain





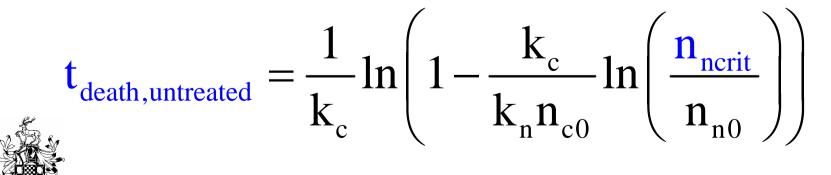
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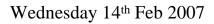


Model of a Patient

• From this we can calculate when a patient will die if untreated:

$$\ln\left(\frac{\mathbf{n}_{\text{ncrit}}}{\mathbf{n}_{n0}}\right) = -\frac{\mathbf{k}_{n}\mathbf{n}_{c0}}{\mathbf{k}_{c}}\left(\exp\left(\mathbf{k}_{c}\mathbf{t}_{\text{death,untreated}}\right) - 1\right)$$

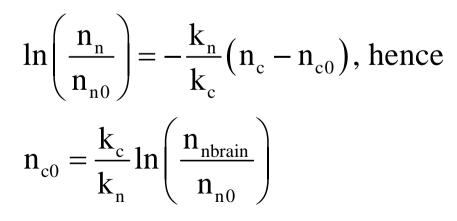






Size of Tumour at Presentation

- n_{c0} is not an independent parameter
- It can be calculated from n_{n0}, k_c, k_n and n_{nbrain} (the number of normal cells in a complete brain)
- The state space is given by





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Age of Tumour at Presentation

- Using the state space
- Given the size of tumour at presentation
- The age of the tumour is...

$$t_{age} = \frac{1}{k_c} ln(n_{c0})$$





Treating the Patient

- Select the patient for a therapy
 - Radical
 - Palliative
 - Best nursing care
- Wait for treatment machine to become available, t_{delay}
- Treat instantaneously







Treating the Patient

$$n_{c}(t_{delay}^{+}) = n_{c}(t_{delay}^{-})x_{s}^{j}$$

- If cell number in tumour < 1 deem tumour to be sterilised
- Otherwise calculate new survival time
- Give j fractions of radiotherapy, each has tumour cell survival fraction x_s
- Assume we know x_s for each patient





Survival After Treatment

- We can calculate how long the normal brain will now take to drop to only n_{crit} cells left.
- Hence we have calculated the patient survival time
- We can impose a clinical trial time and censor surviving patients if necessary





Summary of the Patient

- Tumour growth rate constant, k_c
- Normal brain interaction constant, k_n
- Normal brain at presentation, n_{n0}
- Normal brain at death, n_{crit}
- Delay to start treatment, t_{delay}
- Survival fraction in response to each fraction, x_s





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Populations of Patients

- Monte Carlo Simulation:
- Random generation of each of the parameters that characterise a patient
- Calculate their survival time, etc
- Analyse the statistics for the results
- Input distributions have to be plausible, if not measured.





Populations of Patients

Distributions for

- Normal brain cells at presentation
- Critical cells number left at death
- Tumour doubling time
- Delay to treatment
- Tumour/Normal cell interaction constant
- Survival fraction to a single dose of radiation





Distributions

• Normal distributions

$$p(k) \propto \exp\left(-\left(\frac{k-\overline{k}}{\sigma}\right)^2\right)$$

• Skewed distribution for survival fraction

$$p(x_s) \propto x_s^n (1-x_s)^m \exp(-\alpha x_s)$$



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Constructing a Population of Patients

- Know, estimate or determine by fitting 13 parameters:
- Mean and standard deviations for
 - $-k_c, k_n, n_{n0}, n_{ncrit}, t_{delay}$
- Survival fraction parameters

 $-n, m, \alpha$





Methods for Generating Distributions

- The Normal distribution is easy:
 - Method of Box GEP and Muller ME 'A note on the generation of random normal deviates, Ann. Math. Statist., 29, 610-611, 1958
 - call RANDOM_NUMBER(r)
 - call RANDOM_NUMBER(theta)
 - Theta = two_pi*theta
 - R = SQRT(-two * log(r))
 - x = R*SIN(Theta)*var%std+var%mean
 - y = R*COS(Theta)*var%std+var%mean





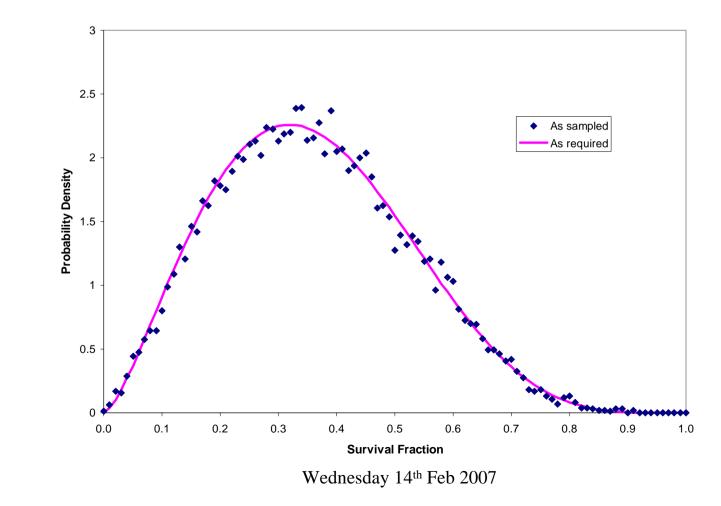
Methods for Generating Distributions

- The Survival Fraction Distribution is not so easy:
 - Use an acceptance/rejection test
 - It is iterative
- Care is required to check that the desired distribution is being generated





Methods for Generating Distributions





Disadvantages of x_s Distribution

- Specify n, m, and $\boldsymbol{\alpha}$
- Have to compute numerically:
 - Mean
 - Variance
 - Skewness
 - Kurtosis etc
- Integrate numerically for the normalisation constant



$$p(x_s) = \mathbf{k}_{x_s} x_s^{n} (1 - x_s)^{m} \exp(-\alpha x_s)$$

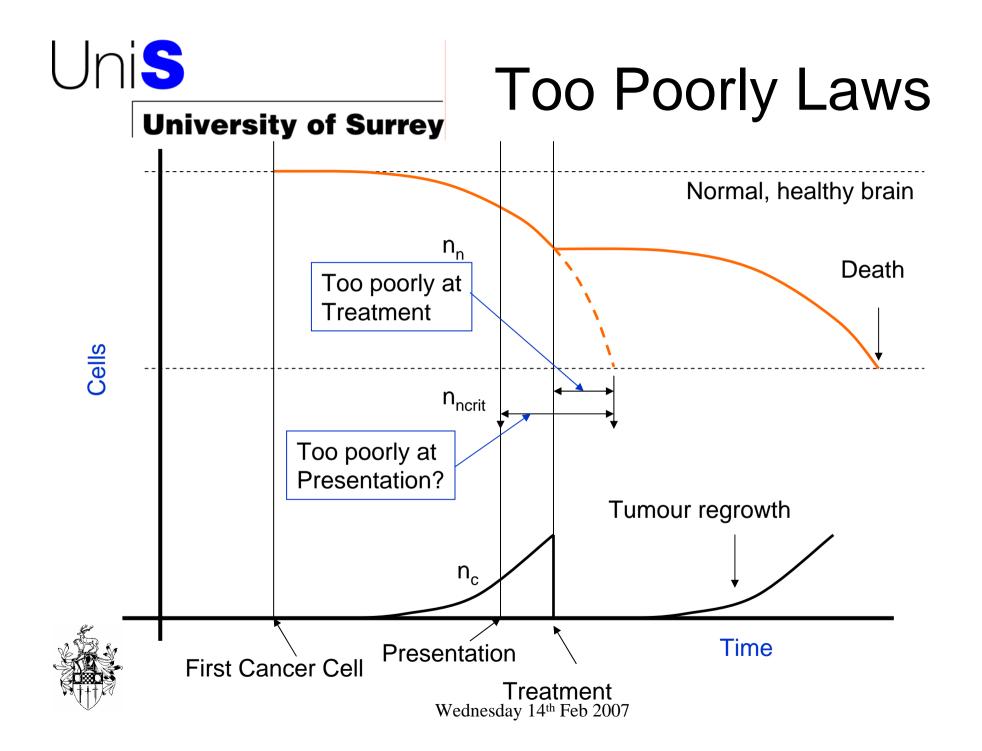
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Clinical Patient Selection

- Some patients, with parameters selected at random, are dead at presentation, die before treatment etc
- Clinicians use WHO performance status to select patients
- We invented the 'too poorly laws'







Modelling Clinicians

- Our model clinicians know when a patient will die if untreated
- Our model clinicians do not know the waiting time to commence treatment
- Both are wrong!
- Both flatter real clinicians!





Treatment Selection

- If $t_{untreated} < t_1$
 - Best nursing care, too poorly for any treatment
- If $t_2 > t_{untreated} > t_1$
 - Palliative treatment
 - Short planning and waiting time
- If t₂<t_{untreated}
 - Radical treatment
 - Longer planning and waiting time





Assessment at Presentation

- Given that we pretend we do not know the waiting time...
- Patients can be too poorly to treat when they arrive for treatment – and they are turned away.
- Conclude...
- A large number of patients may have to be generated in order to get enough to actually treat





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Fitting the Model to Data

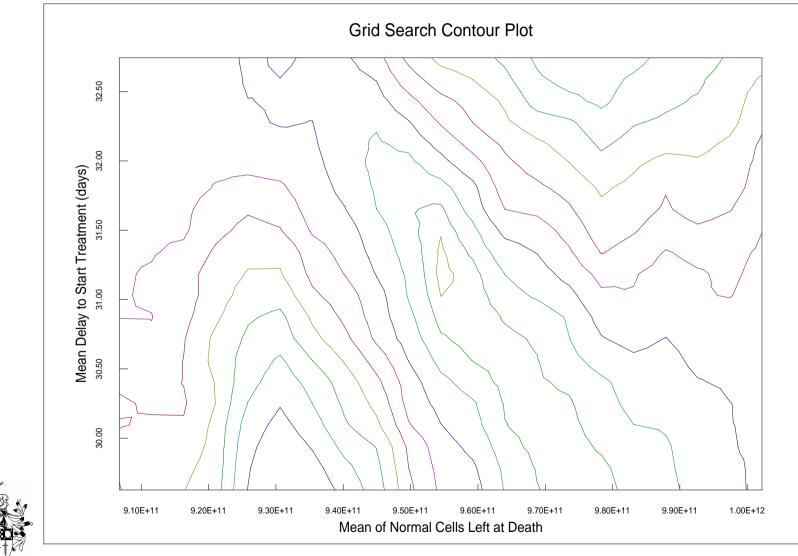
- Some parameters we know ahead of time e.g. distribution of waiting times
- Some have to be obtained by fitting
- Optimisation routines can be very slow and can home in on local minima
- We used a simulated annealing/ folding polygon hybrid.





Problems Fitting

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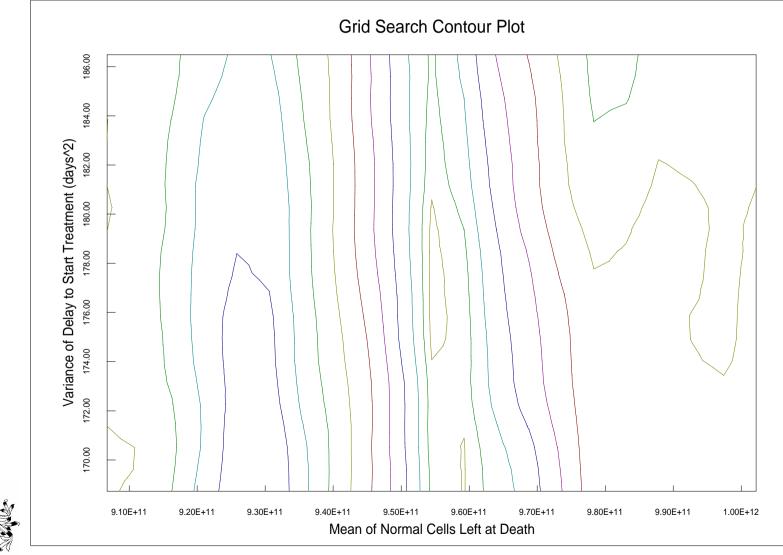


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Problems Fitting

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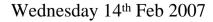


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Sensitivity Analysis

		Relative change in sum of squares of errors from best fit (%)						
	0%	2000%	4000%	6000%	8000%	10000%	12000%	14000%
Mean initial normal cell number - 10	0%	1		I				14003%
Mean initial normal cell number + 10	0% 🛓		2875%					
Mean tumour doubling time - 10	0% 🛓	1560%						
Mean tumour doubling time + 1	0% 🛓	1002%						
Survival fraction parameter n - 10	0% 🛓	687%						
Survival fraction parameter n + 1	0% 🛓	538%						
Mean normal cell number at death + 1	0% 🛓	326%						
Mean delay to start radiotherapy + 1	0% 🛓	289%						
Survival fraction parameter m + 1	0% 🛓	270%						
Survival fraction parameter m - 1	0% 🛓	149%						
Survival fraction parameter alpha - 10	0%]	47%						
Mean normal cell number at death - 1	0%]	45%						
Survival fraction parameter alpha + 1	0%]	39%						
Mean delay to start radiotherapy - 1	0%]:	36%						
Variance of initial normal cell number - 10	0%]:	30%						
Variance of initial normal cell number + 1	0%]	11%						
Variance of delay to start radiotherapy + 1	0%]:	7%						
Variance of normal cell number at death - 1	0%]:	3%						
Variance in delay to start radiotherapy - 1	0%	1%						
Variance of tumour doubling time + 1	0%] [·]	1%						
Mean value of interaction constant - 1	0%] [·]	1%						
Variance of tumour doubling time - 1	0%	0%						
Variance of normal cell number at death + 1	0%](0%						
Mean value of interaction constant + 1	0%](0%						
Variance of value of interaction constant + 1	0%]	0%						
Variance of value of interaction constant - 1	0%](0%						





Recall that this model currently has a log(log()) term, recall...

$$\mathbf{t}_{\text{death,untreated}} = \frac{1}{k_{c}} \ln \left(1 - \frac{k_{c}}{k_{n} n_{c0}} \ln \left(\frac{n_{\text{ncrit}}}{n_{n0}} \right) \right)$$

- So it is not surprising that the sensitivities are widely spread.
- Variances are covariant





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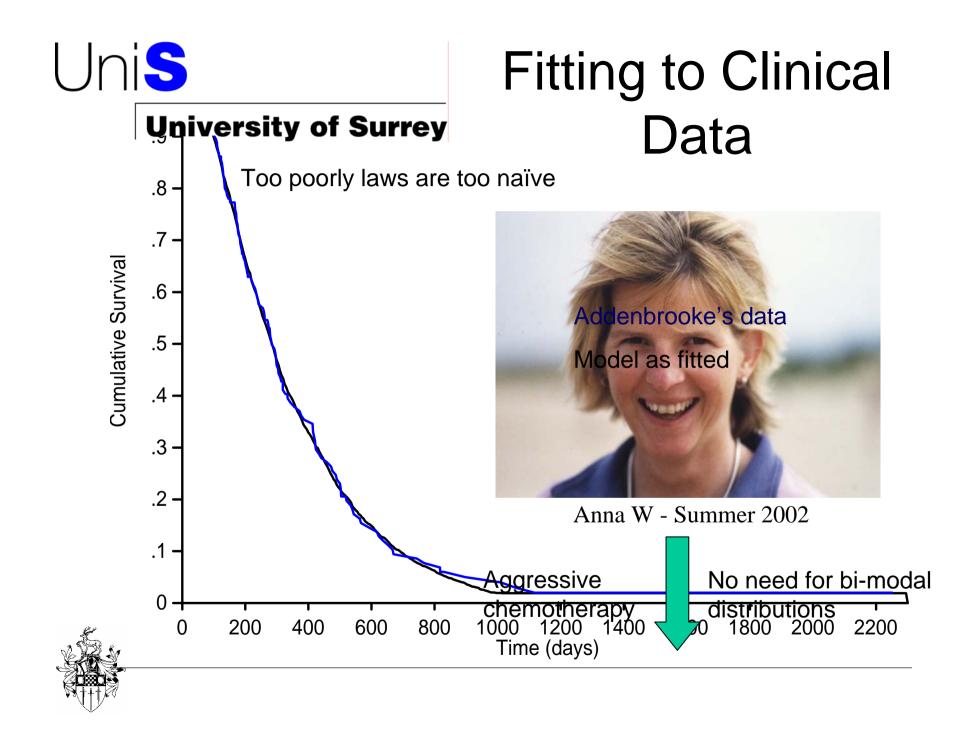




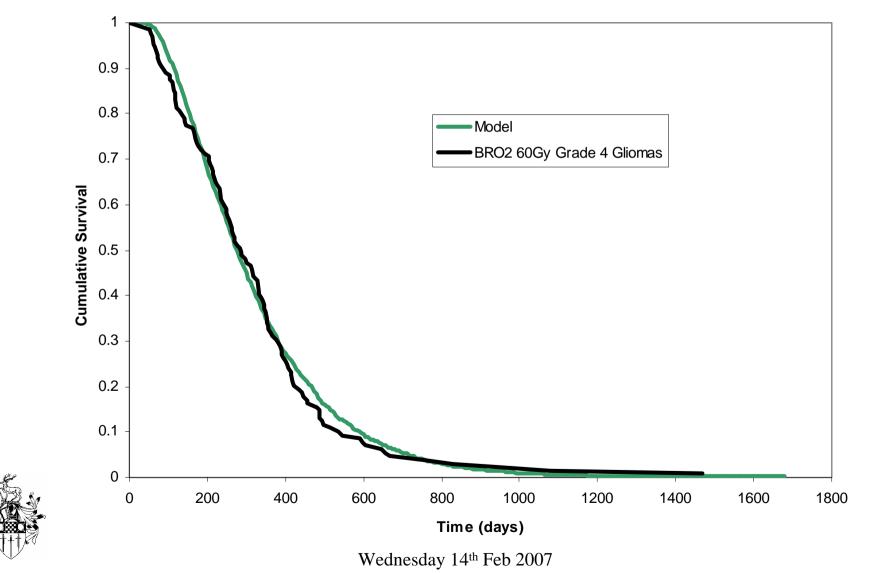
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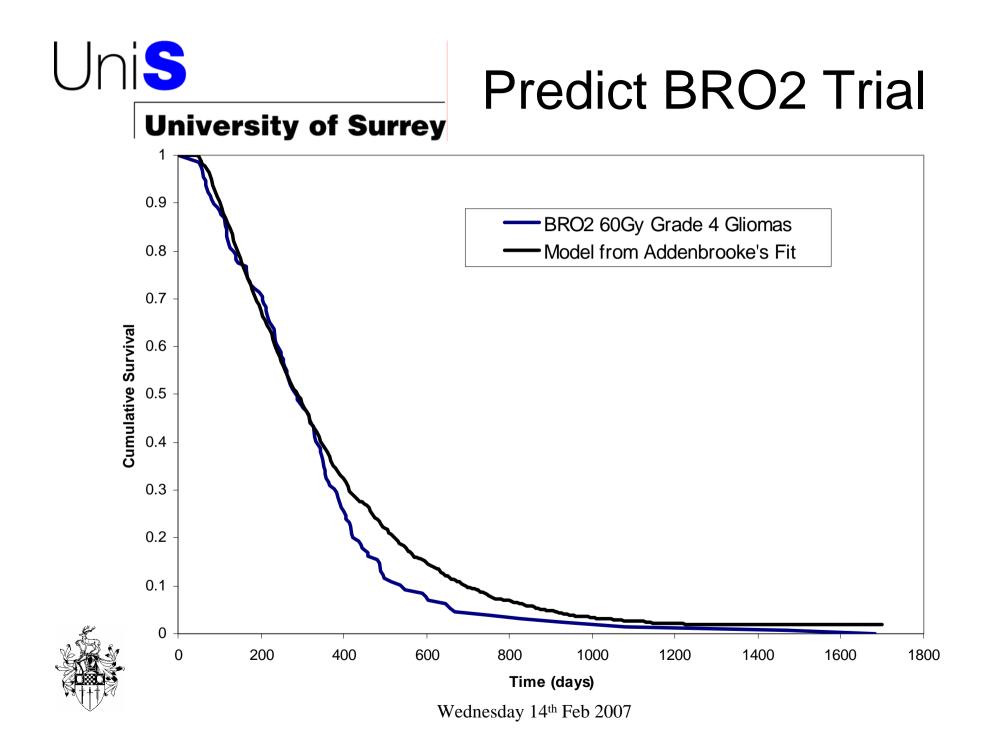
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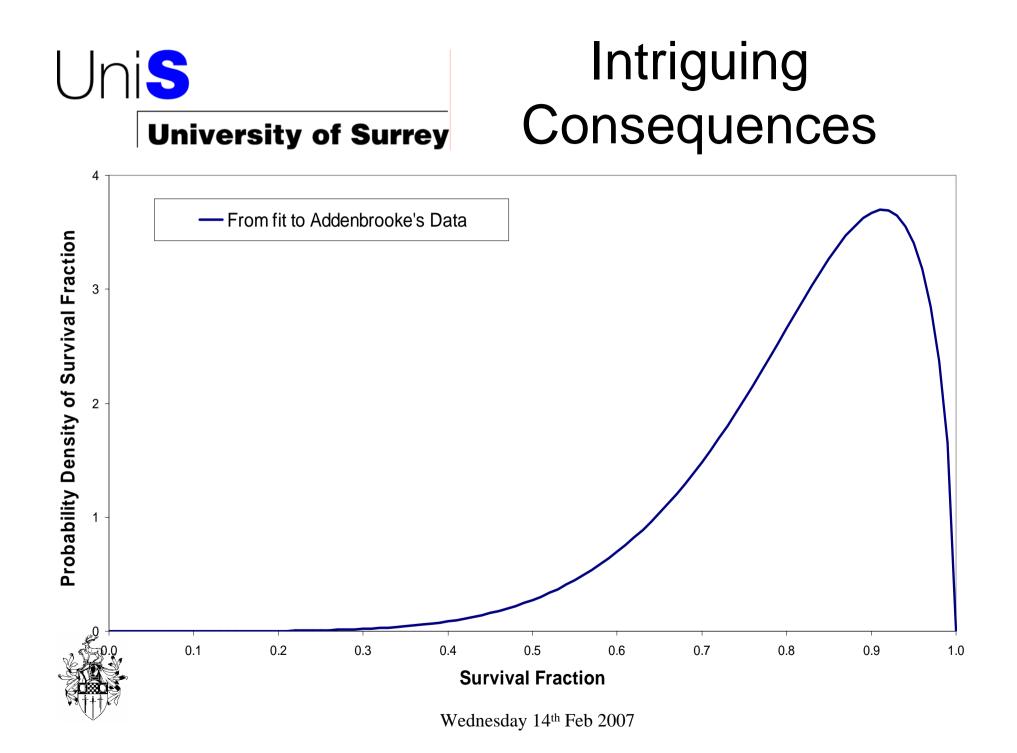








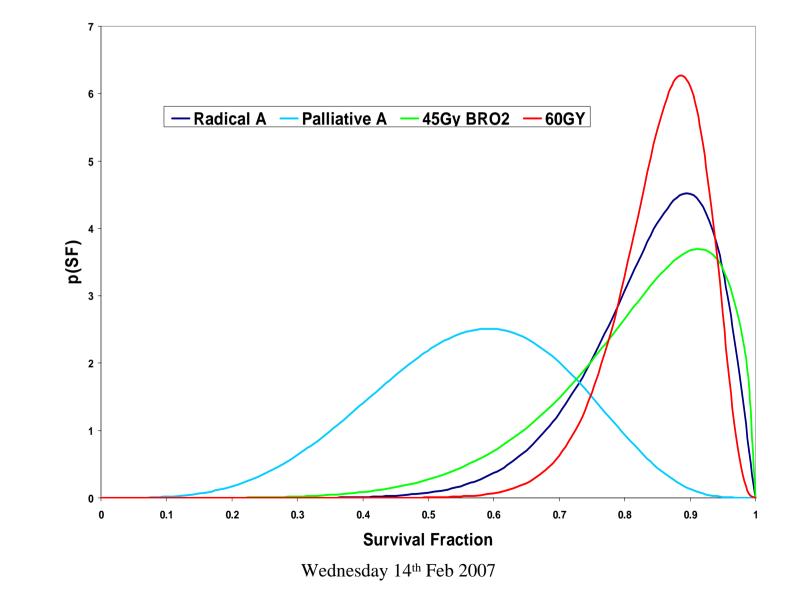






Radical & Palliative

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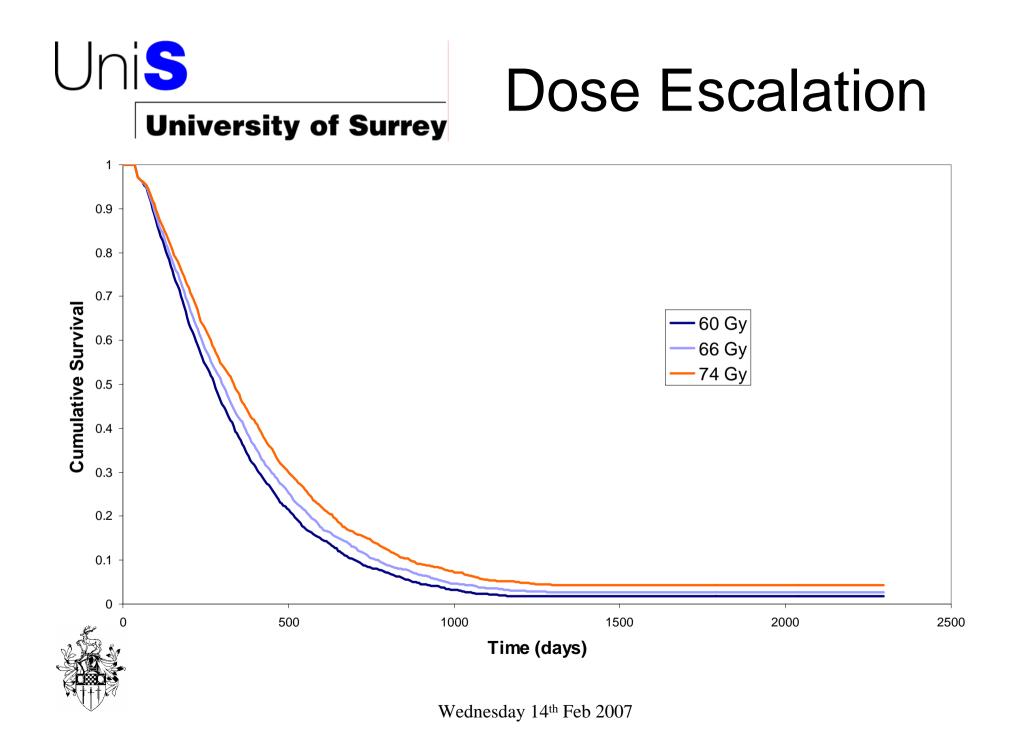


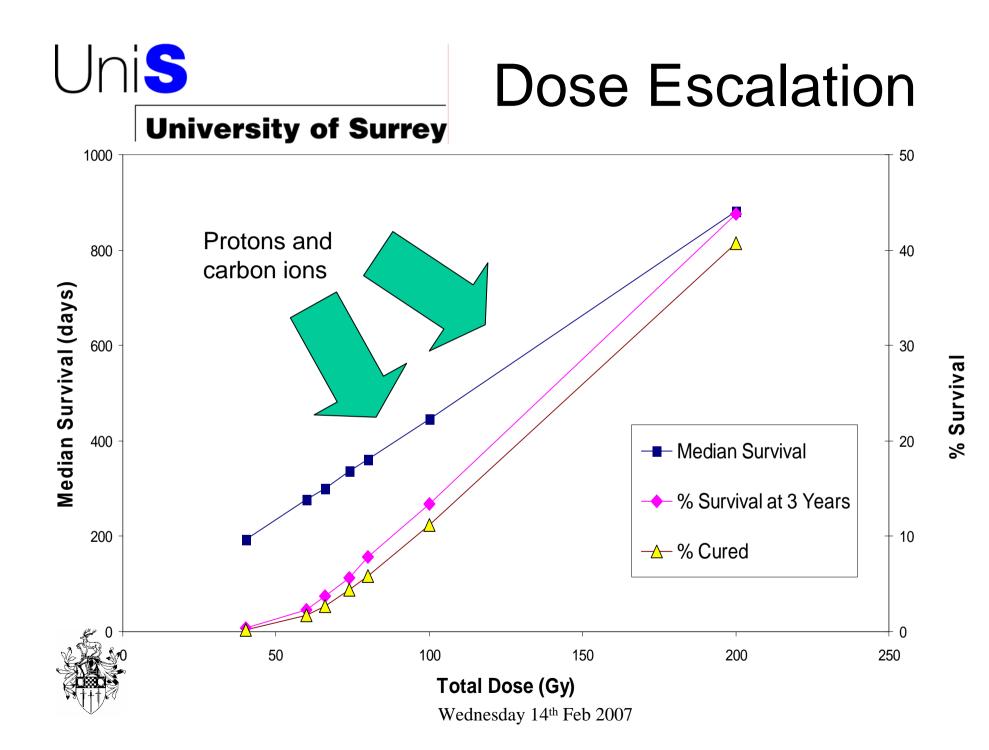


Distribution of SF

- What we predict here is consistent with measured SF at these doses
- The mean α/β is about right
- Can we drive these distributions to the left with the high RBE of carbon ions?
- Can radiosensitising drugs help?







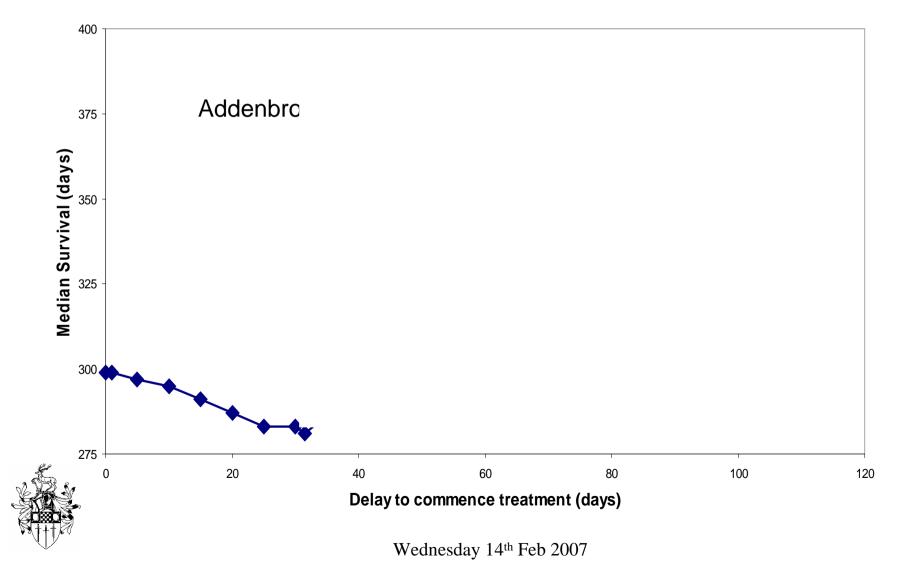


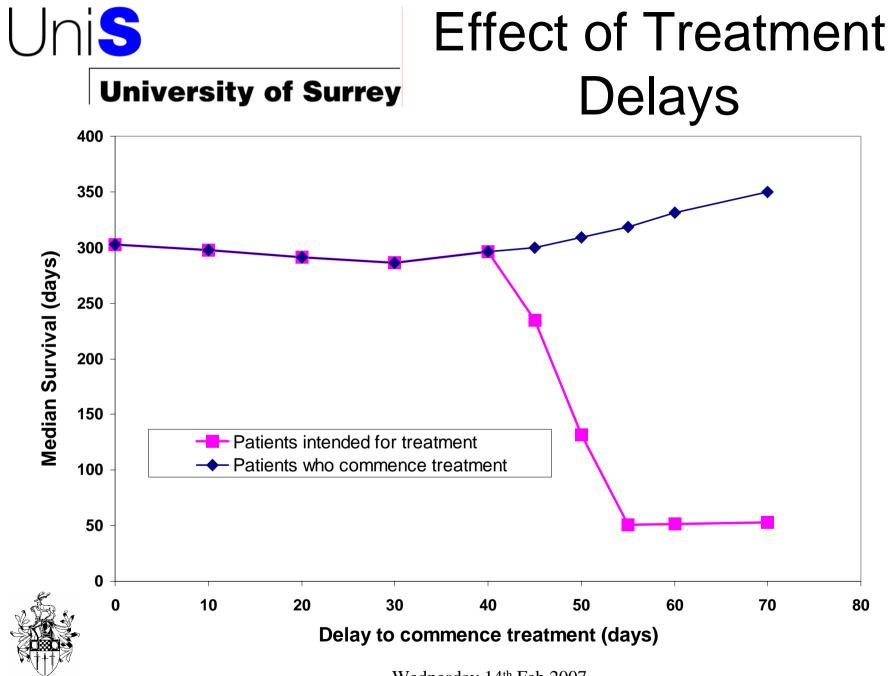
Design of Clinical Trials?

- How many patients do we need to recruit in order that the effects of a given dose escalation can be seen at a conclusive level of statistical significance.
- Can this model help design trials?









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Waiting Times

- Political Issue in UK
- Effects seem to be highly dependent on too poorly laws
- Do and Barton study, but very few others – evidence difficult to interpret
- What effect would a proton facility in the UK have on waiting times?



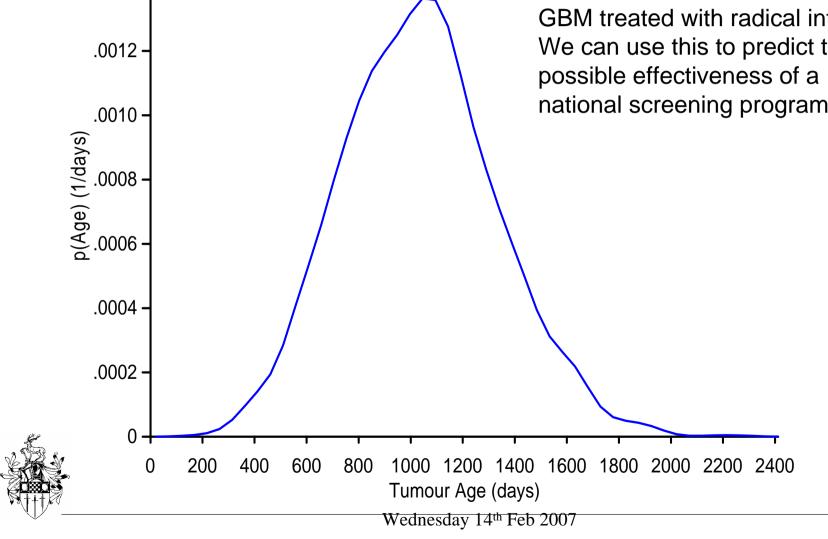


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Tumour Age Distribution

GBM treated with radical intent. We can use this to predict the national screening programme





Tumour Age Distribution

- Very difficult to measure
- Real GBM distribution is probably younger (k_c decreases with size and nutrient limitation)
- Early diagnosis does not seem to help in GBM





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Conclusions

- We have made a start
- Formulating the model has caused us to ask some interesting and difficult questions.
- The model has been used to guide the design of a clinical trial
 - Which is now in progress





- Multi-processor Fortran and Migration to Super-computer
- Optimisation Methodologies to exploit Cluster or Grid capabilities
- Radiobiology, e.g. SF related to dose
- Effect on normal tissue
- Low dose and bystander effect
- Time distributed treatment
- Conventional treatments
- Missing appointments





- Concurrent/Adjuvant chemotherapy
- temozolomide
- "Chemo as a second, isolated treatment"
- Time dependent interactions between the treatments
- Chemo normal brain interactions
- Question: difference between concurrent vs adjuvant chemotherapy





- Grade 3 Gliomas
- Analysis of data on grade 3 from BRO2 and other trials
- Analysis of the effects of chemo
- Clinical and Anatomical Aspects
- Age and WHO performance status
- Normal brain interactions
- Tumour location in brain
- Other tumour cites, e.g. cervix





- Distributions in clinician decision making
- Question: how does this effect Do and Barton waiting times?
- Improved growth kinetics
- Simple diffusion and reaction model
- Cell cycle-based growth kinetics
- Cell cycle-based radiosensitivity
- Novel and enhanced treatment strategies
- New radiotherapy schedules
- Protons and carbon ions
- New drugs (e.g. AQ4N?)





Collaborators

- Neil Burnet
- Raj Jena
- Sarah Jefferies
- Karen Kirkby
- M Eng Projects
 - Ollie Jones









Acknowledgements

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- The Master and Fellows of Fitzwilliam College
- The Addenbrooke's Hospital,
- The Department of Oncology, University of Cambridge,
- Surrey Ion Beam Centre
- School of Engineering, Surrey







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Take-home Message

- Currently, the main value in these models is in the creative process of formulation,
- Perhaps not in the results yet!
- Modelling is a potent catalyst for constructive dialog





Thank you for listening

Questions

"All models are a compromise with reality" Octave Levenspiel



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