Designing Evolutionary Treatment for Metastatic Non-Small Cell Lung Cancer

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Outline

- Cancer Treatment as a Stackelberg Evolutionary Game (SEG)
- SEG Application to Treatment of Metastatic Non-Small Cell Lung Cancer with Immune Checkpoint Inhibition
 - Data
 - Questions
 - Approach to Address One of These Questions
 - Interesting Observations
 - Results
- Conclusions & Future Research
- Discussion Points



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strategies	therapy options	effective strategies of therapy resistance
objectives	patients' quality of life	fitness



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Salvioli, PhD thesis 2020; Salvioli et al, PLOS One, in press; Cunningham et al, JTB 2018, PLOS One 2020, Stankova et al, JAMA Onc 2019,...





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What is the minimal knowledge necessary?

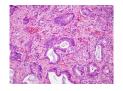




Data on NSCLC treated with Immune Checkpoint Inhibitian:



- Stage 4, anti-PD1 drug Atezolizumab (MPDL3280A)
- Tumor diameter over time, typically few time points
- In our dataset typically 2 or 3 metastases per patient

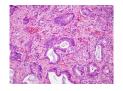




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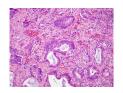




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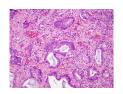
- Based on the initial tumor volume proxy and its trend, can we predict how its volume will change in the future?
- Would adaptive immunotherapy work in cases where tumor keeps growing when treated in the standard way?



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Our modelling approach to answer question 2:

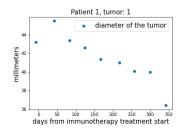


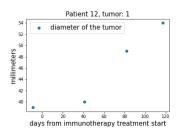


- Fitting the data into minimalistic G-function model
- Optimizing the treatment for cases with tumor growing



Example of data:





- All patients treated with the same immune checkpoint inhibition
- Treatment starts between the first and second data points

Are these different trends a result of treatment-induced resistance and other patient- and tumor-specific factors?

NSCLC model details (Vincent and Brown (2005)):

$$\dot{x} = x G(m, u, x)$$

$$\dot{u} = \sigma \frac{\partial G(m, u, x)}{\partial u}$$

$$G(m, u, x) = r(u) \left(1 - \frac{x}{K}\right) - \frac{m}{k + b u} - d$$

symbol & range	meaning
<i>m</i> ∈ {0, 1}	treatment (on or off)
$u \in [0, 1]$	rate of treatment-
	induced resistance
<i>K</i> > 0	carrying capacity
$x \in [0, K]$	cancer cell population
G(m, u, x)	fitness-generating function
$r_{\text{max}} > 0$	maximal growth rate
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Three different forms of r(u) considered:

- Quadratic cost of resistance: $r(u) = r_{\text{max}}(1 u^2)$
- Linear cost of resistance: $r(u) = r_{max}(1 u)$
- No cost of resistance : $r(u) = r_{max}$



Fitting the model (also for predictions):

- The population of cancer cells is estimated from the diameter.
- Distinguishing 6 groups of tumors: 3 according to initial volume and 2 according to initial trend.
- Fix K, b and σ per group, estimate r_{max} , k and u(0) per patient, d fixed to 0.01.

	Small	Medium	Large
	K = 0.1	K = 2	K = 2
Increasing	<i>b</i> = 100	b = 150	b = 0.9
	σ = 0.01	σ = 0.01	σ = 0.05
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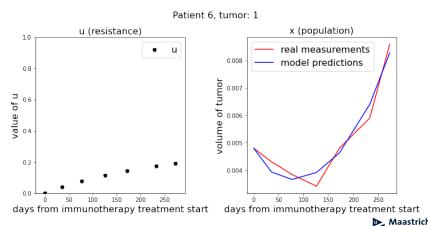
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Jakob Nikolas Kather: "There is no cost of resistance or carrying capacity in cancers I am dealing with"

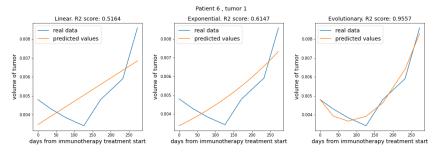


Fitting the model:



Comparison with linear and exponential models:

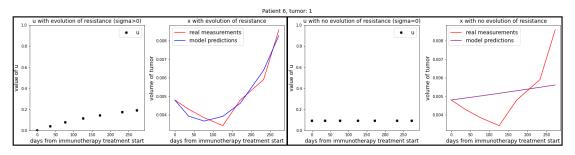
 Mean of the R2-scores: 0.59, 0.72, 0.80 for linear, exponential and evolutionary models, respectively.



Linear, exponential and evolutionary model with quadratic cost of resistance



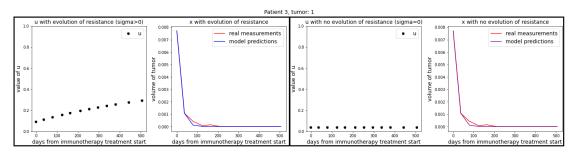
Evolution of treatment-induced resistance only plays an important role in tumors that exhibit a rapid change of trend.



Evolutionary model with quadratic cost of resistance with and without evolution of resistance



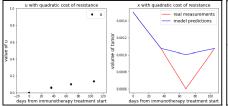
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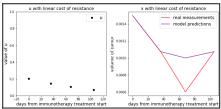


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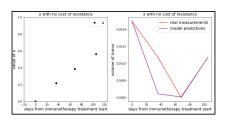


Comparison of the different forms of r(u):





Quadratic and linear cost of resistance



No cost of resistance



2. SEG Application to Treatment of Metastatic NSCLC Optimization

$$\begin{split} m^*(\cdot) &= \arg \min_{m(t) \in \{0,1\}} x(T) \\ \dot{x}(t) &= x(t) \, G(m(t), u(t), x(t)) \\ \dot{u}(t) &= \sigma \, \frac{\partial G(m(t), u(t), x(t))}{\partial u(t)} \\ G(m(t), u(t), x(t)) &= r(u(t)) \left(1 - \frac{x(t)}{K}\right) - \frac{m(t)}{k + b \, u(t)} - d, \quad t \in [0, T] \end{split}$$



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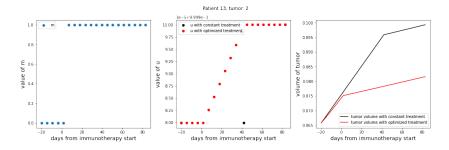
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Preliminary results:

- With linear and/or quadratic cost of resistance, the best is to always treat or not treat at all
- With no cost of resistance, sometimes it is the best to switch between no and standard treatment. We do not understand yet when precisely.
- With the objective min x(T), we can never stop the growth, we can only slow it down



Optimization - preliminary results:







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- For small tumors the error in estimating tumor volume may cause issues with the best fit; it may be that our conclusion would change with better tumor proxy
- Our model has better predictive capabilities than linear/exponential models used until now



4. Discussion Points/Questions for You

- Does resistance to immune checkpoint inhibition in NSCLC carry a cost?
- If yes, is there any way how to estimate it from other measurements (For other projects/cancers, we are exploring whether genomics and histopathology can help us to answer such questions, but here we know too little)?
- What other information can be useful here?
- Would a model with immune cells as active players be a better choice here?



Thank you!





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