Improving Strategic Decision-Making with Early Prediction of Survival Outcomes in Oncology Clinical Trials

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ACOP 11 - 10 November 2020

Collaborators

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With thanks to the patients and investigators who participated in these clinical trials.





<u>All stakeholders:</u> get safe & effective treatments to patients quickly

Drug developers: stage investment of unpromising candidates

Use of modeling & simulation has grown substantially over the past 15 years

- Broad recognition of its utility (e.g., PDUFA VI)
- Impacting a variety of decisions (patient treatment, <u>drug</u> <u>development</u>, regulatory)





How do we leverage existing data to make decisions about new therapies?







Flow of proposed approach







Joint model for tumor size and overall survival

• Data from 4 clinical trials of Pembrolizumab in NSCLC (N > 2500)

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- Chemotherapy (N=720; 28%)
- Pembrolizumab (N=1324; 52%)
- Pembrolizumab + Chemotherapy (N=497; 20%)

Moore model¹ for asymptotic tumor growth: $TS_i(t) = TS_{0,i}(5TS_{max,i})(1 - e^{-k_{g,i}t}) + TS_{i,0}e^{-k_{d,i}t}$

Hazard function with cure fraction:

 $h_i(t|X_i) = a(X_i)e^{\beta * \log(RTS_i(t))}e^{-\lambda t}\lambda$

where $RTS(t) = \frac{TS(t)}{TS_0}$, a and λ are distributional parameters, and X_i is a vector of baseline predictive factors and effects.

1.Moore, H. A New Tumor Dynamics Mathematical Model. *American Conference on Pharmacometrics* (2016):Poster W–29.





Joint model provides good out-of-sample prediction of tumor dynamics and OS





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Modulation of individual TS parameters highlights dynamic impact on OS outcomes.

Survival

1.00

0.75

Depth Modulate



RTS

1.5

Difficult to predict how modulation of >1 parameter will impact OS.

Joint TGD-OS

Model

Moore model¹ for asymptotic tumor growth: $TS_{i}(t) = TS_{i,0}(5TS_{i,max})(1 - e^{-k_{i,g}t}) + TS_{i,0}e^{-k_{i,d}t}$

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Relationship

between

clinical metrics

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Multiple clinical measures could be used to describe tumor response

Effects on depth of response

- Difference in mean *best relative change* from baseline at t≤T.
- Difference in mean *relative change from baseline at t=T*.
- Difference in proportion of patients with best change from baseline at t≤T of ≤ 0, 10, 30, 50%.

Effects on durability of response

 Hazard ratio for <u>time-to-rebound</u> (20% growth from nadir; "tPFS").

T=[18, 27, 36, 45, 54, 63, 72] weeks





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Relationship

between

clinical metrics

Joint TGD-OS

Model

Prior distributions for new theoretical treatments

Using the reference model as a prior distribution for a new therapy, we constrain parameter space by the likely correlations between parameters observed previously

Derivation of Priors in THETA-space for simulation of novel therapies

	TS main effects	Other parameters
Chemo arm	Reference model standard error	Reference model standard error
New arm	Standard deviation = 1 "Pembro effect"	Reference model standard error

10

This gives ~16% probability that the novel therapy has an effect greater than Pembro.



Relationship

between

clinical

measures and

OS

Prior

distribution on

clinical

measures

OS is predicted to improve (HR<1 relative to chemo) when Kg and Tmax decrease, or Kd increases.





Converting THETA-space priors into clinical-space priors







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Modifying priors on clinical measures





Application of model to two hypothetical therapies

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Observed data at 18 weeks from randomly sampled data set of 20 patients/arm.

Tumor Metric Estimate



Application of model to two hypothetical therapies

Posterior

predictive

distribution

for OS HR

Observed clinical measures for new therapy

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Prior distribution or

clinical

measures

80% power to predict survival benefit with summary data from 20 patients/arm at 18 weeks, even with very strong prior distribution.

1.00 0.75 Cumulative Probability comparison Cherno vs cherno 0.50 Combo vs chemo Prior 0.25 **Combination therapy vs chemotherapy** Chemotherapy vs chemotherapy (Negative control) 0.00 -0.5 2.0 1.0 Overall Survival HR **IERCK** RESEARCH GROUP

Conclusions

- M&S enables integration of information across a spectrum of clinical observations
- In non-linear models, prior distributions in parameter space don't result in normal priors in clinical space
- Importance sampling can be used to generate multi-dimensional normal priors.
- Summary level data from a small cohort of patients can be leveraged to simulate expected clinical benefit.
- Further work:
 - Use likelihood profiles to dissect the specific contribution of each metric to OS
 - Understand how the likelihoods of each TS metric change over time
 - Apply this approach in comparator analysis setting with summary level data from literature.



