



Testing a Survival Extrapolation Algorithm for Cancer Immunotherapies: Pass or Fail?

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- Delta Hat conducted all analyses independently
- BMS provided helpful insight and comments

Survival modelling in Health Technology Assessment (HTA)

- Accurately extrapolating survival beyond trial follow-up is essential in HTA
- Different survival models can result in drastically different estimates of treatment benefits



Survival modelling in HTA: Immuno-oncology

- Immuno-oncology is especially affected because survival curves can flatten over time
- "Standard" survival models may not capture this
- "Flexible" survival models (including cure models) may capture the flattening, but are they credible?



Mean survival (years)	Control	10	Difference
Weibull model	0.77	1.45	0.67
Cure model	2.93	8.43	5.50

- Different models can result in drastically different estimates of survival benefit
- This has become a crucial discussion point in many appraisals of immuno-oncology treatments

Selecting survival models for immunotherapies

• Palmer *et al.* 2023 present an algorithm for selecting survival models to inform economic evaluations of cancer immunotherapies



- The algorithm involves 8 steps and 4 questions, involving:
 - Review of external data
 - Assessment of proportional hazards
 - Elicitation of expert beliefs
 - Consideration of turning points in the hazard function and data maturity
 - Evaluation of the possibility of cure
 - Sensitivity analysis

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 $\hat{\mathcal{S}}$ Delta Ha

Testing the Palmer et al. algorithm

CheckMate 649 compared first line nivolumab plus chemotherapy versus chemotherapy alone, in patients with advanced gastric, gastrooesophageal junction, or oesophageal adenocarcinoma

- 12-, 24- and 36-month data-cuts are available
- Pivotal trial in NICE TA 857

12-month data-cut

1.00 Nivolumab+chemotherapy Chemotherapy 0.80 0.60 Proportion 0.40 0.20 0.00 18 21 Time since randomisation (months) Nivolumab+Chemotherapy At-risk 473 438 Censored 0 Died 0 Chemotherapy 21 37 50 78 93 At-risk 482 Censored 0 Died



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Methods

- Applied Palmer *et al.* algorithm to 12- and 24-month data-cuts from CheckMate 649
- Selected preferred survival models based on our application of the algorithm
- Compared predictions to data observed in the 36-month data-cut (concentrating on the Combined Positive Score (CPS) ≥ 5 subgroup)

Aims

- **Applicability.** How straightforward is the algorithm to apply?
- Model selection. Does the algorithm result in survival models that extrapolate survival accurately?
- Implications for HTA. What impact might the algorithm have on the HTA process and decision making?
- **Development.** Would the algorithm benefit from any amendments?

Steps 1-8. Determining candidate survival models

- 1. Review external data [Evidence from a range of RCTs, real world data from the UK, US, Canada and the Netherlands]
- **2.** Assess proportional hazards [Hazard plots from CheckMate 649, external information on treatment effects of IOs]
- **3. Elicit expert beliefs** [Informed by all expert beliefs documented in the appraisal documents for NICE TA857]

4. Consider turning points in the hazard function	
5. Consider the need for flexible survival modelling	Informed by
6. Evaluate the possibility of a cure	Steps 1-3
7. Consider the plausibility and justifiability of a cure assumption	

8. Selection of potentially plausible models...



Steps 1-8. Determining candidate survival models

8. Selection of potentially plausible models...

- Based on responses to all previous steps

lssue	Chemotherapy	Nivolumab + chemotherapy	
Plausible	4 years: 3-7%	May approximately double	
expectations	10 years: 1-4%	survival that is observed with	
proportions		спепистегару.	
proportions		20 years: approximately 3%	
Expectations	Initial increase followed by decrease.		
around hazards	In both treatment groups hazards will converge towards background rates in the long-term (5+ years), but will always remain above background mortality levels.		
	Age-related increase in t term survivors.	the very long-term in longest-	
Treatment effect waning	Uncertain, but in the small percent who live beyond a few years, hazards will equalise		

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Findings: Plausibility criteria

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Findings: Candidate models

- Log normal models
- Log-logistic models
- Generalised gamma models
- Flexible parametric models
- Mixture cure models
- Non-mixture cure models

 →All models to include background mortality and SMR uplift
→All models to be fit independently to treatment groups

Results: Models fit to 12-month data-cut

Compared to Plausibility Criteria

- Only log-logistic and non-mixture cure models with a 15-year cure time-point met our plausibility criteria
- FPMs under-predicted survival at 10 years, and mixture cure models and non-mixture cure models with earlier cure time-points (7-10 years) over-predicted survival compared to our plausibility criteria

Model		Mean survival (50 years)	Survival %			AIC	Cure %
			Year 4	Year 10	Year 20		
12-month data-cut, CPS≥5							
1. Log-logistic, SMR=2.5	Chemo	1.44	6.1%	1.0%	0.1%	829	N/A
	Nivo+chemo	1.99	11.1%	2.0%	0.3%	906	N/A
2. Non-mixture cure (FPM, 4df),	Chemo	1.97	7.4%	3.6%	1.8%	829	5.7%
SMR=2.5, 15-year boundary knot	Nivo+chemo	3.28	15.3%	8.3%	4.1%	909	13.1%

- Log-logistic models predict survival at low end of plausible range
- Non-mixture cure models predict survival at top end of plausible range

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Results: Models fit to 12-month data-cut

Compared to 36-month data

- Models seem pessimistic: especially the log-logistic models
 - For nivolumab + chemo neither model produced survival predictions within the confidence intervals of the observed data at year 5



- Did not change much!
- The same models produced extrapolations that fell within the plausible range (log-logistic and non-mixture cure)
- FPMs still under-predicted survival and mixture cure models still seemingly over-predicted survival
- The log-logistic models seemed very pessimistic compared to the 36-month data-cut
- The non-mixture cure model seemed pessimistic, but did produce survival predictions that were within the confidence intervals of the observed data at year 5
- \rightarrow So similar results to those for models fitted to the 12-month data-cut, just with slightly improved extrapolations (as we would expect)

Applicability

• The Palmer *et al.* algorithm was simple to use and offered a systematic procedure for model selection

Model selection

- Allowed us to successfully narrow down the list of plausible models
- Resulted in models that provided credible extrapolations (though possibly pessimistic?)

Implications for HTA

- The algorithm "front-loads" the work, before models are fitted
- Should reduce disagreement around model choice; reduced need for additional modelling during appraisals

Development

- The Palmer *et al.* algorithm may benefit from some modifications
- The algorithm does not require that plausibility criteria are explicitly defined
- This provides an additional mechanism to ensure preferred models are selected in an unbiased manner

Thank you

