

Modelling Response

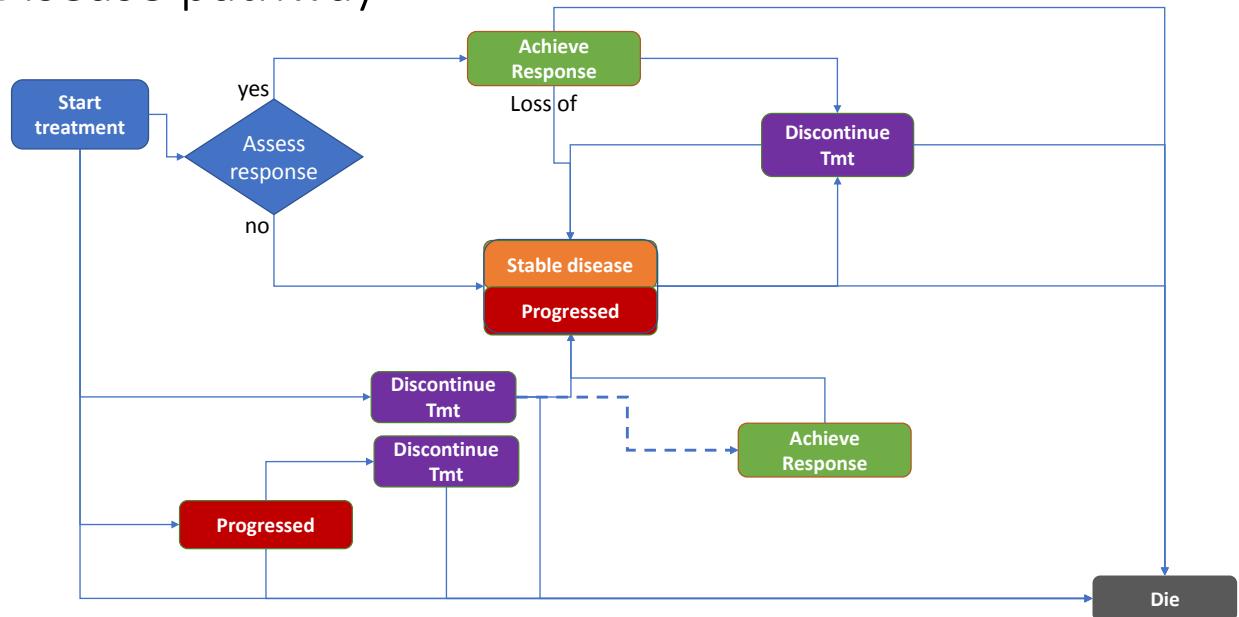
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Model Structures Incorporating Response

1. Partitioned Survival Analysis
 - A. Modified With Response
 - B. With Landmark Analysis
2. Time-to-Event

Disease pathway

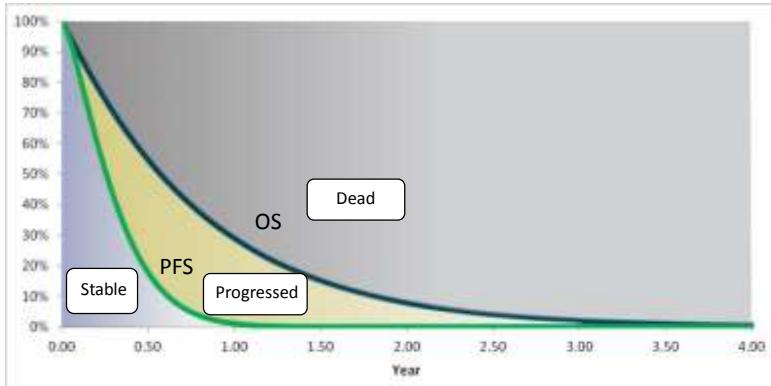


Partitioned Survival Analysis: Classic

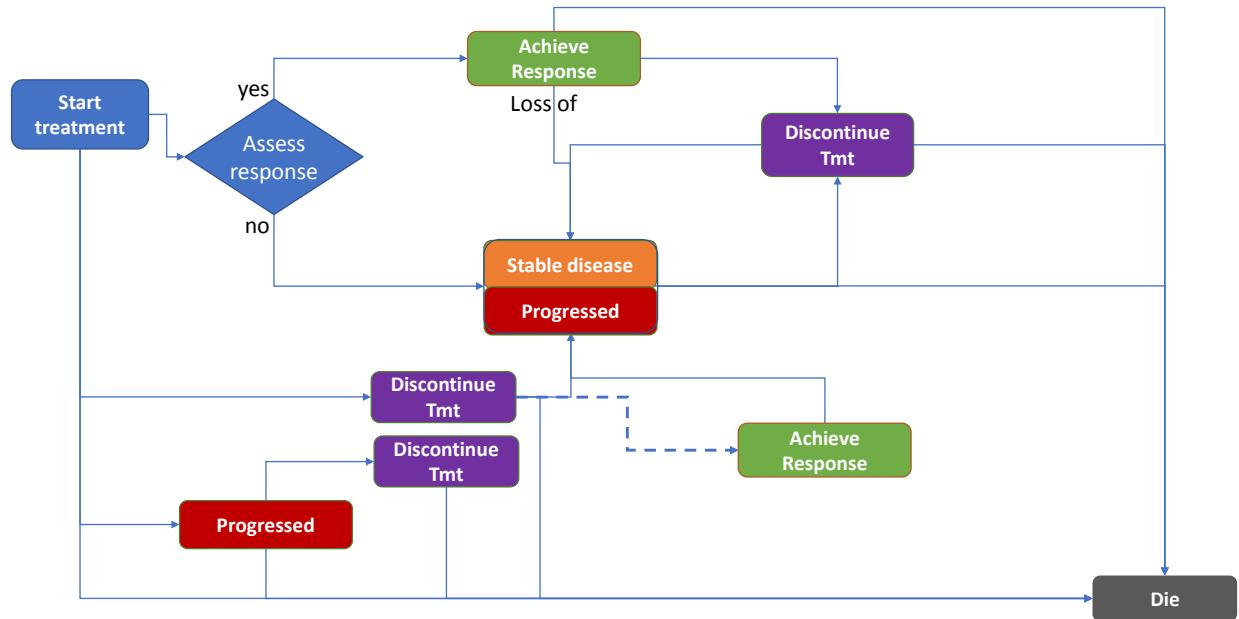
- Most common modeling technique in oncology
- Well received by HTAs and clinicians and easy and quick to implement
- Based on the historically most common clinical trial outcomes: OS and PFS

Underlying assumptions:

- Costs, utilities and survival depend on progression status
- Treatment costs often external to structure with time-to-treatment discontinuation curves.

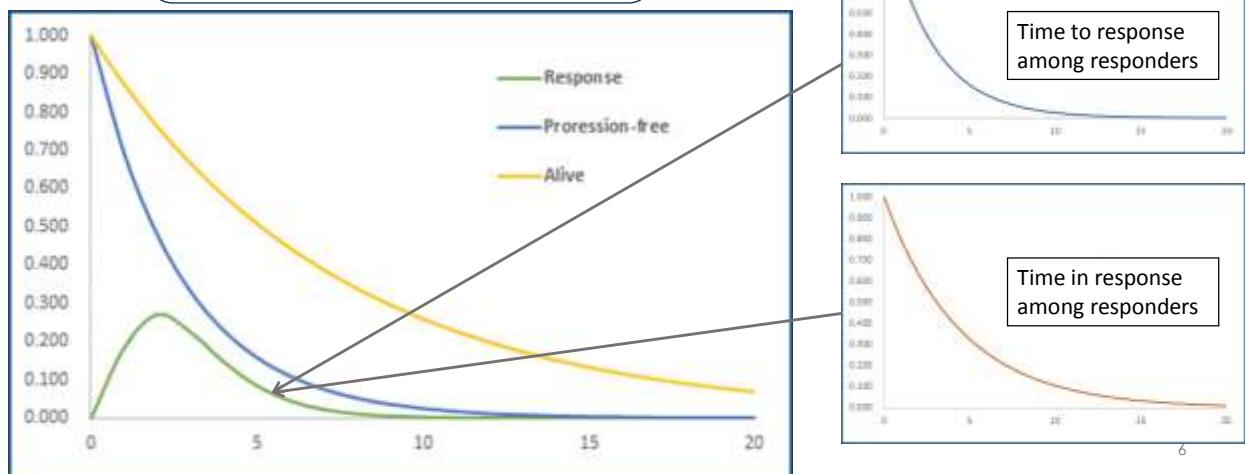


Disease pathway very limited with partition SA

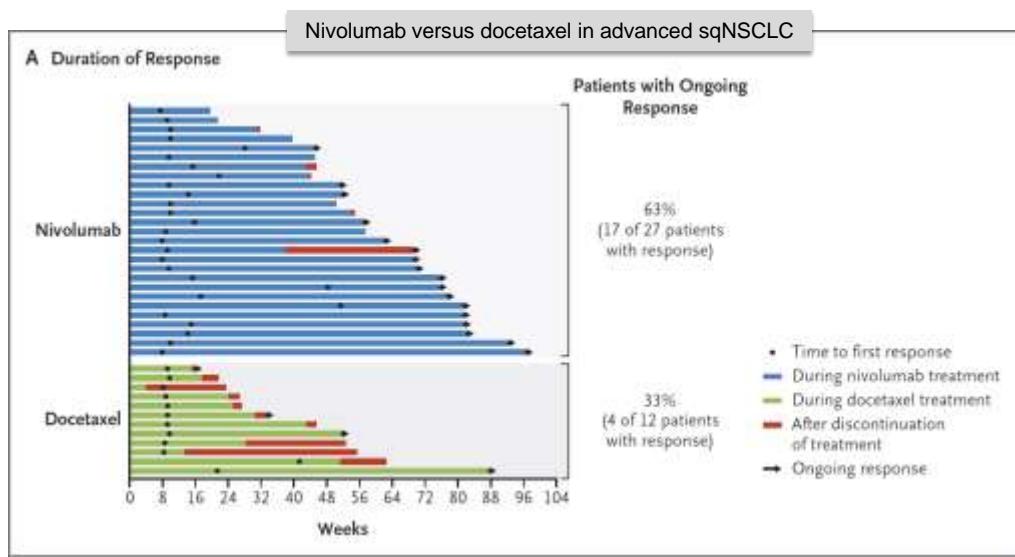


1A. Partitioned Survival Analysis: With Response

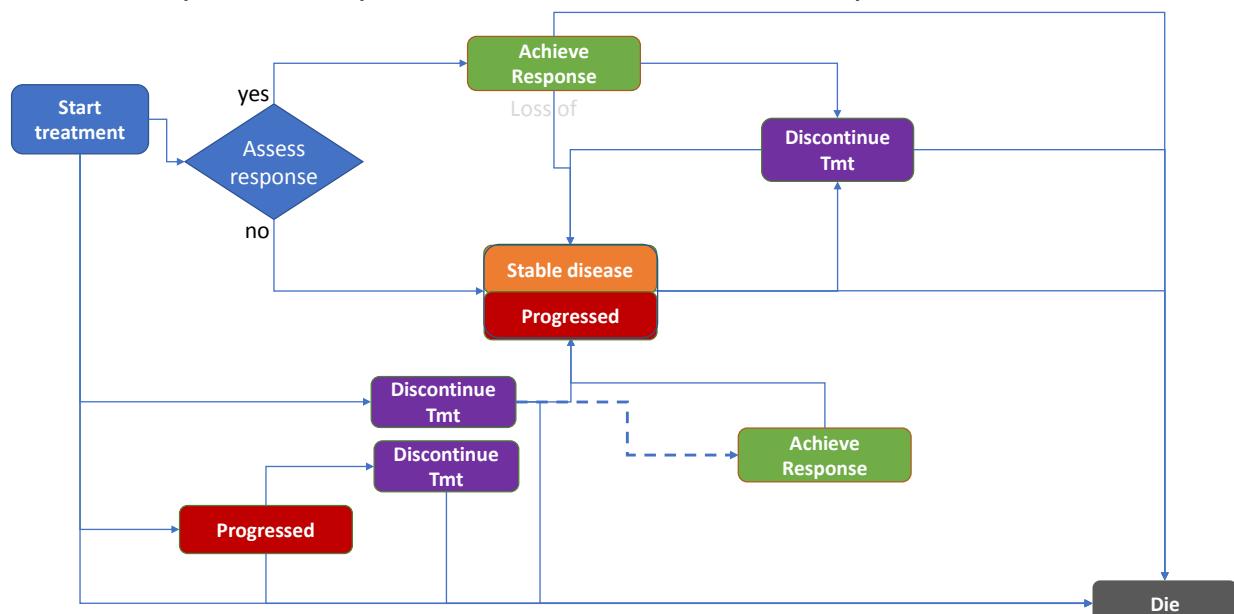
- Accepted structure with the use of response
- Can be extrapolated
- Data are often available
- Treatment costs often external to structure



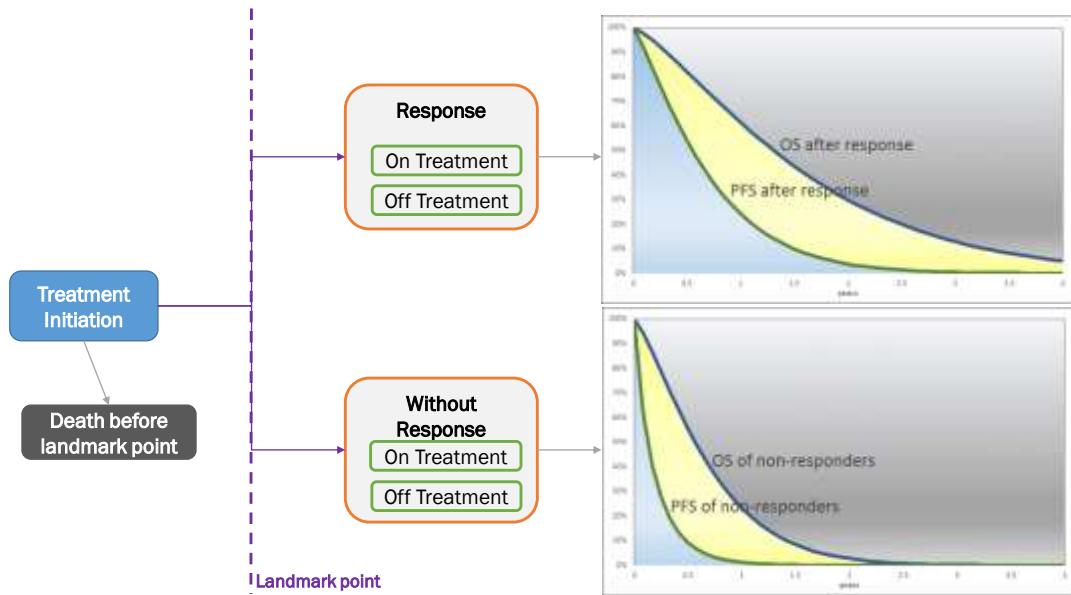
Example of Data for Modified Partitioned SA



Disease pathway still limited with response



1B. Partitioned Survival Analysis with Landmark Analysis



Partitioned Survival Analysis with Landmark Analysis

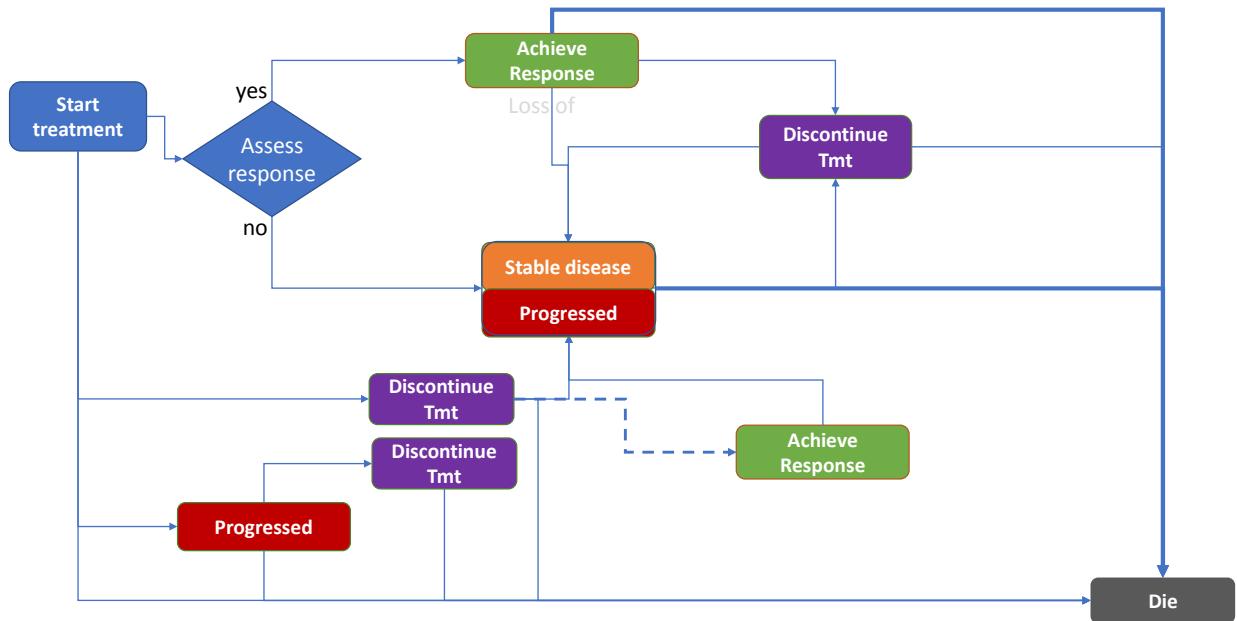
Advantages

- Allows modelling survival according to response
- Might be easier to fit survival, if it follows simpler patterns conditional on response
- Still simple and easy to understand
- Better clinical validity
- Keeps the partitioned survival structure

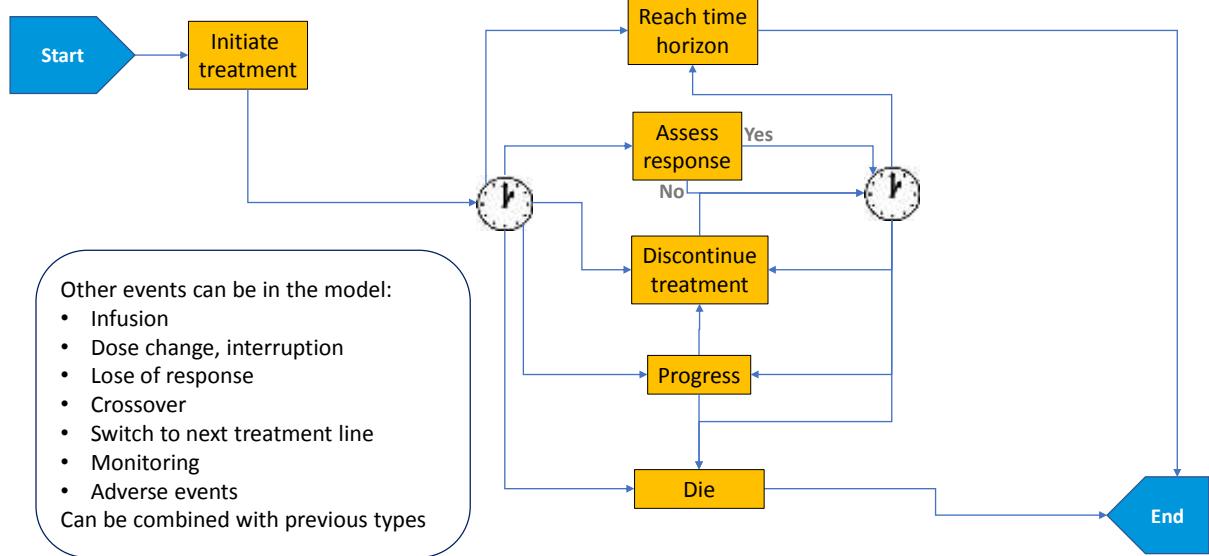
Disadvantages

- Choice of landmark point is subjective
- Results of clinical trials are rarely published in this format
- Due to fewer events, extrapolation of OS after response can be even more uncertain
- Potential for bias: patients surviving until landmark point can differ between responders and non-responders

Landmark analysis further extends pathway



2. Time to Event Model – Full pathway



TTE can model the full disease pathway

