

# Modelling Response

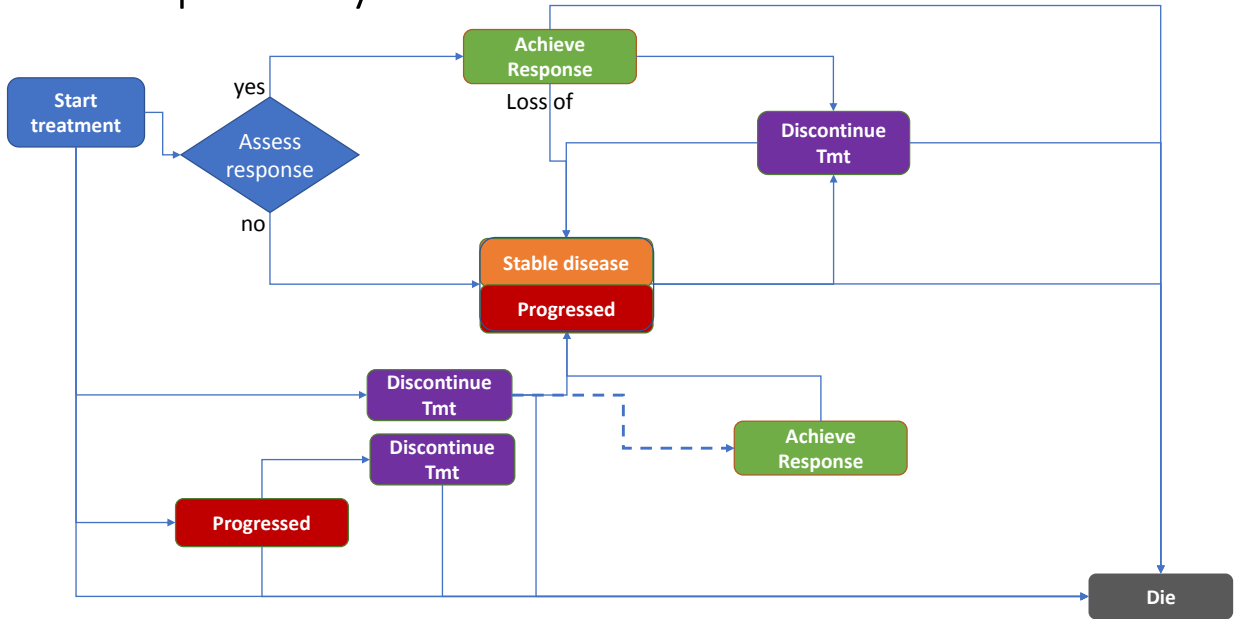
Jaime Caro



## Model Structures Incorporating Response

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

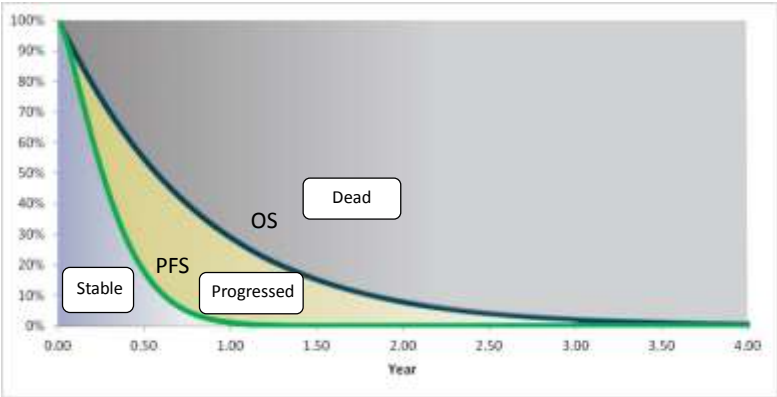
# Disease pathway



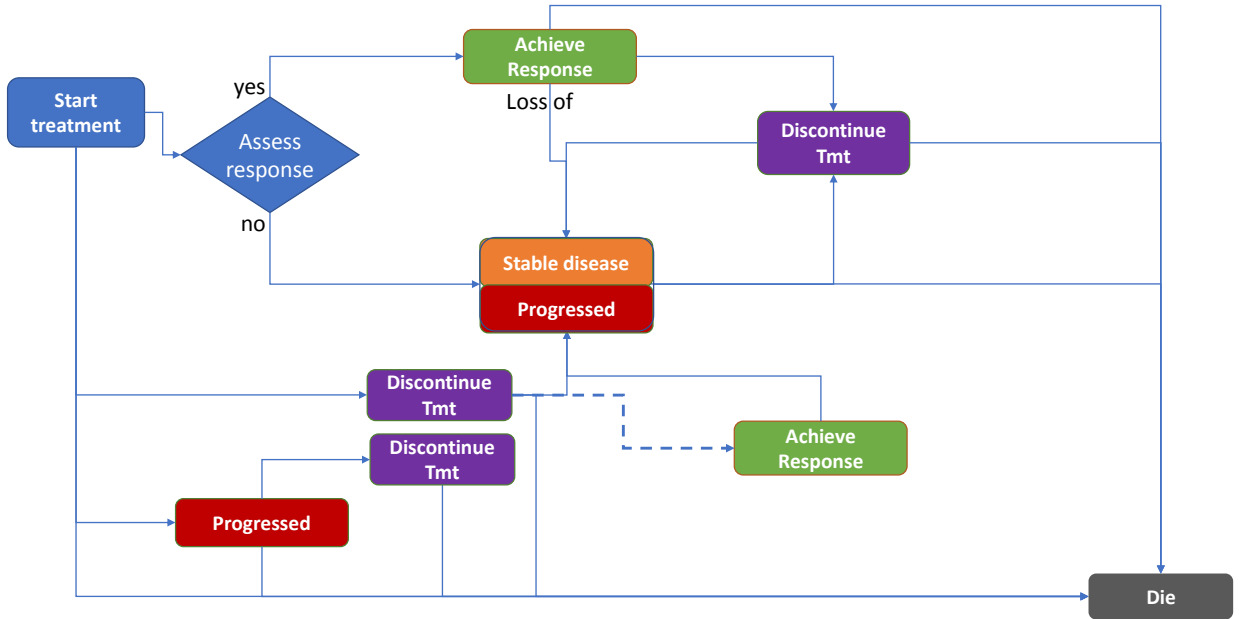
## Partitioned Survival Analysis: Classic

- 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.

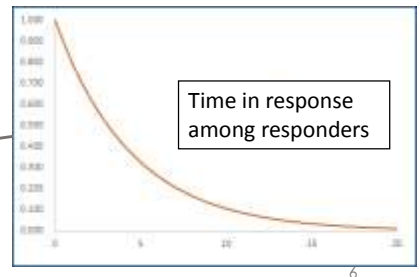
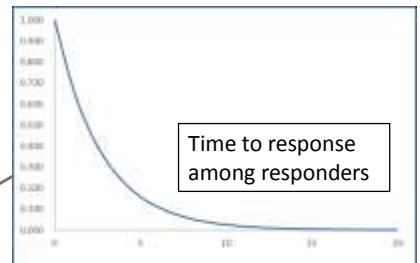
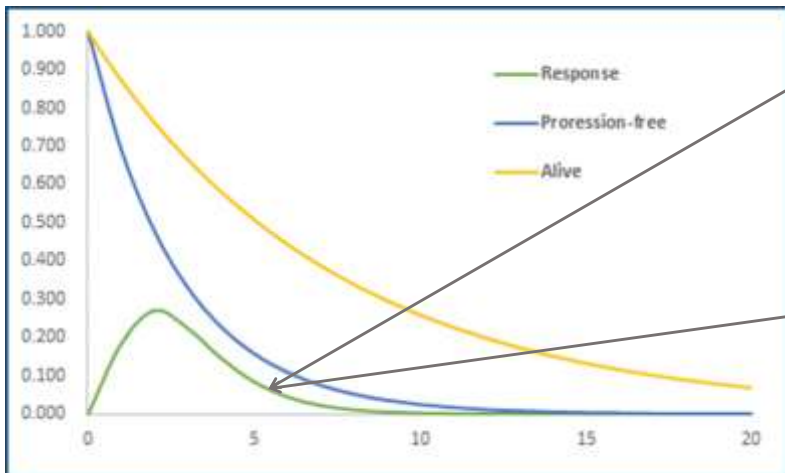


# Disease pathway very limited with partition SA

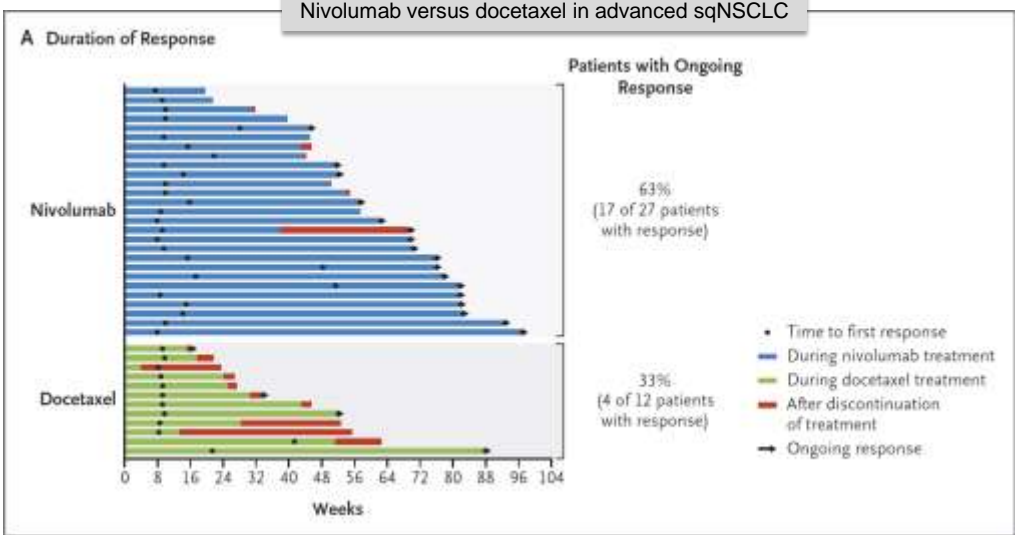


## 1A. Partitioned Survival Analysis: With Response

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

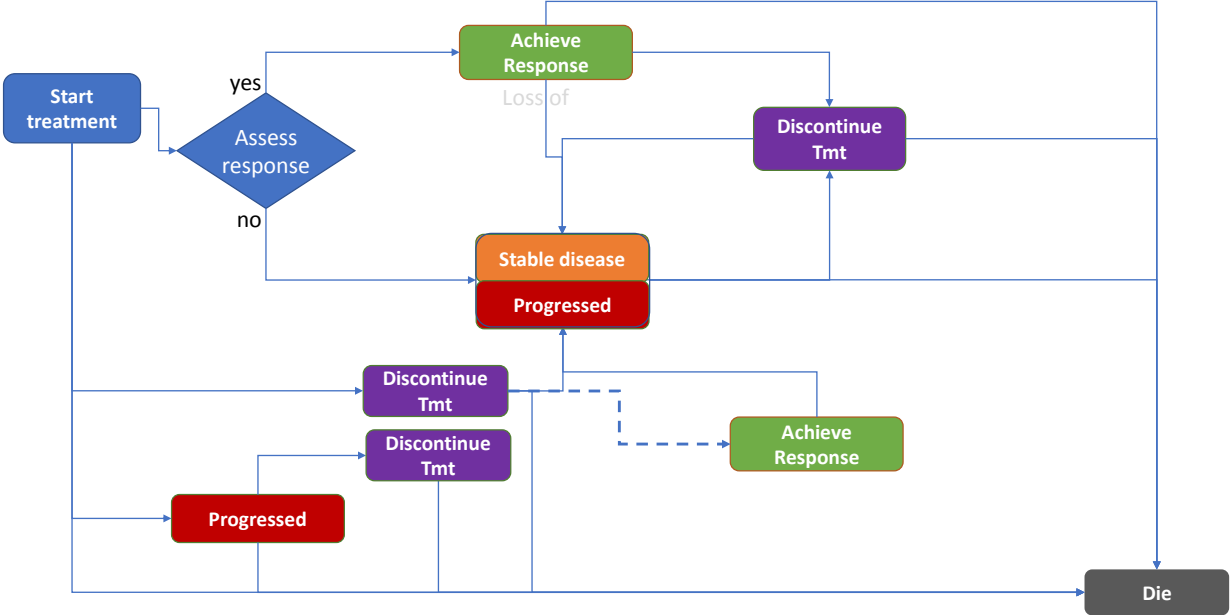


# Example of Data for Modified Partitioned SA

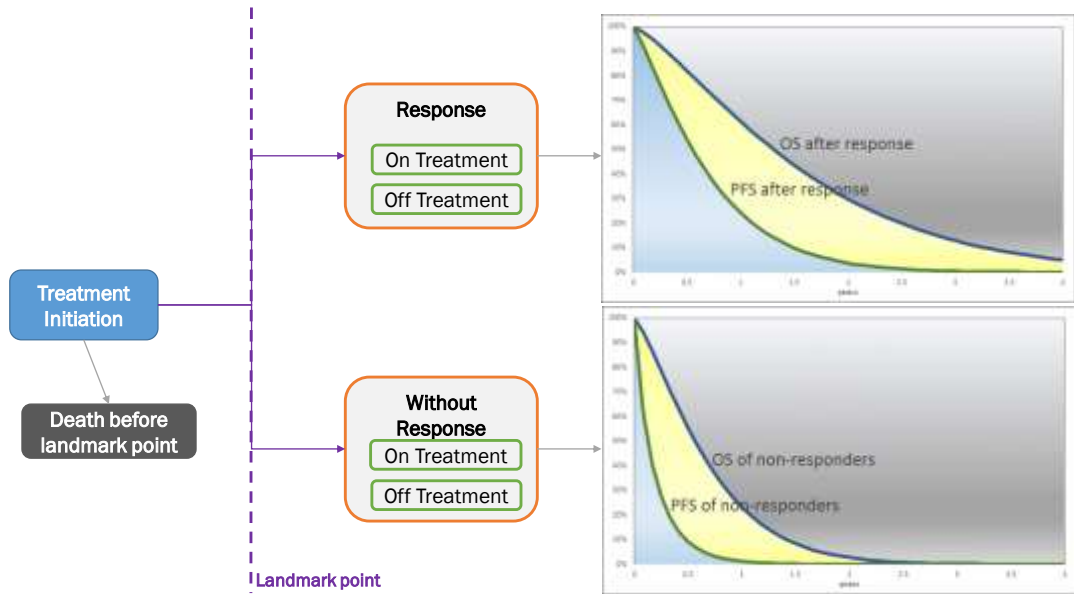


Source: Brahmer et al. 2015

## 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



# TTE can model the full disease pathway

