

# HoLISTIC Multistate Model (MSM) Working Group Meeting

5/10/24

# Outline

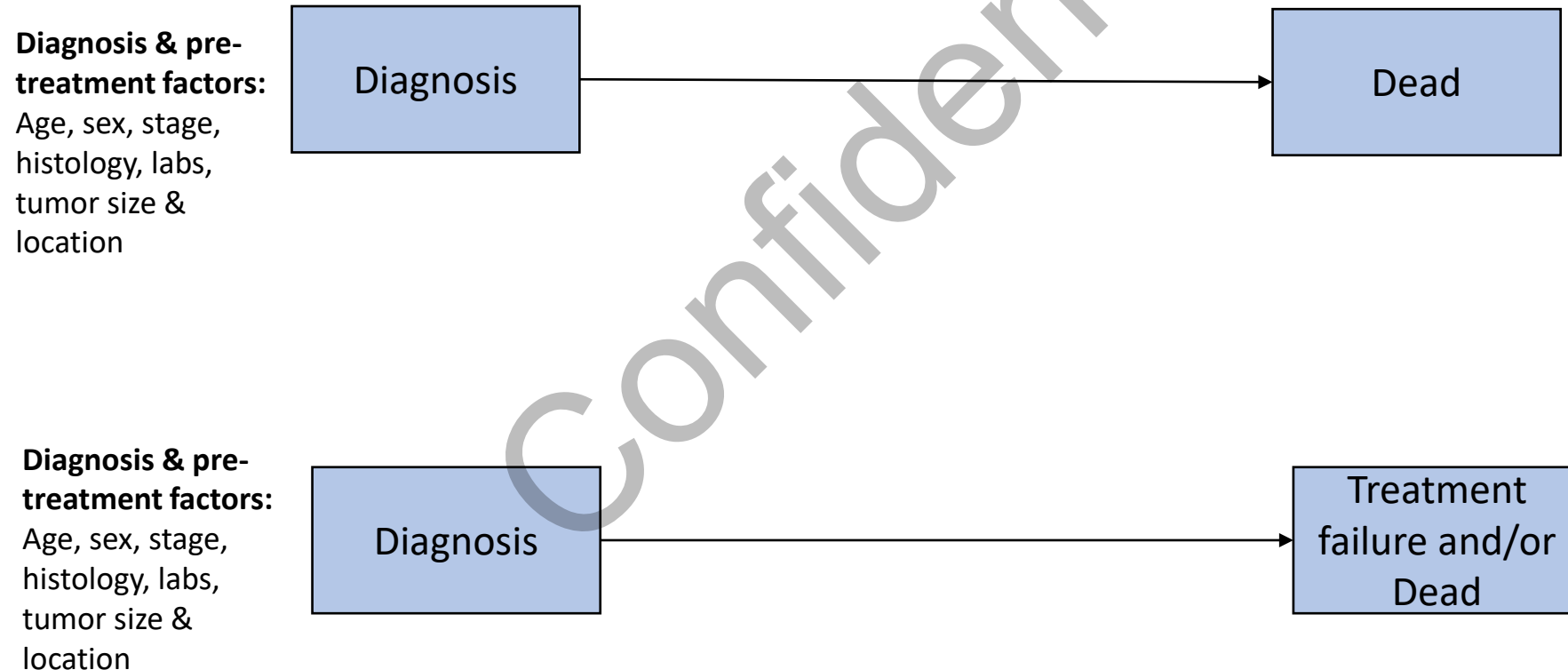
- Big picture & introduction to MSM
- Inclusion/exclusion criteria
- Baseline characteristics & outcomes by treatment arm
- Very preliminary MSM results
- Next steps

*\*Disclaimer: all results are very preliminary as we are still cleaning the data, but we wanted to begin testing the model.*

# Big picture

- Goal of this analysis is to predict outcomes for a future patient based on baseline disease characteristics and treatment
- First MSM iterations (through 5 years) will be based on clinical trials because outcomes are adjudicated, and we have detailed information about treatment (dates, dosages)
  - We will start with time-invariant treatment regimens (ECOG2496, HD2000, HD9601, Stanford V); these all have response based on CT scan
- Separate models for early and advanced stage trials
  - We will start with advanced stage trials

Reminder: we already built prediction models using pre-treatment factors & one outcome per model (A-HIPI, E-HIPI)



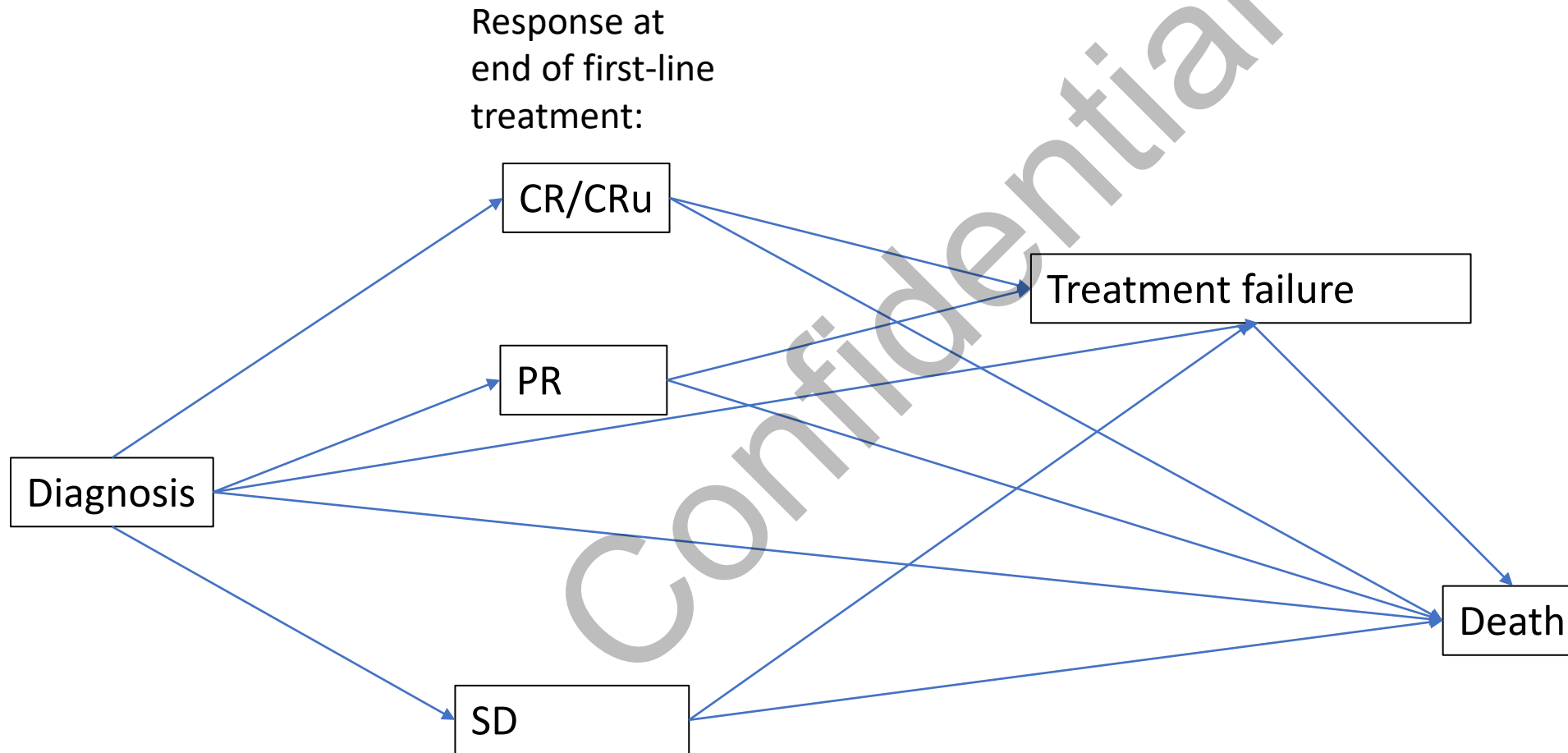
# Advantages of MSM

- Allows us to incorporate different disease states
  - Deals with the issue of non-informative censoring & competing risks
    - *Survival analysis assumes censoring is non-informative or independent, meaning that censored patients have similar outcomes as those who remain in the study.*
    - *Violated when competing events prevent occurrence of an event of interest (e.g., death prevents occurrence of relapse).*
    - *Furthermore, intermediate, non-fatal events that influence risk of future event (e.g., relapse changes risk of death) can also violate this assumption.*
- Estimates transition probabilities between disease states
  - Can have different covariates for each transition
- We will add treatment to the MSM

# 3-state MSM (“illness-death model”)



# MSM incorporating treatment response



CR=complete response; PR=partial response; CRu=complete response-unconfirmed aka complete clinical response; SD=stable disease; PD=progressive disease

# Current inclusion/exclusion for advanced stage MSM

- Inclusion

- cHL
- Age 18 to 65y
- Advanced stage trials with time-invariant treatment regimens
- Stage IIB, III, IV

- Exclusion

- No treatment received
- Extended field RT only
- Those treated with regimens no longer used (e.g., COPP/MOPP-like regimens)
- Those with inevaluable end-of-treatment scans



# Baseline characteristics by treatment arm, n=1579

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	ABVD, n=805	BEACOPP, n=89	Stanford V, n=685
<b>Study, n (%)</b>			
<b>1ECOG2496</b>	407 ( 50.6)	0 ( 0.0)	403 ( 58.8)
<b>5HD2000</b>	95 ( 11.8)	89 (100.0)	0 ( 0.0)
<b>6HD9601</b>	109 ( 13.5)	0 ( 0.0)	99 ( 14.5)
<b>13StanfordV</b>	194 ( 24.1)	0 ( 0.0)	183 ( 26.7)
<b>Age (years), mean (SD)</b>	34.64 (11.62)	33.08 (11.40)	34.85 (11.39)
<b>Female, n (%)</b>	370 ( 46.0)	35 ( 39.3)	305 ( 44.5)
<b>Stage, n (%)</b>			
<b>Stage IIB</b>	206 ( 25.8)	26 ( 29.2)	172 ( 25.2)
<b>Stage III</b>	285 ( 35.8)	44 ( 49.4)	241 ( 35.3)
<b>Stage IV</b>	197 ( 24.7)	19 ( 21.3)	155 ( 22.7)
<b>B symptoms, n (%)</b>	520 ( 66.2)	65 ( 73.0)	420 ( 63.2)

(currently there is variable-level missingness in these tables; multiple imputation not yet been applied)

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	ABVD, n=805	BEACOPP, n=89	Stanford V, n=685
<b>Histology, n (%)</b>			
<b>Lymphocyte depleted</b>	5 ( 0.6)	3 ( 3.4)	4 ( 0.6)
<b>Lymphocyte rich</b>	11 ( 1.4)	3 ( 3.4)	11 ( 1.6)
<b>Mixed cellularity</b>	79 ( 9.8)	7 ( 7.9)	64 ( 9.3)
<b>Nodular sclerosis</b>	550 ( 68.3)	68 ( 76.4)	466 ( 68.0)
<b>NOS</b>	50 ( 6.2)	7 ( 7.9)	50 ( 7.3)
<b>Any bulk<sup>a</sup>, n (%)</b>	400 ( 55.0)	48 ( 53.9)	327 ( 53.9)
<b>WBC count (10<sup>3</sup>/uL), mean (SD)</b>	9.64 (5.28)	10.74 (5.14)	9.65 (5.74)
<b>Lymphocyte count (10<sup>3</sup>/uL), mean (SD)</b>	1.43 (0.77)	All missing	1.35 (0.72)
<b>Hemoglobin (g/dL), mean (SD)</b>	12.08 (1.84)	11.53 (2.01)	11.98 (1.94)
<b>Albumin (g/dL), mean (SD)</b>	3.72 (0.63)	3.57 (0.67)	3.70 (0.66)
<b>ESR (mm/hr) (subset only), mean (SD)</b>	58.62 (36.28)	71.75 (36.85)	58.95 (36.45)

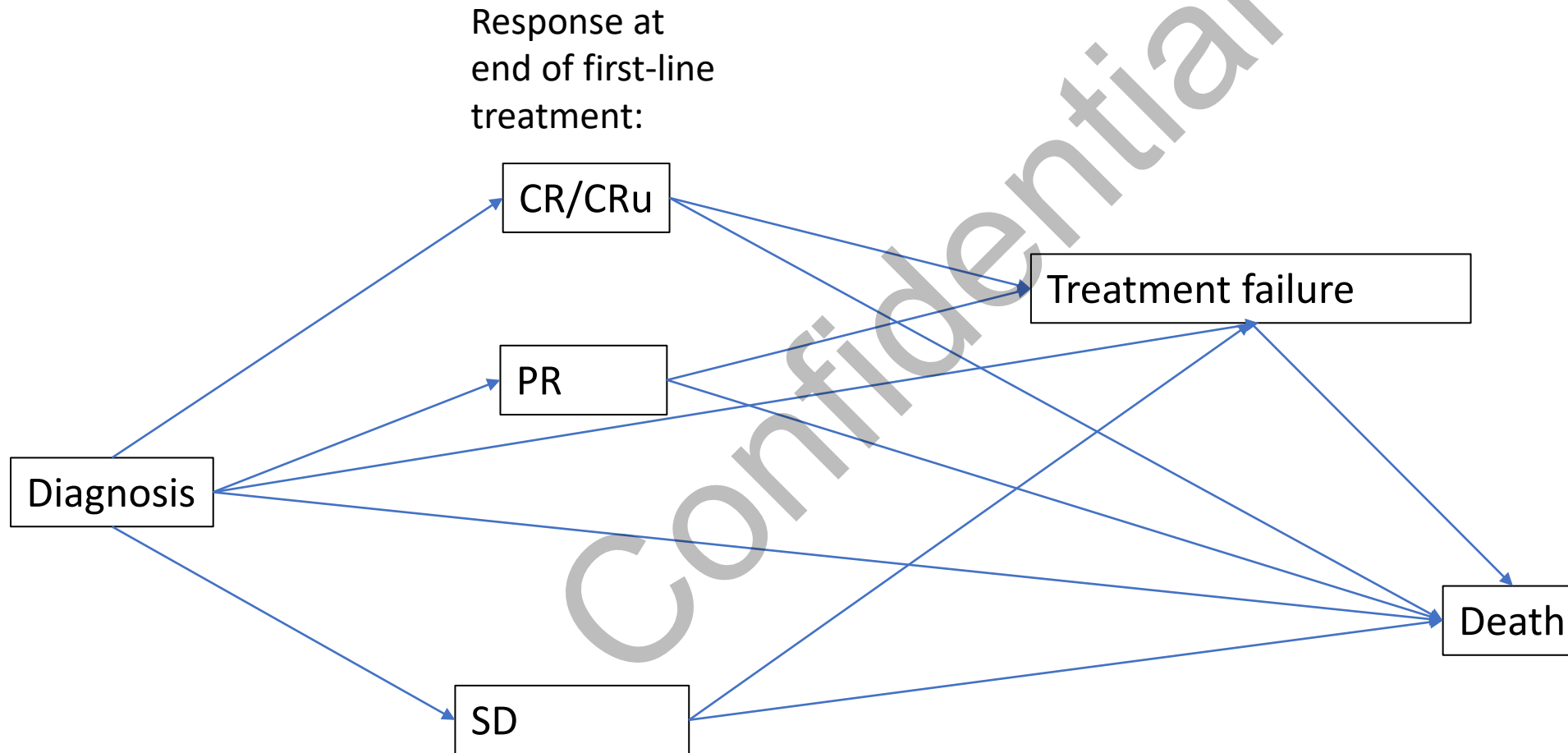
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# Outcomes by treatment arm

	ABVD, n=805	BEACOPP, n=89	Stanford V, n=685
<b>Outcomes</b>			
<b>Treatment response, n (%)</b>			
CR/CRu	527 ( 71.4)	81 ( 95.3)	315 ( 51.1)
PR	141 ( 19.1)	3 ( 3.5)	190 ( 30.8)
SD/PD	70 ( 9.5)	1 ( 1.2)	111 ( 18.0)
<b>No treatment response information</b>	67	4	69
<b>Treatment failure by 5 years, n (%)</b>	149 ( 18.5)	13 ( 14.6)	156 ( 22.8)
<b>Death by 5 years, n (%)</b>	69 ( 8.6)	8 ( 9.0)	71 ( 10.4)

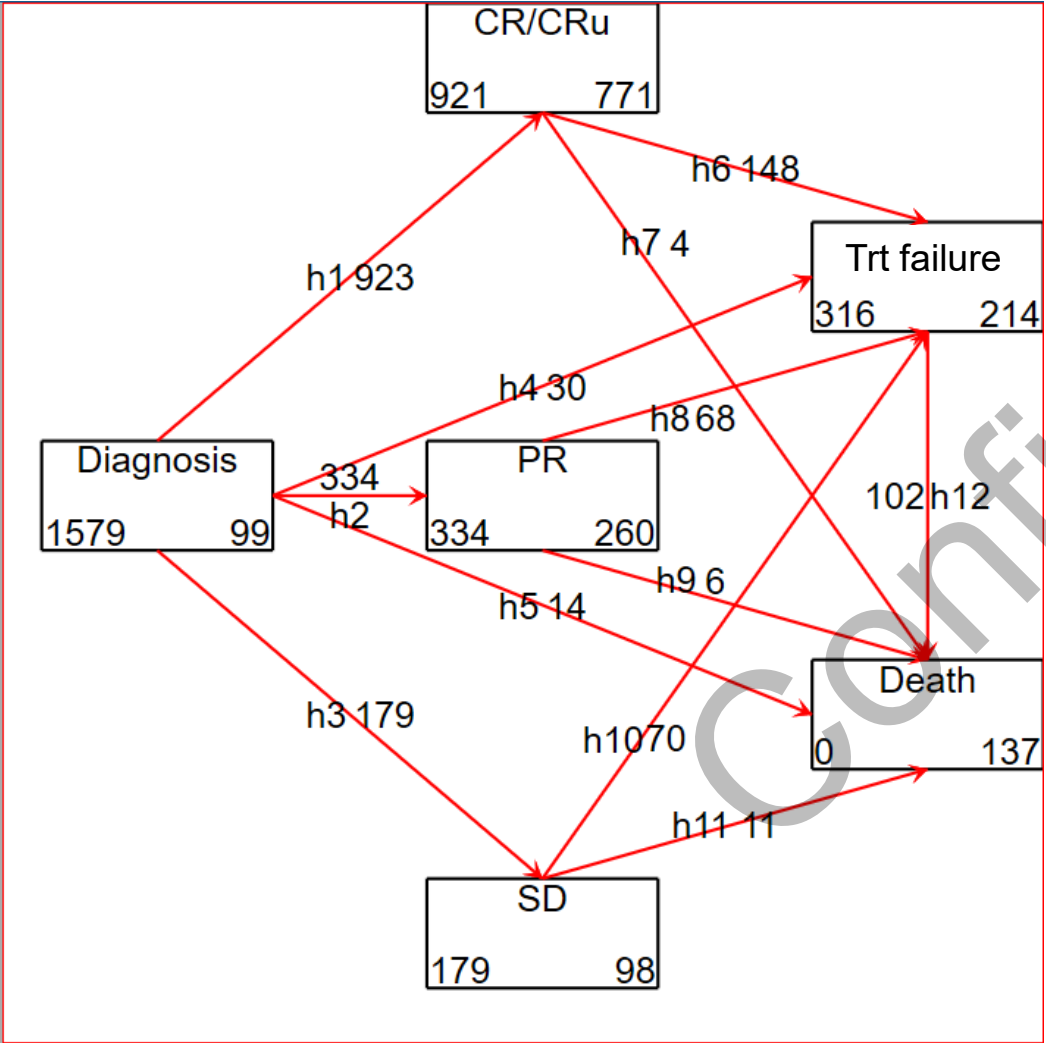
To do: Look at overlap between trt failure & death; look at PFS

# MSM incorporating treatment response



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# MSM incorporating treatment response

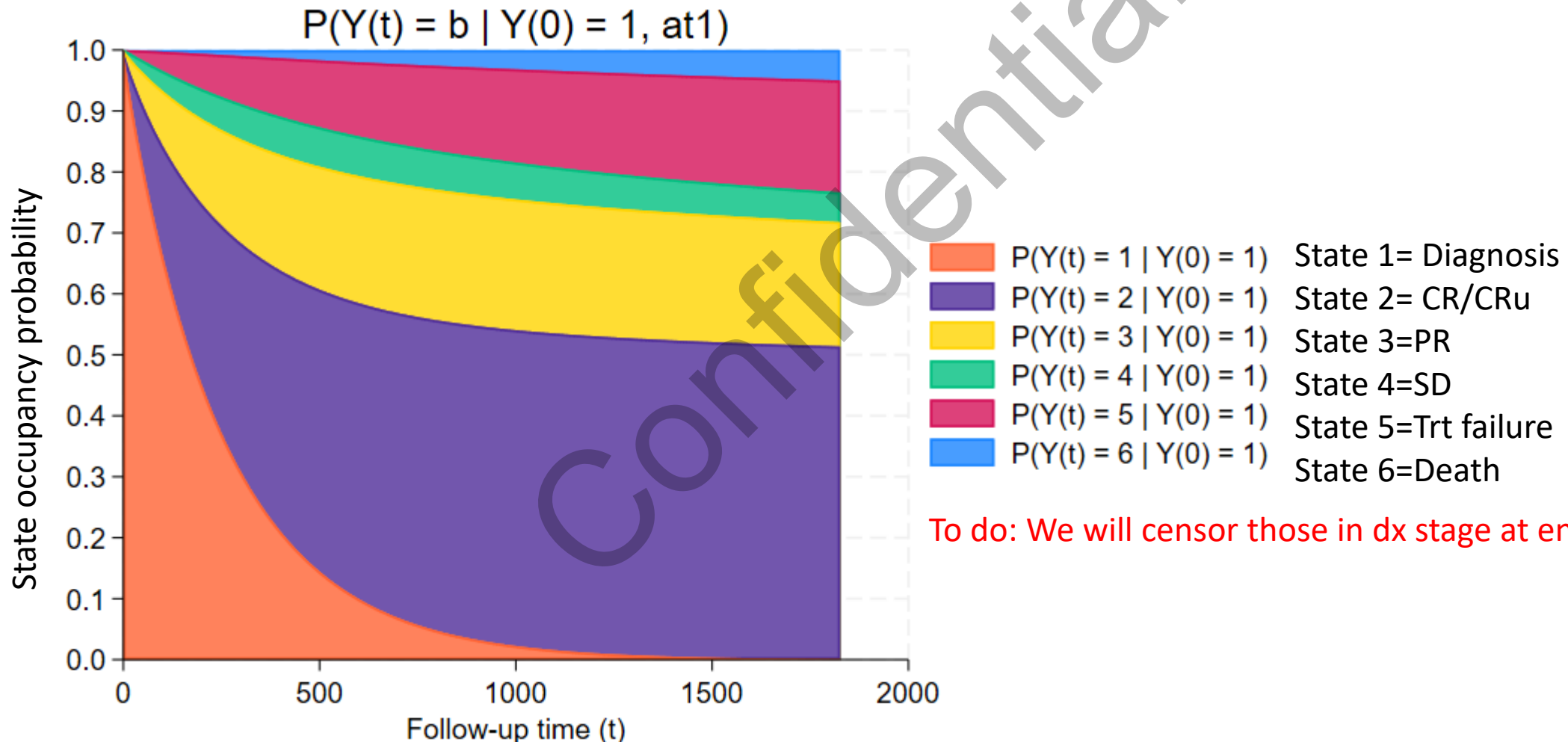


- Challenges:
  - Small Ns for some transitions
  - Estimated hazard ratios based on this model were unstable

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# State occupancy for MSM incorporating treatment response

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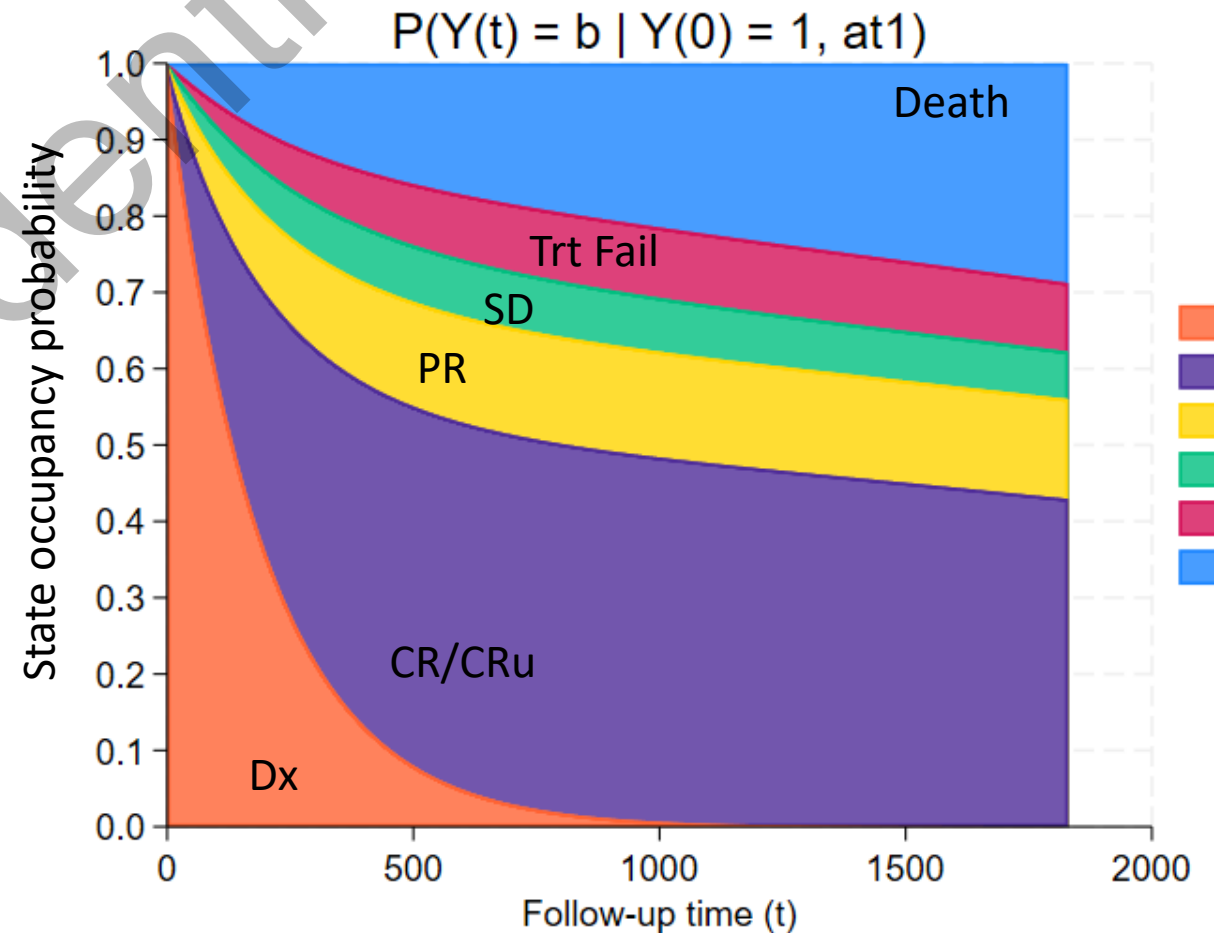
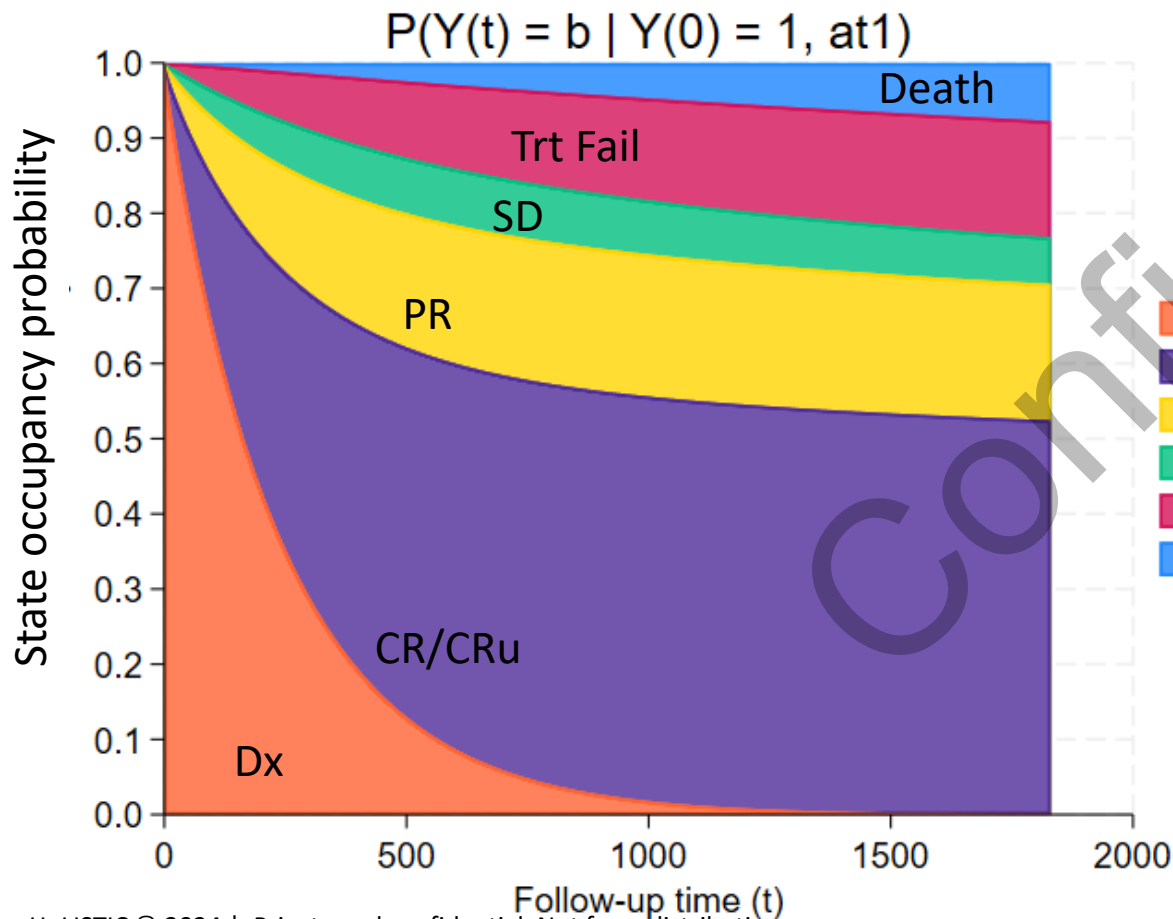
To do: We will censor those in dx stage at end of trt

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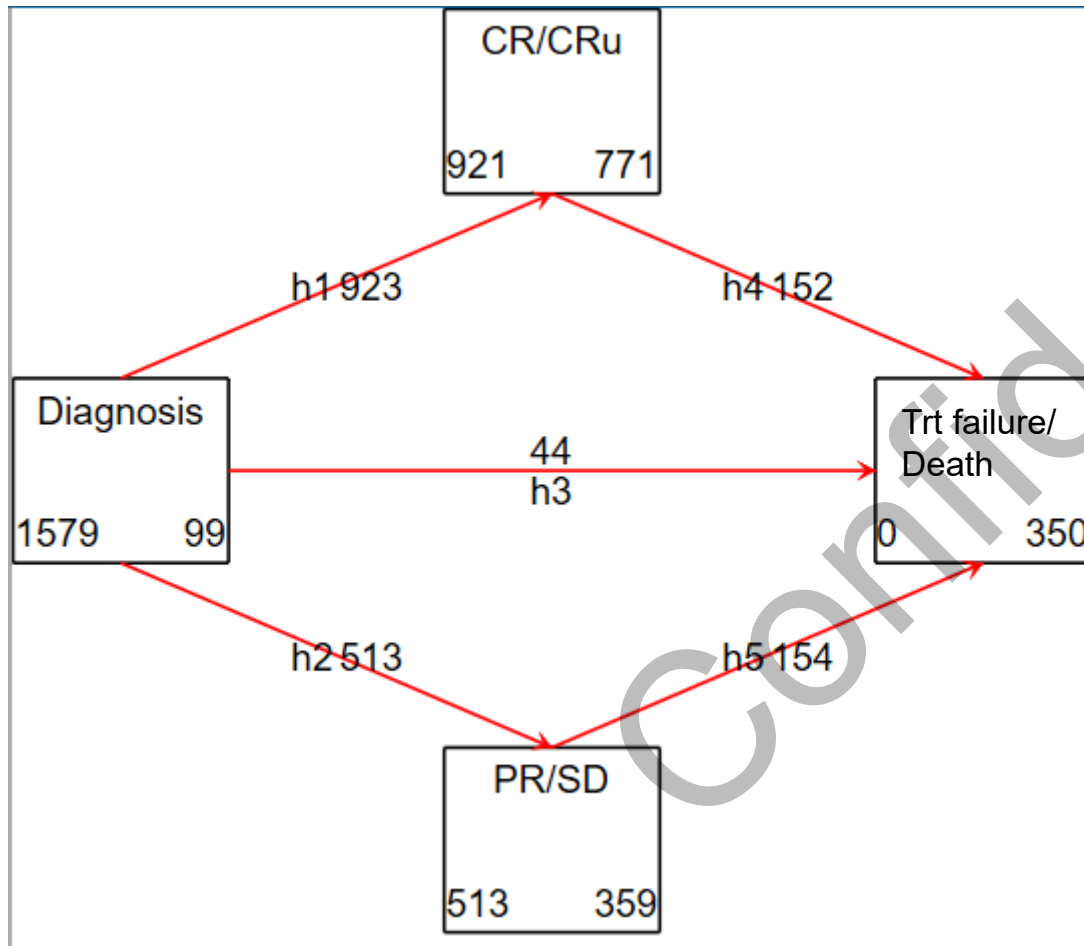
# State occupancy for MSM incorporating treatment response by age

**Age 30** To do: We will censor those in dx stage at end of trt

**Age 65**



# MSM incorporating treatment response & combining some states



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# Next steps

- Near term
  - Continue data cleaning and refining model structure
  - Impute missing data
  - Incorporate other covariates (e.g., age, AHIPI score)
    - Model building strategy; could use AHIPI in first transitions out of dx
  - Prepare ASH abstract (no tables/figures)
- Long term
  - Incorporate trials with PET-adaptive, time-varying treatment regimens
    - RT (y/n)
  - Validation of model results
  - Incorporate relapse/refractory & post-acute adverse effects
    - More detailed treatment variables here (e.g., chemo dose, RT site/dose)

# MSM Members as of May 10, 2024

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