

Modeling Progressive Disease Using Longitudinal Panel Data with Applications to Alzheimer's Disease



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Introduction

- Alzheimer's disease (AD) is a progressive disorder with no cure
- Three major clinical stages: Normal cognition, mild cognitive impairment (MCI), dementia/Alzheimer's disease
- MCI definition is broad and allows MCI - Normal transitions¹
- Often exclude reverters, a source of pathological information
- Best treatment available is to **delay the onset** of later stages
- Identifying individuals with highest risk of progression** is necessary for providing best treatment options and testing interventions in clinical trials
- Predicting the probability an individual will progress along disease pathway helps identify the highest-risk patients
- AD progression is continuous over time**, but assessment is completed at **annual exam**
- Repeated measurements at pre-scheduled times, **panel data**
- Interval-censored data** to investigate **transitions between stages**

Alzheimer's Disease Progression

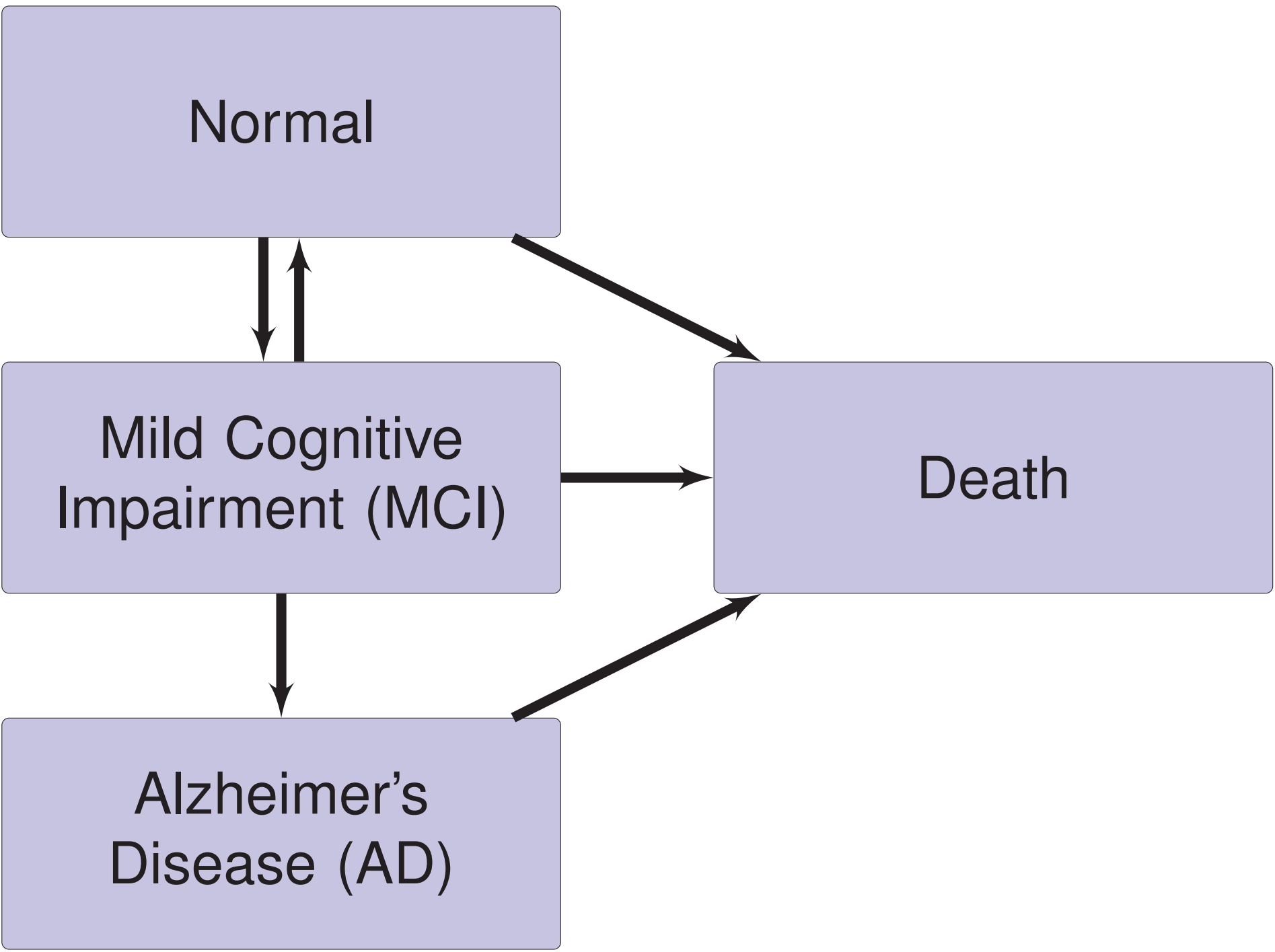


Figure 1: Progression between AD stages

National Alzheimer's Coordinating Center (NACC)

- Uniform Data Set (UDS) defines a common set of clinical observations collected longitudinally on participants at Alzheimer Disease Centers (ADCs)
- Information on **30,000+ participants from 31 ADCs** since 2005
- Majority are white, female, and have more than 3 follow-up visits
- 6329 normal participants, 10.2% converted to MCI or AD, 3% died
- 2740 MCI participants, 36.2% converted to AD, 4.8% died
- Cognitive complaint** is predictor of interest
- 4-levels: "No Complaint" (referent), "Self Complaint", "Informant Complaint", or "Both Complaint"

Observed Transition Paths between Visits Using NACC Data

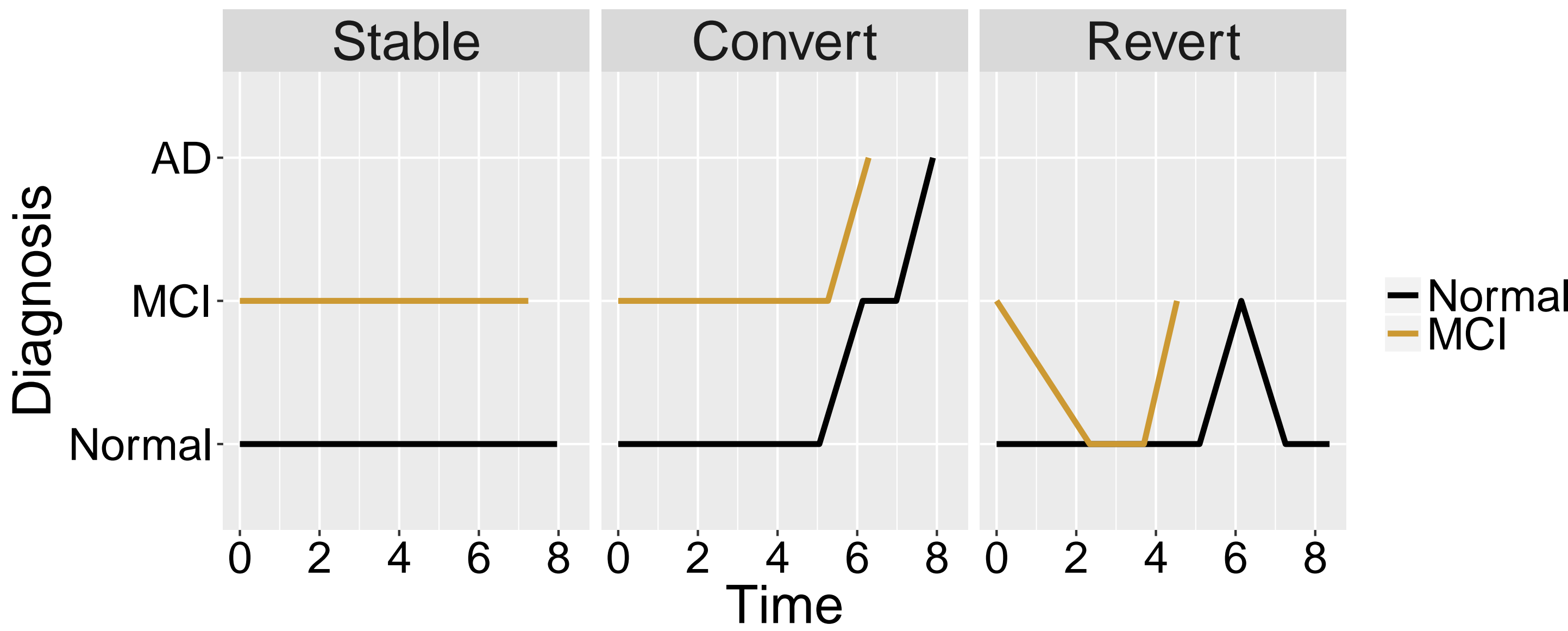


Figure 2: Example Disease Paths of NACC Participants

Multi-state Model Specifications

Figure 1 illustrates a multi-state model in continuous time, though the process is observed in discrete time, introducing **interval-censoring** to the model. We specify the **intensity matrix Q** , the matrix of individual transition intensities for consecutive visits, $q_{rs}(t, z(t))$, for each pair of states (r, s) . The process also depends on $z(t)$, a matrix of covariate values for each individual².

$$q_{rs}(t, z(t)) = \lim_{\delta t \rightarrow 0} P(S(t + \delta t) = s | S(t) = r) / \delta t \quad (1)$$

$$q_{rr} = - \sum_{s \neq r} q_{rs} \quad (2)$$

$$P(t) = \exp(tQ) \quad (3)$$

$$q_{rs}(z(t)) = q(0)_{rs} \exp(\beta_{rs}^T z(t)) \quad (4)$$

- Intensity represents **instantaneous risk** of moving from state r to state s , defined in equation (1)
- Intensities form Q , a matrix whose rows sum to zero, with diagonal entries defined by equation (2)
- Using equation (3) above, transition probabilities can be calculated from Q
- Transition intensities modeled as a function of predictors using equation (4)

Multi-state Model Assumptions

- Markov Assumption
 - Aim: Test Markov assumption by incorporating past history of MCI into model**
 - $q_{rs}(Z_t, \mathcal{F}_t) = q_{rs}(Z_t)$, \mathcal{F}_t is full history
 - Future state depends only on the current state**, not on the full sequence of previous events
- Time-Homogeneous Assumption
 - $P(u, t + u) = P(t)$
 - Transition matrix P is the same at each time point**
 - Transition probabilities at k^{th} step computed as P^k
 - Necessary for panel data to give **closed form of likelihood function** due to interval-censoring

Multi-state Model Results

Models chosen to **compare traditional method** (Model 1) with the **inclusion of reverters** (Model 2), and to compare these methods with **incorporating prior history of MCI** (Model 3) to address the Markov assumption:

- Model 1: Traditional baseline normal longitudinal dataset
- Model 2: Model 1 plus data from MCI participants that revert
- Model 3: Model 2 plus adjustment for prior MCI history

Table 1: Hazard ratios for baseline self complaint only

	Model 1	Model 2	Model 3
Normal - MCI	2.64 (2.10,3.33)	2.38 (1.92,2.96)	1.85 (1.48,2.32)
Normal - Death	0.15 (0.00,10.98)	0.22 (0.02,2.97)	0.30 (0.04,2.46)
MCI - Normal	1.93 (1.27,2.92)	1.18 (0.82,1.71)	1.04 (0.69,1.57)
MCI - AD	1.20 (0.63,2.27)	1.19 (0.63,2.23)	1.48 (0.81,2.71)
MCI - Death	0.30 (0.03,3.18)	0.34 (0.02,4.59)	0.13 (0.004,4.24)
AD - Death	3.18 (0.62,16.24)	2.82 (0.51,15.72)	2.99 (0.63,14.13)

Table 2: Hazard ratios for baseline both complaints

	Model 1	Model 2	Model 3
Normal - MCI	3.50 (2.74,4.47)	3.55 (2.80,4.49)	2.41 (1.87,3.12)
Normal - Death	0.41 (0.07,2.27)	0.59 (0.16,2.17)	0.53 (0.12,2.39)
MCI - Normal	0.99 (0.60,1.63)	1.21 (0.87,1.69)	1.19 (0.81,1.77)
MCI - AD	1.51 (0.81,2.83)	1.03 (0.57,1.87)	1.79 (0.98,3.26)
MCI - Death	0.51 (0.09,2.84)	0.39 (0.08,1.97)	0.33 (0.05,1.98)
AD - Death	1.18 (0.22,6.30)	1.00 (0.20,4.97)	1.08 (0.21,5.43)

Results of note:

- Hazard ratios consistent comparing between levels of complaint
- Model 1 overestimates the hazard ratio for MCI-Normal for self-complaint
- 11.6 (9.1,14.9)**: Hazard ratio for Normal - MCI when adjusting for past history

Discussion

- Advantages**
 - Models the transition intensities between stages, transformed into hazard ratios
 - Flexible choice in Q allows for model approximating biological process
- Disadvantages**
 - Requires assumption that all history is contained in previous diagnosis, **disregards full disease history**
 - All participants must have same initial state
- Previous studies use **cross-sectional methods** which treat transition between normal cognition and MCI the same as MCI to AD
- Potential method: **Partly Conditional Model**³
 - Flexible selection of past history to use in predicting future disease progression
 - Can incorporate baseline normal and MCI participants in same model

References

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