Lifetime risk, prevalence, and incidence estimates from ATN model after optimization

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1 Methods

A multistate model for Alzheimer's disease (AD) dementia using the dichotomous factors amyloidosis (A), tauopathy (T), neurodegeneration (N) and mild cognitive impairment (MCI) is shown in Figure 1. We refer to this model as the ATN model. The transitions $r_{ij}(a)$ are the one-year transition probabilities from state i to state j, conditional on surviving to age a + 1. To calculate lifetime risks of AD dementia, we estimate the transition rates $r_{ij}(a)$ using available information.

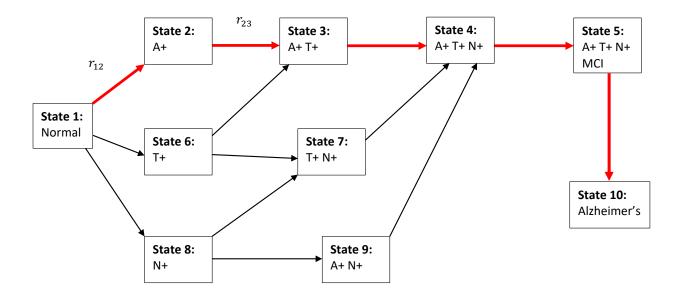


Figure 1: Multistate model for AD dementia with Amyloid (A+), Tauopathy (T+), Neurodegeneration (N+), and mild cognitive impairment (MCI)

1.1 Data and notation

We incorporate three sources of Alzheimer's disease (AD) dementia information into our analysis. These include age- and state-specific conditional prevalence rates $Pc_i(a)$ (Jack Jr et al. 2017), age-specific incidence rates I(a) of AD dementia (Brookmeyer et al. 2007), and age-specific probabilities C(a) of being in a preclinical state (Brookmeyer and Abdalla 2019). The prevalences $Pc_i(a)$ are conditional on being preclinical, i.e. no MCI has developed. The probabilities C(a) were calculated using a multistate model for AD dementia using only the factors A, N, and MCI, which we refer to as the AN model.

We will use the following notation throughout:

- $r_{ij}(a)$ is the one-year transition probability of a person at age a going from state i to state j;
- $Pc_i(a)$ is to the prevalence of state i for individuals age $a, a = 50, \ldots, 95$, conditional on being preclinical and alive (Jack Jr et al. 2017),
- $Pu_i(a)$ is the prevalence of state i for individuals age a, a = 50, ..., 95, conditional only on being alive. These were calculated as $Pu_i(a) = Pc_i(a) * C(a)$, where C(a) is the probability of being preclinical as calculated using the AN model (Brookmeyer and Abdalla 2019);
- I(a) is the incidence of AD dementia for individuals age $a, a = 65, \ldots, 90$ (Brookmeyer et al. 2007); its functional form is $I(a) = .00117 \exp(0.126a)$.
- k_{0ij} and k_{1ij} are transition parameters from state i to state j such that $r_{ij}(a) = k_{0ij} \exp(k_{1ij}a)$

• $\widehat{Pc}_i(\mathbf{k}, a)$, $\widehat{Pu}_i(\mathbf{k}, a)$, and $\widehat{I}(\mathbf{k}, a)$, are estimates of conditional prevalence rates, unconditional prevalence rates, and incidence rates of AD dementia from the multistate model for a given set of transition parameters \mathbf{k}

1.2 Optimizing transition rate parameters

There are 14 total transitions r_{ij} , including 12 preclinical transitions (r_{12}, \ldots, r_{94}) , r_{45} , and $r_{5,10}$. Each transition has log-linear form $r_{ij}(a) = k_{0ij} \exp(k_{1ij}a)$, and we constrain each pair (k_{0ij}, k_{1ij}) with

$$\log(k_{0ij}) < 0$$

$$k_{1ij} > 0$$

$$\log(k_{0ij}) + k_{1ij} \times 95 < 0$$

$$(1)$$

such that every transition $r_{ij}(a) \in (0,1)$ for all ages $a \in (50,\ldots,95)$.

1.2.1 Transitions r_{45} $r_{5,10}$

We assume $r_{5,10}(a) = 0.3$ for all ages a, i.e. $k_{05,10} = 0.3$ and $k_{15,10} = 0$. To estimate the parameters k_{045} and k_{145} we assume that

$$r_{45}(a) = \lambda(a)\widetilde{r}_{45}(a), \tag{2}$$

where $\lambda(a) = \frac{P_4(a)}{P_4(a) + P_9(a)}$ is the proportion of those with both A+N+ that also have T+, and $\tilde{r}_{45}(a)$ is the transition from State 4 (A+N+) to State 5 (A+N+ MCI) in the AN model. The ATN model assumes that only those with A+T+N+ can develop MCI, so we assume that those transitioning to MCI in the AN model were only coming from the subpopulation of A+N+ subjects that also had T+. Having calculated $r_{45}(a)$ using (2), we assume a log-linear form and perform a simple constrained optimization to find the parameters k_{045} and k_{145} using the base constr0ptim function in R.

1.2.2 Preclinical Transitions

Here we use k to represent the vector of 24 parameters $(\log(k_0), k_1)_{(ij)}$ for preclinical transitions (r_{12}, \ldots, r_{94}) . Optimization is with respect to the parameters k. We use the COBYLA optimization algorithm in the R package nloptr, which accepts both linear and nonlinear constraints. In addition to the constraints (1) we

impose constraints such that the sum of transitions out of state i for age a is less than 1 up to age 95

$$\log\left(\sum_{j\in R_i} \exp(\log(k_{0ij}) + k_{1ij} \times 95)\right) \le 0 \quad \forall i \in I$$
(3)

where R_i is the set of states j one can transition to out of state i and I = (1, 2, 3, 4, 6, 7, 8, 9) is the set of preclinical states.

The loss function f to be minimized has two components – one for prevalence and one for incidence

$$f(\mathbf{k}) = \frac{1}{n_1} \sum_{a=65}^{90} \left(\log(\widehat{I}(\mathbf{k}, a)) - \log(I(a)) \right)^2 + \frac{1}{n_2} \sum_{i \in I} \sum_{a=50}^{95} \left(\log(\widehat{P}u_i(\mathbf{k}, a)) - \log(Pu_i(a)) \right)^2.$$
(4)

where $n_1 = 26$ and $n_2 = 8 \times 46$ are the number of elements contributing to the incidence portion and prevalence portion of the loss function, respectively, and I is the set of preclinical states. Because prevalence and incidence are on different scales, we use squared log-ratios instead of squared absolute differences. We weight each portion with the reciprocal of the number of contributing elements to give "equal" weight to the incidence and prevalence portions of the function.

2 Results

2.1 Lifetime Risks

Tables 1 and 2 show the lifetime risks of developing AD dementia for women and men, respectively, for ages $a \in (60, 65, ..., 90)$ and each pre-AD state $i \in (1, ..., 9)$.

Table 1: Lifetime risk of AD for women by age and disease state

			1				1		
Age	Normal	A	A+T	A+T+N	A+T+N+MCI	Т	T+N	N	A+N
60	17.8 %	30.07 %	54.75 %	58.37 %	95.63 %	38.8 %	27.26 %	10.06 %	28.69 %
65	16.49 %	28.4 %	51.96 %	56.18 %	93.57 %	36.72 %	24.09 %	9.46 %	26.7 %
70	14.66 %	25.9 %	47.92 %	52.79 %	90.06 %	33.63 %	20.32 %	8.57 %	23.97 %
75	12.25 %	22.41 %	42.29 %	47.85 %	84.71 %	29.32 %	16.07 %	7.35 %	20.42 %
80	9.28 %	17.8 %	34.74 %	40.83 %	76.19 %	23.59 %	11.52 %	5.76 %	16.02 %
85	6.07 %	12.39 %	25.38 %	31.6 %	63.81 %	16.7 %	7.16 %	3.94 %	11.07 %
90	3.24 %	7.16 %	15.56 %	20.98 %	46.68 %	9.85 %	3.66 %	2.24 %	6.42 %

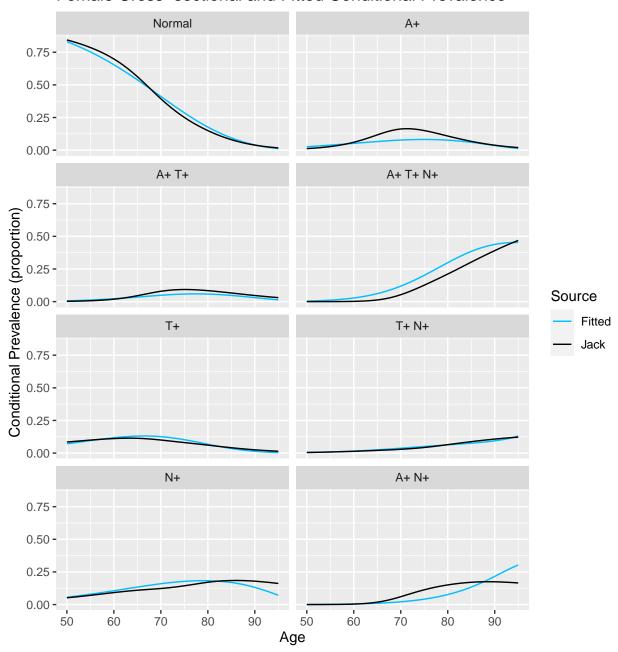
Table 2: Lifetime risk of AD for men by age and disease state

Age	Normal	A	A+T	A+T+N	A+T+N + MCI	Т	T+N	N	A+N
60	12.09 %	21.94 %	45.32 %	49.39 %	92.93 %	29.87 %	20.65 %	6.42 %	21.35 %
65	11.17 %	20.71 %	42.84 %	47.49 %	90.37 %	28.19 %	18.1 %	6.05 %	19.83 %
70	9.82 %	18.72 %	39.04 %	44.27 %	85.99 %	25.55 %	15.04 %	5.45 %	17.63 %
75	8.05 %	15.94 %	33.82 %	39.59 %	79.54 %	21.89 %	11.64 %	4.61 %	14.79 %
80	5.9 %	12.31 %	26.98 %	33.02 %	69.91 %	17.11 %	8.09 %	3.53 %	11.3 %
85	3.66 %	8.21 %	18.97 %	24.79 %	56.66 %	11.62 %	4.81 %	2.31 %	7.53 %
90	1.77 %	4.38 %	10.91 %	15.73 %	40.15 %	6.35 %	2.29 %	1.21 %	4.08 %

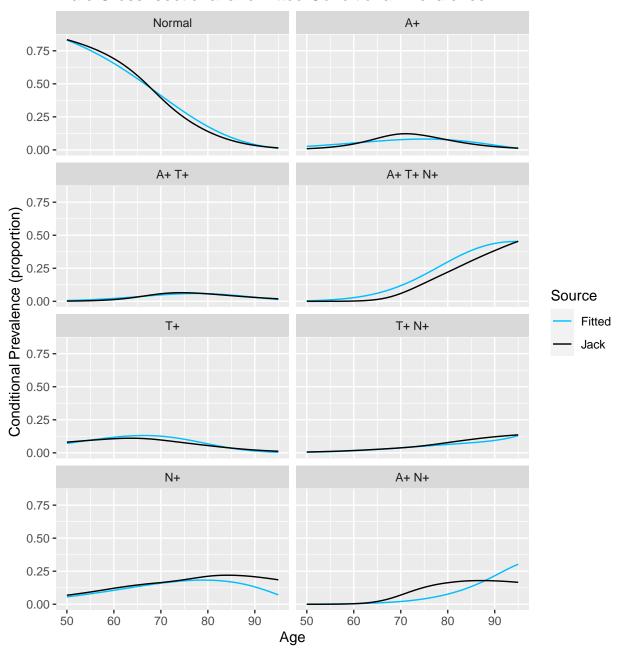
2.2 Conditional Prevalence

Female and male prevalence rates, conditional on being alive and in a preclinical state.

Female Cross-sectional and Fitted Conditional Prevalence



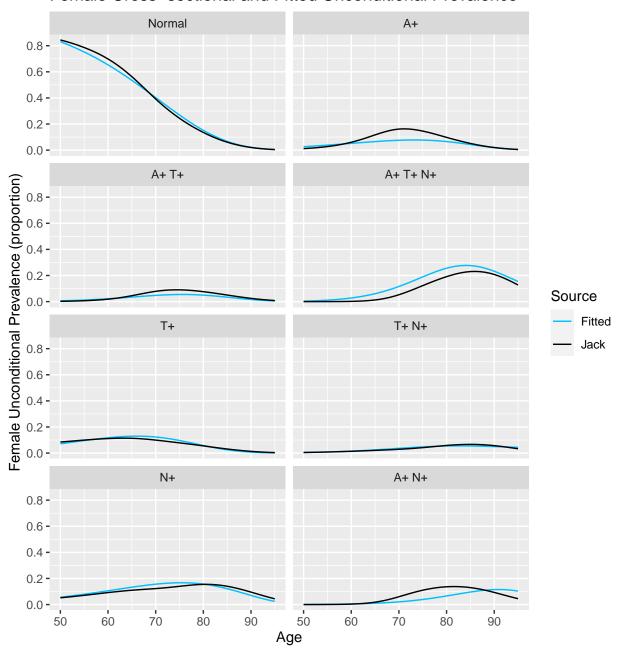
Male Cross-sectional and Fitted Conditional Prevalence



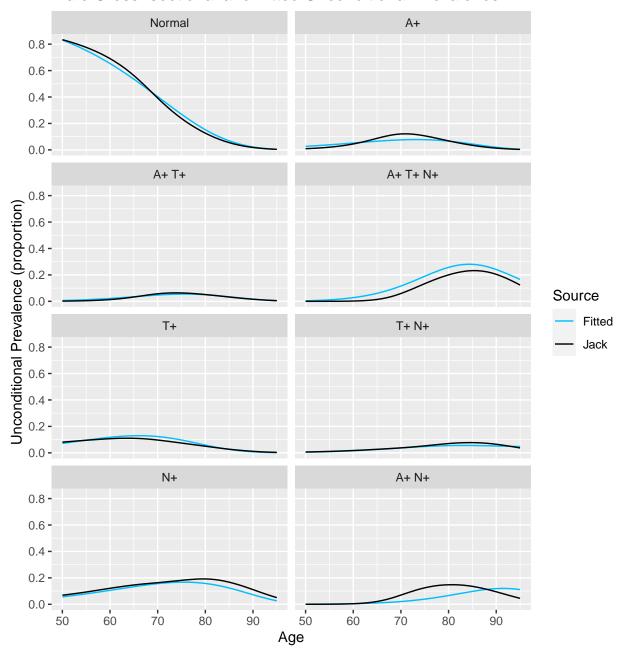
2.3 Unconditional prevalence

Female and male prevalence rates, conditional only on being alive.

Female Cross-sectional and Fitted Unconditional Prevalence



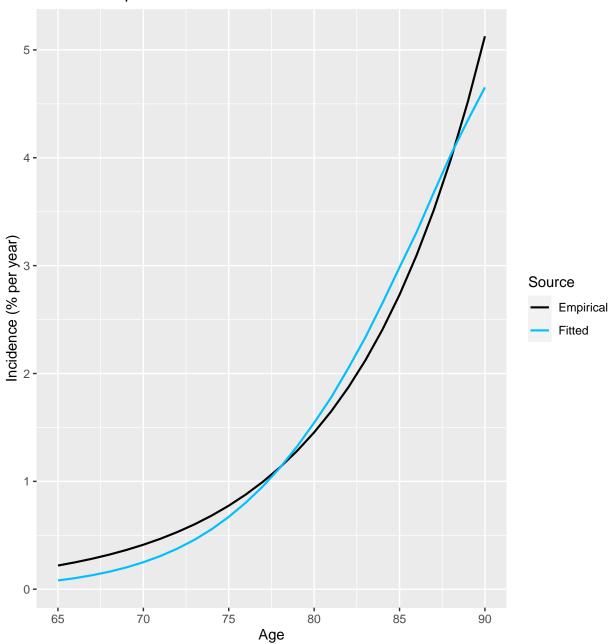
Male Cross-sectional and Fitted Unconditional Prevalence

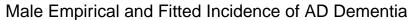


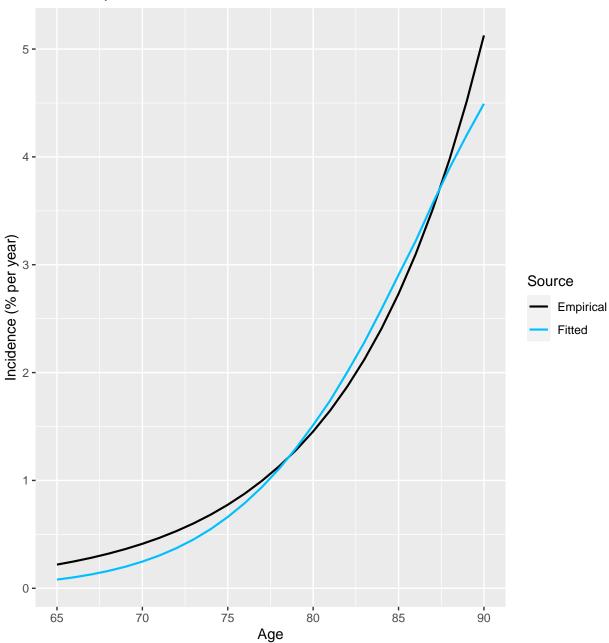
2.4 Incidence of AD

Female and male incidence rates as compared with the empirically estimated rate. Note that the empirical rate is the same in each plot, as it is sex-agnostic.









References

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