

## Mathematical Methods

The purpose of this note is to establish the theoretical basis for extending the work by Ashwood et al. I will first highlight the mathematical model of the hidden Markov model (HMM) used and then briefly cover the reasoning behind using (approximate) leave-one-out cross-validation to obtain the pointwise predictive probability to compare different models as opposed to normalized likelihoods in the case of Ashwood et al.

### The Model

#### Observation Model

We assume that observations are generated from a *Bernoulli*( $p$ ) distribution with the probability  $p$  given by

$$p = \Pr(y_t = 1 \mid z_t = k, u_t, \gamma_k) = \frac{1}{1 + \exp(u_t \gamma_k^\top)}$$

Here,  $y = 1$  indicates a rightward decision and  $y = 0$  indicates a leftward decision,  $z_t$  is the hidden state at time  $t$ ,  $u_t$  represents the vector of predictors at time  $t$  and  $\gamma_k$  is the vector of coefficients for state  $k$ .

Thus,  $p$  is given by a sigmoid function and can be viewed as a logistic regression. The time-varying emission vector  $\tilde{\mathbf{b}}(t)$  reflecting the probability of observing  $y_t$  in state  $j$  is given by

$$b_j(t) = \Pr(y_t \mid z_t = j)$$

Using the probability mass function of the Bernoulli distribution, we can write the observation model as

$$b_j(t) = \begin{cases} \frac{1}{1 + \exp(u_t \gamma_j^\top)} & \text{if } y_t = 1, \\ 1 - \frac{1}{1 + \exp(u_t \gamma_j^\top)} & \text{if } y_t = 0. \end{cases}$$

#### Transition Model

The time-varying transition matrix  $\mathbf{A}(t) \in \mathbb{R}_+^{K \times K}$  is given by

$$a_{i,j}(t) = \Pr(z_t = j \mid z_{t-1} = i, \vec{u}_t) = \text{softmax}(d_{i,j} + u_t \beta_i^\top) = \frac{\exp(d_{i,j} + u_t \beta_i^\top)}{\sum_{k=1}^K \exp(d_{i,k} + u_t \beta_i^\top)}$$

In other words, for each of the  $K$  states, a multinomial logistic regression is used to predict the probability of the next possible  $K$  states. Importantly, while we

allow the intercept to vary between states, the coefficients are fixed and only depend on the previous state ( $z_{t-1} = i$ ).

## Fitting the Model

### The Forward Algorithm

The forward algorithm is used to iteratively compute the probability of an observation sequence given the initial state probabilities  $\pi$ , the transition matrix  $\mathbf{A}(t)$  and the emission vector  $\tilde{\mathbf{b}}(t)$ . By integrating (marginalizing) over all possible state sequences, we obtain the probability of the observation sequence up to time  $t$ , **only** conditional on the hidden state at time  $t$  (in theory, one could just naïvely sum over the probabilities under all possible state sequences, but this is intractable due to combinational explosion).

$$\begin{aligned}\alpha_t(i) &= \Pr(y_1 \dots y_t, z_t = i) \\ \alpha_{t+1}(j) &= \sum_{i=1}^K \alpha_t(i) a_{ij} b_j(t+1)\end{aligned}$$

The initial value  $\alpha_1(i)$  is calculated using  $\pi_i$ .

$$\alpha_{t=1}(i) = \pi_i \cdot b_i(1)$$

The sum over all possible end states at  $t$  is sufficient to compute the likelihood of the observation sequence  $y_1, \dots, y_t$ .

$$L(y_1, \dots, y_t) = \sum_{i=1}^K \alpha_t(i)$$

### Point Estimates

The forward algorithm can be combined with the so-called backward algorithm to compute the probability of a particular latent state at time  $t$  (the process is then fittingly termed forward-backward-algorithm). This information can then be used in an iterative procedure known as expectation maximization (EM) to find the maximum likelihood (ML) or maximum a posteriori (MAP) estimate of the observation sequence. EM works by repeatedly finding the most likely state sequence (E-step) and subsequently maximizing the likelihood by adjusting the other parameters (M-step).

Other options include direct Maximal Likelihood estimation a Variational Bayes estimations.

## **Full Bayesian Inference**

### **Markov Chain Monte Carlo Sampling**

Markov Chain Monte Carlo (MCMC) procedures such as Gibbs's sampling or Hamiltonian Monte Carlo (HMC) can be used to sample from the posterior distributions.

- $K$ : Number of different states
- $P$ : Number of predictors for transitions model.
- $M$ : Number of predictors for observation model.