

Notes on Sequence Modelling

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1 Random Sequence Processes

Consider a random process R that generates arbitrary-length sequences of the form $\vec{R} = (R_1, R_2, \dots, R_N)$, where N is a random variable governing the length of a sequence, and R_t is a random variable governing the value at *stage* t of the sequence. This sequence process is graphically depicted in Figure 1.1.

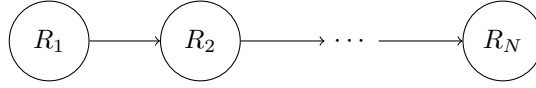


Figure 1.1: A random process R for generating sequences of random length N . The arrows indicate transitions from one stage in the sequence to the next.

We assume that each R_t randomly takes some discrete or continuous value $r_t \in \mathcal{R}$, and hence the probability (or probability density) of observing a particular sequence \vec{r} of length $n = |\vec{r}|$ is given by

$$p(\vec{R} = \vec{r}) = p(N = n) p(R_1 = r_1, \dots, R_n = r_n \mid N = n). \quad (1.1)$$

In practice, this definition presupposes that we know we have observed a *complete* sequence that started at stage 1 and ended at stage n . Suppose instead that the sequence \vec{r} was observed one stage at a time. How do we know if the underlying process has actually terminated, or will instead continue to generate another observed value r_{n+1} ? Similarly, how do we know that the first observed value r_1 was not in fact part of a longer, unobserved sequence of values? We assume that the random process R only ever produces complete sequences, independently of the observation process, which might provide partial or complete sequences of values. Furthermore, if the random process does not signal the start and end of generated sequences, then an observed sequence might actually comprise multiple, contiguously generated subsequences.

In order to handle such difficulties, we consider any arbitrary sequence \vec{r} by default to be *incomplete*, and explicitly denote the corresponding, complete sequence by $\langle \vec{r} \rangle$. We can now introduce the notion of *partially complete* sequences. Thus, a *start sequence* is a generated sequence with an observed (or definite) start (at stage 1) but an unobserved (or indefinite) end, i.e. it might or might not terminate at stage n . This is denoted by $\langle \vec{r}$ if we are truly uncertain as to the termination, or by $\langle \vec{r}]$ if we actually know that the generated sequence does not terminate at stage n . Similarly, an *end sequence* is a generated sequence with an observed end (at stage n) but an unobserved start, i.e. it might or might not have initiated at the observed stage 1. This is denoted by $\vec{r} \rangle$ if we are truly uncertain as to sequence initiation, or by $[\vec{r} \rangle$ if we actually know that the generated sequence was not initiated at stage 1. Clearly, we may also specify the remaining incomplete sequences, namely $[\vec{r}]$, $\vec{r}]$ and $[\vec{r} \rangle$.

Under this augmented notation, knowledge about the start of a sequence can be encapsulated in a random indicator variable ι_{t-1} , which takes on the value 1 if some observed r_t is definitely the first stage in the generated sequence, or the value 0 if it is not. Similarly, the random indicator variable τ_{t+1} takes on the value 1 if r_t is definitely the last stage in the generated sequence, or the value 0 if it is not. In general, these indicators allow us to handle the observation of possibly concatenated, multiple, generated sequences. From now on, however, we shall assume (unless otherwise stated) that we are dealing with a single, contiguous sequence. Thus, notionally, the indicators ι_0 and τ_{n+1} can be thought to correspond to pseudo-stages 0 and $n+1$, such that an arbitrary generated sequence is initiated at stage 0 and terminated at some random stage $N+1$. This augmented random process is depicted in Figure 1.2.

The probability of a given complete sequence $\langle \vec{r} \rangle$ is now defined as

$$p(\langle \vec{r} \rangle) = p(\iota_0 = 1, \tau_1 = 0, R_1 = r_1, \tau_2 = 0, \dots, \tau_n = 0, R_n = r_n, \tau_{n+1} = 1), \quad (1.2)$$

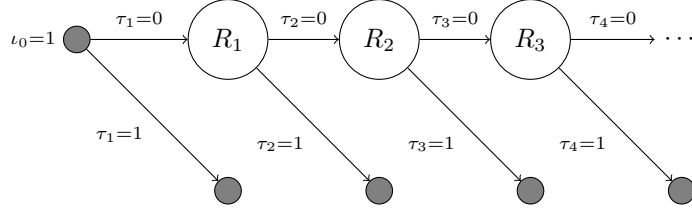


Figure 1.2: A random process for generating complete sequences of random length, with explicit stages for sequence initiation and termination. Multiple arrows exiting from a node indicate different possible (mutually exclusive) stage transition pathways.

such that

$$p(N=n) = p(\iota_0=1, \tau_1=0, \dots, \tau_n=0, \tau_{n+1}=1). \quad (1.3)$$

This has the form of a generalised Bernoulli sequence.

Note that when the context is clear, we may for convenience drop explicit mention of the random variable R_t . Similarly, we may denote $\iota_t = 1$ by ι_t^+ , on the understanding that ι_t^- denotes the negation $\iota_t = 0$. Likewise, we may denote $\tau_t = 1$ by τ_t^+ and $\tau_t = 0$ by τ_t^- . Hence, it is plausible to simplify equation (1.2) as

$$p(\langle \vec{r} \rangle) = p(\iota_0^+, \tau_1^-, r_1, \dots, \tau_n^-, r_n, \tau_{n+1}^+). \quad (1.4)$$

Consequently, we may simplify the explicitly terminated process of Figure 1.2 to more resemble the implicitly terminated process of Figure 1.1; the result is shown in Figure 1.3.

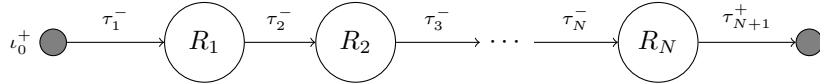


Figure 1.3: A simplified representation of a random process for generating complete sequences of random length N , with explicit stages for sequence initiation and termination, and explicit labelling of non-terminating transitions.

We can now handle both complete and partial sequences by introducing observed indicators $\underline{\iota}$ and $\underline{\tau}$ to correspond to the start-of-sequence and end-of-sequence symbols, respectively. In particular, $\underline{\iota} = 1$ corresponds to ‘ \langle ’ and $\underline{\iota} = 0$ corresponds to ‘ $[\cdot]$ ’; when the start of the sequence is unknown, we let $\underline{\iota} = *$ (see Section 1.1). Likewise, $\underline{\tau} = 1$ corresponds to ‘ \rangle ’, $\underline{\tau} = 0$ corresponds to ‘ $]$ ’, and $\underline{\tau} = *$ corresponds to unknown sequence termination. Hence, in general, we write

$$\begin{aligned} p(\underline{\iota}, \vec{r}, \underline{\tau}) &= p(\iota_0=\underline{\iota}, \tau_1=0, R_1=r_1, \dots, \tau_n=0, R_n=r_n, \tau_{n+1}=\underline{\tau}) \\ &= p(\underline{\iota}, \tau_1^-, r_1, \dots, \tau_n^-, r_n, \underline{\tau}). \end{aligned} \quad (1.5)$$

1.1 Missing Values

The main difference between a complete, generated sequence $\vec{r} = (r_1, \dots, r_n)$ and the observed sequence of values, say¹ $\vec{r} = (\underline{r}_1, \dots, \underline{r}_n)$, is the possibility that some values were unobserved, i.e. arbitrarily *missing* or systematically *hidden*. For convenience, let $\underline{r}_t = *$ denote the case where the observed value of the t -th stage is missing, just like $\underline{\iota} = *$ or $\underline{\tau} = *$ if we do not know whether or not we observed the start or end of the generated sequence, respectively. The ‘ $*$ ’ symbol is just a representational device – its presence has no effect on the computed probabilities, other than to indicate that any associated variable should be marginalised out. Thus, for example:

$$p(\vec{r}) = p(*, \tau_1^-, r_1, \dots, \tau_n^-, r_n, *) = p(\tau_1^-, r_1, \dots, \tau_n^-, r_n). \quad (1.6)$$

In practice, we allow for both observed values and missing values by introducing an indicator function $\delta(\cdot)$, where $\delta(x=y) = 1$ if $x = y$ and $\delta(x=y) = 0$ if $x \neq y$; by definition, we take $\delta(x=*) = 1$. Hence, we

¹We are ignoring the very real problem of aligning the observed values with the generated stages. This difficulty can be partially alleviated under the assumption of stationary distributions, such that each stage behaves like the previous one.

obtain

$$p(\underline{l}, \vec{r}, \underline{\tau}) = \sum_{\iota_0=0}^1 \delta(\iota_0=\underline{l}) \sum_{\tau_{n+1}=0}^1 \delta(\tau_{n+1}=\underline{\tau}) p(\iota_0, \tau_1^-, r_1, \dots, \tau_n^-, r_n, \tau_{n+1}). \quad (1.7)$$

In general, if the domain \mathcal{R} is discrete, then the likelihood of an observed sequence is given by

$$p(\underline{l}, \vec{r}, \underline{\tau}) = \sum_{\iota_0=0}^1 \delta(\iota_0=\underline{l}) \sum_{\tau_{n+1}=0}^1 \delta(\tau_{n+1}=\underline{\tau}) \sum_{r_1 \in \mathcal{R}} \delta(r_1=\underline{r}_1) \cdots \sum_{r_n \in \mathcal{R}} \delta(r_n=\underline{r}_n) p(\iota_0, \tau_1^-, r_1, \dots, \tau_n^-, r_n, \tau_{n+1}). \quad (1.8)$$

Alternatively, if \mathcal{R} is continuous, then the likelihood becomes

$$p(\underline{l}, \vec{r}, \underline{\tau}) = \sum_{\iota_0=0}^1 \delta(\iota_0=\underline{l}) \sum_{\tau_{n+1}=0}^1 \delta(\tau_{n+1}=\underline{\tau}) \int_{\mathcal{R}} \delta(r_1-\underline{r}_1) \cdots \int_{\mathcal{R}} \delta(r_n-\underline{r}_n) p(\iota_0, \tau_1^-, r_1, \dots, \tau_n^-, r_n, \tau_{n+1}) dr_1 dr_2 \cdots dr_n, \quad (1.9)$$

where $\delta(\cdot)$ is now the Dirac delta function, and where, by extension, we define $\delta(x-\ast) = 1$. On the understanding that \sum and $\delta(x=y)$ must be swapped respectively for \int and $\delta(x-y)$ as needed for a continuous or semi-continuous domain, we may henceforth simply utilise the discrete form (1.8) without loss of generality.

1.2 Generic Forward–Backward Algorithm

The likelihood (1.8) of an observed sequence \vec{r} has been written in a computationally inefficient form, but can in practice be efficiently evaluated by nesting the summations, using a modification of the *forward–backward algorithm* to include knowledge of sequence initiation and termination. The precise details of these calculations depend upon the chosen factorisation of the probability model, which is itself a function of the explicit dependencies between various stages in the sequence. Such dependency modelling is dealt with further in Section 2.

Despite not knowing these dependencies in advance, however, the basic form of the forward–backward algorithm can still be formulated. The first requirement is that the sequence process be *causal*, meaning that each stage of a sequence depends only on preceding stages, and never on future stages. This causality allows us to split a generated sequence into two parts at some arbitrary *pivot* stage t , as shown in Figure 1.4. The second requirement is that the dependence on past stages can be limited in scope to some arbitrary *historical* stage s , as also shown.

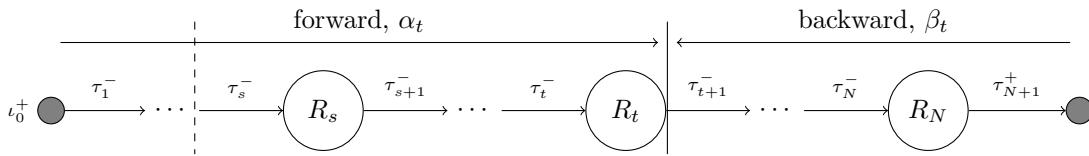


Figure 1.4: *Causality allows the sequence to be partitioned at some pivot stage t , thereby dividing the sequence into past and future stages. Further limitation of past dependencies to some historical stage s defines the active window for one step of the forward–backward algorithm.*

Let us now define the sub-sequence $\vec{r}_{s,t} = (r_s, r_{s+1}, \dots, r_t)$; by definition, $\vec{r}_{s,t} = ()$ if $s > t$. Furthermore, define the concatenation operator ‘ \circ ’, such that $\vec{r}_{s,k} \circ \vec{r}_{k+1,t} = \vec{r}_{s,t}$. Then observe, for a sufficiently long sequence (defined in Section 2), that

$$\begin{aligned} p(\underline{l}, \vec{r}, \underline{\tau}) &= p(\underline{l}, \vec{r}_{1,s-1} \circ \vec{r}_{s,t} \circ \vec{r}_{t+1,n}, \underline{\tau}) \\ &= \sum_{r_s \in \mathcal{R}} \delta(r_s=\underline{r}_s) \cdots \sum_{r_t \in \mathcal{R}} \delta(r_t=\underline{r}_t) p(\underline{l}, \vec{r}_{1,s-1} \circ \vec{r}_{s,t} \circ \vec{r}_{t+1,n}, \underline{\tau}) \\ &= \sum_{r_s \in \mathcal{R}} \delta(r_s=\underline{r}_s) \cdots \sum_{r_t \in \mathcal{R}} \delta(r_t=\underline{r}_t) p(\underline{l}, \vec{r}_{1,s-1} \circ \vec{r}_{s,t}) p(\vec{r}_{t+1,n}, \underline{\tau} \mid \underline{l}, \vec{r}_{1,s-1} \circ \vec{r}_{s,t}) \\ &= \sum_{r_s \in \mathcal{R}} \delta(r_s=\underline{r}_s) \cdots \sum_{r_t \in \mathcal{R}} \delta(r_t=\underline{r}_t) \alpha_t(\vec{r}_{s,t}) \beta_t(\vec{r}_{s,t}), \end{aligned} \quad (1.10)$$

where

$$\alpha_t(\vec{r}_{s,t}) = p(\underline{l}, \vec{r}_{1,s-1} \circ \vec{r}_{s,t}) = p(\underline{l}, \tau_1^-, \underline{r}_1, \dots, \tau_{s-1}^-, \underline{r}_{s-1}, \tau_s^-, r_s, \dots, \tau_t^-, r_t) \quad (1.11)$$

is the forward factor, and

$$\begin{aligned} \beta_t(\vec{r}_{s,t}) &= p(\vec{r}_{t+1,n}, \underline{r} \mid \underline{l}, \vec{r}_{1,s-1} \circ \vec{r}_{s,t}) \\ &= p(\tau_{t+1}^-, \underline{r}_{t+1}, \dots, \tau_n^-, \underline{r}_n, \underline{r} \mid \underline{l}, \tau_1^-, \underline{r}_1, \dots, \tau_{s-1}^-, \underline{r}_{s-1}, \tau_s^-, r_s, \dots, \tau_t^-, r_t) \end{aligned} \quad (1.12)$$

is the backward factor. The entire forward pass of the forward-backward algorithm starts from some initial, historical stage t_0 and progressively computes α_t forward along the sequence for each applicable stage $t_0 \leq t \leq n$. Likewise, the backward pass starts at termination stage, and computes β_t backwards along the sequence for each applicable stage $t_0 \leq t \leq n$. The precise details of these calculations rely upon the nature of the fine-grained dependencies, as discussed in the next section.

2 Markov Sequence Processes

In Section 1, we defined a causal random sequence process R , such that each stage of a sequence, including the termination stage, depends only on the preceding stages. This causal process, depicted in Figure 2.1, is simply the random process from Figure 1.3 with additional, explicit dependencies (in the form of dashed arrows). Hence, under the Markov assumption of conditional independence, the causal sequence process

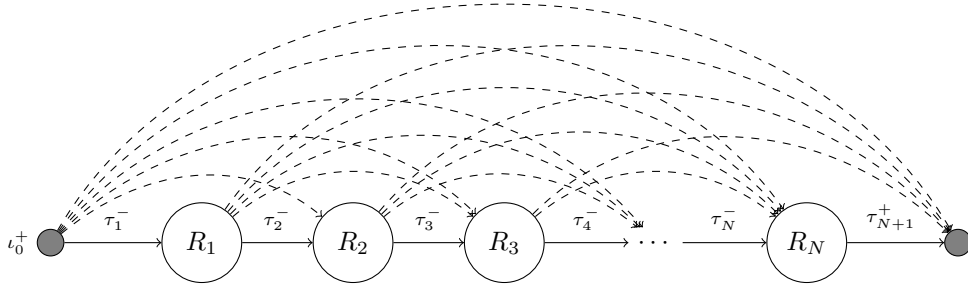


Figure 2.1: A fully-dependent, causal process for generating complete, random sequences of random length N . Solid arrows indicate stage transitions. Both dashed arrows and solid arrows indicate parent-child dependencies, such that each stage is conditionally dependent on the preceding stages.

leads to the fully-dependent conditional model

$$\begin{aligned} p(\iota_0, \vec{r}, \tau_{n+1}) &= p(\iota_0, \tau_1^-, r_1, \tau_2^-, r_2, \dots, \tau_n^-, r_n, \tau_{n+1}) \\ &= p(\iota_0) p(\tau_1^- \mid \iota_0) p(r_1 \mid \iota_0, \tau_1^-) p(\tau_2^- \mid \iota_0, \tau_1^-, r_1) p(r_2 \mid \iota_0, \tau_1^-, r_1, \tau_2^-) \\ &\quad \cdots p(\tau_n^- \mid \iota_0, \dots, \tau_{n-1}^-, r_{n-1}) p(r_n \mid \iota_0, \dots, \tau_n^-) p(\tau_{n+1} \mid \iota_0, \dots, r_n) \\ &= p(\iota_0) \left\{ \prod_{t=1}^n p(\tau_t^-, r_t \mid \iota_0, \vec{r}_{1,t-1}^-, \vec{r}_{1,t-1}) \right\} p(\tau_{n+1} \mid \iota_0, \vec{r}_{1,n}^-, \vec{r}_{1,n}). \end{aligned} \quad (2.1)$$

In practice, this fully-dependent model is considerably simplified by dropping some or even most of the explicit (dashed) dependencies. For example, one might limit the conditionality on past stages to a maximum of m dependencies. This leads to the so-called m -th order Markov model, shown in Figure 2.2. The corresponding likelihood model is given by

$$\begin{aligned} p(\iota_0, \vec{r}, \tau_{n+1}) &= p(\iota_0) \left\{ \prod_{t=1}^m p(\tau_t^-, r_t \mid \iota_0, \vec{r}_{1,t-1}^-, \vec{r}_{1,t-1}) \right\} \\ &\quad \left\{ \prod_{t=m+1}^n p(\tau_t^-, r_t \mid \vec{r}_{t-m,t-1}^-, \vec{r}_{t-m,t-1}) \right\} p(\tau_{n+1} \mid \vec{r}_{n-m+1,n}^-, \vec{r}_{n-m+1,n}), \end{aligned} \quad (2.2)$$

for $n \geq m$.

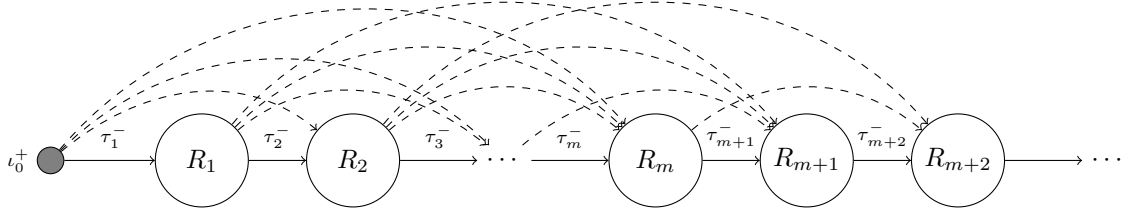


Figure 2.2: An m -th order Markov sequence process of arbitrary length (here $n \geq m + 2$).

An example from the realm of natural language understanding is the lexicographical analysis of the character sequences of words using bigrams (pairs of adjacent characters, corresponding to $m = 1$), and trigrams (triples of adjacent characters, corresponding to $m = 2$), etc. The second-order Markov sequence process, for example, is depicted in Figure 2.3.

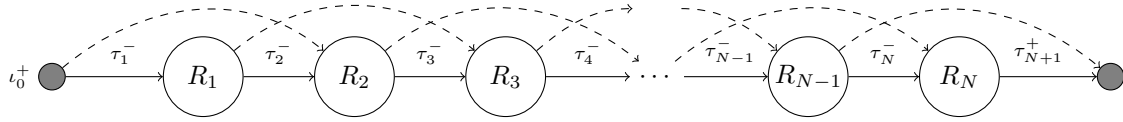


Figure 2.3: A second-order Markov sequence process of random length N .

In the special case of $m = 1$, the first-order Markov model, depicted in Figure 1.3, takes on the especially-simple conditional form of

$$p(\iota_0, \vec{r}, \tau_{n+1}) = p(\iota_0) p(\tau_1^-, r_1 | \iota_0) \left\{ \prod_{t=2}^n p(\tau_t^-, r_t | \tau_{t-1}^-, r_{t-1}) \right\} p(\tau_{n+1} | \tau_n^-, r_n), \quad (2.3)$$

for $n > 0$.

2.1 Markov Forward–Backward Algorithm

The basic description of the generic forward–backward algorithm in Section 1.2 can now be refined under the restriction of the causal sequence process to an m -th order Markov process. Specifically, for any stage $1 \leq t \leq n$, we take the limiting historical stage to be $s = \max(1, t - m + 1)$. Then, for a sufficiently long sequence (i.e. $n \geq m$), the forward factor (1.11) may be computed for stages $t = m, m + 1, \dots, n$ via

$$\begin{aligned} \alpha_t(\vec{r}_{t-m+1,t}) &= p(\underline{\iota}, \tau_1^-, \underline{r}_1, \dots, \tau_{t-m}^-, \underline{r}_{t-m}, \tau_{t-m+1}^-, \underline{r}_{t-m+1}, \dots, \tau_t^-, \underline{r}_t) \\ &= \sum_{\iota_0=0}^1 \delta(\iota_0 = \underline{\iota}) \sum_{r_1 \in \mathcal{R}} \delta(r_1 = \underline{r}_1) \cdots \sum_{r_{t-m} \in \mathcal{R}} \delta(r_{t-m} = \underline{r}_{t-m}) \\ &\quad p(\iota_0) \left\{ \prod_{i=1}^m p(\tau_i^-, r_i | \iota_0, \vec{\tau}_{1,i-1}^-, \vec{r}_{1,i-1}) \right\} \left\{ \prod_{i=m+1}^t p(\tau_i^-, r_i | \vec{\tau}_{i-m,i-1}^-, \vec{r}_{i-m,i-1}) \right\}, \end{aligned} \quad (2.4)$$

from equation (2.2). Furthermore, if $m \leq t < n$ then we may simplify the forward pass by observing that

$$\begin{aligned} \alpha_{t+1}(\vec{r}_{t-m+2,t+1}) &= p(\underline{\iota}, \tau_1^-, \underline{r}_1, \dots, \tau_{t-m+1}^-, \underline{r}_{t-m+1}, \tau_{t-m+2}^-, \underline{r}_{t-m+2}, \dots, \tau_{t+1}^-, \underline{r}_{t+1}) \\ &= \sum_{\iota_0=0}^1 \delta(\iota_0 = \underline{\iota}) \sum_{r_1 \in \mathcal{R}} \delta(r_1 = \underline{r}_1) \cdots \sum_{r_{t-m+1} \in \mathcal{R}} \delta(r_{t-m+1} = \underline{r}_{t-m+1}) \\ &\quad p(\iota_0) \left\{ \prod_{i=1}^m p(\tau_i^-, r_i | \iota_0, \vec{\tau}_{1,i-1}^-, \vec{r}_{1,i-1}) \right\} \left\{ \prod_{i=m+1}^{t+1} p(\tau_i^-, r_i | \vec{\tau}_{i-m,i-1}^-, \vec{r}_{i-m,i-1}) \right\} \\ &= \sum_{r_{t-m+1} \in \mathcal{R}} \delta(r_{t-m+1} = \underline{r}_{t-m+1}) \alpha_t(\vec{r}_{t-m+1,t}) p(\tau_{t+1}^-, r_{t+1} | \vec{\tau}_{t-m+1,t}^-, \vec{r}_{t-m+1,t}). \end{aligned} \quad (2.5)$$

Effectively, this recursive relation comes from moving the size- m active window from stage t to stage $t+1$ and thereby marginalising over the observation \underline{r}_{t-m+1} that has now left the window. Note that for the special case of $m = 1$, this forward pass reduces to

$$\begin{aligned}\alpha_1(r_1) &= \sum_{\iota_0=0}^1 \delta(\iota_0=\underline{\iota}) p(\iota_0) p(\tau_1^-, r_1 | \iota_0), \\ \alpha_t(r_t) &= \sum_{r_{t-1} \in \mathcal{R}} \delta(r_{t-1}=\underline{r}_{t-1}) \alpha_{t-1}(r_{t-1}) p(\tau_t^-, r_t | \tau_{t-1}^-, r_{t-1}) \quad \text{for } t = 2, \dots, n. \end{aligned} \quad (2.6)$$

Similarly, the backward pass is also well-defined for stages $t \geq m$, such that the backward factor (1.12) becomes

$$\begin{aligned}\beta_t(\vec{r}_{t-m+1,t}) &= p(\tau_{t+1}^-, \underline{r}_{t+1}, \dots, \tau_n^-, \underline{r}_n, \underline{\tau} | \tau_{t-m+1}^-, r_{t-m+1}, \dots, \tau_t^-, r_t) \\ &= \sum_{r_{t+1} \in \mathcal{R}} \delta(r_{t+1}=\underline{r}_{t+1}) \cdots \sum_{r_n \in \mathcal{R}} \delta(r_n=\underline{r}_n) \sum_{\tau_{n+1}=0}^1 \delta(\tau_{n+1}=\underline{\tau}_{n+1}) \\ &\quad \left\{ \prod_{i=t+1}^n p(\tau_i^-, r_i | \vec{\tau}_{i-m,i-1}^-, \vec{r}_{i-m,i-1}) \right\} p(\tau_{n+1} | \vec{\tau}_{n-m+1,n}^-, \vec{r}_{n-m+1,n}). \end{aligned} \quad (2.7)$$

Likewise, if $m < t \leq n$ then we may move the window backwards from stage t to stage $t-1$, thereby obtaining the recursive relation

$$\begin{aligned}\beta_{t-1}(\vec{r}_{t-m,t-1}) &= p(\tau_t^-, \underline{r}_t, \dots, \tau_n^-, \underline{r}_n, \underline{\tau} | \tau_{t-m}^-, r_{t-m}, \dots, \tau_{t-1}^-, r_{t-1}) \\ &= \sum_{r_t \in \mathcal{R}} \delta(r_t=\underline{r}_t) \cdots \sum_{r_n \in \mathcal{R}} \delta(r_n=\underline{r}_n) \sum_{\tau_{n+1}=0}^1 \delta(\tau_{n+1}=\underline{\tau}_{n+1}) \\ &\quad \left\{ \prod_{i=t}^n p(\tau_i^-, r_i | \vec{\tau}_{i-m,i-1}^-, \vec{r}_{i-m,i-1}) \right\} p(\tau_{n+1} | \vec{\tau}_{n-m+1,n}^-, \vec{r}_{n-m+1,n}) \\ &= \sum_{r_t \in \mathcal{R}} \delta(r_t=\underline{r}_t) p(\tau_t^-, r_t | \vec{\tau}_{t-m,t-1}^-, \vec{r}_{t-m,t-1}) \beta_t(\vec{r}_{t-m+1,t}). \end{aligned} \quad (2.8)$$

Effectively, moving the size- m active window from stage t to stage $t-1$ allows us to marginalise over the observation \underline{r}_t that has now left the window. For the special case of $m = 1$, this backward pass reduces to

$$\begin{aligned}\beta_n(r_n) &= \sum_{\tau_{n+1}=0}^1 \delta(\tau_{n+1}=\underline{\tau}) p(\tau_{n+1} | \tau_n^-, r_n), \\ \beta_t(r_t) &= \sum_{r_{t+1} \in \mathcal{R}} \delta(r_{t+1}=\underline{r}_{t+1}) p(\tau_{t+1}^-, r_{t+1} | \tau_t^-, r_t) \beta_{t+1}(r_{t+1}) \quad \text{for } t = n-1, \dots, 1. \end{aligned} \quad (2.9)$$

Note that for a short sequence of length $n < m$ we must evaluate the observation likelihood (1.8) directly using equation (2.2). The forward-backward algorithm is of no help in such a case, since the active window overlaps both ends of the sequence, namely $\underline{\iota}$ and $\underline{\tau}$, making the dependencies highly stage specific.

3 Stateful Markov Sequence Processes

Consider the fully-dependent causal process R depicted in Figure 2.1. Suppose now that the random variable R_t at stage t can be decomposed into the tuple $R_t = (S_t, X_t)$, where S_t is a random *state* variable taking values $s_t \in \mathcal{S}$, and X_t is a random *value* variable taking values $x_t \in \mathcal{X}$. We make the common presumption that the stage transitions in the sequence generating process are entirely between states, e.g. from S_{t-1} to S_t . It follows from causation that the value x_t is generated after the state s_t has been determined, i.e. X_t depends upon S_t . Consequently, the fully-dependent stateful sequence process, shown in Figure 3.1, is derived from Figure 2.1 by replacing each node R_t by the pair of nodes S_t and X_t with a dependency from S_t to X_t , such that every *afferent* dependency pointing to R_t becomes two dependencies pointing to S_t and X_t , and every *efferent* dependency pointing from R_t becomes two

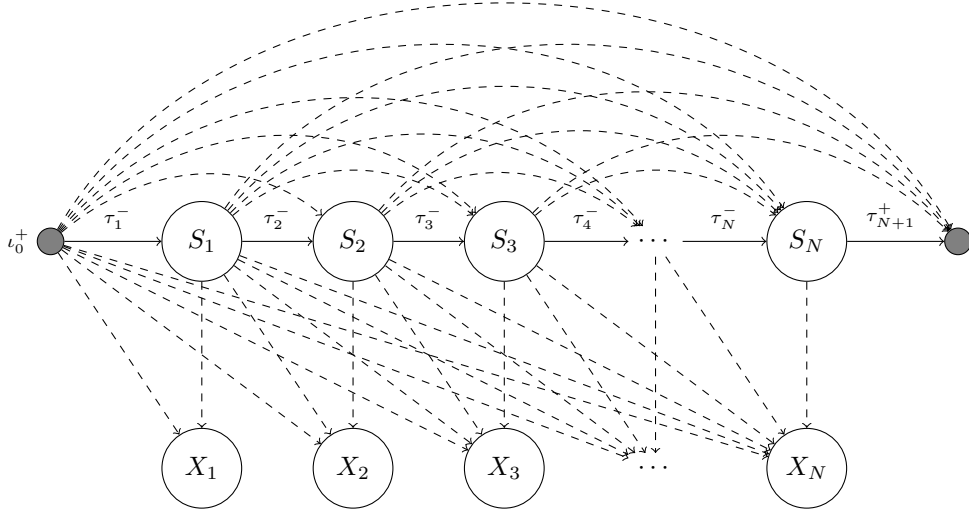


Figure 3.2: A partially-dependent, stateful process of random length N , with strong causal dependencies from states to values.

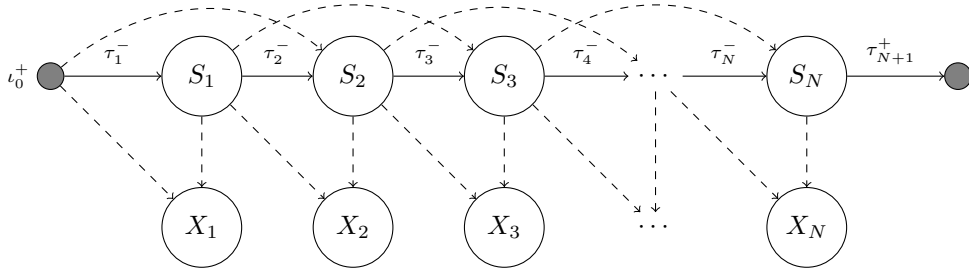


Figure 3.3: An example ($m = 2, \ell = 2$) of an order- (m, ℓ) stateful Markov process of random length N , where state S_t depends on (at most) m previous states, and value X_t depends upon S_t and (at most) $\ell - 1$ previous states.

***** Consider the first-order Markov process R depicted in Figure 1.3. Suppose now that the random variable R_t at stage t can be decomposed into the tuple $R_t = (S_t, X_t)$, where S_t is a random *state* variable taking values $s_t \in \mathcal{S}$, and X_t is a random *value* variable taking values $x_t \in \mathcal{X}$. We make the common presumption that the stage transitions in the sequence generating process are entirely between states, e.g. from S_{t-1} to S_t . It follows from causation that the value is generated after the state has been determined, i.e. X_t depends upon S_t . Keeping to the first-order Markov interpretation of stage-to-stage dependencies leads to the *stateful* process depicted in Figure 3.4, with full cross-dependencies between (S_t, X_t) and (S_{t+1}, X_{t+1}) . Note that the restriction to order $m = 1$ is not really onerous – we may at any time generalise the process by inserting additional (dashed) dependencies (viz Figure 2.1).

For convenience, we may notionally separate the observed states at each stage from the observed values at the corresponding stages, by loosely defining

$$\begin{aligned}
 \vec{r} &= (r_1, r_2, \dots, r_n) \\
 &= ((s_1, x_1), (s_2, x_2), \dots, (s_n, x_n)) \\
 &\equiv (s_1, \dots, s_n) \odot (x_1, \dots, x_n) = (\vec{s}, \vec{x}).
 \end{aligned} \tag{3.4}$$

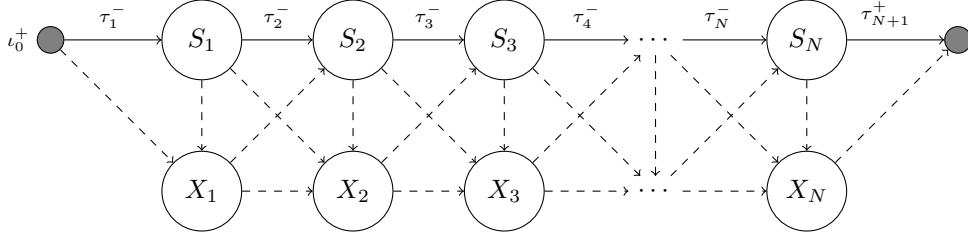


Figure 3.4: A first-order Markov process for generating complete state-value sequences of random length N , with explicit cross-dependencies between adjacent stages.

Consequently, the likelihood of a stateful sequence is now given by

$$\begin{aligned}
 p(\iota_0, \vec{s}, \vec{x}, \tau_{n+1}) &= p(\iota_0, \tau_1^-, s_1, x_1, \dots, \tau_n^-, s_n, x_n, \tau_{n+1}) \\
 &= p(\iota_0) p(\tau_1^-, s_1 | \iota_0) p(x_1 | \iota_0, \tau_1^-, s_1) \\
 &\quad \times \left\{ \prod_{t=2}^n p(\tau_t^-, s_t | \tau_{t-1}^-, s_{t-1}, x_{t-1}) p(x_t | \tau_t^-, s_t, \tau_{t-1}^-, s_{t-1}, x_{t-1}) \right\} \\
 &\quad \times p(\tau_{n+1} | \tau_n^-, s_n, x_n). \tag{3.5}
 \end{aligned}$$

Conditioning the state S_t on both the previous state S_{t-1} and its value X_{t-1} can be useful in some circumstances, e.g. in sequence classification problems. However, due to the increased complexity of such models, it is more usual to further restrict the stateful process by also imposing the first-order Markov assumption at the level of the state-value dependencies themselves. In terms of the process depicted in Figure 3.4, this means retaining only direct node-to-node dependencies, rather than stage-to-stage dependencies. This restricted process is depicted in Figure 3.5. The corresponding sequence model is

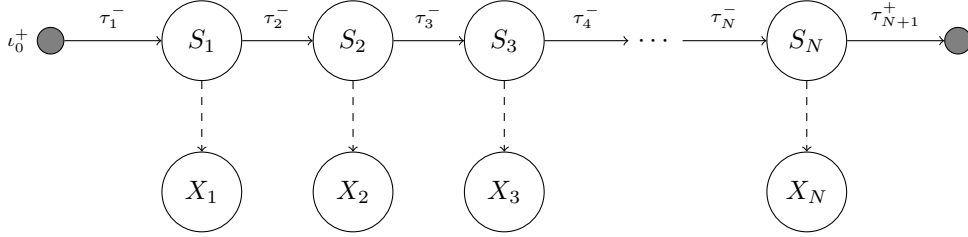


Figure 3.5: A first-order Markov process for generating complete state-value sequences of random length N .

now given by

$$\begin{aligned}
 p(\iota_0, \vec{s}, \vec{x}, \tau_{n+1}) &= p(\iota_0) p(\tau_1^-, s_1 | \iota_0) p(x_1 | \tau_1^-, s_1) \\
 &\quad \times \left\{ \prod_{t=2}^n p(\tau_t^-, s_t | \tau_{t-1}^-, s_{t-1}) p(x_t | \tau_t^-, s_t) \right\} p(\tau_{n+1} | \tau_n^-, s_n). \tag{3.6}
 \end{aligned}$$

3.1 Hidden State Sequences

A special case of the stateful Markov sequence process is the so-called *hidden Markov model* (HMM), where the values of the state sequence \vec{s} are entirely unobserved (and perhaps unobservable). However, more generally we have to consider the possibility that the values of any of the random variables ι_0 , S_t , X_t and τ_{n+1} might or might not have been observed in practice. The extension of equation (1.8) is straightforward for discrete² states and values, namely

$$\begin{aligned}
 p(\underline{\iota}, \underline{\vec{s}}, \underline{\vec{x}}, \underline{\tau}) &= \sum_{\iota_0=0}^1 \delta(\iota_0=\underline{\iota}) \sum_{s_1 \in \mathcal{S}} \delta(s_1=\underline{s}_1) \sum_{x_1 \in \mathcal{X}} \delta(x_1=\underline{x}_1) \cdots \sum_{s_n \in \mathcal{S}} \delta(s_n=\underline{s}_n) \sum_{x_n \in \mathcal{X}} \delta(x_n=\underline{x}_n) \\
 &\quad \sum_{\tau_{n+1}=0}^1 \delta(\tau_{n+1}=\underline{\tau}) p(\iota_0, \tau_1^-, s_1, x_1, \dots, \tau_n^-, s_n, x_n, \tau_{n+1}). \tag{3.7}
 \end{aligned}$$

²Recall from Section 1.1 that the continuous analogue is readily derivable from the discrete model.

This calculation can be simplified further under the first-order state-value model (3.6). For example, the fact that an x_t is unobserved, i.e. $\underline{x}_t = *$, is easily handled, since X_t depends only on S_t in stage t . Specifically, the marginalisation over $x_t \in \mathcal{X}$ affects exactly one term in the model, and consequently we may define $p(* | \tau_t^-, s_t) = p(X_t = * | \tau_t = 0, S_t = s_t) = 1$; hence $p(\underline{x}_t | \tau_t^-, s_t)$ is well defined. Note that this simplification is only applicable when X_t does not depend upon any earlier X_{t-k} ; such higher-order dependencies will require the more general handling of marginalisation above.

Similarly, if the termination marker τ_{n+1} is unobserved, i.e. $\underline{\tau} = *$, then $p(\underline{\tau} | \tau_n^-, s_n)$ is still well defined if we let $p(\tau_{n+1} = * | \tau_n = 0, S_n = s_n) = 1$. However, if the initiation marker ι_0 is unobserved, i.e. $\underline{\iota} = *$, then the marginalisation is more difficult because ι_0 also conditions S_1 via $p(S_1 | \iota_0, \tau_1)$. To allow for this, we define (for order $m = 1$) that

$$p(\underline{\iota}, \tau_1^-, s_1) = \sum_{\iota=0}^1 \delta(\iota = \underline{\iota}) p(\iota_0) p(\tau_1^-, s_1 | \iota_0), \quad (3.8)$$

such that

$$p(\underline{\iota}, \vec{s}, \vec{x}, \underline{\tau}) = p(\underline{\iota}, \tau_1^-, s_1) p(\underline{x}_1 | \tau_1^-, s_1) \left\{ \prod_{t=2}^n p(\tau_t^-, s_t | \tau_{t-1}^-, s_{t-1}) p(\underline{x}_t | s_t) \right\} p(\underline{\tau} | s_n). \quad (3.9)$$

Hence, we obtain

$$p(\underline{\iota}, \vec{s}, \vec{x}, \underline{\tau}) = \sum_{s_1 \in \mathcal{S}} \delta(s_1 = \underline{s}_1) \cdots \sum_{s_n \in \mathcal{S}} \delta(s_n = \underline{s}_n) p(\underline{\iota}, \vec{s}, \vec{x}, \underline{\tau}). \quad (3.10)$$

Note that the conventional HMM formulation is obtained by dropping the $\underline{\iota}$, \vec{s} and $\underline{\tau}$ terms and the $\delta(\cdot)$ indicators.

3.2 Stateful Forward-Backward Algorithm

Under the conventions discussed in the previous section, the forward-backward algorithm for order m stateful Markov sequences takes the form

The observed sequence model (??) can be efficiently evaluated by marginalising over the state of each stage in turn, using a modification of the *forward-backward algorithm* to include knowledge of sequence initiation and termination. The forward pass involves first summing over all terms containing s_1 , then over all remaining terms containing s_2 , and so on up to s_n . This is equivalent to evaluating the reordered model

$$\begin{aligned} p(\underline{\iota}, \vec{s}, \vec{x}, \underline{\tau}) &= \left\{ \sum_{s_n \in \mathcal{S}} \delta(s_n = \underline{s}_n) \cdots \left\{ \sum_{s_2 \in \mathcal{S}} \delta(s_2 = \underline{s}_2) \right. \right. \\ &\quad \left. \left\{ \sum_{s_1 \in \mathcal{S}} \delta(s_1 = \underline{s}_1) p(\underline{\iota}, \tau_1^-, s_1) p(\underline{x}_1 | s_1) p(\tau_2^- | s_1) p(s_2 | \tau_2^-, s_1) \right\} \right. \\ &\quad \left. p(\underline{x}_2 | s_2) p(\tau_3^- | s_2) p(s_3 | \tau_3^-, s_2) \right\} \cdots p(\underline{x}_n | s_n) p(\underline{\tau} | s_n) \Big\}. \end{aligned} \quad (3.11)$$

The forward pass commences with the first stage of the sequence, and includes the inner terms that are dependent on state s_1 , namely

$$\begin{aligned} \alpha_1(s_1) &= p(\underline{\iota}, \tau_1^-, s_1) p(\underline{x}_1 | s_1) \\ &= p(\underline{\iota}, \tau_1^-, s_1, \underline{x}_1) \equiv p(\underline{\iota}, \vec{s}_0 \circ s_1, \vec{x}_1), \end{aligned} \quad (3.12)$$

where the operator ‘ \circ ’ indicates the concatenation of either two elements, two sequences or an element and a sequence, to form a longer sequence. Recall that \vec{s}_0 is an empty sequence.

The next step of the forward pass now moves on to the second stage and includes terms dependent

on state s_2 at that stage:

$$\begin{aligned}
\alpha_2(s_2) &= \left\{ \sum_{s_1 \in \mathcal{S}} \delta(s_1 = \underline{s}_1) \alpha_1(s_1) p(\tau_2^- | s_1) p(s_2 | \tau_2^-, s_1) \right\} p(\underline{x}_2 | s_2) \\
&= \left\{ \sum_{s_1 \in \mathcal{S}} \delta(s_1 = \underline{s}_1) p(\underline{l}, \tau_1^-, s_1, \underline{x}_1) p(\tau_2^- | s_1) p(s_2 | \tau_2^-, s_1) \right\} p(\underline{x}_2 | s_2) \\
&= \left\{ \sum_{s_1 \in \mathcal{S}} \delta(s_1 = \underline{s}_1) p(\underline{l}, \tau_1^-, s_1, \underline{x}_1, \tau_2^-, s_2) \right\} p(\underline{x}_2 | s_2) \\
&= p(\underline{l}, \tau_1^-, \underline{s}_1, \underline{x}_1, \tau_2^-, s_2) p(\underline{x}_2 | s_2) \\
&= p(\underline{l}, \tau_1^-, \underline{s}_1, \underline{x}_1, \tau_2^-, s_2, \underline{x}_2) \equiv p(\underline{l}, \vec{s}_1 \circ s_2, \vec{x}_2). \tag{3.13}
\end{aligned}$$

Recursively, the last step of the forward pass stops at the n -th stage, giving

$$\begin{aligned}
\alpha_n(s_n) &= \left\{ \sum_{s_{n-1} \in \mathcal{S}} \delta(s_{n-1} = \underline{s}_{n-1}) \alpha_{n-1}(s_{n-1}) p(\tau_n^- | s_{n-1}) p(s_n | \tau_n^-, s_{n-1}) \right\} p(\underline{x}_n | s_n) \\
&\equiv p(\underline{l}, \vec{s}_{n-1} \circ s_n, \vec{x}_{n-1}) p(\underline{x}_n | s_n) = p(\underline{l}, \vec{s}_{n-1} \circ s_n, \vec{x}_n). \tag{3.14}
\end{aligned}$$

The remaining terms then give the observed joint likelihood

$$p(\underline{l}, \vec{s}, \vec{x}, \underline{\tau}) = \sum_{s_n \in \mathcal{S}} \delta(s_n = \underline{s}_n) \alpha_n(s_n) p(\underline{\tau} | s_n). \tag{3.15}$$

Conversely, the backward pass involves first summing over all terms containing s_n , then over all remaining terms containing s_{n-1} , and so on down s_1 . This is equivalent to evaluating the reordered model

$$\begin{aligned}
p(\underline{l}, \vec{s}, \vec{x}, \underline{\tau}) &= \left\{ \sum_{s_1 \in \mathcal{S}} \delta(s_1 = \underline{s}_1) p(\underline{l}, \tau_1^-, s_1) p(\underline{x}_1 | s_1) p(\tau_2^- | s_1) \cdots \right. \\
&\quad \left\{ \sum_{s_{n-1} \in \mathcal{S}} \delta(s_{n-1} = \underline{s}_{n-1}) p(s_{n-1} | \tau_{n-1}^-, s_{n-2}) p(\underline{x}_{n-1} | s_{n-1}) p(\tau_n^- | s_{n-1}) \right. \\
&\quad \left. \left. \left\{ \sum_{s_n \in \mathcal{S}} \delta(s_n = \underline{s}_n) p(s_n | \tau_n^-, s_{n-1}) p(\underline{x}_n | s_n) p(\underline{\tau} | s_n) \right\} \right\} \cdots \right\}. \tag{3.16}
\end{aligned}$$

The backward pass commences at the end of the sequence with the termination term that depends on state s_n , namely

$$\beta_n(s_n) = p(\underline{\tau} | s_n). \tag{3.17}$$

The next step in the backward pass then moves backward to the n -th stage and incorporates the terms that depend on state s_{n-1} , including the non-terminating transition from s_{n-1} to s_n

$$\begin{aligned}
\beta_{n-1}(s_{n-1}) &= p(\tau_n^- | s_{n-1}) \left\{ \sum_{s_n \in \mathcal{S}} \delta(s_n = \underline{s}_n) p(s_n | \tau_n^-, s_{n-1}) p(\underline{x}_n | s_n) \beta_n(s_n) \right\} \\
&= p(\tau_n^- | s_{n-1}) \left\{ \sum_{s_n \in \mathcal{S}} \delta(s_n = \underline{s}_n) p(s_n | \tau_n^-, s_{n-1}) p(\underline{x}_n | s_n) p(\underline{\tau} | s_n) \right\} \\
&= p(\tau_n^- | s_{n-1}) \left\{ \sum_{s_n \in \mathcal{S}} \delta(s_n = \underline{s}_n) p(s_n, \underline{x}_n, \underline{\tau} | \tau_n^-, s_{n-1}) \right\} \\
&= p(\tau_n^- | s_{n-1}) p(\underline{s}_n, \underline{x}_n, \underline{\tau} | \tau_n^-, s_{n-1}) \\
&= p(\tau_n^-, \underline{s}_n, \underline{x}_n, \underline{\tau} | s_{n-1}) \equiv p(\circ \vec{s}_n, \vec{x}_n, \underline{\tau} | s_{n-1}). \tag{3.18}
\end{aligned}$$

Observe that we have explicitly included the concatenation operator to indicate the presence of τ_n^- . Symbolically, we may write $p(\vec{r}) = p(\vec{r}_t \circ \vec{r}_{t+1}) = p(\vec{r}_t) p(\vec{r}_{t+1}) = p(\vec{r}_t) p(\circ \vec{r}_{t+1})$.

Recursively, the last step of the backward pass reaches the first stage, giving

$$\begin{aligned}
\beta_1(s_1) &= p(\tau_2^- | s_1) \sum_{s_2 \in \mathcal{S}} \delta(s_2 = \underline{s}_2) p(s_2 | \tau_2^-, s_1) p(\underline{x}_2 | s_2) \beta_2(s_2) \\
&= p(\tau_2^- | s_1) p(\underline{s}_2, \underline{x}_2, \tau_3^-, \dots, \underline{s}_n, \underline{x}_n, \underline{\tau} | \tau_2^-, s_1) \\
&\equiv p(\circ \underline{s}_2, \underline{x}_2, \underline{\tau} | s_1).
\end{aligned} \tag{3.19}$$

The remaining terms then give the observed likelihood

$$p(\underline{\iota}, \vec{s}, \vec{x}, \underline{\tau}) = \sum_{s_1 \in \mathcal{S}} \delta(s_1 = \underline{s}_1) p(\underline{\iota}, \tau_1^-, s_1) \beta_1(s_1). \tag{3.20}$$

In general, we obtain

$$\begin{aligned}
p(\underline{\iota}, \vec{s}, \vec{x}, \underline{\tau}) &= \sum_{s_t \in \mathcal{S}} \delta(s_t = \underline{s}_t) p(\underline{\iota}, \vec{s}_{t-1} \circ s_t \circ \underline{s}_{t+1}, \vec{x}_t \circ \underline{x}_{t+1}, \underline{\tau}) \\
&= \sum_{s_t \in \mathcal{S}} \delta(s_t = \underline{s}_t) p(\underline{\iota}, \vec{s}_{t-1} \circ s_t, \vec{x}_t) p(\circ \underline{s}_{t+1}, \underline{x}_{t+1}, \underline{\tau} | s_t) \\
&= \sum_{s_t \in \mathcal{S}} \delta(s_t = \underline{s}_t) \alpha_t(s_t) \beta_t(s_t),
\end{aligned} \tag{3.21}$$

for all $t = 1, 2, \dots, n$. The case for $t = n$ comes from equations (3.15) and (3.17); alternatively, recall that \tilde{r}_{n+1} is an empty sequence.

4 Discrete-state Sequence Models

Consider the stateful, first-order Markov process depicted by Figure 3.5. Let us now restrict our attention to the class of corresponding sequence models where the state S_t at any stage t may now only take *discrete* values in the set $\mathcal{S} = \{\sigma_1, \sigma_2, \dots, \sigma_S\}$. Hence, the sequence of states may arbitrarily be specified as $\vec{s} = (\sigma_{i_1}, \sigma_{i_2}, \dots, \sigma_{i_n})$, where each $i_t \in \{1, 2, \dots, S\}$. In the event that a particular state S_t is unobserved, we say that the state is *missing* or *hidden*, and denote $i_t = *$ and $s_t = *$. In the situation where all values of \vec{s} are unobserved, the sequence model (3.6) is known as a *hidden-state Markov model* (HMM).

The sequence model (3.6) may now be explicitly conditioned on a general parameter θ that governs the various discrete state distributions. Each term in the model depends directly on the stage index t and indirectly on the state index i_t . Furthermore, each term represents either the initial state, the terminal state, or the non-terminal transitions between states at adjacent stages. Hence, let $\theta = (\Pi, \Gamma, \Omega)$, such that the probability of an arbitrary, observed³ sequence (with no hidden states) is given by

$$p(\underline{\iota}, \vec{s}, \vec{x}, \underline{\tau} | \theta) = \pi_{\underline{\iota}, 1, i_1} o_{1, i_1}(x_1) \left\{ \prod_{t=1}^{n-1} \omega_{0, t, i_t} \Gamma_{t, i_t, i_{t+1}} o_{t+1, i_{t+1}}(x_{t+1}) \right\} \omega_{\underline{\tau}, n, i_n}. \tag{4.1}$$

The initial state S_1 of the sequence at stage $t = 1$ is governed by the parameter $\vec{\pi}$, where

$$\pi_{0, t, i} = p(\iota_{t-1} = 0 | \theta) p(\tau_t = 0 | \iota_{t-1} = 0, \theta) p(S_t = \sigma_i | \iota_{t-1} = 0, \tau_t = 0, \theta), \tag{4.2}$$

$$\pi_{1, t, i} = p(\iota_{t-1} = 1 | \theta) p(\tau_t = 0 | \iota_{t-1} = 1, \theta) p(S_t = \sigma_i | \iota_{t-1} = 1, \tau_t = 0, \theta), \tag{4.3}$$

and

$$\begin{aligned}
\pi_{*, t, i} &= p(\iota_{t-1} = * | \theta) p(\tau_t = 0 | \iota_{t-1} = *, \theta) p(S_t = \sigma_i | \iota_{t-1} = *, \tau_t = 0, \theta) \\
&= p(\tau_t = 0, S_t = \sigma_i | \theta) = \pi_{0, t, i} + \pi_{1, t, i}.
\end{aligned} \tag{4.4}$$

Observe that each state S_t for $t > 1$ is a non-initial state, governed by π_{0, t, i_t} . However, such terms do not explicitly appear in model (4.1), except if $\underline{\iota} \neq 1$, since they are already accounted for by the state transitions. These implicit terms become important when it comes to parameter estimation (see Section 4.3).

³We assume that all observed sequences are non-zero in length, since zero-length sequences are typically unobservable unless the generating process explicitly signals the start and end of each sequence. The modelling of zero-length sequences will require an extra parameter.

The terminal state S_n at stage $t = n$ is likewise governed by the parameter $\vec{\omega}$, where

$$\omega_{0,t,i} = p(\tau_{t+1}=0 \mid S_t=\sigma_i, \theta), \quad (4.5)$$

$$\omega_{1,t,i} = p(\tau_{t+1}=1 \mid S_t=\sigma_i, \theta), \quad (4.6)$$

and

$$\omega_{*,t,i} = p(\tau_{t+1}=* \mid S_t=\sigma_i, \theta) = \omega_{0,t,i} + \omega_{1,t,i} = 1. \quad (4.7)$$

Observe that each state S_t for $t < n$ is a non-terminal state, and is explicitly modelled by the term ω_{0,t,i_t} .

Lastly, the permissible transitions between the states S_t and S_{t+1} of consecutive stages t and $t+1$ are governed by the parameter Γ , where

$$\Gamma_{t,i,j} = p(S_{t+1}=\sigma_j \mid S_t=\sigma_i, \tau_{t+1}=0, \theta). \quad (4.8)$$

Note that the model also includes the likelihood of each observed value x_t at stage t , for $t = 1, 2, \dots, n$. This so-called *data likelihood* is governed by the separate model

$$o_{t,i}(x) = p(X_t=x \mid S_t=\sigma_i, \theta) \quad \forall x \in \mathcal{X}. \quad (4.9)$$

We do not, however, explicitly declare the parameterisation structure of this likelihood model (see Section ?? for a plausible model if X_t takes discrete values). It suffices for our calculations that each $o_{t,i_t}(x_t)$ is available when required.

Finally, note that in the situation where any state in the observed state sequence \vec{s} is hidden, we have to marginalise model (4.1) over each such missing state. Hence, in general, we may define

$$p(\underline{L}, \vec{s}, \vec{x}, \underline{\tau} \mid \theta) = \sum_{i'_1=1}^S \delta(i'_1=i_1) \sum_{i'_2=1}^S \delta(i'_2=i_2) \cdots \sum_{i'_n=1}^S \delta(i'_n=i_n) \left\{ \prod_{t=1}^{n-1} \omega_{0,t,i'_t} \Gamma_{t,i'_t,i'_{t+1}} o_{t+1,i'_{t+1}}(x_{t+1}) \right\} \omega_{\underline{\tau},n,i'_n}, \quad (4.10)$$

where $\delta(\cdot)$ is an indicator function taking the value 1 (or 0) if its argument is true (or false). Note that if S_t is a hidden state, then $i_t = *$ and $\delta(i'_t=*) = 1$ for all $i'_t \in \{1, 2, \dots, S\}$; otherwise, the summation over i'_t collapses to the observed value i_t . The observation likelihood given by model (4.10) can be efficiently computed by an extension of the forward-backward algorithm, described in the next section.

4.1 Modified Forward-Backward Algorithm

The sequence model (4.10) can be efficiently evaluated by marginalising over the state of each stage in turn, using a modification of the *forward-backward algorithm* to include knowledge of sequence initiation and termination. The forward pass involves first summing over all terms containing i'_1 , then over all remaining terms containing i'_2 , and so on up to summing over i'_n . This is equivalent to evaluating the reordered model

$$p(\underline{L}, \vec{s}, \vec{x}, \underline{\tau} \mid \theta) = \left\{ \sum_{i'_n=1}^S \delta(i'_n=i_n) \cdots \left\{ \sum_{i'_2=1}^S \delta(i'_2=i_2) \right. \right. \\ \times \left\{ \sum_{i'_1=1}^S \delta(i'_1=i_1) \pi_{\underline{L},1,i'_1} o_{1,i'_1}(x_1) \omega_{0,1,i'_1} \Gamma_{1,i'_1,i'_2} \right\} \\ \left. \left. o_{2,i'_2}(x_2) \omega_{0,2,i'_2} \Gamma_{2,i'_2,i'_3} \right\} \cdots o_{n,i'_n}(x_n) \omega_{\underline{\tau},n,i'_n} \right\}. \quad (4.11)$$

Conversely, the backward pass reverses the order of evaluation, first summing over all terms containing i'_n , and then over all remaining terms containing i'_{n-1} , and so on down to summing over i'_1 . This is

equivalent to evaluating the reordered model

$$\begin{aligned}
p(\underline{l}, \vec{s}, \vec{x}, \underline{\tau} | \theta) &= \left\{ \sum_{i'_1=1}^S \delta(i'_1 = i_1) \pi_{\underline{l}, 1, i'_1} o_{1, i'_1}(x_1) \omega_{0, 1, i'_1} \cdots \right. \\
&\times \left\{ \sum_{i'_{n-1}=1}^S \delta(i'_{n-1} = i_{n-1}) \Gamma_{n-2, i'_{n-2}, i'_{n-1}} o_{n-1, i'_{n-1}}(x_{n-1}) \omega_{0, n-1, i'_{n-1}} \right. \\
&\times \left. \left. \left\{ \sum_{i'_n=1}^S \delta(i'_n = i_n) \Gamma_{n-1, i'_{n-1}, i'_n} o_{n, i'_n}(x_n) \omega_{\underline{\tau}, n, i'_n} \right\} \cdots \right\} \right\}. \quad (4.12)
\end{aligned}$$

A more efficient mechanism for evaluation comes from making use of the first-order Markov dependencies. Notionally, from the process depicted in Figure 3.5, we may arbitrarily consider the transition from some stage t to stage $t+1$, and partition the sequence into: (i) past values from the initial node up to and including S_t and X_t ; and (ii) future values from S_{t+1} and X_{t+1} up to and including the terminal node. Note that the termination or non-termination of stage t is governed by τ_{t+1} , which is therefore a future value. The Markov dependency then implies that the future values are conditioned only on state S_t via τ_{t+1} and S_{t+1} . Hence, remembering that notionally $s_t = \sigma_{i_t}$, model (4.10) reduces to

$$\begin{aligned}
p(\underline{l}, \vec{s}, \vec{x}, \underline{\tau} | \theta) &= \sum_{i=1}^S p(S_t = \sigma_i, \underline{l}, \vec{s}, \vec{x}, \underline{\tau} | \theta) \\
&= \sum_{i=1}^S \delta(i = i_t) p(\underline{l}, \vec{s}_{t-1} \circ \sigma_i \circ \overleftarrow{s}_{t+1}, \vec{x}, \underline{\tau} | \theta) \\
&= \sum_{i=1}^S \delta(i = i_t) p(\underline{l}, \vec{s}_{t-1} \circ \sigma_i, \vec{x}_t | \theta) p(\downarrow \overleftarrow{s}_{t+1}, \overleftarrow{x}_{t+1}, \underline{\tau} | S_t = \sigma_i, \theta) \\
&= \sum_{i=1}^S \delta(i = i_t) \alpha_{t,i} \beta_{t,i}, \quad (4.13)
\end{aligned}$$

where \circ represents sequence concatenation. The forward step $\alpha_{t,i}$ is defined as

$$\begin{aligned}
\alpha_{t,i} &= p(\underline{l}, \vec{s}_{t-1} \circ \sigma_i, \vec{x}_t | \theta) \\
&= p(\underline{l}, \vec{s}_{t-1} \circ \sigma_i, \vec{x}_{t-1} | \theta) p(X_t = x_t | S_t = \sigma_i, \theta) \\
&= \bar{\alpha}_{t,i} o_{t,i}(x_t), \quad (4.14)
\end{aligned}$$

where $\bar{\alpha}_{t,i}$ is recursively defined as

$$\begin{aligned}
\bar{\alpha}_{t,i} &= p(\underline{l}, \vec{s}_{t-1} \circ \sigma_i, \vec{x}_{t-1} | \theta) \\
&= \sum_{j=1}^S p(S_{t-1} = \sigma_j, \underline{l}, \vec{s}_{t-1} \circ \sigma_i, \vec{x}_{t-1} | \theta) \\
&= \sum_{j=1}^S \delta(j = i_{t-1}) p(\underline{l}, \vec{s}_{t-2} \circ \sigma_j \circ \sigma_i, \vec{x}_{t-1} | \theta) \\
&= \sum_{j=1}^S \delta(j = i_{t-1}) p(\underline{l}, \vec{s}_{t-2} \circ \sigma_j, \vec{x}_{t-1} | \theta) p(\tau_t = 0 | S_{t-1} = \sigma_j, \theta) \\
&\quad \times p(S_t = \sigma_i | \tau_t = 0, S_{t-1} = \sigma_j, \theta) \\
&= \sum_{j=1}^S \delta(j = i_{t-1}) \alpha_{t-1,j} \omega_{0, t-1, j} \Gamma_{t-1, j, i}, \quad (4.15)
\end{aligned}$$

for $t = 2, 3, \dots, n$. The forward pass commences with the first step

$$\begin{aligned}
\alpha_{1,i} &= p(\underline{l}, S_1 = \sigma_i, X_1 = x_1 | \theta) \\
&= p(\iota_0 = \underline{l}) p(\tau_1 = 0 | \iota_0 = \underline{l}, \theta) p(S_1 = \sigma_i | \iota_0 = \underline{l}, \tau_1 = 0, \theta) p(X_1 = x_1 | S_1 = \sigma_i, \theta) \\
&= \pi_{\underline{l}, 1, i} o_{1, i}(x_1). \quad (4.16)
\end{aligned}$$

Hence, observe that α_{2,i'_2} , is just the entire summation over i'_1 from the forward model (4.11). Also note that the standard forward pass derivation commences with the equivalent of $\pi_{*,1,i}$ and does not include the $\delta(\cdot)$ or ω terms.

Conversely to the forward pass, the backward step $\beta_{t,i}$ is defined as

$$\begin{aligned}\beta_{t,i} &= p(\downarrow \tilde{s}_{t+1}, \tilde{x}_{t+1}, \underline{\tau} \mid S_t = \sigma_i, \theta) \\ &= p(\tau_{t+1} = 0 \mid S_t = \sigma_i, \theta) p(\tilde{s}_{t+1}, \tilde{x}_{t+1}, \underline{\tau} \mid \tau_{t+1} = 0, S_t = \sigma_i, \theta) \\ &= \omega_{0,t,i} \bar{\beta}_{t,i},\end{aligned}\tag{4.17}$$

where

$$\begin{aligned}\bar{\beta}_{t,i} &= p(\tilde{s}_{t+1}, \tilde{x}_{t+1}, \underline{\tau} \mid \tau_{t+1} = 0, S_t = \sigma_i, \theta) \\ &= \sum_{j=1}^S p(S_{t+1} = \sigma_j, \tilde{s}_{t+1}, \tilde{x}_{t+1}, \underline{\tau} \mid \tau_{t+1} = 0, S_t = \sigma_i, \theta) \\ &= \sum_{j=1}^S \delta(j = i_{t+1}) p(\sigma_j \circ \tilde{s}_{t+2}, x_{t+1} \circ \tilde{x}_{t+2}, \underline{\tau} \mid \tau_{t+1} = 0, S_t = \sigma_i, \theta) \\ &= \sum_{j=1}^S \delta(j = i_{t+1}) p(S_{t+1} = \sigma_j \mid \tau_{t+1} = 0, S_t = \sigma_i, \theta) p(X_{t+1} = x_{t+1} \mid S_{t+1} = \sigma_j, \theta) \\ &\quad \times p(\downarrow \tilde{s}_{t+2}, \tilde{x}_{t+2}, \underline{\tau} \mid S_{t+1} = \sigma_j, \theta) \\ &= \sum_{j=1}^S \delta(j = i_{t+1}) \Gamma_{t,i,j} o_{t+1,j}(x_{t+1}) \beta_{t+1,j},\end{aligned}\tag{4.18}$$

for $t = n - 1, n - 2, \dots, 1$. The backward pass commences with the first step

$$\beta_{n,i} = p(\tau_{n+1} = \underline{\tau} \mid S_n = \sigma_i, \theta) = \omega_{\underline{\tau},n,i}.\tag{4.19}$$

Observe that $\bar{\beta}_{n-1,i'_{n-1}}$ is just the entire summation over i'_n from the backward model (4.12). Also note that the standard backward pass derivation commences with the equivalent of $\omega_{*,n,i} = 1$, and does not include the $\delta(\cdot)$ or ω terms.

4.2 Posterior Prediction

Given an observed sequence with one or more missing values, it is useful to be able to predict the probable values of the missing variables. For stateful Markov sequences, this typically means predicting the state S_t at some (or each) stage t . Alternatively, one might wish to predict a future value of S_{t+1} or X_{t+1} given a partially observed sequence. The forward-backward algorithm of Section 4.1 enables all of these calculations.

For instance, from equation (4.13), the posterior probabilities of state S_t given an observed sequence are computed as

$$\begin{aligned}\gamma_{t,i} &= p(S_t = \sigma_i \mid \underline{\ell}, \vec{s}, \vec{x}, \underline{\tau}, \theta) \\ &= \frac{p(S_t = \sigma_i, \underline{\ell}, \vec{s}, \vec{x}, \underline{\tau} \mid \theta)}{p(\underline{\ell}, \vec{s}, \vec{x}, \underline{\tau} \mid \theta)} \\ &= \frac{\delta(i = i_t) \alpha_{t,i} \beta_{t,i}}{\sum_{i'=1}^S \delta(i' = i_t) \alpha_{t,i'} \beta_{t,i'}}.\end{aligned}\tag{4.20}$$

Observe that $\gamma_{t,i}$ reduces to $\delta(i = i_t)$ in the special case where $s_t = \sigma_{i_t}$ is known.

Similarly, we may predict the next state S_{n+1} in a given observed sequence of length $n = |\vec{x}|$ via

$$\begin{aligned}p(\downarrow \sigma_i \mid \underline{\ell}, \vec{s}, \vec{x}, \underline{\tau}, \theta) &= p(\tau_{n+1} = 0, S_{n+1} = \sigma_i \mid \underline{\ell}, \vec{s}, \vec{x}, \underline{\tau}, \theta) \\ &= \frac{p(\tau_{n+1} = 0, S_{n+1} = \sigma_i, \underline{\ell}, \vec{s}, \vec{x}, \underline{\tau} \mid \theta)}{p(\underline{\ell}, \vec{s}, \vec{x}, \underline{\tau} \mid \theta)} \\ &= \delta(\underline{\tau} = 0) \frac{p(\underline{\ell}, \vec{s} \circ \sigma_i, \vec{x} \mid \theta)}{p(\underline{\ell}, \vec{s}, \vec{x}, \underline{\tau} \mid \theta)} \\ &= \delta(\underline{\tau} = 0) \frac{\alpha_{n+1,i}}{\sum_{i'=1}^S \delta(i' = i_n) \alpha_{n,i'} \beta_{n,i'}},\end{aligned}\tag{4.21}$$

from equation (4.15). Consequently, we may also predict the future value of X_{t+1} via

$$\begin{aligned}
p(\downarrow x \mid \underline{l}, \vec{s}, \vec{x}, \underline{\tau}, \theta) &= \sum_{i=1}^S p(\downarrow \sigma_i, x \mid \underline{l}, \vec{s}, \vec{x}, \underline{\tau}, \theta) \\
&= \sum_{i=1}^S p(\downarrow \sigma_i \mid \underline{l}, \vec{s}, \vec{x}, \underline{\tau}, \theta) p(X_{n+1} = x \mid S_{n+1} = \sigma_i, \theta) \\
&= \delta(\underline{\tau} = 0) \frac{\sum_{i=1}^S \bar{\alpha}_{n+1,i} o_{n+1,i}(x)}{\sum_{i'=1}^S \delta(i' = i_n) \alpha_{n,i'} \beta_{n,i'}}.
\end{aligned} \tag{4.22}$$

Proceeding to predicting stage transitions, the forward-backward calculations also enable us to compute the posterior probabilities of the joint states of stages t and $t+1$ via

$$\begin{aligned}
\xi_{t,i,j} &= p(S_t = \sigma_i, S_{t+1} = \sigma_j \mid \underline{l}, \vec{s}, \vec{x}, \underline{\tau}, \theta) \\
&= \frac{p(S_t = \sigma_i, S_{t+1} = \sigma_j, \underline{l}, \vec{s}, \vec{x}, \underline{\tau} \mid \theta)}{p(\underline{l}, \vec{s}, \vec{x}, \underline{\tau} \mid \theta)} \\
&= \delta(i = i_t) \delta(j = i_{t+1}) \frac{p(\underline{l}, \vec{s}_{t-1} \circ \sigma_i \circ \sigma_j \circ \vec{s}_{t+2}, \vec{x}, \underline{\tau} \mid \theta)}{p(\underline{l}, \vec{s}, \vec{x}, \underline{\tau} \mid \theta)} \\
&= \delta(i = i_t) \delta(j = i_{t+1}) \frac{\alpha_{t,i} \omega_{0,t,i} \Gamma_{t,i,j} o_{t+1,j}(x_{t+1}) \beta_{t+1,j}}{\sum_{i'=1}^S \delta(i' = i_t) \alpha_{t,i'} \beta_{t,i'}},
\end{aligned} \tag{4.23}$$

since

$$\begin{aligned}
p(\underline{l}, \vec{s}_{t-1} \circ \sigma_i \circ \sigma_j \circ \vec{s}_{t+2}, \vec{x}, \underline{\tau} \mid \theta) &= p(\underline{l}, \vec{s}_{t-1} \circ \sigma_i, \vec{x}_t \mid \theta) p(\downarrow \sigma_j, x_{t+1} \mid S_t = \sigma_i, \theta) \\
&\quad \times p(\downarrow \vec{s}_{t+2}, \vec{x}_{t+2}, \underline{\tau} \mid S_{t+1} = \sigma_j, \theta) \\
&= \alpha_{t,i} \omega_{0,t,i} \Gamma_{t,i,j} o_{t+1,j}(x_{t+1}) \beta_{t+1,j},
\end{aligned} \tag{4.24}$$

from the forward pass (4.14) and the backward pass (4.17). Observe that

$$\begin{aligned}
\gamma_{t,i} &= p(S_t = \sigma_i \mid \underline{l}, \vec{s}, \vec{x}, \underline{\tau}, \theta) \\
&= \sum_{j=1}^S p(S_t = \sigma_i, S_{t+1} = \sigma_j \mid \underline{l}, \vec{s}, \vec{x}, \underline{\tau}, \theta) = \sum_{j=1}^S \xi_{t,i,j},
\end{aligned} \tag{4.25}$$

from equations (4.20) and (4.23).

Finally, the modified forward-backward algorithm also allows us to predict the start and/or end of partially observed sequences. For instance, at the start of a sequence we can predict

$$\begin{aligned}
p(\iota_0 = \underline{l}', S_1 = \sigma_i \mid \underline{l}, \vec{s}, \vec{x}, \underline{\tau}, \theta) &= \frac{p(\iota_0 = \underline{l}', S_1 = \sigma_i, \underline{l}, \vec{s}, \vec{x}, \underline{\tau} \mid \theta)}{p(\underline{l}, \vec{s}, \vec{x}, \underline{\tau} \mid \theta)} \\
&= \delta(\underline{l}' = \underline{l}) \delta(i = i_1) \frac{p(\underline{l}', \sigma_i \circ \vec{s}_2, \vec{x}, \underline{\tau} \mid \theta)}{p(\underline{l}, \vec{s}, \vec{x}, \underline{\tau} \mid \theta)} \\
&= \delta(\underline{l}' = \underline{l}) \delta(i = i_1) \frac{\pi_{\underline{l}',1,i} o_{1,i}(x_1) \beta_{1,i}}{\sum_{i'=1}^S \delta(i' = i_1) \alpha_{1,i'} \beta_{1,i'}} \\
&= \delta(\underline{l}' = \underline{l}) \gamma_{1,i} \frac{\pi_{\underline{l}',1,i}}{\alpha_{1,i}} = \gamma_{1,i} \kappa_{\underline{l}',1,i},
\end{aligned} \tag{4.26}$$

where

$$\kappa_{\underline{l}',1,i} = p(\iota_0 = \underline{l}' \mid S_1 = \sigma_i, \underline{l}, \vec{s}, \vec{x}, \underline{\tau}, \theta) = \begin{cases} \delta(\underline{l}' = \underline{l}) & \text{if } \underline{l} = 0 \text{ or } 1 \\ \frac{\pi_{\underline{l}',1,i}}{\pi_{0,1,i} + \pi_{1,1,i}} & \text{if } \underline{l} = * \end{cases}, \tag{4.27}$$

from equations (4.13), (4.16) and (4.4). It then follows that

$$p(\iota_0 = \underline{l}' \mid \underline{l}, \vec{s}, \vec{x}, \underline{\tau}, \theta) = \sum_{i=1}^S p(\iota_0 = \underline{l}', S_1 = \sigma_i \mid \underline{l}, \vec{s}, \vec{x}, \underline{\tau}, \theta) = \sum_{i=1}^S \gamma_{1,i} \kappa_{\underline{l}',1,i}. \tag{4.28}$$

Likewise, at the end of a sequence we can predict

$$\begin{aligned}
p(\tau_{n+1}=\tau', S_n=\sigma_i | \underline{l}, \vec{s}, \vec{x}, \underline{\tau}, \theta) &= \frac{p(\tau_{n+1}=\tau', S_n=\sigma_i, \underline{l}, \vec{s}, \vec{x}, \underline{\tau} | \theta)}{p(\underline{l}, \vec{s}, \vec{x}, \underline{\tau} | \theta)} \\
&= \delta(\underline{\tau}'=\underline{\tau}) \delta(i=i_n) \frac{p(\underline{l}, \vec{s}_{n-1} \circ \sigma_i, \vec{x}, \underline{\tau}' | \theta)}{p(\underline{l}, \vec{s}, \vec{x}, \underline{\tau} | \theta)} \\
&= \delta(\underline{\tau}'=\underline{\tau}) \delta(i=i_n) \frac{\alpha_{n,i} \omega_{\underline{\tau}',n,i}}{\sum_{i'=1}^S \delta(i'=i_n) \alpha_{n,i'} \beta_{n,i'}} \\
&= \delta(\underline{\tau}'=\underline{\tau}) \gamma_{n,i} \frac{\omega_{\underline{\tau}',n,i}}{\beta_{n,i}} = \gamma_{n,i} \zeta_{\underline{\tau}',n,i}, \tag{4.29}
\end{aligned}$$

where

$$\zeta_{\underline{\tau}',n,i} = p(\tau_{n+1}=\tau' | S_n=\sigma_i, \underline{l}, \vec{s}, \vec{x}, \underline{\tau}, \theta) = \begin{cases} \delta(\underline{\tau}'=\underline{\tau}) & \text{if } \underline{\tau} = 0 \text{ or } 1 \\ \omega_{\underline{\tau}',n,i} & \text{if } \underline{\tau} = * \end{cases}, \tag{4.30}$$

from equations (4.13) and (4.19). It then follows that

$$p(\tau_{n+1}=\tau' | \underline{l}, \vec{s}, \vec{x}, \underline{\tau}, \theta) = \sum_{i=1}^S p(\tau_{n+1}=\tau', S_n=\sigma_i | \underline{l}, \vec{s}, \vec{x}, \underline{\tau}, \theta) = \sum_{i=1}^S \gamma_{n,i} \zeta_{\underline{\tau}',n,i}. \tag{4.31}$$

An example of the use of these posterior predictions is given in Section 4.4, when estimating the model parameters from observations with missing data.

4.3 Posterior Parameter Estimation with Known Data

We desire to estimate the model parameter $\theta = (\Pi, \Gamma, \Omega)$ given an ordered set $\mathbb{V} = \{\vec{v}^{(d)}\}_{d=1}^D$ of observed state and value sequences, where each observation takes the form of $\vec{v}^{(d)} = (\underline{l}^{(d)}, \vec{s}^{(d)}, \vec{x}^{(d)}, \underline{\tau}^{(d)})$. As before, we assume that $\vec{x}^{(d)}$ is a contiguous sequence of observed values with no missing values, whereas each ‘observed’ state s_t might either be known, i.e. $s_t = \sigma_{i_t}$, or missing, i.e. $s_t = *$ and $i_t = *$. Similarly, the sequence initiation and termination markers, $\underline{l}^{(d)}$ and $\underline{\tau}^{(d)}$ respectively, might also be known or unknown. In this section, let us suppose that each $\vec{v}^{(d)}$ is entirely known. The case of hidden data is analysed in the next section.

Due to the typical shortage of observed data, let us additionally assume that the distributions for the sub-parameters are stationary in time; that is, $\Gamma_{t,i,j} \equiv \Gamma_{i,j}$ for any stage t , and likewise $\pi_{\underline{l},t,i} \equiv \omega_{\underline{l},i}$, $\omega_{\underline{\tau},t,i} \equiv \omega_{\underline{\tau},i}$ and $o_{t,i}(x) \equiv o_i(x)$. Then, from equation (4.1), we obtain the likelihood of the d -th observed sequence as

$$p(v^{(d)} | \theta) = \pi_{\underline{l}^{(d)}, i_1^{(d)}} o_{i_1^{(d)}}(x_1^{(d)}) \left\{ \prod_{t=1}^{n^{(d)}-1} \omega_{0, i_t^{(d)}} \Gamma_{i_t^{(d)}, i_{t+1}^{(d)}} o_{i_{t+1}^{(d)}}(x_{t+1}^{(d)}) \right\} \omega_{\underline{\tau}^{(d)}, i_{n^{(d)}}^{(d)}}, \tag{4.32}$$

where $n^{(d)} = |\vec{x}^{(d)}|$, and the log-likelihood as

$$\ell(v^{(d)} | \theta) = \log \pi_{\underline{l}^{(d)}, i_1^{(d)}} + \sum_{t=1}^{n^{(d)}-1} \log \omega_{0, i_t^{(d)}} \Gamma_{i_t^{(d)}, i_{t+1}^{(d)}} + \sum_{t=1}^{n^{(d)}} \log o_{i_t^{(d)}}(x_{i_t^{(d)}}) + \log \omega_{\underline{\tau}^{(d)}, i_{n^{(d)}}^{(d)}}. \tag{4.33}$$

Now, under the assumption that the observed sequences are independent, the log-likelihood of the observed data is given by

$$L(\theta) = \log p(\mathbb{V} | \theta) = \log \prod_{d=1}^D p(v^{(d)} | \theta) = \sum_{d=1}^D \ell(v^{(d)}, \theta). \tag{4.34}$$

Hence, to estimate θ we maximise the log-likelihood subject to the necessary (Lagrangian) constraints

on the sub-parameters. Starting with the state transitions, we maximise

$$F_{\Gamma}(\theta) = \sum_{d=1}^D \sum_{t=1}^{n^{(d)}-1} \log \Gamma_{i_t^{(d)}, i_{t+1}^{(d)}} - \sum_{i=1}^S \lambda_i \left(\sum_{j=1}^S \Gamma_{i,j} - 1 \right) \quad (4.35)$$

$$\begin{aligned} \Rightarrow \frac{\partial F_{\Gamma}(\theta)}{\partial \Gamma_{i,j}} &= \sum_{d=1}^D \sum_{t=1}^{n^{(d)}-1} \delta(i = i_t^{(d)}) \delta(j = i_{t+1}^{(d)}) \frac{1}{\Gamma_{i,j}} - \lambda_i = 0 \text{ when } \theta = \hat{\theta} \\ \Rightarrow \hat{\lambda}_i &= \sum_{j=1}^S \sum_{d=1}^D \sum_{t=1}^{n^{(d)}-1} \delta(i = i_t^{(d)}) \delta(j = i_{t+1}^{(d)}) = \sum_{d=1}^D \sum_{t=1}^{n^{(d)}-1} \delta(i = i_t^{(d)}) \\ \Rightarrow \hat{\Gamma}_{i,j} &= \frac{\sum_{d=1}^D \sum_{t=1}^{n^{(d)}-1} \delta(i = i_t^{(d)}) \delta(j = i_{t+1}^{(d)})}{\sum_{d=1}^D \sum_{t=1}^{n^{(d)}-1} \delta(i = i_t^{(d)})}. \end{aligned} \quad (4.36)$$

Observe that this estimate corresponds to counting all the transitions from state i to state j across all the data, and then normalising these counts by the sum over j .

Similarly, for sequence termination or non-termination, we maximise

$$F_{\Omega}(\theta) = \sum_{d=1}^D \left\{ \sum_{t=1}^{n^{(d)}-1} \log \omega_{0, i_t^{(d)}} + \log \omega_{\tau^{(d)}, i_{n^{(d)}}^{(d)}} \right\} - \sum_{i=1}^S \lambda_i (\omega_{0,i} + \omega_{1,i} - 1) \quad (4.37)$$

$$\begin{aligned} \Rightarrow \frac{\partial F_{\Omega}(\theta)}{\partial \omega_{0,i}} &= \sum_{d=1}^D \left\{ \sum_{t=1}^{n^{(d)}-1} \frac{\delta(i = i_t^{(d)})}{\omega_{0,i}} + \frac{\delta(\tau^{(d)}=0) \delta(i = i_{n^{(d)}}^{(d)})}{\omega_{0,i}} \right\} - \lambda_i, \\ \frac{\partial F_{\Omega}(\theta)}{\partial \omega_{1,i}} &= \sum_{d=1}^D \left\{ \frac{\delta(\tau^{(d)}=1) \delta(i = i_{n^{(d)}}^{(d)})}{\omega_{1,i}} \right\} - \lambda_i. \end{aligned} \quad (4.38)$$

Hence, by multiplying the two derivatives by $\omega_{0,i}$ and $\omega_{1,i}$, respectively, adding the terms and setting the result to zero, we obtain

$$\begin{aligned} \hat{\lambda}_i &= \sum_{d=1}^D \sum_{t=1}^{n^{(d)}} \delta(i = i_t^{(d)}) \\ \Rightarrow \hat{\omega}_{0,i} &= \frac{\sum_{d=1}^D \left\{ \sum_{t=1}^{n^{(d)}-1} \delta(i = i_t^{(d)}) + \delta(\tau^{(d)}=0) \delta(i = i_{n^{(d)}}^{(d)}) \right\}}{\sum_{d=1}^D \sum_{t=1}^{n^{(d)}} \delta(i = i_t^{(d)})}, \\ \hat{\omega}_{1,i} &= \frac{\sum_{d=1}^D \delta(\tau^{(d)}=1) \delta(i = i_{n^{(d)}}^{(d)})}{\sum_{d=1}^D \sum_{t=1}^{n^{(d)}} \delta(i = i_t^{(d)})}. \end{aligned} \quad (4.39)$$

Observe that this latter estimate corresponds to counting the various terminal states over all observed sequences, and then normalising these counts by the overall count of each state. Also note that we have assumed that $\tau^{(d)}$ is known; unfortunately, these estimates will be inaccurate if $\tau^{(d)}$ is unknown, since they ascribe equal weight to $\tau^{(d)} = 0$ and $\tau^{(d)} = 1$ regardless of $v^{(d)}$. The correct estimates in the case of missing data will be analysed in the next section.

Finally, for sequence initiation or non-initiation, we recall the comment made in Section 4 that each stage transition is both explicitly a non-terminal transition and implicitly a non-initial transtion; that is, each state transition $\Gamma_{t,i,j}$ also implies a sequence non-initiation $\pi_{0,t+1,j}$. Hence, from equation (4.33), we maximise the function

$$F_{\Pi}(\theta) = \sum_{d=1}^D \left\{ \log \pi_{\tau^{(d)}, i_1^{(d)}} + \sum_{t=2}^{n^{(d)}} \log \pi_{0, i_t^{(d)}} \right\} - \lambda \left(\sum_{i=1}^S \{\pi_{0,i} + \pi_{1,i}\} - 1 \right) \quad (4.40)$$

$$\begin{aligned} \Rightarrow \frac{\partial F_{\Pi}(\theta)}{\partial \pi_{0,i}} &= \sum_{d=1}^D \left\{ \frac{\delta(\tau^{(d)}=0) \delta(i_1^{(d)}=i)}{\pi_{0,i}} + \sum_{t=2}^{n^{(d)}} \frac{\delta(i_t^{(d)}=i)}{\pi_{0,i}} \right\} - \lambda, \\ \frac{\partial F_{\Pi}(\theta)}{\partial \pi_{1,i}} &= \sum_{d=1}^D \left\{ \frac{\delta(\tau^{(d)}=1) \delta(i_1^{(d)}=i)}{\pi_{1,i}} \right\} - \lambda. \end{aligned} \quad (4.41)$$

Thus, by multiplying the two derivatives by $\pi_{0,i}$ and $\pi_{1,i}$, respectively, adding and summing the terms over i , and setting the result to zero, we obtain

$$\begin{aligned}
\hat{\lambda} &= \sum_{i=1}^S \sum_{d=1}^D \sum_{t=1}^{n^{(d)}} \delta(i_t^{(d)}=i) = \sum_{d=1}^D n^{(d)} \\
\Rightarrow \hat{\pi}_{0,i} &= \frac{\sum_{d=1}^D \left\{ \delta(\underline{l}^{(d)}=0) \delta(i_1^{(d)}=i) + \sum_{t=2}^{n^{(d)}} \delta(i_t^{(d)}=i) \right\}}{\sum_{d=1}^D n^{(d)}}, \\
\hat{\pi}_{1,i} &= \frac{\sum_{d=1}^D \delta(\underline{l}^{(d)}=1) \delta(i_1^{(d)}=i)}{\sum_{d=1}^D n^{(d)}}. \tag{4.42}
\end{aligned}$$

Observe that this latter estimate corresponds to counting the various initial states over all observed sequences, and then normalising these counts by the overall count of all states. Also note that these estimates are inaccurate if \underline{l} is unknown; the correct estimates are derived in the next section.

4.4 Posterior Parameter Estimation with Missing Data

In contrast to Section 4.3, suppose now that any or all values of $\underline{l}^{(d)}$, $\underline{\tau}^{(d)}$ and $\bar{s}^{(d)}$ may be unknown when observing the d -th sequence $v^{(d)}$. The basic procedure is then to first estimate these missing values from the observed data \mathbb{V} , and then to estimate the most likely model parameter value $\hat{\theta}$ given \mathbb{V} and the missing values. This is the principle of the *expectation-maximisation* (EM) algorithm, which underlies the modified *Baum-Welch* parameter estimation algorithm derived here.

Suppose we let $\mathbb{Z} = \{z^{(d)}\}_{d=1}^D$ denote the ordered set of missing values corresponding to the observed values $\mathbb{V} = \{v^{(d)}\}_{d=1}^D$, where $z^{(d)} = (\underline{l}^{(d)}, \bar{s}^{(d)}, \underline{\tau}^{(d)})$; that is, notionally \mathbb{Z} contains the true (but still unknown) values missing from \mathbb{V} . Hence, we take an expectation of the log-likelihood over all possible values of \mathbb{Z} , namely⁴

$$\begin{aligned}
Q(\theta) &= E_{\mathbb{Z}|\mathbb{V},\theta} [\log p(\mathbb{Z}, \mathbb{V} | \theta)] \\
&= E_{\mathbb{Z}|\mathbb{V},\theta} \left[\sum_{d=1}^D \log p(z^{(d)}, v^{(d)} | \theta) \right] \\
&= \sum_{d=1}^D E_{\mathbb{Z}|\mathbb{V},\theta} \left[\ell(\underline{l}^{(d)}, \bar{s}^{(d)}, \underline{\tau}^{(d)}; \theta) \right] \\
&= \sum_{d=1}^D \sum_{\bar{l}=0}^1 \sum_{\bar{i}_1=1}^S \cdots \sum_{\bar{i}_{n^{(d)}}=1}^S \sum_{\bar{\tau}=0}^1 p(\bar{l}, \bar{s}, \bar{\tau} | \underline{l}^{(d)}, \bar{s}^{(d)}, \underline{\tau}^{(d)}, \theta) \ell(\bar{l}, \bar{s}, \bar{\tau}; \theta) \\
&= \sum_{d=1}^D \sum_{\bar{l}=0}^1 \sum_{\bar{i}_1=1}^S \cdots \sum_{\bar{i}_{n^{(d)}}=1}^S \sum_{\bar{\tau}=0}^1 p(z | v^{(d)}, \theta) \ell(\bar{v}^{(d)}; \theta), \tag{4.43}
\end{aligned}$$

where $z = (\bar{l}, \bar{s}, \bar{\tau})$ and $\bar{v}^{(d)} = (\bar{l}, \bar{s}, \bar{\tau})$. In principle, the optimal parameter value $\hat{\theta}$ is estimated by maximising this expected log-likelihood subject to parameter constraints.

In practice, it is difficult to optimise this nonlinear expression analytically. A feasible alternative is to iteratively apply the EM algorithm:

1. *Expectation step*: Compute the expected log-likelihood conditioned on a known parameter estimate $\hat{\theta}_k$, namely

$$\begin{aligned}
Q(\theta, \hat{\theta}_k) &= E_{\mathbb{Z}|\mathbb{V},\hat{\theta}_k} [\log p(\mathbb{Z}, \mathbb{V} | \theta)] \\
&= \sum_{d=1}^D \sum_{\bar{l}=0}^1 \sum_{\bar{i}_1=1}^S \cdots \sum_{\bar{i}_{n^{(d)}}=1}^S \sum_{\bar{\tau}=0}^1 p(z | v^{(d)}, \hat{\theta}_k) \ell(\bar{v}^{(d)}; \theta). \tag{4.44}
\end{aligned}$$

⁴Other expectations are possible, e.g. over the joint distribution $\mathbb{Z}, \mathbb{V} | \theta$. This latter produces macro-averaged parameter estimates of the form $\sum_{d=1}^D \phi^{(d)} / \sum_{d=1}^D \psi^{(d)}$, whereas the discriminative distribution $\mathbb{Z} | \mathbb{V}, \theta$ often leads to micro-averaged estimates of the form $\sum_{d=1}^D [\phi^{(d)} / \psi^{(d)}] / D$.

2. *Maximisation step:* Obtain the optimal parameter estimate $\hat{\theta}_{k+1}$ that maximises the conditional expected log-likelihood, namely

$$\hat{\theta}_{k+1} = \arg \max_{\theta} Q(\theta, \hat{\theta}_k). \quad (4.45)$$

These two steps are iterated until $\hat{\theta}_k$ has converged to a value $\hat{\theta}^*$ that maximises $Q(\hat{\theta}^*) = Q(\hat{\theta}^*, \hat{\theta}^*)$.

Following the methodology of Section 4.3, we now break the optimisation of $Q(\theta, \hat{\theta})$ down into separate maximisation problems over each sub-parameter. For instance, we iteratively estimate the state transitions Γ by optimising

$$\begin{aligned} Q_{\Gamma}(\theta, \hat{\theta}) &= \sum_{d=1}^D \sum_{\bar{l}=0}^1 \sum_{\bar{i}_1=1}^S \cdots \sum_{\substack{\bar{i}_{n^{(d)}}=1 \\ \bar{\tau}=0}}^S \sum_{\bar{\tau}=0}^1 \sum_{t=1}^{n^{(d)}-1} p(z | v^{(d)}, \hat{\theta}) \log \Gamma_{\bar{i}_t, \bar{i}_{t+1}} \\ &= \sum_{d=1}^D \sum_{t=1}^{n^{(d)}-1} \sum_{i=1}^S \sum_{j=1}^S p(S_t = \sigma_i, S_{t+1} = \sigma_j | v^{(d)}, \hat{\theta}) \log \Gamma_{i,j} \\ &= \sum_{d=1}^D \sum_{t=1}^{n^{(d)}-1} \sum_{i=1}^S \sum_{j=1}^S \hat{\xi}_{t,i,j}^{(d)} \log \Gamma_{i,j}, \end{aligned} \quad (4.46)$$

subject to the appropriate constraints. Note that use has been made of equation (4.23). Hence, borrowing the Lagrangian constraints from equation (4.35), we estimate the value $\hat{\Gamma}'$ that maximises

$$\begin{aligned} F_{\Gamma}(\theta, \hat{\theta}) &= \sum_{d=1}^D \sum_{t=1}^{n^{(d)}-1} \sum_{i=1}^S \sum_{j=1}^S \hat{\xi}_{t,i,j}^{(d)} \log \Gamma_{i,j} - \sum_{i=1}^S \lambda_i \left(\sum_{j=1}^S \Gamma_{i,j} - 1 \right) \\ \Rightarrow \frac{\partial F_{\Gamma}(\theta, \hat{\theta})}{\partial \Gamma_{i,j}} &= \sum_{d=1}^D \sum_{t=1}^{n^{(d)}-1} \frac{\hat{\xi}_{t,i,j}^{(d)}}{\Gamma_{i,j}} - \lambda_i = 0 \text{ when } \theta = \hat{\theta}' \\ \Rightarrow \hat{\lambda}'_i &= \sum_{d=1}^D \sum_{t=1}^{n^{(d)}-1} \sum_{j=1}^S \hat{\xi}_{t,i,j}^{(d)} = \sum_{d=1}^D \sum_{t=1}^{n^{(d)}-1} \hat{\gamma}_{t,i} \\ \Rightarrow \hat{\Gamma}'_{i,j} &= \frac{\sum_{d=1}^D \sum_{t=1}^{n^{(d)}-1} \hat{\xi}_{t,i,j}^{(d)}}{\sum_{d=1}^D \sum_{t=1}^{n^{(d)}-1} \hat{\gamma}_{t,i}} \end{aligned} \quad (4.48)$$

from equation (4.25).

Similarly, we iteratively estimate the sequence initiation distributions $\pi_{0,i}$ and $\pi_{1,i}$ by optimising

$$\begin{aligned} Q_{\Pi}(\theta, \hat{\theta}) &= \sum_{d=1}^D \sum_{\bar{l}=0}^1 \sum_{\bar{i}_1=1}^S \cdots \sum_{\substack{\bar{i}_{n^{(d)}}=1 \\ \bar{\tau}=0}}^S \sum_{\bar{\tau}=0}^1 p(z | v^{(d)}, \hat{\theta}) \left\{ \log \pi_{\bar{l}, \bar{i}_1} + \sum_{t=2}^{n^{(d)}} \log \pi_{0, \bar{i}_t} \right\} \\ &= \sum_{d=1}^D \left\{ \sum_{\bar{l}=0}^1 \sum_{\bar{i}_1=1}^S p(\iota_0 = \bar{l}, S_1 = \sigma_{\bar{i}_1} | v^{(d)}, \hat{\theta}) \log \pi_{\bar{l}, \bar{i}_1} \right. \\ &\quad \left. + \sum_{t=2}^{n^{(d)}} \sum_{\bar{i}_t=1}^S p(S_t = \sigma_{\bar{i}_t} | v^{(d)}, \hat{\theta}) \log \pi_{0, \bar{i}_t} \right\} \\ &= \sum_{d=1}^D \sum_{i=1}^S \left\{ \sum_{\bar{l}'=0}^1 \hat{\gamma}_{1,i}^{(d)} \hat{\kappa}_{\bar{l}',1,i}^{(d)} \log \pi_{\bar{l}',i} + \sum_{t=2}^{n^{(d)}} \hat{\gamma}_{t,i}^{(d)} \log \pi_{0,i} \right\} \end{aligned} \quad (4.49)$$

subject to the appropriate constraints. Note that we have utilised equations (4.23) and (4.26). Hence,

borrowing the Lagrangian constraint of equation (4.40), we maximise

$$\begin{aligned}
F_{\Pi}(\theta, \hat{\theta}) &= \sum_{d=1}^D \sum_{i=1}^S \left\{ \sum_{\ell'=0}^1 \hat{\gamma}_{1,i}^{(d)} \hat{\kappa}_{\ell',1,i}^{(d)} \log \pi_{\ell',i} + \sum_{t=2}^{n^{(d)}} \hat{\gamma}_{t,i} \log \pi_{0,i} \right\} \\
&\quad - \lambda \left(\sum_{i=1}^S \{ \pi_{0,i} + \pi_{1,i} \} - 1 \right) \\
\Rightarrow \frac{\partial F_{\Pi}(\theta, \hat{\theta})}{\partial \pi_{0,i}} &= \sum_{d=1}^D \left\{ \frac{\hat{\gamma}_{1,i}^{(d)} \hat{\kappa}_{0,1,i}^{(d)}}{\pi_{0,i}} + \sum_{t=2}^{n^{(d)}} \frac{\hat{\gamma}_{t,i}}{\pi_{0,i}} \right\} - \lambda = 0 \text{ when } \theta = \hat{\theta}', \\
\frac{\partial F_{\Pi}(\theta, \hat{\theta})}{\partial \pi_{1,i}} &= \sum_{d=1}^D \frac{\hat{\gamma}_{1,i}^{(d)} \hat{\kappa}_{1,1,i}^{(d)}}{\pi_{1,i}} - \lambda = 0 \text{ when } \theta = \hat{\theta}' \\
\Rightarrow \hat{\lambda}' &= \sum_{d=1}^D \sum_{i=1}^S \left\{ \hat{\gamma}_{1,i}^{(d)} \hat{\kappa}_{0,1,i}^{(d)} + \hat{\gamma}_{1,i}^{(d)} \hat{\kappa}_{1,1,i}^{(d)} + \sum_{t=2}^{n^{(d)}} \hat{\gamma}_{t,i}^{(d)} \right\} = \sum_{d=1}^D \sum_{i=1}^S \sum_{t=1}^{n^{(d)}} \hat{\gamma}_{t,i}^{(d)} = \sum_{d=1}^D n^{(d)} \quad (4.51)
\end{aligned}$$

which leads to

$$\begin{aligned}
\hat{\pi}'_{0,i} &= \frac{\sum_{d=1}^D \left\{ \hat{\gamma}_{1,i}^{(d)} \hat{\kappa}_{0,1,i}^{(d)} + \sum_{t=2}^{n^{(d)}} \hat{\gamma}_{t,i}^{(d)} \right\}}{\sum_{d=1}^D n^{(d)}}, \\
\hat{\pi}'_{1,i} &= \frac{\sum_{d=1}^D \hat{\gamma}_{1,i}^{(d)} \hat{\kappa}_{1,1,i}^{(d)}}{\sum_{d=1}^D n^{(d)}}. \quad (4.52)
\end{aligned}$$

Finally, we iteratively estimate the sequence termination distributions $\omega_{0,i}$ and $\omega_{1,i}$ by optimising

$$\begin{aligned}
Q_{\Omega}(\theta, \hat{\theta}) &= \sum_{d=1}^D \sum_{\bar{\ell}=0}^1 \sum_{\bar{i}_1=1}^S \cdots \sum_{\bar{i}_{n^{(d)}}=1}^S \sum_{\bar{\tau}=0}^1 p(z | v^{(d)}, \hat{\theta}) \left\{ \sum_{t=1}^{n^{(d)}-1} \log \omega_{0,\bar{i}_t} + \log \omega_{\bar{\tau}, \bar{i}_{n^{(d)}}} \right\} \\
&= \sum_{d=1}^D \left\{ \sum_{t=1}^{n^{(d)}-1} \sum_{\bar{i}_t=1}^S p(S_t = \sigma_{\bar{i}_t}^- | v^{(d)}, \hat{\theta}) \log \omega_{0,\bar{i}_t} \right. \\
&\quad \left. + \sum_{\bar{i}_{n^{(d)}}=1}^S \sum_{\bar{\tau}=0}^1 p(\tau_{n^{(d)}+1} = \bar{\tau}, S_{n^{(d)}} = \sigma_{\bar{i}_{n^{(d)}}}^- | v^{(d)}, \hat{\theta}) \log \omega_{\bar{\tau}, \bar{i}_{n^{(d)}}} \right\} \\
&= \sum_{d=1}^D \sum_{i=1}^S \left\{ \sum_{t=1}^{n^{(d)}-1} \hat{\gamma}_{t,i}^{(d)} \log \omega_{0,i} + \sum_{\tau'=0}^1 \hat{\gamma}_{n^{(d)},i}^{(d)} \hat{\zeta}_{\tau',n^{(d)},i}^{(d)} \log \omega_{\tau',i} \right\} \quad (4.53)
\end{aligned}$$

subject to the appropriate constraints. Note that we have utilised equations (4.23) and (4.29). Hence,

borrowing the Lagrangian constraint of equation (4.37), we maximise

$$\begin{aligned}
F_{\Omega}(\theta, \hat{\theta}) &= \sum_{d=1}^D \sum_{i=1}^S \left\{ \sum_{t=1}^{n^{(d)}-1} \hat{\gamma}_{t,i}^{(d)} \log \omega_{0,i} + \sum_{\tau'=0}^1 \hat{\gamma}_{n^{(d)},i}^{(d)} \hat{\zeta}_{\tau',n^{(d)},i}^{(d)} \log \omega_{\tau',i} \right\} \\
&\quad - \sum_{i=1}^S \lambda_i (\omega_{0,i} + \omega_{1,i} - 1) \\
\Rightarrow \frac{\partial F_{\Omega}(\theta, \hat{\theta})}{\partial \omega_{0,i}} &= \sum_{d=1}^D \left\{ \sum_{t=1}^{n^{(d)}-1} \frac{\hat{\gamma}_{t,i}^{(d)}}{\omega_{0,i}} + \frac{\hat{\gamma}_{n^{(d)},i}^{(d)} \hat{\zeta}_{0,n^{(d)},i}^{(d)}}{\omega_{0,i}} \right\} - \lambda_i = 0 \text{ when } \theta = \hat{\theta}, \\
\frac{\partial F_{\Omega}(\theta, \hat{\theta})}{\partial \omega_{1,i}} &= \sum_{d=1}^D \left\{ \frac{\hat{\gamma}_{n^{(d)},i}^{(d)} \hat{\zeta}_{1,n^{(d)},i}^{(d)}}{\omega_{1,i}} \right\} - \lambda_i = 0 \text{ when } \theta = \hat{\theta}' \\
\Rightarrow \hat{\lambda}'_i &= \sum_{d=1}^D \left\{ \sum_{t=1}^{n^{(d)}-1} \hat{\gamma}_{t,i}^{(d)} + \hat{\gamma}_{n^{(d)},i}^{(d)} \hat{\zeta}_{0,n^{(d)},i}^{(d)} + \hat{\gamma}_{n^{(d)},i}^{(d)} \hat{\zeta}_{1,n^{(d)},i}^{(d)} \right\} = \sum_{d=1}^D \sum_{t=1}^{n^{(d)}} \hat{\gamma}_{t,i}^{(d)} \\
\Rightarrow \hat{\omega}'_{0,i} &= \frac{\sum_{d=1}^D \left\{ \sum_{t=1}^{n^{(d)}-1} \hat{\gamma}_{t,i}^{(d)} + \hat{\gamma}_{n^{(d)},i}^{(d)} \hat{\zeta}_{0,n^{(d)},i}^{(d)} \right\}}{\sum_{d=1}^D \sum_{t=1}^{n^{(d)}} \hat{\gamma}_{t,i}^{(d)}}, \\
\hat{\omega}'_{1,i} &= \frac{\sum_{d=1}^D \hat{\gamma}_{n^{(d)},i}^{(d)} \hat{\zeta}_{1,n^{(d)},i}^{(d)}}{\sum_{d=1}^D \sum_{t=1}^{n^{(d)}} \hat{\gamma}_{t,i}^{(d)}} = 1 - \hat{\omega}'_{0,i}.
\end{aligned} \tag{4.54}$$

$$\tag{4.55}$$

Observe in comparison to equation (4.39) for known data that each certainty, represented by a $\delta(\cdot)$ term, has now been replaced by a corresponding posterior probability.