# Advanced Machine Learning

Exam Training
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2017-10-17

### **Hidden Markov Models**

```
library(HMM)
## Hidden variables (true positions)
states <- 1:10
0, 0.5, 0.5, 0, 0, 0, 0, 0, 0, 0,
                          0, 0, 0.5, 0.5, 0, 0, 0, 0, 0, 0,
                          0, 0, 0, 0.5, 0.5, 0, 0, 0, 0, 0,
                          0, 0, 0, 0, 0.5, 0.5, 0, 0, 0, 0,
                          0, 0, 0, 0, 0, 0.5, 0.5, 0, 0, 0,
                          0, 0, 0, 0, 0, 0.5, 0.5, 0, 0,
                          0, 0, 0, 0, 0, 0, 0.5, 0.5, 0,
                          0, 0, 0, 0, 0, 0, 0, 0.5, 0.5,
                          0.5, 0, 0, 0, 0, 0, 0, 0, 0, 0.5),
                        byrow=TRUE, nrow=length(states), ncol=length(states))
## Emission variables (observed positions)
symbols <- 1:10
0.2, 0.2, 0.2, 0.2, 0, 0, 0, 0, 0, 0.2,
                        0.2, 0.2, 0.2, 0.2, 0.2, 0, 0, 0, 0, 0,
                        0, 0.2, 0.2, 0.2, 0.2, 0.2, 0, 0, 0, 0,
                        0, 0, 0.2, 0.2, 0.2, 0.2, 0.2, 0, 0, 0,
                        0, 0, 0, 0.2, 0.2, 0.2, 0.2, 0.2, 0, 0,
                        0, 0, 0, 0, 0.2, 0.2, 0.2, 0.2, 0.2, 0,
                        0, 0, 0, 0, 0.2, 0.2, 0.2, 0.2, 0.2,
                        0.2, 0, 0, 0, 0, 0.2, 0.2, 0.2, 0.2,
                        0.2, 0.2, 0, 0, 0, 0, 0.2, 0.2, 0.2,
                       byrow=TRUE, nrow=length(states), ncol=length(states))
start_probs <- rep(1, length(states)) / length(states)</pre>
robot_hmm <- initHMM(states, symbols,</pre>
                   startProbs=start_probs,
                   transProbs=transition_probs,
                   emissionProbs=emission_probs)
set.seed(123)
samples_hmm <- simHMM(robot_hmm, 10)</pre>
compute_filtered_probs <- function(hmm, observations) {</pre>
```

```
log_probs <- HMM::forward(hmm, observations)</pre>
    probs <- prop.table(exp(log_probs), 2)</pre>
    probs
}
get_most_probable_states_by_filtered <- function(hmm, observations, states) {</pre>
    probs <- compute_filtered_probs(hmm, observations)</pre>
    most probable states <- as.numeric(apply(probs, 2, function(x) {</pre>
        states[which.max(x)]
    }))
    most_probable_states
}
compute_smoothed_probs <- function(hmm, observations) {</pre>
    probs <- HMM::posterior(hmm, observations)</pre>
    probs
}
get_most_probable_states_by_smoothed <- function(hmm, observations, states) {</pre>
    probs <- compute_smoothed_probs(hmm, observations)</pre>
    most_probable_states <- as.numeric(apply(probs, 2, function(x) {</pre>
        states[which.max(x)]
    }))
    {\tt most\_probable\_states}
}
get_most_probable_path_by_viterbi <- function(hmm, observations) {</pre>
    most_probable_path <- HMM::viterbi(hmm, observations)</pre>
    most_probable_path
}
get_accuracy_filtered <- function(hmm, samples, states) {</pre>
    predicted_states <- get_most_probable_states_by_filtered(hmm, samples$observation, states)</pre>
    sum(predicted_states == samples$states) / length(predicted_states)
}
get_accuracy_smoothed <- function(hmm, samples, states) {</pre>
    predicted_states <- get_most_probable_states_by_smoothed(hmm, samples$observation, states)</pre>
    sum(predicted_states == samples$states) / length(predicted_states)
}
get_accuracy_viterbi <- function(hmm, samples, states) {</pre>
    predicted_states <- get_most_probable_path_by_viterbi(hmm, samples$observation)</pre>
    sum(predicted_states == samples$states) / length(predicted_states)
}
sample_states <- samples_hmm$states</pre>
sample_obs <- samples_hmm$observation</pre>
most_probable_states_filtered <- get_most_probable_states_by_filtered(robot_hmm, sample_obs, states)</pre>
most_probable_states_smoothed <- get_most_probable_states_by_smoothed(robot_hmm, sample_obs, states)</pre>
most_probable_path <- get_most_probable_path_by_viterbi(robot_hmm, sample_obs)</pre>
```

### Forward-Backward

```
emission_density <- function(x, z) {</pre>
    return(emission_probs[z, x])
}
transition density <- function(z, previous z) {</pre>
    return(transition_probs[previous_z, z])
prior_density <- function(z) {</pre>
    return(1 / length(states))
myforward_backward <- function(x, states, prior_density, transition_density, emission_density) {
    alphas <- myforward(x, states, prior_density, transition_density, emission_density)</pre>
    betas <- mybackward(x, states, prior_density, transition_density, emission_density)</pre>
    list(alpha=alphas, beta=betas)
}
myforward <- function(x, states, prior_density, transition_density, emission_density) {</pre>
    T <- length(x)
    alphas <- matrix(NA, ncol=length(states), nrow=T)</pre>
    for (state in states) {
        alphas[1, state] <- emission_density(x[1], state) * prior_density(state)
    for (t in 2:T) {
        for (state in states) {
            inner_sum <- 0
            for (previous_state in states) {
                 inner_sum <- inner_sum + alphas[t - 1, previous_state] *</pre>
                     transition_density(state, previous_state)
            alphas[t, state] <- emission_density(x[t], state) * inner_sum</pre>
        }
    }
    alphas
}
mybackward <- function(x, states, prior_density, transition_density, emission_density) {</pre>
    T <- length(x)
    betas <- matrix(NA, ncol=length(states), nrow=T)</pre>
    betas[T, ] <- 1
    for (t in (T - 1):1) {
        for (state in states) {
            inner_sum <- 0
```

```
for (next_state in states) {
                inner_sum <- inner_sum + betas[t + 1, next_state] *</pre>
                    emission_density(x[t + 1], next_state) * transition_density(next_state, state)
            }
            betas[t, state] <- inner_sum</pre>
        }
   }
   betas
}
alphabeta <- myforward backward(x=sample obs,
                               states = states,
                               prior_density=prior_density,
                               emission_density = emission_density,
                                transition_density = transition_density)
a <- alphabeta$alpha
b <- alphabeta$beta
filtering <- a / rowSums(a)
filtering
         [,1] [,2] [,3]
                             [,4]
                                       [,5]
                                                [,6]
                                                            [,7]
#>
  [1,]
                     0 0.2000000 0.2000000 0.2000000 0.20000000 0.2000000
                      0 0.1428571 0.2857143 0.2857143 0.28571429 0.0000000
#> \[ \( \( \) 2. \( \) \]
#> [3,]
                 0
                     0 0.1250000 0.3750000 0.5000000 0.00000000 0.0000000
           0
#> [4,]
           0
                0
                     0 0.0625000 0.2500000 0.4375000 0.25000000 0.0000000
#> [5,] 0
                   0 0.0000000 0.0000000 0.4230769 0.42307692 0.1538462
                   0 0.0000000 0.0000000 0.2291667 0.45833333 0.3125000
#> [6,]
         0 0
#> [7,]
         0 0
                    0 0.0000000 0.0000000 0.1145833 0.34375000 0.3854167
#> [8,]
         0 0
                    0 0.0000000 0.0000000 0.0000000 0.24309392 0.3867403
#> [9,]
                     0 0.0000000 0.0000000 0.0000000 0.15714286 0.4071429
           0
#> [10,]
                0
                      0 0.0000000 0.0000000 0.0000000 0.07857143 0.2821429
#>
              [,9]
#> [1,] 0.0000000 0.00000000
#> [2,] 0.0000000 0.00000000
#> [3,] 0.0000000 0.00000000
#> [4,] 0.0000000 0.00000000
#> [5,] 0.0000000 0.00000000
#> [6,] 0.0000000 0.00000000
#> [7,] 0.1562500 0.00000000
#> [8,] 0.2872928 0.08287293
#> [9,] 0.4357143 0.00000000
#> [10,] 0.4214286 0.21785714
t(compute_filtered_probs(robot_hmm, sample_obs))
#>
       states
#> index 1 2 3
                                5
                                          6
     1 0 0 0 0.2000000 0.2000000 0.20000000 0.20000000 0.20000000
      2 0 0 0 0.1428571 0.2857143 0.2857143 0.28571429 0.0000000 0.0000000
#>
      3 0 0 0 0 0.1250000 0.3750000 0.5000000 0.00000000 0.0000000 0.0000000
```

```
4 0 0 0 0.0625000 0.2500000 0.4375000 0.25000000 0.0000000 0.0000000
#>
      5 0 0 0 0.0000000 0.0000000 0.4230769 0.42307692 0.1538462 0.0000000
     6 0 0 0 0.0000000 0.0000000 0.2291667 0.45833333 0.3125000 0.0000000
#>
     7 0 0 0 0 0.0000000 0.0000000 0.1145833 0.34375000 0.3854167 0.1562500
#>
     8 0 0 0 0 0.0000000 0.0000000 0.0000000 0.24309392 0.3867403 0.2872928
#>
#>
     9 0 0 0 0.0000000 0.0000000 0.0000000 0.15714286 0.4071429 0.4357143
#>
      10 0 0 0 0.0000000 0.0000000 0.07857143 0.2821429 0.4214286
#>
#> index
                10
     1 0.00000000
#>
#>
     2 0.00000000
#>
     3 0.00000000
     4 0.00000000
#>
     5 0.00000000
#>
     6 0.00000000
#>
#>
     7 0.00000000
#>
     8 0.08287293
     9 0.00000000
     10 0.21785714
smoothing <- a * b / rowSums(a * b)</pre>
smoothing
#>
         [,1] [,2] [,3]
                             [,4]
                                       [,5]
                                                  [,6]
                                                             [,7]
                                                                        Γ.87
   [1,]
                     0 0.45000000 0.4107143 0.13928571 0.00000000 0.00000000
   [2,]
                     0 0.17857143 0.5428571 0.27857143 0.00000000 0.00000000
#> [3,]
           0
                     0 0.04642857 0.3964286 0.55714286 0.00000000 0.00000000
                0
#> [4,]
                     0 0.00000000 0.1857143 0.60000000 0.21428571 0.00000000
#> [5,]
         0 0
                    0 0.00000000 0.0000000 0.51071429 0.43214286 0.05714286
         0 0
#> [6,]
                     0 0.00000000 0.0000000 0.23571429 0.55000000 0.21428571
#> [7,]
         0 0
                   0 0.00000000 0.0000000 0.07857143 0.47142857 0.39642857
                   0 0.00000000 0.0000000 0.00000000 0.31428571 0.50000000
#>
  [8,]
  [9,]
                   0 0.00000000 0.0000000 0.00000000 0.15714286 0.40714286
#>
           0
                0
#> [10,]
              0
                     0 0.00000000 0.0000000 0.00000000 0.07857143 0.28214286
#>
              [,9]
  [1,] 0.00000000 0.0000000
#> [2,] 0.00000000 0.0000000
#> [3,] 0.00000000 0.0000000
#> [4,] 0.00000000 0.0000000
#> [5,] 0.00000000 0.0000000
#> [6,] 0.00000000 0.0000000
#> [7,] 0.05357143 0.0000000
#> [8,] 0.18571429 0.0000000
#> [9,] 0.43571429 0.0000000
#> [10,] 0.42142857 0.2178571
t(compute_smoothed_probs(robot_hmm, sample_obs))
       states
#> index 1 2 3
                                5
                                           6
     1 0 0 0 0.45000000 0.4107143 0.13928571 0.00000000 0.00000000
     2 0 0 0 0.17857143 0.5428571 0.27857143 0.00000000 0.00000000
     3 0 0 0 0.04642857 0.3964286 0.55714286 0.00000000 0.00000000
#>
     4 0 0 0 0.00000000 0.1857143 0.60000000 0.21428571 0.00000000
      5 0 0 0 0.00000000 0.0000000 0.51071429 0.43214286 0.05714286
```

```
#> 6 0 0 0 0.00000000 0.0000000 0.23571429 0.55000000 0.21428571
#> 7 0 0 0 0.00000000 0.0000000 0.07857143 0.47142857 0.39642857
   8 0 0 0 0.00000000 0.0000000 0.00000000 0.31428571 0.50000000
#> 9 0 0 0 0.00000000 0.00000000 0.15714286 0.40714286
    #>
#>
    states
#> index
               9
#>
   1 0.00000000 0.0000000
#>
   2 0.00000000 0.0000000
   3 0.00000000 0.0000000
#>
#>
   4 0.00000000 0.0000000
#> 5 0.00000000 0.0000000
#> 6 0.00000000 0.0000000
#> 7 0.05357143 0.0000000
#> 8 0.18571429 0.0000000
#> 9 0.43571429 0.0000000
#> 10 0.42142857 0.2178571
```

### Viterbi

```
myviterbi <- function(x, states, prior_density, transition_density, emission_density) {</pre>
    T <- length(x)
    weights <- matrix(NA, ncol=length(states), nrow=T)</pre>
    weights[1, ] <- log(prior_density(states)) + log(emission_density(x[1], states))</pre>
    paths <- matrix(NA, ncol=length(states), nrow=T)</pre>
    for (t in 1:(T-1)) {
        for (state in states) {
            weights[t + 1, state] <- log(emission_density(x[t + 1], state)) +</pre>
                 max(log(transition_density(state, states)) + weights[t, states])
            paths[t + 1, state] <- which.max(log(transition_density(state, states)) +</pre>
                                                weights[t, states])
        }
    }
    path <- rep(NA, T)</pre>
    path[T] <- which.max(weights[T,])</pre>
    for (t in (T - 1):1) {
        path[t] \leftarrow paths[t + 1, path[t + 1]]
    path
}
myviterbi(x=sample_obs,
          states=states,
          prior_density=prior_density,
          emission_density = emission_density,
          transition_density = transition_density)
#> [1] 4 4 4 5 6 6 6 7 7 7
HMM::viterbi(robot_hmm, sample_obs)
#> [1] 4 4 4 5 6 6 6 7 7 7
```

# State Space Models

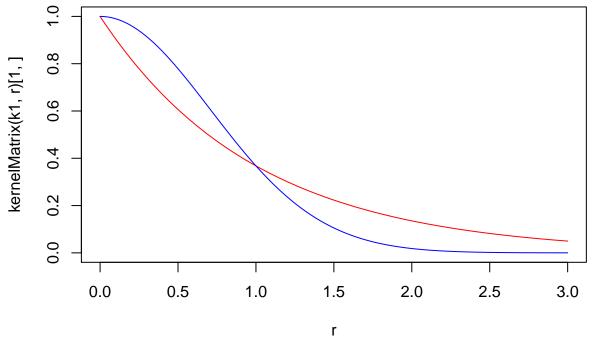
### Kalman Filter

### Weather Forecast System

```
library(HMM)
## States are (1 day ago, 2 days ago)
## 1: (sunny, sunny), 2: (sunny, rainy), 3: (rainy, sunny), 4: (rainy, rainy)
states <- 1:4
transition_probs <- matrix(c(0.75, 0, 0.25, 0,
                             0.5, 0, 0.5, 0,
                             0, 0.5, 0, 0.5,
                             0, 0.25, 0, 0.75),
                           nrow=length(states), ncol=length(states),
                           byrow=TRUE)
emission_probs <- matrix(c(0.9, 0, 0.1, 0,
                           0, 0.9, 0, 0.1,
                           0.1, 0, 0.9, 0,
                           0, 0.1, 0, 0.9),
                         nrow=length(states), ncol=length(states),
                         byrow=TRUE)
robot_hmm <- initHMM(states, states, 1 / length(states), transition_probs, emission_probs)</pre>
set.seed(12345)
simHMM(robot_hmm, 10)
#> $states
#> [1] 4 2 3 4 4 4 4 4 4 2
#> $observation
#> [1] 4 2 3 4 4 4 4 4 4 4
```

# Gaussian Processes

```
library(MASS)
library(kernlab)
## gamma-exponential kernel
k <- function(sigmaf = 1, ell = 1, gamma = 2)</pre>
{
    rval <- function(x, y = NULL)</pre>
         r = sqrt(crossprod(x-y))
         return(sigmaf^2*exp(-(r/ell)^gamma))
    class(rval) <- "kernel"</pre>
    return(rval)
}
k1 \leftarrow k(gamma=1)
k2 \leftarrow k(gamma=2)
r \leftarrow seq(0, 3, by=0.01)
plot(r, kernelMatrix(k1, r)[1, ], type="l", col="red", ylim=c(0, 1))
lines(r, kernelMatrix(k2, r)[1, ], type="l", col="blue")
```



```
x <- seq(0, 2, by=0.01)
n <- length(x)
set.seed(12345)
s11 <- mvrnorm(n=1, mu=rep(0, n), Sigma=kernelMatrix(k1, x))</pre>
```

```
s12 <- mvrnorm(n=1, mu=rep(0, n), Sigma=kernelMatrix(k1, x))
s13 <- mvrnorm(n=1, mu=rep(0, n), Sigma=kernelMatrix(k1, x))

s21 <- mvrnorm(n=1, mu=rep(0, n), Sigma=kernelMatrix(k2, x))
s22 <- mvrnorm(n=1, mu=rep(0, n), Sigma=kernelMatrix(k2, x))
s23 <- mvrnorm(n=1, mu=rep(0, n), Sigma=kernelMatrix(k2, x))

old <- par(mfrow=c(1, 2))
plot(x, s11, type="l", col="red", ylim=c(-5, 5))
lines(x, s12, col="blue")
lines(x, s13, col="green")

plot(x, s21, type="l", col="red", ylim=c(-5, 5))
lines(x, s22, col="blue")
lines(x, s23, col="green")</pre>
```

