Advanced Machine Learning

Lab 4

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Assignment 1

```
sample_transition_component <- function() {</pre>
    sample(1:3, size=1, prob=rep(1, 3) / 3)
}
sample_emission_component <- function() {</pre>
    sample(1:3, size=1, prob=rep(1, 3) / 3)
}
sample_initial_state <- function() {</pre>
    runif(1, 0, 100)
}
sample_transition <- function(z) {</pre>
    component <- sample_transition_component()</pre>
    mu \leftarrow ifelse(component == 1, z, ifelse(component == 2, z + 1, z + 2))
    rnorm(1, mean=mu, sd=1)
}
sample_emission <- function(z, sigma=1) {</pre>
    component <- sample_emission_component()</pre>
    mu \leftarrow ifelse(component == 1, z, ifelse(component == 2, z - 1, z + 1))
    rnorm(1, mean=mu, sd=sigma)
}
generate_states_and_emissions <- function(n, sigma) {</pre>
    states \leftarrow rep(0, n)
    emissions <- rep(0, n)
    initial_state <- sample_initial_state()</pre>
    current_state <- initial_state</pre>
    for (i in 1:n) {
         current_emission <- sample_emission(current_state, sigma)</pre>
         states[i] <- current_state</pre>
         emissions[i] <- current_emission</pre>
         current_state <- sample_transition(current_state)</pre>
    }
    list(states=states, emissions=emissions)
}
```

```
get_emission_density_given_state <- function(emission, state, sigma=1) {</pre>
    (dnorm(emission, mean=state, sd=sigma) +
     dnorm(emission, mean=state - 1, sd=sigma) +
     dnorm(emission, mean=state + 1, sd=sigma)) / 3
}
get_sampling_weights <- function(emission, particles, sigma=1) {</pre>
    unnormalized weights <- sapply(particles, function(state) {
        get_emission_density_given_state(emission, state, sigma)
    unnormalized_weights / sum(unnormalized_weights)
}
particle_filtering <- function(x, nparticles, sigma) {</pre>
    steps <- length(x)
    particles <- matrix(NA, nrow=steps, ncol=nparticles)</pre>
    weights <- matrix(NA, nrow=steps, ncol=nparticles)</pre>
    particles[1, ] <- runif(nparticles, 0, 100)</pre>
    weights[1, ] <- get_sampling_weights(x[1], particles[1, ], sigma=sigma)</pre>
    for (i in 2:steps) {
        predictions <- sample(particles[i - 1, ], size=nparticles,</pre>
                                replace=TRUE, prob=weights[i - 1, ])
        particles[i, ] <- sapply(predictions, sample_transition)</pre>
        weights[i, ] <- get_sampling_weights(x[i], particles[i, ], sigma=sigma)</pre>
    list(particles=particles, weights=weights)
}
get_expected_states <- function(particles, weights) {</pre>
    rowSums(weights * particles)
}
plot_everything <- function(states, emissions, particles, expected) {</pre>
    nparticles <- ncol(particles)</pre>
    old <- par(mfrow=c(2, 1))
    plot(states, type="1", col="blue",
         main="Blue=State, Red=Emission",
         xlab="state",
         ylim=c(min(c(states, emissions)), max(c(states, emissions))))
    lines(emissions, col="red")
    plot(1:nparticles, states, type="l", col="blue",
         main="Blue=State, Orange=Expected",
         xlab="state",
         ylim=c(min(c(states, expected)), max(c(states, expected))))
    lines(1:nparticles, expected, col="orange")
    par(old)
    plot_helper <- function(idx) {</pre>
        p <- particles[idx,]</pre>
```

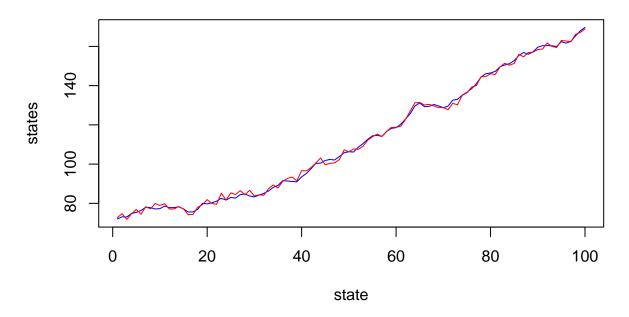
```
ex <- expected[idx]</pre>
        s <- states[idx]
        em <- emissions[idx]</pre>
        hist(p, breaks=20,
             main=paste("Step", idx, sep=" "),
             xlab="state", xlim=c(min(c(p, ex, s, em)), max(c(p, ex, s, em))))
        abline(v=s, col="blue", lwd=2)
        abline(v=em, col="red", lwd=2)
        abline(v=ex, col="orange", lwd=2)
    }
    old <- par(mfrow=c(2, 2), oma=c(0, 0, 2, 0))
    plot_helper(1)
    plot_helper(33)
    plot_helper(66)
    plot_helper(100)
    title(main="Blue=State, Red=Emission, Orange=Expected",outer=T)
    par(old)
}
```

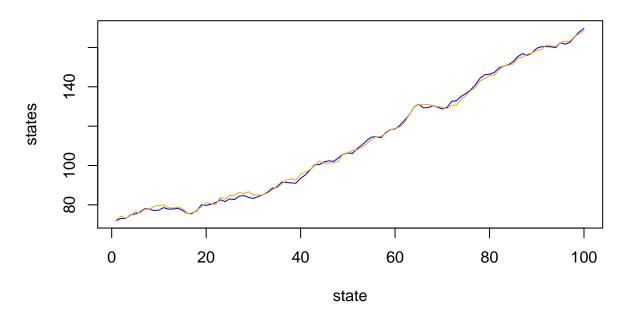
Sigma 1

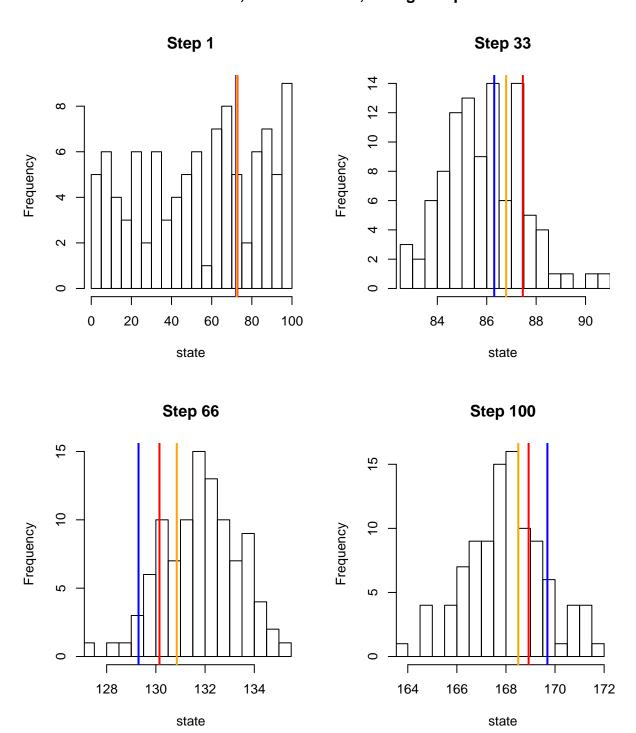
```
sigma <- 1
set.seed(12345)
samples <- generate_states_and_emissions(100, sigma)

nparticles <- 100
particles <- particle_filtering(samples$emissions, nparticles, sigma)

expected <- get_expected_states(particles$particles, particles$weights)</pre>
```





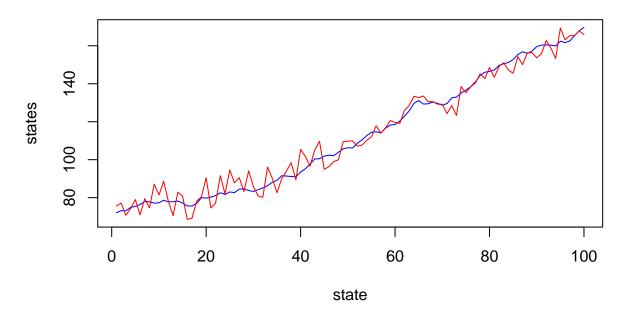


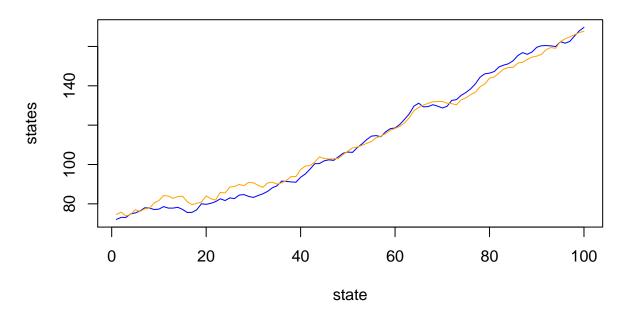
Sigma 5

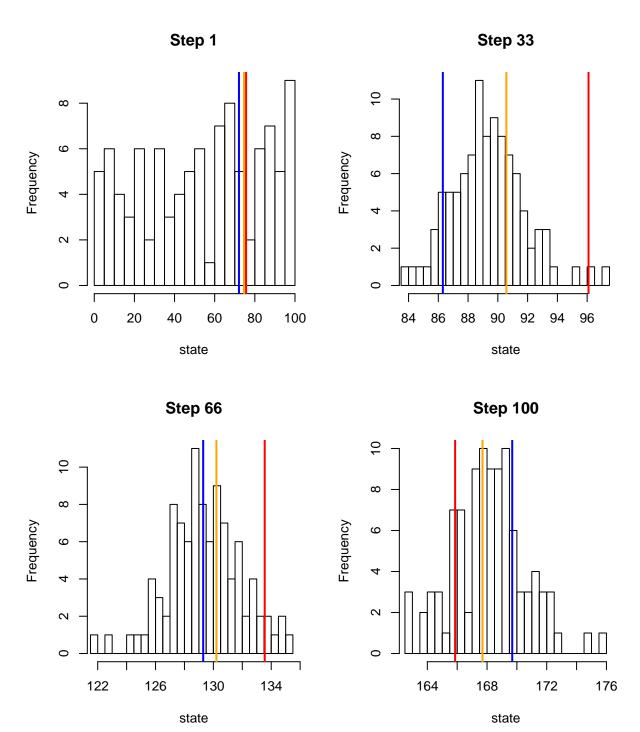
```
sigma <- 5
set.seed(12345)
samples <- generate_states_and_emissions(100, sigma)

nparticles <- 100
particles <- particle_filtering(samples$emissions, nparticles, sigma)

expected <- get_expected_states(particles$particles, particles$weights)</pre>
```







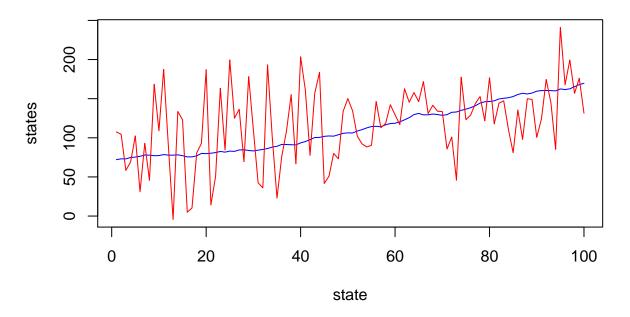
Sigma 50

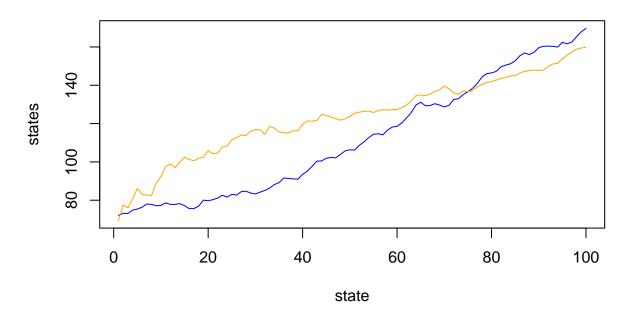
```
sigma <- 50

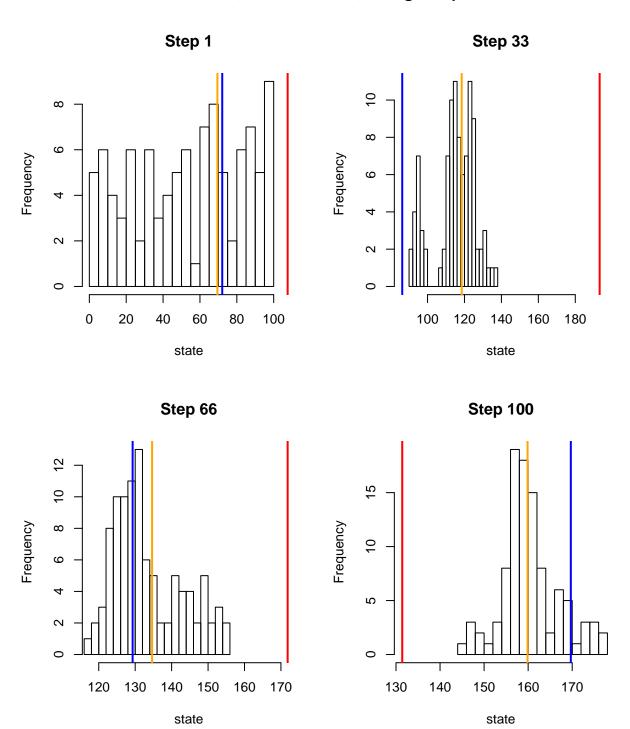
set.seed(12345)
samples <- generate_states_and_emissions(100, sigma)

nparticles <- 100
particles <- particle_filtering(samples$emissions, nparticles, sigma)

expected <- get_expected_states(particles$particles, particles$weights)</pre>
```





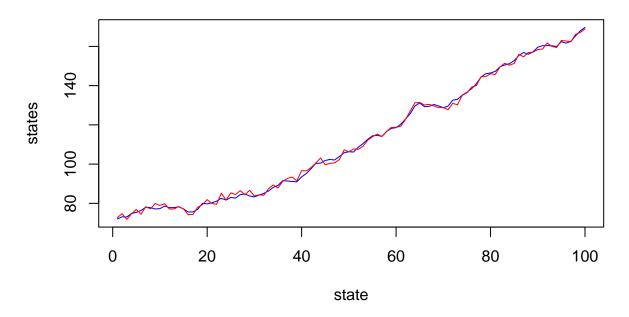


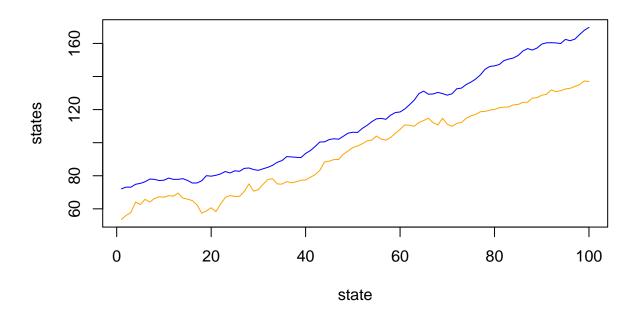
Analysis

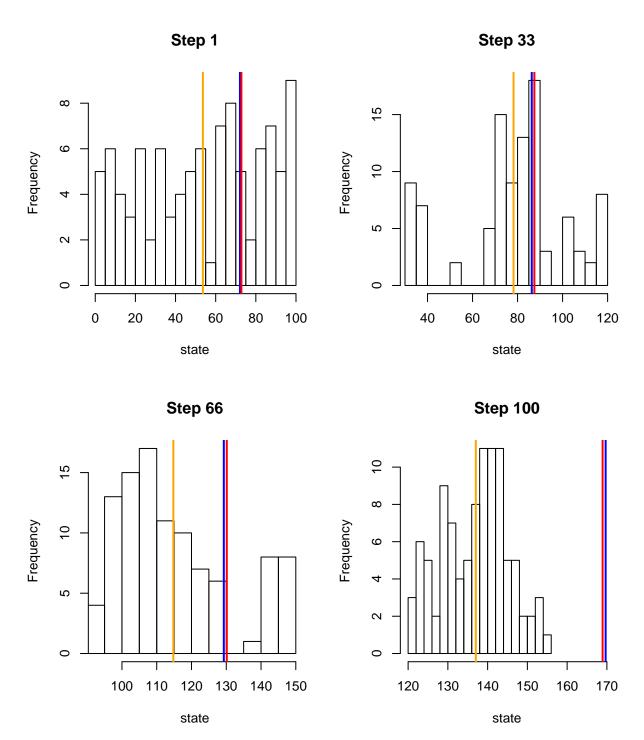
Increasing the variance in the emission model will obviously make the readings less reliable. Since the particles are based on the readings they will also become less accurate. However, the particle estimations are still pretty good which indicate that particle filter can handle high uncertainty in the sensor readings and still be useful.

Sigma 1 without weight correction

```
get_uniform_sampling_weights <- function(emission, particles) {</pre>
    rep(1, length(particles)) / length(particles)
particle_filtering_without_correction <- function(x, nparticles, sigma) {</pre>
    steps <- length(x)</pre>
    particles <- matrix(NA, nrow=steps, ncol=nparticles)</pre>
    weights <- matrix(NA, nrow=steps, ncol=nparticles)</pre>
    particles[1, ] <- runif(nparticles, 0, 100)</pre>
    weights[1, ] <- get_uniform_sampling_weights(x[1], particles[1, ])</pre>
    for (i in 2:steps) {
        predictions <- sample(particles[i - 1, ], size=nparticles,</pre>
                                replace=TRUE, prob=weights[i - 1, ])
        particles[i, ] <- sapply(predictions, sample_transition)</pre>
        weights[i, ] <- get_uniform_sampling_weights(x[i], particles[i, ])</pre>
    }
    list(particles=particles, weights=weights)
sigma <- 1
set.seed(12345)
samples <- generate_states_and_emissions(100, sigma)</pre>
nparticles <- 100
particles <- particle_filtering_without_correction(samples$emissions, nparticles, sigma)</pre>
expected <- get_expected_states(particles$particles, particles$weights)</pre>
```







Without using weight correction we do not use our emission data to learn better estimations of the true states. All particles becomes equally good no matter how likely or unlikely they correspond to the sensor readings.