

Homework 3

Summaries of provided readings

Article 1: “Hidden Markov Model for Stock Trading,” Nguyet Nguyen
(<https://www.mdpi.com/2227-7072/6/2/36/pdf>)

Researchers developed a hidden Markov model (HMM) to forecast closing prices of stocks. Four criteria (AIC, BIC, HQC, and CAIC) were used to select the optimal number of hidden states in the HMM. The close, open, high, and low prices of stocks were used as observations in the HMM. The researchers then tested their model against the historical average return (HAR) model.

In their findings, the researchers noted that four hidden states were optimal for their HMM. Their model predicted S&P 500 monthly prices accurately. It also consistently generated a greater profit than the HAR model when simulating S&P 500 trading.

Article 2: “Gene finding and the Hidden Markov models”
(https://www.cs.us.es/~fran/students/julian/gene_finding/gene_finding.html)

The aim of this project was to identify genes and segmentation in genomes. Preliminarily, researchers looked at prokaryotes, which are predominantly protein-coding, to see where open reading frames occurred. They classified longer open reading frames as more likely to contain genes.

Then they moved to eukaryotes, which contain introns and exons that are not protein coding and complicate genome sequencing. They applied hidden Markov models to determine the segmentation of the genome to then narrow in on segments that could contain genes.

Article clustering kernel on Kaggle: <https://www.kaggle.com/maksimeren/covid-19-literature-clustering>

The author of this project used PCA and LDA to visualize the topic clustering of research articles about COVID-19. After visualizing the distinct clusters, they looked into each of them to assign topics. Topics ranged from surveillance to therapeutics.