CSE 515 Multimedia and Web Databases

Phase #1 (Due Sept 21st 2014, midnight)

Description: In this project, you will experiment with

- epidemic simulation data sets and
- time series.

This project phase will be performed by each group member; but, you will get group grades.

- Download the sample data files from the project directory. See also the attached data description file.
- Task 1: Implement a program which creates an *epidemic words* dictionary: Given a directory, dir, window length, w, a shift length, h, and a resolution, r,
 - 1. for each data file, $f \in dir$, in the given directory, the program
 - (a) normalizes the data file to values between 0.0 and 1.0,
 - (b) quantizes the entries into r levels by dividing the range 0 to 1 into r Gaussian bands (with parameters: mean $\mu = 0$ and standard deviation $\sigma = 0.25$)
 - the length of the i^{th} band is equal to

$$length_i = \frac{\int_{(i-1)/r}^{i/r} Gaussian_{(\mu=0.0,\sigma=0.25)}(x) \delta x}{\int_0^1 Gaussian_{(\mu=0.0,\sigma=0.25)}(x) \delta x}$$

- for each level, the center of the band is used as its representative.
- (c) for each state, s, the program
 - i. moves a w-length window on the corresponding time series (by shifting it h time units at a time), and
 - ii. for each window, starting at t, writes the pair $\langle idx, \vec{win} \rangle$, where
 - $-idx = \langle f, s, t \rangle$ and
 - $w\bar{i}n$ is quantized content of the w-length window starting at time t, into a given file, $epidemic_word_file$.
- Task 2: Implement a program which reads a connectivity graph, G, a file, $epidemic_word_file$. Given a weight, $0 \le \alpha \le 1$, the program does the following: for each pair

$$w_i = \langle idx_i, \vec{win_i} \rangle,$$

the program

1. writes the pair

$$w_{avq,i} = \langle idx_i, \vec{win}_{avq,i} \rangle,$$

where

$$\vec{win}_{avg,i} = \left(\alpha \times \vec{win}_i\right) + \left((1 - \alpha) \times AVG\left\{\vec{win}_j \mid \left(s_j \in 1HN(G, idx_i.s)\right) \land \left(idx_j = \langle idx_i.f, s_j, idx_i.t \rangle\right)\right\}\right)$$

into the file epidemic_word_file_avq.

2. writes the pair

$$w_{diff,i} = \langle idx_i, w\vec{i}n_{diff,i} \rangle,$$

where

$$\vec{win}_{diff,i} = \left(\vec{win}_i - \left(AVG\left\{\vec{win}_j \mid \left(s_j \in 1HN(G, idx_i.s)\right) \land \left(idx_j = \langle idx_i.f, s_j, idx_i.t \rangle\right)\right\}\right)\right) \div \vec{win}_i$$

and where \div denotes element-wise division operation, into the file *epidemic_word_file_diff*.

Above, 1HN(G,s) denotes the set of 1-hop neighbors of the state s according to the connectivity matrix G.

- Task 3: Given a word $\langle idx, \vec{win} \rangle$, let the length (norm-2), $|\vec{win}|$, of the vector, \vec{win} , denote the strength of the word. Implement a program which lets the user
 - select an epidemic simulation file, f, and view it in the form of a heat map, and highlight (based on the user's choice)
 - * the two windows, corresponding to f, with the highest and lowest strengths in the file $epidemic_word_file$,
 - * the two windows, corresponding to f, with the highest and lowest strengths in the file epidemic_word_file_avg,
 - * the two windows, corresponding to f, with the highest and lowest strengths in the file epidemic_word_file_diff.

For each window, $\langle idx, w in \rangle$, please visualize the state idx.s and the states in the 1-hop neighborhood of idx.s.

Deliverables:

- Your code (properly commented) and a README file.
- Your outputs for the provided sample inputs.
- A short report describing your work and the results.

Please place your code in a directory titled "Code", the outputs to a directory called "Outputs", and your report in a directory called "Report"; zip or tar all off them together and submit it through the digital dropbox.

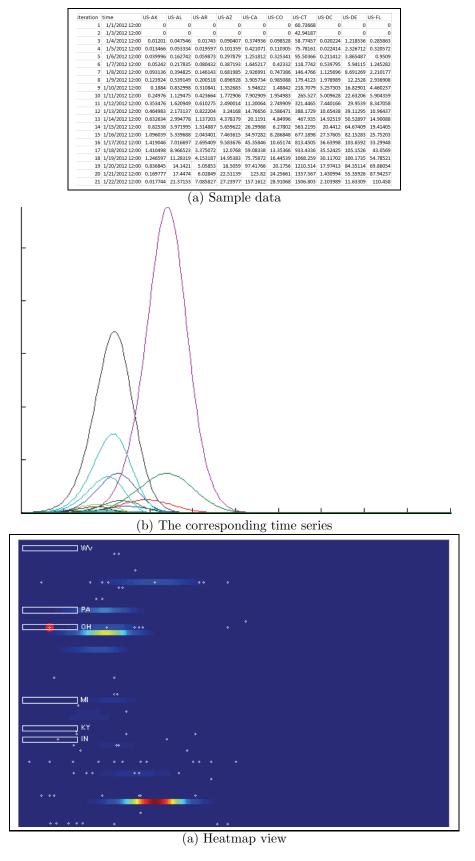


Figure 1: A sample visualization showing a heatmap and a highlighted window: the state is "OH" and its 1-hop neighbors are "WV", "PA", "MI", "KY", and "IN"