Pertinent reading for Problem #1:

R. Briandet, E.K. Kemsley, and R.H. Wilson, "Discrimination of *Arabica* and *Robusta* in Instant Coffee by Fourier Transform Infrared Spectroscopy and Chemometrics," *J. Agric. Food Chem.* 44 (1) 170 – 174 (1996)

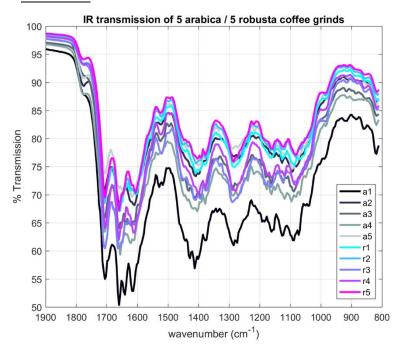


As you progress into this problem, here's a reminder of what your giant data matrix looks like!

In this problem, you are given 10 bags of coffee beans:

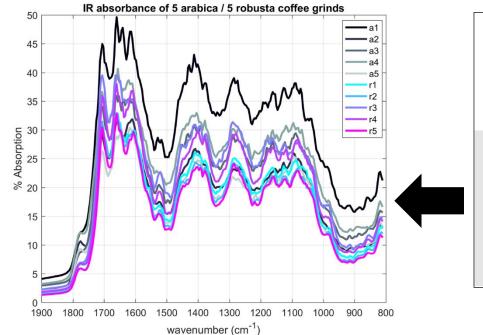
5 arabica beans (*C. arabica*) 5 robusta beans (*C. Canephora v. Robusta*)

and you want to know whether you can use infrared (IR) spectroscopy to differentiate between the 2 kinds of beans. To satisfy your curiosity, you used the FTIR machine in the basement of 44 Cummington and measured the IR spectrum of the 10 coffee grinds near the "fingerprint" (1/wavelength) region between 800 – 1900 cm⁻¹. Figure 1a is what your IR <u>transmission</u> results look like, where Figure 1b reports the same data plotted in IR <u>absorbance</u>



<u>Figure 1a</u>: The IR transmission spectra for:

5 Arabica grinds (a1 – a5) 5 Robusta grinds (r1 – r5)



<u>Figure 1b</u>: The same data as in Figure 1a,but plotted in term of IR <u>absorbance</u>

The file:

"Problem2_coffee_FTIR_data.txt"

will be the <u>absorbance</u> data set shown here !!

<u>Your Tasks for this section:</u> As usual, please turn in all <u>*.m matlab scripts</u>, plots, or requested echoed values (in diary files) shaded in yellow.

1. Using matlab's *textread* function, load in 10 columns of IR data stored in "Problem2_coffee_FTIR_data.txt." Since reading in 10 columns worth of stuff will make the command really long, you can use the 3 - period ellipsis notation "..." to join several lines of code into 1 single line of command. For instance, I would spread out the *textread* command into 4 lines of code:

[wavenumber, arabica1, arabica2, arabica3, arabica4, arabica5, ...

robusta1, robusta2, robusta3, robusta4, robusta5] ...

= textread('Problem1_coffee_FTIR_data.txt', ...

'%f%f%f%f%f%f%f%f%f%f%f%f, 'headerlines', 3)

- ** Note: There are 3 lines of headers in this file, and also:
 - a) The 1st data column = wavenumber (units = cm^{-1})
 - b) The next 5 columns = IR absorption data from 5 different sources of arabica beans
 - c) The last 5 columns = IR absorption data from 5 different sources of robusta beans
- 2. Build a giant data matrix X by joining the 10 columns of IR data side-by-side. The matrix X should be in the form:

$$X = \left[egin{array}{c|c} arabica1 & arabica2 \ data \end{array} \right] \left[egin{array}{c|c} arabica5 \ data \end{array} \right] \left[egin{array}{c|c} robusta1 \ data \end{array} \right] \left[egin{array}{c|c} robusta2 \ data \end{array} \right] \left[egin{array}{c|c} robusta5 \ data \end{array} \right]$$

- 3. Using matlab's *svd* function, calculate the orthogonal matrices U, V, and the singular value matrix Σ . Please echo the following:
 - a) The smaller eigenvector matrix V.
 - b) The 10 non-zero singular values σ_1 , σ_2 , \cdots , σ_{10} that's inside your Σ matrix. Notice that matlab has automatically ranked them in descending order!

- 4. Using matlab's *semilogy* function, plot all the singular values σ_p on the y-axis versus a dummy index on the x-axis that represents the pth singular value of your data matrix X. Also, add grid lines to your plot!
- 5. Recall the equation for SVD spectral decompositions for your data matrix *X*:

Using *pcolor*, plot the *values of the 4 following matrices (X and the 3 SVD spectra)* and save them as *.jpg files. Check out Figure 2 on the next page to see an example of what these 4 plots should look like!

SVD spectrum	Matrix	[cmin cmax]
		for the caxis command (% absorbance colors)
Original data	X	[0 50]
#1 (Dominant)	$\sigma_1 \left(u_1 {v_1}^T ight)$	[0 50]
#2 (Secondary)	$\sigma_2 (u_2 v_2^T)$	[0 10] (increases the data contrast)
#3 (Tertiary)	$\sigma_3 \left(u_3 {v_3}^T ight)$	[0 10]

For each of these plots:

- a) Using the *set* command, set the 'EdgeColor' to 'none' to remove the black borders for each pixels.
- b) Please color-label the plot using *colorbar*, and also, limit the color values using *caxis*([cmin cmax]) command. Then, add an "% absorbance" label to your colorbar (check out the example matlab script to see how to do this).
- c) Reverse the y-axis values by invoking the command:

set(gca, 'YDir', 'reverse')

- d) The title of the original data matrix X should say, "Original data X." The title for each SVD spectrum plot should say, "SVD submatrix p with $\sigma_p = xxxx \cdot xxxxx$ ", where:
 - i) p = 1, 2, or 3
 - ii) σ_p = Greek letter with a dyanmically changing 'p' value
 - iii) $xxxx.xxxx = The actual singular value \sigma_p in you S matrix$
 - ** <u>Hint</u>: You should use the <u>strcat</u> and <u>num2str</u> functions to do this!!

 Again, check out the example matlab script to see how to do this efficiently!
- e) Next, the x-axis should have labels "a1 a2 a3 a4 a5 r1 r2 r3 r4 r5" aligned with each pixel columns. This corresponds to the 10 different coffee bean types. Consult the example matlab script on how to do this.
- f) Moreover, the y-axis should be labeled as "wavenumber (cm⁻¹)." This corresponds to the wavenumber used in our IR scans.

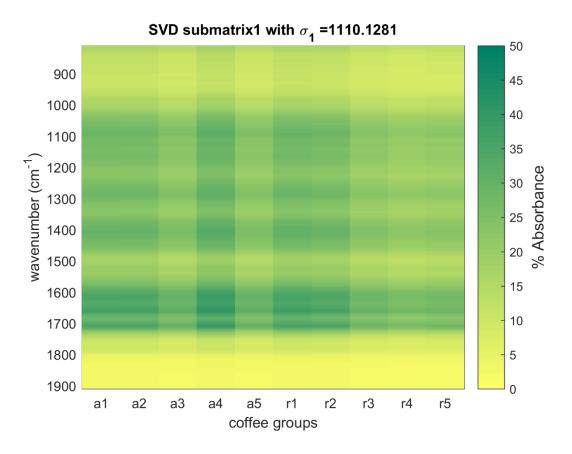


Figure 2: An example of what your SVD submatrix poolor plots should look like!

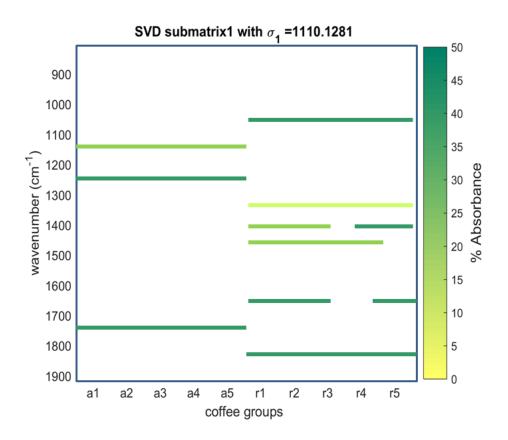
Task #2: Analyze the SVD component that can "separate" arabica vs. robusta beans

Yay! You've actually done the hardest part of this problem! =)

Now, stare at each of the 3 SVD submatrix plots for a minute. You'll notice that 1 of those 3 plots will exhibit the following characteristics:

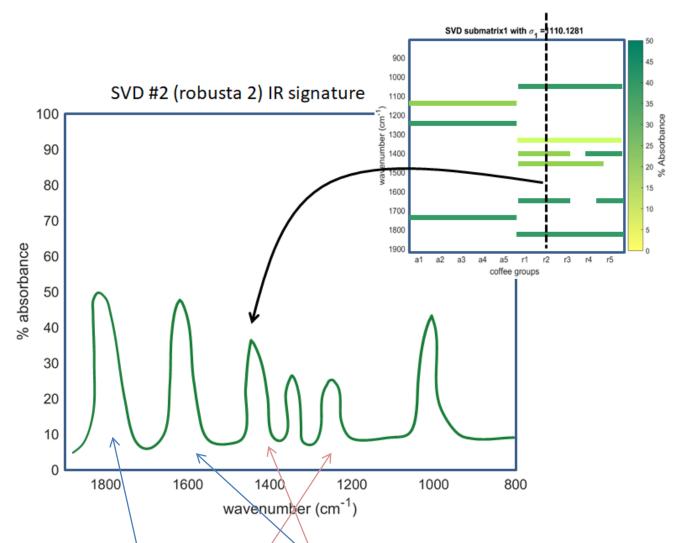
- a) The vertical SVD patterns of the arabica group (a1 a5) will look similar to each other
- b) The vertical SVD patterns of the robusta group (r1 r5) will also look similar to each other
- c) Group (a1-a5) looks completely different than Group (r1 r5).

An example of what you may see is depicted in Figure 3.

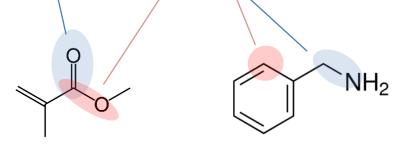


<u>Figure 3</u>: An example in which the vertical SVD patterns of the arabica beans (left, a1-a5) can be differentiated from the robusta beans (right, r1-r5).

Yes !!! This is the SVD component that we want! Now, we are ready to extract the robusta-specific IR signature from the SVD submatrix in Figure 3. As depicted in Figure 4, we will take a vertical slice of data from either coffee beans r1 or r2 that best represents the robusta group's IR signature. Then, we can plot the resulting IR absorbance versus the wavenumber.



<u>Figure 4</u>: We can extract coffee bean r2's IR spectrum from the SVD submatrix to get a qualitative sense of the chemical signatures that could differentiate arabica beans vs. robusta beans. After



If you were a coffee biochemist, your job is now is to stare at the IR signature in Figure 4 and try to identify:

- 1) Any possible molecules that could be in your robusta beans
- 2) Any "dominant" IR signatures from various functional groups within these molecules

<u>Your Tasks for this section:</u> As usual, please turn in all <u>*.m matlab scripts</u>, plots, or requested echoed values (in diary files) shaded in yellow.

- 1. Stare at your SVD submatrix plots σ_1 ($u_1v_1^T$), σ_2 ($u_2v_2^T$), and σ_3 ($u_3v_3^T$) for a little bit. Identify the one that exhibits patterns that could best differentiate between the 2 coffee bean groups.
- 2. From your chosen SVD submatrix, choose the best robusta bean type that best represents the entire family of robusta IR spectra. Cookie-cut out that data column from your SVD matrix.
- 3. Using matlab's *plot* function, plot the absorbance vs. wavenumber spectrum from your chosen data column in step 2 and save it as a *.jpg file. Again, the format of your plot should look very similar to Figure 4. On that plot, you should have:
 - a) Title = "SVD # p, robusta m IR signature", where:

$$\left\{ \begin{array}{ll} p = The & \sigma_p\left(u_pv_p^T\right) & \textit{SVD submatrix you chose} \\ \\ m = The & m^{th} & \textit{robusta bean you chose that best represents the robusta group} \end{array} \right. .$$

- b) X-axis label = "wavenumber", where it is shown in <u>reverse axis order</u> by using: set(gca, 'XDir', 'reverse')
- c) Y-axis label = "% Absorbance"
- d) Add grids to your plot. This will be important for visualizing your data in the next step.
- 4. Now, read the accompanied research paper by *R. Briandet, et. al.* carefully. In the text, the authors suggests that robusta beans may be differentiated from arabica beans because of an increase in IR signatures from 2 chemical ingredients. Please identify these 2 chemicals (type the answer in your diary file).

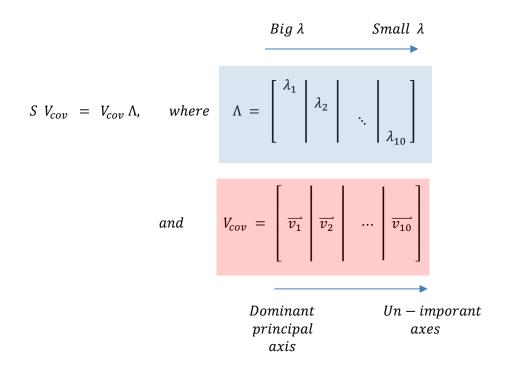
<u>Hint</u>: Your plotted IR signature should look really similar to one of the figures in that paper, and also, your plot should contain a linear combination of the peaks that are associated with these 2 chemicals !! =)

Task #3: Confirm your "best IR signature" findings using the old-school covariance S

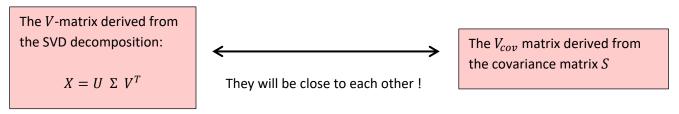
Recall that if we had a giant data matrix X, we can immediately calculate the covariance matrix S:

$$X = \begin{array}{ccc} Data & Subtract the means \\ of each column of data \\ matrix & T \end{array} \longrightarrow \begin{array}{ccc} Deviation & S = \frac{1}{n-1} & D^TD \end{array}$$

And once you know the eigenvectors and eigenvalues of S (and ranked them in order of descending values of λ), you can identify the "dominant," secondary, tertiary... etc. principal components of your data:



Now, if you recall from Problem 1's overview file:



(The main difference is that V_{cov} contains the zero-meaned data matrix D instead of the raw data matrix X)

<u>Your Tasks for this section:</u> As usual, please turn in all *.m matlab scripts, plots, or requested echoed values (in diary files) shaded in yellow.

- 1. From your original data matrix *X*, calculate (no need to echo these):
 - a) The deviation matrix D:

$$D = \begin{bmatrix} \overrightarrow{d_1} & \overrightarrow{d_2} & \cdots & \overrightarrow{d_8} & \overrightarrow{d_9} & \overrightarrow{d_{10}} \end{bmatrix}, \quad \text{where} \quad \overrightarrow{d_p} = \overrightarrow{x_p} - \text{mean}(\overrightarrow{x_p})$$

b) The covariance matrix *S*:

$$S = \frac{1}{n-1} D^T D$$

2. Then, using matlab's eig function, find the eigenvalue matrix Λ and the eigenvector matrix V_{cov} from the covariance matrix S and echo them in your diary. Remember, Λ and V_{cov} should obey the eigenvalue equation:

$$S V_{cov} = V_{cov} \Lambda$$

** Look at this matrix, and note that the first 3 eigenvalues λ_1 , λ_2 and , λ_3 are much larger than the other ones !

3. Normalize all entries in V_{cov} such that <u>the smallest entry in each column is 1.000</u>. Echo $V_{Normalized}$ in your diary. Here's a quick example: If V_{cov} was a 3×3 matrix:

$$V_{cov} = \begin{bmatrix} 2.828 & 2.828 & 1.500 \\ 1.414 & -14.14 & -1.500 \\ 0.707 & 0.707 & -0.500 \end{bmatrix} \xrightarrow{\textit{Make the smallest entry} = 1} V_{normalized} = \begin{bmatrix} 4 & 4 & 3 \\ 2 & -10 & -3 \\ 1 & 1 & -1 \end{bmatrix}$$

Part #4: Interpretation of your PCA results from $V_{normalized}$

As you stare at your $V_{Normalized}$ matrix, notice the following characteristics:

- Every entry within the <u>dominant-eigenvalue's corresponding eigenvector</u> (otherwise known as the **1**st **Principal Axis**) have the <u>same signs</u> right now
- Moreover, these entries are all small in magnitude
- Then, notice that for other eigenvectors, in particular the one that corresponds to the 2nd-most-dominant eigenvalue (ie. the 2nd Principal Axis), the entries within that vector are similar in magnitude..... but some have different signs.

Again, let's suppose V_{cov} was a 3×3 matrix (yours will be a 10×10 matrix !!), where the eigenvectors $\overrightarrow{v_p}$ are listed in descending order with respect to the eigenvalue of S. Once we've calculated $V_{normalized}$, you can actually visualize the principal axes characteristics $\underline{by\ eye\ !!!!!!!}$

$$V_{cov} = \begin{bmatrix} 2.828 & 2.828 & 1.500 \\ 1.414 & -14.14 & -1.500 \\ 0.707 & 0.707 & -0.500 \end{bmatrix} \xrightarrow{smallest\ entry=1} V_{normalized} = \begin{bmatrix} 4 & 4 & 3 \\ 2 & -10 & -3 \\ 1 & -1 & -1 \end{bmatrix}$$

The relative weight factors in the 1st PC will usually give you a qualitative measure of the "similarities" between different coffee beans.

Usually, the 1st PC is not so useful in differentiating between different "groups" of data columns =(The 2nd and 3rd PCs are usually the more interesting ones. When there are a lot of sign changes, they will usually tell you:

- a) Is one or more <u>GROUPS</u> of photos (or variables) differ from other GROUPS?
- b) Are there <u>super-oddballs</u>, <u>or super-"opposite-signed" oddballs</u> within your entire collections of photos? You will see instances of this when you have a huge +/- magnitude change between 2 or 3 entries, and the same time, all other entries remain small in magnitude.

The reason why you want to do this will only be apparent when you check out Johnson & Wichern, Example 8.5, pp. 451-452. Here's that example copied from p. 452 of the book. My commentaries are on the side...

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and

$$\mathbf{R} = \begin{bmatrix} 1.000 & .632 & .511 & .115 & .155 \\ .632 & 1.000 & .574 & .322 & .213 \\ .511 & .574 & 1.000 & .183 & .146 \\ .115 & .322 & .183 & 1.000 & .683 \\ .155 & .213 & .146 & .683 & 1.000 \end{bmatrix}$$

$$z_1 = \frac{x_1 - \bar{x}_1}{\sqrt{s_{11}}}, z_2 = \frac{x_2 - \bar{x}_2}{\sqrt{s_{22}}}, \dots, z_5 = \frac{x_5 - \bar{x}_5}{\sqrt{s_{55}}}$$

Our eigenvector matrix (derived from the covariance matrix S, or equivalently, from matrix R)

$$V_{cov} = \left[\begin{array}{c|c} \overrightarrow{e_1} & \overrightarrow{e_2} & \overrightarrow{e_3} & \overrightarrow{e_4} & \overrightarrow{e_5} \end{array} \right]$$

We note that **R** is the covariance matrix of the standardized observations
$$\Lambda = \begin{bmatrix} \lambda_1 \\ \lambda_2 \\ \lambda_3 \end{bmatrix}$$
 $\lambda_4 \begin{bmatrix} \lambda_4 \\ \lambda_5 \end{bmatrix}$

The eigenvalues and corresponding normalized eigenvectors of R, determined by a computer, are

The sum of these Eigenvalues = 5

The 1st PC explains 48.7% of the data Variations

Adding a 2nd PC will gain another 28.1% increase in explaining our data variations.

 $\hat{\lambda}_1 = 2.437$, $\hat{\mathbf{e}}_1' = [.469, .532, .465, .387, .361]$ $\hat{\lambda}_2 - 1.407$, $\hat{\epsilon}_2^t = [-.368, -.236, -.315, .585, .606]$ $\hat{\lambda}_3 = .501,$ $\hat{\mathbf{e}}_3' = [-.604, -.136, .772, .093, -.109]$ $\hat{\lambda}_4 = .400,$ $\hat{\mathbf{e}}_4' = [.363, -.629, .289, -.381, .493]$

Using the standardized variables, we obtain the first two sample principal components:

 $\hat{\lambda}_5 = .255, \quad \hat{\mathbf{e}}_5' = [.384, -.496, .071, .595, -.498]$

$$\hat{y}_1 = \hat{\mathbf{e}}_1' \mathbf{z} = .469z_1 + .532z_2 + .465z_3 + .387z_4 + .361z_5$$

$$\hat{y}_2 = \hat{\mathbf{e}}_2' \mathbf{z} = -.368z_1 - .236z_2 - .315z_3 + .585z_4 + .606z_5$$

These components, which account for

$$\left(\frac{\hat{\lambda}_1 + \hat{\lambda}_2}{p}\right) 100\% = \left(\frac{2.437 + 1.407}{5}\right) 100\% = 77\%$$

of the total (standardized) sample variance, have interesting interpretations. The first component is a roughly equally weighted sum, or "index," of the five stocks. This component might be called a general stock-market component, or, simply, a market component.

The second component represents a contrast between the banking stocks (JP Morgan, Citibank, Wells Fargo) and the oil stocks (Royal Dutch Shell, Exxon-Mobil). It might be called an industry component. Thus, we see that most of the variation in these stock returns is due to market activity and uncorrelated industry activity. This interpretation of stock price behavior also has been suggested by King [12].

The remaining components are not easy to interpret and, collectively, represent variation that is probably specific to each stock. In any event, they do not explain much of the total sample variance.

PC #1 (not useful...)

PC #2 (useful !!)

The 3 sign changes in the original coordinates $(z_i \sim x_i)$ signify that the experimental variables z_1 , z_2 , and z_3 are maybe fundamentally different than z₄ and z₅.

 $Z_1 \sim X_1 = JP Morgan$

 $Z_2 \sim X_2 = Citybank$

 $Z_3 \sim X_3 = \text{Wells Fargo}$

 $Z_4 \sim X_4 = Shell$

 $Z_5 \sim X_5 = Exxon Mobil$

<u>Your Tasks for this section:</u> As usual, please turn in all <u>*.m matlab scripts</u>, plots, or requested echoed values (in diary files) shaded in yellow.

I would like to <u>examine the first 3 PCs</u> (the largest 3 eigenvalues and their corresponding eigenvectors). To be specific, I want to know the following:

1. What % of the total variance can be explained by taking the 1st three PC's?

- 2. In the 2nd PC eigenvector, please <u>use your intuition</u> and give a plausible explanation to:
 - Why 5 out of the 10 entries in appear to have a "-" sign and
 - Why do the other 5 entries appear to have a "+" sign?



By answering this question, you have just proved (statistically) that:

